# TeNPy 

Release 0.8.0

## TeNPy Developers

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## Part I

## User guide

TeNPy (short for 'Tensor Network Python') is a Python library for the simulation of strongly correlated quantum systems with tensor networks.
The philosophy of this library is to get a new balance of a good readability and usability for new-comers, and at the same time powerful algorithms and fast development of new algorithms for experts. For good readability, we include an extensive documentation next to the code, both in Python doc strings and separately as user guides, as well as simple example codes and even toy codes, which just demonstrate various algorithms (like TEBD and DMRG) in ~100 lines per file.

## HOW DO I GET SET UP?

If you have the conda package manager, you can install the latest released version of TeNPy with:

```
conda install --channel=conda-forge physics-tenpy
```

Further details and alternative methods can be found the file doc/INSTALL.rst. The latest version of the source code can be obtained from https://github.com/tenpy/tenpy.

## HOW TO READ THE DOCUMENTATION

The documentation is available online at https://tenpy.readthedocs.io/. The documentation is roughly split in two parts: on one hand the full "reference" containing the documentation of all functions, classes, methods, etc., and on the other hand the "user guide" containing some introductions with additional explanations and examples.

The documentation is based on Python's docstrings, and some additional *. rst files located in the folder doc/ of the repository. All documentation is formated as reStructuredText, which means it is quite readable in the source plain text, but can also be converted to other formats. If you like it simple, you can just use intective python help(), Python IDEs of your choice or jupyter notebooks, or just read the source. Moreover, the documentation gets converted into HTML using Sphinx, and is made available online at https://tenpy.readthedocs.io/. The big advantages of the (online) HTML documentation are a lot of cross-links between different functions, and even a search function. If you prefer yet another format, you can try to build the documentation yourself, as described in doc/contr/build_doc.rst.

## HELP - I LOOKED AT THE DOCUMENTATION, BUT I DON'T UNDERSTAND HOW ...?

We have set up a community forum at https://tenpy.johannes-hauschild.de/, where you can post questions and hopefully find answers. Once you got some experience with TeNPy, you might also be able to contribute to the community and answer some questions yourself ;-) We also use this forum for official annoucements, for example when we release a new version.

## I FOUND A BUG

You might want to check the github issues, if someone else already reported the same problem. To report a new bug, just open a new issue on github. If you already know how to fix it, you can just create a pull request :) If you are not sure whether your problem is a bug or a feature, you can also ask for help in the TeNPy forum.

## CITING TENPY

When you use TeNPy for a work published in an academic journal, you can cite this paper to acknowledge the work put into the development of TeNPy. (The license of TeNPy does not force you, however.) For example, you could add the sentence "Calculations were performed using the TeNPy Library (version X. X. X) \cite $\{$ tenpy $\}$. " in the acknowledgements or in the main text.

The corresponding BibTex Entry would be the following (the \}requires\usepackage\{hyperref\}intheLaTeXpreamble.):undefinedundefinedundefinedundefinedundefinedundefinedundefinedundefinedundefinedundefinedundefinedundefinedundefinedundefinedundefinedundefinedundefined

```
@Article{tenpy,
    title={{Efficient numerical simulations with Tensor Networks: Tensor Network
->Python (TeNPy)}},
    author={Johannes Hauschild and Frank Pollmann},
    journal={SciPost Phys. Lect. Notes},
    pages={5},
    year={2018},
    publisher={SciPost},
    doi={10.21468/SciPostPhysLectNotes.5},
    url={https://scipost.org/10.21468/SciPostPhysLectNotes.5},
    archiveprefix={arXiv},
    eprint={1805.00055},
    note={Code available from \url{https://github.com/tenpy/tenpy}},
}
```

To keep us motivated, you can also include your work into the list of papers using TeNPy.

## ACKNOWLEDGMENT

This work was funded by the U.S. Department of Energy, Office of Science, Office of Basic Energy Sciences, Materials Sciences and Engineering Division under Contract No. DE-AC02-05- CH11231 through the Scientific Discovery through Advanced Computing (SciDAC) program (KC23DAC Topological and Correlated Matter via Tensor Networks and Quantum Monte Carlo).

## LICENSE

The code is licensed under GPL-v3.0 given in the file LICENSE of the repository, in the online documentation readable at https://tenpy.readthedocs.io/en/latest/install/license.html.

## INSTALLATION INSTRUCTIONS

With the [conda] package manager you can install python with:

```
conda install --channel=conda-forge physics-tenpy
```

More details and tricks in Installation with conda from conda-forge.
If you don't have conda, but you have [pip], you can:

```
pip install physics-tenpy
```

More details for this method can be found in Installation from PyPi with pip.
We also have a bunch of optional Extra requirements, which you don't have to install to use TeNPy, but you might want to.

The method with the minimal requirements is to just download the source and adjust the PYTHONPATH, as described in Installation from source. This is also the recommended way if you plan to modify parts of the source.

### 8.1 Installation with conda from conda-forge

We provide a package for the [conda] package manager in the conda-forge channel, so you can install TeNPy as:

```
conda install --channel=conda-forge physics-tenpy
```

Following the recommondation of conda-forge, you can also make conda-forge the default channel as follows:

```
conda config --add channels conda-forge
conda config --set channel_priority strict
```

If you have done this, you don't need to specify the --channel=conda-forge explicitly.

Note: The numpy package provided by the conda-forge channel by default uses openblas on linux. As outlined in the conda forge docs, you can switch to MKL using:

```
conda install "libblas=*=*mkl"
```

Warning: If you use the conda-forge channe and don't pin BLAS to the MKL version as outlined in the above version, but nevertheless have mkl-devel installed during compilation of TeNPy, this can have crazy effects on the number of threads used: numpy will call openblas and open up \$OMP_NUM_THREADS - 1 new threads, while

MKL called from tenpy will open another \$MKL_NUM_THREADS - 1 threads, making it very hard to control the number of threads used!

Moreover, it is actually recommended to create a separate environment. To create a conda environment with the name tenpy, where the TeNPy package (called physics-tenpy) is installed:

```
conda create --name tenpy --channel=conda-forge physics-tenpy
```

In that case, you need to activate the environment each time you want to use the package with:

```
conda activate tenpy
```

The big advantage of this approach is that it allows multiple version of software to be installed in parallel, e.g., if one of your projects requires python $>=3.8$ and another one requires an old library which doesn't support that. Further info can be found in the conda documentation.

### 8.2 Installation from PyPi with pip

### 8.2.1 Preparation: install requirements

If you have the [conda] package manager from anaconda, you can just download the environment.yml file (using the conda-forge channel, or the environment_other.yml for all other channels) out of the repository and create a new environment (called tenpy, if you don't speficy another name) for TeNPy with all the required packages:

```
conda env create -f environment.yml
```

conda activate tenpy

Further information on conda environments can be found in the conda documentation. Note that installing conda also installs a version of [pip].

Alternatively, if you only have [pip] (and not [conda]), install the required packages with the following command (after downloading the requirements.txt file from the repository):

```
pip install -r requirements.txt
```

Note: Make sure that the pip you call corresponds to the python version you want to use. (One way to ensure this is to use python -m pip instead of a simple pip.) Also, you might need to use the argument --user to install the packages to your home directory, if you don't have sudo rights. (Using --user with conda's pip is discouraged, though.)

Warning: It might just be a temporary problem, but I found that the pip version of numpy is incompatible with the python distribution of anaconda. If you have installed the intelpython or anaconda distribution, use the conda packagemanager instead of pip for updating the packages whenever possible!

### 8.2.2 Installing the latest stable TeNPy package

Now we are ready to install TeNPy. It should be as easy as (note the different package name - 'tenpy' was taken!)
pip install physics-tenpy

Note: If the installation fails, don't give up yet. In the minimal version, tenpy requires only pure Python with somewhat up-to-date NumPy and SciPy. See Installation from source.

### 8.2.3 Installation of the latest version from Github

To get the latest development version from the github main branch, you can use:

```
pip install git+git://github.com/tenpy/tenpy.git
```

This should already have the lastest features described in [latest]. Disclaimer: this might sometimes be broken, although we do our best to keep to keep it stable as well.

### 8.2.4 Installation from the downloaded source folder

Finally, if you downloaded the source and want to modify parts of the source, You can also install TeNPy with in development version with --editable:

```
cd $HOME/tenpy # after downloading the source, got to the repository
pip install --editable .
```


### 8.2.5 Uninstalling a pip-installed version

In all of the above cases, you can uninstall tenpy with:

```
pip uninstall physics-tenpy
```


### 8.3 Updating to a new version

Before you update, take a look at the Release Notes, which lists the changes, fixes, and new stuff. Most importantly, it has a section on backwards incompatible changes (i.e., changes which may break your existing code) along with information how to fix it. Of course, we try to avoid introducing such incompatible changes, but sometimes, there's no way around them. If you skip some intermediate version(s) for the update, read also the release notes of those!

How to update depends a little bit on the way you installed TeNPy. Of course, you have always the option to just remove the TeNPy files (possibly with a pip uninstall physics-tenpy or conda uninstall physics-tenpy), and to start over with downloading and installing the newest version.

### 8.3.1 When installed with conda

When you installed TeNPy with [conda], you just need to activate the corresponding environment (e.g. conda activate tenpy) and do a:

```
conda update physics-tenpy
```


### 8.3.2 When installed with pip

When you installed TeNPy with [pip], you just need to do a:

```
pip install --upgrade physics-tenpy
```


### 8.3.3 When installed from source

If you used git clone . . . to download the repository, you can update to the newest version using [git]. First, briefly check that you didn't change anything you need to keep with git status. Then, do a git pull to download (and possibly merge) the newest commit from the repository.

Note: If some Cython file (ending in . pyx) got renamed/removed (e.g., when updating from v0.3.0 to v0.4.0), you first need to remove the corresponding binary files. You can do so with the command bash cleanup. sh.

Furthermore, whenever one of the cython files (ending in . pyx) changed, you need to re-compile it. To do that, simply call the command bash . / compile again. If you are unsure whether a cython file changed, compiling again doesn't hurt.

To summarize, you need to execute the following bash commands in the repository:

```
# O) make a backup of the whole folder
git status # check the output whether you modified some files
git pull
bash ./cleanup.sh # (confirm with 'y')
bash ./compile.sh
```


### 8.4 Installation from source

### 8.4.1 Minimal Requirements

This code works with a minimal requirement of pure Python $>=3.6$ and somewhat recent versions of NumPy and SciPy.

### 8.4.2 Getting the source

The following instructions are for (some kind of) Linux, and tested on Ubuntu. However, the code itself should work on other operating systems as well (in particular MacOS and Windows).

The offical repository is at https://github.com/tenpy/tenpy.git. To get the latest version of the code, you can clone it with [git] using the following commands:

```
git clone https://github.com/tenpy/tenpy.git $HOME/TeNPy
cd $HOME/TeNPy
```

Note: Adjust \$HOME/TeNPy to the path wherever you want to save the library.

Optionally, if you don't want to contribute, you can checkout the latest stable release:

```
git tag # this prints the available version tags
git checkout v0.3.0 # or whatever is the lastest stable version
```

Note: In case you don't have [git] installed, you can download the repository as a ZIP archive. You can find it under releases, or the latest development version.

### 8.4.3 Minimal installation: Including tenpy into PYTHONPATH

The python source is in the directory tenpy/ of the repository. This folder tenpy/ should be placed in (one of the folders of) the environment variable PYTHONPATH. On Linux, you can simply do this with the following line in the terminal:

```
export PYTHONPATH=$HOME/TeNPy
```

(If you have already a path in this variable, separate the paths with a colon :.) However, if you enter this in the terminal, it will only be temporary for the terminal session where you entered it. To make it permanently, you can add the above line to the file $\$ \mathrm{HOME} / \mathrm{b}$ bashrc. You might need to restart the terminal session or need to relogin to force a reload of the $\sim /$. bashrc.

Whenever the path is set, you should be able to use the library from within python:

```
>>> import tenpy
/home/johannes/postdoc/2021-01-TenPy-with-MKL/TeNPy/tenpy/tools/optimization.py:308: 
\hookrightarrowUserWarning: Couldn't load compiled cython code. Code will run a bit slower.
warnings.warn("Couldn't load compiled cython code. Code will run a bit slower.")
>>> tenpy.show_config()
tenpy 0.7.2.dev130+76c5b7f (not compiled),
git revision 76c5b7fe46df3e2241d85c47cbced3400caad05a using
python 3.9.1 | packaged by conda-forge | (default, Jan 10 2021, 02:55:42)
[GCC 9.3.0]
numpy 1.19.5, scipy 1.6.0
```

tenpy. show_config() prints the current version of the used TeNPy library as well as the versions of the used python, numpy and scipy libraries, which might be different on your computer. It is a good idea to save this data (given as string in tenpy.version.version_summary along with your data to allow to reproduce your results exactly.

If you got a similar output as above: congratulations! You can now run the codes :)

### 8.4.4 Compilation of np_conserved

At the heart of the TeNPy library is the module tenpy.linalg.np_conseved, which provides an Array class to exploit the conservation of abelian charges. The data model of python is not ideal for the required book-keeping, thus we have implemented the same np_conserved module in Cython. This allows to compile (and thereby optimize) the corresponding python module, thereby speeding up the execution of the code. While this might give a significant speed-up for code with small matrix dimensions, don't expect the same speed-up in cases where most of the CPU-time is already spent in matrix multiplications (i.e. if the bond dimension of your MPS is huge).

To compile the code, you first need to install Cython

```
conda install cython # when using anaconda, or
pip install --upgrade Cython # when using pip
```

Moreover, you need a C++ compiler. For example, on Ubuntu you can install sudo apt-get install build_essential, or on Windows you can download MS Visual Studio 2015. If you use anaconda, you can also use conda install -c conda-forge cxx-compiler.

After that, go to the root directory of TeNPy (\$HOME/TeNPy) and simply run

```
bash ./compile.sh
```

Note: There is no need to compile if you installed TeNPy directly with conda or pip. (You can verify this with tenpy.show_config() as illustrated below.)

Note that it is not required to separately download (and install) Intel MKL: the compilation just obtains the includes from numpy. In other words, if your current numpy version uses MKL (as the one provided by anaconda), the compiled TeNPy code will also use it.

After a successful compilation, the warning that TeNPy was not compiled should go away:

```
>>> import tenpy
>>> tenpy.show_config()
tenpy 0.7.2.dev130+76c5b7f (compiled without HAVE_MKL),
git revision 76c5b7fe46df3e2241d85c47cbced3400caad05a using
python 3.9.1 | packaged by conda-forge | (default, Jan 10 2021, 02:55:42)
[GCC 9.3.0]
numpy 1.19.5, scipy 1.6.0
```

Note: For further optimization options, e.g. how to link against MKL, look at Extra requirements and tenpy. tools.optimization.

### 8.4.5 Quick-setup of a development environment with conda

You can use the following bash commands to setup a new conda environment called tenpy_dev (call it whatever you want!) and install TeNPy in there in a way which allows editing TeNPy's python code and still have it available everywhere in the conda environment:

```
git clone https://github.com/tenpy/tenpy TeNPy
cd TeNPy
conda env create -f environment.yml -n tenpy_dev
conda activate tenpy_dev
pip install -e .
```


### 8.5 Extra requirements

We have some extra requirements that you don't need to install to use TeNPy, but that you might find usefull to work with. TeNPy does not import the following libraries (at least not globally), but some functions might expect arguments behaving like objects from these libraries.

Note: If you created a [conda] environment with conda env create -f environment.yml, all the extra requirements below should already be installed:) (However, apip install $-r$ requirements.tyt does not install all of them.)

### 8.5.1 Matplotlib

The first extra requirement is the [matplotlib] plotting library. Some functions expect a matplotlib.axes. Axes instance as argument to plot some data for visualization.

### 8.5.2 Intel's Math Kernel Library (MKL)

If you want to run larger simulations, we recommend the use of Intel's MKL. It ships with a Lapack library, and uses optimization for Intel CPUs. Moreover, it uses parallelization of the LAPACK/BLAS routines, which makes execution much faster. As of now, the library itself supports no other way of parallelization.
If you don't have a python version which is built against MKL, we recommend using [conda] or directly intelpython. Conda has the advantage that it allows to use different environments for different projects. Both are available for Linux, Mac and Windows; note that you don't even need administrator rights to install it on linux. Simply follow the (straight-forward) instructions of the web page for the installation. After a successfull installation, if you run python interactively, the first output line should state the python version and contain Anaconda or Intel Corporation, respectively.

If you have a working conda package manager, you can install the numpy build against MKL with:

```
conda install mkl mkl-devel numpy scipy
```

The mkl-devel package is required for linking against MKL, i.e. for compiling the Cython code. As outlined in /doc/install/conda, on Linux/Mac you also need to pin blas to use MKL with the following line, if you use the `conda-forge` channel:

```
conda install "libblas=*=*mkl"
```

Note: MKL uses different threads to parallelize various BLAS and LAPACK routines. If you run the code on a cluster, make sure that you specify the number of used cores/threads correctly. By default, MKL uses all the available CPUs, which might be in stark contrast than what you required from the cluster. The easiest way to set the used threads is using the environment variable MKL_NUM_THREADS (or OMP_NUM_THREADS). For a dynamic change of the used threads, you might want to look at process.

## Compile linking agains MKL

When you compile the Cython files of TeNPy, you have the option to explicilty link against MKL, such that e.g. tenpy. Iinalg.np_conserved.tensordot () is guaranteed to call the corresponding dgemm or zgemm function in the BLAS from MKL. To link against MKL, the MKL library including the headers must be available during the compilation of TeNPy's Cython files. If you have the MKL library installed, you can export the environemnt variable MKLROOT to point to the root folder. Alternatively, TeNPy will recognise if you are in an active conda environment and have the $m k l$ and $m k l-d e v e l ~ c o n d a ~ p a c k a g e s ~ i n s t a l l e d ~ d u r i n g ~ c o m p i l a t i o n . ~ I n ~ t h i s ~ c a s e, ~ i t ~ w i l l ~ l i n k ~ a g a i n s t ~ t h e ~ M K L ~$ provided as conda package.

```
tenpy.show_config() indicates whether you linked successfully against MKL:
```

```
>>> import tenpy
>>> tenpy.show_config()
tenpy 0.7.2.dev130+76c5b7f (compiled with HAVE_MKL),
git revision 76c5b7fe46df3e2241d85c47cbced3400caad05a using
python 3.9.1 | packaged by conda-forge | (default, Jan 10 2021, 02:55:42)
[GCC 9.3.0]
numpy 1.19.5, scipy 1.6.0
```


### 8.5.3 HDF5 file format support

We support exporting data to files in the [HDF5] format through the python interface of the h5py [https://docs.h5py.org/en/stable/](https://docs.h5py.org/en/stable/) package, see Saving to disk: input/output for more information. However, that requires the installation of the HDF5 library and h5py.

### 8.5.4 YAML parameter files

The tenpy.tools.params.Config class supports reading and writing YAML files, which requires the package pyyaml; pip install pyyaml.

### 8.5.5 Tests

To run the tests, you need to install pytest, which you can for example do with pip install pytest. For information how to run the tests, see Checking the installation.

### 8.6 Checking the installation

The first check of whether tenpy is installed successfully, is to try to import it from within python:

```
>>> import tenpy
```

Note: If this raises a warning Couldn't load compiled cython code. Code will run a bit slower., something went wrong with the compilation of the Cython parts (or you didn't compile at all). While the code might run slower, the results should still be the same.

The function tenpy.show_config() prints information about the used versions of tenpy, numpy and scipy, as well on the fact whether the Cython parts were compiled and could be imported.

As a further check of the installation you can try to run (one of) the python files in the examples/ subfolder; hopefully all of them should run without error.

You can also run the automated testsuite with pytest to make sure everything works fine. If you have pytest installed, you can go to the tests folder of the repository, and run the tests with:

```
cd tests
pytest
```

In case of errors or failures it gives a detailed traceback and possibly some output of the test. At least the stable releases should run these tests without any failures.

If you can run the examples but not the tests, check whether pytest actually uses the correct python version.
The test suite is also run automatically by github actions and with travis-ci, results can be inspected here.

### 8.7 TeNPy developer team

```
The following people are part of the TeNPy developer team and contributed a
\hookrightarrowsignificant amount.
The full list of contributors can be obtained from the git repository with ``git*
\hookrightarrowshortlog -sn``.
Johannes Hauschild tenpy@johannes-hauschild.de
Frank Pollmann
Michael P. Zaletel
Maximilian Schulz
Leon Schoonderwoerd
Kévin Hémery
Samuel Scalet
Markus Drescher
Wilhelm Kadow
Gunnar Moeller
Jakob Unfried
Further, the code is based on an earlier version of the library, mainly developed by
Frank Pollmann, Michael P. Zaletel and Roger S. K. Mong.
```


### 8.8 License

The source code documented here is published under a GPL v3 license, which we include below.

```
        GNU GENERAL PUBLIC LICENSE
    Version 3, 29 June 2007
Copyright (C) 2007 Free Software Foundation, Inc. <https://fsf.org/>
Everyone is permitted to copy and distribute verbatim copies
of this license document, but changing it is not allowed.
    Preamble
    The GNU General Public License is a free, copyleft license for
software and other kinds of works.
```

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## CHAPTER

NINE

## RELEASE NOTES

The project adheres semantic versioning.
All notable changes to the project should be documented in the changelog. The most important things should be summarized in the release notes.

The changes in [latest] are implemented in the latest development version on github, but not yet released.
Changes compared to previous TeNPy highlights the most important changes compared to the other, previously developed (closed source) TeNPy version.

## 9.1 [latest]

### 9.1.1 Release Notes

TODO: Summarize the most important changes

### 9.1.2 Changelog

## Backwards incompatible changes

- nothing yet


## Added

- nothing yet


## Changed

- nothing yet


## Fixed

- nothing yet


## 9.2 [latest] - 2021-02-19

### 9.2.1 Release Notes

First of all: We have optimized the cython parts such that they can now link directly against MKL and have been optimized for the case of small blocks inside charge-conserved tensors. During compilation, TeNPy now checks whether MKL is available, and then directly links against it. This changed the depencies: in particular, when you created a conda environment for TeNPy, it is highly recommended to start off with a new one based on the environment.yml file. If you want to continue using the existing conda environment, you need to conda install mkl-devel before compilation. Additionally, when you use the conda-forge channel of conda, you should pin blas to use MKL by conda install libblas $=\star=\star \mathrm{mkl}$.

Another great reason to update are simulation classes and a console script tenpy-run to allow running and even resuming a simulation when it aborted! See /intro/simulation for details.

Further, there is a big change in verbosity: we switched to using Python's default logging mechanism. This implies that by default you don't get any output besides error messages and warning any more, at least not in pre-simulation setups. See Logging and terminal output on how to get the output back, and what to change in your code.

Finally, note that the default (stable) git branch was renamed from master to main.

### 9.2.2 Changelog

## Backwards incompatible changes

- Drop official support for Python 3.5.
- tenpy.linalg.np_conserved.from_ndarray () : raise ValueError instead of just a warning in case of the wrong non-zero blocks. This behaviour can be switched back with the new argument raise_wrong_sector.
- Argument $v 0$ of tenpy.networks.mps.MPS.TransferMatrix.eigenvectors() is renamed to $v 0 \_n p c ; v 0$ now serves for non-np_conserved guess.
- Default parameters for lattice initialization in the following classes changed. In particular, the $b c \_M P S$ parameter now defaults to 'finite'.
- tenpy.models.hofstadter. HofstadterFermions
- tenpy.models.hofstadter. HofstadterBosons
- tenpy.models.toric_code.ToricCode
- Renamed tenpy.algorithms.tebd.Engine to tenpy.algorithms.tebd.TEBDEngine and tenpy.algorithms.tdvp.Engine to tenpy.algorithms.tdvp.TDVPEngine to have unique algorithm class-names.
- When running, no longer print stuff by default. Instead, we use Python's logging mechanism. To enable printing again, you need to configure the logging to print on "INFO" level (which is the default when running from command line)
As part of this big change in the way verbosity is handled, there were many minor changes: - rename Config.print_if_verbose to $\log ()$ - deprecate the verbose class argument of the Config - deprecate the verbose class attribute of all classes (if they had it). - change argument names of params ().


## Added

- Simulation class Simulation and subclasses as a new extra layer for handling the general setup.
- Command line script tenpy-run and run_simulation () for setting up a simulation.
- entanglement_entropy_segment2()
- apply_product_op()
- tenpy.linalg.sparse.FlatLinearOperator.eigenvectors() and eigenvectors() to unify code from tenpy.networks.mps.TransferMatrix.eigenvectors() and tenpy. linalg.lanczos.lanczos_arpack().
- tenpy.tools.misc.group_by_degeneracy()
- tenpy.tools.fit.entropy_profile_from_CFT() and tenpy.tools.fit. central_charge_from_S_profile()
- tenpy.networks.site.Site.multiply_operators() as a variant of multiply_op_names() accepting both string and npc arrays.
- tenpy.tools.events.EventHandler() to simplify call-backs e.g. for measurement codes during an algorithms.
- tenpy.tools.misc.find_subclass () to recursively find subclasses of a given base class by the name. This function is now used e.g. to find lattice classes given the name, hence supporting user-defined lattices defined outside of TeNPy.
- tenpy.tools.misc.get_recursive() and set_recursive() for nested data strucutres, e.g., parameters.
- tenpy.tools.misc.flatten () to turn a nested data structure into a flat one.
- tenpy. networks.mps. InitialStateBuilder to simplify building various initial states.
- Common base class tenpy. algorithms.Algorithm for all algorithms.
- Common base class tenpy. algorithms.TimeEvolutionAlgorithm for time evolution algorithms.
- tenpy.models.lattice.Lattice. Lu as a class attribute.
- tenpy.models.lattice.Lattice.find_coupling_pairs() to automatically find coupling pairs of 'nearest_neighbors' etc..
- tenpy.models. Iattice. Helicallattice allowing to have a much smaller MPS unit cell by shifting the boundary conditions around the cylinder.
- tenpy.networks.purification_mps.PurificationMPS.from_infiniteT_canonical() for a canonical ensemble.


## Changed

- For finite DMRG, DMRGEngine.N_sweeps_check now defaults to 1 instead of 10 (which is still the default for infinite MPS).
- Merge tenpy.linalg.sparse.FlatLinearoperator.npc_to_flat_all_sectors() into npc_to_flat(), merge tenpy.linalg.sparse.FlatLinearOperator. flat_to_npc_all_sectors() into flat_to_npc().
- Change the chinfo. names of the specific Site classes to be more consistent and clear.
- Add the more powerful tenpy.networks.site.set_common_charges() to replace tenpy. networks.site.multi_sites_combine_charges().
- Allow swap_op='autoInv' for tenpy.networks.mps.MPS.swap_sites() and explain the idea of the swap_op.
- The tenpy.models.model.CouplingMPOModel.init_lattice() now respects new class attributes default_lattice and force_default_lattice.
- Support additional priority argument for get_order_grouped (), issue \#122.
- Warn if one of the $a d d_{-} *$ methods of the CouplingMPOModel gets called after initialization.


## Fixed

- Sign error for the couplings of the tenpy.models.toric_code. ToricCode.
- The form of the eigenvectors returned by tenpy.networks.mps.TransferMatrix. eigenvectors () was dependent on the charge_sector given in the initialization; we try to avoid this now (if possible).
- The charge conserved by SpinHalfFermionSite (cons_Sz='parity') was weird.
- Allow to pass npc Arrays as Arguments to expectation_value_multi_sites() and other correlation functions (issue \#116).
- tenpy.tools.hdf5_io did not work with h5py version $>=(3,0)$ due to a change in string encoding (issue \#117).
- The overall phase for the returned $W$ from compute_K () was undefined.
- tenpy.networks.mpo.MPO.expectation_value () didn't work with max_range=0
- The default trunc_par for tenpy.networks.mps.MPS.swap_sites(), permute_sites() and compute_K () was leading to too small chi for intial MPS with small chi.
- issue \#120 Lattice with different sites in the unit cell.
- Index offset in tenpy.networks.mps.MPS.expectation_value_term() for the sites to be used.
- issue \#121 tenpy. networks.mps.MPS.correlation_length() worked with charge_sector=0, but included additional divergent value with charge_sector=[0].
- Some MPS methods (correlation function, expectation value, ...) raised an error for negative site indices even for infinite MPS.
- Warn if we add terms to a couplingMPOMOdel after initialization


## 9.3 [0.7.2]-2020-10-09

### 9.3.1 Release Notes

We've added a list of all papers using (and citing) TeNPy, see Papers using TeNPy. Feel free to include your own works!

And a slight simplicifation, which might affect your code: using the MultiCouplingModel is no longer necessary, just use the tenpy.models.model. CouplingModel directly.

### 9.3.2 Changelog

## Backwards incompatible changes

- Deprecated the tenpy.models.model.MultiCouplingModel. The functionality is fully merged into the CouplingModel, no need to subclass the MultiCouplingModel anymore.
- The Kagome lattice did not include all next_next_nearest_neighbors. (It had only the ones across the hexagon, missing those maiking up a bow-tie.)
- Combined arguments onsite_terms and coupling_terms of tenpy.networks.mpo.mPOGraph. from_terms () into a single argument terms.


## Added

- Allow to include jupyter notebooks into the documentation; collect example notebooks in [TeNPyNotebooks].
- term_correlation_function_right() and term_correlation_function_left() for correlation functions with more than one operator on each end.
- tenpy.networks.terms.ExponentiallyDecayingTerms for constructing MPOs with exponential decay, and tenpy.networks.model.CouplingModel. add_exponentially_decaying_coupling() for using it. This closes issue \#78.

Fixed

- The IrregularLattice used the 'default' order of the regular lattice instead of whatever the order of the regular lattice was.
- charge_variance () did not work for more than 1 charge.


## 9.4 [0.7.1] - 2020-09-04

### 9.4.1 Release Notes

This is just a minor fix to allow building the conda package

## 9.5 [0.7.0]-2020-09-04

### 9.5.1 Release Notes

The big new feature is the implementation of the W_I and W_II method for approximating exponentials of an MPO with an MPO, and MPS compression / MPO application to an MPS, to allow time evolution with ExpMPOEvolution().

### 9.5.2 Changelog

## Backwards incompatible changes

- Remove argument leg0 from build_MPO.
- Remove argument leg0 from from_grids, instead optionally give all legs as argument.
- Moved/renamed the module tenpy.algorithms.mps_sweeps to tenpy.algorithms. mps_common. The old mps_sweeps still exists for compatibility, but raises a warning upon import.
- Moved/renamed the module tenpy.algorithms.purification_tebd to tenpy.algorithms.purification (for the PurificaitonTEBD and PurificationTEBD2) and tenpy.algorithms.disentangler (for the disentanglers).


## Added

- VariationalCompression and VariationalApplyMPO for variational compression
- PurificationApplyMPO and PurificationTwoSiteU for variational compression with purifications.
- Argument insert_all_id for tenpy.networks.mpo.MPOGraph.from_terms() and from_term_list()
- implemented the IrregularLattice.
- extended user guide on lattices, Details on the lattice geometry.
- Function to approximate a decaying function by a sum of exponentials.
- spatial_inversion () to perform an explicit spatial inversion of the MPS.


## Changed

- By default, for an usual MPO define $I d L$ and $I d R$ on all bonds. This can generate "dead ends" in the MPO graph of finite systems, but it is useful for the make_WI/make_WII for MPO-exponentiation.
- tenpy.models.lattice.Lattice.plot_basis() now allows to shade the unit cell and shift the origin of the plotted basis.
- Don't use bc_shift in tenpy.models.lattice.Lattice.plot_couplings() any more - it lead to confusing figures. Instead, the new keyword wrap=True allows to directly connect all sites. This is done to avoid confusing in combination with plot_bc_identified().
- Error handling of non-zero qtotal for TransferMatrix.


## Fixed

- Removed double counting of chemical potential terms in the BosonicHaldaneModel and FermionicHaldaneModel.
- Wrong results of tenpy.networks.mps.MPS.get_total_charge() with only_physical_legs=True.
- tenpy.models.Iattice.Lattice.plot_bc_identified() had a sign error for the bc_shift.
- calc_H_MPO_from_bond () didn't work for charges with blocks $>1$.
- TEBD: keep qtotal of the B tensors constant
- order model parameter was read out but not used in tenpy.models.model.CouplingMPOModel. init_lattice() for 1D lattices.


## 9.6 [0.6.1]-2020-05-18

### 9.6.1 Release Notes

This only is a follow-up release to [0.6.0]-2020-05-16. It fixes a small bug in the examples/c_tebd.py and some roundoff problems in the tests.

It is now possible to install TeNPy with the conda package manager:

```
conda install --channel=conda-forge physics-tenpy
```


## 9.7 [0.6.0] - 2020-05-16

### 9.7.1 Release Notes

This release contains a major update of the documentation, which is now hosted by "Read the Docs" at https://tenpy. readthedocs.io/. Update your bookmark :-)

Apart from that, this release introduces a format how to save and load data (in particular TeNPy classes) to HDF5 files. See Saving to disk: input/output for more details. To use that feature, you need to install the h5py package (and therefore some version of the HDF5 library). This is easy with anaconda, conda install h5py, but might be cumbersome on your local computing cluster. (However, many university computing clusters have some version of HDF5 installed already. Check with your local sysadmin.)

Moreover, we changed how we read out parameter dictionaries - instead of the get_parameter() function, we have now a Config class which behaves like a dictionary, you can simpy use options.get (key, default) for model parameters - as you would do for a python dictionary.

### 9.7.2 Changelog

## Backwards incompatible changes

- Created a class Config to replace Python-native parameter dictionaries and add some useful functionality. Old code using tenpy.tools.params.get_parameter() and tenpy.tools.params. unused_parameters () still works as before, but raises a warning, and should be replaced. For example, if you defined your own models, you should replace calls get_parameter (model_params, "key ", "default_value", "ModelName") with model_params.get("key", "default_value ") , the latter syntax being what you would use for a normal python dictionary as well.
- Renamed the following class parameter dictionaries to simply options for more consitency. Old code using the class attributes should still work (since we provide property aliases), but raises warnings. Note that this affects also derived classes (for example the TwoSiteDMRGEngine).

```
- tenpy.algorithms.dmrg.DMRGEngine.DMRG_params (was already renamed to en-
    gine_params in versin 0.5.0)
- tenpy.algorithms.mps_common.Sweep.engine_params
- tenpy.algorithms.tebd.Engine.TEBD_params
```

```
- tenpy.algorithms.tdvp.Engine.TDVP_params
- tenpy.linalg.lanczos.Lanczos
```

- Changed the arguments of tenpy.models.model.MultiCouplingModel(): We replaced the three arguments $u 0$, op0 and other_op with other_ops $=[(\mathrm{u} 1, \mathrm{op} 1, \mathrm{dx} 1)$, (op2, u2, dx2), ...] by single, equivalent argment ops which should now read ops=[(op0, dx0, u0), (op1, dx1, u1), $(o p 2, d x 2, u 2), \ldots]$, where $d x 0=[0] *$ lat.dim. Note the changed order inside the tuple! Old code (which specifies opstr and category as keyword argument, if at all) still works as before, but raises a warning, and should be replaced. Since tenpy.lattice.Lattice.possible_multi_couplings() used similar arguments, they were changed as well.
- Don't save $H_{-} M P O \_g r a p h$ as model attribute anymore - this also wasn't documented.
- Renamed the truncation parameter symmetry_tol to degeneracy_tol and make the criterion more reasonable by not checking $\log \left(S_{i} / S_{j}\right)<\log \left(\right.$ symmetry $\left._{t} o l\right)$, but simply $\log \left(S_{i} / S_{j}\right)<$ degeneracy $_{t} o l$. The latter makes more sense, as it is equivalent to $\left(S_{i}-S_{j}\right) / S_{j}<\exp \left(\right.$ degeneracy $\left._{t} o l\right)-1=$ degeneracy $_{t} o l+$ $\mathcal{O}\left(\right.$ degeneracy $_{t}$ ol $\left.^{2}\right)$.
- Deprecated tenpy.networks.mps.MPS.increase_L() in favor of the newly added tenpy. networks.mps.MPS.enlarge_mps_unit_cell() (taking factor instead of new_L=factor*L as argument).
- tenpy.networks.mps.MPS.correlation_function() now auto-determines whether a JordanWigner string is necessary. If any of the given operators is directly an npc Array, it will now raise an error; set autoJW=False in that case.
- Instead of "monkey-patching" matvec of the tenpy.algorithms.mps_common.EffectiveH for the case that ortho_to_envs is not empty, we defined a proper class NpcLinearOperatorWrapper, which serves as baseclass for OrthogonalNpcLinearOperator. The argument ortho_to_envs has been removed from EffectiveH.
- Switch order of the sites in the unit cell for the DualSquare, and redefine what the "default" order means. This is a huge optimization of DMRG, reducing the necessary MPS bond dimension for the ground state to the optimal $2^{L-1}$ on each bond.
- Deprecated the Lanczos funciton/class argument orthogonal_to of in LanczosGroundState. Instead, one can use the OrthogonalNpcLinearOperator.
- Deprecation warning for changing the default argument of shift_ket for non-zero shift_bra of the TransferMatrix.


## Added

- tenpy.networks.mpo.MPO.variance () to calculate the variance of an MPO against a finite MPS.
- Classmethod tenpy.networks.MPS.from_lat_product_state() to initialize an MPS from a product state given in lattice coordinates (independent of the order of the lattice).
- argument plus_hc for tenpy.models.model.CouplingModel.add_onsite(), tenpy. models.model.CouplingModel.add_coupling(), and tenpy.models.model. MultiCouplingModel.add_multi_coupling() to simplify adding the hermitian conjugate terms.
- parameter explicit_plus_hc for MPOMOdel, CouplingModel and MPO, to reduce MPO bond dimension by not storing Hermitian conjugate terms, but computing them at runtime.
- tenpy.models.model.CouplingModel.add_local_term() for adding a single term to the lattice, and still handling Jordan-Wigner strings etc.
- tenpy.networks.site.Site.get_hc_opname() and hc_ops to allow getting the hermitian conjugate operator (name) of the onsite operators.
- tenpy.tools.hdf5_io with convenience functions for import and output with pickle, as well as an implementation allowing to save and load objects to HDF5 files in the format specified in Saving to disk: input/output.
- human-readable boundary_conditions property in Lattice.
- save_hdf5 and load_hdf5 methods to support saving/loading to HDF5 for the following classes (and their subclasses): - ChargeInfo-LegCharge - LegPipe - Array - MP S - MPO-Lattice
- tenpy.networks.mps.MPSEnvironment.get_initialization_data() for a convenient way of saving the necessary parts of the environment after an DMRG run.
- Method enlarge_mps_unit_cell for the following classes: - MPS - MPO-Lattice - Model, MPOModel, NearestNeighborModel
- tenpy.tools.misc.to_iterable_of_len() for convenience of handling arguments.
- tenpy.models.lattice.Lattice.mps2lat_values_masked() as generalization of tenpy. models.lattice.Lattice.mps2lat_values().
- tenpy. Iinalg.sparse. OrthogonalNpcLinearOperator to orthogonalize against vectors.
- tenpy.linalg.sparse. ShiftNpcLinearOperator to add a constant.
- tenpy. Iinalg.sparse.SumNpcLinearOperator which serves e.g. to add the h.c. during the matvec (in combination with the new tenpy.linalg. sparse. NpcLinearOperator.adjoint ()).
- tenpy.algorithms.mps_common.make_eff_H() to simplify implementations of prepare_update().
- attribute options for the Model.
- tenpy.networks.mps.MPS.roll_mps_unit_cell().


## Changed

- DEFAULT DMRG paramter 'diag_method' from 'lanczos' to 'default', which is the same for large bond dimensions, but performs a full exact diagonalization if the effective Hamiltonian has small dimensions. The threshold introduced is the new DMRG parameter 'max_N_for_ED'.
- DEFAULT parameter charge_sector=None instead of charge_sector=0 in tenpy.networks. mps.MPS.overlap () to look for eigenvalues of the transfer matrix in all charge sectors, and not assume that it's the 0 sector.
- Derive the following classes (and their subclasses) from the new Hdf5Exportable to support saving to HDF5: - Site-Terms - OnsiteTerms-CouplingTerms-Model, i.e., all model classes.
- Instead of just defining to_matrix and adjoint for EffectiveH, define the interface directly for NpcLinearOperator.
- Try to keep the charge block structure as far as possible for add_charge () and drop_charge ()


## Fixed

- Adjust the default DMRG parameter min_sweeps if chi_list is set.
- Avoid some unnecessary transpositions in MPO environments for MPS sweeps (e.g. in DMRG).
- sort (bunch=True) could return un-bunched Array, but still set the bunched flag.
- LegPipe did not initialize self.bunched correctly.
- issue \#98: Error of calling psi.canonical_form() directly after disabling the DMRG mixer.
- svd() with full_matrices=True gave wrong charges.
- tenpy.linalg.np_conserved.Array.drop_charge() and tenpy.lina.np_conserved. Array.drop_charge () did not copy over labels.
- wrong pairs for the fifth_nearest_neighbors of the Honeycomb.
- Continue in tenpy.algorithms.dmrg.full_diag_effH () with a warning instaed of raising an Error, if the effective Hamltonian is zero.
- correlation_length (): check for hermitian Flag might have raised and Error with new numpy warnings
- correlation_function() did not respect argument str_on_first=False.
- tenpy.networks.mps.MPS.get_op() worked unexpected for infinite $b c$ with incomensurate self.L and len(op_list).
- tenpy.networks.mps.MPS.permute_sites() did modify the given perm.
- issue \#105 Unintended side-effects using lanczos_params.verbose in combination with orthogonal_to
- issue \#108 tenpy.linalg.sparse.FlatLinearOperator._matvec() changes self. _charge_sector


## 9.8 [0.5.0]-2019-12-18

### 9.8.1 Backwards incompatible changes

- Major rewriting of the DMRG Engines, see issue \#39 and issue \#85 for details. The EngineCombine and EngineFracture have been combined into a single TwoSiteDMRGEngine with an The run function works as before. In case you have directly used the EngineCombine or EngineFracture, you should update your code and use the TwoSiteEngine instead.
- Moved init_LP and init_RP method from MPS into MPSEnvironment and MPOEnvironment.


### 9.8.2 Changed

- Addition/subtraction of Array: check whether the both arrays have the same labels in differnt order, and in that case raise a warning that we will transpose in the future.
- Made tenpy. Iinalg.np_conserved.Array.get_block() public (previously tenpy.linalg. np_conserved.Array._get_block).
- groundstate() now returns a tuple (E0, psi0) instead of just psi0. Moreover, the argument charge_sector was added.
- Simplification in the Lattice: Instead of having separate arguments/attributes/functions for 'nearest_neighbors', 'next_nearest_neighbors', 'next_next_nearest_neighbors' and possibly (Honeycomb) even 'fourth_nearest_neighbors', 'fifth_nearest_neighbors', collect them in a dictionary called pairs. Old call structures still allowed, but deprecated.
- issue \#94: Array addition and inner () should reflect the order of the labels, if they coincided. Will change the default behaviour in the future, raising FutureWarning for now.
- Default parameter for DMRG params: increased precision by setting $P$ _tol_min down to the maximum of $1 . e-30$, lanczos_params['svd_min']**2 * P_tol_to_trunc, lanczos_params['trunc_cut']**2 * P_tol_to_trunc by default.


### 9.8.3 Added

- tenpy.algorithms.mps_common with the Sweep class and EffectiveH to be a OneSiteH or TwoSiteH.
- Single-Site DMRG with the SingleSiteDMRG.
- Example function in examples/c_tebd.py how to run TEBD with a model originally having next-nearest neighbors.
- increase_L () to allow increasing the unit cell of an MPS.
- Additional option order='folded' for the Chain.
- tenpy.algorithms.exact_diag.ExactDiag.from_H_mpo() wrapper as replacement for tenpy.networks.mpo.MPO.get_full_hamiltonian() and tenpy.networks.mpo.MPO. get_grouped_mpo (). The latter are now deprecated.
- Argument max_size to limit the matrix dimension in ExactDiag.
- tenpy.linalg.sparse.FlatLinearOperator.from_guess_with_pipe() to allow quickly converting matvec functions acting on multi-dimensional arrays to a FlatLinearOperator by combining the legs into a LegPipe.
- tenpy.tools.math.speigsh() for hermitian variant of speigs()
- Allow for arguments 'LA', 'SA' in argsort().
- tenpy.linalg.lanczos.lanczos_arpack() as possiple replacement of the self-implemented lanczos function.
- tenpy.algorithms.dmrg.full_diag_effH() as another replacement of lanczos().
- The new DMRG parameter 'diag_method' allows to select a method for the diagonalization of the effective Hamiltonian. See tenpy.algorithms.dmrg.DMRGEngine.diag() for details.
- dtype attribute in EffectiveH.
- tenpy.linalg.charges.LegCharge.get_qindex_of_charges() to allow selecting a block of an Array from the charges.
- tenpy.algorithms.mps_common.EffectiveH.to_matrix to allow contracting an EffectiveH to a matrix, as well as metadata tenpy.linalg.sparse.NpcLinearoperator.acts_on and tenpy. algorithms.mps_common.EffectiveH.N.
- argument only_physical_legs in tenpy.networks.mps.MPS.get_total_charge()


### 9.8.4 Fixed

- MPO expectation_value () did not work for finite systems.
- Calling compute_K () repeatedly with default parameters but on states with different chi would use the chi of the very first call for the truncation parameters.
- allow MPSEnvironment and MPOEnvironment to have MPS/MPO with different length
- group_sites () didn't work correctly in some situations.
- matvec_to_array () returned the transposed of A.
- tenpy.networks.mps.MPS.from_full() messed up the form of the first array.
- issue \#95: blowup of errors in DMRG with update_env $>0$. Turns out to be a problem in the precision of the truncation error: TruncationErroreps was set to 0 if it would be smaller than machine precision. To fix it, I added from_S ().


## 9.9 [0.4.1]-2019-08-14

### 9.9.1 Backwards incompatible changes

- Switch the sign of the BoseHubbardModel and FermiHubbardModel to hopping and chemical potential having negative prefactors. Of course, the same adjustment happens in the BoseHubbardChain and FermiHubbardChain.
- moved BoseHubbardModel and BoseHubbardChain as well as FermiHubbardModel and FermiHubbardChain into the new module tenpy.models.hubbard.
- Change arguments of coupling_term_handle_JW() and multi_coupling_term_handle_JW() to use strength and sites instead of op_needs_JW.
- Only accept valid identifiers as operator names in add_op ().


### 9.9.2 Changed

- grid_concat () allows for None entries (representing zero blocks).
- from_full () allows for 'segment' boundary conditions.
- apply_local_op() allows for n-site operators.


### 9.9.3 Added

- max_range attribute in MPO and MPOGraph.
- is_hermitian()
- Nearest-neighbor interaction in BoseHubbardModel
- multiply_op_names() to replace ' '.join(op_names) and allow explicit compression/multiplication.
- order_combine_term() to group operators together.
- dagger () of MPO's (and to implement that also flip_charges_qconj()).
- has_label () to check if a label exists
- qr_li() and rq_li()
- Addition of MPOs
- 3 additional examples for chern insulators in examples/chern_insulators/.
- FermionicHaldaneModel and BosonicHaldaneModel.
- from_MPOMOdel () for initializing nearest-neighbor models after grouping sites.


### 9.9.4 Fixed

- issue \#36: long-range couplings could give IndexError.
- issue \#42: Onsite-terms in FermiHubbardModel were wrong for lattices with non-trivial unit cell.
- Missing a factor 0.5 in GUE ().
- Allow TermList to have terms with multiple operators acting on the same site.
- Allow MPS indices outside unit cell in mps2lat_idx() and lat2mps_idx().
- expectation_value () did not work for n-site operators.


### 9.10 [0.4.0]-2019-04-28

### 9.10.1 Backwards incompatible changes

- The argument order of tenpy.models.lattice.Lattice could be a tuple (priority, snake_winding) before. This is no longer valid and needs to be replaced by ("standard", snake_winding, priority).
- Moved the boundary conditions bc_coupling from the tenpy.models.model.CouplingModel into the tenpy.models.Iattice.Lattice (as bc). Using the parameter bc_coupling will raise a FutureWarning, one should set the boundary conditions directly in the lattice.
- Added parameter permute (True by default) in tenpy.networks.mps.MPS.from_product_state () and tenpy.networks.mps.MPS.from_Bflat(). The resulting state will therefore be independent of the "conserve" parameter of the Sites - unlike before, where the meaning of the p_state argument might have changed.
- Generalize and rename tenpy.networks.site.DoubleSite to tenpy.networks.site. GroupedSite, to allow for an arbitrary number of sites to be grouped. Arguments site0, site1, label0, label1 of the __init__ can be replaced with [site0, site1], [label0, label1] and op0, op1 of the kronecker_product with [op0, op1] ; this will recover the functionality of the DoubleSite.
- Restructured callstructure of Mixer in DMRG, allowing an implementation of other mixers. To enable the mixer, set the DMRG parameter "mixer" to True or 'DensityMatrixMixer' instead of just 'Mixer'.
- The interaction parameter in the tenpy.models.bose_hubbbard_chain.BoseHubbardModel (and tenpy.models.bose_hubbbard_chain. BoseHubbardChain) did not correspond to $U / 2 N(N-1)$ as claimed in the Hamiltonian, but to $U N^{2}$. The correcting factor $1 / 2$ and change in the chemical potential have been fixed.
- Major restructuring of tenpy.linalg.np_conserved and tenpy.linalg.charges. This should not break backwards-compatibility, but if you compiled the cython files, you need to remove the old binaries in the source directory. Using bash cleanup. sh might be helpful to do that, but also remove other files within the repository, so be careful and make a backup beforehand to be on the save side. Afterwards recompile with bash compile.sh.
- Changed structure of tenpy.models.model.CouplingModel.onsite_terms and tenpy. models.model. CouplingModel. coupling_terms: Each of them is now a dictionary with category strings as keys and the newly introduced tenpy. networks.terms.OnsiteTerms and tenpy. networks.terms. CouplingTerms as values.
- tenpy.models.model.CouplingModel.calc_H_onsite() is deprecated in favor of new methods.
- Argument raise_op2_left of tenpy.models.model.CouplingModel.add_coupling() is deprecated.


### 9.10.2 Added

- tenpy.networks.mps.MPS.canonical_form_infinite().
- tenpy.networks.mps.MPS.expectation_value_term(), tenpy.networks. mps.MPS.expectation_value_terms_sum() and tenpy.networks.mps.MPS. expectation_value_multi_sites() for expectation values of terms.
- tenpy.networks.mpo.MPO.expectation_value() for an MPO.
- tenpy. Iinalg.np_conserved.Array.extend() and tenpy.Iinalg.charges.LegCharge. extend(), allowing to extend an Array with zeros.
- DMRG parameter 'orthogonal_to' allows to calculate excited states for finite systems.
- possibility to change the number of charges after creating LegCharges/Arrays.
- more general way to specify the order of sites in a tenpy.models. Iattice. Lattice.
- new tenpy.models.lattice. Triangular, tenpy.models.lattice. Honeycomb and tenpy. models.lattice. Kagome lattice
- a way to specify nearest neighbor couplings in a Lattice, along with methods to count the number of nearest neighbors for sites in the bulk, and a way to plot them (plot_coupling() and friends)
- tenpy.networks.mpo.MPO.from_grids() to generate the MPO from a grid.
- tenpy.models.model. MultiCouplingModel for couplings involving more than 2 sites.
- request \#8: Allow shift in boundary conditions of CouplingModel.
- Allow to use state labels in tenpy. networks.mps.MPS.from_product_state().
- tenpy.models.model. CouplingMPOModel structuring the default initialization of most models.
- Allow to force periodic boundary conditions for finite MPS in the CouplingMPOModel. This is not recommended, though.
- tenpy.models.model.NearestNeighborModel.calc_H_MPO_from_bond() and tenpy. models.model.MPOModel.calc_H_bond_from_MPO() for conversion of H_bond into H_MPO and vice versa.
- tenpy.algorithms.tebd.RandomUnitaryEvolution for random unitary circuits
- Allow documentation links to github issues, arXiv, papers by doi and the forum with e.g. :issue:`5`, :arxiv:`1805.00055`, :doi:`10.21468/SciPostPhysLectNotes.5`, :forum:`3`
- tenpy.models.model.CouplingModel.coupling_strength_add_ext_flux() for adding hoppings with external flux.
- tenpy.models.model.CouplingModel.plot_coupling_terms() to visualize the added coupling terms.
-tenpy.networks.terms.OnsiteTerms, tenpy.networks.terms. CouplingTerms, tenpy. networks.terms.MultiCouplingTerm containing the of terms for the CouplingModel and MultiCouplingModel. This allowed to add the category argument to add_onsite, add_coupling and add_multi_coupling.
- tenpy. networks.terms. TermList as another (more human readable) representation of terms with conversion from and to the other $*$ Term classes.
- tenpy.networks.mps.MPS.init_LP() and tenpy.networks.mps.MPS.init_RP() to initialize left and right parts of an Environment.
-tenpy.networks.mpo.MPOGraph.from_terms() and tenpy.networks.mpo.MPOGraph. from_term_list().
- argument charge_sector in tenpy. networks.mps.MPS.correlation_length().


### 9.10.3 Changed

- moved toycodes from the folder examples/ to a new folder toycodes / to separate them clearly.
- major remodelling of the internals of tenpy. linalg.np_conserved and tenpy.linalg.charges.
- Introduced the new module tenpy/linalg/_npc_helper.pyx which contains all the Cython code, and gets imported by
- Array now rejects addition/subtraction with other types
- Array now rejects multiplication/division with non-scalar types
- By default, make deep copies of npc Arrays.
- Restructured lanczos into a class, added time evolution calculating exp (A*dt) | psi0>
- Warning for poorly conditioned Lanczos; to overcome this enable the new parameter reortho.
- Simplified call strucutre of extend(), and extend().
- Restructured tenpy.algorithms.dmrg:
- run() is now just a wrapper around the new run(), run(psi, model, pars) is roughly equivalent to eng = EngineCombine(psi, model, pars); eng.run().
- Added init_env() and reset_stats() to allow a simple restart of DMRG with slightly different parameters, e.g. for tuning Hamiltonian parameters.
- Call canonical_form () for infinite systems if the final state is not in canonical form.
- Changed default values for some parameters:
- set trunc_params ['chi_max'] = 100. Not setting a chi_max at all will lead to memory problems. Disable DMRG_params ['chi_list'] = None by default to avoid conflicting settings.
- reduce to mixer_params['amplitude'] = 1.e-5. A too strong mixer screws DMRG up pretty bad.
- increase Lanczos_params['N_cache'] = N_max (i.e., keep all states)
- set DMRG_params['P_tol_to_trunc'] = 0.05 and provide reasonable ..._min and ..._max values.
- increased (default) DMRG accuracy by setting DMRG_params['max_E_err'] = $1 . e-8$ and DMRG_params['max_S_err'] = 1.e-5.
- don't check the (absolute) energy for convergence in Lanczos.
- set DMRG_params ['norm_tol'] = 1.e-5 to check whether the final state is in canonical form.
- Verbosity of get_parameter () reduced: Print parameters only for verbosity $>=1$. and default values only for verbosity $>=2$.
- Don't print the energy during real-time TEBD evolution - it's preserved up to truncation errors.
- Renamed the SquareLattice class to tenpy.models. Iattice. Square for better consistency.
- auto-determine whether Jordan-Wigner strings are necessary in add_coupling().
- The way the labels of npc Arrays are stored internally changed to a simple list with None entries. There is a deprecated propery setter yielding a dictionary with the labels.
- renamed first_LP and last_RP arguments of MPSEnvironment and MPOEnvironment to init_LP and init_RP.
- Testing: insetad of the (outdated) nose, we now use pytest [https://pytest.org](https://pytest.org) for testing.


### 9.10.4 Fixed

- issue \#22: Serious bug in tenpy. linalg.np_conserved.inner(): if do_conj=True is used with non-zero qtotal, it returned 0 . instead of non-zero values.
- avoid error in tenpy.networks.mps.MPS.apply_local_op()
- Don't carry around total charge when using DMRG with a mixer
- Corrected couplings of the FermionicHubbardChain
- issue \#2: memory leak in cython parts when using intelpython/anaconda
- issue \#4: incompatible data types.
- issue \#6: the CouplingModel generated wrong Couplings in some cases
- issue \#19: Convergence of energy was slow for infinite systems with N_sweeps_check=1
- more reasonable traceback in case of wrong labels
- wrong dtype of npc.Array when adding/subtracting/... arrays of different data types
- could get wrong H_bond for completely decoupled chains.
- SVD could return outer indices with different axes
- tenpy.networks.mps.MPS.overlap () works now for MPS with different total charge (e.g. after psi. apply_local_op(i, 'Sp')).
- skip existing graph edges in MPOGraph.add() when building up terms without the strength part.


### 9.10.5 Removed

- Attribute chinfo of Lattice.


### 9.11 [0.3.0] - 2018-02-19

This is the first version published on github.

### 9.11.1 Added

- Cython modules for np_conserved and charges, which can optionally be compiled for speed-ups
- tools.optimization for dynamical optimization
- Various models.
- More predefined lattice sites.
- Example toy-codes.
- Network contractor for general networks


### 9.11.2 Changed

- Switch to python3


### 9.11.3 Removed

- Python 2 support.


### 9.12 [0.2.0] - 2017-02-24

- Compatible with python2 and python3 (using the 2 to 3 tool).
- Development version.
- Includes TEBD and DMRG.


### 9.13 Changes compared to previous TeNPy

This library is based on a previous (closed source) version developed mainly by Frank Pollmann, Michael P. Zaletel and Roger S. K. Mong. While allmost all files are completely rewritten and not backwards compatible, the overall structure is similar. In the following, we list only the most important changes.

### 9.13.1 Global Changes

- syntax style based on PEP8. Use \$>yapf -r -i . / to ensure consitent formatting over the whole project. Special comments \# yapf: disable and \# yapf: enable can be used for manual formatting of some regions in code.
- Following PEP8, we distinguish between 'private' functions, indicated by names starting with an underscore and to be used only within the library, and the public API. The puplic API should be backwards-compatible with different releases, while private functions might change at any time.
- all modules are in the folder tenpy to avoid name conflicts with other libraries.
- withing the library, relative imports are used, e.g., from ..tools.math import (toiterable, tonparray) Exception: the files in tests/ and examples/ run as $\qquad$ main $\qquad$ and can't use relative imports

Files outside of the library (and in tests/, examples/) should use absolute imports, e.g. import tenpy. algorithms.tebd

- renamed tenpy/mps/ to tenpy/networks, since it containes various tensor networks.
- added Site describing the local physical sites by providing the physical LegCharge and onsite operators.


### 9.13.2 np_conserved

- pure python, no need to compile!
- in module tenpy. linalg instead of algorithms/linalg.
- moved functionality for charges to charges
- Introduced the classes ChargeInfo (basically the old q_number, and mod_q) and LegCharge (the old qind, qconj).
- Introduced the class LegPipe to replace the old leg_pipe. It is derived from LegCharge and used as a leg in the array class. Thus any inherited array (after tensordot etc still has all the necessary information to split the legs. (The legs are shared between different arrays, so it's saved only once in memory)
- Enhanced indexing of the array class to support slices and 1D index arrays along certain axes
- more functions, e.g. grid_outer()


### 9.13.3 TEBD

- Introduced TruncationError for easy handling of total truncation error.
- some truncation parameters are renamed and may have a different meaning, e.g. svd_max -> svd_min has no ' $\log ^{\prime}$ ' in the definition.


### 9.13.4 DMRG

- separate Lanczos module in tenpy/linalg/. Strangely, the old version orthoganalized against the complex conjugates of orthogonal_to (contrary to it's doc string!) (and thus calculated 'theta_o' as bra, not ket).
- cleaned up, provide prototypes for DMRG engine and mixer.


### 9.13.5 Tools

- added tenpy.tools.misc, which contains 'random stuff' from old tools.math like to_iterable and to_array (renamed to follow PEP8, documented)
- moved stuff for fitting to tenpy.tools.fit
- enhanced tenpy.tools.string.vert_join() for nice formatting
- moved (parts of) old cluster/omp.py to tenpy.tools.process
- added tenpy.tools.params for a simplified handling of parameter/arguments for models and/or algorithms. Similar as the old models.model.set_var, but use it also for algorithms. Also, it may modify the given dictionary.


## INTRODUCTIONS

The following documents are meant as introductions to various topics relevant to TeNPy.
If you are new to TeNPy, read the Overview.

### 10.1 Overview

### 10.1.1 Repository

The root directory of the git repository contains the following folders:
tenpy The actual source code of the library. Every subfolder contains an __init__. py file with a summary what the modules in it are good for. (This file is also necessary to mark the folder as part of the python package. Consequently, other subfolders of the git repo should not include a $\qquad$ init $\qquad$ . py file.)
toycodes Simple toy codes completely independet of the remaining library (i.e., codes in tenpy/). These codes should be quite readable and intend to give a flavor of how (some of) the algorithms work.
examples Some example files demonstrating the usage and interface of the library.
doc A folder containing the documentation: the user guide is contained in the $*$. rst files. The online documentation is autogenerated from these files and the docstrings of the library. This folder contains a make file for building the documentation, run make help for the different options. The necessary files for the reference in doc/ reference can be auto-generated/updated with make src2html.
tests Contains files with test routines, to be used with pytest. If you are set up correctly and have pytest installed, you can run the test suite with pytest from within the tests/folder.
build This folder is not distributed with the code, but is generated by setup.py (or compile.sh, respectively). It contains compiled versions of the Cython files, and can be ignored (and even removed without loosing functionality).

### 10.1.2 Code structure: getting started

There are several layers of abstraction in TeNPy. While there is a certain hierarchy of how the concepts build up on each other, the user can decide to utilize only some of them. A maximal flexibility is provided by an object oriented style based on classes, which can be inherited and adjusted to individual demands.

The following figure gives an overview of the most important modules, classes and functions in TeNPy. Gray backgrounds indicate (sub)modules, yellow backgrounds indicate classes. Red arrows indicate inheritance relations, dashed black arrows indicate a direct use. (The individual models might be derived from the NearestNeighborModel depending on the geometry of the lattice.) There is a clear hierarchy from high-level algorithms in the tenpy. algorithms module down to basic operations from linear algebra in the tenpy. linalg module.


## Most basic level: linear algebra

Note: See Charge conservation with np_conserved for more information on defining charges for arrays.

The most basic layer is given by in the linalg module, which provides basic features of linear algebra. In particular, the np_conserved submodule implements an Array class which is used to represent the tensors. The basic interface of np_conserved is very similar to that of the NumPy and SciPy libraries. However, the Array class implements abelian charge conservation. If no charges are to be used, one can use 'trivial' arrays, as shown in the following example code.

```
"""Basic use of the `Array` class with trivial arrays."""
# Copyright 2019-2021 TeNPy Developers, GNU GPLv3
import tenpy.linalg.np_conserved as npc
M = npc.Array.from_ndarray_trivial([[0., 1.], [1., 0.]])
v = npc.Array.from_ndarray_trivial([2., 4. + 1.j])
v[0] = 3. # set indiviual entries like in numpy
print(" | v> =", v.to_ndarray())
# |v> = [ 3.+0.j 4.+1.j]
M_v = npc.tensordot(M, v, axes=[1, 0])
print("M|v> =", M_v.to_ndarray())
# M/v> = [ 4.+1.j 3.+0.j]
print("<v|M|v> =", npc.inner(v.conj(), M_v, axes='range'))
# <v|M|v> = (24+0j)
```

The number and types of symmetries are specified in a ChargeInfo class. An Array instance represents a tensor satisfying a charge rule specifying which blocks of it are nonzero. Internally, it stores only the non-zero blocks of the tensor, along with one LegCharge instance for each leg, which contains the charges and sign qconj for each leg. We can combine multiple legs into a single larger LegPipe, which is derived from the LegCharge and stores all the information necessary to later split the pipe.

The following code explicitly defines the spin-1/2 $S^{+}, S^{-}, S^{z}$ operators and uses them to generate and diagonalize the two-site Hamiltonian $H=\vec{S} \cdot \vec{S}$. It prints the charge values (by default sorted ascending) and the eigenvalues of H .

```
"""Explicit definition of charges and spin-1/2 operators."""
# Copyright 2019-2021 TeNPy Developers, GNU GPLv3
import tenpy.linalg.np_conserved as npc
# consider spin-1/2 with Sz-conservation
chinfo = npc.ChargeInfo([1]) # just a U(I) charge
# charges for up, down state
p_leg = npc.LegCharge.from_qflat(chinfo, [[1], [-1]])
Sz = npc.Array.from_ndarray([[0.5, 0.], [0., -0.5]], [p_leg, p_leg.conj()])
Sp = npc.Array.from_ndarray([[0., 1.], [0., 0.]], [p_leg, p_leg.conj()])
Sm = npc.Array.from_ndarray([[0., 0.], [1., 0.]], [p_leg, p_leg.conj()])
Hxy = 0.5 * (npc.outer(Sp, Sm) + npc.outer(Sm, Sp))
Hz = npc.outer(Sz, Sz)
H = Hxy + Hz
# here, H has 4 legs
H.iset_leg_labels(["s1", "t1", "s2", "t2"])
H = H.combine_legs([["s1", "s2"], ["t1", "t2"]], qconj=[+1, -1])
```

```
# here, H has 2 legs
print(H.legs[0].to_qflat().flatten())
# prints [\begin{array}{lllll}{-2}&{0}&{0}&{2}\end{array}]
E, U = npc.eigh(H) # diagonalize blocks individually
print(E)
# [ [ 0.25 -0.75 0.25 0.25]
```


## Sites for the local Hilbert space and tensor networks

The next basic concept is that of a local Hilbert space, which is represented by a Site in TeNPy. This class does not only label the local states and define the charges, but also provides onsite operators. For example, the SpinHalfSite provides the $S^{+}, S^{-}, S^{z}$ operators under the names 'Sp', 'Sm', 'Sz', defined as Array instances similarly as in the code above. Since the most common sites like for example the SpinSite (for general spin $\mathrm{S}=0.5,1,1.5, \ldots$ ), BosonSite and FermionSite are predefined, a user of TeNPy usually does not need to define the local charges and operators explicitly. The total Hilbert space, i.e, the tensor product of the local Hilbert spaces, is then just given by a list of Site instances. If desired, different kinds of Site can be combined in that list. This list is then given to classes representing tensor networks like the MPS and MPO. The tensor network classes also use Array instances for the tensors of the represented network.

The following example illustrates the initialization of a spin- $1 / 2$ site, an MP $S$ representing the Neel state, and an MPO representing the Heisenberg model by explicitly defining the $W$ tensor.

```
"""Initialization of sites, MPS and MPO."""
# Copyright 2019-2021 TeNPy Developers, GNU GPLv3
from tenpy.networks.site import SpinHalfSite
from tenpy.networks.mps import MPS
from tenpy.networks.mpo import MPO
spin = SpinHalfSite(conserve="Sz")
print(spin.Sz.to_ndarray())
# [[ 0.5 0. ]
# [0. -0.5]]
N = 6 # number of sites
sites = [spin] * N # repeat entry of list N times
pstate = ["up", "down"] * (N // 2) # Neel state
psi = MPS.from_product_state(sites, pstate, bc="finite")
print("<Sz> =", psi.expectation_value("Sz"))
# <Sz> = [llllll}0.5-0.5 0.5 -0.5]
print("<Sp_i Sm_j> =", psi.correlation_function("Sp", "Sm"), sep="\n")
# <Sp_i Sm_j> =
# [[1. 0. 0.0.0.0.]
# [0. 0. 0. 0. 0. 0.]
# [0.0. 1. 0. 0. 0.]
# [0.0.0.0.0.0.]
# [0.0.0.0. 1. 0.]
# [0.0.0.0.0.0.]]
# define an MPO
Id, Sp, Sm, Sz = spin.Id, spin.Sp, spin.Sm, spin.Sz
J, Delta, hz = 1., 1., 0.2
W_bulk = [[Id, Sp, Sm, Sz, -hz * Sz], [None, None, None, None, 0.5 * J * Sm],
    [None, None, None, None, 0.5 * J * Sp], [None, None, None, None, J * Delta,
\hookrightarrow* Sz],
```

```
    [None, None, None, None, Id]]
W_first = [W_bulk[0]] # first row
W_last = [[row[-1]] for row in W_bulk] # last column
Ws = [W_first] + [W_bulk] * (N - 2) + [W_last]
H = MPO.from_grids([spin] * N, Ws, bc='finite', IdL=0, IdR=-1)
print("<psi|H|psi> =", H.expectation_value(psi))
# <psi|H/psi> = -1.25
```


## Models

Note: See Models for more information on sites and how to define and extend models on your own.

Technically, the explicit definition of an MPO is already enough to call an algorithm like DMRG in dmrg. However, writing down the $W$ tensors is cumbersome especially for more complicated models. Hence, TeNPy provides another layer of abstraction for the definition of models, which we discuss first. Different kinds of algorithms require different representations of the Hamiltonian. Therefore, the library offers to specify the model abstractly by the individual onsite terms and coupling terms of the Hamiltonian. The following example illustrates this, again for the Heisenberg model.

```
"""Definition of a model: the XXZ chain."""
# Copyright 2019-2021 TeNPy Developers, GNU GPLv3
from tenpy.networks.site import SpinSite
from tenpy.models.lattice import Chain
from tenpy.models.model import CouplingModel, NearestNeighborModel, MPOModel
class XXZChain(CouplingModel, NearestNeighborModel, MPOModel):
    def __init__(self, L=2, S=0.5, J=1., Delta=1., hz=0.):
        spin = SpinSite(S=S, conserve="Sz")
        # the lattice defines the geometry
        lattice = Chain(L, spin, bc="open", bc_MPS="finite")
        CouplingModel.__init__(self, lattice)
        # add terms of the Hamiltonian
        self.add_coupling(J * 0.5, 0, "Sp", 0, "Sm", 1) # Sp_i Sm_{i+1}
        self.add_coupling(J * 0.5, 0, "Sp", 0, "Sm", -1) # Sp_i Sm_{i-1}
        self.add_coupling(J * Delta, 0, "Sz", 0, "Sz", 1)
        # (for site dependent prefactors, the strength can be an array)
        self.add_onsite(-hz, 0, "Sz")
        # finish initialization
        # generate MPO for DMRG
        MPOModel.__init__(self, lat, self.calc_H_MPO())
        # generate H_bond for TEBD
        NearestNeighborModel.__init__(self, lat, self.calc_H_bond())
```

While this generates the same MPO as in the previous code, this example can easily be adjusted and generalized, for example to a higher dimensional lattice by just specifying a different lattice. Internally, the MPO is generated using a finite state machine picture. This allows not only to translate more complicated Hamiltonians into their corresponding MPOs, but also to automate the mapping from a higher dimensional lattice to the 1D chain along which the MPS winds. Note that this mapping introduces longer-range couplings, so the model can no longer be defined to be a NearestNeighborModel suited for TEBD if another lattice than the Chain is to be used. Of course, many
commonly studied models are also predefined. For example, the following code initializes the Heisenberg model on a kagome lattice; the spin liquid nature of the ground state of this model is highly debated in the current literature.

```
"""Initialization of the Heisenberg model on a kagome lattice."""
# Copyright 2019-2021 TeNPy Developers, GNU GPLv3
from tenpy.models.spins import SpinModel
model_params = {
    "S": 0.5, # Spin 1/2
    "lattice": "Kagome",
    "bc_MPS": "infinite",
    "bc_y": "cylinder",
    "Ly": 2, # defines cylinder circumference
    "conserve": "Sz", # use Sz conservation
    "Jx": 1.,
    "Jy": 1.,
    "Jz": 1. # Heisenberg coupling
}
model = SpinModel(model_params)
```


## Algorithms

Another layer is given by algorithms like DMRG and TEBD. Using the previous concepts, setting up a simulation running those algorithms is a matter of just a few lines of code. The following example runs a DMRG simulation, see $d m r g$, exemplary for the transverse field Ising model at the critical point.

```
"""Call of (finite) DMRG."""
# Copyright 2019-2021 TeNPy Developers, GNU GPLv3
from tenpy.networks.mps import MPS
from tenpy.models.tf_ising import TFIChain
from tenpy.algorithms import dmrg
N = 16 # number of sites
model = TFIChain({"L": N, "J": 1., "g": 1., "bc_MPS": "finite"})
sites = model.lat.mps_sites()
psi = MPS.from_product_state(sites, ['up'] * N, "finite")
dmrg_params = {"trunc_params": {"chi_max": 100, "svd_min": 1.e-10}, "mixer": True}
info = dmrg.run(psi, model, dmrg_params)
print("E =", info['E'])
# E = -20.01638790048513
print("max. bond dimension =", max(psi.chi))
# max. bond dimension = 27
```

The switch from DMRG to iDMRG in TeNPy is simply accomplished by a change of the parameter "bc_MPS" from "finite" to "infinite", both for the model and the state. The returned $E$ is then the energy density per site. Due to the translation invariance, one can also evaluate the correlation length, here slightly away from the critical point.

```
"""Call of infinite DMRG."""
# Copyright 2019-2021 TeNPy Developers, GNU GPLv3
from tenpy.networks.mps import MPS
from tenpy.models.tf_ising import TFIChain
from tenpy.algorithms import dmrg
```

(continued from previous page)

```
N = 2 # number of sites in unit cell
model = TFIChain({"L": N, "J": 1., "g": 1.1, "bc_MPS": "infinite"})
sites = model.lat.mps_sites()
psi = MPS.from_product_state(sites, ['up'] * N, "infinite")
dmrg_params = {"trunc_params": {"chi_max": 100, "svd_min": 1.e-10}, "mixer": True}
info = dmrg.run(psi, model, dmrg_params)
print("E =", info['E'])
# E = -1.342864022725017
print("max. bond dimension =", max(psi.chi))
# max. bond dimension = 56
print("corr. length =", psi.correlation_length())
# corr. length = 4.915809146764157
```

Running time evolution with TEBD requires an additional loop, during which the desired observables have to be measured. The following code shows this directly for the infinite version of TEBD.

```
"""Call of (infinite) TEBD."""
# Copyright 2019-2021 TeNPy Developers, GNU GPLv3
from tenpy.networks.mps import MPS
from tenpy.models.tf_ising import TFIChain
from tenpy.algorithms import tebd
M = TFIChain({"L": 2, "J": 1., "g": 1.5, "bc_MPS": "infinite"})
psi = MPS.from_product_state(M.lat.mps_sites(), [0] * 2, "infinite")
tebd_params = {
    "order": 2,
    "delta_tau_list": [0.1, 0.001, 1.e-5],
    "max_error_E": 1.e-6,
    "trunc_params": {
        "chi_max": 30,
        "svd_min": 1.e-10
    }
}
eng = tebd.TEBDEngine(psi, M, tebd_params)
eng.run_GS() # imaginary time evolution with TEBD
print("E =", sum(psi.expectation_value(M.H_bond)) / psi.L)
print("final bond dimensions: ", psi.chi)
```


## Simulations

The top-most layer is given by Simulations. A simulation wraps the whole setup of initializing the Model, MPS and Algorithm classes, running the algorithm, possibly performing measurements, and finally saving results to disk, if desired. It provides some extra functionality like the ability to resume an interrupted simulation, e.g. if your job got killed on the cluster due to runtime limitis.

Ideally, the Simulation (sub) class represents the whole Simulation from start to end, giving re-producable results depending only on the parameters given to it.

### 10.2 Charge conservation with np_conserved

The basic idea is quickly summarized: By inspecting the Hamiltonian, you can identify symmetries, which correspond to conserved quantities, called charges. These charges divide the tensors into different sectors. This can be used to infer for example a block-diagonal structure of certain matrices, which in turn speeds up SVD or diagonalization a lot. Even for more general (non-square-matrix) tensors, charge conservation imposes restrictions which blocks of a tensor can be non-zero. Only those blocks need to be saved, which ultimately (= for large enough arrays) leads to a speedup of many routines, e.g., tensordot.
This introduction covers our implementation of charges; explaining mathematical details of the underlying symmetry is beyond its scope. We refer you to the corresponding chapter in our [TeNPyNotes] for a more general introduction of the idea (also stating the "charge rule" introduced below). [[singh2010]] explains why it works form a mathematical point of view, [[singh2011]] has the focus on a $U(1)$ symmetry and might be easier to read.

### 10.2.1 What you really need to know about np_conserved

The good news is: It is not necessary to understand all the details explained in the following sections if you just want to use TeNPy for "standard" simulations like TEBD and DMRG. In praxis, you will likely not have to define the charges by yourself. For most simulations using TeNPy, the charges are initially defined in the Site; and there are many pre-defined sites like the :class:SpinHalfSite, which you can just use. The sites in turn are initialized by the Model class you are using (see also Models). From there, all the necessary charge information is automatically propagated along with the tensors.

However, you should definitely know a few basic facts about the usage of charge conservation in TeNPy:

- Instead of using numpy arrays, tensors are represented by the Array class. This class is defined in np_conserved (the name standing for "numpy with charge conservation"). Internally, it stores only nonzero blocks of the tensor, which are "compatible" with the charges of the indices. It has to have a well defined overall charge qtotal. This expludes certain operators (like $S^{x}$ for Sz conservation) and MPS which are a superpositions of states in different charge sectors.
- There is a class ChargeInfo holding the general information what kind of charges we have, and a LegCharge for the charge data on a given leg. The leg holds a flag qconj which is +1 or -1 , depending on whether the leg goes into the tensor (representing a vector space) or out of the tensor (representing the corresponding dual vector space).
- Besides the array class methods, there are a bunch of functions like tensordot(), svd() or eigh () to manipulate tensors. These function have a very similar call structure as the corresponding numpy functions, but they act on our tensor Array class, and preserve the block structure (and exploit it for speed, wherever possible).
- The only allowed "reshaping" operations for those tensors are to combine legs and to split previously combined legs. See the correspoding section below.
- It is convenient to use string labels instead of numbers to refer to the various legs of a tensor. The rules how these labels change during the various operations are also described a section below.


### 10.2.2 Introduction to combine_legs, split_legs and LegPipes

Often, it is necessary to "combine" multiple legs into one: for example to perfom a SVD, a tensor needs to be viewed as a matrix. For a flat array, this can be done with np. reshape, e.g., if $A$ has shape $(10,3,7)$ then $B=n p$. reshape ( $\mathrm{A},(30,7$ ) ) will result in a (view of the) array with one less dimension, but a "larger" first leg. By default (order='C'), this results in

```
B[i*3 + j , k] == A[i, j, k] for i in range(10) for j in range(3) for k in range(7)
```

While for a np.array, also a reshaping (10, 3, 7) $\rightarrow$ ( $2,21,5$ ) would be allowed, it does not make sense physically. The only sensible "reshape" operation on an Array are

1) to combine multiple legs into one leg pipe (LegPipe) with combine_legs (), or
2) to split a pipe of previously combined legs with split_legs ().

Each leg has a Hilbert space, and a representation of the symmetry on that Hilbert space. Combining legs corresponds to the tensor product operation, and for abelian groups, the corresponding "fusion" of the representation is the simple addition of charge.

Fusion is not a lossless process, so if we ever want to split the combined leg, we need some additional data to tell us how to reverse the tensor product. This data is saved in the class LegPipe, derived from the LegCharge and used as new leg. Details of the information contained in a LegPipe are given in the class doc string.
The rough usage idea is as follows:

1) You can call combine_legs () without supplying any LegPipes, combine_legs will then make them for you.

Nevertheless, if you plan to perform the combination over and over again on sets of legs you know to be identical [with same charges etc, up to an overall -1 in qconj on all incoming and outgoing Legs] you might make a LegPipe anyway to save on the overhead of computing it each time.
2) In any way, the resulting Array will have a LegPipe as a LegCharge on the combined leg. Thus, it - and all tensors inheriting the leg (e.g. the results of svd, tensordot etc.) - will have the information how to split the LegPipe back to the original legs.
3) Once you performed the necessary operations, you can call split_legs (). This uses the information saved in the LegPipe to split the legs, recovering the original legs.
For a LegPipe, conj () changes qconj for the outgoing pipe and the incoming legs. If you need a LegPipe with the same incoming qconj, use outer_conj().

### 10.2.3 Leg labeling

It's convenient to name the legs of a tensor: for instance, we can name legs $0,1,2$ to be 'a', 'b', 'c': $T_{i_{a}, i_{b}, i_{c}}$. That way we don't have to remember the ordering! Under tensordot, we can then call
$\mathrm{U}=$ npc.tensordot $(\mathrm{S}, \mathrm{T}$, axes $=$ [ [...], ['b'] ] )
without having to remember where exactly ' $b$ ' is. Obviously $U$ should then inherit the name of its legs from the uncontracted legs of $S$ and $T$. So here is how it works:

- Labels can only be strings. The labels should not include the characters . or ?. Internally, the labels are stored as dict a.labels $=\{$ label: leg_position, ... \}. Not all legs need a label.
- To set the labels, call
A.set_labels(['a', 'b', None, 'c', ... ])
which will set up the labeling \{'a': 0, 'b': 1, 'c': $3 \ldots\}$.
- (Where implemented) the specification of axes can use either the labels or the index positions. For instance, the call tensordot (A, B, [['a', 2, 'c'], [...]]) will interpret 'a' and 'c' as labels (calling get_leg_indices () to find their positions using the dict) and 2 as 'the 2 nd leg'. That's why we require labels to be strings!
- Labels will be intelligently inherited through the various operations of np_conserved.
- Under transpose, labels are permuted.
- Under tensordot, labels are inherited from uncontracted legs. If there is a collision, both labels are dropped.
- Under combine_legs, labels get concatenated with a . delimiter and sourrounded by brackets. Example: let a.labels = \{'a': 1, 'b': 2, 'c': 3\}. Then if b = a. combine_legs([[0, 1], [2]]), it will have b.labels = \{'(a.b)': 0, '(c)': 1 \}. If some sub-leg of a combined leg isn't named, then a '? \#' label is inserted (with \# the leg index), e.g., 'a.?0.c'.
- Under split_legs, the labels are split using the delimiters (and the '?\#' are dropped).
_ Under conj, iconj: take 'a' -> 'a*', 'a*' -> 'a', and'(a.(b*.c))' -> '(a*. (b. C*) ) '
- Under $s v d$, the outer labels are inherited, and inner labels can be optionally passed.
- Under pinv, the labels are transposed.


### 10.2.4 Indexing of an Array

Although it is usually not necessary to access single entries of an Array, you can of course do that. In the simplest case, this is something like $A[0,2,1]$ for a rank-3 Array A. However, accessing single entries is quite slow and usually not recommended. For small Arrays, it may be convenient to convert them back to flat numpy arrays with to_ndarray().

On top of that very basic indexing, Array supports slicing and some kind of advanced indexing, which is however different from the one of numpy arrarys (described here). Unlike numpy arrays, our Array class does not broadcast existing index arrays - this would be terribly slow. Also, np.newaxis is not supported, since inserting new axes requires additional information for the charges.

Instead, we allow just indexing of the legs independent of each other, of the form $\mathrm{A}[\mathrm{i} 0, \mathrm{i} 1, \ldots \mathrm{l}$. If all indices i0, i1, ... are integers, the single corresponding entry (of type dtype) is returned.

However, the individual 'indices' io for the individual legs can also be one of what is described in the following list. In that case, a new Array with less data (specified by the indices) is returned.

The 'indices' can be:

- an int: fix the index of that axis, return array with one less dimension. See also take_slice ().
- a slice (None) or :: keep the complete axis
- an Ellipsis or . . . : shorthand for slice (None) for missing axes to fix the len
- an 1D bool ndarray mask: apply a mask to that axis, see iproject ().
- a slice (start, stop, step) or start:stop: step: keep only the indices specified by the slice. This is also implemented with iproject.
- an 1D int ndarray mask: keep only the indices specified by the array. This is also implemented with iproject.

For slices and 1D arrays, additional permuations may be perfomed with the help of permute ().

If the number of indices is less than rank, the remaining axes remain free, so for a rank 4 Array A, A[i0, i1] == A[i0, i1, ...] == A[i0, i1, :, :].
Note that indexing always copies the data - even if int contains just slices, in which case numpy would return a view. However, assigning with $A[:,[3,5], 3]=B$ should work as you would expect.

Warning: Due to numpy's advanced indexing, for 1D integer arrays a 0 and a1 the following holds
A[a0, al].to_ndarray() ==A.to_ndarray()[np.ix_(a0, a1)] ! = A.to_ndarray()[a0, a1]
For a combination of slices and arrays, things get more complicated with numpys advanced indexing. In that case, a simple np.ix_(...) doesn't help any more to emulate our version of indexing.

### 10.2.5 Details of the np_conserved implementation

## Notations

Lets fix the notation of certain terms for this introduction and the doc-strings in np_conserved. This might be helpful if you know the basics from a different context. If you're new to the subject, keep reading even if you don't understand each detail, and come back to this section when you encounter the corresponding terms again.

A Array is a multi-dimensional array representing a tensor with the entries:

$$
T_{a_{0}, a_{1}, \ldots a_{\text {rank }-1}} \quad \text { with } \quad a_{i} \in\left\{0, \ldots, n_{i}-1\right\}
$$

Each leg $a_{i}$ corresponds the a vector space of dimension $n \_i$.
An index of a leg is a particular value $a_{i} \in\left\{0, \ldots, n_{i}-1\right\}$.
The rank is the number of legs, the shape is $\left(n_{0}, \ldots, n_{\text {rank-1 }}\right)$.
We restrict ourselfes to abelian charges with entries in $\mathbb{Z}$ or in $\mathbb{Z}_{m}$. The nature of a charge is specified by $m$; we set $m=1$ for charges corresponding to $\mathbb{Z}$. The number of charges is refered to as qnumber as a short hand, and the collection of $m$ for each charge is called qmod. The qnumber, qmod and possibly descriptive names of the charges are saved in an instance of ChargeInfo.

To each index of each leg, a value of the charge(s) is associated. A charge block is a contiguous slice corresponding to the same charge(s) of the leg. A qindex is an index in the list of charge blocks for a certain leg. A charge sector is for given charge(s) is the set of all qindices of that charge(s). A leg is blocked if all charge sectors map one-to-one to qindices. Finally, a leg is sorted, if the charges are sorted lexiographically. Note that a sorted leg is always blocked. We can also speak of the complete array to be blocked by charges or legcharge-sorted, which means that all of its legs are blocked or sorted, respectively. The charge data for a single leg is collected in the class LegCharge. A LegCharge has also a flag qconj, which tells whether the charges point inward ( +1 ) or outward ( -1 ). What that means, is explained later in Which entries of the npc Array can be non-zero?.

For completeness, let us also summarize also the internal structure of an Array here: The array saves only non-zero blocks, collected as a list of np.array in self._data. The qindices necessary to map these blocks to the original leg indices are collected in self._qdata An array is said to be qdata-sorted if its self._qdata is lexiographically sorted. More details on this follow later. However, note that you usually shouldn't access _qdata and _data directly this is only necessary from within tensordot, $s v d$, etc. Also, an array has a total charge, defining which entries can be non-zero - details in Which entries of the npc Array can be non-zero?.

Finally, a leg pipe (implemented in LegPipe) is used to formally combine multiple legs into one leg. Again, more details follow later.

## Physical Example

For concreteness, you can think of the Hamiltonian $H=-t \sum_{<i, j>}\left(c_{i}^{\dagger} c_{j}+H . c.\right)+U n_{i} n_{j}$ with $n_{i}=c_{i}^{\dagger} c_{i}$. This Hamiltonian has the global $U(1)$ gauge symmetry $c_{i} \rightarrow c_{i} e^{i \phi}$. The corresponding charge is the total number of particles $N=\sum_{i} n_{i}$. You would then introduce one charge with $m=1$.

Note that the total charge is a sum of local terms, living on single sites. Thus, you can infer the charge of a single physical site: it's just the value $q_{i}=n_{i} \in \mathbb{N}$ for each of the states.

Note that you can only assign integer charges. Consider for example the spin $1 / 2$ Heisenberg chain. Here, you can naturally identify the magnetization $S^{z}=\sum_{i} S_{i}^{z}$ as the conserved quantity, with values $S_{i}^{z}= \pm \frac{1}{2}$. Obviously, if $S^{z}$ is conserved, then so is $2 S^{z}$, so you can use the charges $q_{i}=2 S_{i}^{z} \in\{-1,+1\}$ for the down and $u p$ states, respectively. Alternatively, you can also use a shift and define $q_{i}=S_{i}^{z}+\frac{1}{2} \in\{0,1\}$.
As another example, consider BCS like terms $\sum_{k}\left(c_{k}^{\dagger} c_{-k}^{\dagger}+H . c.\right)$. These terms break the total particle conservation, but they preserve the total parity, i.e., $N \bmod 2$ is conserved. Thus, you would introduce a charge with $m=2$ in this case.

In the above examples, we had only a single charge conserved at a time, but you might be lucky and have multiple conserved quantities, e.g. if you have two chains coupled only by interactions. TeNPy is designed to handle the general case of multiple charges. When giving examples, we will restrict to one charge, but everything generalizes to multiple charges.

## The different formats for LegCharge

As mentioned above, we assign charges to each index of each leg of a tensor. This can be done in three formats: qflat, as qind and as qdict. Let me explain them with examples, for simplicity considereing only a single charge (the most inner array has one entry for each charge).
qflat form: simply a list of charges for each index. An example:
qflat $=[[-2],[-1],[-1],[0],[0],[0],[0],[3],[3]]$
This tells you that the leg has size 9, the charges for are [-2], [-1], [-1], ..., [3]((%5Cmathrm%7BL%7D=100)) for the indices 0, $1,2,3, \ldots, 8$. You can identify four charge $\operatorname{blocks}$ slice (0, 1), slice(1, 3), slice (3, 7), slice (7, 9) in this example, which have charges [-2], [-1], [0], [3]((%5Cmathrm%7BL%7D=100)). In other words, the indices 1,2 (which are in slice ( 1,3 ) ) have the same charge value [-1]. A qindex would just enumerate these blocks as $0,1,2,3$.
qind form: a 1D array slices and a 2D array charges. This is a more compact version than the qfat form: the slices give a partition of the indices and the charges give the charge values. The same example as above would simply be:

```
slices = [0, 1, 3, 7, 9]
charges = [[-2], [-1], [0], [3]]
```

Note that slices includes 0 as first entry and the number of indices (here 9) as last entries. Thus it has len block_number + 1, where block_number (given by block_number) is the number of charge blocks in the leg, i.e. a qindex runs from 0 to block_number -1 . On the other hand, the 2D array charges has shape (block_number, qnumber), where qnumber is the number of charges (given by quimber).

In that way, the qind form maps an qindex, say qi, to the indices slice (slices [qi], slices [qi+1]) and the charge(s) charges [qi].
qdict form: a dictionary in the other direction than qind, taking charge tuples to slices. Again for the same example:

```
{(-2,): slice(0, 1),
    (-1,): slice(1, 3),
    (0,) : slice(3, 7),
    (3,) : slice(7, 9)}
```

Since the keys of a dictionary are unique, this form is only possible if the leg is completely blocked.
The LegCharge saves the charge data of a leg internally in qind form, directly in the attribute slices and charges. However, it also provides convenient functions for conversion between from and to the qflat and qdict form.
The above example was nice since all charges were sorted and the charge blocks were 'as large as possible'. This is however not required.

The following example is also a valid qind form:

```
slices = [0, 1, 3, 5, 7, 9]
charges = [[-2], [-1], [0], [0], [3]]
```

This leads to the same qflat form as the above examples, thus representing the same charges on the leg indices. However, regarding our Arrays, this is quite different, since it diveds the leg into 5 (instead of previously 4) charge blocks. We say the latter example is not bunched, while the former one is bunched.

To make the different notions of sorted and bunched clearer, consider the following (valid) examples:

| charges | bunched | sorted | blocked |
| :--- | :--- | :--- | :--- | :--- |
| $[[-2],[-1],[0],[1],[3]]$ | True | True | True |
| $[[-2],[-1],[0],[0],[3]]$ | False | True | False |
| $[[-2],[0],[-1],[1],[3]]$ | True | False | True |
| $[[-2],[0],[-1],[0],[3]]$ | True | False | False |

If a leg is bunched and sorted, it is automatically blocked (but not vice versa). See also below for further comments on that.

## Which entries of the npc Array can be non-zero?

The reason for the speedup with np_conserved lies in the fact that it saves only the blocks 'compatible' with the charges. But how is this 'compatible' defined?

Assume you have a tensor, call it $T$, and the LegCharge for all of its legs, say $a, b, c, \ldots$.
Remeber that the LegCharge associates to each index of the leg a charge value (for each of the charges, if qnumber $>$ 1). Let a.to_qflat () [ia] denote the charge(s) of index ia for leg a, and similar for other legs.

In addition, the LegCharge has a flag qcon $j$. This flag qconj is only a sign, saved as +1 or -1 , specifying whether the charges point 'inward' $(+1$, default) or 'outward' $(-1)$ of the tensor.

Then, the total charge of an entry $\mathrm{T}[\mathrm{ia}, \mathrm{ib}, \mathrm{ic}, \ldots .$.$] of the tensor is defined as:$

```
qtotal[ia, ib, ic, ...] = a.to_qflat()[ia] * a.qconj + b.to_qflat()[ib] * b.qconj + c.
\hookrightarrowto_qflat()[ic] * c.qconj + ... modulo qmod
```

The rule which entries of the a Array can be non-zero (i.e., are 'compatible' with the charges), is then very simple:

## Rule for non-zero entries

An entry ia, ib, ic, ... of a Array can only be non-zero, if qtotal[ia, ib, ic, ...] matches the unique qtotal attribute of the class.

In other words, there is a single total charge . qtotal attribute of array. All indices ia, ib, ic, ... for which the above defined qtotal[ia, ib, ic, ...] matches this total charge, are said to be compatible with the charges and can be non-zero. All other indices are incompatible with the charges and must be zero.

In case of multiple charges, qnumber $>1$, is a straigth-forward generalization: an entry can only be non-zero if it is compatible with each of the defined charges.

## The pesky qconj - contraction as an example

Why did we introduce the qcon $j$ flag? Remember it's just a sign telling whether the charge points inward or outward. So whats the reasoning?

The short answer is, that LegCharges actually live on bonds (i.e., legs which are to be contracted) rather than individual tensors. Thus, it is convenient to share the LegCharges between different legs and even tensors, and just adjust the sign of the charges with qconj.

As an example, consider the contraction of two tensors, $C_{i a, i c}=\sum_{i b} A_{i a, i b} B_{i b, i c}$. For simplicity, say that the total charge of all three tensors is zero. What are the implications of the above rule for non-zero entries? Or rather, how can we ensure that $C$ complies with the above rule? An entry $C[i a, i c]$ will only be non-zero, if there is an ib such that both $A[i a, i b]$ and $B[i b, i c]$ are non-zero, i.e., both of the following equations are fullfilled:

```
A.qtotal == A.legs[0].to_qflat()[ia] * A.legs[0].qconj + A.legs[1].to_qflat()[ib] * A.
\hookrightarrowegs[1].qconj modulo qmod
B.qtotal == B.legs[0].to_qflat()[ib] * B.legs[0].qconj + B.legs[1].to_qflat()[ic] * B.
\hookrightarrowlegs[1].qconj modulo qmod
```

(A. legs [0] is the LegCharge saving the charges of the first leg (with index ia) of A.)

For the uncontracted legs, we just keep the charges as they are:

```
C.legs = [A.legs[0], B.legs[1]]
```

It is then straight-forward to check, that the rule is fullfilled for $C$, if the following condition is met:

```
A.qtotal + B.qtotal - C.qtotal == A.legs[1].to_qflat()[ib] A.b.qconj + B.legs[0].to_
\hookrightarrowqflat()[ib] B.b.qconj modulo qmod
```

The easiest way to meet this condition is (1) to require that A.b and B.b share the same charges b.to_qflat (), but have opposite qconj, and (2) to define C.qtotal $=\mathrm{A} . q \operatorname{total}+\mathrm{B}$. qtotal. This justifies the introduction of qconj: when you define the tensors, you have to define the LegCharge for the $b$ only once, say for A. legs [1]. For B.legs [0] you simply use A. legs [1].conj() which creates a copy of the LegCharge with shared slices and charges, but opposite qconj. As a more impressive example, all 'physical' legs of an MPS can usually share the same LegCharge (up to different qconj if the local Hilbert space is the same). This leads to the following convention:

## Convention

When an npc algorithm makes tensors which share a bond (either with the input tensors, as for tensordot, or amongst the output tensors, as for SVD), the algorithm is free, but not required, to use the same LegCharge for the tensors
sharing the bond, without making a copy. Thus, if you want to modify a LegCharge, you must make a copy first (e.g. by using methods of LegCharge for what you want to acchive).

## Assigning charges to non-physical legs

From the above physical examples, it should be clear how you assign charges to physical legs. But what about other legs, e.g, the virtual bond of an MPS (or an MPO)?

The charge of these bonds must be derived by using the 'rule for non-zero entries', as far as they are not arbitrary. As a concrete example, consider an MPS on just two spin $1 / 2$ sites:
$\square$
The two legs $p$ are the physical legs and share the same charge, as they both describe the same local Hilbert space. For better distincition, let me label the indices of them by $\uparrow=0$ and $\downarrow=1$. As noted above, we can associate the charges 1 ( $p=\uparrow$ ) and -1 $(p=\downarrow)$, respectively, so we define:

```
chinfo = npc.ChargeInfo([1], ['2*Sz'])
p = npc.LegCharge.from_qflat(chinfo, [1, -1], qconj=+1)
```

For the qcon j signs, we stick to the convention used in our MPS code and indicated by the arrows in above 'picture': physical legs are incoming ( $q c o n j=+1$ ), and from left to right on the virtual bonds. This is acchieved by using [ $p$, $\mathrm{x}, \mathrm{y} . \operatorname{conj()]}$ as legs for $A$, and $[p, y, z \cdot \operatorname{conj}()]$ for $B$, with the default $q c o n j=+1$ for all $p, x, y$, $z: y \cdot c o n j()$ has the same charges as $y$, but opposite qcon $j=-1$.

The legs $x$ and $z$ of an $L=2$ MPS, are 'dummy' legs with just one index 0 . The charge on one of them, as well as the total charge of both $A$ and $B$ is arbitrary (i.e., a gauge freedom), so we make a simple choice: total charge 0 on both arrays, as well as for $x=0, \mathrm{x}=$ npc.LegCharge.from_qflat (chinfo, [0], qconj=+1).

The charges on the bonds $y$ and $z$ then depend on the state the MPS represents. Here, we consider a singlet $\psi=(\mid \uparrow \downarrow$ $\rangle-|\downarrow \uparrow\rangle) / \sqrt{2}$ as a simple example. A possible MPS representation is given by:

```
A[up, :, :] = [[1/2.**0.5, 0]] B[up, :, :] = [[0], [-1]]
A[down, :, :] = [[0, 1/2.**0.5]] B[down, :, :] = [[1], [0]]
```

There are two non-zero entries in $A$, for the indices $(a, x, y)=(\uparrow, 0,0)$ and $(\downarrow, 0,1)$. For $(a, x, y)=(\uparrow, 0,0)$, we want:

```
A.qtotal = 0 = p.to_qflat()[up] * p.qconj + x.to_qflat()[0] * x.qconj + y.conj().to_
\hookrightarrowqflat()[0] * y.conj().qconj
    =1 * (+1) + 0 * (+1) + y.conj().to_
\hookrightarrowqflat()[0] * (-1)
```

This fixes the charge of $\mathrm{y}=0$ to 1 . A similar calculation for $(a, x, y)=(\downarrow, 0,1)$ yields the charge -1 for $\mathrm{y}=1$. We have thus all the charges of the leg $y$ and can define $y=n p c . L e g C h a r g e . f r o m \_q f l a t(c h i n f o, ~[1, ~-1]$, $q c o n j=+1$ ).

Now take a look at the entries of B. For the non-zero entry $(b, y, z)=(\uparrow, 1,0)$, we want:

```
B.qtotal = 0 = p.to_qflat()[up] * p.qconj + y.to_qflat()[1] * y.qconj + z.conj().to_
->qflat()[0] * z.conj().qconj
    = 1 * (+1) + (-1) * (+1) + z.conj().to_
```

$\rightarrow$ qflat () [0] * (-1)

This implies the charge 0 for $z=0$, thus $z=n p c$. LegCharge.form_qflat (chinfo, [0], qconj=+1). Finally, note that the rule for $(b, y, z)=(\downarrow, 0,0)$ is automatically fullfilled! This is an implication of the fact that the singlet has a well defined value for $S_{a}^{z}+S_{b}^{z}$. For other states without fixed magnetization (e.g., $|\uparrow \uparrow\rangle+|\downarrow \downarrow\rangle$ ) this would not be the case, and we could not use charge conservation.

As an exercise, you can calculate the charge of $z$ in the case that A. qtotal=5, B. qtotal $=-1$ and charge 2 for $x=0$. The result is -2 .

Note: This section is meant be an pedagogical introduction. In you program, you can use the functions detect_legcharge () (which does exactly what's described above) or detect_qtotal () (if you know all LegCharges, but not qtotal).

## Array creation

## Direct creation

Making an new Array requires both the tensor entries (data) and charge data.
The default initialization $a=$ Array (...) creates an empty Array, where all entries are zero (equivalent to zeros ()). (Non-zero) data can be provided either as a dense np.array to from_ndarray (), or by providing a numpy function such as np.random, np.ones etc. to from_func ().

In both cases, the charge data is provided by one ChargeInfo, and a LegCharge instance for each of the legs.

Note: The charge data instances are not copied, in order to allow it to be shared between different Arrays. Consequently, you must make copies of the charge data, if you manipulate it directly. (However, methods like sort () do that for you.)

## Indirect creation by manipulating existing arrays

Of course, a new Array can also created using the charge data from exisiting Arrays, for example with zeros_like () or creating a (deep or shallow) copy (). Further, there are many higher level functions like tensordot () or svd(), which also return new Arrays.

## Complete blocking of Charges

While the code was designed in such a way that each charge sector has a different charge, the code should still run correctly if multiple charge sectors (for different qindex) correspond to the same charge. In this sense Array can act like a sparse array class to selectively store subblocks. Algorithms which need a full blocking should state that explicitly in their doc-strings. (Some functions (like svd and eigh) require complete blocking internally, but if necessary they just work on a temporary copy returned by as_completely_blocked () ).

If you expect the tensor to be dense subject to charge constraints (as for MPS), it will be most efficient to fully block by charge, so that work is done on large chunks.

However, if you expect the tensor to be sparser than required by charge (as for an MPO), it may be convenient not to completely block, which forces smaller matrices to be stored, and hence many zeroes to be dropped. Nevertheless,
the algorithms were not designed with this in mind, so it is not recommended in general. (If you want to use it, run a benchmark to check whether it is really faster!)

If you haven't created the array yet, you can call sort () (with bunch=True) on each LegCharge which you want to block. This sorts by charges and thus induces a permution of the indices, which is also returned as an 1D array perm. For consistency, you have to apply this permutation to your flat data as well.

Alternatively, you can simply call sort_legcharge () on an existing Array. It calls sort () internally on the specified legs and performs the necessary permutations directly to (a copy of) self. Yet, you should keep in mind, that the axes are permuted afterwards.

## Internal Storage schema of npc Arrays

The actual data of the tensor is stored in _data. Rather than keeping a single np.array (which would have many zeros in it), we store only the non-zero sub blocks. So _data is a python list of np.array's. The order in which they are stored in the list is not physically meaningful, and so not guaranteed (more on this later). So to figure out where the sub block sits in the tensor, we need the _qdat a structure (on top of the LegCharges in legs).

Consider a rank 3 tensor T , with the first leg like:

```
legs[0].slices = np.array([0, 1, 4, ...])
legs[0].charges = np.array([[-2], [1], ...])
```

Each row of charges gives the charges for a charge block of the leg, with the actual indices of the total tensor determined by the slices. The qindex simply enumerates the charge blocks of a lex. Picking a qindex (and thus a charge block) from each leg, we have a subblock of the tensor.

For each (non-zero) subblock of the tensor, we put a (numpy) ndarray entry in the _data list. Since each subblock of the tensor is specified by rank qindices, we put a corresponding entry in _qdata, which is a 2D array of shape (\#stored_blocks, rank). Each row corresponds to a non-zero subblock, and there are rank columns giving the corresponding qindex for each leg.

Example: for a rank 3 tensor we might have:

```
T._data = [t1, t2, t3, t4, ...]
T._qdata = np.array([[3, 2, 1],
    [1, 1, 1],
    [4, 2, 2],
    [2, 1, 2],
    ... ])
```

The third subblock has an ndarray $t 3$, and qindices [ $\left.\begin{array}{lll}4 & 2 & 2\end{array}\right]$ for the three legs.

- To find the position of $t 3$ in the actual tensor you can use get_slice ():
T.legs[0].get_slice(4), T.legs[1].get_slice(2), T.legs[2].get_slice(2)

The function leg.get_charges(qi) simply returns slice(leg.slices[qi], leg. slices [qi+1])

- To find the charges of t 3 , we an use get_charge ():

```
T.legs[0].get_charge(2), T.legs[1].get_charge(2), T.legs[2].get_charge(2)
```

The function leg.get_charge (qi) simply returns leg.charges [qi]*leg.qconj.

Note: Outside of $n p$ _conserved, you should use the API to access the entries. If you really need to iterate over all blocks of an Array T, try for (block, blockslices, charges, qindices) in $T$ :
do_something().

The order in which the blocks stored in _datal_qdata is arbitrary (although of course _data and _qdata must be in correspondence). However, for many purposes it is useful to sort them according to some convention. So we include a flag ._qdata_sorted to the array. So, if sorted (with isort_qdata(), the _qdata example above goes to

```
_qdata = np.array([[1, 1, 1],
    [3, 2, 1],
    [2, 1, 2],
    [4, 2, 2],
    ... ])
```

Note that $n p$.lexsort chooses the right-most column to be the dominant key, a convention we follow throughout.
If _qdata_sorted == True, _qdata and _data are guaranteed to be lexsorted. If _qdata_sorted == False, there is no gaurantee. If an algorithm modifies _qdata, it must set_qdata_sorted = False (unless it gaurantees it is still sorted). The routine sort_qdata () brings the data to sorted form.

### 10.2.6 See also

- The module tenpy. Iinalg.np_conserved should contain all the API needed from the point of view of the algorithms. It contians the fundamental Array class and functions for working with them (creating and manipulating).
- The module tenpy. Iinalg. charges contains implementations for the charge structure, for example the classes ChargeInfo, LegCharge, and LegPipe. As noted above, the 'public' API is imported to (and accessible from) np_conserved.


### 10.2.7 A full example code for spin-1/2

Below follows a full example demonstrating the creation and contraction of Arrays. (It's the file $a \_n p \_c o n s e r v e d . p y ~$ in the examples folder of the tenpy source.)

```
"""An example code to demonstrate the usage of :class:`~tenpy.linalg.np_conserved.
\hookrightarrowArray`.
This example includes the following steps:
1) create Arrays for an Neel MPS
2) create an MPO representing the nearest-neighbour AFM Heisenberg Hamiltonian
3) define 'environments' left and right
4) contract MPS and MPO to calculate the energy
5) extract two-site hamiltonian ``H2`` from the MPO
6) calculate '`exp(-1.j*dt*H2)`` by diagonalization of H2
7) apply ``exp(H2)`` to two sites of the MPS and truncate with svd
Note that this example uses only np_conserved, but no other modules.
Compare it to the example `b_mps.py`,
which does the same steps using a few predefined classes like MPS and MPO.
"""
# Copyright 2018-2021 TeNPy Developers, GNU GPLv3
import tenpy.linalg.np_conserved as npc
import numpy as np
```

(continues on next page)

```
# model parameters
Jxx, Jz = 1., 1.
L = 20
dt = 0.1
cutoff = 1.e-10
print("Jxx={Jxx}, Jz={Jz}, L={L:d}".format(Jxx=Jxx, Jz=Jz, L=L))
print("1) create Arrays for an Neel MPS")
    vL ->--B-->- vR
# l
# ^
# p
# create a ChargeInfo to specify the nature of the charge
chinfo = npc.ChargeInfo([1], ['2*Sz']) # the second argument is just a descriptive
->name
# create LegCharges on physical leg and even/odd bonds
p_leg = npc.LegCharge.from_qflat(chinfo, [[1], [-1]]) # charges for up, down
v_leg_even = npc.LegCharge.from_qflat(chinfo, [[0]])
v_leg_odd = npc.LegCharge.from_qflat(chinfo, [[1]])
B_even = npc.zeros([v_leg_even, v_leg_odd.conj(), p_leg],
    labels=['vL', 'vR', 'p']) # virtual left/right, physical
B_odd = npc.zeros([v_leg_odd, v_leg_even.conj(), p_leg], labels=['vL', 'vR', 'p'])
B_even[0, 0, 0] = 1. # up
B_odd[0, 0, 1] = 1. # down
Bs = [B_even, B_odd] * (L // 2) + [B_even] * (L % 2) # (right-canonical)
Ss = [np.ones(1)] * L # Ss[i] are singular values between Bs[i-1] and BS[i]
# Side remark:
# An MPS is expected to have non-zero entries everywhere compatible with the charges.
# In general, we recommend to use `sort_legcharge` (or `as_completely_blocked`)
# to ensure complete blocking. (But the code will also work, if you don't do it.)
# The drawback is that this might introduce permutations in the indices of single
->legs,
# which you have to keep in mind when converting dense numpy arrays to and from npc.
\rightarrow A r r a y s .
print("2) create an MPO representing the AFM Heisenberg Hamiltonian")
## nc
# create physical spin-1/2 operators Sz, S+, S-
Sz = npc.Array.from_ndarray([[0.5, 0.], [0., -0.5]], [p_leg, p_leg.conj()], labels=['p
```


(continued from previous page)

```
Sp = npc.Array.from_ndarray([[0., 1.], [0., 0.]], [p_leg, p_leg.conj()], labels=['p',
G'p*'])
Sm = npc.Array.from_ndarray([[0., 0.], [1., 0.]], [p_leg, p_leg.conj()], labels=['p',
G'p*'])
Id = npc.eye_like(Sz, labels=Sz.get_leg_labels()) # identity
mpo_leg = npc.LegCharge.from_qflat(chinfo, [[0], [2], [-2], [0], [0]])
W_grid = [[Id, Sp, Sm, Sz, None ],
    [None, None, None, None, 0.5 * Jxx * Sm],
    [None, None, None, None, 0.5 * Jxx * Sp],
    [None, None, None, None, Jz * Sz ],
    [None, None, None, None, Id ]] # yapf:disable
W = npc.grid_outer(W_grid, [mpo_leg, mpo_leg.conj()], grid_labels=['wL', 'wR'])
# wL/wR = virtual left/right of the MPO
Ws = [W] * L
print("3) define 'environments' left and right")
# .---->- vR vL ->-----
# l l
# envL->- wR wL ->-envR
# l l
# .---->- vR* vL*->-----
envL = npc.zeros([W.get_leg('wL').conj(), Bs[0].get_leg('vL').conj(), Bs[0].get_leg(
↔'vL')],
    labels=['wR', 'vR', 'vR*'])
envL[0, :, :] = npc.diag(1., envL.legs[1])
envR = npc.zeros([W.get_leg('wR').conj(), Bs[-1].get_leg('vR').conj(), Bs[-1].get_leg(
\hookrightarrow'vR')],
    labels=['wL', 'vL', 'vL*'])
envR[-1, :, :] = npc.diag(1., envR.legs[1])
print("4) contract MPS and MPO to calculate the energy <psi|H|psi>")
contr = envL
for i in range(L):
    # contr labels: wR, vR, vR*
    contr = npc.tensordot(contr, Bs[i], axes=('vR', 'vL'))
    # wR, vR*, vR, p
    contr = npc.tensordot(contr, Ws[i], axes=(['p', 'wR'], ['p*', 'wL']))
    # vR*, vR, wR, p
    contr = npc.tensordot(contr, Bs[i].conj(), axes=(['p', 'vR*'], ['p*', 'vL*']))
    # vR, wR, vR*
    # note that the order of the legs changed, but that's no problem with labels:
    # the arrays are automatically transposed as necessary
E = npc.inner(contr, envR, axes=(['vR', 'wR', 'vR*'], ['vL', 'wL', 'vL*']))
print("E =", E)
print("5) calculate two-site hamiltonian ``H2`` from the MPO")
# label left, right physical legs with p, q
W0 = W.replace_labels(['p', 'p*'], ['p0', 'p0*'])
W1 = W.replace_labels(['p', 'p*'], ['p1', 'p1*'])
H2 = npc.tensordot(w0, W1, axes=('wR', 'wL')).itranspose(['wL', 'wR', 'p0', 'p1', 'p0*
\hookrightarrow', 'p1*'])
H2 = H2[0, -1] # (If H has single-site terms, it's not that simple anymore)
```

(continues on next page)

```
print("H2 labels:", H2.get_leg_labels())
print("6) calculate exp(H2) by diagonalization of H2")
# diagonalization requires to view H2 as a matrix
H2 = H2.combine_legs([('p0', 'p1'), ('p0*', 'p1*')], qconj=[+1, -1])
print("labels after combine_legs:", H2.get_leg_labels())
E2, U2 = npc.eigh(H2)
print("Eigenvalues of H2:", E2)
U_expE2 = U2.scale_axis(np.exp(-1.j * dt * E2), axis=1) # scale_axis ~= apply an_
\hookrightarrowdiagonal matrix
exp_H2 = npc.tensordot(U_expE2, U2.conj(), axes=(1, 1))
exp_H2.iset_leg_labels(H2.get_leg_labels())
exp_H2 = exp_H2.split_legs() # by default split all legs which are `LegPipe`
# (this restores the originial labels ['po', 'p1', 'p0*', 'p1*'] of `H2` in `exp_H2`)
print("7) apply exp(H2) to even/odd bonds of the MPS and truncate with svd")
# (this implements one time step of first order TEBD)
for even_odd in [0, 1]:
    for i in range(even_odd, L - 1, 2):
        B_L = Bs[i].scale_axis(Ss[i], 'vL').ireplace_label('p', 'p0')
        B_R = Bs[i + 1].replace_label('p', 'pl')
        theta = npc.tensordot(B_L, B_R, axes=('vR', 'vL'))
        theta = npc.tensordot(exp_H2, theta, axes=(['p0*', 'p1*'], ['p0', 'p1']))
        # view as matrix for SVD
        theta = theta.combine_legs([('vL', 'p0'), ('p1', 'vR')], new_axes=[0, 1], b
qconj=[+1, -1])
    # now theta has labels '(vL.p0)', '(pl.vR)'
    U, S, V = npc.svd(theta, inner_labels=['vR', 'vL'])
    # truncate
    keep = S > cutoff
    S = S[keep]
    invsq = np.linalg.norm(S)
    Ss[i + 1] = S / invsq
    U = U.iscale_axis(S / invsq, 'vR')
    Bs[i] = U.split_legs('(vL.p0)').iscale_axis(Ss[i]**(-1), 'vL').ireplace_label(
@'p0', 'p')
    Bs[i + 1] = V.split_legs('(pl.vR)').ireplace_label('pl', 'p')
print("finished")
```


### 10.3 Models

### 10.3.1 What is a model?

Abstractly, a model stands for some physical (quantum) system to be described. For tensor networks algorithms, the model is usually specified as a Hamiltonian written in terms of second quantization. For example, let us consider a spin-1/2 Heisenberg model described by the Hamiltonian

$$
H=J \sum_{i} S_{i}^{x} S_{i+1}^{x}+S_{i}^{y} S_{i+1}^{y}+S_{i}^{z} S_{i+1}^{z}
$$

Note that a few things are defined more or less implicitly.

- The local Hilbert space: it consists of Spin-1/2 degrees of freedom with the usual spin-1/2 operators $S^{x}, S^{y}, S^{z}$.
- The geometric (lattice) strucuture: above, we spoke of a 1D "chain".
- The boundary conditions: do we have open or periodic boundary conditions? The "chain" suggests open boundaries, which are in most cases preferable for MPS-based methods.
- The range of $i$ : How many sites do we consider (for a 2D system: in each direction)?

Obviously, these things need to be specified in TeNPy in one way or another, if we want to define a model.
Ultimately, our goal is to run some algorithm. However, different algorithm requires the model and Hamiltonian to be specified in different forms. We have one class for each such required form. For example dmrg requires an MPOMOdel, which contains the Hamiltonian written as an MPO. So a new model class suitable for DMRG should have this general structure:

```
class MyNewModel(MPOModel):
    def __init__(self, model_params):
        lattice = somehow_generate_lattice(model_params)
        H_MPO = somehow_generate_MPO(lattice, model_params)
        # initialize MPOModel
        MPOModel.__init___(self, lattice, H_MPO)
```

On the other hand, if we want to evolve a state with tebd we need a NearestNeighborModel, in which the Hamiltonian is written in terms of two-site bond-terms to allow a Suzuki-Trotter decomposition of the time-evolution operator:

```
class MyNewModel2(NearestNeighborModel):
    """General strucutre for a model suitable for TEBD."""
    def __init__(self, model_params):
        lattice = somehow_generate_lattice(model_params)
        H_bond = somehow_generate_H_bond(lattice, model_params)
        # initialize MPOModel
        NearestNeighborModel.__init__(self, lattice, H_bond)
```

Of course, the difficult part in these examples is to generate the $\mathrm{H} \_\mathrm{MPO}$ and $\mathrm{H} \_$bond in the required form. If you want to write it down by hand, you can of course do that. But it can be quite tedious to write every model multiple times, just because we need different representations of the same Hamiltonian. Luckily, there is a way out in TeNPy: the CouplingModel. Before we describe this class, let's discuss the background of the Site and Site class.

### 10.3.2 The Hilbert space

The local Hilbert space is represented by a Site (read its doc-string!). In particular, the Site contains the local LegCharge and hence the meaning of each basis state needs to be defined. Beside that, the site contains the local operators - those give the real meaning to the local basis. Having the local operators in the site is very convenient, because it makes them available by name for example when you want to calculate expectation values. The most common sites (e.g. for spins, spin-less or spin-full fermions, or bosons) are predefined in the module tenpy.networks. site, but if necessary you can easily extend them by adding further local operators or completely write your own subclasses of Site.

The full Hilbert space is a tensor product of the local Hilbert space on each site.

Note: The LegCharge of all involved sites need to have a common ChargeInfo in order to allow the contraction of tensors acting on the various sites. This can be ensured with the function set_common_charges ().

An example where set_common_charges () is needed would be a coupling of different types of sites, e.g., when a tight binding chain of fermions is coupled to some local spin degrees of freedom. Another use case of this function would be a model with a $\$ \mathrm{U}(1) \$$ symmetry involving only half the sites, say $\sum_{i=0}^{L / 2} n_{2 i}$.

Note: If you don't know about the charges and np_conserved yet, but want to get started with models right away, you can set conserve=None in the existing sites or use leg = tenpy.linalg.np_conserved.LegCharge. from_trivial (d) for an implementation of your custom site, where $d$ is the dimension of the local Hilbert space. Alternatively, you can find some introduction to the charges in the Charge conservation with np_conserved.

### 10.3.3 The geometry : lattice class

The geometry is usually given by some kind of lattice structure how the sites are arranged, e.g. implicitly with the sum over nearest neighbours $\sum_{\langle i, j>}$. In TeNPy, this is specified by a Lattice class, which contains a unit cell of a few Site which are shifted periodically by its basis vectors to form a regular lattice. Again, we have pre-defined some basic lattices like a Chain, two chains coupled as a Ladder or 2D lattices like the Square, Honeycomb and Kagome lattices; but you are also free to define your own generalizations.
MPS based algorithms like DMRG always work on purely 1D systems. Even if our model "lives" on a 2D lattice, these algorithms require to map it onto a 1D chain (probably at the cost of longer-range interactions). This mapping is also done by the lattice by defining the order (order) of the sites.

Note: Further details on the lattice geometry can be found in Details on the lattice geometry.

### 10.3.4 The CouplingModel: general structure

The CouplingModel provides a general, quite abstract way to specify a Hamiltonian of couplings on a given lattice. Once initialized, its methods add_onsite() and add_coupling() allow to add onsite and coupling terms repeated over the different unit cells of the lattice. In that way, it basically allows a straight-forward translation of the Hamiltonian given as a math forumla $H=\sum_{i} A_{i} B_{i+d x}+\ldots$ with onsite operators $A, B, \ldots$ into a model class.
The general structure for a new model based on the CouplingModel is then:

```
class MyNewModel3(CouplingModel,MPOModel,NearestNeighborModel):
    def __init__(self, ...):
        ... # follow the basic steps explained below
```

In the initialization method __init__(self, . . .) of this class you can then follow these basic steps:
0 . Read out the parameters.

1. Given the parameters, determine the charges to be conserved. Initialize the LegCharge of the local sites accordingly.
2. Define (additional) local operators needed.
3. Initialize the needed Site.

Note: Using pre-defined sites like the SpinHalfSite is recommended and can replace steps 1-3.
4. Initialize the lattice (or if you got the lattice as a parameter, set the sites in the unit cell).
5. Initialize the CouplingModel with CouplingModel.__init__(self, lat).
6. Use add_onsite() and add_coupling() to add all terms of the Hamiltonian. Here, the pairs of the lattice can come in handy, for example:

```
self.add_onsite(-np.asarray(h), 0, 'Sz')
for u1, u2, dx in self.lat.pairs['nearest_neighbors']:
    self.add_coupling(0.5*J, u1, 'Sp', u2, 'Sm', dx, plus_hc=True)
    self.add_coupling( J, u1, 'Sz', u2, 'Sz', dx)
```

Note: The method add_coupling() adds the coupling only in one direction, i.e. not switching $i$ and $j$ in a $\sum_{\langle i, j\rangle}$. If you have terms like $c_{i}^{\dagger} c_{j}$ or $S_{i}^{+} S_{j}^{-}$in your Hamiltonian, you need to add it in both directions to get a Hermitian Hamiltonian! The easiest way to do that is to use the plus_hc option of add_onsite () and add_coupling (), as we did for the $J / 2\left(S_{i}^{+} S_{j}^{-}+h . c.\right)$ terms of the Heisenberg model above. Alternatively, you can add the hermitian conjugate terms explicitly, see the examples in add_coupling () for more details.

Note that the strength arguments of these functions can be (numpy) arrays for site-dependent couplings. If you need to add or multipliy some parameters of the model for the strength of certain terms, it is recommended use np. asarray beforehand - in that way lists will also work fine.
7. Finally, if you derived from the MPOMOdel, you can call calc_H_MPO () to build the MPO and use it for the initialization as MPOModel.__init__(self, lat, self.calc_H_MPO()).
8. Similarly, if you derived from the NearestNeighborModel, you can call calc_H_bond () to initialze it as NearestNeighborModel.__init__(self, lat, self.calc_H_bond()). Calling self. calc_H_bond () will fail for models which are not nearest-neighbors (with respect to the MPS ordering), so you should only subclass the NearestNeighborModel if the lattice is a simple Chain.

Note: The method add_coupling() works only for terms involving operators on 2 sites. If you have couplings involving more than two sites, you can use the add_multi_coupling() instead. A prototypical example is the exactly solvable ToricCode.

The code of the module tenpy.models.xxz_chain is included below as an illustrative example how to implement a Model. The implementation of the XXZChain directly follows the steps outline above. The XXZChain2 implements the very same model, but based on the CouplingMPOModel explained in the next section.

```
"""Prototypical example of a 1D quantum model: the spin-1/2 XXZ chain.
The XXZ chain is contained in the more general :class:`~tenpy.models.spins.SpinChain`;
 the idea of
this module is more to serve as a pedagogical example for a model.
"""
# Copyright 2018-2021 TeNPy Developers, GNU GPLv3
import numpy as np
from .lattice import Site, Chain
from .model import CouplingModel, NearestNeighborModel, MPOModel, CouplingMPOModel
from ..linalg import np_conserved as npc
from ..tools.params import asConfig
from ..networks.site import SpinHalfSite # if you want to use the predefined site
__all__ = ['XXZChain', 'XXZChain2']
class XXZChain(CouplingModel, NearestNeighborModel, MPOModel):
    r"""Spin-1/2 XXZ chain with Sz conservation.
```

```
    The Hamiltonian reads:
    .. math ::
        H=\sum_i \mathtt{Jxx}/2 (S^{+}_i S^{-}_{i+1} + S^{-}_i S^{+}_{i+1})
            + \mathtt{Jz} S^z_i S^\mp@subsup{_}{-}{\prime}{i+1} \\
            - \sum_i \mathtt{hz} S^z_i
    All parameters are collected in a single dictionary `model_params`, which
    is turned into a :class:`~tenpy.tools.params.Config` object.
    Parameters
    model_params : :class:`~tenpy.tools.params.Config`
        Parameters for the model. See :cfg:config:`XXZChain` below.
    Options
    --------
    .. cfg:config :: xxzChain
        :include: CouplingMPOModel
        L : int
        Length of the chain.
    Jxx, Jz, hz : float | array
            Coupling as defined for the Hamiltonian above.
    bc_MPS : {'finite' / 'infinte'}
            MPS boundary conditions. Coupling boundary conditions are chosen_
->appropriately.
    """
    def __init__(self, model_params):
        # 0) read out/set default parameters
        model_params = asConfig(model_params, "XXZChain")
        L = model_params.get('L', 2)
        Jxx = model_params.get('Jxx', 1.)
        Jz = model_params.get('Jz', 1.)
        hz = model_params.get('hz', 0.)
        bc_MPS = model_params.get('bc_MPS', 'finite')
        # 1-3):
        USE_PREDEFINED_SITE = False
        if not USE_PREDEFINED_SITE:
        # I) charges of the physical leg. The only time that we actually define
Charges!
            leg = npc.LegCharge.from_qflat(npc.ChargeInfo([1], ['2*Sz']), [1, -1])
            # 2) onsite operators
            Sp = [[0., 1.], [0., 0.]]
            Sm = [[0., 0.], [1., 0.]]
            Sz = [[0.5, 0.], [0., -0.5]]
            # (Can't define Sx and Sy as onsite operators: they are incompatible with
Gz charges.)
            # 3) local physical site
            site = Site(leg, ['up', 'down'], Sp=Sp, Sm=Sm, Sz=Sz)
        else:
            # there is a site for spin-1/2 defined in TeNPy, so just we can just use
->it
        # replacing steps 1-3)
        site = SpinHalfSite(conserve='Sz')
    # 4) lattice
```

```
bc = 'periodic' if bc_MPS == 'infinite' else 'open'
lat = Chain(L, site, bc=bc, bc_MPS=bc_MPS)
# 5) initialize CouplingModel
CouplingModel.__init__(self, lat)
# 6) add terms of the Hamiltonian
# (u is always 0 as we have only one site in the unit cell)
self.add_onsite(-hz, 0, 'Sz')
self.add_coupling(Jxx * 0.5, 0, 'Sp', 0, 'Sm', 1, plus_hc=True)
# the `plus_hc=True` adds the h.c. term
# see also the examples tenpy.models.model.CouplingModel.add_coupling
self.add_coupling(Jz, 0, 'Sz', 0, 'Sz', 1)
# 7) initialize H_MPO
MPOModel.__init__(self, lat, self.calc_H_MPO())
# 8) initialize H_bond (the order of 7/8 doesn't matter)
NearestNeighborModel.__init__(self, lat, self.calc_H_bond())
class XXZChain2(CouplingMPOModel, NearestNeighborModel):
    """Another implementation of the Spin-1/2 XXZ chain with Sz conservation.
    This implementation takes the same parameters as the :class:`XXZChain`, but isu
@implemented
    based on the :class:`~tenpy.models.model.CouplingMPOModel`.
    Parameters
    ----------
    model_params : dict | :class:`~tenpy.tools.params.Config
        See :cfg:config:`XXZChain
    """
    default_lattice = "Chain"
    force_default_lattice = True
    def init_sites(self, model_params):
    return SpinHalfSite(conserve='Sz') # use predefined Site
    def init_terms(self, model_params):
        # read out parameters
        Jxx = model_params.get('Jxx', 1.)
        Jz = model_params.get('Jz', 1.)
        hz = model_params.get('hz', 0.)
        # add terms
        for u in range(len(self.lat.unit_cell)):
            self.add_onsite(-hz, u, 'Sz')
            for u1, u2, dx in self.lat.pairs['nearest_neighbors']:
                self.add_coupling(Jxx * 0.5, u1, 'Sp', u2, 'Sm', dx, plus_hc=True)
                self.add_coupling(Jz, u1, 'Sz', u2, 'Sz', dx)
```


### 10.3.5 The easiest way: the CouplingMPOModel

Since many of the basic steps above are always the same, we don't need to repeat them all the time. So we have yet another class helping to structure the initialization of models: the CouplingMPOModel. The general structure of this class is like this:

```
class CouplingMPOModel(CouplingModel,MPOModel):
    default_lattice = "Chain"
    "
    def ___init__(self, model_param):
        # ... follows the basic steps 1-8 using the methods
        lat = self.init_lattice(self, model_param) # for step 4
        # ...
        self.init_terms(self, model_param) # for step 6
            # ...
    def init_sites(self, model_param):
        # You should overwrite this in most cases to ensure
        # getting the site(s) and charge conservation you want
        site = SpinSite(...) # or FermionSite, BosonSite, ...
        return site # (or tuple of sites)
    def init_lattice(self, model_param):
        sites = self.init_sites(self, model_param) # for steps 1-3
        # and then read out the class attribute `default_lattice`,
        # initialize an arbitrary pre-defined lattice
        # using model_params['lattice']
        # and enure it's the default lattice if the class attribute
        # `force_default_lattice` is True.
    def init_terms(self, model_param):
        # does nothing.
        # You should overwrite this
```

The XXZChain2 included above illustrates, how it can be used. You need to implement steps 1-3) by overwriting the method init_sites () Step 4) is performed in the method init_lattice (), which initializes arbitrary 1D or 2D lattices; by default a simple 1D chain. If your model only works for specific lattices, you can overwrite this method in your own class. Step 6) should be done by overwriting the method init_terms (). Steps 5,7,8 and calls to the init_... methods for the other steps are done automatically if you just call the CouplingMPOModel. __init__(self, model_param).

The XXZChain and XXZChain2 work only with the Chain as lattice, since they are derived from the NearestNeighborModel. This allows to use them for TEBD in 1D (yeah!), but we can't get the MPO for DMRG on (for example) a Square lattice cylinder - although it's intuitively clear, what the Hamiltonian there should be: just put the nearest-neighbor coupling on each bond of the 2D lattice.

It's not possible to generalize a NearestNeighborModel to an arbitrary lattice where it's no longer nearest Neigbors in the MPS sense, but we can go the other way around: first write the model on an arbitrary 2D lattice and then restrict it to a 1D chain to make it a NearestNeighborModel.

Let me illustrate this with another standard example model: the transverse field Ising model, implemented in the module tenpy.models.tf_ising included below. The TFIModel works for arbitrary 1D or 2D lattices. The TFIChain is then taking the exact same model making a NearestNeighborModel, which only works for the 1D chain.
"""Prototypical example of a quantum model: the transverse field Ising model.
(continues on next page)

```
                                    (continued from previous page)
Like the :class:`~tenpy.models.xxz_chain.XXZChain`, the transverse field ising chain
:class:`TFIChain` is contained in the more general :class:`~tenpy.models.spins.
\hookrightarrowSpinChain`;
the idea is more to serve as a pedagogical example for a 'model'.
We choose the field along z to allow to conserve the parity, if desired.
"""
# Copyright 2018-2021 TeNPy Developers, GNU GPLv3
import numpy as np
from .model import CouplingMPOModel, NearestNeighborModel
from .lattice import Chain
from ..tools.params import asConfig
from ..networks.site import SpinHalfSite
__all__ = ['TFIModel', 'TFIChain']
class TFIModel(CouplingMPOModel):
    r"""Transverse field Ising model on a general lattice.
    The Hamiltonian reads:
    .. math ::
        H = - \sum_{\langle i,j\rangle, i < j} \mathtt{J} \sigma^x_i \sigma^x_{j}
            - \sum_{i} \mathtt{g} \sigma^z_i
    Here, :math:`\langle i,j \rangle, i< j` denotes nearest neighbor pairs, each pair,
\hookrightarrowappearing
    exactly once.
    All parameters are collected in a single dictionary `model_params`, which
    is turned into a :class:`~tenpy.tools.params.Config` object.
    Parameters
    model_params : :class:`~tenpy.tools.params.Config`
        Parameters for the model. See :cfg:config:`TFIModel` below.
    Options
    -------
    .. cfg:config :: TFIModel
        :include: CouplingMPOModel
        conserve : None / 'parity'
            What should be conserved. See :class:`~tenpy.networks.Site.SpinHalfSite`.
        J, g : float / array
            Coupling as defined for the Hamiltonian above.
    " ""
    def init_sites(self, model_params):
        conserve = model_params.get('conserve', 'parity')
        assert conserve != 'Sz'
        if conserve == 'best':
            conserve = 'parity'
            self.logger.info("%s: set conserve to %s", self.name, conserve)
        site = SpinHalfSite(conserve=conserve)
```

```
    return site
    def init_terms(self, model_params)
    J=np.asarray(model_params.get('J', 1.))
    g = np.asarray(model_params.get('g', 1.))
    for u in range(len(self.lat.unit_cell)):
        self.add_onsite(-g, u, 'Sigmaz')
    for ul, u2, dx in self.lat.pairs['nearest_neighbors']:
        self.add_coupling(-J, ul, 'Sigmax', u2, 'Sigmax', dx)
    # done
class TFIChain(TFIModel, NearestNeighborModel):
    """The :class:`TFIModel` on a Chain, suitable for TEBD.
    See the :class:`TFIModel` for the documentation of parameters.
    " " "
    default_lattice = Chain
    force_default_lattice = True
```


### 10.3.6 Automation of Hermitian conjugation

As most physical Hamiltonians are Hermitian, these Hamiltonians are fully determined when only half of the mutually conjugate terms is defined. For example, a simple Hamiltonian:

$$
H=\sum_{\langle i, j\rangle, i<j}-\mathrm{J}\left(c_{i}^{\dagger} c_{j}+c_{j}^{\dagger} c_{i}\right)
$$

is fully determined by the term $c_{i}^{\dagger} c_{j}$ if we demand that Hermitian conjugates are included automatically. In TeNPy, whenever you add a coupling using add_onsite(), add_coupling(), or add_multi_coupling(), you can use the optional argument plus_hc to automatically create and add the Hermitian conjugate of that coupling term - as shown above.

Additionally, in an MPO, explicitly adding both a non-Hermitian term and its conjugate increases the bond dimension of the MPO, which increases the memory requirements of the MPOEnvironment. Instead of adding the conjugate terms explicitly, you can set a flag explicit_plus_hc in the MPOCouplingModel parameters, which will ensure two things:

1. The model and the MPO will only store half the terms of each Hermitian conjugate pair added, but the flag explicit_plus_hc indicates that they represent self + h.c.. In the example above, only the term $c_{i}^{\dagger} c_{j}$ would be saved.
2. At runtime during DMRG, the Hermitian conjugate of the (now non-Hermitian) MPO will be computed and applied along with the MPO, so that the effective Hamiltonian is still Hermitian.

Note: The model flag explicit_plus_hc should be used in conjunction with the flag plus_hc in add_coupling () or add_multi_coupling (). If plus_hc is False while explicit_plus_hc is True the MPO bond dimension will not be reduced, but you will still pay the additional computational cost of computing the Hermitian conjugate at runtime.

Thus, we end up with several use cases, depending on your preferences. Consider the FermionModel. If you do not care about the MPO bond dimension, and want to add Hermitian conjugate terms manually, you would set model_par['explicit_plus_hc'] = False and write:

```
self.add_coupling(-J, u1, 'Cd', u2, 'C', dx)
self.add_coupling(np.conj(-J), u2, 'Cd', u1, 'C', -dx)
```

If you wanted to save the trouble of the extra line of code (but still did not care about MPO bond dimension), you would keep the model_par, but instead write:

```
self.add_coupling(-J, u1, 'Cd', u2, 'C', dx, plus_hc=True)
```

Finally, if you wanted a reduction in MPO bond dimension, you would need to set model_par['explicit_plus_hc'] = True, and write:

```
self.add_coupling(-J, u1, 'Cd', u2, 'C', dx, plus_hc=True)
```


### 10.3.7 Non-uniform terms and couplings

The CouplingModel-methods add_onsite(), add_coupling(), and add_multi_coupling() add a sum over a "couplig" term shifted by lattice vectors. However, some models are not that "uniform" over the whole lattice.

First of all, you might have some local term that gets added only at one specific location in the lattice. You can add such a term for example with add_local_term().

Second, if you have irregular lattices, take a look at the corresponding section in Details on the lattice geometry.
Finally, note that the argument strength for the add_onsite, add_coupling, and add_multi_coupling methods can not only be a numpy scalar, but also a (numpy) array. In general, the sum performed by the methods runs over the given term shifted by lattice vectors as far as possible to still fit the term into the lattice.

For the add_onsite () case this criterion is simple: there is exactly one site in each lattice unit cell with the $u$ specified as separate argument, so the correct shape for the strength array is simply given by Ls. For example, if you want the defacto standard model studied for many-body localization, a Heisenberg chain with random , uniform onsite field $h_{i}^{z} \in[-W, W]$,

$$
H=J \sum_{i=0}^{L-1} \vec{S}_{i} \cdot \vec{S}_{i+1}-\sum_{i=0}^{L} h_{i}^{z} S_{i}^{z}
$$

you can use the SpinChain with the following model parameters:

```
L = 30 # or whatever you like...
W = 5. # MBL transition at W_c ~= 3.5 J
model_params = {
    'L': L,
    'Jx': 1., 'Jy': 1., 'Jz': 1.,
    'hz': 2.*W*(np.random.random(L) - 0.5), # random values in [-W, W], shape (L,)
    'conserve': 'best',
}
M = tenpy.models.spins.SpinChain(model_params)
```

For add_coupling() and add_multi_coupling(), things become a little bit more complicated, and the correct shape of the strength array depends not only on the Ls but also on the boundary conditions of the lattice. Given a term, you can call coupling_shape () and multi_coupling_shape () to find out the correct shape for strength. To avoid any ambiguity, the shape of the strength always has to fit, at least after a tiling performed by to_array().

For example, consider the Su-Schrieffer-Heeger model, a spin-less FermionCha in with hopping strength alternating between two values, say $t 1$ and $t 2$. You can generete this model for example like this:

```
L = 30 # or whatever you like...
t1, t2 = 0.5, 1.5
t_array = np.array([(t1 if i % 2 == 0 else t2) for i in range(L-1)])
model_params = {
    'L': L,
    't': t_array,
    'V': 0., 'mu': 0., # just free fermions, but you can generalize...
    'conserve': 'best'
}
M = tenpy.models.fermions.FermionChain(model_params)
```


### 10.3.8 Some random remarks on models

- Needless to say that we have also various predefined models under tenpy.models.
- Of course, an MPO is all you need to initialize a MPOModel to be used for DMRG; you don't have to use the CouplingModel or CouplingMPOModel. For example an exponentially decaying long-range interactions are not supported by the coupling model but straight-forward to include to an MPO, as demonstrated in the example examples/mpo_exponentially_decaying.py.
- If you want to debug or double check that the you added the correct terms to a CouplingMPOModel, you can print the terms with print (M.all_coupling_terms().to_TermList()). This will
- If the model of your interest contains Fermions, you should read the Fermions and the Jordan-Wigner transformation.
- We suggest writing the model to take a single parameter dictionary for the initialization, as the CouplingMPOModel does. The CouplingMPOModel converts the dictionary to a dict-like Config with some additional features before passing it on to the init_lattice, init_site, ... methods. It is recommended to read out providing default values with model_params.get("key", default_value), see get().
- When you write a model and want to include a test that it can be at least constructed, take a look at tests/ test_model.py.


### 10.4 Simulations

Simulations provide the highest-level interface in TeNPy. They represent one simulation from start (initializing the various classes from given parameters) to end (saving the results to a file). The idea is that they contain the full package of code that you run by a job on a computing cluster. (You don't have to stick to that rule, of course.) In fact, any simulation can be run from the command line, given only a parameter file as input, like this:

```
python -m tenpy -c SimulationClassName parameters.yml
# or alternatively, if tenpy is installed correctly:
tenpy-run -c SimulationClassName parameters.yml
```

Of course, you should replace SimulationClassName with the class name of the simulation class you want to use, for example GroundStateSearch or RealTimeEvolution. For more details, see tenpy. run_commandline().

In some cases, this might not be enough, and you want to do some pre- or post-processing, or just do something a litte bit differently during the simulation. In that case, you can also define your own simulation class (as subclass of one the existing ones).

### 10.5 Details on the lattice geometry

The Lattice class defines the geometry of the system. In the basic form, it represents a unit cell of a few sites repeated in one or multiple directions. Moreover, it maps this higher-dimensional geometry to a one-dimensional chain for MPS-based algorithms.

### 10.5.1 Visualization

A plot of the lattice can greatly help to understand which sites are connected by what couplings. The methods plot_* of the Lattice can do a good job for a quick illustration. Let's look at the Honeycomb lattice as an example.

```
import matplotlib.pyplot as plt
from tenpy.models import lattice
plt.figure(figsize=(5, 6))
ax = plt.gca()
lat = lattice.Honeycomb(Lx=4, Ly=4, sites=None, bc='periodic')
lat.plot_coupling(ax)
lat.plot_order(ax, linestyle=':')
lat.plot_sites(ax)
lat.plot_basis(ax, origin=-0.5*(lat.basis[0] + lat.basis[1]))
ax.set_aspect('equal')
ax.set_xlim(-1)
ax.set_ylim(-1)
plt.show()
```

In this case, the unit cell (shaded green) consists of two sites, which for the purpose of plotting we just set to sites=None; in general you should specify instances of Site for that. The unit cell gets repeated in the directions given by the lattice basis (green arrows at the unit cell boundary). Hence, we can label each site by a lattice index ( $x, y, u$ ) in this case, where $x$ in range (Lx), $y$ in range (Ly) specify the translation of the unit cell and $u$ in range (len (unit_cell)), here $u$ in [ 0,1$]$, specifies the index within the unit cell.

### 10.5.2 How an MPS winds through the lattice: the order

For MPS-based algorithms, we need to map a 2D lattice like the one above to a 1D chain. The red, dashed line in the plot indicates how an MPS winds through the 2D lattice. The MPS index $i$ is a simple enumeration of the sites along this line, shown as numbers next to the sites in the plot. The methods mps2lat_idx () and lat2mps_idx () map indices of the MPS to and from indices of the lattice.

The MPS class itself is (mostly) agnostic of the underlying geometry. For example, expectation_value () will return a 1D array of the expectation value on each site indexed by the MPS index $i$. If you have a two-dimensional lattice, you can use mps2lat_values () to map this result to a 2D array index by the lattice indices.

A suitable order is critical for the efficiency of MPS-based algorithms. On one hand, different orderings can lead to different MPO bond-dimensions, with direct impact on the complexity scaling. On the other hand, it influences how much entanglement needs to go through each bonds of the underlying MPS, e.g., the ground strate to be found in DMRG, and therefore influences the required MPS bond dimensions. For the latter reason, the "optimal" ordering can not be known a priori and might even depend on your coupling parameters (and the phase you are in). In the end, you can just try different orderings and see which one works best.

The simplets way to change the order is to use a non-default value for the initialization parameter order of the Lattice class. This gets passed on to ordering(), which you an override in a custom lattice class to define new possible orderings. Alternatively, you can go the most general way and simply set the attribute order to be a 2D numpy array with lattice indices as rows, in the order you want.


```
import matplotlib.pyplot as plt
from tenpy.models import lattice
Lx, Ly = 3, 3
fig, axes = plt.subplots(2, 2, figsize=(7, 8))
lat1 = lattice.Honeycomb(Lx, Ly, sites=None, bc='periodic') # default order
lat2 = lattice.Honeycomb(Lx, Ly, sites=None, bc='periodic',
    order="Cstyle") # first method to change order
# alternative: directly set "Cstyle" order
lat3 = lattice.Honeycomb(Lx, Ly, sites=None, bc='periodic')
lat3.order = lat2.ordering("Cstyle") # now equivalent to lat2
# general: can apply arbitrary permutation to the order
lat4 = lattice.Honeycomb(Lx, Ly, sites=None, bc='periodic',
    order="Cstyle")
old_order = lat4.order
permutation = []
for i in range(0, len(old_order), 2):
    permutation.append(i+1)
    permutation.append(i)
lat4.order = old_order[permutation, :]
for lat, label, ax in zip([lat1, lat2, lat3, lat4],
    ["order='default'",
    "order='Cstyle'",
    "order='Cstyle'",
    "custom permutation"],
    axes.flatten()):
    lat.plot_coupling(ax)
    lat.plot_sites(ax)
    lat.plot_order(ax, linestyle=':', linewidth=2.)
    ax.set_aspect('equal')
    ax.set_title('order = ' + repr(label))
plt.show()
```


### 10.5.3 Boundary conditions

The Lattice defines the boundary conditions $b c$ in each direction. It can be one of the usual 'open' or 'periodic' in each direcetion.

On top of that, there is the $b c_{-} M P S$ boundary condition of the MPS, one of 'finite', 'segment', 'infinite'. For an 'infinite' MPS, the whole lattice is repeated in the direction of the first basis vector of the lattice. For bc_MPS='infinite', the first direction should always be 'periodic', but you can also define a lattice with bc_MPS='finite', bc=['periodic', 'perioid'] for a finite system on the torus. This is discouraged, though, because the ground state MPS will require the squared bond dimension for the same precision in this case!

For two (or higher) dimensional lattices, e.g for DMRG on an infinite cylinder, you can also specify an integer shift instead of just saying 'periodic': Rolling the 2D lattice up into a cylinder, you have a degree of freedom which sites to connect. This is illustrated in the figure below for a Square lattice with bc=['periodic', shift] for shift in $[-1,0,1]$ (different columns). In the first row, the orange markers indicate a pair of identified sites (see plot_bc_shift ()). The dashed orange line indicates the direction of the cylinder axis. The line where the cylinder is "cut open" therefore winds around the the cylinder for a non-zero shift. (A similar thing happens even for shift $=0$ for more complicated lattices with non-orthogonal basis.) In the second row, we directly draw lines between



order = 'custom permutation'

all sites connected by nearest-neighbor couplings, as they appear in the MPO.


### 10.5.4 Irregular Lattices

The Irregularlattice allows to add or remove sites from/to an existing regular lattice. The doc-string of IrregularLattice contains several examples, let us consider another one here, where we use the IrregularLattice to "fix" the boundary of the Honeycomb lattice: when we use "open" boundary conditions for a finite system, there are two sites (on the lower left, and upper right), wich are not included into any hexagonal. The following example shows how to remove them from the system:

```
import matplotlib.pyplot as plt
from tenpy.models import lattice
Lx, Ly = 3, 3
fig, axes = plt.subplots(1, 2, sharex=True, sharey=True, figsize=(6, 4))
reg_lat = lattice.Honeycomb(Lx=Lx, Ly=Ly, sites=None, bc='open')
irr_lat = lattice.IrregularLattice(reg_lat, remove=[[0, 0, 0], [-1, -1, 1]])
for lat, label, ax in zip([reg_lat, irr_lat],
    ["regular", "irregular"],
    axes.flatten()) :
    lat.plot_coupling(ax)
    lat.plot_order(ax, linestyle=':')
    lat.plot_sites(ax)
    ax.set_aspect('equal')
    ax.set_title(label)
plt.show()
```



### 10.6 Logging and terminal output

By default, calling (almost) any function in TeNPy will not print output, appart from error messages, tracebacks, and warnings. Instead, we use Python's logging module to allow fine-grained redirecting of status messages etc.

Of course, when you get an error message, you should be concerned to find out what it is about and how to fix it. (If you believe it is a bug, report it.) Warnings can be reported either using warnings. warn (. . . ) or with the logging mechanism logger. warn (...). The former is used for warnings about things in your setup that you should fix. The latter give you notifications about bad things that can happen in calculations, e.g. bad conditioning of a matrix, but there is not much you can do about it. Those warnings indicate that you should take your results with a grain of salt and carefully double-check them.

### 10.6.1 Configuring logging

If you also want to see status messages (e.g. during a DMRG run whenever a checkpoint is reached), you can use set the logging level to logging.INFO with the following, basic setup:

```
import logging
logging.basicConfig(level=logging.INFO)
```

We use this snippet in our examples to activate the printing of info messages to the standard output stream. For really detailed output, you can even set the level to logging.DEBUG. logging.basicConfig() also takes a filename argument, which allows to redirect the output to a file instead of stdout. Note that you should call basicConfig only once; subsequent calls have no effect.

More detailed configurations can be made through logging. config. For example, the following both prints log messages to stdout and saves them to`ouput_filename.log`:

```
import logging.config
conf = {
    'version': 1
    'disable_existing_loggers': False,
    'formatters': {'custom': {'format': '%(levelname)-8s: %(message)s'}},
    'handlers': {'to_file': {'class': 'logging.FileHandler',
                            'filename': 'output_filename.log',
                            formatter': 'custom',
                            'level': 'INFO',
                            mode': 'a'},
            'to_stdout': {'class': 'logging.StreamHandler',
                'formatter': 'custom',
                'level': 'INFO',
                'stream': 'ext://sys.stdout'}},
    'root': {'handlers': ['to_stdout', 'to_file'], 'level': 'DEBUG'},
}
logging.config.dictConfig(conf)
```

Note: Whether you use logging.config.fileConfig() or the logging.config.dictConfig(), make sure that you also set disable_existing_loggers=False. Otherwise, it will not work as expected in the case where you import tenpy before setting up the logging.

To also capture warnings, you might also want to call logging. captureWarnings ().
In fact, the above is the default configuration used by tenpy.tools.misc.setup_logging (). If you use a Simulation class, it will automatically call setup_logging () for you, saving the log to the same filename as the Simulation.output_filename but with a .log ending. Moreover, you can easily adjust the log levels with simple parameters, for example with the following configuration (using [yaml] notation):

```
logging_params:
    to_stdout: # nothing in yaml -> None in python => no logging to stdout
    to_file: INFO
    log_levels:
        tenpy.tools.params : WARNING # suppres INFO/DEBUG output for any logging of
\hookrightarrowparameters
```

Of course, you can also explicilty call the setup_logging () yourself, if you don't use the Simulation classes:

```
tenpy.tools.misc.setup_logging({'to_stdout': None, 'to_file': 'INFO', 'filename': 'my_
\hookrightarrowlog.txt',
    'log_levels': {'tenpy.tools.params': 'WARNING'}})
```


### 10.6.2 How to write your own logging (and warning) code

Of course, you can still use simple print (. . .) statements in your code, and they will just appear on your screen. In fact, this is one of the benefits of logging: you can make sure that you only get the print statements you have put yourself, and at the same time redirect the logging messages of tenpy to a file, if you want.

However, these print (. . .) statements are not re-directed to the log-files. Therefore, if you write your own subclasses like Models, I would recommended that you also use the loggers instead of simple print statements. You can read the official logging tutorial for details, but it's actually straight-forward, and just requires at most two steps.

1. If necessary, import the necessary modules and create a logger at the top of your module:
```
import warnings
import logging
logger = logging.getLogger(__name__)
```

Note: Most TeNPy classes that you might want to subclass, like models, algorithm engines or simulations, provide a Logger as self.logger class attribute. In that case you can even skip this step and just use self.logger instead of logger in the snippets below.
2. Inside your funtions/methods/..., make calls like this:

```
if is_likely_bad(options['parameter']):
    # this can be fixed by the user!
    warnings.warn("This is a bad parameter, you shouldn't do this!")
if "old_parameter" in options:
    warnings.warn("Use `new_parameter` instead of `old_parameter`", FutureWarning,
@ 2)
logger.info("starting some lengthy calculation")
n_steps = do_calculation()
if something_bad_happened():
    # the user can't do anything about it
    logger.warn("Something bad happend")
logger.info("calculation finished after %od steps", n_steps)
```

You can use printf-formatting for the arguments of logger.debug(...), logger.info(...), logger. warn (...), as illustrated in the last line.
In summary, instead of just print("do X") statements, use self.logger.info("do X") inside TeNPy classes, or just logger.info("do X") for the module-wide logger, which you can initialize right at the top of your file with the import statements. If you have non-string arguments, add a formatter string, e.g. replace print(max(psi.chi)) with logger.info("ஃd", max(psi.chi)), or even better, logger.info( "max (chi) $=\% \mathrm{~d}$ ", $\max (\mathrm{psi} . \mathrm{chi}))$. For genereic types, use $" \% \mathrm{~s}$ " or $" \% \mathrm{r}$ ", which converts the other arguments to strings with $\operatorname{str}(\ldots$ ) or repr (. . . ) , respectively.

### 10.7 Parameters and options

(We use parameter and option synonymously.)
Standard simulations in TeNPy can be defined by just set of options collected in a dictionary (possibly containing other parameter dictionaries). It can be convenient to represent these options in a [yaml] file, say parameters. yml, which might look like this:

```
output_filename : params_output.h5
overwrite_output : True
model_class : SpinChain
model_params :
    L : 14
    bc_MPS : finite
initial_state_params:
    method : lat_product_state
    product_state : [[up], [down]]
```

```
algorithm_class: TwoSiteDMRG
algorithm_params:
    trunc_params:
        chi_max: 120
        svd_min: 1e.-8
    max_sweeps: 10
    mixer : True
```

Note that the default values and even the allowed/used option names often depend on other parameters. For example, the model_class parameter above given to a Simulation selects a model class, and diffent model classes might have completely different parameters. This gives you freedom to define your own parameters when you implement a model, but it also makes it a little bit harder to keep track of allowed values.

In the TeNPy documentation, we use the Options sections of doc-strings to define parameters that are read out. Each documented parameter is attributed to one set of parameters, called "config", and managed in a Config class at runtime. The above example represents the config for a Simulation, with the model_params representing the config given as options to the model for initialization. Sometimes, there is also a structure of one config including the parameters from another one: For example, the generic parameters for time evolution algorithms, TimeEvolutionAlgorithm are included into the TEBDEngine config, similarly to the sub-classing used.

During runtime, the Config class logs the first use of any parameter (with DEBUG log-level, if the default is used, and with INFO log-level, if it is non-default). Moreover, the default is saved into the parameter dictionary. Hence, it will contain the full set of all used parameters, default and non-default, at the end of a simulation, e.g., in the sim_params of the results returned by tenpy. simulations. Simulation.run().

You can find a list of all the different configs in the cfg-config-index, and a list of all parameters in cfg-option-index.
If you add extra options to your configuration that TeNPy doesn't read out by the end of the simulation, it will issue a warning. Getting such a warnings is an indicator for a typo in your configuration, or an option being in the wrong config dictionary.

### 10.8 Saving to disk: input/output

### 10.8.1 Using pickle

A simple and pythonic way to store data of TeNPy arrays is to use pickle from the Python standard library. Pickle allows to store (almost) arbitrary python objects, and the Array is no exception (and neither are other TeNPy classes).

Say that you have run DMRG to get a ground state $p s i$ as an MPS. With pickle, you can save it to disk as follows:

```
import pickle
with open('my_psi_file.pkl', 'wb') as f:
    pickle.dump(psi, f)
```

Here, the with . . . : structure ensures that the file gets closed after the pickle dump, and the 'wb ' indicates the file opening mode "write binary". Reading the data from disk is as easy as ( ' rb' for reading binary):

```
with open('my_psi_file.pkl', 'rb') as f:
    psi = pickle.load(f)
```

Note: It is a good (scientific) practice to include meta-data to the file, like the parameters you used to generate that state. Instead of just the psi, you can simply store a dictionary containing psi and other data, e.g., data = \{'psi':
psi, 'dmrg_params': dmrg_params, 'model_params': model_params\}. This can save you a lot of pain, when you come back looking at the files a few month later and forgot what you've done to generate them!

In some cases, compression can significantly reduce the space needed to save the data. This can for example be done with gzip (as well in the Python standard library). However, be warned that it might cause longer loading and saving times, i.e. it comes at the penalty of more CPU usage for the input/output. In Python, this requires only small adjustments:

```
import pickle
import gzip
# to save:
with gzip.open('my_data_file.pkl', 'wb') as f:
    pickle.dump(data, f)
# and to load:
with gzip.open('my_data_file.pkl', 'rb') as f:
    data = pickle.load(data, f)
```


### 10.8.2 Using HDF5 with h5py

While pickle is great for simple input/output of python objects, it also has disadvantages. The probably most dramatic one is the limited portability: saving data on one PC and loading it on another one might fail! Even exporting data from Python 2 to load them in Python 3 on the same machine can give quite some troubles. Moreover, pickle requires to load the whole file at once, which might be unnecessary if you only need part of the data, or even lead to memory problems if you have more data on disk than fits into RAM.

Hence, we support saving to HDF5 files as an alternative. The h5py package provides a dictionary-like interface for the file/group objects with numpy-like data sets, and is quite easy to use. If you don't know about HDF5, read the quickstart of the h5py documentation (and this guide).

The implementation can be found in the tenpy.tools.hdf5_io module with the Hdf5Saver and Hdf5Loader classes and the wrapper functions save_to_hdf5(), load_from_hdf5().

The usage is very similar to pickle:

```
import h5py
from tenpy.tools import hdf5_io
data = {"psi": psi, # e.g. an MPS
        "model": my_model,
        "parameters": {"L": 6, "g": 1.3}}
with h5py.File("file.h5", 'w') as f:
    hdf5_io.save_to_hdf5(f, data)
# ...
with h5py.File("file.h5", 'r') as f:
    data = hdf5_io.load_from_hdf5(f)
    # or for partial reading:
    pars = hdf5_io.load_from_hdf5(f, "/parameters")
```

Warning: Like loading a pickle file, loading data from a manipulated HDF5 file with the functions described has the potential to cause arbitrary code execution. Only load data from trusted sources!

Note: The hickle package imitates the pickle functionality while saving the data to HDF5 files. However, since it aims to be close to pickle, it results in a more complicated data structure than we want here.

Note: To use the export/import features to HDF5, you need to install the h5py python package (and hence some version of the HDF5 library).

## Data format specification for saving to HDF5

This section motivates and defines the format how we save data of TeNPy-defined classes. The goal is to have the save_to_hdf5() function for saving sufficiently simple enough python objects (supported by the format) to disk in an HDF5 file, such that they can be reconstructed with the load_from_hdf5 () function, as outlined in the example code above.

## Guidelines of the format:

0 . Store enough data such that load_from_hdf5 () can reconstruct a copy of the object (provided that the save did not fail with an error).

1. Objects of a type supported by the HDF5 datasets (with the h5py interface) should be directly stored as h5py Dataset. Such objects are for example numpy arrays (of non-object dtype), scalars and strings.
2. Allow to save (nested) python lists, tuples and dictionaries with values (and keys) which can be saved.
3. Allow user-defined classes to implement a well-defined interface which allows to save instances of that class, hence extending what data can be saved. An instance of a class supporting the interface gets saved as an HDF5 Group. Class attributes are stored as entries of the group, metadata like the type should be stored in HDF5 attributes, see attributes.
4. Simple and intuitive, human-readable structure for the HDF5 paths. For example, saving a simple dictionary \{'a': np.arange (10), 'b': 123.45\} should result in an HDF5 file with just the two data sets /a and $/ \mathrm{b}$.
5. Allow loading only a subset of the data by specifying the path of the HDF5 group to be loaded. For the above example, specifying the path $/ \mathrm{b}$ should result in loading the float 123.45 , not the array.
6. Avoid unnecessary copies if the same python object is referenced by different names, e.g, for the data \{'c': large_obj, 'd': large_obj\} with to references to the same large_obj, save it only once and use HDF5 hard-links such that / c and / d are the same HDF5 dataset/group. Also avoid the copies during the loading, i.e., the loaded dictionary should again have two references to a single object large_obj. This is also necessary to allow saving and loading of objects with cyclic references.
The full format specification is given by the what the code in hdf5_io does... Since this is not trivial to understand, let me summarize it here:

- Following 1), simple scalars, strings and numpy arrays are saved as Dataset. Other objects are saved as a HDF5 Group, with the actual data being saved as group members (as sub-groups and sub-datasets) or as attributes (for metadata or simple data).
- The type of the object is stored in the HDF5 attribute 'type ', which is one of the global REPR_* variables in tenpy.tools.hdf5_io. The type determines the format for saving/loading of builtin types (list, ...)
- Userdefined classes which should be possible to export/import need to implement the methods save_hdf5 and from_hdf5 as specified in Hdf5Exportable. When saving such a class, the attribute 'type' is automatically set to 'instance ', and the class name and module are saved under the attributes 'module' and 'class'. During loading, this information is used to automatically import the module, get the class and
call the classmethod $f r o m \_h d f 5$ for reconstruction. This can only work if the class definition already exists, i.e., you can only save class instances, not classes itself.
- For most (python) classes, simply subclassing Hdf5Exportable should work to make the class exportable. The latter saves the contents of $\qquad$ dict $\qquad$ , with the extra attribute 'format' specifying whether the dictionary is "simple" (see below.).
- The None object is saved as a group with the attribute 'type' being 'None' and no subgroups.
- For iterables (list, tuple and set), we simple enumerate the entries and save entries as group members under the names '0', '1', '2', ..., and a maximum 'len' attribute.
- The format for dictionaries depends on whether all keys are "simple", which we define as being strings which are valid path names in HDF5, see valid_hdf5_path_component (). Following 4), the keys of a simple dictionary are directly used as names for group members, and the values being whatever object the group member represents.
- Partial loading along 5) is possible by directly specifying the subgroup or the path to load_from_hdf5 ().
- Guideline 6) is ensured as much as possible. However, there is a bug/exception: tuples with cyclic references are not re-constructed correctly; the inner objects will be lists instead of tuples (but with the same object entries).

Finally, we have to mention that many TeNPy classes are Hdffexportable. In particular, the Array supports this. To see what the exact format for those classes is, look at the save_hdf5 and from_hdf5 methods of those classes.

Note: There can be multiple possible output formats for the same object. The dictionary - with the format for simple keys or general keys - is such an example, but userdefined classes can use the same technique in their from_hdf5 method. The user might also explicitly choose a "lossy" output format (e.g. "flat" for np_conserved Arrays and LegCharges).

Tip: The above format specification is quite general and not bound to TeNPy. Feel free to use it in your own projects ;-) To separate the development, versions and issues of the format clearly from TeNPy, we maintain the code for it in a separate git repository, https://github.com/tenpy/hdf5_io

### 10.9 Fermions and the Jordan-Wigner transformation

The Jordan-Wigner tranformation maps fermionic creation- and annihilation operators to (bosonic) spin-operators.

### 10.9.1 Spinless fermions in 1D

Let's start by explicitly writing down the transformation. With the Pauli matrices $\sigma_{j}^{x, y, z}$ and $\sigma_{j}^{ \pm}=\left(\sigma_{j}^{x} \pm \mathrm{i} \sigma_{j}^{y}\right) / 2$ on each site, we can map

$$
\begin{aligned}
n_{j} & \leftrightarrow\left(\sigma_{j}^{z}+1\right) / 2 \\
c_{j} & \leftrightarrow(-1)^{\sum_{l<j} n_{l}} \sigma_{j}^{-} \\
c_{j}^{\dagger} & \leftrightarrow(-1)^{\sum_{l<j} n_{l}} \sigma_{j}^{+}
\end{aligned}
$$

The $n_{l}$ in the second and third row are defined in terms of Pauli matrices according to the first row. We do not interpret the Pauli matrices as spin-1/2; they have nothing to do with the spin in the spin-full case. If you really want to interpret them physically, you might better think of them as hard-core bosons ( $b_{j}=\sigma_{j}^{-}, b_{j}^{\dagger}=\sigma_{j}^{+}$), with a spin of the fermions mapping to a spin of the hard-core bosons.

Note that this transformation maps the fermionic operators $c_{j}$ and $c_{j}^{\dagger}$ to global operators; although they carry an index $j$ indicating a site, they actually act on all sites $l<=j$ ! Thus, clearly the operators $C$ and Cd defined in the Fermionsite do not directly correspond to $c_{j}$ and $c_{j}^{\dagger}$. The part $(-1)^{\sum_{l<j} n_{l}}$ is called Jordan-Wigner string and in the Fermionsite is given by the local operator $J W:=(-1)^{n_{l}}$ acting all sites $l<j$. Since this important, let me stress it again:

Warning: The fermionic operator $c_{j}$ (and similar $c_{j}^{\dagger}$ ) maps to a global operator consisting of the Jordan-Wigner string built by the local operator JW on sites $1<j$ and the local operator C (or Cd, respectively) on site $j$.

On the sites itself, the onsite operators C and Cd in the Fermionsite fulfill the correct anti-commutation relation, without the need to include JW strings. The JW string is necessary to ensure the anti-commutation for operators acting on different sites.

Written in terms of onsite operators defined in the FermionSite, with the $i$-th entry entry in the list acting on site $i$, the relations are thus:

```
["JW", ..., "JW", "C", "Id", ..., "Id"] # for the annihilation operator
["JW", ..., "JW", "Cd", "Id", ..., "Id"] # for the creation operator
```

Note that "JW" squares to the identity, "JW JW" == "Id", which is the reason that the Jordan-wigner string completely cancels in $n_{j}=c_{j}^{\dagger} c_{j}$. In the above notation, this can be written as:

```
["JW", ..., "JW", "Cd", "Id", ..., "Id"] * ["JW", ..., "JW", "C", "Id", ..., "Id"]
== ["JW JW", ..., "JW JW", "Cd C", "Id Id", ..., "Id Id"] # by definition of,
\hookrightarrowthe tensorproduct
== ["Id", ..., "Id", "N", "Id", ..., "Id"] # by definition off
\hookrightarrowthe local operators
# ("X Y" stands for the local operators X and Y applied on the same site. We assume,
\hookrightarrowthat the "Cd" and "C" on the first line act on the same site.)
```

For a pair of operators acting on different sites, JW strings have to be included for every site between the operators. For example, taking i $<j, c_{i}^{\dagger} c_{j} \leftrightarrow \sigma_{i}^{+}(-1)^{\sum_{i<=l<j} n_{l}} \sigma_{j}^{-}$. More explicitly, for $j=i+2$ we get:

```
["JW", ..., "JW", "Cd", "Id", "Id", "Id", ..., "Id"] * ["JW", ..., "JW", "JW", "JW",
\hookrightarrow"C", "Id", ..., "Id"]
== ["JW JW", ..., "JW JW", "Cd JW", "Id JW", "Id C", ..., "Id"]
== ["Id", ..., "Id", "Cd JW", "JW", "C", ..., "Id"]
```

In other words, the Jordan-Wigner string appears only in the range i $<=1<j$, i.e. between the two sites and on the smaller/left one of them. (You can easily generalize this rule to cases with more than two $c$ or $c^{\dagger}$.)

This last line (as well as the last line of the previous example) can be rewritten by changing the order of the operators Cd JW to "JW Cd " $==-\mathrm{Cd}$ ". (This is valid because either site $i$ is occupied, yielding a minus sign from the JW , or it is empty, yielding a 0 from the Cd.)
This is also the case for $j<i$, say $j=i-2: c_{i}^{\dagger} c_{j} \leftrightarrow(-1)^{\sum_{j<=l<i} n_{l}} \sigma_{i}^{+} \sigma_{j}^{-}$. As shown in the following, the JW again appears on the left site, but this time acting after C :

```
["JW", ..., "JW", "JW", "JW", "Cd", "Id", ..., "Id"] * ["JW", ..., "JW", "C", "Id",
\hookrightarrow"Id", "Id", ..., "Id"]
== ["JW JW", ..., "JW JW", "JW C", "JW", "Cd Id", ..., "Id"]
== ["Id", ..., "Id", "JW C", "JW", "Cd", ..., "Id"]
```


### 10.9.2 Higher dimensions

For an MPO or MPS, you always have to define an ordering of all your sites. This ordering effectifely maps the higherdimensional lattice to a 1D chain, usually at the expence of long-range hopping/interactions. With this mapping, the Jordan-Wigner transformation generalizes to higher dimensions in a straight-forward way.

### 10.9.3 Spinful fermions



As illustrated in the above picture, you can think of spin- $1 / 2$ fermions on a chain as spinless fermions living on a ladder (and analogous mappings for higher dimensional lattices). Each rung (a blue box in the picture) forms a SpinHalffermionSite which is composed of two FermionSite (the circles in the picture) for spin-up and spin-down. The mapping of the spin- $1 / 2$ fermions onto the ladder induces an ordering of the spins, as the final result must again be a one-dimensional chain, now containing both spin species. The solid line indicates the convention for the ordering, the dashed lines indicate spin-preserving hopping $c_{s, i}^{\dagger} c_{s, i+1}+h . c$. and visualize the ladder structure. More generally, each species of fermions appearing in your model gets a separate label, and its Jordan-Wigner string includes the signs $(-1)^{n_{l}}$ of all species of fermions to the 'left' of it (in the sense of the ordering indicated by the solid line in the picture).

In the case of spin- $1 / 2$ fermions labeled by $\uparrow$ and $\downarrow$ on each site, the complete mapping is given (where $j$ and $l$ are indices of the FermionSite):

$$
\begin{aligned}
& n_{\uparrow, j} \leftrightarrow\left(\sigma_{\uparrow, j}^{z}+1\right) / 2 \\
& n_{\downarrow, j} \leftrightarrow\left(\sigma_{\downarrow, j}^{z}+1\right) / 2 \\
& c_{\uparrow, j} \leftrightarrow(-1)^{\sum_{l<j} n_{\uparrow, l}+n_{\downarrow, l}} \sigma_{\uparrow, j}^{-} \\
& c_{\uparrow, j}^{\dagger} \leftrightarrow(-1)^{\sum_{l<j} n_{\uparrow, l}+n_{\downarrow, l}} \sigma_{\uparrow, j}^{+} \\
& c_{\downarrow, j} \leftrightarrow(-1)^{\sum_{l<j} n_{\uparrow, l}+n_{\downarrow, l}}(-1)^{n_{\uparrow, j}} \sigma_{\downarrow, j}^{-} \\
& c_{\downarrow, j}^{\dagger} \leftrightarrow \leftrightarrow(-1)^{\sum_{l<j} n_{\uparrow, l}+n_{\downarrow, l}}(-1)^{n_{\uparrow, j}} \sigma_{\downarrow, j}^{+}
\end{aligned}
$$

In each of the above mappings the operators on the right hand sides commute; we can rewrite $(-1)^{\sum_{l<j} n_{\uparrow, l}+n_{\downarrow, l}}=$ $\prod_{l<j}(-1)^{n_{\uparrow, l}}(-1)^{n_{\downarrow, l}}$, which resembles the actual structure in the code more closely. The parts of the operator acting in the same box of the picture, i.e. which have the same index $j$ or $l$, are the 'onsite' operators in the SpinHalfFermionSite: for example JW on site $j$ is given by $(-1)^{n_{\uparrow, j}}(-1)^{n_{\downarrow, j}}, \mathrm{Cu}$ is just the $\sigma_{\uparrow, j}^{-}$, Cdu is $\sigma_{\uparrow, j}^{+}$, Cd is $(-1)^{n_{\uparrow, j}} \sigma_{\downarrow, j}^{-}$. and Cdd is $(-1)^{n_{\uparrow, j}} \sigma_{\downarrow, j}^{+}$. Note the asymmetry regarding the spin in the definition of the onsite operators: the spin-down operators include Jordan-Wigner signs for the spin-up fermions on the same site. This asymetry stems from the ordering convention introduced by the solid line in the picture, according to which the spin-up site is "left" of the spin-down site. With the above definition, the operators within the same SpinHalfFermionSite fulfill the expected commutation relations, for example " $\mathrm{Cu} \mathrm{Cdd} "==-\mathrm{Cdd} \mathrm{Cu} "$, but again the JW on sites left of the operator pair is crucial to get the correct commutation relations globally.

Warning: Again, the fermionic operators $c_{\downarrow, j}, c_{\downarrow, j}^{\dagger}, c_{\downarrow, j}, c_{\downarrow, j}^{\dagger}$ correspond to global operators consisting of the Jordan-Wigner string built by the local operator JW on sites $1<j$ and the local operators ' Cu ', 'Cdu', 'Cd', 'Cdd' on site j.

Written explicitly in terms of onsite operators defined in the FermionSite, with the $j$-th entry entry in the list acting on site $j$, the relations are:

```
["JW", ..., "JW", "Cu", "Id", ..., "Id"] # for the annihilation operator spin-up
["JW", ..., "JW", "Cd", "Id", ..., "Id"] # for the annihilation operator spin-down
["JW", ..., "JW", "Cdu", "Id", ..., "Id"] # for the creation operator spin-up
["JW", ..., "JW", "Cdd", "Id", ..., "Id"] # for the creation operator spin-down
```

As you can see, the asymmetry regaring the spins in the definition of the local onsite operators " Cu ", " Cd ", " Cdu ", "Cdd" lead to a symmetric definition in the global sense. If you look at the definitions very closely, you can see that in terms like ["Id", "Cd JW", "JW", "Cd"] the Jordan-Wigner sign $(-1)^{n_{\uparrow}, 2}$ appears twice (namely once in the definition of " Cd " and once in the "JW" on site 2 ) and could in principle be canceled, however in favor of a simplified handling in the code we do not recommend you to cancel it. Similar, within a spinless FermionSite, one can simplify "Cd JW" == "Cd" and "JW C" == "C", but these relations do not hold in the SpinHalfSite, and for consistency we recommend to explicitly keep the "JW" operator string even in nearest-neighbor models where it is not strictly necessary.

### 10.9.4 How to handle Jordan-Wigner strings in practice

There are only a few pitfalls where you have to keep the mapping in mind: When building a model, you map the physical fermionic operators to the usual spin/bosonic operators. The algorithms don't care about the mapping, they just use the given Hamiltonian, be it given as MPO for DMRG or as nearest neighbor couplings for TEBD. Only when you do a measurement (e.g. by calculating an expectation value or a correlation function), you have to reverse this mapping. Be aware that in certain cases, e.g. when calculating the entanglement entropy on a certain bond, you cannot reverse this mapping (in a straightforward way), and thus your results might depend on how you defined the Jordan-Wigner string.

Whatever you do, you should first think about if (and how much of) the Jordan-Wigner string cancels. For example for many of the onsite operators (like the particle number operator N or the spin operators in the SpinHalfFermionSite) the Jordan-Wigner string cancels completely and you can just ignore it both in onsiteterms and couplings. In case of two operators acting on different sites, you typically have a Jordan-Wigner string inbetween (e.g. for the $c_{i}^{\dagger} c_{j}$ examples described above and below) or no Jordan-Wigner strings at all (e.g. for densitydensity interactions $n_{i} n_{j}$ ). In fact, the case that the Jordan Wigner string on the left of the first non-trivial operator does not cancel is currently not supported for models and expectation values, as it usually doesn't appear in practice. For terms involving more operators, things tend to get more complicated, e.g. $c_{i}^{\dagger} c_{j}^{\dagger} c_{k} c_{l}$ with $i<j<k<l$ requires a Jordan-Wigner string on sites $m$ with $i \leq m<j$ or $k \leq m<l$, but not for $j<m<k$.

Note: TeNPy keeps track of which onsite operators need a Jordan-Wigner string in the Site class, specifically in need_JW_string and op_needs_JW (). Hence, when you define custom sites or add extra operators to the sites, make sure that op_needs_JW () returns the expected results.

When building a model the Jordan-Wigner strings need to be taken into account. If you just specify the $H_{-} M P O$ or $H \_b o n d$, it is your responsibility to use the correct mapping. However, if you use the add_coupling() method of the CouplingModel, (or the generalization add_multi_coupling() for more than 2 operators), TeNPy can use the information from the Site class to automatically add Jordan-Wigner strings as needed. Indeed, with the default argument op_string=None, add_coupling will automatically check whether the operators need Jordan-Wigner strings and correspondlingly set op_string='JW', str_on_first=True, if necessary. For add_multi_coupling, you cann't even explicitly specify the correct Jordan-Wigner strings, but you must use op_string=None, from which it will automatically determine where Jordan-Wigner strings are needed.

Obviously, you should be careful about the convention which of the operators is applied first (in a physical sense as an operator acting on a state), as this corresponds to a sign of the prefactor. Read the doc-strings of add_coupling () add_multi_coupling() for details.
As a concrete example, let us specify a hopping $\sum_{i}\left(c_{i}^{\dagger} c_{i+1}+h . c.\right)=\sum_{i}\left(c_{i}^{\dagger} c_{i+1}+c_{i}^{\dagger} c_{i-1}\right)$ in a 1D chain of FermionSite with add_coupling(). The recommended way is just:

```
self.add_coupling(strength, 0, 'Cd', 0, 'C', 1, plus_hc=True)
```

If you want to specify both the Jordan-Wigner string and the h.c. term explicitly, you can use:

```
self.add_coupling(strength, 0, 'Cd', 0, 'C', 1, op_string='JW', str_on_first=True)
self.add_coupling(strength, 0, 'Cd', 0, 'C', -1, op_string='JW', str_on_first=True)
```

Slightly more complicated, to specify the hopping $\sum_{\langle i, j\rangle, s}\left(c_{s, i}^{\dagger} c_{s, j}+h . c.\right)$ in the Fermi-Hubbard model on a 2D square lattice, we could use:

```
for (dx, dy) in [(1, 0), (0, 1)]:
    self.add_coupling(strength, 0, 'Cdu', 0, 'Cu', (dx, dy), plus_hc=True) # spin up
    self.add_coupling(strength, 0, 'Cdd', 0, 'Cd', (dx, dy), plus_hc=True) # spin_s
cown
# or without `plus_hc`
for (dx, dy) in [(1, 0), (-1, 0), (0, 1), (0, -1)]: # include -dx !
    self.add_coupling(strength, 0, 'Cdu', 0, 'Cu', (dx, dy)) # spin up
    self.add_coupling(strength, 0, 'Cdd', 0, 'Cd', (dx, dy)) # spin down
# or specifying the 'JW' string explicitly
for (dx, dy) in [(1, 0), (-1, 0), (0, 1), (0, -1)]:
    self.add_coupling(strength, 0, 'Cdu', 0, 'Cu', (dx, dy), 'JW', True) # spin up
    self.add_coupling(strength, 0, 'Cdd', 0, 'Cd', (dx, dy), 'JW', True) # spin down
```

The most important functions for doing measurements are probably expectation_value() and correlation_function (). Again, if all the Jordan-Wigner strings cancel, you don't have to worry about them at all, e.g. for many onsite operators or correlation functions involving only number operators. If you build multi-site operators to be measured by expectation_value, take care to include the Jordan-Wigner string correctly.

Some MPS methods like correlation_function(), expectation_value_term() and expectation_value_terms_sum() automatically add Jordan-Wignder strings (at least with default arguments). Other more low-level functions like expectation_value_multi_sites() don't do it. Hence, you should always watch out during measurements, if the function used needs special treatment for Jordan-Wigner strings.

### 10.10 Protocol for using (i)DMRG

While this documentation contains extensive guidance on how to interact with the tenpy, it is often unclear how to approach a physics question using these methods. This page is an attempt to provide such guidance, describing a protocol on how to go from a model implementation to an answered question.

The basic workflow for an (i)DMRG project is as follows, with individual steps expanded on later where necessary.

1. Confirm the correctness of the model implementation.
2. Run some low-effort tests to see whether the question seems answerable.
3. If the tests are successful, run production-quality simulations. This will be entirely particular to the project you're working on.
4. Confirm that your results are converged.

### 10.10.1 Confirming the model is correct

Although TeNPy makes model implementation much easier than constructing the MPO by hand, one should still ensure that the MPO represents the intended model faithfully. There are several possible ways to do this. Firstly, for sufficiently small system sizes, one can contract the entire MPO into a matrix, and inspect the matrix elements. In TeNPy, this can be done using get_full_hamiltonian (). These should reproduce the analytical Hamiltonian up to machine precision, or any other necessary cut-off (e.g., long-range interactions may be truncated at some finite distance).

Secondly, if the model basis allows it, one can construct (product state) MPSs for known eigenstates of the model and evaluate whether these reproduce the correct eigenvalues upon contraction with the MPO.

Finally, one can sometimes construct a basis of single- or even two-particle MPSs in some basis, and evaluate the MPO on this basis to get a representation of the single- and two-particle Hamiltonian. If the model contains only single- and two-body terms, this latter approach should reproduce all terms in the Hamiltonian.

### 10.10.2 Low-effort tests

As not every state can be accurately represented by an MPS, some results are outside the reach of (i)DMRG. To prevent wasting considerable numerical resources on a fruitless project, it is recommended to run some low-effort trials first, and see whether any indication of the desired result can be found. If so, one can then go on to more computationally expensive simulations. If not, one should evaluate:

1. Whether there is a mistake in the model or simulation set-up,
2. Whether a slightly more computationally expensive test would potentially yield a result, or
3. Whether your approach is unfortunately out of reach of (i)DMRG.

To set up low-effort trials, one should limit system size, bond dimension and the range of interactions, as well as (if possible) target a non-critical region of phase space. All these measures reduce the size of and/or entanglement entropy needing to be captured by the MPS, which yields both memory and run time advantages. Of course, one introduces a trade-off between computational cost and accuracy, which is why one should be careful to not put too much faith into results obtained at this stage.

### 10.10.3 Detecting convergence issues

Ensuring that the results of an (i)DMRG simulation are well-converged and thus reliable is a hugely important part of any (i)DMRG study. Possible indications that there might be a convergence issue include:

1. The simulation shows a non-monotonous decrease of energy, and/or a non-monotonous increase of entanglement entropy. An increase of energy or decrease of entanglement entropy on subsequent steps within a sweep, or between subsequent sweeps, are particularly suspicious.
2. The simulation does not halt because it reached a convergence criterion, but because it reached its maximum number of sweeps.
3. Results vary wildly under small changes of parameters. In particular, if a small change in bond dimension yields a big change in results, one should be suspicious of the data.

### 10.10.4 Combating convergence issues

To combat convergence issues of the (i)DMRG algorithm, several strategies (short of switching to a different method) can be attempted:

1. Ensure that there are no errors in the model (see above) or the simulation set-up.
2. Increase the maximum bond dimension.
3. Ramp up the maximum bond dimension during simulation, rather than starting at the highest value. I.e., define a schedule wherein the first $N_{\text {sweeps }}$ sweeps run at some $\chi_{1}<\chi_{\max }$, the next $N_{\text {sweeps }}$ at $\chi_{1}<\chi_{2}<\chi_{\text {max }}$, etc. This can be done through the chi_list option of the DMRGEngine. You should also make sure that the max_hours option is set to sufficiently long runtimes.
4. Increase the maximum number of sweeps the algorithm is allowed to make, through the max_sweeps option of the DMRGEngine.
5. Change the Mixer settings to in- or decrease the effects of the mixer.
6. Change convergence criteria. This will not overcome convergence issues in itself, but can help fine tune the (i)DMRG simulation if it takes a long time to converge (relax the convergence constraints), or if the simulation finishes too soon (tighten the constraints). Criteria to consider are max_E_err and max_S_err, in DMRGEngine.
7. Increase the minimum number of sweeps taken by the algorithm. Again, this will not resolve issues due to bad convergence, but might prevent bad results due to premature convergence. This can be done through the min_sweeps option of the DMRGEngine.
8. Change the size and shape of the MPS unit cell (where possible), in case an artificially enforced translational invariance prevents the algorithm from finding a true ground state which is incommensurate with this periodicity. For example, a chain system which has a true ground state that is periodic in three sites, will not be accurately represented by a two-site MPS unit cell, as the latter enforces two-site periodicity.
In some instances, it is essentially unavoidable to encounter convergence issues. In particular, a simulation of a critical state can cause problems with (i)DMRG convergence, as these states violate the area law underlying an accurate MPS approximation. In these cases, one should acknowledge the difficulties imposed by the method and take care to be very careful in interpreting the data.

## ELEVEN

## EXAMPLES

### 11.1 Toycodes

These toycodes are meant to give you a flavor of the different algorithms, while keeping the codes as readable and simple as possible. The scripts are included in the [TeNPySource] repository in the folder toycodes/, but not part of the basic TeNpy library; the only requirements to run them are Python 3, Numpy, and Scipy.

### 11.1.1 a_mps.py

on github.

```
"""Toy code implementing a matrix product state."""
# Copyright 2018-2021 TeNPy Developers, GNU GPLv3
import numpy as np
from scipy.linalg import svd
# if you get an error message "LinAlgError: SVD did not converge",
# uncomment the following line. (This requires TeNPy to be installed.)
# from tenpy.linalg.svd_robust import svd # (works like scipy.linalg.svd)
class SimpleMPS:
    """Simple class for a matrix product state.
    We index sites with `i` from 0 to L-1; bond `i` is left of site `i`.
    We *assume* that the state is in right-canonical form.
    Parameters
    ----------
    Bs, Ss, bc:
        Same as attributes.
    Attributes
    BS : list of np.Array[ndim=3]
            The 'matrices' in right-canonical form, one for each physical site
            (within the unit-cell for an infinite MPS).
            Each `B[i]` has legs (virtual left, physical, virtual right), in short ``vL iv
GR``
    Ss : list of np.Array[ndim=1]
            The Schmidt values at each of the bonds, `'Ss[i]`` is left of ``Bs[i]``.
    bc : 'infinite', 'finite'
```

```
        Boundary conditions.
    L : int
    Number of sites (in the unit-cell for an infinite MPS).
    nbonds : int
    Number of (non-trivial) bonds: L-1 for 'finite' boundary conditions
    """
    def ___init__(self, Bs, Ss, bc='finite'):
    assert bc in ['finite', 'infinite']
    self.Bs = Bs
    self.Ss = Ss
    self.bc = bc
    self.L = len(Bs)
    self.nbonds = self.L - 1 if self.bc == 'finite' else self.L
    def copy(self):
    return SimpleMPS([B.copy() for B in self.Bs], [S.copy() for S in self.Ss],u
\hookrightarrowself.bc)
    def get_thetal(self, i):
    """Calculate effective single-site wave function on sites i in mixedu
\hookrightarrowcanonical form.
    The returned array has legs ``vL, i, vR`` (as one of the Bs).
    """
    return np.tensordot(np.diag(self.Ss[i]), self.Bs[i], [1, 0]) # vL [vL'],-
\hookrightarrow[vL] i vR
    def get_theta2(self, i):
    """Calculate effective two-site wave function on sites i,j=(i+1) in mixed_
\hookrightarrowcanonical form.
    The returned array has legs ``vL, i, j, vR``.
    """
    j = (i + 1) % self.L
    return np.tensordot(self.get_thetal(i), self.Bs[j], [2, 0]) # vL i [vR],\sqcup
\hookrightarrow[VL] j vR
    def get_chi(self):
    """Return bond dimensions."""
    return [self.Bs[i].shape[2] for i in range(self.nbonds)]
    def site_expectation_value(self, op):
    """Calculate expectation values of a local operator at each site."""
    result = []
    for i in range(self.L):
        theta = self.get_thetal(i) # vL i vR
        op_theta = np.tensordot(op, theta, axes=[1, 1]) # i [i*], vL [i] vR
        result.append(np.tensordot(theta.conj(), op_theta, [[0, 1, 2], [1, 0, ५
@2]]))
        # [vL*] [i*] [vR*], [i] [vL] [vR]
    return np.real_if_close(result)
    def bond_expectation_value(self, op):
    """Calculate expectation values of a local operator at each bond."""
    result = []
    for i in range(self.nbonds):
        theta = self.get_theta2(i) # vL i j vR
```

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```
    op_theta = np.tensordot(op[i], theta, axes=[[2, 3], [1, 2]])
        # i j [i*] [j*], vL [i] [j] vR
        result.append(np.tensordot(theta.conj(), op_theta, [[0, 1, 2, 3], [2, 0,-
@1, 3]]))
            # [vL*] [i*] [j*] [vR*], [i] [j] [vL] [vR]
    return np.real_if_close(result)
    def entanglement_entropy(self):
    """Return the (von-Neumann) entanglement entropy for a bipartition at any of,
\hookrightarrowthe bonds."""
    bonds = range(1, self.L) if self.bc == 'finite' else range(0, self.L)
    result = []
    for i in bonds:
        S = self.Ss[i].copy()
        S[S < 1.e-20] = 0. # 0*log(0) should give 0; avoid warning or NaN.
        S2 = S * S
            assert abs(np.linalg.norm(S) - 1.) < 1.e-14
            result.append(-np.sum(S2 * np.log(S2)))
    return np.array(result)
    def correlation_length(self):
    """Diagonalize transfer matrix to obtain the correlation length."""
    import scipy.sparse.linalg.eigen.arpack as arp
    assert self.bc == 'infinite' # works only in the infinite case
    B = self.Bs[0] # vL i vR
    chi = B.shape[0]
    T = np.tensordot(B, np.conj(B), axes=[1, 1]) # vL [i] vR, vL* [i*] vR*
    T = np.transpose(T, [0, 2, 1, 3]) # vL vL* vR vR*
    for i in range(1, self.L):
        B = self.Bs[i]
        T = np.tensordot(T, B, axes=[2, 0]) # vL vL* [vR] vR*, [vL] i vR
        T = np.tensordot(T, np.conj(B), axes=[[2, 3], [0, 1]])
        # vL vL* [vR*] [i] vR, [vL*] [i*] vR*
        T = np.reshape(T, (chi**2, chi**2))
        # Obtain the 2nd largest eigenvalue
        eta = arp.eigs(T, k=2, which='LM', return_eigenvectors=False, ncv=20)
        return -self.L / np.log(np.min(np.abs(eta)))
def init_FM_MPS(L, d, bc='finite'):
    """Return a ferromagnetic MPS (= product state with all spins up)"""
    B = np.zeros([1, d, 1], dtype=float)
    B[0, 0, 0] = 1.
    S = np.ones([1], dtype=float)
    Bs = [B.copy() for i in range(L)]
    Ss = [S.copy() for i in range(L)]
    return SimpleMPS(Bs, Ss, bc)
def split_truncate_theta(theta, chi_max, eps):
    """Split and truncate a two-site wave function in mixed canonical form.
    Split a two-site wave function as follows::
        vL --(theta)-- vR => vL -- (A)--diag(S)--(B)-- vR
            l l l l i
```

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```
Afterwards, truncate in the new leg (labeled ``vC``).
Parameters
----------
theta : np.Array[ndim=4]
    Two-site wave function in mixed canonical form, with legs ``vL, i, j, vR``.
chi_max : int
    Maximum number of singular values to keep
eps : float
    Discard any singular values smaller than that.
Returns
-------
A : np.Array[ndim=3]
    Left-canonical matrix on site i, with legs ``vL, i, vC``
S : np.Array[ndim=1]
    Singular/Schmidt values.
B : np.Array[ndim=3]
    Right-canonical matrix on site j, with legs ``vC, j, vR``
"""
chivL, dL, dR, chivR = theta.shape
theta = np.reshape(theta, [chivL * dL, dR * chivR])
X, Y, Z = svd(theta, full_matrices=False)
# truncate
chivC = min(chi_max, np.sum(Y > eps))
piv = np.argsort(Y) [::-1][:chivC] # keep the largest `chivC` singular values
X, Y, Z = X[:, piv], Y[piv], Z[piv, :]
# renormalize
S = Y / np.linalg.norm(Y) # == Y/sqrt(sum(Y**2))
# split legs of X and Z
A = np.reshape(X, [chivL, dL, chivC])
B = np.reshape(Z, [chivC, dR, chivR])
return A, S, B
```


### 11.1.2 b_model.py

on github.

```
"""Toy code implementing the transverse-field ising model."""
# Copyright 2018-2021 TeNPy Developers, GNU GPLv3
import numpy as np
class TFIModel:
    """Simple class generating the Hamiltonian of the transverse-field Ising model.
    The Hamiltonian reads
    .. math ::
        H=-J \\sum_{i} \\sigma^x_i \\sigma^x_{i+1} - g \\sum_{i} \\sigma^z_i
    Parameters
    ----------
    L : int
        Number of sites.
```

```
    J, g : float
        Coupling parameters of the above defined Hamiltonian.
    bc : 'infinite', 'finite'
    Boundary conditions.
    Attributes
    ------
    L : int
        Number of sites.
    bc : 'infinite', 'finite'
        Boundary conditions.
    sigmax, sigmay, sigmaz, id :
        Local operators, namely the Pauli matrices and identity.
    H_bonds : list of np.Array[ndim=4]
        The Hamiltonian written in terms of local 2-site operators, ``H = sum_i H_
\hookrightarrowbonds[i]``.
        Each ``H_bonds[i]`` has (physical) legs (i out, (i+1) out, i in, (i+1) in),
        in short ``i j i* j*``.
    H_mpo : lit of np.Array[ndim=4]
        The Hamiltonian written as an MPO.
        Each ``H_mpo[i]`` has legs (virutal left, virtual right, physical out,七
\hookrightarrowphysical in),
        in short ``wL wR i i*``.
    """
    def __init__(self, L, J, g, bc='finite'):
        assert bc in ['finite', 'infinite']
        self.L, self.d, self.bc = L, 2, bc
        self.J, self.g = J, g
        self.sigmax = np.array([[0., 1.], [1., 0.]])
        self.sigmay = np.array([[0., -1j], [1j, 0.]])
        self.sigmaz = np.array([[1., 0.], [0., -1.]])
        self.id = np.eye(2)
        self.init_H_bonds()
        self.init_H_mpo()
    def init_H_bonds(self):
        """Initialize `H_bonds` hamiltonian.
        Called by __init__().
        """
        sx, sz, id = self.sigmax, self.sigmaz, self.id
        d = self.d
        nbonds = self.L - 1 if self.bc == 'finite' else self.L
        H_list = []
        for i in range(nbonds):
            gL = gR = 0.5 * self.g
            if self.bc == 'finite':
                if i == 0:
                    gL = self.g
                if i + 1 == self.L - 1:
                    gR = self.g
            H_bond = -self.J * np.kron(sx, sx) - gL * np.kron(sz, id) - gR * np.
Gron(id, sz)
            # H_bond has legs ``i, j, i*, j*`
            H_list.append(np.reshape(H_bond, [d, d, d, d]))
        self.H_bonds = H_list
```

```
# (note: not required for TEBD)
def init_H_mpo(self):
        """Initialize `H_mpo` Hamiltonian.
        Called by__init__().
        " ""
        w_list = []
        for i in range(self.L):
            w = np.zeros((3, 3, self.d, self.d), dtype=float)
            w[0, 0] = w[2, 2] = self.id
            w[0, 1] = self.sigmax
            w[0, 2] = -self.g * self.sigmaz
            w[1, 2] = -self.J * self.sigmax
            w_list.append(w)
        self.H_mpo = w_list
```


### 11.1.3 c_tebd.py

on github.

```
"""Toy code implementing the time evolving block decimation (TEBD)."""
# Copyright 2018-2021 TeNPy Developers, GNU GPLv3
import numpy as np
from scipy.linalg import expm
from a_mps import split_truncate_theta
def calc_U__bonds(H_bonds, dt):
    """Given the H_bonds, calculate ``U_bonds[i] = expm(-dt*H_bonds[i])``.
    Each local operator has legs (i out, (i+1) out, i in, (i+1) in), in short ``i ju
\hookrightarrowi* j*``.
    Note that no imaginary 'i' is included, thus real `dt' means 'imaginary time'_
\hookrightarrowevolution!
    """
    d = H_bonds[0].shape[0]
    U_bonds = []
    for H in H_bonds:
        H = np.reshape(H, [d * d, d * d])
        U = expm(-dt * H)
        U_bonds.append(np.reshape(U, [d, d, d, d]))
    return U__bonds
def run_TEBD(psi, U_bonds, N_steps, chi_max, eps):
    """Evolve for `N_steps` time steps with TEBD."""
    Nbonds = psi.L - 1 if psi.bc == 'finite' else psi.L
    assert len(U_bonds) == Nbonds
    for n in range(N_steps):
        for k in [0, 1]: # even, odd
            for i__bond in range(k, Nbonds, 2):
                update_bond(psi, i_bond, U_bonds[i_bond], chi_max, eps)
    # done
```

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```
def update_bond(psi, i, u_bond, chi_max, eps):
    """Apply `U_bond` acting on i,j=(i+1) to `psi`."""
    j = (i + 1) % psi.L
    # construct theta matrix
    theta = psi.get_theta2(i) # vL i j vR
    # apply U
    Utheta = np.tensordot(U_bond, theta, axes=([2, 3], [1, 2])) # i j [i*] [j*], vL_
\hookrightarrow[i] [j] vR
    Utheta = np.transpose(Utheta, [2, 0, 1, 3]) # vL i j vR
    # split and truncate
    Ai, Sj, Bj = split_truncate_theta(Utheta, chi_max, eps)
    # put back into MPS
    Gi = np.tensordot(np.diag(psi.Ss[i]**(-1)), Ai, axes=[1, 0]) # vL [vL*], [vL] iv
GC
    psi.Bs[i] = np.tensordot(Gi, np.diag(Sj), axes=[2, 0]) # vL i [vC], [vC] vC
    psi.Ss[j] = Sj # vC
    psi.Bs[j] = Bj # vC j vR
def example_TEBD_gs_tf_ising_finite(L, g):
    print("finite TEBD, imaginary time evolution, transverse field Ising")
    print("L={L:d},g={g:.2f}".format (L=L, g=g))
    import a_mps
    import b_model
    M = b_model.TFIModel(L=L, J=1., g=g, bc='finite')
    psi = a_mps.init_FM_MPS(M.L, M.d, M.bc)
    for dt in [0.1, 0.01, 0.001, 1.e-4, 1.e-5]:
        U_bonds = calc_U_bonds(M.H_bonds, dt)
        run_TEBD(psi, U_bonds, N_steps=500, chi_max=30, eps=1.e-10)
        E = np.sum(psi.bond_expectation_value(M.H_bonds))
        print("dt = {dt:.5f}: E = {E:.13f}".format(dt=dt, E=E))
    print("final bond dimensions: ", psi.get_chi())
    mag_x = np.sum(psi.site_expectation_value(M.sigmax))
    mag_z = np.sum(psi.site_expectation_value(M.sigmaz))
    print("magnetization in X = {mag_x:.5f}".format(mag_x=mag_x))
    print("magnetization in Z = {mag_z:.5f}".format(mag_z=mag_z))
    if L < 20: # compare to exact result
        from tfi_exact import finite_gs_energy
        E_exact = finite_gs_energy(L, 1., g)
        print("Exact diagonalization: E = {E:.l3f}".format(E=E_exact))
        print("relative error: ", abs((E - E_exact) / E_exact))
    return E, psi, M
def example_TEBD_gs_tf_ising_infinite(g):
    print("infinite TEBD, imaginary time evolution, transverse field Ising")
    print("g={g:.2f}".format (g=g))
    import a_mps
    import b_model
    M = b_model.TFIModel(L=2, J=1., g=g, bc='infinite')
    psi = a_mps.init_FM_MPS(M.L, M.d, M.bc)
    for dt in [0.1, 0.01, 0.001, 1.e-4, 1.e-5]:
        U_bonds = calc_U_bonds (M.H_bonds, dt)
        run_TEBD(psi, U_bonds, N_steps=500, chi_max=30, eps=1.e-10)
        E = np.mean(psi.bond_expectation_value(M.H_bonds))
        print("dt = {dt:.5f}: E (per site) = {E:.13f}".format(dt=dt, E=E))
```

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```
    print("final bond dimensions: ", psi.get_chi())
    mag_x = np.mean(psi.site_expectation_value(M.sigmax))
    mag_z = np.mean(psi.site_expectation_value(M.sigmaz))
    print("<sigma_x> = {mag_x:.5f}".format(mag_x=mag_x))
    print("<sigma_z> = {mag_z:.5f}".format(mag_z=mag_z))
    print("correlation length:", psi.correlation_length())
    # compare to exact result
    from tfi_exact import infinite_gs_energy
    E_exact = infinite_gs_energy(1., g)
    print("Analytic result: E (per site) = {E:.l3f}".format(E=E_exact))
    print("relative error: ", abs((E - E_exact) / E_exact))
    return E, psi, M
def example_TEBD_tf_ising_lightcone(L, g, tmax, dt):
    print("finite TEBD, real time evolution, transverse field Ising")
    print("L={L:d}, g={g:.2f}, tmax={tmax:.2f}, dt={dt:.3f}".format(L=L,g=g, b
\hookrightarrowtmax=tmax, dt=dt))
    # find ground state with TEBD or DMRG
    # E, psi, M = example_TEBD_gS_tf_ising_finite(L, g)
    from d_dmrg import example_DMRG_tf_ising_finite
    E, psi, M = example_DMRG_tf_ising_finite(L, g)
    iO = L // 2
    # apply sigmaz on site iO
    SzB = np.tensordot(M.sigmaz, psi.Bs[i0], axes=[1, 1]) # i [i*], vL [i] vR
    psi.Bs[i0] = np.transpose(SzB, [1, 0, 2]) # vL i vR
    U_bonds = calc_U_bonds(M.H_bonds, 1.j * dt) # (imaginary dt -> realtime_
Gevolution)
    S = [psi.entanglement_entropy()]
    Nsteps = int(tmax / dt + 0.5)
    for n in range(Nsteps):
        if abs((n * dt + 0.1) % 0.2 - 0.1) < 1.e-10:
                print("t = {t:.2f}, chi =".format(t=n * dt), psi.get_chi())
            run_TEBD(psi, U_bonds, 1, chi_max=50, eps=1.e-10)
            S.append(psi.entanglement_entropy())
    import matplotlib.pyplot as plt
    plt.figure()
    plt.imshow(S[::-1],
                vmin=0.,
                aspect='auto',
                interpolation='nearest',
                extent=(0, L - 1., -0.5 * dt, (Nsteps + 0.5) * dt))
    plt.xlabel('site $i$')
    plt.ylabel('time $t/J$')
    plt.ylim(0., tmax)
    plt.colorbar().set_label('entropy $S$')
    filename = 'c_tebd_lightcone_{g:.2f}.pdf'.format(g=g)
    plt.savefig(filename)
    print("saved " + filename)
if name == " main ":
    example_TEBD_gs_tf_ising_finite(L=10, g=1.)
    print("-" * 100)
    example_TEBD_gs_tf_ising_infinite(g=1.5)
    print("-" * 100)
    example_TEBD_tf_ising_lightcone(L=20, g=1.5, tmax=3., dt=0.01)
```


### 11.1.4 d_dmrg.py

on github.

```
"""Toy code implementing the density-matrix renormalization group (DMRG)."""
# Copyright 2018-2021 TeNPy Developers, GNU GPLv3
import numpy as np
from a_mps import split_truncate_theta
import scipy.sparse
import scipy.sparse.linalg.eigen.arpack as arp
class SimpleHeff(scipy.sparse.linalg.LinearOperator):
    """Class for the effective Hamiltonian.
    To be diagonalized in `SimpleDMRGEnginge.update_bond`. Looks like this::
    lovL* 
    """
    def __init__(self, LP, RP, W1, W2):
        self.LP = LP # vL wL* vL*
        self.RP = RP # vR* wR* vR
        self.W1 = W1 # wL wC i i*
        self.W2 = W2 # wC wR j j*
        chi1, chi2 = LP.shape[0], RP.shape[2]
        d1, d2 = W1.shape[2], W2.shape[2]
        self.theta_shape = (chi1, d1, d2, chi2) # vL i j vR
        self.shape = (chi1 * d1 * d2 * chi2, chi1 * d1 * d2 * chi2)
        self.dtype = W1.dtype
    def __matvec(self, theta):
        """Calculate |theta'> = H_eff |theta>.
        This function is used by :func:scipy.sparse.linalg.eigen.arpack.eigsh` to_
\hookrightarrowdiagonalize
    the effective Hamiltonian with a Lanczos method, withouth generating the full_
\hookrightarrowmatrix."""
    x = np.reshape(theta, self.theta_shape) # vL i j vR
    x = np.tensordot(self.LP, x, axes=(2, 0)) # vL wL* [vL*], [vL] i j vR
    x = np.tensordot(x, self.W1, axes=([1, 2], [0, 3])) # vL [wL*] [i] j vR, b
\hookrightarrow[wL] WC i [i*]
    x = np.tensordot(x, self.W2, axes=([3, 1], [0, 3])) # vL [j] VR [WC] i, [WC],
\hookrightarrowwR j [j*]
    x = np.tensordot(x, self.RP, axes=([1, 3], [0, 1])) # vL [vR] i [wR] j,u
\hookrightarrow[vR*] [WR*] vR
    x = np.reshape(x, self.shape[0])
    return x
class SimpleDMRGEngine:
    """DMRG algorithm, implemented as class holding the necessary data.
```

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```
    Parameters
    psi, model, chi_max, eps:
        See attributes
    Attributes
    psi : SimpleMPS
    The current ground-state (approximation).
    model :
    The model of which the groundstate is to be calculated.
    chi_max, eps:
        Truncation parameters, see :func:`a_mps.split_truncate_theta`.
    LPS, RPS : list of np.Array[ndim=3]
    Left and right parts ("environments") of the effective Hamiltonian.
        ``IPs[i]`` is the contraction of all parts left of site `i` in the network
\hookrightarrow<psi|H|psi>``,
    and similar ``RPs[i]`` for all parts right of site `i`.
    Each ``LPs[i]`` has legs ``vL wL* vL*``, ``RPS[i]`` has legs ``vR* wR* vR``
    " ""
    def __init__(self, psi, model, chi_max, eps):
        assert psi.L == model.L and psi.bc == model.bc # ensure compatibility
        self.H_mpo = model.H_mpo
        self.psi = psi
        self.LPs = [None] * psi.L
        self.RPs = [None] * psi.L
        self.chi_max = chi_max
        self.eps = eps
        # initialize left and right environment
        D = self.H_mpo[0].shape[0]
        chi = psi.Bs[0].shape[0]
        LP = np.zeros([chi, D, chi], dtype=float) # vL wL* vL*
        RP = np.zeros([chi, D, chi], dtype=float) # vR* wR* vR
        LP[:, 0, :] = np.eye(chi)
        RP[:, D - 1, :] = np.eye(chi)
        self.LPs[0] = LP
        self.RPs[-1] = RP
        # initialize necessary RPs
        for i in range(psi.L - 1, 1, -1):
        self.update_RP(i)
    def sweep(self):
    # sweep from left to right
    for i in range(self.psi.nbonds - 1):
        self.update_bond(i)
    # sweep from right to left
    for i in range(self.psi.nbonds - 1, 0, -1):
        self.update_bond(i)
    def update_bond(self, i):
    j = (i + 1) % self.psi.L
    # get effective Hamiltonian
    Heff = SimpleHeff(self.LPs[i], self.RPs[j], self.H_mpo[i], self.H_mpo[j])
    # Diagonalize Heff, find ground state `theta`
    theta0 = np.reshape(self.psi.get_theta2(i), [Heff.shape[0]]) # initial guess
    e, v = arp.eigsh(Heff, k=1, which='SA', return_eigenvectors=True, v0=theta0)
```

```
    theta = np.reshape(v[:, 0], Heff.theta_shape)
    # split and truncate
    Ai, Sj, Bj = split_truncate_theta(theta, self.chi_max, self.eps)
    # put back into MPS
    Gi = np.tensordot(np.diag(self.psi.Ss[i]**(-1)), Ai, axes=[1, 0]) # vL [vL*],
\hookrightarrow [vL] i vC
    self.psi.Bs[i] = np.tensordot(Gi, np.diag(Sj), axes=[2, 0]) # vL i [vC],\smile
G[vC*] vC
    self.psi.Ss[j] = Sj # vC
    self.psi.Bs[j] = Bj # vC j vR
    self.update_LP(i)
    self.update_RP(j)
    def update_RP(self, i):
    """Calculate RP right of site `i-I` from RP right of site `i`."""
    j = (i - 1) % self.psi.L
    RP = self.RPs[i] # vR* wR* vR
    B = self.psi.Bs[i] # vL i vR
    BC = B.conj() # vL* i* vR*
    W = self.H_mpo[i] # wL wR i i*
    RP = np.tensordot(B, RP, axes=[2, 0]) # vL i [vR], [vR*] wR* vR
    RP = np.tensordot(RP, W, axes=[[1, 2], [3, 1]]) # vL [i] [wR*] vR, wL [wR] i_
G[i*]
    RP = np.tensordot(RP, BC, axes=[[1, 3], [2, 1]]) # vL [vR] wL [i], vL* [i*]_
\hookrightarrow vR*]
    self.RPs[j] = RP # vL wL vL* (== vR* wR* vR on site i-1)
    def update_LP(self, i):
    """Calculate LP left of site `i+1` from LP left of site `i`."""
    j = (i + 1) % self.psi.L
    LP = self.LPs[i] # vL wL vL*
    B = self.psi.Bs[i] # vL i vR
    G = np.tensordot(np.diag(self.psi.Ss[i]), B, axes=[1, 0]) # vL [vL*], [vL] iv
vR
    A = np.tensordot(G, np.diag(self.psi.Ss[j]**-1), axes=[2, 0]) # vL i [vR],u
G[vR*] vR
    Ac = A.conj() # vL* i* vR*
    W = self.H_mpo[i] # wL wR i i*
    LP = np.tensordot(LP, A, axes=[2, 0]) # vL wL* [vL*], [VL] i vR
    LP = np.tensordot(W, LP, axes=[[0, 3], [1, 2]]) # [wL] wR i [i*], vL [wL*]_
\hookrightarrow[i] vR
    LP = np.tensordot(Ac, LP, axes=[[0, 1], [2, 1]]) # [vL*] [i*] vR*, wR [i]u
G[VL] vR
    self.LPs[j] = LP # vR* wR vR (== vL wL* vL* on site i+1)
def example_DMRG_tf_ising_finite(L, g):
    print("finite DMRG, transverse field Ising")
    print("L={L:d}, g={g:.2f}".format(L=L, g=g))
    import a_mps
    import b_model
    M = b_model.TFIModel(L=L, J=1., g=g, bc='finite')
    psi = a_mps.init_FM_MPS(M.L, M.d, M.bc)
    eng = SimpleDMRGEngine(psi, M, chi_max=30, eps=1.e-10)
    for i in range(10):
        eng.sweep()
        E = np.sum(psi.bond_expectation_value(M.H_bonds))
```

```
        print("sweep {i:2d}: E = {E:.13f}".format(i=i + 1, E=E))
    print("final bond dimensions: ", psi.get_chi())
    mag_x = np.sum(psi.site_expectation_value(M.sigmax))
    mag_z = np.sum(psi.site_expectation_value(M.sigmaz))
    print("magnetization in X = {mag_x:.5f}".format(mag_x=mag_x))
    print("magnetization in Z = {mag_z:. 5f}".format(mag_z=mag_z))
    if L < 20: # compare to exact result
        from tfi_exact import finite_gs_energy
        E_exact = finite_gs_energy(L, 1., g)
        print("Exact diagonalization: E = {E:.l3f}".format(E=E_exact))
        print("relative error: ", abs((E - E_exact) / E_exact))
    return E, psi, M
def example_DMRG_tf_ising_infinite(g):
    print("infinite DMRG, transverse field Ising")
    print("g={g:.2f}".format(g=g))
    import a_mps
    import b_model
    M = b_model.TFIModel(L=2, J=1., g=g, bc='infinite')
    psi = a_mps.init_EM_MPS(M.L, M.d, M.bc)
    eng = SimpleDMRGEngine(psi, M, chi_max=20, eps=1.e-14)
    for i in range(20):
        eng.sweep()
        E = np.mean(psi.bond_expectation_value(M.H_bonds))
        print("sweep {i:2d}: E (per site) = {E:.l3f}".format(i=i + 1, E=E))
    print("final bond dimensions: ", psi.get_chi())
    mag_x = np.mean(psi.site_expectation_value(M.sigmax))
    mag_z = np.mean(psi.site_expectation_value(M.sigmaz))
    print("<sigma_x> = {mag_x:.5f}".format(mag_x=mag_x))
    print("<sigma_z> = {mag_z:.5f}".format(mag_z=mag_z))
    print("correlation length:", psi.correlation_length())
    # compare to exact result
    from tfi_exact import infinite_gs_energy
    E_exact = infinite_gs_energy(1., g)
    print("Analytic result: E (per site) = {E:.I3f}".format(E=E_exact))
    print("relative error: ", abs((E - E_exact) / E_exact))
    return E, psi, M
if ___name___ == "___main__"":
    example_DMRG_tf_ising_finite(L=10, g=1.)
    print("-" * 100)
    example_DMRG_tf_ising_infinite(g=1.5)
```


### 11.1.5 tfi_exact.py

on github.

```
"""Provides exact ground state energies for the transverse field ising model for,
Comparison.
The Hamiltonian reads
.. math : :
    H=-J \\sum_{i} \\sigma^x_i \\sigma^x_{i+1} - g \\sum_{i} \\sigma^z_i
```

```
" " "
# Copyright 2019-2021 TeNPy Developers, GNU GPLV3
import numpy as np
import scipy.sparse as sparse
import scipy.sparse.linalg.eigen.arpack as arp
import warnings
import scipy.integrate
def finite_gs__energy(L, J, g):
        """For comparison: obtain ground state energy from exact diagonalization.
    Exponentially expensive in L, only works for small enough `L` <~ 20.
    " " "
    if L >= 20:
            warnings.warn("Large L: Exact diagonalization might take a long time!")
    # get single site operaors
    sx = sparse.csr_matrix(np.array([[0., 1.], [1., 0.]]))
    sz = sparse.csr_matrix(np.array([[1., 0.], [0., -1.]]))
    id = sparse.csr_matrix(np.eye(2))
    sx_list = [] # sx_list[i] = kron([id, id, ..., id, sx, id, .... id])
    sz_list = []
    for i_site in range(L):
            x_ops = [id] * L
            z_ops = [id] * L
            x_ops[i_site] = sx
            z_ops[i_site] = sz
            X = x_ops[0]
            Z = z_ops[0]
            for j in range(1, L):
                X = sparse.kron(X, x_ops[j], 'csr')
                Z = sparse.kron(Z, z_ops[j], 'csr')
            sx_list.append(X)
            sz_list.append(Z)
    H_xx = sparse.csr_matrix((2**L, 2**L))
    H_z = sparse.csr_matrix((2**L, 2**L))
    for i in range(L - 1):
            H_xx = H_xx + sx_list[i] * sx_list[(i + 1) % L]
    for i in range(L):
        H_z = H_z + sz_list[i]
    H = -J * H_xx - g * H_z
    E, V = arp.eigsh(H, k=1, which='SA', return_eigenvectors=True, ncv=20)
    return E[0]
def infinite_gs_energy(J, g):
    """For comparison: Calculate groundstate energy density from analytic formula.
    The analytic formula stems from mapping the model to free fermions, see P. Pfeuty,
GThe one-
    dimensional Ising model with a transverse field, Annals of Physics 57, p. 79,
G(1970). Note that
    we use Pauli matrices compared this reference using spin-1/2 matrices and replace
->the sum_k ->
    integral dk/2pi to obtain the result in the N -> infinity limit.
    """
```

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```
    def f(k, lambda_):
        return np.sqrt(1 + lambda_**2 + 2 * lambda_ * np.cos(k))
    E0_exact = -g / (J * 2. * np.pi) * scipy.integrate.quad(f, -np.pi, np.pi, args=(J |
\hookrightarrow/ g, )) [0]
    return E0_exact
```


### 11.2 Python scripts

These example scripts illustrate the very basic interface for calling TeNPy. They are included in the [TeNPySource] repository in the folder examples/, we include them here in the documentation for reference. You need to install TeNPy to call them (see Installation instructions), but you can copy them anywhere before execution. (Some scripts include other files from the same folder, though; copy those as well.)

### 11.2.1 a_np_conserved.py

on github.

```
"""An example code to demonstrate the usage of :class:`~tenpy.linalg.np_conserved.
\hookrightarrowArray`.
This example includes the following steps:
1) create Arrays for an Neel MPS
2) create an MPO representing the nearest-neighbour AFM Heisenberg Hamiltonian
3) define 'environments' left and right
4) contract MPS and MPO to calculate the energy
5) extract two-site hamiltonian ``H2`` from the MPO
6) calculate ``exp(-1.j*dt*H2)`` by diagonalization of H2
7) apply ``exp(H2)`` to two sites of the MPS and truncate with svd
Note that this example uses only np_conserved, but no other modules.
Compare it to the example `b_mps.py`,
which does the same steps using a few predefined classes like MPS and MPO.
" ""
# Copyright 2018-2021 TeNPy Developers, GNU GPLv3
import tenpy.linalg.np_conserved as npc
import numpy as np
# model parameters
Jxx, Jz = 1., 1.
L=20
dt = 0.1
cutoff = 1.e-10
print("Jxx={Jxx}, Jz={Jz}, L={L:d}".format(Jxx=Jxx, Jz=Jz, L=L))
print("1) create Arrays for an Neel MPS")
# vL ->--B-->- vR
# l
# l
```

```
# p
# create a ChargeInfo to specify the nature of the charge
chinfo = npc.ChargeInfo([1], ['2*Sz']) # the second argument is just a descriptive,
@ame
# create LegCharges on physical leg and even/odd bonds
p_leg = npc.LegCharge.from_qflat(chinfo, [[1], [-1]]) # charges for up, down
v_leg_even = npc.LegCharge.from_qflat(chinfo, [[0]])
v_leg_odd = npc.LegCharge.from_qflat(chinfo, [[1]])
B_even = npc.zeros([v_leg_even, v_leg_odd.conj(), p_leg],
    labels=['vL', 'vR', 'p']) # virtual left/right, physical
B_odd = npc.zeros([v_leg_odd, v_leg_even.conj(), p_leg], labels=['vL', 'vR', 'p'])
B_even[0, 0, 0] = 1. # up
B_odd[0, 0, 1] = 1. # down
Bs = [B_even, B_odd] * (L // 2) + [B_even] * (L % 2) # (right-canonical)
Ss = [np.ones(1)] * L # Ss[i] are singular values between Bs[i-l] and Bs[i]
# Side remark:
# An MPS is expected to have non-zero entries everywhere compatible with the charges.
# In general, we recommend to use `sort_legcharge` (or `as_completely_blocked`)
# to ensure complete blocking. (But the code will also work, if you don't do it.)
# The drawback is that this might introduce permutations in the indices of singleu
\hookrightarrowlegs,
# which you have to keep in mind when converting dense numpy arrays to and from npc.
AArrays.
print("2) create an MPO representing the AFM Heisenberg Hamiltonian")
```



```
# create physical spin-1/2 operators Sz, St, S-
\hookrightarrow', 'p*'])
Sp = npc.Array.from_ndarray([[0., l.], [0., 0.]], [p_leg, p_leg.conj()], labels=['p',
@'p*'])
Sm = npc.Array.from_ndarray([[0., 0.], [1., 0.]], [p_leg, p_leg.conj()], labels=['p',
\hookrightarrow'p*'])
Id = npc.eye_like(Sz, labels=Sz.get_leg_labels()) # identity
mpo_leg = npc.LegCharge.from_qflat(chinfo, [[0], [2], [-2], [0], [0]])
W_grid = [[Id, Sp, Sm, Sz, None ],
    [None, None, None, None, 0.5 * Jxx * Sm],
    [None, None, None, None, 0.5 * Jxx * Sp],
    [None, None, None, None, Jz * Sz ],
    [None, None, None, None, Id ]] # yapf:disable
```

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```
            (continued from previous page)
W = npc.grid_outer(W_grid, [mpo_leg, mpo_leg.conj()], grid_labels=['wL', 'wR'])
# wL/wR = virtual left/right of the MPO
Ws = [W] * L
print("3) define 'environments' left and right")
# .---->- vR VL ->----.
# 1
# envL->- wR wL ->-envR
# 1 |
# .---->- vR* VL*->-----
envL = npc.zeros([W.get_leg('wL').conj(), Bs[0].get_leg('vL').conj(), Bs[0].get_leg(
G'vL')],
    labels=['wR', 'vR', 'vR*'])
envL[0, :, :] = npc.diag(1., envL.legs[1])
envR = npc.zeros([W.get_leg('wR').conj(), Bs[-1].get_leg('vR').conj(), Bs[-1].get_leg(
\hookrightarrow'vR')],
            labels=['wL', 'vL', 'vL*'])
envR[-1, :, :] = npc.diag(1., envR.legs[1])
print("4) contract MPS and MPO to calculate the energy <psi|H|psi>")
contr = envL
for i in range(L):
    # contr labels: wR, vR, vR*
    contr = npc.tensordot(contr, Bs[i], axes=('vR', 'vL'))
    # wR, vR*, vR, p
    contr = npc.tensordot(contr, Ws[i], axes=(['p', 'wR'], ['p*', 'wL']))
    # vR*, vR, wR, p
    contr = npc.tensordot(contr, Bs[i].conj(), axes=(['p', 'vR*'], ['p*', 'vL*']))
    # vR, wR, vR*
    # note that the order of the legs changed, but that's no problem with labels:
    # the arrays are automatically transposed as necessary
E = npc.inner(contr, envR, axes=(['vR', 'wR', 'vR*'], ['vL', 'wL', 'vL*']))
print("E =", E)
print("5) calculate two-site hamiltonian ``H2`` from the MPO")
# label left, right physical legs with p, q
W0 = W.replace_labels(['p', 'p*'], ['p0', 'p0*'])
W1 = W.replace_labels(['p', 'p*'], ['p1', 'pl*'])
H2 = npc.tensordot(W0, W1, axes=('wR', 'wL')).itranspose(['wL', 'wR', 'p0', 'p1', 'p0*
\hookrightarrow', 'p1*'])
H2 = H2[0, -1] # (If H has single-site terms, it's not that simple anymore)
print("H2 labels:", H2.get_leg_labels())
print("6) calculate exp(H2) by diagonalization of H2")
# diagonalization requires to view H2 as a matrix
H2 = H2.combine_legs([('p0', 'p1'), ('p0*', 'p1*')], qconj=[+1, -1])
print("labels after combine_legs:", H2.get_leg_labels())
E2, U2 = npc.eigh(H2)
print("Eigenvalues of H2:", E2)
U_expE2 = U2.scale_axis(np.exp(-1.j * dt * E2), axis=1) # scale_axis ~= apply an_
\hookrightarrowdiagonal matrix
exp_H2 = npc.tensordot(U_expE2, U2.conj(), axes=(1, 1))
exp_H2.iset_leg_labels(H2.get_leg_labels())
exp_H2 = exp_H2.split_legs() # by default split all legs which are `LegPipe
```

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```
# (this restores the originial labels ['pO', 'p1', 'pO*', 'pl*'] of `H2` in `exp_H2`)
print("7) apply exp(H2) to even/odd bonds of the MPS and truncate with svd")
# (this implements one time step of first order TEBD)
for even_odd in [0, 1]:
    for i in range(even_odd, L - 1, 2):
        B_L = Bs[i].scale_axis(Ss[i], 'vL').ireplace_label('p', 'p0')
        B_R = Bs[i + 1].replace_label('p', 'pl')
        theta = npc.tensordot(B_L, B_R, axes=('vR', 'vL'))
        theta = npc.tensordot(exp_H2, theta, axes=(['p0*', 'p1*'], ['p0', 'p1']))
        # view as matrix for SVD
        theta = theta.combine_legs([('vL', 'p0'), ('p1', 'vR')], new_axes=[0, 1], b
qconj=[+1, -1])
            # now theta has labels '(vL.p0)', '(pl.vR)'
            U, S, V = npc.svd(theta, inner_labels=['vR', 'vL'])
            # truncate
            keep = S > cutoff
            S = S[keep]
            invsq = np.linalg.norm(S)
            Ss[i + 1] = S / invsq
            U = U.iscale_axis(S / invsq, 'vR')
            Bs[i] = U.split_legs('(vL.p0)').iscale_axis(Ss[i]**(-1), 'vL').ireplace_label(
\hookrightarrow'p0', 'p')
    Bs[i + 1] = V.split_legs('(pl.vR)').ireplace_label('pl', 'p')
print("finished")
```


### 11.2.2 b_mps.py

on github.

```
"""Simplified version of `a_np_conserved.py` making use of other classes (like MPS, 
\hookrightarrowMPO).
This example includes the following steps:
1) create Arrays for an Neel MPS
2) create an MPO representing the nearest-neighbour AFM Heisenberg Hamiltonian
3) define 'environments' left and right
4) contract MPS and MPO to calculate the energy
5) extract two-site hamiltonian ``H2`` from the MPO
6) calculate ``exp(-1.j*dt*H2)`` by diagonalization of H2
7) apply ``exp(H2)`` to two sites of the MPS and truncate with svd
Note that this example performs the same steps as `a_np_conserved.py`,
but makes use of other predefined classes except npc.
" ""
# Copyright 2018-2021 TeNPy Developers, GNU GPLv3
import tenpy.linalg.np_conserved as npc
import numpy as np
# some more imports
from tenpy.networks.site import SpinHalfSite
from tenpy.models.lattice import Chain
from tenpy.networks.mps import MPS
from tenpy.networks.mpo import MPO, MPOEnvironment
```

```
from tenpy.algorithms.truncation import svd_theta
# model parameters
Jxx, Jz = 1., 1.
L = 20
dt = 0.1
cutoff = 1.e-10
print("Jxx={Jxx}, Jz={Jz}, L={L:d}".format(Jxx=Jxx, Jz=Jz, L=L))
print("1) create Arrays for an Neel MPS")
site = SpinHalfSite(conserve='Sz') # predefined charges and Sp,Sm,Sz operators
p_leg = site.leg
chinfo = p_leg.chinfo
# make lattice from unit cell and create product state MPS
lat = Chain(L, site, bc_MPS='finite')
state = ["up", "down"] * (L // 2) + ["up"] * (L % 2) # Neel state
print("state = ", state)
psi = MPS.from_product_state(lat.mps_sites(), state, lat.bc_MPS)
print("2) create an MPO representing the AFM Heisenberg Hamiltonian")
# predefined physical spin-1/2 operators Sz, S+, S-
Sz, Sp, Sm, Id = site.Sz, site.Sp, site.Sm, site.Id
mpo_leg = npc.LegCharge.from_qflat(chinfo, [[0], [2], [-2], [0], [0]])
W_grid = [[Id, Sp, Sm, Sz, None ],
    [None, None, None, None, 0.5 * Jxx * Sm],
    [None, None, None, None, 0.5 * Jxx * Sp],
    [None, None, None, None, Jz * Sz ],
    [None, None, None, None, Id ]] # yapf:disable
W = npc.grid_outer(W_grid, [mpo_leg, mpo_leg.conj()], grid_labels=['wL', 'wR'])
# wL/wR = virtual left/right of the MPO
Ws = [W] * L
Ws[0] = W[:1, :]
Ws[-1] = W[:, -1:]
H = MPO(psi.sites, Ws, psi.bc, IdL=0, IdR=-1)
print("3) define 'environments' left and right")
# this is automatically done during initialization of MPOEnvironment
env = MPOEnvironment (psi, H, psi)
envL = env.get_LP(0)
envR = env.get_RP(L - 1)
print("4) contract MPS and MPO to calculate the energy <psi|H|psi>")
E = env.full_contraction(L - 1)
print("E =", E)
print("5) calculate two-site hamiltonian ``H2`` from the MPO")
# label left, right physical legs with p, q
W0 = H.get_W(0).replace_labels(['p', 'p*'], ['p0', 'p0*'])
W1 = H.get_W(1).replace_labels(['p', 'p*'], ['p1', 'p1*'])
H2 = npc.tensordot(w0, W1, axes=('wR', 'wL')).itranspose(['wL', 'wR', 'p0', 'p1', 'p0*
↔', 'p1*'])
```

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```
H2 = H2[H.IdL[0], H.IdR[2]] # (If H has single-site terms, it's not that simple
->anymore)
print("H2 labels:", H2.get_leg_labels())
print("6) calculate exp(H2) by diagonalization of H2")
# diagonalization requires to view H2 as a matrix
H2 = H2.combine_legs([('p0', 'p1'), ('p0*', 'p1*')], qconj=[+1, -1])
print("labels after combine_legs:", H2.get_leg_labels())
E2, U2 = npc.eigh(H2)
print("Eigenvalues of H2:", E2)
U_expE2 = U2.scale_axis(np.exp(-1.j * dt * E2), axis=1) # scale_axis ~= apply a
\hookrightarrowdiagonal matrix
exp_H2 = npc.tensordot(U_expE2, U2.conj(), axes=(1, 1))
exp_H2.iset_leg_labels(H2.get_leg_labels())
exp_H2 = exp_H2.split_legs() # by default split all legs which are `LegPipe`
# (this restores the originial labels ['po', 'p1', 'po*', 'pl*'] of `H2` in `exp_H2`)
# alternative way: use :func:`~tenpy.linalg.np_conserved.expm`
exp_H2_alternative = npc.expm(-1.j * dt * H2).split_legs()
assert (npc.norm(exp_H2_alternative - exp_H2) < 1.e-14)
print("7) apply exp(H2) to even/odd bonds of the MPS and truncate with svd")
# (this implements one time step of first order TEBD)
trunc_par = {'svd_min': cutoff, 'trunc_cut': None}
for even_odd in [0, 1]:
    for i in range(even_odd, L - 1, 2):
        theta = psi.get_theta(i, 2) # handles canonical form (i.e. scaling with 'S')
        theta = npc.tensordot(exp_H2, theta, axes=(['p0*', 'p1*'], ['p0', 'p1']))
        # view as matrix for SVD
        theta = theta.combine_legs([('vL', 'p0'), ('p1', 'vR')], new_axes=[0, 1], ५
Gconj=[+1, -1])
        # now theta has labels '(vL.p0)', '(pl.vR)'
        U, S, V, err, invsq = svd_theta(theta, trunc_par, inner_labels=['vR', 'vL'])
        psi.set_SR(i, S)
        A_L = U.split_legs('(vL.p0)').ireplace_label('p0', 'p')
        B_R = V.split_legs('(pl.vR)').ireplace_label('pl', 'p')
        psi.set_B(i, A_L, form='A') # left-canonical form
        psi.set_B(i + 1, B_R, form='B') # right-canonical form
print("finished")
```


### 11.2.3 c_tebd.py

on github.

```
"""Example illustrating the use of TEBD in tenpy.
The example functions in this class do the same as the ones in `toycodes/c_tebd.py`,
\hookrightarrowbut make use
of the classes defined in tenpy.
"""
# Copyright 2018-2021 TeNPy Developers, GNU GPLv3
import numpy as np
from tenpy.networks.mps import MPS
```

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```
from tenpy.models.tf_ising import TFIChain
from tenpy.algorithms import tebd
def example_TEBD_gs_tf_ising_finite(L, g):
    print("finite TEBD, imaginary time evolution, transverse field Ising")
    print("L={L:d},g={g:.2f}".format(L=L, g=g))
    model_params = dict(L=L, J=1., g=g, bc_MPS='finite', conserve=None)
    M = TFIChain(model_params)
    product_state = ["up"] * M.lat.N_sites
    psi = MPS.from_product_state(M.lat.mps_sites(), product_state, bc=M.lat.bc_MPS)
    tebd_params = {
        'order': 2,
        'delta_tau_list': [0.1, 0.01, 0.001, 1.e-4, 1.e-5],
        'N_steps': 10,
        'max_error_E': 1.e-6,
        'trunc_params': {
            'chi_max': 30,
            'svd_min': 1.e-10
        },
    }
    eng = tebd.TEBDEngine(psi, M, tebd_params)
    eng.run_GS() # the main work...
    # expectation values
    E = np.sum(M.bond_energies(psi)) # M.bond_energies() works only a for_
\hookrightarrowNearestNeighborModel
    # alternative: directly measure E2 = np.sum(psi.expectation_value(M.H_bond[1:]))
    print("E = {E:.l3f}".format(E=E))
    print("final bond dimensions: ", psi.chi)
    mag_x = np.sum(psi.expectation_value("Sigmax"))
    mag_z = np.sum(psi.expectation_value("Sigmaz"))
    print("magnetization in X = {mag_x:.5f}".format(mag_x=mag_x))
    print("magnetization in Z = {mag_z:.5f}".format(mag_z=mag_z))
    if L < 20: # compare to exact result
        from tfi_exact import finite_gs_energy
        E_exact = finite_gs_energy(L, 1., g)
        print("Exact diagonalization: E = {E:.l3f}".format(E=E_exact))
        print("relative error: ", abs((E - E_exact) / E_exact))
    return E, psi, M
def example_TEBD_gs_tf_ising_infinite(g):
    print("infinite TEBD, imaginary time evolution, transverse field Ising")
    print("g={g:.2f}".format (g=g))
    model_params = dict(L=2, J=1., g=g, bc_MPS='infinite', conserve=None)
    M = TFIChain(model_params)
    product_state = ["up"] * M.lat.N_sites
    psi = MPS.from_product_state(M.lat.mps_sites(), product_state, bc=M.lat.bc_MPS)
    tebd_params = {
        'order': 2,
        'delta_tau_list': [0.1, 0.01, 0.001, 1.e-4, 1.e-5],
        'N_steps': 10,
        'max_error_E': 1.e-8,
        'trunc_params': {
            'chi_max': 30,
            'svd_min': 1.e-10
```

```
        },
    }
    eng = tebd.TEBDEngine(psi, M, tebd_params)
    eng.run_GS() # the main work...
    E = np.mean(M.bond_energies(psi)) # M.bond_energies() works only a for_
\hookrightarrowNearestNeighborModel
    # alternative: directly measure E2 = np.mean(psi.expectation_value(M.H_bond))
    print("E (per site) = {E:.I3f}".format (E=E))
    print("final bond dimensions: ", psi.chi)
    mag_x = np.mean(psi.expectation_value("Sigmax"))
    mag_z = np.mean(psi.expectation_value("Sigmaz"))
    print("<sigma_x> = {mag_x:.5f}".format(mag_x=mag_x))
    print("<sigma_z> = {mag_z:.5f}".format(mag_z=mag_z))
    print("correlation length:", psi.correlation_length())
    # compare to exact result
    from tfi_exact import infinite_gs_energy
    E_exact = infinite_gs_energy(1., g)
    print("Analytic result: E (per site) = {E:.I3f}".format(E=E_exact))
    print("relative error: ", abs((E - E_exact) / E_exact))
    return E, psi, M
def example_TEBD_tf_ising_lightcone(L, g, tmax, dt):
    print("finite TEBD, real time evolution")
    print("L={L:d}, g={g:.2f}, tmax={tmax:.2f}, dt={dt:.3f}".format(L=L, g=g, ,
\hookrightarrowtmax=tmax, dt=dt))
    # find ground state with TEBD or DMRG
    # E, psi, M = example_TEBD_gS_tf_ising_finite(L, g)
    from d_dmrg import example_DMRG_tf_ising_finite
    print("(run DMRG to get the groundstate)")
    E, psi, M = example_DMRG_tf_ising_finite(L, g)
    print("(DMRG finished)")
    iO = L // 2
    # apply sigmaz on site iO
    psi.apply_local_op(i0, 'Sigmaz', unitary=True)
    dt_measure = 0.05
    # tebd.TEBDEngine makes 'N_steps' steps of 'dt' at once;
    # for second order this is more efficient.
    tebd_params = {
        'order': 2,
        'dt': dt,
        'N_steps': int(dt_measure / dt + 0.5),
        'trunc_params': {
            'chi_max': 50,
            'svd_min': 1.e-10,
            'trunc_cut': None
        },
    }
    eng = tebd.TEBDEngine(psi, M, tebd_params)
    S = [psi.entanglement_entropy()]
    for }n\mathrm{ in range(int(tmax / dt_measure + 0.5)):
        eng.run()
        S.append(psi.entanglement_entropy())
    import matplotlib.pyplot as plt
    plt.figure()
    plt.imshow(S[::-1],
        vmin=0.,
```

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```
                                    (continued from previous page)
                                    aspect='auto'
                                    interpolation='nearest',
                                    extent=(0, L - 1., -0.5 * dt_measure, eng.evolved_time + 0.5 * dt_
\hookrightarrowmeasure))
    plt.xlabel('site $i$')
    plt.ylabel('time $t/J$')
    plt.ylim(0., tmax)
    plt.colorbar().set_label('entropy $S$')
    filename = 'c_tebd_lightcone_{g:.2f}.pdf'.format(g=g)
    plt.savefig(filename)
    print("saved " + filename)
def example_TEBD_gs_tf_ising_next_nearest_neighbor(L, g, Jp):
    from tenpy.models.spins_nnn import SpinChainNNN2
    from tenpy.models.model import NearestNeighborModel
    print("finite TEBD, imaginary time evolution, transverse field Ising next-nearest,
↔neighbor")
    print("L={L:d}, g={g:.2f}, Jp={Jp:.2f}".format(L=L, g=g, Jp=Jp))
    model_params = dict(
        L=L,
        Jx=1.,
        Jy=0.,
        Jz=0.,
        Jxp=Jp,
        Jyp=0.,
        Jzp=0.,
        hz=g,
        bc_MPS='finite',
        conserve=None,
    )
    # we start with the non-grouped sites, but next-nearest neighbor interactions,u
\hookrightarrowbuilding the MPO
    M = SpinChainNNN2 (model_params)
    product_state = ["up"] * M.lat.N_sites
    psi = MPS.from_product_state(M.lat.mps_sites(), product_state, bc=M.lat.bc_MPS)
    # now we group each to sites ...
    psi.group_sites(n=2) # ... in the state
    M.group_sites(n=2) # ... and model
    # now, M has only 'nearest-neighbor' interactions with respect to the grouped_
@sites
    # thus, we can convert the MPO into H_bond terms:
    M_nn = NearestNeighborModel.from_MPOModel(M) # hence, we can initialize H_bond_
\hookrightarrowfrom the MPO
    # now, we continue to run TEBD as before
    tebd_params = {
        'order': 2,
        'delta_tau_list': [0.1, 0.01, 0.001, 1.e-4, 1.e-5],
        'N_steps': 10,
        'max_error_E': 1.e-6,
        'trunc_params': {
            'chi_max': 30,
            'svd_min': 1.e-10
        },
    }
```

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```
    eng = tebd.TEBDEngine(psi, M_nn, tebd_params) # use M_nn and grouped psi
    eng.run_GS() # the main work...
    # expectation values:
    E = np.sum(M_nn.bond_energies(psi)) # bond_energies() works only a for_
\hookrightarrowNearestNeighborModel
    print("E = {E:.I3f}".format(E=E))
    print("final bond dimensions: ", psi.chi)
    # we can split the sites of the state again for an easier evaluation of
\hookrightarrowexpectation values
    psi.group_split()
    mag_x = 2. * np.sum(psi.expectation_value("Sx")) # factor of 2 for Sx vs Sigmax
    mag_z = 2. * np.sum(psi.expectation_value("Sz"))
    print("magnetization in X = {mag_x:.5f}".format(mag_x=mag_x))
    print("magnetization in Z = {mag_z:.5f}".format(mag_z=mag_z))
    return E, psi, M
if ___name___ == "___main___":
    import logging
    logging.basicConfig(level=logging.INFO)
    example_TEBD_gs_tf_ising_finite(L=10, g=1.)
    print("=" * 100, '', ''', "=" * 100, sep='\n')
    example_TEBD_gs_tf_ising_infinite(g=1.5)
    print("=" * 100, '', '', "=" * 100, sep='\n')
    example_TEBD_tf_ising_lightcone(L=20, g=1.5, tmax=3., dt=0.01)
    print("=" * 100, '', ''', "=" * 100, sep='\n')
    example_TEBD_gs_tf_ising_next_nearest_neighbor(L=10, g=1.0, Jp=0.1)
```


### 11.2.4 d_dmrg.py

on github.

```
"""Example illustrating the use of DMRG in tenpy.
The example functions in this class do the same as the ones in `toycodes/d_dmrg.py`,
but make use of the classes defined in tenpy.
" ""
# Copyright 2018-2021 TeNPy Developers, GNU GPLv3
import numpy as np
from tenpy.networks.mps import MPS
from tenpy.models.tf_ising import TFIChain
from tenpy.models.spins import SpinModel
from tenpy.algorithms import dmrg
def example_DMRG_tf_ising_finite(L, g):
    print("finite DMRG, transverse field Ising model")
    print("L={L:d}, g={g:.2f}".format (L=L, g=g))
    model_params = dict(L=L, J=1., g=g, bc_MPS='finite', conserve=None)
    M = TFIChain(model_params)
    product_state = ["up"] * M.lat.N_sites
    psi = MPS.from_product_state(M.lat.mps_sites(), product_state, bc=M.lat.bc_MPS)
```

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```
    dmrg_params = {
        'mixer': None, # setting this to True helps to escape local minima
        'max_E_err': 1.e-10,
        'trunc_params': {
            'chi_max': 30,
            'svd_min': 1.e-10
        },
    'combine': True
    }
    info = dmrg.run(psi, M, dmrg_params) # the main work...
    E = info['E']
    print("E = {E:.I3f}".format(E=E))
    print("final bond dimensions: ", psi.chi)
    mag_x = np.sum(psi.expectation_value("Sigmax"))
    mag_z = np.sum(psi.expectation_value("Sigmaz"))
    print("magnetization in X = {mag_x:.5f}".format(mag_x=mag_x))
    print("magnetization in Z = {mag_z:.5f}".format(mag_z=mag_z))
    if L < 20: # compare to exact result
        from tfi_exact import finite_gs_energy
        E_exact = finite_gs_energy(L, 1., g)
        print("Exact diagonalization: E = {E:.l3f}".format(E=E_exact))
    print("relative error: ", abs((E - E_exact) / E_exact))
    return E, psi, M
def example_1site_DMRG_tf_ising_finite(L, g):
    print("single-site finite DMRG, transverse field Ising model")
    print("L={L:d}, g={g:.2f}".format (L=L, g=g))
    model_params = dict(L=L, J=1., g=g, bc_MPS='finite', conserve=None)
    M = TFIChain(model_params)
    product_state = ["up"] * M.lat.N_sites
    psi = MPS.from_product_state(M.lat.mps_sites(), product_state, bc=M.lat.bc_MPS)
    dmrg_params = {
    'mixer': True, # setting this to True is essential for the l-site algorithm
\hookrightarrowto work.
        'max_E_err': 1.e-10,
        'trunc_params': {
            'chi_max': 30,
            'svd_min': 1.e-10
        },
        'combine': False,
            'active_sites': 1 # specifies single-site
    }
    info = dmrg.run(psi, M, dmrg_params)
    E = info['E']
    print("E = {E:.I3f}".format(E=E))
    print("final bond dimensions: ", psi.chi)
    mag_x = np.sum(psi.expectation_value("Sigmax"))
    mag_z = np.sum(psi.expectation_value("Sigmaz"))
    print("magnetization in X = {mag_x:.5f}".format(mag_x=mag_x))
    print("magnetization in Z = {mag_z:.5f}".format(mag_z=mag_z))
    if L < 20: # compare to exact result
        from tfi_exact import finite_gs_energy
        E_exact = finite_gs_energy(L, 1., g)
        print("Exact diagonalization: E = {E:.l3f}".format(E=E_exact))
        print("relative error: ", abs((E - E_exact) / E_exact))
    return E, psi, M
```

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```
def example_DMRG_tf_ising_infinite(g) :
    print("infinite DMRG, transverse field Ising model")
    print("g={g:.2f}".format (g=g))
    model_params = dict(L=2, J=1., g=g, bc_MPS='infinite', conserve=None)
    M = TFIChain(model_params)
    product_state = ["up"] * M.lat.N_sites
    psi = MPS.from_product_state(M.lat.mps_sites(), product_state, bc=M.lat.bc_MPS)
    dmrg_params = {
        'mixer': True, # setting this to True helps to escape local minima
        'trunc_params': {
            'chi_max': 30,
            'svd_min': 1.e-10
        },
        'max_E_err': 1.e-10,
    }
    # Sometimes, we want to call a 'DMRG engine' explicitly
    eng = dmrg.TwoSiteDMRGEngine(psi, M, dmrg_params)
    E, psi = eng.run() # equivalent to dmrg.run() up to the return parameters.
    print("E = {E:.l3f}".format(E=E))
    print("final bond dimensions: ", psi.chi)
    mag_x = np.mean(psi.expectation_value("Sigmax"))
    mag_z = np.mean(psi.expectation_value("Sigmaz"))
    print("<sigma_x> = {mag_x:.5f}".format(mag_x=mag_x))
    print("<sigma_z> = {mag_z:.5f}".format(mag_z=mag_z))
    print("correlation length:", psi.correlation_length())
    # compare to exact result
    from tfi_exact import infinite_gs_energy
    E_exact = infinite_gs_energy(1., g)
    print("Analytic result: E (per site) = {E:.l3f}".format(E=E_exact))
    print("relative error: ", abs((E - E_exact) / E_exact))
    return E, psi, M
def example_1site_DMRG_tf_ising_infinite(g):
    print("single-site infinite DMRG, transverse field Ising model")
    print("g={g:.2f}".format (g=g))
    model_params = dict(L=2, J=1., g=g, bc_MPS='infinite', conserve=None)
    M = TFIChain(model_params)
    product_state = ["up"] * M.lat.N_sites
    psi = MPS.from_product_state(M.lat.mps_sites(), product_state, bc=M.lat.bc_MPS)
    dmrg_params = {
        'mixer': True, # setting this to True is essential for the I-site algorithm,
\hookrightarrowto work.
            'trunc_params': {
            'chi_max': 30,
            'svd_min': 1.e-10
        },
        'max_E_err': 1.e-10,
        'combine': True
    }
    eng = dmrg.SingleSiteDMRGEngine(psi, M, dmrg_params)
    E, psi = eng.run() # equivalent to dmrg.run() up to the return parameters.
    print("E = {E:.13f}".format (E=E))
    print("final bond dimensions: ", psi.chi)
    mag_x = np.mean(psi.expectation_value("Sigmax"))
```

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```
    mag_z = np.mean(psi.expectation_value("Sigmaz"))
    print("<sigma_x> = {mag_x:.5f}".format(mag_x=mag_x))
    print("<sigma_z> = {mag_z:.5f}".format(mag_z=mag_z))
    print("correlation length:", psi.correlation_length())
    # compare to exact result
    from tfi_exact import infinite_gs_energy
    E_exact = infinite_gs_energy(1., g)
    print("Analytic result: E (per site) = {E:.l3f}".format(E=E_exact))
    print("relative error: ", abs((E - E_exact) / E_exact))
def example_DMRG_heisenberg_xxz_infinite(Jz, conserve='best'):
    print("infinite DMRG, Heisenberg XXZ chain")
    print("Jz={Jz:.2f}, conserve={conserve!r}".format(Jz=Jz, conserve=conserve))
    model_params = dict(
        L=2,
        S=0.5, # spin 1/2
        Jx=1.,
        Jy=1.,
        Jz=Jz, # couplings
        bc_MPS='infinite',
        conserve=conserve)
    M = SpinModel(model_params)
    product_state = ["up", "down"] # initial Neel state
    psi = MPS.from_product_state(M.lat.mps_sites(), product_state, bc=M.lat.bc_MPS)
    dmrg_params = {
        'mixer': True, # setting this to True helps to escape local minima
        'trunc_params': {
            'chi_max': 100,
            'svd_min': 1.e-10,
        },
        'max_E_err': 1.e-10,
    }
    info = dmrg.run(psi, M, dmrg_params)
    E = info['E']
    print("E = {E:.l3f}".format(E=E))
    print("final bond dimensions: ", psi.chi)
    Sz = psi.expectation_value("Sz") # Sz instead of Sigma z: spin-1/2 operators!
    mag_z = np.mean(Sz)
    print("<S_z> = [{Sz0:.5f}, {Sz1:.5f}]; mean ={mag_z:.5f}".format(Sz0=Sz[0],
                                    Sz1=Sz[1],
                                    mag_z=mag_z))
    # note: it's clear that mean(\langleSz>) is 0: the model has Sz conservation!
    print("correlation length:", psi.correlation_length())
    corrs = psi.correlation_function("Sz", "Sz", sitesl=range(10))
    print("correlations <Sz_i Sz_j> =")
    print(corrs)
    return E, psi, M
if name == " main ":
    import logging
    logging.basicConfig(level=logging.INFO)
    example_DMRG_tf_ising_finite(L=10, g=1.)
    print("-" * 100)
    example_1site_DMRG_tf_ising_finite(L=10, g=1.)
    print("-" * 100)
```

```
example_DMRG_tf_ising_infinite(g=1.5)
print("-" * 100)
example_1site_DMRG_tf_ising_infinite(g=1.5)
print("-" * 100)
example_DMRG_heisenberg_xxz_infinite(Jz=1.5)
```


### 11.2.5 e_tdvp.py

on github.

```
"""Example illustrating the use of TDVP in tenpy.
As of now, we have TDVP only for finite systems. The call structure is quite similar,
\hookrightarrowto TEBD.A
difference is that we can run one-site TDVP or two-site TDVP. In the former, the bond,
@dimension can
not grow; the latter allows to grow the bond dimension and hence requires ab
\hookrightarrowtruncation.
"""
# Copyright 2019-2021 TeNPy Developers, GNU GPLv3
import numpy as np
import tenpy.linalg.np_conserved as npc
import tenpy.models.spins
import tenpy.networks.mps as mps
import tenpy.networks.site as site
from tenpy.algorithms import tdvp
from tenpy.networks.mps import MPS
import copy
def run_out_of_equilibrium():
    L = 10
    chi = 5
    delta_t = 0.1
    model_params = {
        'L': L,
        'S': 0.5,
        'conserve': 'Sz',
        'Jz': 1.0,
        'Jy': 1.0,
        'JX': 1.0,
            'hx': 0.0,
            'hy': 0.0,
            'hz': 0.0,
            'muJ': 0.0,
            'bc_MPS': 'finite',
    }
    heisenberg = tenpy.models.spins.SpinChain(model_params)
    product_state = ["up"] * (L // 2) + ["down"] * (L - L // 2)
    # starting from a domain-wall product state which is not an eigenstate of the
\hookrightarrowHeisenberg model
    psi = MPS.from_product_state(heisenberg.lat.mps_sites(),
                        product_state,
                        bc=heisenberg.lat.bc_MPS,
```

```
            form='B')
    tdvp_params = {
        'start_time': 0,
        'dt': delta_t,
        'trunc_params': {
            'chi_max': chi,
            'svd_min': 1.e-10,
            'trunc_cut': None
        }
    }
    tdvp_engine = tdvp.TDVPEngine(psi, heisenberg, tdvp_params)
    times = []
    S_mid = []
    for i in range(30):
        tdvp_engine.run_two_sites(N_steps=1)
        times.append(tdvp_engine.evolved_time)
        S_mid.append(psi.entanglement_entropy(bonds=[L // 2])[0])
        for i in range(30):
        tdvp_engine.run_one_site(N_steps=1)
        #psi_2=copy.deepcopy(psi)
        #psi_2.canonical_form()
        times.append(tdvp_engine.evolved_time)
        S_mid.append(psi.entanglement_entropy(bonds=[L // 2])[0])
    import matplotlib.pyplot as plt
    plt.figure()
    plt.plot(times, S_mid)
    plt.xlabel('t')
    plt.ylabel('S')
    plt.axvline(x=3.1, color='red')
    plt.text(0.0, 0.0000015, "Two sites update")
    plt.text(3.1, 0.0000015, "One site update")
    plt.show()
if n_me_== " main ":
    import logging
    logging.basicConfig(level=logging.INFO)
    run_out_of_equilibrium()
```


### 11.2.6 purification.py

on github.

```
from tenpy.models.tf_ising import TFIChain
from tenpy.networks.purification_mps import PurificationMPS
from tenpy.algorithms.purification import PurificationTEBD, PurificationApplyMPO
def imag_tebd(L=30, beta_max=3., dt=0.05, order=2, bc="finite"):
    model_params = dict(L=L, J=1., g=1.2)
    M = TFIChain(model_params)
    psi = PurificationMPS.from_infiniteT(M.lat.mps_sites(), bc=bc)
    options = {
        'trunc_params': {
```

```
            'chi_max': 100,
            'svd_min': 1.e-8
        },
        'order': order,
        'dt': dt,
        'N_steps': 1
    }
    beta = 0.
    eng = PurificationTEBD(psi, M, options)
    Szs = [psi.expectation_value("Sz")]
    betas = [0.]
    while beta < beta_max:
        beta += 2. * dt # factor of 2: |psi> ~= exp^{-dt H}, but rho = |psi><psi|
        betas.append(beta)
        eng.run_imaginary(dt) # cool down by dt
        Szs.append(psi.expectation_value("Sz")) # and further measurements...
    return {'beta': betas, 'Sz': Szs}
def imag_apply_mpo(L=30, beta_max=3., dt=0.05, order=2, bc="finite", approx="II"):
    model_params = dict(L=L, J=1., g=1.2)
    M = TFIChain(model_params)
    psi = PurificationMPS.from_infiniteT(M.lat.mps_sites(), bc=bc)
    options = {'trunc_params': {'chi_max': 100, 'svd_min': 1.e-8}}
    beta = 0.
    if order == 1:
        Us = [M.H_MPO.make_U(-dt, approx)]
    elif order == 2:
            Us = [M.H_MPO.make_U(-d * dt, approx) for d in [0.5 + 0.5j, 0.5 - 0.5j]]
    eng = PurificationApplyMPO(psi, Us[0], options)
    Szs = [psi.expectation_value("Sz")]
    betas = [0.]
    while beta < beta_max:
        beta += 2. * dt # factor of 2: |psi> ~= exp^{-dt H}, but rho = |psi><psi|
        betas.append (beta)
        for U in Us:
            eng.init_env(U) # reset environment, initialize new copy of psi
            eng.run() # apply U to psi
        Szs.append(psi.expectation_value("Sz")) # and further measurements...
    return {'beta': betas, 'Sz': Szs}
if ___name___ == "___main___":
    import logging
    logging.basicConfig(level=logging.INFO)
    data_tebd = imag_tebd()
    data_mpo = imag_apply_mpo()
    import numpy as np
    from matplotlib.pyplot import plt
    plt.plot(data_mpo['beta'], np.sum(data_mpo['Sz'], axis=1), label='MPO')
    plt.plot(data_tebd['beta'], np.sum(data_tebd['Sz'], axis=1), label='TEBD')
    plt.xlabel(r'$\beta$')
    plt.ylabel(r'total $S^z$')
    plt.show()
```


### 11.2.7 tfi_exact.py

on github.

```
"""Provides exact ground state energies for the transverse field ising model for,
@comparison.
The Hamiltonian reads
.. math : :
    H = - J \\sum_{i} \\sigma^x_i \\sigma^x_{i+1} - g \\sum_{i} \\sigma^z_i
"""
# Copyright 2019-2021 TeNPy Developers, GNU GPLv3
import numpy as np
import scipy.sparse as sparse
import scipy.sparse.linalg.eigen.arpack as arp
import warnings
import scipy.integrate
def finite_gs_energy(L, J, g):
    """For comparison: obtain ground state energy from exact diagonalization.
    Exponentially expensive in L, only works for small enough `L` <~ 20.
    " ""
    if L >= 20:
        warnings.warn("Large L: Exact diagonalization might take a long time!")
    # get single site operaors
    sx = sparse.csr_matrix(np.array([[0., 1.], [1., 0.]]))
    sz = sparse.csr_matrix(np.array([[1., 0.], [0., -1.]]))
    id = sparse.csr_matrix(np.eye(2))
    sx_list = [] # sx_list[i] = kron([id, id, ..., id, sx, id, .... id])
    sz_list = []
    for i_site in range(L):
        x_ops = [id] * L
        z_ops = [id] * L
        x_ops[i_site] = sx
        z_ops[i__site] = sz
        X = x_ops[0]
        Z = z_ops[0]
        for j in range(1, L):
            X = sparse.kron(X, x_ops[j], 'csr')
            Z = sparse.kron(Z, z_ops[j], 'csr')
        sx_list.append(X)
        sz_list.append(Z)
    H_xx = sparse.csr_matrix((2**L, 2**L))
    H_z = sparse.csr_matrix((2**L, 2**L))
    for i in range(L - 1):
        H_xx = H_xx + sx_list[i] * sx_list[(i + 1) % L]
    for i in range(L):
        H_z = H_z + sz_list[i]
    H = -J * H_xx - g * H_z
    E, V = arp.eigsh(H, k=1, which='SA', return_eigenvectors=True, ncv=20)
    return E[O]
def infinite_gs_energy(J, g):
    """For comparison: Calculate groundstate energy density from analytic formula.
```

(continues on next page)

```
    The analytic formula stems from mapping the model to free fermions, see P. Pfeuty,
\hookrightarrow The one-
    dimensional Ising model with a transverse field, Annals of Physics 57, p. 79
\hookrightarrow(1970). Note that
    we use Pauli matrices compared this reference using spin-1/2 matrices and replace,
\hookrightarrowthe sum_k ->
    integral dk/2pi to obtain the result in the N -> infinity limit.
    """
    def f(k, lambda_):
        return np.sqrt(1 + lambda_**2 + 2 * lambda_ * np.cos(k))
    E0_exact = -g / (J * 2. * np.pi) * scipy.integrate.quad(f, -np.pi, np.pi, args=(J, |
@/ g, ))[0]
    return E0_exact
```


### 11.2.8 z_exact_diag.py

on github.

```
"""A simple example comparing DMRG output with full diagonalization (ED).
Sorry that this is not well documented! ED is meant to be used for debugging only ;)
"""
# Copyright 2018-2021 TeNPy Developers, GNU GPLv3
import tenpy.linalg.np_conserved as npc
from tenpy.models.xxz_chain import XXZChain
from tenpy.networks.mps import MPS
from tenpy.algorithms.exact_diag import ExactDiag
from tenpy.algorithms import dmrg
def example_exact_diagonalization(L, Jz):
    xxz_pars = dict(L=L, Jxx=1., Jz=Jz, hz=0.0, bc_MPS='finite')
    M = XXZChain(xxz_pars)
    product_state = ["up", "down"] * (xxz_pars['L'] // 2) # this selects a charge 
sector!
    psi_DMRG = MPS.from_product_state(M.lat.mps_sites(), product_state)
    charge_sector = psi_DMRG.get_total_charge(True) # ED charge sector should match
    ED = ExactDiag(M, charge_sector=charge_sector, max_size=2.e6)
    ED.build_full_H_from_mpo()
    # ED.build_full_H_from_bonds() # whatever you prefer
    print("start diagonalization")
    ED.full_diagonalization() # the expensive part for large L
    EO_ED, psi_ED = ED.groundstate() # return the ground state
    print("psi_ED =", psi_ED)
    print("run DMRG")
    dmrg.run(psi_DMRG, M, {'verbose': 0}) # modifies psi_DMRG in place!
    # first way to compare ED with DMRG: convert MPS to ED vector
    psi_DMRG_full = ED.mps_to_full(psi_DMRG)
```

(continues on next page)

```
    print("psi_DMRG_full =", psi__DMRG_full)
    ov = npc.inner(psi_ED, psi_DMRG_full, axes='range', do_conj=True)
    print("<psi_ED|psi__DMRG_full> =", ov)
    assert (abs(abs(ov) - 1.) < 1.e-13)
    # second way: convert ED vector to MPS
    psi_ED_mps = ED.full_to_mps(psi_ED)
    ov2 = psi_ED_mps.overlap(psi_DMRG)
    print("<psi_ED_mps|psi_DMRG> =", ov2)
    assert (abs(abs(ov2) - 1.) < 1.e-13)
    assert (abs(ov - ov2) < 1.e-13)
    # -> advantage: expectation_value etc. of MPS are available!
    print("<Sz> =", psi_ED_mps.expectation_value('Sz'))
if ___name___ == "__main__":
    example_exact_diagonalization(10, 1.)
```


### 11.3 Jupyter Notebooks

This is a collection of [jupyter] notebooks from the [TeNPyNotebooks] repository. You need to install TeNPy to execute them (see Installation instructions), but you can copy them anywhere before execution. Note that some of them might take a while to run, as they contain more extensive examples.

### 11.3.1 A first TEBD Example

Like examples/c_tebd.py, this notebook shows the basic interface for TEBD. It initalized the transverse field Ising model $H=J X X+g Z$ at the critical point $J=g=1$, and an MPS in the all-up state $|\uparrow \cdots \uparrow\rangle$. It then performs a real-time evolution with TEBD and measures a few observables. This setup correspond to a global quench from $g=\infty$ to $g=1$.

```
[1]: import numpy as np
import scipy
import matplotlib.pyplot as plt
import matplotlib
[2]: import tenpy
import tenpy.linalg.np_conserved as npc
from tenpy.algorithms import tebd
from tenpy.networks.mps import MPS
from tenpy.models.tf_ising import TFIChain
[3]: L = 30
[4]: model_params = {
    'J': 1. , 'g': 1., # critical
    'L': L,
    'bc_MPS': 'finite',
}
M = TFIChain(model_params)
```

Reading 'bc_MPS'='finite' for config TFIChain
Reading 'L'=30 for config TFIChain
Reading 'J'=1.0 for config TFIChain
Reading 'g'=1.0 for config TFIChain
[5]: psi = MPS.from_lat_product_state(M.lat, [['up']])
[6]: tebd_params $=$ \{
'N_steps': 1,
'dt': 0.1,
'order': 4,
'trunc_params': \{'chi_max': 100, 'svd_min': 1.e-12\}
\}
eng = tebd.Engine(psi, M, tebd_params)
Subconfig 'trunc_params'=Config(<3 options>, 'trunc_params') for config TEBD

```
[7]: def measurement(eng, data):
```

    keys \(=\) ['t', 'entropy', 'Sx', 'Sz', 'corr_XX', 'corr_ZZ', 'trunc_err']
    if data is None:
            data \(=\operatorname{dict}([(k,[])\) for \(k\) in keys])
    data['t'].append (eng.evolved_time)
    data['entropy'].append (eng.psi.entanglement_entropy())
    data['Sx'].append (eng.psi.expectation_value('Sigmax'))
    data['Sz'].append(eng.psi.expectation_value('Sigmaz'))
    data['corr_XX'].append(eng.psi.correlation_function('Sigmax', 'Sigmax'))
    data['trunc_err']. append (eng.trunc_err.eps)
    return data
    [8]: data = measurement(eng, None)
[9]: while eng.evolved_time < 5.:
eng.run()
measurement (eng, data)
Reading 'dt'=0.1 for config TEBD
Reading 'N_steps'=1 for config TEBD
Reading 'order'=4 for config TEBD
Calculate $U$ for $\left\{' o r d e r ': 4, ~ ' d e l t a \_t ': ~ 0.1, ~ ' t y p e \_e v o ': ~ ' r e a l ', ~ ' E \_o f f s e t ': ~ N o n e, ~\right.$
$\hookrightarrow ' t a u ': 0.1\}$
--> time=0.100, max_chi=6, Delta_S=5.5243e-02, $S=0.0552432967$, since last update: 0.5
$\hookrightarrow S$
$-->$ time $=0.200$, max_chi=6, Delta_S=1.0453e-01, $S=0.1597705686$, since last update: 0.5 ,
$\hookrightarrow S$
--> time=0.300, max_chi=8, Delta_S=1.1368e-01, $S=0.2734471407$, since last update: 0.5
$\hookrightarrow S$
$-->$ time $=0.400$, max_chi=10, Delta_S=1.0226e-01, $S=0.3757082127$, since last update: 0.
$\rightarrow 5 \mathrm{~s}$
$-->$ time $=0.500$, max_chi=12, Delta_S $=8.2474 e-02, S=0.4581821173$, since last update: 0 .
$\hookrightarrow 5$ s
--> time=0.600, max_chi=12, Delta_S=6.3393e-02, $S=0.5215746922$, since last update: 0.
$\hookrightarrow 5 \mathrm{~s}$
$-->$ time $=0.700$, max_chi=15, Delta_S=5.0837e-02, $S=0.5724119006$, since last update: 0.
$\hookrightarrow 5 \mathrm{~s}$
--> time=0.800, max_chi=18, Delta_S=4.7046e-02, $S=0.6194580889$, since last update: 0 .
$\hookrightarrow 5 \mathrm{~s}$
(continues on next page)
(continued from previous page)
--> time=0.900, max_chi=20, Delta_S=5.0606e-02, $S=0.6700636890$, since last update: 0 . $\hookrightarrow 5$ s
--> time=1.000, max_chi=20, Delta_S=5.7462e-02, $S=0.7275254979$, since last update: 0. $\hookrightarrow 5 \mathrm{~s}$
$-->$ time=1.100, max_chi=24, Delta_S=6.3115e-02, $S=0.7906402648$, since last update: 0 . $\hookrightarrow 5 \mathrm{~s}$
--> time=1.200, max_chi=26, Delta_S=6.4779e-02, $S=0.8554189184$, since last update: 0 . $\hookrightarrow 5 \mathrm{~s}$
--> time=1.300, max_chi=30, Delta_S=6.2269e-02, $S=0.9176882429$, since last update: 0 . $\hookrightarrow 5 \mathrm{~s}$
--> time=1.400, max_chi=34, Delta_S=5.7451e-02, $S=0.9751394599$, since last update: 0. $\rightarrow 5 \mathrm{~s}$
--> time=1.500, max_chi=40, Delta_S=5.2899e-02, $S=1.0280386118$, since last update: 0 . $\rightarrow 5 \mathrm{~s}$
--> time=1.600, max_chi=40, Delta_S=5.0543e-02, $S=1.0785817862$, since last update: 0. $\rightarrow 5 \mathrm{~s}$
--> time=1.700, max_chi=46, Delta_S=5.0852e-02, $S=1.1294340516$, since last update: 0 . $\rightarrow 5 \mathrm{~s}$
--> time=1.800, max_chi=52, Delta_S=5.2841e-02, $S=1.1822748827$, since last update: 0. $\hookrightarrow 7 \mathrm{~s}$
--> time=1.900, max_chi=57, Delta_S=5.4804e-02, $S=1.2370787276$, since last update: 0 . $\hookrightarrow 9 \mathrm{~s}$
--> time=2.000, max_chi=62, Delta_S=5.5311e-02, $S=1.2923895877$, since last update: 0 . $\leftrightarrow 9 \mathrm{~s}$
$-->$ time=2.100, max_chi=71, Delta_S=5.3906e-02, $S=1.3462960135$, since last update: 1 . $\leftrightarrow 0 \quad \mathrm{~s}$
--> time=2.200, max_chi=80, Delta_S=5.1209e-02, $S=1.3975050668$, since last update: 1 . $\hookrightarrow 2 \mathrm{~s}$
$-->$ time=2.300, max_chi=85, Delta_S=4.8446e-02, $S=1.4459514729$, since last update: 1 . $\hookrightarrow 4 \mathrm{~s}$
--> time=2.400, max_chi=95, Delta_S=4.6732e-02, $S=1.4926837515$, since last update: 1. $\rightarrow 4$ s
--> time=2.500, max_chi=100, Delta_S=4.6486e-02, $S=1.5391700161$, since last update: 1 . $\hookrightarrow 8 \mathrm{~s}$
--> time=2.600, max_chi=100, Delta_S=4.7282e-02, $S=1.5864521162$, since last update: 1 . $\hookrightarrow 9 \mathrm{~s}$
$-->$ time=2.700, max_chi=100, Delta_S=4.8160e-02, $S=1.6346120966$, since last update: 1 . $\hookrightarrow 9 \mathrm{~s}$
--> time=2.800, max_chi=100, Delta_S=4.8202e-02, $S=1.6828138289$, since last update: 1 . $\hookrightarrow 6 \mathrm{~s}$
--> time=2.900, max_chi=100, Delta_S=4.7044e-02, $S=1.7298576742$, since last update: 1 . $\hookrightarrow 7 \mathrm{~s}$
--> time=3.000, max_chi=100, Delta_S=4.5033e-02, $S=1.7748910539$, since last update: 1 . $\hookrightarrow 6 \mathrm{~s}$
$-->$ time=3.100, max_chi=100, Delta_S=4.2960e-02, $S=1.8178514255$, since last update: 1 . $\hookrightarrow 5$ s
--> time=3.200, max_chi=100, Delta_S=4.1575e-02, $S=1.8594265450$, since last update: 1. $\hookrightarrow 4 \mathrm{~s}$
$-->$ time=3.300, max_chi=100, Delta_S=4.1182e-02, $S=1.9006080872$, since last update: 1 . $\hookrightarrow 4 \mathrm{~s}$
$-->$ time=3.400, max_chi=100, Delta_S=4.1503e-02, $S=1.9421110677$, since last update: 1. $\hookrightarrow 4 \mathrm{~s}$
$-->$ time=3.500, max_chi=100, Delta_S=4.1873e-02, S=1.9839839097, since last update: 1. $\hookrightarrow 6 \mathrm{~s}$
$-->$ time=3.600, max_chi=100, Delta_S=4.1645e-02, $S=2.0256293591$, since last update: 1 . $\hookrightarrow 6 \mathrm{~s}$
$-->$ time=3.700, max_chi=100, Delta_S=4.0570e-02, $S=2.0661996178$, since last update: 1 .
$\hookrightarrow 8 \mathrm{~s}$
(continues on next page)
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```
--> time=3.800, max_chi=100, Delta_S=3.8897e-02, S=2.1050963078, since last update: 1.
\hookrightarrow s
--> time=3.900, max_chi=100, Delta_S=3.7189e-02, S=2.1422855970, since last update: 2.
\hookrightarrow s
--> time=4.000, max_chi=100, Delta_S=3.5994e-02, S=2.1782792652, since last update: 1.
\hookrightarrow s
--> time=4.100, max_chi=100, Delta_S=3.5534e-02, S=2.2138131347, since last update: 1.
\hookrightarrow s
--> time=4.200, max_chi=100, Delta_S=3.5597e-02, S=2.2494096463, since last update: 1.
\hookrightarrow s
--> time=4.300, max_chi=100, Delta_S=3.5673e-02, S=2.2850830917, since last update: 1.
\hookrightarrow s
--> time=4.400, max_chi=100, Delta_S=3.5272e-02, S=2.3203546259, since last update: 1.
\hookrightarrow S
--> time=4.500, max_chi=100, Delta_S=3.4188e-02, S=2.3545423452, since last update: 1.
\hookrightarrow4 s
--> time=4.600, max_chi=100, Delta_S=3.2596e-02, S=2.3871383468, since last update: 1.
\hookrightarrow S
--> time=4.700, max_chi=100, Delta_S=3.0915e-02, S=2.4180531110, since last update: 1.
\leftrightarrow 6 ~ s
--> time=4.800, max_chi=100, Delta_S=2.9546e-02, S=2.4475988717, since last update: 1.
\hookrightarrow s
--> time=4.900, max_chi=100, Delta_S=2.8630e-02, S=2.4762292780, since last update: 1.
\hookrightarrow s
--> time=5.000, max_chi=100, Delta_S=2.7962e-02, S=2.5041913364, since last update: 1.
\hookrightarrow s
--> time=5.100, max_chi=100, Delta_S=2.7099e-02, S=2.5312904883, since last update: 1.
\hookrightarrow s
```

[10]:

```
plt.plot(data['t'], np.array(data['entropy'])[:, L//2])
plt.xlabel('time $t$')
plt.ylabel('entropy $S$')
```

[10]: Text(0, 0.5, 'entropy \$S\$')


The growth of $S$ linear in time is typical for a global quench and to be expected from the quasi-particle picture
[11](matplotlib.legend.Legend%20at%200x7f1018760e80):

```
plt.plot(data['t'], np.sum(data['Sx'], axis=1), label="X")
plt.plot(data['t'], np.sum(data['Sz'], axis=1), label="Z")
plt.xlabel('time $t$')
plt.ylabel('magnetization')
plt.legend(loc='best')
```



The strict conservation of X being zero is ensured by charge conservation, because X changes the parity sector.
Nevertheless, the XX correlation function can be nontrivial:

```
[12]:
corrs = np.array(data['corr_XX'])
tmax = data['t'][-1]
x = np.arange(L)
cmap = matplotlib.cm.viridis
for i, t in list(enumerate(data['t'])):
    if i == 0 or i == len(data['t']) - 1:
            label = '{t:.2f}'.format (t=t)
    else:
            label = None
    plt.plot(x, corrs[i, L//2, :], color=cmap(t/tmax), label=label)
plt.xlabel(r'time $t$')
plt.ylabel(r'correlations $\langle X_i X_{j:d}\rangle$'.format(j=L//2))
plt.yscale('log')
plt.ylim(1.e-6, 1.)
plt.legend()
plt.show()
```



The output of the run showed that we gradually increased the bond dimension and only reached the maximum chi around $t=2.5$. At this point we start to truncate significantly, because we cut off the tail whatever the singular values are. This is clearly visible if we plot the truncation error vs. time. Note the log-scale, though: if you are fine with an error of say 1 permille for expectation values, you can still go on for a bit more!
[13]: plt.plot(data['t'], data['trunc_err'])
plt.yscale('log')
\#plt.ylim(1.e-15, 1.)
plt.xlabel('time \$t\$')
plt.ylabel('truncation error')
[13]:
Text (0, 0.5, 'truncation error')

[ ]:

### 11.3.2 A first finite DMRG Example

Like examples/d_dmrg.py, this notebook shows the basic interface for DMRG. It initalized the transverse field Ising model $H=J X X+g Z$ at the critical point $J=g=1$, and a finite MPS in the all-up state $|\uparrow \cdots \uparrow\rangle$. It then runs DMRG to find the ground state. Finally, we look at the profile of the entanglement-cuts.

```
[1]: import numpy as np
import scipy
import matplotlib.pyplot as plt
[2]: import tenpy
import tenpy.linalg.np_conserved as npc
from tenpy.algorithms import dmrg
from tenpy.networks.mps import MPS
from tenpy.models.tf_ising import TFIChain
```

[4]: model_params = \{
'J': 1. , 'g': 1., \# critical
'L': L,
'bc_MPS': 'finite',
\}
$\mathrm{M}=$ TFIChain(model_params)
Reading 'bc_MPS'='finite' for config TFIChain
Reading 'L'=100 for config TFIChain
Reading 'J'=1.0 for config TFIChain
Reading 'g'=1.0 for config TFIChain
[5]: psi = MPS.from_lat_product_state(M.lat, [['up']])
[6]: dmrg_params = \{
'mixer': None, \# setting this to True helps to escape local minima
'max_E_err': 1.e-10,
'trunc_params': \{
'chi_max': 100,
'svd_min': 1.e-10,
\},
'verbose': True,
'combine': True
\}
eng = dmrg.TwoSiteDMRGEngine(psi, M, dmrg_params)
E, psi = eng.run() \# the main work; modifies psi in place
Reading 'combine'=True for config TwoSiteDMRGEngine
Subconfig 'trunc_params'=Config(<3 options>, 'trunc_params') for config
$\rightarrow$ TwoSiteDMRGEngine
Reading 'max_E_err'=1e-10 for config TwoSiteDMRGEngine
Reading 'mixer'=None for config TwoSiteDMRGEngine

sweep 1, age = 100
Energy $=-126.9290280127265333, \mathrm{~S}=0.3766760098039985$, norm_err $=1.2 \mathrm{e}-01$
Current memory usage 97.4 MB , time elapsed: 2.3 s
Delta $\mathrm{E}=$ nan, Delta $\mathrm{S}=$ nan (per sweep)
max_trunc_err $=0.0000 \mathrm{e}+00$, max_E_trunc $=1.8474 \mathrm{e}-13$

```
MPS bond dimensions: [2, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 
\leftrightarrow4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4,ப
\hookrightarrow4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4,七
\leftrightarrow4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 2]
========================================================================================
sweep 2, age = 100
Energy = -126.9618018107068309, S = 0.5108558254154341, norm_err = 7.0e-03
Current memory usage 99.5 MB, time elapsed: 5.6 s
Delta E = -3.2774e-02, Delta S = 1.3418e-01 (per sweep)
max_trunc_err = 0.0000e+00, max_E_trunc = 1.9895e-13
MPS bond dimensions: [2, 4, 8, 16, 16, 16, 16, 16, 16, 16, 16, 16, 16, 16, 16, 16, 16,
46, 16, 16, 16, 16, 16, 16, 16, 16, 16, 16, 16, 16, 16, 16, 16, 16, 16, 16, 16, 16,
416, 16, 16, 16, 16, 16, 16, 16, 16, 16, 16, 16, 16, 16, 16, 16, 16, 16, 16, 16, 16,
\leftrightarrow16, 16, 16, 16, 16, 16, 16, 16, 16, 16, 16, 16, 16, 16, 16, 16, 16, 16, 16, 16, 16,
\hookrightarrow 16, 16, 16, 16, 16, 16, 16, 16, 16, 16, 16, 16, 16, 16, 16, 16, 8, 4, 2]
==================================================================================
sweep 3, age = 100
Energy = -126.9618767383882130, S = 0.5277643082687588, norm_err = 1.1e-05
Current memory usage 107.0 MB, time elapsed: 10.1 s
Delta E = -7.4928e-05, Delta S = 1.6908e-02 (per sweep)
max_trunc_err = 4.7789e-20, max_E_trunc = 2.8422e-13
MPS bond dimensions: [2, 4, 8, 16, 27, 35, 39, 43, 46, 49, 49, 52, 55, 60, 62, 64, 64,
4 64, 64, 64, 64, 64, 64, 64, 64, 64, 64, 64, 64, 64, 64, 64, 64, 64, 64, 64, 64, 64,
\hookrightarrow 64, 64, 64, 64, 64, 64, 64, 64, 64, 64, 64, 64, 64, 64, 64, 64, 64, 64, 64, 64, 64,
\hookrightarrow 64, 64, 64, 64, 64, 64, 64, 64, 64, 64, 64, 64, 64, 64, 64, 64, 64, 64, 64, 64, 64,
4 64, 64, 61, 59, 60, 55, 51, 47, 44, 42, 42, 38, 35, 32, 23, 16, 8, 4, 2]
```

sweep 4, age $=100$
Energy $=-126.9618767396799086, S=0.5277994024302479$, norm_err $=5.8 e-10$
Current memory usage 115.4 MB , time elapsed: 16.9 s
Delta $\mathrm{E}=-1.2917 \mathrm{e}-09$, Delta $\mathrm{S}=3.5094 \mathrm{e}-05$ (per sweep)
max_trunc_err $=1.3388 \mathrm{e}-18$, max_E_trunc $=2.8422 \mathrm{e}-13$
MPS bond dimensions: $[2,4,8,16,22,30,34,36,41,45,47,48,52,54,57,61,61$,
$\leftrightarrow 63,64,66,67,70,71,72,74,76,76,77,79,79,80,82,83,83,83,84,84,84$,
$\hookrightarrow 84,86,86,89,88,89,89,88,88,89,88,88,88,87,86,87,87,87,87,86,86$,
$\hookrightarrow 85,85,84,84,84,83,83,83,83,81,80,79,77,77,77,75,73,71,70,67,66$,
$\hookrightarrow 64,62,61,61,57,54,52,48,47,45,41,36,34,30,22,16,8,4,2]$
$=================================================================================$
sweep 5, age $=100$
Energy $=-126.9618767396792691, S=0.5277994025033066$, norm_err $=2.1 e-13$
Current memory usage 115.4 MB , time elapsed: 20.4 s
Delta $E=6.3949 e-13$, Delta $S=7.3059 e-11$ (per sweep)
max_trunc_err $=6.5986 \mathrm{e}-20$, max_E_trunc $=3.1264 \mathrm{e}-13$
MPS bond dimensions: $[2,4,8,16,22,30,34,36,41,45,47,48,52,54,57,61,61$,
$\hookrightarrow 63,64,66,67,71,71,74,77,77,78,79,81,81,81,83,83,84,85,85,85,85$,
$\leftrightarrow 86,88,89,90,90,90,90,90,90,91,91,91,90,90,90,90,90,90,90,90,89$,
$\hookrightarrow 88,86,86,85,85,85,84,83,83,81,81,81,79,78,78,77,73,71,71,67,66$,
$\hookrightarrow 64,63,61,61,57,54,52,48,47,45,41,36,34,30,22,16,8,4,2]$
$================================================================================$
DMRG finished after 5 sweeps.
total size $=100$, maximum chi $=91$

## Expectation Values

```
[7]: # the ground state energy was directly returned by dmrg.run()
print("ground state energy = ", E)
# there are other ways to extract the energy from psi:
E1 = M.H_MPO.expectation_value(psi) # based on the MPO
E2 = np.sum(M.bond_energies(psi)) # based on bond terms of H, works only for a
\hookrightarrowNearestNeighborModel
assert abs(E-E1) < 1.e-10 and abs (E-E2) < 1.e-10
ground state energy = -126.96187673967927
[8]: # onsite expectation values
X = psi.expectation_value("Sigmax")
Z = psi.expectation_value("Sigmaz")
x = np.arange(psi.L)
plt.figure()
plt.plot(x, Z, label="Z")
plt.plot(x, X, label="X") # note: it's clear that this is zero due to charge,
->conservation!
plt.xlabel("site")
plt.ylabel("onsite expectation value")
plt.legend()
plt.show()
```



```
[9]:
```

```
# correlation functions
```


# correlation functions

i0 = psi.L // 4 \# for fixed `i`
i0 = psi.L // 4 \# for fixed `i`
j = np.arange(i0 + 1, psi.L)
j = np.arange(i0 + 1, psi.L)
XX = psi.term_correlation_function_right([("Sigmax", 0)], [("Sigmax", 0)], i_L=i0, j_
XX = psi.term_correlation_function_right([("Sigmax", 0)], [("Sigmax", 0)], i_L=i0, j_
\hookrightarrowR=j)
\hookrightarrowR=j)
XX_disc = XX - X[i0] * X[j]
XX_disc = XX - X[i0] * X[j]
ZZ = psi.term_correlation_function_right([("Sigmaz", 0)], [("Sigmaz", 0)], i_L=i0, j_
ZZ = psi.term_correlation_function_right([("Sigmaz", 0)], [("Sigmaz", 0)], i_L=i0, j_
GR=j)
GR=j)
ZZ_disc = ZZ - Z[i0] * Z[j]
ZZ_disc = ZZ - Z[i0] * Z[j]
dx = j - i0

```
dx = j - i0
```

(continues on next page)

```
plt.figure()
plt.plot(dx, XX_disc, label="X X")
plt.plot(dx, ZZ_disc, label="Z Z")
plt.xlabel(r"distance $|i-j|$")
plt.ylabel(r"disconnected correlations $\langle A_i A_j\rangle - \langle A_i \rangle\
\hookrightarrowlangle A_j\rangle$")
plt.legend()
plt.loglog()
plt.show()
```



We find power-law decaying correlations, as expected for a critical model. For a gapped model, we would expect exponentially decaying correlations.

We now look at the entanglement entropy. The transverse-field Ising model is critical at $g=J$. Conformal field theory, Calabrese,Cardy 2004, predicts an entanglement entropy profile of
where : math : ' $c=0.5$ 'isthecentralcharge, : math : 'a'isthelatticespacing(weset : math : ‘a=1'), : math : ' $L$ 'isthetotalsize

```
[12]:
```

```
S = psi.entanglement_entropy()
```

S = psi.entanglement_entropy()
bonds = np.arange(0.5, psi.L-1)
bonds = np.arange(0.5, psi.L-1)
plt.plot(bonds, S, 'o', label="S")
plt.plot(bonds, S, 'o', label="S")

# preform fit to extract the central charge

# preform fit to extract the central charge

central_charge, const, res = tenpy.tools.fit.central_charge_from_S_profile(psi)
central_charge, const, res = tenpy.tools.fit.central_charge_from_S_profile(psi)
fit = tenpy.tools.fit.entropy_profile_from_CFT(bonds + 0.5, psi.L, central_charge,_
fit = tenpy.tools.fit.entropy_profile_from_CFT(bonds + 0.5, psi.L, central_charge,_
Gconst)
Gconst)
print(f"extraced central charge {central_charge:.5f} with residuum {res:.2e}")
print(f"extraced central charge {central_charge:.5f} with residuum {res:.2e}")
print("(Expect central charge = 0.5 for the transverse field Ising model.)")
print("(Expect central charge = 0.5 for the transverse field Ising model.)")
plt.plot(bonds, fit, label=f"fit with $c={central_charge:.3f}$")
plt.plot(bonds, fit, label=f"fit with $c={central_charge:.3f}$")
plt.xlabel("bond")
plt.xlabel("bond")
plt.ylabel("entanglement entropy $S$")

```
plt.ylabel("entanglement entropy $S$")
```

```
plt.legend()
plt.show()
extraced central charge 0.50788 with residuum 1.95e-09
(Expect central charge = 0.5 for the transverse field Ising model.)
```


[ ]:

### 11.3.3 Lattices: visualization and further examples

This notebook demonstrates a few ways to visualize lattices and couplings inside the lattice.
[1]:

```
import tenpy
import numpy as np
import matplotlib.pyplot as plt
from tenpy.models import lattice
```

In the following, we will focus on the Honeycomb lattice as an example with a non-trivial unit cell. If you want to try it out yourself for a different lattice, simply adjust the following alias and re-run all the cells below:
[3]((%5Cmathrm%7BL%7D=100)): MyLattice = lattice. Honeycomb
$\mathrm{Lu}=$ MyLattice.Lu $\#=2=$ the number of sites in the unit cell
fig_args = dict(figsize=(7, 5), dpi=150) \# make figures a bit larger

## plotting the lattice itself

To get started, let's recall that a lattice consists of a unit cell that is repeated in the directions of the basis. The following plot visualizes the first unit cell and basis and plots the sites in the whole lattice. For the Honeycomb lattice, we have two different sites in the unit cell, which get visualized by different markers.

```
[4]: lat = MyLattice(5, 4, sites=None, bc='periodic')
```



We can also plot the nearest- and next-nearest-neighbor bonds:

```
[6]: plt.figure(**fig_args)
```

ax = plt.gca()
lat.plot_sites(ax)
lat.plot_coupling (ax)
lat.plot_coupling(ax, lat.pairs['next_nearest_neighbors'], linestyle=':', color='r')
ax.set_aspect('equal')


If you have a 1D MPS, it's winding through the lattice is defined by the order (See the userguide with "Details on the lattice geometry"). You can plot it as follows:

```
plt.figure(**fig_args)
ax = plt.gca()
lat.plot_sites(ax)
lat.plot_order(ax)
```



## Visually verifying pairs of couplings by plotting them

In this section, we visually verify that the lattice pairs like the nearest_neighbors are indeed what they claim to be. To acchieve this, we first get all the possible distances of them, plot circles with these distances and lines connecting the points for each distance.

Then, you have to stare at the plots and verify that these couplings include all pairs you want to have.

```
[8]: lat = MyLattice(12, 12, sites=None, bc='periodic') # the lattice to plot
    lat_pairs = lat.pairs # the coupling pairs to plot
[9]: # get distances of the couplings
    dist_pair = {}
    for pair in lat_pairs:
        #print(pair)
        dist = None
        for u1, u2, dx in lat_pairs[pair]:
            d = lat.distance(u1, u2, dx)
            #print(ul, u2, dx, d)
            if dist is None:
                dist = d
            dist_pair[d] = pair
        else:
            assert abs(dist-d) < 1.e-14
dists = sorted(dist_pair.keys())
if len(dists) != len(lat_pairs):
    raise ValueError("no unique mapping dist -> pair")
```

[10]:

```
print("(dist) (pairs)")
for d in dists:
    print("{0:.6f} {I}".format(d, dist_pair[d]))
(dist) (pairs)
0.577350 nearest_neighbors
1.000000 next_nearest_neighbors
1.154701 next_next_nearest_neighbors
1.527525 fourth_nearest_neighbors
1.732051 fifth_nearest_neighbors
```

[11](matplotlib.legend.Legend%20at%200x7f1018760e80):

```
colors = [plt.cm.viridis(r/dists[-1]) for r in dists]
centers = np.array([[3, 3, 0], [8, 3, 1], [3, 8, 2], [8, 8, 3]]) # one center for_
\hookrightarroweach site in the unit cell
us = list(range(Lu))
fig = plt.figure(**fig_args)
ax = plt.gca()
lat.plot_sites(ax, markersize=1.3)
for u, center in zip(us, centers):
    center = lat.position(center)
    for r, c in zip(dists, colors):
        circ = plt.Circle(center, r, fill=False, color=c)
        ax.add_artist(circ)
ax.set_aspect (1.)
t = ax.set_title("distances: " + ' '.join(['{0:.2f}'.format(d) for d in dists]))
```

[12]:
distances: 0.581 .001 .151 .531 .73

[13]:

```
for dist in dists:
    pair_name = dist_pair[dist]
    print(dist, pair_name)
    pairs = lat.pairs[pair_name]
    fig = plt.figure(**fig_args)
    ax = plt.gca()
    lat.plot_sites(ax, markersize=1.3)
    for u, center in zip(us, centers):
        center = lat.position(center)
        for r, c in zip(dists, colors):
```

(continued from previous page)

```
            circ = plt.Circle(center, r, fill=False, color=c)
            ax.add_artist(circ)
    pairs_with_reverse = pairs + [(u2, u1, -np.array(dx)) for u1, u2, dx in pairs]
    for u1, u2, dx in pairs_with_reverse:
        print(u1, u2, dx, lat.distance(u1, u2, dx))
        start = centers[u1]
    end = start.copy()
    end[-1] = u2
    end[:-1] = start[:-1] + dx
    x1, y1 = lat.position(start)
    x2, y2 = lat.position(end)
    ax.arrow(x1, y1, x2-x1, y2 - y1)
for u in us:
    x, y = lat.position(centers[u])
    number = lat.count_neighbors(u, pair_name)
    ax.text(x, y, str(number), color='r')
ax.set_aspect(1.)
ax.set_title(pair_name + ' distance = {0:.3f}'.format(dist))
```

```
0.5773502691896258 nearest_neighbors
```

0.5773502691896258 nearest_neighbors
0 1 [0 0] 0.5773502691896258
0 1 [0 0] 0.5773502691896258
1 0 [1 0] 0.5773502691896257
1 0 [1 0] 0.5773502691896257
1 0 [0 1] 0.5773502691896258
1 0 [0 1] 0.5773502691896258
1 0 [0 0] 0.5773502691896258
1 0 [0 0] 0.5773502691896258
0 1 [-1 0] 0.5773502691896257
0 1 [-1 0] 0.5773502691896257
0 1 [ 0 -1] 0.5773502691896258
0 1 [ 0 -1] 0.5773502691896258
0.9999999999999999 next_nearest_neighbors
0.9999999999999999 next_nearest_neighbors
0 0 [1 0] 0.9999999999999999
0 0 [1 0] 0.9999999999999999
0 0 [0 1] 1.0
0 0 [0 1] 1.0
0 0 [ 1 -1] 0.9999999999999999
0 0 [ 1 -1] 0.9999999999999999
1 1 [1 0] 0.9999999999999999
1 1 [1 0] 0.9999999999999999
1 1 [0 1] 1.0
1 1 [0 1] 1.0
1 [ [ 1 -1] 0.9999999999999999
1 [ [ 1 -1] 0.9999999999999999
0 0 [-1 0] 0.9999999999999999
0 0 [-1 0] 0.9999999999999999
0 0 [ 0 -1] 1.0
0 0 [ 0 -1] 1.0
0 0 [-1 1] 0.9999999999999999
0 0 [-1 1] 0.9999999999999999
1 1 [-1 0] 0.9999999999999999
1 1 [-1 0] 0.9999999999999999
1 1 [ 0 -1] 1.0
1 1 [ 0 -1] 1.0
1 1 [-1 1] 0.9999999999999999
1 1 [-1 1] 0.9999999999999999
1.1547005383792515 next_next_nearest_neighbors
1.1547005383792515 next_next_nearest_neighbors
1 0 [1 1] 1.1547005383792515
1 0 [1 1] 1.1547005383792515
0 1 [-1 1] 1.1547005383792515
0 1 [-1 1] 1.1547005383792515
0 1 [ 1 -1] 1.1547005383792515
0 1 [ 1 -1] 1.1547005383792515
0 1 [-1 -1] 1.1547005383792515
0 1 [-1 -1] 1.1547005383792515
1 0 [ 1 -1] 1.1547005383792515
1 0 [ 1 -1] 1.1547005383792515
1 0 [-1 1] 1.1547005383792515
1 0 [-1 1] 1.1547005383792515
1.5275252316519468 fourth_nearest_neighbors
1.5275252316519468 fourth_nearest_neighbors
0 1 [0 1] 1.5275252316519468
0 1 [0 1] 1.5275252316519468
0 1 [1 0] 1.5275252316519465
0 1 [1 0] 1.5275252316519465
0 1 [ 1 -2] 1.5275252316519465
0 1 [ 1 -2] 1.5275252316519465
0 1 [ 0 -2] 1.5275252316519468
0 1 [ 0 -2] 1.5275252316519468
0 1 [-2 0] 1.5275252316519465
0 1 [-2 0] 1.5275252316519465
0 1 [-2 1] 1.5275252316519465
0 1 [-2 1] 1.5275252316519465
1 0 [ 0 -1] 1.5275252316519468
1 0 [ 0 -1] 1.5275252316519468
1 0 [-1 0] 1.5275252316519465
1 0 [-1 0] 1.5275252316519465
1 0 [l-1 2] 1.5275252316519465
1 0 [l-1 2] 1.5275252316519465
1 0 [0 2] 1.5275252316519468

```
1 0 [0 2] 1.5275252316519468
```

```
1 0 [2 0] 1.5275252316519465
1 0 [ 2 -1] 1.5275252316519465
1.7320508075688772 fifth_nearest_neighbors
0 0 [1 1] 1.7320508075688772
0 0 [ 2 -1] 1.7320508075688772
0 0 [l-1 2] 1.7320508075688772
1 1 [1 1] 1.7320508075688772
1 [ [ 2 -1] 1.7320508075688772
1 1 [-1 2] 1.7320508075688772
0 0 [-1 -1] 1.7320508075688772
0 0 [l-2 1] 1.7320508075688772
O 0 [ 1 -2] 1.7320508075688772
1 1 [-1 -1] 1.7320508075688772
1 1 [-2 1] 1.7320508075688772
1 [ [ -2] 1.7320508075688772
```



next_next_nearest_neighbors distance $=1.155$

fourth_nearest_neighbors distance $=1.528$

fifth_nearest_neighbors distance $=1.732$


### 11.3.4 Toric Code

This notebook shows how to implement Kitaev's Toric Code model on an infinite cylinder geometry. The Hamiltonian of the Toric Code is:

$$
\mathcal{H}=-\sum_{p} B_{p}-\sum_{s} A_{s}
$$

where $B_{p}=\prod_{i \in p} \sigma_{i}^{z}$ denotes the product of $\sigma^{z}$ around an elementary plaquette and $A_{s}=\prod_{i \in s} \sigma_{i}^{x}$ denotes the product of $\sigma^{x}$ on the four links sticking out of a site of the lattice. The $B_{p}$ and $A_{s}$ operators commute with each other and have eigenvalues $\pm 1$, hence the ground state will have eigenvalue +1 for each of them.

On an infinite cylinder, the model exhibits a four-fold degeneracy of the ground state, characterized by the different values $\pm 1$ of the incontractible Wilson and t'Hooft loops. These loop operators are defined by the products

$$
W=\prod_{\text {vert links }} \sigma^{z} \quad H=\prod_{\text {hor links }} \sigma^{x}
$$

around the cylinder.
TeNPy already implements this hamiltonian in tenpy.models.toric_code.ToricCode. This notebook demonstrates how to extend the existing model by additinoal terms in the Hamiltonian and convenient functions of the loop operators for simplified measurements. Then we run iDMRG to obtain the four different ground states and check that they are indeed orthogonal and degenerate.

```
import numpy as np
import scipy
import matplotlib.pyplot as plt
np.set_printoptions(precision=5, suppress=True, linewidth=120)
import tenpy
import tenpy.linalg.np_conserved as npc
from tenpy.algorithms.dmrg import TwoSiteDMRGEngine
from tenpy.networks.mps import MPS
from tenpy.networks.terms import TermList
from tenpy.models.toric_code import ToricCode, DualSquare
from tenpy.models.lattice import Square
```

Let's plot the DualSquare lattice first to get the geometry clear. The unit cell consists of two sites, the blue one on the vertical links $(u=0)$, and the orange one on the horizontal links $(u=1)$. We also plot lines illustrating where the Wilson and t'Hooft loops go.

```
[3]: plt.figure(figsize=(7, 5))
ax = plt.gca()
lat = DualSquare(4, 4, None, bc='periodic')
sq = Square(4, 4, None, bc='periodic')
sq.plot_coupling(ax, linewidth=3.)
ax.plot([2., 2.], [-0.5, 4.3], 'r:', linewidth=5., label="Wilson")
ax.plot([2.5, 2.5], [-0.5, 4.3], 'b:', linewidth=5., label="t'Hooft")
lat.plot_sites(ax)
lat.plot_basis(ax, origin=-0.25*(lat.basis[0] + lat.basis[1]))
ax.set_aspect('equal')
ax.set_xlim(-1, 6)
ax.set_ylim(-1)
ax.legend()
plt.show()
```



## Extending the existing model

We will implement additional parameters which allow to optionally add terms for the Wilson and t'Hooft loop operators $W, H$ defined above,

$$
H \rightarrow H-\mathrm{J}_{\mathrm{WL}} W-\mathrm{J}_{\mathrm{HL}} H
$$

Note that we only want to add a single loop for each of them, not one at each possible starting point. Hence, the correct method is add_local_term, not add_multi_coupling_term or add_coupling_term.
The sign of $\mathrm{J}_{\mathrm{WL}}$, $\mathrm{J}_{\mathrm{HL}}$ will allow us to select sectors with $\langle\psi| W|\psi\rangle= \pm 1$ and $\langle\psi| H|\psi\rangle= \pm 1$, respectively. Note that so far the constraints commute with the Hamiltonian and with each other. However, this is no longer the case if we add another global field $H \rightarrow H-\mathrm{h} \sum_{i} \sigma_{i}^{z}$, which makes the Hamiltonian no longer exactly solvable.
[4]:

```
class ExtendedToricCode(ToricCode):
    def init_terms(self, model_params):
    ToricCode.init_terms(self, model_params) # add terms of the original_
\hookrightarrowToricCode model
    Ly = self.lat.shape[1]
    J_WL = model_params.get('J_WL', 0.)
    J_HL = model_params.get('J_HL', 0.)
    # unit-cell indices:
    # u=0: vertical links
    # u=1: horizontal links
    # Wilson Loop
    x, u = 0, 0 # vertical links
    self.add_local_term(-J_WL, [('Sigmaz', [x, y, u]) for y in range(Ly)])
    # t'Hooft Loop
    x, u = 0, 1 # vertical links
    self.add_local_term(-J_HL, [('Sigmax', [x, y, u]) for y in range(Ly)])
```

(continues on next page)

```
    h = model_params.get('h', 0.)
    for u in range(2):
        self.add_onsite(-h, u, 'Sigmaz')
    def wilson_loop_y(self, psi):
    """Measure wilson loop around the cylinder."""
    Ly = self.lat.shape[1]
    x, u = 0, 0 # vertical links
    W = TermList.from_lattice_locations(self.lat, [[("Sigmaz", [x, y, u]) for y in
->range (Ly)] ])
    return psi.expectation_value_terms_sum(W) [0]
    def hooft_loop_y(self, psi):
    """Measure t'Hooft loop around the cylinder."""
    Ly = self.lat.shape[1]
    x, u = 0, 1 # horizontal links
    H = TermList.from_lattice_locations(self.lat, [[("Sigmax",[x, y, u]) for y in
\hookrightarrowrange(Ly)] ])
    return psi.expectation_value_terms_sum(H) [0]
```


## iDMRG

We now run the iDMRG algorithm for the four different sectors to obtain the four degenerate ground states of the system.

To reliably get the 4 different ground states, we will add the energetic constraints for the loops. In this case, the original Hamiltonian will completely commute with the loop operators, such that you directly get the correct ground state. However, this is a very peculiar case - in general, e.g. when you add the external field with $h \neq 0$, this will no longer be the case. We hence demonstrate a more "robust" way of getting the four different ground states (at least if you have an idea which operator distinguishes them): For each sector, we run DMRG twice: first with the energy penalties for the $W, H$ loops to get an initial $|\psi\rangle$ in the desired sector. Then we run DMRG again for the clean toric code model, without the $W, H$ added, to make sure we have a ground state of the clean model.

```
[5]: dmrg_params = {
    'mixer': True,
    'trunc_params': {'chi_max': 64,
                'svd_min': 1.e-8},
    'max_E_err': 1.e-8,
    'max_S_err': 1.e-7,
    'N_sweeps_check': 4,
    'max_sweeps':24,
    'verbose': 0,
}
model_params = {
    'Lx': 1, 'Ly': 4, # Ly is set below
    'bc_MPS': "infinite",
    'conserve': None,
    'verbose': 0,
}
def run_DMRG(Ly, J_WL, J_HL, h=0.):
    print("="*80)
    print(f"Start iDMRG for Ly={Ly:d}, J_WL={J_WL:.2f}, J__HL={J_HL:.2f}, h={h:.2f}")
```

```
model_params_clean = model_params.copy()
model_params_clean['Ly'] = Ly
model_params_clean['h'] = h
model_clean = ExtendedToricCode(model_params_clean)
model_params_seed = model_params_clean.copy()
model_params_seed['J_WL'] = J_WL
model_params_seed['J_HL'] = J_HL
model = ExtendedToricCode(model_params_seed)
psi = MPS.from_lat_product_state(model.lat, [[["up"]]])
eng = TwoSiteDMRGEngine(psi, model, dmrg_params)
EO, psi = eng.run()
WL = model.wilson_loop_y(psi)
HL = model.hooft_loop_y(psi)
print(f"after first DMRG run: <psi|W|psi> = {WL: .3f}")
print(f"after first DMRG run: <psi|H|psi> = {HL: .3f}")
print(f"after first DMRG run: E (including W/H loops) = {E0:.10f}")
EO_clean = model_clean.H_MPO.expectation_value(psi)
print(f"after first DMRG run: E (excluding W/H loops) = {E0_clean:.10f}")
# switch to model without Wilson/Hooft loops
eng.init_env(model=model_clean)
E1, psi = eng.run()
WL = model_clean.wilson_loop_y(psi)
HL = model_clean.hooft_loop_y(psi)
print(f"after second DMRG run: <psi|W|psi> = {WL: . 3f}")
print(f"after second DMRG run: <psi|H|psi> = {HL: . 3f }")
print(f"after second DMRG run: E (excluding W/H loops) = {E1:.10f}")
print("max chi: ", max(psi.chi))
return {'psi': psi,
    'model': model_clean,
    'EO': EO, 'EO_clean': E0_clean, 'E': E1,
    'WL': WL, 'HL': HL}
print("="*80)
```

[6]: results_loops $=$ \{\}

```
for J_WL in [-5., 5.]:
```

    for J_HL in [-5., 5.]:
        results_loops[(J_WL, J_HL)] = run_DMRG(4, J_WL, J_HL)
    
Start iDMRG for $L y=4, J \_W L=-5.00$, J_HL=-5.00, $h=0.00$
after first DMRG run: <psi|W|psi> $=-1.000$
after first DMRG run: <psi|H|psi> $=-1.000$
after first DMRG run: E (including W/H loops) $=-2.2500000000$
after first DMRG run: E (excluding W/H loops) $=-1.0000000000$
after second DMRG run: <psi|W|psi> $=-1.000$
after second DMRG run: <psi|H|psi> $=-1.000$
after second DMRG run: E (excluding W/H loops) $=-1.0000000000$
max chi: 8

Start iDMRG for $L y=4, ~ J \_W L=-5.00$, J_HL=5.00, h=0.00
after first DMRG run: <psi|W|psi> $=-1.000$
(continues on next page)

```
after first DMRG run: <psi|H|psi> = 1.000
after first DMRG run: E (including W/H loops) = -2.2500000000
after first DMRG run: E (excluding W/H loops) = -1.0000000000
after second DMRG run: <psi|W|psi> = -1.000
after second DMRG run: <psi|H|psi> = 1.000
after second DMRG run: E (excluding W/H loops) = -1.0000000000
max chi: 8
=======================================================================================
Start iDMRG for Ly=4, J_WL=5.00, J_HL=-5.00, h=0.00
after first DMRG run: <psi|W|psi> = 1.000
after first DMRG run: <psi|H|psi> = -1.000
after first DMRG run: E (including W/H loops) = -2.2500000000
after first DMRG run: E (excluding W/H loops) = -1.0000000000
after second DMRG run: <psi|W|psi> = 1.000
after second DMRG run: <psi|H|psi> = -1.000
after second DMRG run: E (excluding W/H loops) = -1.0000000000
max chi: 8
======================================================================================
Start iDMRG for Ly=4, J_WL=5.00, J_HL=5.00, h=0.00
after first DMRG run: <psi|W|psi> = 1.000
after first DMRG run: <psi|H|psi> = 1.000
after first DMRG run: E (including W/H loops) = -2.2500000000
after first DMRG run: E (excluding W/H loops) = -1.0000000000
after second DMRG run: <psi|W|psi> = 1.000
after second DMRG run: <psi|H|psi> = 1.000
after second DMRG run: E (excluding W/H loops) = -1.0000000000
max chi: 8
```

As we can see from the output, the idea worked and we indeed get the desired expectation values for the W/H loops. Also, we can directly see that the ground states are degenerate in energy.
To check that we found indeed four orhogonal ground states, let's calculate mutual overlaps $\left\langle\psi_{i} \mid \psi_{j}\right\rangle$ for $i, j \in$ range(4):
[7]:

```
psi_list = [res['psi'] for res in results_loops.values()]
overlaps= [[psi_i.overlap(psi_j) for psi_j in psi_list] for psi_i in psi_list]
print("overlaps")
print(np.array(overlaps))
```

```
overlaps
[[ 1.+0.j 0.+0.j -0.+0.j 0.-0.j]
    [ 0.+0.j 1.+0.j 0.+0.j -0.+0.j]
    [-0.+0.j 0.+0.j 1.+0.j 0.+0.j]
    [ 0.-0.j -0.+0.j 0.+0.j 1.+0.j]]
```

The correlation length is almost vanishing, as it should be:
[8]: print("Correlation lengths: ")
model $=$ results_loops [(+5., +5.)]['model']
print(np.array([psi.correlation_length() / model.lat.N_sites_per_ring for psi in psi_
↔list]))

Correlation lengths:
$\left[\begin{array}{llll}0.0294 & 0.02924 & 0.02915 & 0.02876\end{array}\right]$

Intermezzo: Let's quickly check for a single case that this still works for non-zero $h=0.1$, where the t'Hooft loop no longer commutes with the Hamiltonian:

```
res_h = run_DMRG(4, +5., -5., 0.1)
====================================================================================
Start iDMRG for Ly=4, J_WL=5.00, J_HL=-5.00, h=0.10
after first DMRG run: <psi|W|psi> = 1.000
after first DMRG run: <psi|H|psi> = -1.000
after first DMRG run: E (including W/H loops) = -2.2516147149
after first DMRG run: E (excluding W/H loops) = -1.0018727133
after second DMRG run: <psi|W|psi> = 1.000
after second DMRG run: <psi|H|psi> = -0.995
after second DMRG run: E (excluding W/H loops) = -1.0025283892
max chi: 64
```

The t'Hooft loop no longer is exactly 1 in the ground state, since it no longer commutes with the Hamiltonian, but it is still close to one, so it still distinguishes the degenerate states. As expected, the local perturbation of the field can not destry the global, topological nature of the ground state (manifold).

## Topological entanglement entropy

Back to $h=0$. For the toric code, the half-cylinder entanglement entropy should be $\left(L_{y}-1\right) \log 2$. Indeed, we get the correct value at bond 0 :
[10]:

```
print("Entanglement entropies / log(2)")
print(np.array([psi.entanglement_entropy()[0]/np.log(2) for psi in psi_list]))
Entanglement entropies / log(2)
[3. 3. 3. 3.]
```

Recall that bond 0 is usually a sensible cut for infinite DMRG on a common lattice. This is especially important if you want to extract the topological entanglement entropy $\gamma$ from a fit $S\left(L_{y}\right)=\alpha L_{y}-\gamma$, as corners in the cut can also contribute to $\gamma$.

Let us demonstrate the extraction of $\gamma=\log (2)$ for the toric code:
[11](matplotlib.legend.Legend%20at%200x7f1018760e80):

```
model_params['order'] = 'Cstyle' # The effect doesn't appear with the "default",u
\hookrightarrowordering for the toric code.
# This is also a hint that you need bond 0: you want something independent of what u
Gorder you choose
# inside the MPS unit cell.
results_Ly = {}
for Ly in range(2, 7):
    results_Ly[Ly] = run_DMRG(Ly, 5., 5.)
```


Start iDMRG for $L y=2$, J_WL=5.00, J_HL=5.00, h=0.00
after first DMRG run: <psi|W|psi> = 1.000
after first DMRG run: <psi|H|psi> = 1.000
after first DMRG run: E (including $W / H$ loops) $=-3.5000000000$
after first DMRG run: E (excluding W/H loops) $=-1.0000000000$
after second DMRG run: <psi|W|psi> $=1.000$
after second DMRG run: <psi|H|psi> $=1.000$
after second DMRG run: E (excluding W/H loops) $=-1.0000000000$
max chi: 4

Start iDMRG for $L y=3, J \_W L=5.00$, J_HL=5.00, $h=0.00$
(continues on next page)

```
(continued from previous page)
```

```
after first DMRG run: <psi|W|psi> = 1.000
```

after first DMRG run: <psi|W|psi> = 1.000
after first DMRG run: <psi|H|psi> = 1.000
after first DMRG run: <psi|H|psi> = 1.000
after first DMRG run: E (including W/H loops) = -2.6666666667
after first DMRG run: E (including W/H loops) = -2.6666666667
after first DMRG run: E (excluding W/H loops) = -1.0000000000
after first DMRG run: E (excluding W/H loops) = -1.0000000000
after second DMRG run: <psi|W|psi> = 1.000
after second DMRG run: <psi|W|psi> = 1.000
after second DMRG run: <psi|H|psi> = 1.000
after second DMRG run: <psi|H|psi> = 1.000
after second DMRG run: E (excluding W/H loops) = -1.0000000000
after second DMRG run: E (excluding W/H loops) = -1.0000000000
max chi: 8
max chi: 8
========================================================================================
========================================================================================
Start iDMRG for Ly=4, J_WL=5.00, J_HL=5.00, h=0.00
Start iDMRG for Ly=4, J_WL=5.00, J_HL=5.00, h=0.00
after first DMRG run: <psi|W|psi> = 1.000
after first DMRG run: <psi|W|psi> = 1.000
after first DMRG run: <psi|H|psi> = 1.000
after first DMRG run: <psi|H|psi> = 1.000
after first DMRG run: E (including W/H loops) = -2.2500000000
after first DMRG run: E (including W/H loops) = -2.2500000000
after first DMRG run: E (excluding W/H loops) = -1.0000000000
after first DMRG run: E (excluding W/H loops) = -1.0000000000
after second DMRG run: <psi|W|psi> = 1.000
after second DMRG run: <psi|W|psi> = 1.000
after second DMRG run: <psi|H|psi> = 1.000
after second DMRG run: <psi|H|psi> = 1.000
after second DMRG run: E (excluding W/H loops) = -1.0000000000
after second DMRG run: E (excluding W/H loops) = -1.0000000000
max chi: 16
max chi: 16
========================================================================================
========================================================================================
Start iDMRG for Ly=5, J_WL=5.00, J_HL=5.00, h=0.00
Start iDMRG for Ly=5, J_WL=5.00, J_HL=5.00, h=0.00
after first DMRG run: <psi|W|psi> = 1.000
after first DMRG run: <psi|W|psi> = 1.000
after first DMRG run: <psi|H|psi> = 1.000
after first DMRG run: <psi|H|psi> = 1.000
after first DMRG run: E (including W/H loops) = -2.0000000000
after first DMRG run: E (including W/H loops) = -2.0000000000
after first DMRG run: E (excluding W/H loops) = -1.0000000000
after first DMRG run: E (excluding W/H loops) = -1.0000000000
after second DMRG run: <psi|W|psi> = 1.000
after second DMRG run: <psi|W|psi> = 1.000
after second DMRG run: <psi|H|psi> = 1.000
after second DMRG run: <psi|H|psi> = 1.000
after second DMRG run: E (excluding W/H loops) = -1.0000000000
after second DMRG run: E (excluding W/H loops) = -1.0000000000
max chi: 32
max chi: 32
=========================================================================================
=========================================================================================
Start iDMRG for Ly=6, J_WL=5.00, J_HL=5.00, h=0.00
Start iDMRG for Ly=6, J_WL=5.00, J_HL=5.00, h=0.00
after first DMRG run: <psi|W|psi> = 1.000
after first DMRG run: <psi|W|psi> = 1.000
after first DMRG run: <psi|H|psi> = 1.000
after first DMRG run: <psi|H|psi> = 1.000
after first DMRG run: E (including W/H loops) = -1.8333333333
after first DMRG run: E (including W/H loops) = -1.8333333333
after first DMRG run: E (excluding W/H loops) = -1.0000000000
after first DMRG run: E (excluding W/H loops) = -1.0000000000
after second DMRG run: <psi|W|psi> = 1.000
after second DMRG run: <psi|W|psi> = 1.000
after second DMRG run: <psi|H|psi> = 1.000
after second DMRG run: <psi|H|psi> = 1.000
after second DMRG run: E (excluding W/H loops) = -1.0000000000
after second DMRG run: E (excluding W/H loops) = -1.0000000000
max chi: 64
max chi: 64
[12]:

```
```

Lys = np.array(list(results_Ly.keys()))

```
Lys = np.array(list(results_Ly.keys()))
S0s = np.array([res['psi'].entanglement_entropy()[0] for res in results_Ly.values()])
S0s = np.array([res['psi'].entanglement_entropy()[0] for res in results_Ly.values()])
Sls = np.array([res['psi'].entanglement_entropy()[2] for res in results_Ly.values()])
Sls = np.array([res['psi'].entanglement_entropy()[2] for res in results_Ly.values()])
plt.figure(figsize=(7, 5))
plt.figure(figsize=(7, 5))
ax = plt.gca()
ax = plt.gca()
ax.plot(Lys, S0s / np.log(2), 'bo', label='data at bond 0')
ax.plot(Lys, S0s / np.log(2), 'bo', label='data at bond 0')
ax.plot(Lys, S1s / np.log(2), 'rs', label='data at wrong bond')
ax.plot(Lys, S1s / np.log(2), 'rs', label='data at wrong bond')
Lys_all = np.arange(0., max(Lys) + 2.)
Lys_all = np.arange(0., max(Lys) + 2.)
ax.plot(Lys_all, Lys_all - 1., '-', label='expected: $L_y - 1$')
ax.plot(Lys_all, Lys_all - 1., '-', label='expected: $L_y - 1$')
ax.axhline(0., linestyle='--', color='k')
ax.axhline(0., linestyle='--', color='k')
ax.set_xlabel('$L_y$')
ax.set_xlabel('$L_y$')
ax.set_ylabel('$S / log(2)$')
ax.set_ylabel('$S / log(2)$')
ax.legend()
ax.legend()
plt.show()
```

plt.show()

```


As expected, We find the scaling \(S\left(L_{y}\right)=\log (2) L_{y}-\log (2)\) at bond 0 , indicating that \(\gamma=\log (2)\). Cutting the MPS at bond 1 (red points) is a "wired" cut of the cylinder and does not show the expected scaling.

\section*{TWELVE}

\section*{TROUBLESHOOTING AND FAQ}

I updated to a new version and now I get and error/warning. Take a look at the section "Backwards incomptatible changes" in /changelog of the corresponding versions since when you updated.

Where did all the output go? Take a look at Logging and terminal output.
What are possible parameters for ...? See Parameters and options.
How can I set the number of threads TeNPy is using? Most algorithms in TeNPy don't use any parallelization besides what the underlying BLAS provides, so that depends on how you installed TeNPy, numpy and scipy! Using for example an export OMP_NUM_THREADS=4 should limit it to 4 threads under usual setups, but you might also want to export MKL_NUM_THREADS=4 instead, if you are sure that you are using MKL.

Why is TeNPy not respecting MKL_NUM_THREADS? It might be that it is not using MKL. On linux, check whether you have installed a pip version of numpy or scipy in \$HOME/.local/lib/python3.* Those packages do not use MKL - you would need to install numpy and scipy from conda. If you use the conda-forge channel as recommended in the installation, also make sure that you select the BLAS provided by MKL, see the note in install/conda.rst.

\subsection*{12.1 I get an error when...}
... I try to measure \(S x\) _i \(S x\) _j correlations in a state with \(S z\) conseration. Right now this is not possible. See the basic facts in Charge conservation with np_conserved.

\subsection*{12.2 I get a warning about ...}
... an unused parameter. Make sure that you don't have a typo and that it is in the right parameter set! Also, check the logging output whether the parameter was actually used. For further details, see Parameters and options

\section*{LITERATURE AND REFERENCES}

This is a (by far non-exhaustive) list of some references for the various ideas behind the code. They can be cited like this:
- [TeNPyNotes] for TeNPy/software related sources
- [[white 1992]] (lowercase first-author + year) for entries from literature.bib.

\subsection*{13.1 TeNPy related sources}
[TeNPyNotes] are lecture notes, meant as an introduction to tensor networks (focusing on MPS), and introduced TeNPy to the scientific community by giving examples how to call the algorithms in TeNPy. [TeNPySource] is the location of the source code, and the place where you can report bugs. We have split example notebooks into [TeNPyNotebooks] to keep the git history of the original repository clean. [TeNPyDoc] is where the documentation is hosted online. [TeNPyForum] is the place where you can ask questions and look for help when you are stuck with implementing something.

\subsection*{13.2 Software-related}

The following links are not physics-related, but are good to know if you want to work with TeNPy (or more generally Python).

\subsection*{13.3 General reading}
[[schollwoeck2011]] is an extensive introduction to MPS, DMRG and TEBD with lots of details on the implementations, and a classic read, although a bit lengthy. Our [TeNPyNotes] are a shorter summary of the important concepts, similar as [[orus2014]]. [[paeckel2019]] is a very good, recent review focusing on time evolution with MPS. The lecture notes of [[eisert2013]] explain the area law as motivation for tensor networks very well. PEPS are for example reviewed in [[verstraete2008]], [[eisert2013]] and [[orus2014]]. [[stoudenmire2012]] reviews the use of DMRG for 2D systems. [[cirac2009]] discusses the different groups of tensor network states.

\subsection*{13.4 Algorithm developments}
[[white1992], [white1993]] is the invention of DMRG, which started everything. [[vidal2004]] introduced TEBD. [[white2005]] and [[hubig2015]] solved problems for single-site DMRG. [[mcculloch2008]] was a huge step forward to solve convergence problems for infinite DMRG. [[singh2010], [singh2011]] explain how to incorporate Symmetries. [[haegeman2011]] introduced TDVP, again explained more accessible in [[haegeman2016]]. [[zaletel2015]] is another standard method for time-evolution with long-range Hamiltonians. [[karrasch2013]] gives some tricks to do finitetemperature simulations (DMRG), which is a bit extended in [[hauschild2018a]]. [[vidal2007]] introduced MERA. The scaling \(S=c / 6 \log (\chi)\) at a 1D critical point is explained in [[pollmann2009]].

\subsection*{13.5 References}

\section*{PAPERS USING TENPY}

This page collects papers using (and citing) the TeNPy library, both as an inspiration what can be done, as well as to keep track of the usage, such that we can see how useful our work is to the community. It keeps us motivated!

To include your own work, you can either fill out this template on github, or you can directly add your citation in this Zotero online library (and notify us about it or just wait).

Entries in the following list are sorted by year-author.
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\section*{CONTRIBUTING}

There are lots of things where you can help, even if you don't wont to dig deep into the source code. You are welcome to do any of the following things, all of them are very helpful!
- Report bugs and problems, such that they can be fixed.
- Implement new models.
- Update and extend the documentation.
- Provide examples of how to use TeNPy.
- Give feedback on how you like TeNPy and what you would like to see improved.
- Help fixing bugs.
- Help fixing minor issues.
- Extend the functionality by implementing new functions, methods, and algorithms.

The code is maintained in a git repository, the official repository is on github. Even if you're not yet on the developer team, you can still submit pull requests on github. If you're unsure how or what to do, you can ask for help in the [TeNPyForum]. If you want to become a member of the developer team, just ask ;-)

\section*{Thank You!}

\subsection*{15.1 Coding Guidelines}

To keep consistency, we ask you to comply with the following guidelines for contributions. However, these are just guidelines - it still helps if you contribute something, even if doesn't follow these rules ;-)
- Use a code style based on PEP 8. The git repo includes a config file. style.yapf for the python package yapf. yapf is a tool to auto-format code, e.g., by the command yapf -i some/file (-i for "in place"). We run yapf on a regular basis on the github main branch. If your branch diverged, it might help to run yapf before merging.

Note: Since no tool is perfect, you can format some regions of code manually and enclose them with the special comments \# yapf: disable and \# yapf: enable.
- Every function/class/module should be documented by its doc-string, see PEP 257. We auto-format the docstrings with docformatter on a regular basis.

Additional documentation for the user guide is in the folder doc/.
The documentation uses reStructuredText. If you are new to reStructuredText, read this introduction. We use the numpy style for doc-strings (with the napoleon extension to sphinx). You can read abouth them in these

Instructions for the doc strings. In addition, you can take a look at the following example file. Helpful hints on top of that:
```

r"""<- this r makes me a raw string, thus '\' has no special meaning.
Otherwise you would need to escape backslashes, e.g. in math formulas.
You can include cross references to classes, methods, functions, modules like
:class:` ~tenpy.linalg.np_conserved.Array`, :meth:` ~tenpy.linalg.np_conserved. 4Array.to_ndarray`,
:func:`tenpy.tools.math.toiterable`, :mod:`tenpy.linalg.np_conserved`.
The ~ in the beginning makes only the last part of the name appear in thes
@enerated documentation.
Documents of the userguide can be referenced with :doc:`/intro_npc` even from,
sinside the doc-strings.
You can also cross-link to other documentations, e.g. :class:`numpy.ndarray`, :
func`scipy.linalg.svd` and :mod: will work.
Moreover, you can link to github issues, arXiv papers, dois, and topics in thes
Community forum with
e.g. :issue:`5`, :arxiv:`1805.00055`, :doi:`10.1000/1` and :forum:`3`.
Citations from the literature list can be cited as :cite:`whitelg92` using thev
\hookrightarrowbibtex key.
Write inline formulas as :math:`H /\Psi\rangle = E /\Psi\rangle` or displayedr
\hookrightarrowequations as
.. math : :
e^{i\pi}+1=0
In doc-strings, math can only be used in the Notes section.
To refer to variables within math, use '\mathtt{varname}'.
.. todo : :
This block can describe things which need to be done and is automaticallyu
\hookrightarrowincluded in a section of :doc:`todo`.
" " "

```
- Use relative imports within TeNPy. Example:
```

from ..linalg import np_conserved as npc

```
- Use the python package pytest for testing. Run it simply with pytest in tests/. You should make sure that all tests run through, before you git push back into the public repo. Long-running tests are marked with the attribute slow; for a quick check you can also run pytest \(-m\) "not slow".

We have set up github actions to automatically run the tests.
- Reversely, if you write new functions, please also include suitable tests!
- During development, you might introduce \# TODO comments. But also try to remove them again later! If you're not \(100 \%\) sure that you will remove it soon, please add a doc-string with a . . todo : : block, such that we can keep track of it.

Unfinished functions should raise NotImplementedError().
- Summarize the changes you have made in the Changelog under [latest].
- If you want to try out new things in temporary files: any folder named playground is ignored by git.
- If you add a new toycode or example: add a reference to include it in the documentation.
- We've created a sphinx extensions for documenting config-option dictionaries. If a class takes a dictionary of options, we usually call it options, convert it to a Config at the very beginning of the _init_ with asConfig(), save it as self.options, and document it in the class doc-string with a . . cfg:config : : directive. The name of the config should usually be the class-name (if that is sufficiently unique), or for algorithms directly the common name of the algorithm, e.g. "DMRG"; use the same name for the use the same name for the documentation of the . cfg:config : : directive as for the Config class instance. Attributes which are simply read-out options should be documented by just referencing the options with the :cfg:option:`configname.optionname` role.

\subsection*{15.2 Bulding the documentation}

You can use Sphinx to generate the full documentation in various formats (including HTML or PDF) yourself, as described in the following.

First, I will assume that you downloaded the [TeNPySource] repository with:
```

git clone --recursive https://github.com/tenpy/tenpy

```

This includes the [TeNPyNotebooks] as a git submodule; you might need to git submodule update if it is out of date.
Building the documentation requires a few more packages (including Sphinx). The recommended way is to create a separate conda environment for it with:
```

conda env create -f doc/environment.yml \# make sure to use the file from the doc/s
\hookrightarrowsubfolder!
conda activate tenpydoc

```

Alternatively, you can use pip and pip install -r doc/requirements.txt, but this will not be able to install all dependencies: some packages like Graphviz are not available from pip alone.

Afterwards, simply go to the folder doc / and run the following command:
```

make html

```

This should generate the html documentation in the folder doc/sphinx_build/html. Open this folder (or to be precise: the file index.html in it) in your webbroser and enjoy this and other documentation beautifully rendered, with cross links, math formulas and even a search function. Other output formats are available as other make targets, e.g., make latexpdf.

Note: Building the documentation with sphinx requires loading the TeNPy modules. The conf.py adjusts the python sys.path to include the /tenpy folder from root directory of the git repository. It will not use the cython-compiled parts.

\subsection*{15.3 To-Do list}

You can check https://github.com/tenpy/tenpy/issues for things to be done.
The following list is auto-generated by sphinx, extracting . . todo : : blocks from doc-strings of the code.

Todo: TDVP is currently not implemented with the sweep class.
(The original entry is located in/home/docs/checkouts/readthedocs.org/user_builds/tenpy/checkouts/v0.8.0/tenpy/algorithms/mps_comn of tenpy.algorithms.mps_common.Sweep, line 6.)

\section*{Todo:}
- implement or wrap netcon.m, a function to find optimal contractionn sequences (arXiv:1304.6112)
- improve helpfulness of Warnings
- _do_trace: trace over all pairs of legs at once. need the corresponding npc function first.
(The original entry is located in /home/docs/checkouts/readthedocs.org/user_builds/tenpy/checkouts/v0.8.0/tenpy/algorithms/network_cc of tenpy.algorithms.network_contractor, line 10.)

Todo: This is still a beta version, use with care. The interface might still change.
(The original entry is located in /home/docs/checkouts/readthedocs.org/user_builds/tenpy/checkouts/v0.8.0/tenpy/algorithms/tdvp.py:do of tenpy.algorithms.tdvp, line 12.)

Todo: long-term: Much of the code is similar as in DMRG. To avoid too much duplicated code, we should have a general way to sweep through an MPS and updated one or two sites, used in both cases.
(The original entry is located in/home/docs/checkouts/readthedocs.org/user_builds/tenpy/checkouts/v0.8.0/tenpy/algorithms/tdvp.py:do of tenpy.algorithms.tdvp, line 16.)

Todo: add further terms (e.g. \(c^{\wedge}\) dagger \(c^{\wedge}\) dagger + h.c. ) to the Hamiltonian.
(The original entry is located in /home/docs/checkouts/readthedocs.org/user_builds/tenpy/checkouts/v0.8.0/tenpy/models/fermions_spin of tenpy.models.fermions_spinless, line 3.)

Todo: WARNING: These models are still under development and not yet tested for correctness. Use at your own risk! Replicate known results to confirm models work correctly. Long term: implement different lattices. Long term: implement variable hopping strengths \(\mathrm{J}_{\mathrm{X}}\), Jy.
(The original entry is located in /home/docs/checkouts/readthedocs.org/user_builds/tenpy/checkouts/v0.8.0/tenpy/models/hofstadter.py:d of tenpy.models.hofstadter, line 3.)

Todo: This is a naive, expensive implementation contracting the full network. Try to follow arXiv:1711.01104 for a better estimate; would that even work in the infinite limit?
(The original entry is located in/home/docs/checkouts/readthedocs.org/user_builds/tenpy/checkouts/v0.8.0/tenpy/networks/mpo.py:docs of tenpy.networks.mpo.MPO.variance, line 5.)

Todo: might be useful to add a "cleanup" function which removes operators cancelling each other and/or unused states. Or better use a 'compress' of the MPO?
(The original entry is located in/home/docs/checkouts/readthedocs.org/user_builds/tenpy/checkouts/v0.8.0/tenpy/networks/mpo.py:docs of tenpy.networks.mpo.MPOGraph, line 17.)

Todo: Make more general: it should be possible to specify states as strings.
(The original entry is located in /home/docs/checkouts/readthedocs.org/user_builds/tenpy/checkouts/v0.8.0/tenpy/networks/mps.py:docs of tenpy.networks.mps.build_initial_state, line 14.)

Todo: test, provide more.
(The original entry is located in /home/docs/checkouts/readthedocs.org/user_builds/tenpy/checkouts/v0.8.0/tenpy/simulations/measurem of tenpy.simulations.measurement, line 7.)

Todo: provide examples, give user guide
(The original entry is located in/home/docs/checkouts/readthedocs.org/user_builds/tenpy/checkouts/v0.8.0/tenpy/simulations/simulation of tenpy.simulations.simulation, line 8.)

Todo: For memory caching with big MPO environments, we need a Hdf5Cacher clearing the memo's every now and then (triggered by what?).
(The original entry is located in /home/docs/checkouts/readthedocs.org/user_builds/tenpy/checkouts/v0.8.0/tenpy/tools/hdf5_io.py:docst of tenpy.tools.hdf5_io, line 65.)

\section*{Part II}

\section*{Reference}

\section*{TENPY MAIN MODULE}
- full name: tenpy
- parent module: tenpy
- type: module

\section*{Submodules}
\begin{tabular}{ll}
\hline algorithms & A collection of algorithms such as TEBD and DMRG. \\
\hline linalg & Linear-algebra tools for tensor networks. \\
\hline models & Definition of the various models. \\
\hline networks & Definitions of tensor networks like MPS and MPO. \\
\hline tools & \begin{tabular}{l} 
A collection of tools: mostly short yet quite useful func- \\
tions.
\end{tabular} \\
\hline version & Access to version of this library. \\
\hline
\end{tabular}

\section*{Module description}

TeNPy - a Python library for Tensor Network Algorithms
TeNPy is a library for algorithms working with tensor networks, e.g., matrix product states and -operators, designed to study the physics of strongly correlated quantum systems. The code is intended to be accessible for newcommers and yet powerful enough for day-to-day research.
\begin{tabular}{ll}
\hline run_simulation([simulation_class_name,...\(])\) & Run the simulation with a simulation class. \\
\hline console_main() & Command line interface. \\
\hline show_config() & Print information about the version of tenpy and used \\
& libraries. \\
\hline
\end{tabular}

\section*{16.1 run_simulation}
- full name: tenpy.run_simulation
- parent module: tenpy
- type: function
tenpy.run_simulation (simulation_class_name='GroundStateSearch', simulation_class_kwargs=None, **simulation_params)
Run the simulation with a simulation class.

\section*{Parameters}
- simulation_class_name (str) - The name of a (sub)class of Simulation to be used for running the simulaiton.
- simulation_class_kwargs (dict / None) - A dictionary of keyword-arguments to be used for the initializing the simulation.
- **simulation_params - Further keyword arguments as documented in the corresponding simulation class, see :cfg:config`Simulation`.

Returns The results from running the simulation, i.e., what tenpy.simulations. Simulation.run() returned.

Return type results

\section*{16.2 console_main}
- full name: tenpy.console_main
- parent module: tenpy
- type: function
tenpy.console_main()
Command line interface.
See also run_simulation() for the python interface running a simulation.
When tenpy is installed correctly via pip/conda, a command line script called tenpy-run is set up, which calls this function, i.e., you can do the following in the terminal:
```

tenpy-run --help

```

Equivalently, you can also invoke the tenpy module from your python interpreter like this:
python -m tenpy --help
```

usage: tenpy-run [-h] [--import-module MODULE] [--sim-class SIM_CLASS] [--option_
๑KEY VALUE]
[--version]
[parameters_file]
Command line interface to run a TeNPy simulation.
positional arguments:
parameters_file

```
(continued from previous page)
```

            A yaml (*.yml) file with the simulation parameters/options.
    optional arguments:
-h, --help
show this help message and exit
--import-module MODULE, -i MODULE
Import the given python MODULE before setting up the simulation. This is,
\hookrightarrowuseful if the
module contains user-defined subclasses. Use python-style names like
\hookrightarrow`numpy` without the
.py ending.
--sim-class SIM_CLASS, -c SIM_CLASS
selects the Simulation (sub)class, e.g. 'GroundStateSearch' (default) or
'RealTimeEvolution'.
--option KEY VALUE, -o KEY VALUE
Allows overwriting some options from the yaml files. KEY can be recursives
separated by
``/``, e.g. ``algorithm_params/trunc_params/chi``. VALUE is initialized_
@by python's
``eval(VALUE)`` with `np`, `scipy` and `tenpy` defined. Thus ``'1.2'`` and
`''2.*np.pi*1.j/6'` or `''np.linspace(0., 1., 6)'` will work if you
\leftrightarrowinclude the quotes
on the command line to ensure that the VALUE is passed as a single
\hookrightarrowargument.
--version, -v
show program's version number and exit
Examples
-_------
In the simplest case, you just give a single yaml file with all the parameters ass
\hookrightarrowargument:
tenpy-run my_params.yml
If you implemented a custom simulation class called ``MyGreatSimulation`` in a
file ``my_simulations.py``, you can use it like this:
tenpy-run -i my_simulations -c MyGreatSimulation my_params.yml
Further, you can overwrite one or multiple options of the parameters file:
tenpy-run my_params.yml -o output_filename '"output.h5"' -o model_params/Jz 2.
Note that string values for the options require double quotes on the command line.

```

\section*{16.3 show_config}
- full name: tenpy.show_config
- parent module: tenpy
- type: function
tenpy.show_config()
Print information about the version of tenpy and used libraries.
The information printed is tenpy.version.version_summary.
```

tenpy.__version__ = '0.8.0'
hard-coded version string
tenpy.__full_version__ = '0.8.0'
full version from git description, and numpy/scipy/python versions

```

\section*{Submodules}
\begin{tabular}{ll}
\hline algorithms & A collection of algorithms such as TEBD and DMRG. \\
\hline linalg & Linear-algebra tools for tensor networks. \\
\hline models & Definition of the various models. \\
\hline networks & Definitions of tensor networks like MPS and MPO. \\
\hline simulations & Simulation setup. \\
\hline tools & \begin{tabular}{l} 
A collection of tools: mostly short yet quite useful func- \\
tions.
\end{tabular} \\
\hline version & Access to version of this library. \\
\hline
\end{tabular}

ALGORITHMS
- full name: tenpy.algorithms
- parent module: tenpy
- type: module

\section*{Module description}

A collection of algorithms such as TEBD and DMRG.

\section*{Submodules}
\begin{tabular}{ll}
\hline algorithm & This module contains some base classes for algorithms. \\
\hline truncation & Truncation of Schmidt values. \\
\hline tebd & Time evolving block decimation (TEBD). \\
\hline mps_common & 'Sweep' algorithm and effective Hamiltonians. \\
\hline dmrg & Density Matrix Renormalization Group (DMRG). \\
\hline tdvp & Time Dependant Variational Principle (TDVP) with \\
& MPS (finite version only). \\
\hline purification & Algorithms for using Purification. \\
\hline mpo_evolution & Time evolution using the WI or WII approximation of \\
& the time evolution operator. \\
\hline network_contractor & Network Contractor. \\
\hline exact_diag & Full diagonalization (ED) of the Hamiltonian. \\
\hline
\end{tabular}

\section*{17.1 algorithm}
- full name: tenpy.algorithms.algorithm
- parent module: tenpy.algorithms
- type: module

\section*{Classes}

\begin{tabular}{ll}
\hline Algorithm(psi, model, options, \({ }^{*}[\), resume_data \(\left.]\right)\) & \begin{tabular}{l} 
Base class and common interface for a tensor-network \\
based algorithm in TeNPy.
\end{tabular} \\
\hline \begin{tabular}{l} 
TimeEvolutionAlgorithm(psi, model, options,
\end{tabular} & Common interface for (real) time evolution algorithms. \\
\(\left.{ }^{*}\right)\)
\end{tabular}

\section*{Module description}

This module contains some base classes for algorithms.

\section*{17.2 truncation}
- full name: tenpy.algorithms.truncation
- parent module: tenpy.algorithms
- type: module

\section*{Classes}


\subsection*{17.2.1 TruncationError}
- full name: tenpy.algorithms.truncation.TruncationError
- parent module: tenpy.algorithms.truncation
- type: class

\section*{Inheritance Diagram}


\section*{Methods}
\begin{tabular}{ll}
\hline TruncationError.__init__([eps, ov]) & Initialize self. \\
\hline TruncationError.copy() & Return a copy of self. \\
\hline \begin{tabular}{l} 
TruncationError.from_S(S_discarded[, \\
norm_old])
\end{tabular} & \begin{tabular}{l} 
Construct TruncationError from discarded singular val- \\
ues.
\end{tabular} \\
\hline \begin{tabular}{l} 
TruncationError. from_hdf5(hdf5_loader, \\
h5gr,...)
\end{tabular} & Load instance from a HDF5 file. \\
\hline \begin{tabular}{l} 
TruncationError.from_norm(norm_new[, \\
norm_old])
\end{tabular} & \begin{tabular}{l} 
Construct TruncationError from norm after and before \\
the truncation.
\end{tabular} \\
\hline \begin{tabular}{ll} 
TruncationError.save_hdf5(hdf5_saver, h5gr, \\
\(\ldots)\)
\end{tabular} & Export self into a HDF5 file. \\
\hline
\end{tabular}

\section*{Class Attributes and Properties}
TruncationError.ov_err_ Error 1.-ov of the overlap with the correct state.
class tenpy.algorithms.truncation. TruncationError (eps=0.0,ov=1.0)
Bases: tenpy.tools.hdf5_io.Hdf5Exportable
Class representing a truncation error.
The default initialization represents "no truncation".

Warning: For imaginary time evolution, this is not the error you are interested in!

\section*{Parameters}
- eps (float) - See below.
- ov (float) - See below.
eps
The total sum of all discared Schmidt values squared. Note that if you keep singular values up to 1.e-14 (= a bit more than machine precision for 64bit floats), eps is on the order of 1.e-28 (due to the square)!

Type float
ov
A lower bound for the overlap \(\left|\left\langle\psi_{\text {trunc }} \mid \psi_{\text {correct }}\right\rangle\right|^{2}\) (assuming normalization of both states). This is probably the quantity you are actually interested in. Takes into account the factor 2 explained in the section on Errors in the TEBD Wikipedia article <https://en.wikipedia.org/wiki/Time-evolving_block_decimation>.

Type float
copy ()
Return a copy of self.
classmethod from_norm (norm_new, norm_old \(=1.0\) )
Construct TruncationError from norm after and before the truncation.

\section*{Parameters}
- norm_new (float) - Norm of Schmidt values kept, \(\sqrt{\sum_{a k e p t} \lambda_{a}^{2}}\) (before renormalization).
- norm_old (float) - Norm of all Schmidt values before truncation, \(\sqrt{\sum_{a} \lambda_{a}^{2}}\).
classmethod from_S (S_discarded, norm_old=None)
Construct TruncationError from discarded singular values.

\section*{Parameters}
- S_discarded (1D numpy array) - The singular values discarded.
- norm_old (float) - Norm of all Schmidt values before truncation, \(\sqrt{\sum_{a} \lambda_{a}^{2}}\). Default (None) is 1 .
property ov_err
Error 1.-ov of the overlap with the correct state.
classmethod from_hdf5 (hdf5_loader, h5gr, subpath)
Load instance from a HDF5 file.
This method reconstructs a class instance from the data saved with save_hdf5 ().

\section*{Parameters}
- hdf5_loader (Hdf5Loader) - Instance of the loading engine.
- h5gr (Group) - HDF5 group which is represent the object to be constructed.
- subpath (str) - The name of \(h 5 g r\) with a '/' in the end.

Returns obj - Newly generated class instance containing the required data.
Return type cls
```

save_hdf5 (hdf5_saver, h5gr, subpath)

```

Export self into a HDF5 file.
This method saves all the data it needs to reconstruct self with from_hdf5 ().
This implementation saves the content of __ dict__ with save_dict_content (), storing the format under the attribute 'format'.

\section*{Parameters}
- hdf5_saver (Hdf5Saver) - Instance of the saving engine.
- h5gr (: class`Group ) - HDF5 group which is supposed to represent self.
- subpath (str) - The name of \(h 5 g r\) with a '/' in the end.

\section*{Functions}
\begin{tabular}{ll}
\hline Svd_theta(theta, trunc_par[, qtotal_LR, ...]) & \begin{tabular}{l} 
Performs SVD of a matrix theta \((=\) the wavefunction) \\
and truncates it.
\end{tabular} \\
\hline truncate(S, options) & \begin{tabular}{l} 
Given a Schmidt spectrum \(S\), determine which values to \\
keep.
\end{tabular} \\
\hline
\end{tabular}

\subsection*{17.2.2 svd_theta}
- full name: tenpy.algorithms.truncation.svd_theta
- parent module: tenpy.algorithms.truncation
- type: function
tenpy.algorithms.truncation.svd_theta(theta, trunc_par, qtotal_LR=[None, None], inner_labels \(=\left[\right.\) ' \(v R^{\prime}\), 'vL'])
Performs SVD of a matrix theta (= the wavefunction) and truncates it.
Perform a singular value decomposition (SVD) with svd() and truncates with truncate(). The result is an approximation theta \(\sim=\) tensordot(U.scale_axis(S*renormalization, 1), VH, axes=1)

\section*{Parameters}
- theta (Array, shape ( \(\mathrm{M}, \mathrm{N}\) ) ) - The matrix, on which the singular value decomposition (SVD) is performed. Usually, theta represents the wavefunction, such that the SVD is a Schmidt decomposition.
- trunc_par (dict) - truncation parameters as described in truncate ().
- qtotallR ( (charges, charges)) - The total charges for the returned \(U\) and \(V H\).
- inner_labels ((string, string)) - Labels for the \(U\) and \(V H\) on the newlycreated bond.

\section*{Returns}
- U (Array) - Matrix with left singular vectors as columns. Shape (M, M) or (M, K) depending on full_matrices.
- \(\mathbf{S}\) (lD ndarray) - The singluar values of the array. If no cutoff is given, it has lenght min (M, \(\mathrm{N})\). Normalized to np. linalg. norm (S) ==1.
- VH (Array) - Matrix with right singular vectors as rows. Shape (N, N) or (K, N) depending on full_matrices.
- \(\mathbf{e r r}\) (TruncationError) - The truncation error introduced.
- renormalization (float) - Factor, by which S was renormalized.

\section*{Module description}

Truncation of Schmidt values.
Often, it is necessary to truncate the number of states on a virtual bond of an MPS, keeping only the state with the largest Schmidt values. The function truncate () picks exactly those from a given Schmidt spectrum \(\lambda_{a}\), depending on some parameters explained in the doc-string of the function.

Further, we provide TruncationError for a simple way to keep track of the total truncation error.
The SVD on a virtual bond of an MPS actually gives a Schmidt decomposition \(|\psi\rangle=\sum_{a} \lambda_{a}\left|L_{a}\right\rangle\left|R_{a}\right\rangle\) where \(\left|L_{a}\right\rangle\) and \(\left|R_{a}\right\rangle\) form orthonormal bases of the parts left and right of the virtual bond. Let us assume that the state is properly normalized, \(\langle\psi \mid \psi\rangle=\sum_{a} \lambda_{a}^{2}=1\). Assume that the singular values are ordered descending, and that we keep the first \(\chi_{c}\) of the initially \(\chi\) Schmidt values.
Then we decompose the untruncated state as \(|\psi\rangle=\sqrt{1-\epsilon}\left|\psi_{t r}\right\rangle+\sqrt{\epsilon}\left|\psi_{t r}^{\perp}\right\rangle\) where \(\left|\psi_{t r}\right\rangle=\frac{1}{\sqrt{1-\epsilon}} \sum_{a<\chi_{c}} \lambda_{a}\left|L_{a}\right\rangle\left|R_{a}\right\rangle\) is the truncated state kept (normalized to 1), \(\left|\psi_{t r}\right\rangle=\frac{1}{\sqrt{\epsilon}} \sum_{a>=\chi_{c}} \lambda_{a}\left|L_{a}\right\rangle\left|R_{a}\right\rangle\) is the discarded part (orthogonal to the kept part) and the truncation error of a single truncation is defined as \(\epsilon=1-\left|\left\langle\psi \mid \psi_{t r}\right\rangle\right|^{2}=\sum_{a>=\chi_{c}} \lambda_{a}^{2}\).

Warning: For imaginary time evolution (e.g. with TEBD), you try to project out the ground state. Then, looking at the truncation error defined in this module does not give you any information how good the found state coincides with the actual ground state! (Instead, the returned truncation error depends on the overlap with the initial state, which is arbitrary \(>0\) )

Warning: This module takes only track of the errors coming from the truncation of Schmidt values. There might be other sources of error as well, for example TEBD has also an discretisation error depending on the chosen time step.

\section*{17.3 tebd}
- full name: tenpy.algorithms.tebd
- parent module: tenpy.algorithms
- type: module

\section*{Classes}

\begin{tabular}{ll}
\hline Engine(psi, model, options) & Deprecated old name of TEBDEngine. \\
\hline RandomUnitaryEvolution(psi, options) & \begin{tabular}{l} 
Evolution of an MPS with random two-site unitaries in \\
a TEBD-like fashion.
\end{tabular} \\
\hline TEBDEngine(psi, model, options, **kwargs) & Time Evolving Block Decimation (TEBD) algorithm. \\
\hline
\end{tabular}

\subsection*{17.3.1 Engine}
- full name: tenpy.algorithms.tebd.Engine
- parent module: tenpy.algorithms.tebd
- type: class

\section*{Inheritance Diagram}


\section*{Methods}
\begin{tabular}{ll}
\hline Engine.__init__(psi, model, options) & Initialize self. \\
\hline Engine.calc_U(order, delta_t \([\), type_evo,...]) & \begin{tabular}{l} 
Calculate self.U_bond from self. \\
bond_eig_\{vals, vecs \(\}.\)
\end{tabular} \\
\hline Engine.get_resume_data() & \begin{tabular}{l} 
Return necessary data to resume a run () interrupted at \\
a checkpoint.
\end{tabular} \\
\hline Engine.resume_run() & Resume a run that was interrupted. \\
\hline Engine.run() & Run TEBD real time evolution by N_steps**dt. \\
\hline Engine.run_GS() & \begin{tabular}{l} 
TEBD algorithm in imaginary time to find the ground \\
state.
\end{tabular} \\
\hline Engine.suzuki_trotter_decomposition(orderReturns list of necessary steps for the suzuki trotter de- \\
\(\ldots\) composition.
\end{tabular}

\section*{Class Attributes and Properties}

Engine.TEBD_params
\begin{tabular}{ll}
\hline Engine.trunc_err_bonds & truncation error introduced on each non-trivial bond. \\
\hline Engine.verbose &
\end{tabular}
class tenpy.algorithms.tebd.Engine (psi, model, options)
Bases: tenpy.algorithms.tebd.TEBDEngine
Deprecated old name of TEBDEngine.
calc_U (order, delta_t, type_evo='real', E_offset=None)
Calculate self.U_bond from self.bond_eig_\{vals, vecs\}.
This function calculates
- U_bond = exp(-i dt (H_bond-E_offset_bond)) for type_evo='real', or
- U_bond \(=\exp (-d t\) H_bond) for type_evo='imag'.

For first order (in delta_t), we need just one \(d t=\) delta_t. Higher order requires smaller \(d t\) steps, as given by suzuki_trotter_time_steps().

\section*{Parameters}
- order (int) - Trotter order calculated U_bond. See update for more information.
- delta_t (float) - Size of the time-step used in calculating U_bond
- type_evo ('imag' | 'real')-Determines whether we perform real or imaginary time-evolution.
- E_offset (None | list of float) - Possible offset added to H_bond for realtime evolution.
get_resume_data()
Return necessary data to resume a run () interrupted at a checkpoint.
At a checkpoint, you can save psi, model and options along with the data returned by this function. When the simulation aborts, you can resume it using this saved data with:
```

eng = AlgorithmClass(psi, model, options, resume_data=resume_data)
eng.resume_run(resume_data)

```

An algorithm which doesn't support this should override resume_run to raise an Error.
Returns resume_data - Dictionary with necessary data (apart from copies of psi, model, options) that allows to continue the simulation from where we are now.
Return type dict
resume_run()
Resume a run that was interrupted.
In case we saved an intermediate result at a checkpoint, this function allows to resume the run () of the algorithm (after re-initialization with the resume_data). Since most algorithms just have a while loop with break conditions, the default behaviour implemented here is to just call run ().
run()
Run TEBD real time evolution by \(N\) _steps \({ }^{`}\) ` \(d t\).
run_GS ()
TEBD algorithm in imaginary time to find the ground state.

Note: It is almost always more efficient (and hence advisable) to use DMRG. This algorithms can nonetheless be used quite well as a benchmark and for comparison.
option TEBDEngine.delta_tau_list: list
A list of floats: the timesteps to be used. Choosing a large timestep delta_tau introduces large (Trotter) errors, but a too small time step requires a lot of steps to reach exp (-tau H) --> \(\mid p s i 0><\) psi0|. Therefore, we start with fairly large time steps for a quick time evolution until convergence, and the gradually decrease the time step.
option TEBDEngine.order: int
Order of the Suzuki-Trotter decomposition.
option TEBDEngine.N_steps: int
Number of steps before measurement can be performed
static suzuki_trotter_decomposition (order, \(N_{-}\)steps)
Returns list of necessary steps for the suzuki trotter decomposition.
We split the Hamiltonian as \(H=H_{\text {even }}+H_{o d d}=H[0]+H[1]\). The Suzuki-Trotter decomposition is an approximation \(\exp (t H) \approx \operatorname{prod}_{(j, k) \in S T} \exp (d[j] t H[k])+O\left(t^{\text {order }+1}\right)\).

Parameters order (1, 2, 4, '4_opt') - The desired order of the Suzuki-Trotter decomposition. Order 1 approximation is simply \(e^{A} a^{B}\). Order 2 is the "leapfrog" \(e^{\wedge}\{A / 2\} e^{\wedge} B e^{\wedge}\{A / 2\}\). Order 4 is the fourth-order from [[suzuki1991]] (also referenced in [[schollwoeck2011]]), and '4_opt' gives the optmized version of Equ. (30a) in [[barthel2020]].
Returns ST_decomposition - Indices j, \(k\) of the time-steps \(d=\) suzuki_trotter_time_step (order) and the decomposition of \(H\). They are chosen such that a subsequent application of \(\exp (d[j] t H[k])\) to a given state |psi> yields (exp (N_steps \(\left.t H[k])+O\left(N \_s t e p s t^{\wedge}\{o r d e r+1\}\right)\right) \mid p s i>\).

Return type list of (int, int)
static suzuki_trotter_time_steps (order)
Return time steps of U for the Suzuki Trotter decomposition of desired order.
See suzuki_trotter_decomposition () for details.
Parameters order (int) - The desired order of the Suzuki-Trotter decomposition.
Returns time_steps - We need \(U=\exp \left(-i \quad H \_\{e v e n / o d d\}\right.\) delta_t * dt) for the \(d t\) returned in this list.

Return type list of float
```

property trunc_err_bonds

```
truncation error introduced on each non-trivial bond.
update (N_steps)
Evolve by N_steps * U_param['dt'].
Parameters N_steps (int) - The number of steps for which the whole lattice should be updated.
Returns trunc_err - The error of the represented state which is introduced due to the truncation during this sequence of update steps.

\section*{Return type TruncationError}
update_bond (i, U_bond)
Updates the B matrices on a given bond.
Function that updates the B matrices, the bond matrix s between and the bond dimension chi for bond i . The correponding tensor networks look like this:


\section*{Parameters}
- i (int) - Bond index; we update the matrices at sites i-1, i.
- U_bond (Array) - The bond operator which we apply to the wave function. We expect labels 'p0', 'p1', 'p0*', 'p1*'.
Returns trunc_err - The error of the represented state which is introduced by the truncation during this update step.

\section*{Return type TruncationError}
update_bond_imag ( \(i, U \_\)bond \()\)
Update a bond with a (possibly non-unitary) \(U \_\)bond.
Similar as update_bond (); but after the SVD just keep the \(A, S, B\) canonical form. In that way, one can sweep left or right without using old singular values, thus preserving the canonical form during imaginary time evolution.

\section*{Parameters}
- i (int) - Bond index; we update the matrices at sites i-1, i.
- U_bond (Array) - The bond operator which we apply to the wave function. We expect labels 'p0', 'p1', 'p0*', 'p1*'.

Returns trunc_err - The error of the represented state which is introduced by the truncation during this update step.
Return type TruncationError
update_imag ( \(N\) _steps)
Perform an update suitable for imaginary time evolution.
Instead of the even/odd brick structure used for ordinary TEBD, we 'sweep' from left to right and right to left, similar as DMRG. Thanks to that, we are actually able to preserve the canonical form.

Parameters N_steps (int) - The number of steps for which the whole lattice should be updated.

Returns trunc_err - The error of the represented state which is introduced due to the truncation during this sequence of update steps.
Return type TruncationError
update_step ( \(U \_i d x \_d t\),odd)
Updates either even or odd bonds in unit cell.
Depending on the choice of \(p\), this function updates all even ( E , odd=False, 0 ) or odd ( \(O\) ) ( \(\mathrm{odd}=\) True, 1 ) bonds:


Note that finite boundary conditions are taken care of by having Us [0] = None.

\section*{Parameters}
- U_idx_dt (int) - Time step index in self._U, evolve with Us[i] = self. U[U_idx_dt][i] at bond (i-1,i).
- odd (bool/int) - Indication of whether to update even (odd=False, 0) or even (odd=True, 1) sites

Returns trunc_err - The error of the represented state which is introduced due to the truncation during this sequence of update steps.

\section*{Return type TruncationError}

\section*{Module description}

Time evolving block decimation (TEBD).
The TEBD algorithm (proposed in [[vidal2004]]) uses a trotter decomposition of the Hamiltonian to perform a time evoltion of an MPS. It works only for nearest-neighbor hamiltonians (in tenpy given by a NearestNeighborModel), which can be written as \(H=H^{\text {even }}+H^{\text {odd }}\), such that \(H^{\text {even }}\) contains the the terms on even bonds (and similar \(H^{\text {odd }}\) the terms on odd bonds). In the simplest case, we apply first \(U=\exp \left(-i * d t * H^{\text {even }}\right)\), then \(U=\exp \left(-i * d t * H^{\text {odd }}\right)\) for each time step \(d t\). This is correct up to errors of \(O\left(d t^{2}\right)\), but to evolve until a time \(T\), we need \(T / d t\) steps, so in total it is only correct up to error of \(O(T * d t)\). Similarly, there are higher order schemata (in dt) (for more details see TEBDEngine.update () ).

Remember, that bond \(i\) is between sites ( \(i-1, i\) ), so for a finite MPS it looks like:


After each application of a \(U i\), the MPS needs to be truncated - otherwise the bond dimension chi would grow indefinitely. A bound for the error introduced by the truncation is returned.

If one chooses imaginary \(d t\), the exponential projects (for sufficiently long 'time' evolution) onto the ground state of the Hamiltonian.

Note: The application of DMRG is typically much more efficient than imaginary TEBD! Yet, imaginary TEBD might
be usefull for cross-checks and testing.

\section*{17.4 mps_common}
- full name: tenpy.algorithms.mps_common
- parent module: tenpy.algorithms
- type: module

\section*{Classes}

\begin{tabular}{ll}
\hline EffectiveH(env, i0[, combine, move_right]) & \begin{tabular}{l} 
Prototype class for local effective Hamiltonians used in \\
sweep algorithms.
\end{tabular} \\
\hline OneSiteH(env, i0[, combine, move_right]) & \begin{tabular}{l} 
Class defining the one-site effective Hamiltonian for \\
Lanczos.
\end{tabular} \\
\hline Sweep(psi, model, options, *[, resume_data]) & Prototype class for a 'sweeping' algorithm. \\
\hline TwoSiteH(env, i0[, combine, move_right]) & \begin{tabular}{l} 
Class defining the two-site effective Hamiltonian for \\
Lanczos.
\end{tabular} \\
\hline \begin{tabular}{ll} 
VariationalApplyMPO(psi, U_MPO, options[,, & \begin{tabular}{l} 
Variational compression for applying an MPO to an \\
MPS (in place).
\end{tabular} \\
\hline V.]) & Variational compression of an MPS (in place). \\
\hline
\end{tabular}
\end{tabular}

\subsection*{17.4.1 EffectiveH}
- full name: tenpy.algorithms.mps_common.EffectiveH
- parent module: tenpy.algorithms.mps_common
- type: class

\section*{Inheritance Diagram}


\section*{Methods}
\begin{tabular}{ll}
\hline EffectiveH.__init__(env, i0[, combine, ...]) & Initialize self. \\
\hline EffectiveH.adjoint() & Return the hermitian conjugate of self \\
\hline EffectiveH.combine_theta(theta) & \begin{tabular}{l} 
Combine the legs of theta, such that it fits to how we \\
combined the legs of self.
\end{tabular} \\
\hline EffectiveH.matvec(vec) & Calculate the action of the operator on a vector vec. \\
\hline EffectiveH.to_matrix() & Contract self to a matrix. \\
\hline
\end{tabular}

\section*{Class Attributes and Properties}
```

EffectiveH.acts_on

```
EffectiveH.length
class tenpy.algorithms.mps_common.EffectiveH (env, iO, combine=False, move_right=True) Bases: tenpy.linalg.sparse.NpcLinearOperator

Prototype class for local effective Hamiltonians used in sweep algorithms.
As an example, the local effective Hamiltonian for a two-site (DMRG) algorithm looks like:
\(\square\)
(continues on next page)
where HO and H 1 are MPO tensors.

\section*{Parameters}
- env (MPOEnvironment) - Environment for contraction <psi|H|psi>.
- i0 (int) - Index of the active site if length=1, or of the left-most active site if length \(>1\).
- combine (bool, optional) - Whether to combine legs into pipes as far as possible. This reduces the overhead of calculating charge combinations in the contractions.
- move_right (bool, optional) - Whether the sweeping algorithm that calls for an EffectiveH is moving to the right.

\section*{length}

Number of (MPS) sites the effective hamiltonian covers. NB: Class attribute.
Type int
dtype
The data type of the involved arrays.
Type np.dtype
N
Contracting self with as_matrix() will result in an \(N^{`} x^{`} N\) matrix .
Type int

\section*{acts_on}

Labels of the state on which self acts. NB: class attribute. Overwritten by normal attribute, if combine.
Type list of str

\section*{combine}

Whether to combine legs into pipes as far as possible. This reduces the overhead of calculating charge combinations in the contractions.

Type bool
move_right
Whether the sweeping algorithm that calls for an EffectiveH is moving to the right.
Type bool
combine_theta (theta)
Combine the legs of theta, such that it fits to how we combined the legs of self.
Parameters theta (Array) - Wave function to apply the effective Hamiltonian to, with uncombined legs.
Returns theta - Wave function with labels as given by self.acts_on.

\section*{Return type Array}
```

adjoint()

```

Return the hermitian conjugate of self
If self is hermitian, subclasses can choose to implement this to define the adjoint operator of self.
matvec (vec)
Calculate the action of the operator on a vector vec.

Note that we don't require vec to be one-dimensional. However, for square operators we require that the result of matvec has the same legs (in the same order) as vec such that they can be added. Note that this excludes a non-trivial qtotal for square operators.
```

to_matrix()

```

Contract self to a matrix.
If self represents an operator with very small shape, e.g. because the MPS bond dimension is very small, an algorithm might choose to contract self to a single tensor.

Returns matrix - Contraction of the represented operator.
Return type Array

\subsection*{17.4.2 OneSiteH}
- full name: tenpy.algorithms.mps_common.OneSiteH
- parent module: tenpy.algorithms.mps_common
- type: class

\section*{Inheritance Diagram}


\section*{Methods}
\begin{tabular}{ll}
\hline \begin{tabular}{l} 
OneSiteH.__init__(env, i0[, combine, \\
move_right])
\end{tabular} & Initialize self. \\
\hline OneSiteH.adjoint() & Return the hermitian conjugate of self. \\
\hline OneSiteH.combine_Heff() & \begin{tabular}{l} 
Combine LP and RP with W to form LHeff and RHeff, \\
depending on the direction.
\end{tabular} \\
\hline OneSiteH.combine_theta(theta) & \begin{tabular}{l} 
Combine the legs of theta, such that it fits to how we \\
combined the legs of self.
\end{tabular} \\
\hline OneSiteH.matvec(theta) & Apply the effective Hamiltonian to theta. \\
\hline &
\end{tabular}

Table 13 - continued from previous page
OneSiteH.to_matrix() Contract self to a matrix.

\section*{Class Attributes and Properties}
```

OneSiteH.acts_on

```
OneSiteH.length
class tenpy.algorithms.mps_common. OneSiteH (env, iO, combine=False, move_right=True)
    Bases: tenpy.algorithms.mps_common. EffectiveH

Class defining the one-site effective Hamiltonian for Lanczos.
The effective one-site Hamiltonian looks like this:


If combine is True, we define either LHeff as contraction of \(L P\) with \(W\) (in the case move_right is True) or RHeff as contraction of \(R P\) and \(W\).

\section*{Parameters}
- env (MPOEnvironment) - Environment for contraction <psi|H|psi>.
- i0 (int) - Index of the active site if length \(=1\), or of the left-most active site if length \(>1\).
- combine (bool) - Whether to combine legs into pipes. This combines the virtual and physical leg for the left site (when moving right) or right side (when moving left) into pipes. This reduces the overhead of calculating charge combinations in the contractions, but one matvec () is formally more expensive, \(O\left(2 d^{3} \chi^{3} D\right)\). Is originally from the wosite method; unclear if it works well for 1 site.
- move_right (bool) - Whether the the sweep is moving right or left for the next update.

\section*{length}

Number of (MPS) sites the effective hamiltonian covers.
Type int
acts_on
Labels of the state on which self acts. NB: class attribute. Overwritten by normal attribute, if combine.
Type list of str
combine, move_right
See above.
Type bool
LHeff, RHeff
Only set if combine, and only one of them depending on move_right. If move_right was True, LHeff is set with labels ' (vR*.p0) ', 'wR', ' (vR.p0*) ' for bra, MPO, ket; otherwise RHeff is set with labels '(p0*.vL)', 'wL', '(p0, vL*)'

Type Array
LP, W0, RP
Tensors making up the network of self.
Type Array
matvec (theta)
Apply the effective Hamiltonian to theta.
Parameters theta (Array) - Labels: vL, p0, vR if combine=False, (vL.p0), vR or vL, ( \(p 0 . \mathrm{vR}\) ) if True (depending on the direction of movement)
Returns Product of theta and the effective Hamiltonian.
Return type theta Array
combine_Heff()
Combine LP and RP with W to form LHeff and RHeff, depending on the direction.
In a move to the right, we need LHeff. In a move to the left, we need RHeff. Both contain the same W.
combine_theta (theta)
Combine the legs of theta, such that it fits to how we combined the legs of self.
Parameters theta (Array) - Wave function with labels 'vL', 'p0', 'p1', 'vR'
Returns theta - Wave function with labels 'vL', 'p0', 'p1', 'vR'
Return type Array
to_matrix()
Contract self to a matrix.
adjoint()
Return the hermitian conjugate of self.

\subsection*{17.4.3 TwoSiteH}
- full name: tenpy.algorithms.mps_common.TwoSiteH
- parent module: tenpy.algorithms.mps_common
- type: class

\section*{Inheritance Diagram}


\section*{Methods}
\begin{tabular}{|c|c|}
\hline TwoSiteH.__init__(env, i0[,
move_right \(]\) ) & combine, Initialize self. \\
\hline TwoSiteH.adjoint() & Return the hermitian conjugate of self. \\
\hline TwoSiteH.combine_Heff() & Combine LP and RP with W to form LHeff and RHeff. \\
\hline IwoSiteH.combine_theta(theta) & Combine the legs of theta, such that it fits to how we combined the legs of self. \\
\hline TwoSiteH.matvec(theta) & Apply the effective Hamiltonian to theta. \\
\hline IwoSiteH.to_matrix() & Contract self to a matrix. \\
\hline
\end{tabular}

\section*{Class Attributes and Properties}

\section*{TwoSiteH.acts_on}

TwoSiteH.length
class tenpy.algorithms.mps_common.TwoSiteH (env, iO, combine=False, move_right=True)
Bases: tenpy.algorithms.mps_common.EffectiveH
Class defining the two-site effective Hamiltonian for Lanczos.
The effective two-site Hamiltonian looks like this:
\(\square\)

If combine is True, we define LHeff and RHeff, which are the contractions of \(L P\) with \(W 0\), and \(R P\) with \(W 1\), respectively.

\section*{Parameters}
- env (MPOEnvironment) - Environment for contraction <psi|H|psi>.
- i0 (int) - Index of the active site if length=1, or of the left-most active site if length>1.
- combine (bool) - Whether to combine legs into pipes. This combines the virtual and physical leg for the left site (when moving right) or right side (when moving left) into pipes. This reduces the overhead of calculating charge combinations in the contractions, but one matvec () is formally more expensive, \(O\left(2 d^{3} \chi^{3} D\right)\).
- move_right (bool) - Whether the the sweep is moving right or left for the next update.

\section*{combine}

Whether to combine legs into pipes. This combines the virtual and physical leg for the left site and right site into pipes. This reduces the overhead of calculating charge combinations in the contractions, but one matvec () is formally more expensive, \(O\left(2 d^{3} \chi^{3} D\right)\).

Type bool

\section*{length}

Number of (MPS) sites the effective hamiltonian covers.
Type int

\section*{acts_on}

Labels of the state on which self acts. NB: class attribute. Overwritten by normal attribute, if combine.
Type list of str

\section*{LHeff}

Left part of the effective Hamiltonian. Labels ' (vR*.p0)', 'wR', '(vR.p0*)' for bra, MPO, ket.

Type Array
RHeff
Right part of the effective Hamiltonian. Labels '(p1*.vL)', 'wL', '(p1.vL*)' for ket, MPO, bra.

Type Array
LP, W0, W1, RP
Tensors making up the network of self.
Type Array
matvec (theta)
Apply the effective Hamiltonian to theta.
Parameters theta (Array) - Labels: vL, p0, p1, vR if combine=False, (vL.p0), ( \(\mathrm{p} 1 . \mathrm{vR}\) ) if True

Returns Product of theta and the effective Hamiltonian.
Return type theta Array
combine_Heff()
Combine LP and RP with W to form LHeff and RHeff.
Combine LP with W0 and RP with W1 to get the effective parts of the Hamiltonian with piped legs.

\section*{combine_theta (theta)}

Combine the legs of theta, such that it fits to how we combined the legs of self.
Parameters theta (Array) - Wave function with labels 'vL', 'p0', 'p1', 'vR'
Returns theta-Wave function with labels 'vL', 'po', 'p1', 'vR'
Return type Array
to_matrix()
Contract self to a matrix.
adjoint()
Return the hermitian conjugate of self.

\section*{Module description}
'Sweep' algorithm and effective Hamiltonians.
Many MPS-based algorithms use a 'sweep' structure, wherein local updates are performed on the MPS tensors sequentially, first from left to right, then from right to left. This procedure is common to DMRG, TDVP, sequential time evolution, etc.
Another common feature of these algorithms is the use of an effective local Hamiltonian to perform the local updates. The most prominent example of this is probably DMRG, where the local MPS object is optimized with respect to the rest of the MPS-MPO-MPS network, the latter forming the effective Hamiltonian.

The Sweep class attempts to generalize as many aspects of 'sweeping' algorithms as possible. EffectiveH and its subclasses implement the effective Hamiltonians mentioned above. Currently, effective Hamiltonians for 1-site and 2 -site optimization are implemented.

The VariationalCompression and VariationalApplyMPO implemented here also directly use the Sweep class.

\section*{17.5 dmrg}
- full name: tenpy.algorithms.dmrg
- parent module: tenpy.algorithms
- type: module

\section*{Classes}

\begin{tabular}{ll}
\hline DMRGEngine(psi, model, options, **kwargs) & \begin{tabular}{l} 
DMRG base class with common methods for the \\
TwoSiteDMRG and SingleSiteDMRG.
\end{tabular} \\
\hline DensityMatrixMixer(options) & Mixer based on density matrices. \\
\hline EngineCombine(psi, model, DMRG_params) & \begin{tabular}{l} 
Engine which combines legs into pipes as far as possi- \\
ble.
\end{tabular} \\
\hline EngineFracture(psi, model, DMRG_params) & Engine which keeps the legs separate. \\
\hline Mixer(options) & Base class of a general Mixer. \\
\hline SingleSiteDMRGEngine(psi, model, options,...) & Engine for the single-site DMRG algorithm. \\
\hline SingleSiteMixer(options) & Mixer for single-site DMRG. \\
\hline \begin{tabular}{ll} 
TwoSiteDMRGEngine(psi, \\
\(* *\) kwargs)
\end{tabular} & model, \\
\hline TwoSiteMixer(options) & Engine for the two-site DMRG algorithm. \\
\hline
\end{tabular}

\subsection*{17.5.1 DensityMatrixMixer}
- full name: tenpy.algorithms.dmrg.DensityMatrixMixer
- parent module: tenpy.algorithms.dmrg
- type: class

\section*{Inheritance Diagram}


\section*{Methods}
\begin{tabular}{|c|c|}
\hline DensityMatrixMixer.__init__(options) & Initialize self. \\
\hline DensityMatrixMixer.get_xL(wL_leg, Id_L, Id_R) & Generate the coupling of the MPO legs for the reduced density matrix. \\
\hline DensityMatrixMixer.get_xR(wR_leg, Id_L, Id_R) & Generate the coupling of the MPO legs for the reduced density matrix. \\
\hline DensityMatrixMixer.mix_rho_I(engine, theta,...) & Calculated mixed reduced density matrix for left site. \\
\hline DensityMatrixMixer.mix_rho_R(engine, theta, ...) & Calculated mixed reduced density matrix for left site. \\
\hline DensityMatrixMixer.perturb_svd(engine, ...) & Mix extra terms to theta and perform an SVD. \\
\hline DensityMatrixMixer. update_amplitude(sweeps) & Update the amplitude, possibly disable the mixer. \\
\hline
\end{tabular}
class tenpy.algorithms.dmrg.DensityMatrixMixer (options)
Bases: tenpy.algorithms.dmrg.Mixer
Mixer based on density matrices.
This mixer constructs density matrices as described in the original paper [[white2005]].
perturb_svd (engine, theta, i0, update_LP, update_RP)
Mix extra terms to theta and perform an SVD.
We calculate the left and right reduced density using the mixer (which might include applications of \(H\) ). These density matrices are diagonalized and truncated such that we effectively perform a svd for the case mixer.amplitude=0.

\section*{Parameters}
- engine (SingleSiteDMRGEnginel TwoSiteDMRGEngine) - The DMRG engine calling the mixer.
- theta (Array) - The optimized wave function, prepared for svd.
- i0 (int) - Site index; theta lives on i0, i0+1.
- update_LP (bool) - Whether to calculate the next env.LP [i0+1].
- update_RP (bool) - Whether to calculate the next env.RP [i0].

\section*{Returns}
- U (Array) - Left-canonical part of theta. Labels '(vL.p0)', 'vR'.
- S (1D ndarray I 2D Array) - Without mixer just the singluar values of the array; with mixer it might be a general matrix; see comment above.
- VH (Array) - Right-canonical part of theta. Labels 'vL', '(p1.vR)'.
- \(\mathbf{e r r}\) (TruncationError) - The truncation error introduced.
mix_rho_L (engine, theta, i0, mix_enabled)
Calculated mixed reduced density matrix for left site.
Pictorially:


\section*{Parameters}
- engine (DMRGEngine) - The DMRG engine calling the mixer.
- theta (Array) - Ground state of the effective Hamiltonian, prepared for svd.
- i0 (int) - Site index; theta lives on i0, i0+1.
- mix_enabled (bool) - Whether we should perturb the density matrix.

Returns rho_L - A (hermitian) square array with labels ' (vL.p0)', '(vL*.p0*)', Mainly the reduced density matrix of the left part, but with some additional mixing.

\section*{Return type Array}
mix_rho_R (engine, theta, i0, mix_enabled)
Calculated mixed reduced density matrix for left site.
Pictorially:



\section*{Parameters}
- engine (DMRGEngine) - The DMRG engine calling the mixer.
- theta (Array) - Ground state of the effective Hamiltonian, prepared for svd.
- i0 (int) - Site index; theta lives on i0, i0+1.
- mix_enabled (bool) - Whether we should perturb the density matrix.

Returns rho_R - A (hermitian) square array with labels '(p1.vR)', '(p1*.vR*)'. Mainly the reduced density matrix of the right part, but with some additional mixing.

Return type Array
get_xR \(\left(w R \_l e g, I d \_L, I d \_R\right)\)
Generate the coupling of the MPO legs for the reduced density matrix.

\section*{Parameters}
- wR_leg (LegCharge) - LegCharge to be connected to.
- IdL (int I None) - Index within the leg for which the MPO has only identities to the left.
- IdR (int I None) - Index within the leg for which the MPO has only identities to the right.

\section*{Returns}
- mixed_xR (Array) - Connection of the MPOs on the right for the reduced density matrix rhoL. Labels ('wL', 'wL*').
- add_separate_Id (bool) - If Id_L is None, we can't include the identity into mixed_xR, so it has to be added directly in mix_rho_L ().
get_xL (wL_leg, Id_L,Id_R)
Generate the coupling of the MPO legs for the reduced density matrix.

\section*{Parameters}
- wL_leg (LegCharge) - LegCharge to be connected to.
- Id_L (int I None) - Index within the leg for which the MPO has only identities to the left.
- Id_R (int I None) - Index within the leg for which the MPO has only identities to the right.

\section*{Returns}
- mixed_xL (Array) - Connection of the MPOs on the left for the reduced density matrix rhoR. Labels ('wR', 'wR*').
- add_separate_Id (bool) - If Id_R is None, we can't include the identity into mixed_xL, so it has to be added directly in mix_rho_R().
update_amplitude (sweeps)
Update the amplitude, possibly disable the mixer.

Parameters sweeps (int) - The number of performed sweeps, to check if we need to disable the mixer.
Returns mixer - Returns self if we should continue mixing, or None, if the mixer should be disabled.

Return type Mixer I None

\subsection*{17.5.2 EngineCombine}
- full name: tenpy.algorithms.dmrg.EngineCombine
- parent module: tenpy.algorithms.dmrg
- type: class

Inheritance Diagram


\section*{Methods}
\begin{tabular}{ll}
\hline \begin{tabular}{l} 
EngineCombine.__init__(psi, \\
DMRG_params)
\end{tabular} & model,
\end{tabular} Initialize self.

\section*{Class Attributes and Properties}

EngineCombine.DMRG_params
EngineCombine.engine_params
EngineCombine.n_optimize the number of sites to be optimized over at once.
EngineCombine. verbose
```

class tenpy.algorithms.dmrg.EngineCombine(psi, model,DMRG_params)

```
    Bases: tenpy.algorithms.dmrg.TwoSiteDMRGEngine

Engine which combines legs into pipes as far as possible.
This engine combines the virtual and physical leg for the left site and right site into pipes. This reduces the overhead of calculating charge combinations in the contractions, but one matvec () is formally more expensive, \(O\left(2 d^{3} \chi^{3} D\right)\).

Deprecated since version 0.5.0: Directly use the TwoSiteDMRGEngine with the DMRG parameter combine=True.

\section*{DefaultMixer}
alias of tenpy.algorithms.dmrg.DensityMatrixMixer

\section*{EffectiveH}
alias of tenpy.algorithms.mps_common.TwoSiteH

\section*{diag (theta_guess)}

Diagonalize the effective Hamiltonian represented by self.
option DMRGEngine.max_N_for_ED: int
Maximum matrix dimension of the effective hamiltonian up to which the 'default' diag_method uses ED instead of Lanczos.
option DMRGEngine.diag_method: str
One of the folloing strings:
'default' Same as 'lanczos' for large bond dimensions, but if the total dimension of the effective Hamiltonian does not exceed the DMRG parameter 'max_N_for_ED' it uses 'ED_block'.
'lanczos’ lanczos () Default, the Lanczos implementation in TeNPy.
'arpack' lanczos_arpack() Based on scipy.linalg.sparse.eigsh(). Slower than 'lanczos', since it needs to convert the npc arrays to numpy arrays during each matvec, and possibly does many more iterations.
'ED_block' full_diag_effH() Contract the effective Hamiltonian to a (large!) matrix and diagonalize the block in the charge sector of the initial state. Preserves the charge sector of the explicitly conserved charges. However, if you don't preserve a charge explicitly, it can break it. For example if you use a SpinChain(\{'conserve': 'parity'\}), it could change the total "Sz", but not the parity of 'Sz'.
'ED_all’ full_diag_effH() Contract the effective Hamiltonian to a (large!) matrix and diagonalize it completely. Allows to change the charge sector even for explicitly conserved charges. For example if you use a SpinChain ( \(\{\) 'conserve': 'Sz'\}), it can change the total "Sz".

Parameters theta_guess (Array) - Initial guess for the ground state of the effective Hamiltonian.

\section*{Returns}
- E0 (float) - Energy of the found ground state.
- theta (Array) - Ground state of the effective Hamiltonian.
- \(\mathbf{N}(\) int \()\) - Number of Lanczos iterations used. -1 if unknown.
- ov_change (float) - Change in the wave function 1. abs (<theta_guess|theta_diag>)
environment_sweeps ( \(N_{-}\)sweeps)
Perform \(N \_\)sweeps sweeps without optimization to update the environment.
Parameters N_sweeps (int) - Number of sweeps to run without optimization
get_resume_data()
Return necessary data to resume a run () interrupted at a checkpoint.
At a checkpoint, you can save psi, model and options along with the data returned by this function. When the simulation aborts, you can resume it using this saved data with:
```

eng = AlgorithmClass(psi, model, options, resume_data=resume_data)
eng.resume_run(resume_data)

```

An algorithm which doesn't support this should override resume_run to raise an Error.
Returns resume_data - Dictionary with necessary data (apart from copies of psi, model, options) that allows to continue the simulation from where we are now.

\section*{Return type dict}
```

get_sweep_schedule()

```

Define the schedule of the sweep.
One 'sweep' is a full sequence from the leftmost site to the right and back. Only those \(L P\) and \(R P\) that can be used later should be updated.

Returns schedule - Schedule for the sweep. Each entry is (i0, move_right, (update_LP, update_RP) ), where \(i O\) is the leftmost of the self.EffectiveH. length sites to be updated in update_local (), move_right indicates whether the next \(i 0\) in the schedule is rigth (True) of the current one, and update_LP, update_RP indicate whether it is necessary to update the \(L P\) and \(R P\). The latter are chosen such that the environment is growing for infinite systems, but we only keep the minimal number of environment tensors in memory.

Return type iterable of (int, bool, (bool, bool))
init_env (model=None, resume_data=None)
(Re-)initialize the environment.
This function is useful to (re-)start a Sweep with a slightly different model or different (engine) parameters. Note that we assume that we still have the same psi. Calls reset_stats ().

\section*{Parameters}
- model (MPOModel) - The model representing the Hamiltonian for which we want to find the ground state. If None, keep the model used before.
- resume_data (None / dict) - Given when resuming a simulation, as returned by get_resume_data().

\section*{Options}

Deprecated since version 0.6.0: Options \(L P, L P \_a g e, R P\) and \(R P\) _age are now collected in a dictionary init_env_data with different keys init_LP, init_RP, age_LP, age_RP
Deprecated since version 0.8.0: Instead of passing the init_env_data as a option, it should be passed as dict entry of resume_data.
```

option Sweep.init_env_data: dict
Dictionary as returned by self.env.get_initialization_data() from
get_initialization_data().
option Sweep.orthogonal_to: list of MPS
List of other matrix product states to orthogonalize against. Works only for finite systems. This parameter can be used to find (a few) excited states as follows. First, run DMRG to find the ground

```
state and then run DMRG again while orthogonalizing against the ground state, which yields the first excited state (in the same symmetry sector), and so on.
option Sweep.start_env: int
Number of sweeps to be performed without optimization to update the environment.
Raises ValueError - If the engine is re-initialized with a new model, which legs are incompatible with those of hte old model.
make_eff_H()
Create new instance of self.EffectiveH at self.i0 and set it to self.eff_H.
mixed_svd (theta)
Get (truncated) \(B\) from the new theta (as returned by diag).
The goal is to split theta and truncate it:
\(\square\)
Without a mixer, this is done by a simple svd and truncation of Schmidt values.
With a mixer, the state is perturbed before the SVD. The details of the perturbation are defined by the Mixer class.

Note that the returned \(S\) is a general (not diagonal) matrix, with labels 'vL', 'vR'.
Parameters theta (Array) - The optimized wave function, prepared for svd.

\section*{Returns}
- U (Array) - Left-canonical part of theta. Labels '(vL.p0)', 'vR'.
- S (1D ndarray I 2D Array) - Without mixer just the singluar values of the array; with mixer it might be a general matrix with labels 'vL', 'vR'; see comment above.
- VH (Array) - Right-canonical part of theta. Labels 'vL', '(p1.vR)'.
- err (TruncationError) - The truncation error introduced.
mixer_activate()
Set self.mixer to the class specified by options['mixer'].
option TwoSiteDMRGEngine.mixer: str | class | bool
Chooses the Mixer to be used. A string stands for one of the mixers defined in this module, a class is used as custom mixer. Default (None) uses no mixer, True uses DensityMatrixMixer for the 2-site case and SingleSiteMixer for the 1-site case.
option TwoSiteDMRGEngine.mixer_params: dict Mixer parameters as described in Mixer.
mixer_cleanup ()
Cleanup the effects of a mixer.
A sweep () with an enabled Mixer leaves the MPS psi with 2D arrays in \(S\). To recover the originial form, this function simply performs one sweep with disabled mixer.
property n_optimize
the number of sites to be optimized over at once.
Indirectly set by the class attribute EffectiveH and it's length. For example, TwoSiteDMRGEngine uses the TwoSiteH and hence has n_optimize=2, while the SingleSiteDMRGEngine has n_optimize=1.
plot_sweep_stats (axes=None, xaxis='time', yaxis='E', y_exact=None, \({ }^{* *} k w a r g s\) )
Plot sweep_stats to display the convergence with the sweeps.

\section*{Parameters}
- axes (matplotlib.axes.Axes) - The axes to plot into. Defaults to matplotlib. pyplot.gca()
- xaxis (key of sweep_stats) - Key of sweep_stats to be used for the x-axis and \(y\)-axis of the plots.
- yaxis (key of sweep_stats) - Key of sweep_stats to be used for the x-axis and \(y\)-axis of the plots.
- y_exact (float) - Exact value for the quantity on the \(y\)-axis for comparison. If given, plot abs ((y-y_exact)/y_exact) on a log-scale yaxis.
- **kwargs - Further keyword arguments given to axes.plot (....).
plot_update_stats (axes, xaxis='time', yaxis='E', y_exact=None, **kwargs)
Plot update_stats to display the convergence during the sweeps.

\section*{Parameters}
- axes (matplotlib.axes.Axes) - The axes to plot into. Defaults to matplotlib. pyplot.gca()
- xaxis ('N_updates' | 'sweep' | keys of update_stats) - Key of update_stats to be used for the x-axis of the plots. 'N_updates' is just enumerating the number of bond updates, and 'sweep' corresponds to the sweep number (including environment sweeps).
- yaxis ('E' I keys of update_stats) - Key of update_stats to be used for the \(y\)-axisof the plots. For ' \(E\) ', use the energy (per site for infinite systems).
- y_exact (float) - Exact value for the quantity on the \(y\)-axis for comparison. If given, plot abs ((y-y_exact) /y_exact) on a log-scale yaxis.
- **kwargs - Further keyword arguments given to axes.plot (....).
post_update_local (update_data)
Perform post-update actions.
Compute truncation energy, remove \(L P / R P\) that are no longer needed and collect statistics.
Parameters update_data (dict) - What was returned by update_local().
prepare_svd (theta)
Transform theta into matrix for svd.
prepare_update()
Prepare self for calling update_local() on sites i0 : i0+n_optimize.
Returns theta - Current best guess for the ground state, which is to be optimized. Labels are 'vL', 'p0', 'p1', 'vR', or combined versions of it (if self.combine). For single-site DMRG, the 'p1' label is missing.

\section*{Return type Array}
reset_stats (resume_data=None)
Reset the statistics, useful if you want to start a new sweep run.
option DMRGEngine.chi_list: dict | None
A dictionary to gradually increase the chi_max parameter of trunc_params. The key defines starting from which sweep chi_max is set to the value, e.g. \(\{0: 50,20: 100\}\) uses chi_max=50 for
the first 20 sweeps and chi_max=100 afterwards. Overwrites trunc_params['chi_list']. By default (None) this feature is disabled.
option DMRGEngine.sweep_0: int
The number of sweeps already performed. (Useful for re-start).
resume_run ()
Resume a run that was interrupted.
In case we saved an intermediate result at a checkpoint, this function allows to resume the run () of the algorithm (after re-initialization with the resume_data). Since most algorithms just have a while loop with break conditions, the default behaviour implemented here is to just call run ().
run()
Run the DMRG simulation to find the ground state.

\section*{Returns}
- \(\mathbf{E}(\) float \()\) - The energy of the resulting ground state MPS.
- psi (MPS) - The MPS representing the ground state after the simluation, i.e. just a reference to psi.

\section*{Options}
option DMRGEngine.diag_method: str Method to be used for diagonalzation, default 'default'. For possible arguments see DMRGEngine.diag().
option DMRGEngine.E_tol_to_trunc: float
It's reasonable to choose the Lanczos convergence criteria 'E_tol' not many magnitudes lower than the current truncation error. Therefore, if \(E_{-}\)tol_to_trunc is not None, we update E_tol of lanczos_params to max_E_trunc*E_tol_to_trunc, restricted to the interval [E_tol_min, E_tol_max], where max_E_trunc is the maximal energy difference due to truncation right after each Lanczos optimization during the sweeps.
option DMRGEngine.E_tol_max: float See E_tol_to_trunc
option DMRGEngine.E_tol_min: float See E_tol_to_trunc
option DMRGEngine.max_E_err: float
Convergence if the change of the energy in each step satisfies -Delta \(E / \max (|E|, 1)<\) max_E_err. Note that this is also satisfied if Delta E > 0, i.e., if the energy increases (due to truncation).
option DMRGEngine.max_hours: float
If the DMRG took longer (measured in wall-clock time), 'shelve' the simulation, i.e. stop and return with the flag shelve=True.
option DMRGEngine.max_S_err: float Convergence if the relative change of the entropy in each step satisfies |Delta \(S \| / \mathrm{S}<\) max_S_err
option DMRGEngine.max_sweeps: int Maximum number of sweeps to be performed.
option DMRGEngine.min_sweeps: int
Minimum number of sweeps to be performed. Defaults to \(1.5 *\) N_sweeps_check.
option DMRGEngine.N_sweeps_check: int
Number of sweeps to perform between checking convergence criteria and giving a status update.
option DMRGEngine.norm_tol: float
After the DMRG run, update the environment with at most norm_tol_iter sweeps until np. linalg. norm(psi.norm_err()) < norm_tol.
option DMRGEngine.norm_tol_iter: float
Perform at most norm_tol_iter \({ }^{*}\) `update_env sweeps to converge the norm error below norm_tol. If the state is not converged after that, call canonical_form () instead.
option DMRGEngine.P_tol_to_trunc: float
It's reasonable to choose the Lanczos convergence criteria ' \(P\) _tol' not many magnitudes lower than the current truncation error. Therefore, if \(P_{-}\)tol_to_trunc is not None, we update \(P_{-}\)tol of lanczos_params to max_trunc_err*P_tol_to_trunc, restricted to the interval [P_tol_min, \(P_{-}\)tol_max], where max_trunc_err is the maximal truncation error (discarded weight of the Schmidt values) due to truncation right after each Lanczos optimization during the sweeps.
option DMRGEngine.P_tol_max: float
See \(P_{-}\)tol_to_trunc
option DMRGEngine.P_tol_min: float
See P_tol_to_trunc
option DMRGEngine.update_env: int
Number of sweeps without bond optimizaiton to update the environment for infinite boundary conditions, performed every \(N_{-}\)sweeps_check sweeps.
\(\boldsymbol{s e t} \mathbf{B}(U, S, V H)\)
Update the MPS with the U, S, VH returned by self.mixed_svd.

\section*{Parameters}
- U (Array) - Left and Right-canonical matrices as returned by the SVD.
- VH (Array) - Left and Right-canonical matrices as returned by the SVD.
- S (1D array I 2D Array) - The middle part returned by the SVD, theta \(=\mathrm{U}\) S VH. Without a mixer just the singular values, with enabled mixer a 2D array.
sweep (optimize=True, meas_E_trunc=False)
One 'sweep' of a the algorithm.
Iteratate over the bond which is optimized, to the right and then back to the left to the starting point.

\section*{Parameters}
- optimize (bool, optional) - Whether we actually optimize to find the ground state of the effective Hamiltonian. (If False, just update the environments).
- meas_E_trunc (bool, optional) - Whether to measure truncation energies.

\section*{Returns}
- max_trunc_err (float) - Maximal truncation error introduced.
- max_E_trunc (None I float) - None if meas_E_trunc is False, else the maximal change of the energy due to the truncation.
update_LP ( \(U\) )
Update left part of the environment.
We always update the environment at site \(i 0+1\) : this environment then contains the site where we just performed a local update (when sweeping right).

Parameters U (Array) - The U as returned by the SVD, with combined legs, labels 'vL. po', 'vR'.
update_RP (VH)
Update right part of the environment.
We always update the environment at site i 0 : this environment then contains the site where we just performed a local update (when sweeping left).

Parameters VH (Array) - The VH as returned by SVD, with combined legs, labels 'vL', '(vR.p1)'.
update_local (theta, optimize=True)
Perform site-update on the site \(i 0\).

\section*{Parameters}
- theta (Array) - Initial guess for the ground state of the effective Hamiltonian.
- optimize (bo○l) - Wheter we actually optimize to find the ground state of the effective Hamiltonian. (If False, just update the environments).

\section*{Returns}
update_data - Data computed during the local update, as described in the following:
E0 [float] Total energy, obtained before truncation (if optimize=True), or after truncation (if optimize=False) (but never None).
\(\mathbf{N}\) [int] Dimension of the Krylov space used for optimization in the lanczos algorithm. 0 if optimize=False.
age [int] Current size of the DMRG simulation: number of physical sites involved into the contraction.

U, VH: Array \(U\) and \(V H\) returned by mixed_svd().
ov_change: float Change in the wave function 1. - abs (<theta_guess|theta>) induced by diag (), not including the truncation!

\section*{Return type dict}

\subsection*{17.5.3 EngineFracture}
- full name: tenpy.algorithms.dmrg.EngineFracture
- parent module: tenpy.algorithms.dmrg
- type: class

\section*{Inheritance Diagram}


\section*{Methods}
\begin{tabular}{|c|c|}
\hline EngineFracture.__init_(psi, model, DMRG_params) & Initialize self. \\
\hline EngineFracture.diag(theta_guess) & Diagonalize the effective Hamiltonian represented by self. \\
\hline EngineFracture.environment_sweeps(N_sw & eper)form \(N_{-}\)sweeps sweeps without optimization to update the environment. \\
\hline EngineFracture.get_resume_data() & Return necessary data to resume a run () interrupted at a checkpoint. \\
\hline EngineFracture.get_sweep_schedule() & Define the schedule of the sweep. \\
\hline EngineFracture.init_env([model, sume_data]) & (Re-)initialize the environment. \\
\hline EngineFracture.make_eff_H() & Create new instance of self.EffectiveH at self.iO and set it to self.eff_H. \\
\hline EngineFracture.mixed_svd(theta) & Get (truncated) \(B\) from the new theta (as returned by diag). \\
\hline EngineFracture.mixer_activate() & Set self.mixer to the class specified by options['mixer']. \\
\hline EngineFracture.mixer_cleanup() & Cleanup the effects of a mixer. \\
\hline ```
EngineFracture.plot_sweep_stats([axes,
...])
``` & Plot sweep_stats to display the convergence with the sweeps. \\
\hline
\end{tabular}

Table 21 - continued from previous page
\begin{tabular}{|c|c|}
\hline \[
\begin{aligned}
& \text { EngineFracture.plot_update_stats(axes[, } \\
& \ldots .] \text { ) }
\end{aligned}
\] & Plot update_stats to display the convergence during the sweeps. \\
\hline \multicolumn{2}{|l|}{EngineFracture.post_update_local(update_dalatform post-update actions.} \\
\hline EngineFracture.prepare_svd(theta) & Transform theta into matrix for svd. \\
\hline EngineFracture.prepare_update() & Prepare self for calling update_local () on sites io : i0+n_optimize. \\
\hline EngineFracture.reset_stats([resume_data]) & Reset the statistics, useful if you want to start a new sweep run. \\
\hline EngineFracture.resume_run() & Resume a run that was interrupted. \\
\hline EngineFracture.run() & Run the DMRG simulation to find the ground state. \\
\hline EngineFracture.set_B(U, S, VH) & Update the MPS with the U, S, VH returned by self.mixed_svd. \\
\hline EngineFracture.sweep([optimize, meas_E_trunc]) & One 'sweep' of a the algorithm. \\
\hline EngineFracture.update_LP(U) & Update left part of the environment. \\
\hline EngineFracture.update_RP(VH) & Update right part of the environment. \\
\hline EngineFracture.update_local(theta[, optimize]) & Perform site-update on the site io. \\
\hline
\end{tabular}

\section*{Class Attributes and Properties}
\begin{tabular}{l}
\hline EngineFracture.DMRG_params \\
\hline EngineFracture.engine_params \\
\hline EngineFracture.n_optimize \\
\hline EngineFracture.verbose
\end{tabular}
class tenpy.algorithms.dmrg. EngineFracture (psi, model, DMRG_params)
Bases: tenpy.algorithms.dmrg.TwoSiteDMRGEngine
Engine which keeps the legs separate.
Due to a different contraction order in matvec (), this engine might be faster than EngineCombine, at least for large physical dimensions and if the MPO is sparse. One matvec () is \(O\left(2 \chi^{3} d^{2} W+2 \chi^{2} d^{3} W^{2}\right)\).
Deprecated since version 0.5.0: Directly use the TwoSiteDMRGEngine with the DMRG parameter combine=False.

\section*{DefaultMixer}
alias of tenpy.algorithms.dmrg.DensityMatrixMixer

\section*{EffectiveH}
alias of tenpy.algorithms.mps_common. TwoSiteH
diag (theta_guess)
Diagonalize the effective Hamiltonian represented by self.
option DMRGEngine.max_N_for_ED: int
Maximum matrix dimension of the effective hamiltonian up to which the 'default' diag_method uses ED instead of Lanczos.
option DMRGEngine.diag_method: str

One of the folloing strings:
'default' Same as 'lanczos' for large bond dimensions, but if the total dimension of the effective Hamiltonian does not exceed the DMRG parameter 'max_N_for_ED' it uses 'ED_block'.
'lanczos' Ianczos () Default, the Lanczos implementation in TeNPy.
'arpack' lanczos_arpack() Based on scipy.linalg.sparse.eigsh(). Slower than 'lanczos', since it needs to convert the npc arrays to numpy arrays during each matvec, and possibly does many more iterations.
'ED_block' full_diag_effH() Contract the effective Hamiltonian to a (large!) matrix and diagonalize the block in the charge sector of the initial state. Preserves the charge sector of the explicitly conserved charges. However, if you don't preserve a charge explicitly, it can break it. For example if you use a SpinChain(\{'conserve': 'parity'\}), it could change the total "Sz", but not the parity of 'Sz'.
'ED_all' full_diag_effH() Contract the effective Hamiltonian to a (large!) matrix and diagonalize it completely. Allows to change the charge sector even for explicitly conserved charges. For example if you use a SpinChain ( \(\{\) 'conserve': 'Sz'\}), it can change the total "Sz".

Parameters theta_guess (Array) - Initial guess for the ground state of the effective Hamiltonian.

\section*{Returns}
- E0 (float) - Energy of the found ground state.
- theta (Array) - Ground state of the effective Hamiltonian.
- \(\mathbf{N}(\) int \()\) - Number of Lanczos iterations used. -1 if unknown.
- ov_change (float) - Change in the wave function 1. abs (<theta_guess|theta_diag>)
environment_sweeps ( \(N\) _sweeps)
Perform \(N_{-}\)sweeps sweeps without optimization to update the environment.
Parameters N_sweeps (int) - Number of sweeps to run without optimization
get_resume_data()
Return necessary data to resume a run () interrupted at a checkpoint.
At a checkpoint, you can save psi, model and options along with the data returned by this function. When the simulation aborts, you can resume it using this saved data with:
```

eng = AlgorithmClass(psi, model, options, resume_data=resume_data)
eng.resume_run(resume_data)

```

An algorithm which doesn't support this should override resume_run to raise an Error.
Returns resume_data - Dictionary with necessary data (apart from copies of psi, model, options) that allows to continue the simulation from where we are now.

Return type dict
get_sweep_schedule()
Define the schedule of the sweep.
One 'sweep' is a full sequence from the leftmost site to the right and back. Only those \(L P\) and \(R P\) that can be used later should be updated.

Returns schedule - Schedule for the sweep. Each entry is (i0, move_right, (update_LP, update_RP)), where \(i 0\) is the leftmost of the self.EffectiveH. length sites to be updated in update_local (), move_right indicates whether the next \(i 0\) in the schedule is rigth (True) of the current one, and update_LP, update_RP indicate
whether it is necessary to update the \(L P\) and \(R P\). The latter are chosen such that the environment is growing for infinite systems, but we only keep the minimal number of environment tensors in memory.
Return type iterable of (int, bool, (bool, bool))
init_env (model=None, resume_data=None)
(Re-)initialize the environment.
This function is useful to (re-)start a Sweep with a slightly different model or different (engine) parameters.
Note that we assume that we still have the same psi. Calls reset_stats ().

\section*{Parameters}
- model (MPOModel) - The model representing the Hamiltonian for which we want to find the ground state. If None, keep the model used before.
- resume_data (None / dict) - Given when resuming a simulation, as returned by get_resume_data().

\section*{Options}

Deprecated since version 0.6.0: Options \(L P, L P \_\)age, \(R P\) and \(R P\) _age are now collected in a dictionary init_env_data with different keys init_LP, init_RP, age_LP, age_RP
Deprecated since version 0.8.0: Instead of passing the init_env_data as a option, it should be passed as dict entry of resume_data.
option Sweep.init_env_data: dict
Dictionary as returned by self.env.get_initialization_data() from get_initialization_data().
option Sweep.orthogonal_to: list of MPS
List of other matrix product states to orthogonalize against. Works only for finite systems. This parameter can be used to find (a few) excited states as follows. First, run DMRG to find the ground state and then run DMRG again while orthogonalizing against the ground state, which yields the first excited state (in the same symmetry sector), and so on.
option Sweep.start_env: int Number of sweeps to be performed without optimization to update the environment.

Raises ValueError - If the engine is re-initialized with a new model, which legs are incompatible with those of hte old model.
make_eff_H()
Create new instance of self.EffectiveH at self.i0 and set it to self.eff_ \(H\).

\section*{mixed_svd (theta)}

Get (truncated) \(B\) from the new theta (as returned by diag).
The goal is to split theta and truncate it:


Without a mixer, this is done by a simple svd and truncation of Schmidt values.
With a mixer, the state is perturbed before the SVD. The details of the perturbation are defined by the Mixer class.

Note that the returned \(S\) is a general (not diagonal) matrix, with labels 'vL', 'vR'.

Parameters theta (Array) - The optimized wave function, prepared for svd.

\section*{Returns}
- \(\mathbf{U}\) (Array) - Left-canonical part of theta. Labels '(vL.p0)', 'vR'.
- S (1D ndarray I 2D Array) - Without mixer just the singluar values of the array; with mixer it might be a general matrix with labels 'vL', 'vR'; see comment above.
- VH (Array) - Right-canonical part of theta. Labels 'vL', '(p1.vR)'.
- \(\mathbf{e r r}\) (TruncationError) - The truncation error introduced.
mixer_activate()
Set self.mixer to the class specified by options['mixer'].
option TwoSiteDMRGEngine.mixer: str | class | bool
Chooses the Mixer to be used. A string stands for one of the mixers defined in this module, a class is used as custom mixer. Default (None) uses no mixer, True uses DensityMatrixMixer for the 2-site case and SingleSiteMixer for the 1-site case.
option TwoSiteDMRGEngine.mixer_params: dict Mixer parameters as described in Mixer.
mixer_cleanup ()
Cleanup the effects of a mixer.
A sweep () with an enabled Mixer leaves the MPS psi with 2D arrays in S. To recover the originial form, this function simply performs one sweep with disabled mixer.

\section*{property n_optimize}
the number of sites to be optimized over at once.
Indirectly set by the class attribute EffectiveH and it's length. For example, TwoSiteDMRGEngine uses the TwoSiteH and hence has n_optimize=2, while the SingleSiteDMRGEngine has n_optimize=1.
plot_sweep_stats (axes=None, xaxis='time', yaxis='E', y_exact=None, **kwargs)
Plot sweep_stats to display the convergence with the sweeps.

\section*{Parameters}
- axes (matplotlib.axes.Axes) - The axes to plot into. Defaults to matplotlib. pyplot.gca()
- xaxis (key of sweep_stats) - Key of sweep_stats to be used for the x-axis and \(y\)-axis of the plots.
- yaxis (key of sweep_stats) - Key of sweep_stats to be used for the x-axis and \(y\)-axis of the plots.
- y_exact ( \(f\) loat) - Exact value for the quantity on the y-axis for comparison. If given, plot abs ((y-y_exact)/y_exact) on a log-scale yaxis.
- **kwargs - Further keyword arguments given to axes.plot (....).
plot_update_stats (axes, xaxis='time', yaxis='E', y_exact=None, **kwargs)
Plot update_stats to display the convergence during the sweeps.

\section*{Parameters}
- axes (matplotlib.axes.Axes) - The axes to plot into. Defaults to matplotlib. pyplot.gca()
- xaxis ('N_updates' | 'sweep' | keys of update_stats) - Key of update_stats to be used for the x-axis of the plots. 'N_updates' is just enumerating the number of bond updates, and 'sweep' corresponds to the sweep number (including environment sweeps).
- yaxis ('E' I keys of update_stats) - Key of update_stats to be used for the \(y\)-axisof the plots. For ' \(E\) ', use the energy (per site for infinite systems).
- Y_exact (float) - Exact value for the quantity on the \(y\)-axis for comparison. If given, plot abs ((y-y_exact) /y_exact) on a log-scale yaxis.
- **kwargs - Further keyword arguments given to axes.plot (....).
```

post_update_local (update_data)

```

Perform post-update actions.
Compute truncation energy, remove \(L P / R P\) that are no longer needed and collect statistics.
Parameters update_data (dict) - What was returned by update_local().

\section*{prepare_svd (theta)}

Transform theta into matrix for svd.
```

prepare_update()

```

Prepare self for calling update_local() on sites i0 : i0+n_optimize.
Returns theta - Current best guess for the ground state, which is to be optimized. Labels are 'vL', 'p0', 'p1', 'vR', or combined versions of it (if self.combine). For single-site DMRG, the ' p 1 ' label is missing.
Return type Array
reset_stats (resume_data=None)
Reset the statistics, useful if you want to start a new sweep run.
option DMRGEngine.chi_list: dict | None
A dictionary to gradually increase the chi_max parameter of trunc_params. The key defines starting from which sweep chi_max is set to the value, e.g. \(\{0: 50,20: 100\}\) uses chi_max=50 for the first 20 sweeps and chi_max=100 afterwards. Overwrites trunc_params['chi_list']'. By default (None) this feature is disabled.
option DMRGEngine.sweep_0: int
The number of sweeps already performed. (Useful for re-start).

\section*{resume_run()}

Resume a run that was interrupted.
In case we saved an intermediate result at a checkpoint, this function allows to resume the run () of the algorithm (after re-initialization with the resume_data). Since most algorithms just have a while loop with break conditions, the default behaviour implemented here is to just call run ().
run()
Run the DMRG simulation to find the ground state.

\section*{Returns}
- \(\mathbf{E}(\) float \()\) - The energy of the resulting ground state MPS.
- psi (MPS) - The MPS representing the ground state after the simluation, i.e. just a reference to psi.

\section*{Options}
option DMRGEngine.diag_method: str
Method to be used for diagonalzation, default 'default'. For possible arguments see DMRGEngine. diag().
option DMRGEngine.E_tol_to_trunc: float
It's reasonable to choose the Lanczos convergence criteria 'E_tol' not many magnitudes lower than the current truncation error. Therefore, if \(E_{-}\)tol_to_trunc is not None, we update E_tol of lanczos_params to max_E_trunc*E_tol_to_trunc, restricted to the interval [E_tol_min, E_tol_max], where max_E_trunc is the maximal energy difference due to truncation right after each Lanczos optimization during the sweeps.
option DMRGEngine.E_tol_max: float See E_tol_to_trunc
option DMRGEngine.E_tol_min: float See E_tol_to_trunc
option DMRGEngine.max_E_err: float Convergence if the change of the energy in each step satisfies -Delta \(E / \max (|E|, 1)<\) max_E_err. Note that this is also satisfied if Delta \(E>0\), i.e., if the energy increases (due to truncation).
option DMRGEngine.max_hours: float
If the DMRG took longer (measured in wall-clock time), 'shelve' the simulation, i.e. stop and return with the flag shelve=True.
option DMRGEngine.max_S_err: float Convergence if the relative change of the entropy in each step satisfies |Delta \(\mathrm{S} \mid / \mathrm{S}<\) max_S_err
option DMRGEngine.max_sweeps: int Maximum number of sweeps to be performed.
option DMRGEngine.min_sweeps: int Minimum number of sweeps to be performed. Defaults to \(1.5 *\) N_sweeps_check.
option DMRGEngine.N_sweeps_check: int Number of sweeps to perform between checking convergence criteria and giving a status update.
option DMRGEngine.norm_tol: float After the DMRG run, update the environment with at most norm_tol_iter sweeps until np.linalg. norm(psi.norm_err()) < norm_tol.
option DMRGEngine.norm_tol_iter: float Perform at most norm_tol_iter`* update_env sweeps to converge the norm error below norm_tol. If the state is not converged after that, call canonical_form () instead.
option DMRGEngine.P_tol_to_trunc: float
It's reasonable to choose the Lanczos convergence criteria ' \(P\) _tol' not many magnitudes lower than the current truncation error. Therefore, if \(P_{-}\)tol_to_trunc is not None, we update \(P_{-}\)tol of lanczos_params to max_trunc_err*P_tol_to_trunc, restricted to the interval [P_tol_min, \(P_{-}\)tol_max], where max_trunc_err is the maximal truncation error (discarded weight of the Schmidt values) due to truncation right after each Lanczos optimization during the sweeps.
```

option DMRGEngine.P_tol_max: float
See P_tol_to_trunc
option DMRGEngine.P_tol_min: float
See P_tol_to_trunc

```
option DMRGEngine.update_env: int
Number of sweeps without bond optimizaiton to update the environment for infinite boundary conditions, performed every \(N_{\text {_sweeps_check sweeps. }}^{\text {swe }}\)
\(\boldsymbol{s e t} \mathbf{B}(U, S, V H)\)
Update the MPS with the U, S, VH returned by self.mixed_svd.

\section*{Parameters}
- U (Array) - Left and Right-canonical matrices as returned by the SVD.
- VH (Array) - Left and Right-canonical matrices as returned by the SVD.
- S (1D array I 2D Array) - The middle part returned by the SVD, theta \(=\mathrm{U} S \mathrm{VH}\). Without a mixer just the singular values, with enabled mixer a 2D array.
sweep (optimize=True, meas_E_trunc=False)
One 'sweep' of a the algorithm.
Iteratate over the bond which is optimized, to the right and then back to the left to the starting point.

\section*{Parameters}
- optimize (bool, optional) - Whether we actually optimize to find the ground state of the effective Hamiltonian. (If False, just update the environments).
- meas_E_trunc (bool, optional) - Whether to measure truncation energies.

\section*{Returns}
- max_trunc_err (float) - Maximal truncation error introduced.
- max_E_trunc (None I float) - None if meas_E_trunc is False, else the maximal change of the energy due to the truncation.
update_LP \((U)\)
Update left part of the environment.
We always update the environment at site \(i 0+1\) : this environment then contains the site where we just performed a local update (when sweeping right).

Parameters U (Array) - The U as returned by the SVD, with combined legs, labels 'vL. p0', 'vR'.
update_RP (VH)
Update right part of the environment.
We always update the environment at site i 0 : this environment then contains the site where we just performed a local update (when sweeping left).

Parameters VH (Array) - The VH as returned by SVD, with combined legs, labels 'vL', '(vR.p1)'.
```

update_local (theta, optimize=True)

```

Perform site-update on the site io.

\section*{Parameters}
- theta (Array) - Initial guess for the ground state of the effective Hamiltonian.
- optimize (b○○1) - Wheter we actually optimize to find the ground state of the effective Hamiltonian. (If False, just update the environments).

\section*{Returns}
update_data - Data computed during the local update, as described in the following:

E0 [float] Total energy, obtained before truncation (if optimize=True), or after truncation (if optimize=False) (but never None).
\(\mathbf{N}\) [int] Dimension of the Krylov space used for optimization in the lanczos algorithm. 0 if optimize=False.
age [int] Current size of the DMRG simulation: number of physical sites involved into the contraction.

U, VH: Array \(U\) and \(V H\) returned by mixed_svd().
ov_change: float Change in the wave function 1. - abs (<theta_guess|theta>) induced by diag(), not including the truncation!
Return type dict

\subsection*{17.5.4 SingleSiteMixer}
- full name: tenpy.algorithms.dmrg.SingleSiteMixer
- parent module: tenpy.algorithms.dmrg
- type: class

\section*{Inheritance Diagram}


\section*{Methods}
\begin{tabular}{ll}
\hline SingleSiteMixer.__init__(options) & Initialize self. \\
\hline SingleSiteMixer.perturb_svd(engine, theta, & Mix extra terms to theta and perform an SVD. \\
\(\ldots\)..) & \\
\hline SingleSiteMixer.subspace_expand(engine, & \begin{tabular}{l} 
Expand the MPS subspace, to allow the bond dimension \\
to increase.
\end{tabular} \\
\begin{tabular}{ll} 
SingleSiteMixer. & Update the amplitude, possibly disable the mixer. \\
update_amplitude(sweeps) & \\
\hline
\end{tabular}
\end{tabular}
class tenpy.algorithms.dmrg.SingleSiteMixer (options)
Bases: tenpy.algorithms.dmrg.Mixer
Mixer for single-site DMRG.

Performs a subspace expansion following [[hubig2015]].
perturb_svd (engine, theta, i0, move_right, next_B)
Mix extra terms to theta and perform an SVD.
We calculate the left and right reduced density matrix using the mixer (which might include applications of \(H\) ). These density matrices are diagonalized and truncated such that we effectively perform a svd for the case mixer. amplitude \(=0\).

\section*{Parameters}
- engine (DMRGEngine) - The DMRG engine calling the mixer.
- theta (Array) - The optimized wave function, prepared for svd.
- i0 (int) - The site index where theta lives.
- move_right (bool) - Whether we move to the right (True) or left (False).
- next_B (Array) - The subspace expansion requires to change the tensor on the next site as well. If move_right, it should correspond to engine.psi.get_B(i0+1, form='B'). If not move_right, it should correspond to engine.psi.get_B(i0-1, form='A').

\section*{Returns}
- U (Array) - Left-canonical part of tensordot(theta, next_B). Labels '(vL.p0)', 'vR'.
- S (1D ndarray) - (Perturbed) singular values on the new bond (between theta and next_B).
- VH (Array) - Right-canonical part of tensordot(theta, next_B). Labels 'vL', '(p1. vR)'.
- err (TruncationError) - The truncation error introduced.
subspace_expand (engine, theta, \(i 0\), move_right, next_B)
Expand the MPS subspace, to allow the bond dimension to increase.
This is the subspace expansion following [[hubig2015]].

\section*{Parameters}
- engine (DMRGEngine) - The DMRG engine calling the mixer.
- theta (Array) - Optimized guess for the ground state of the effective local Hamiltonian.
- io (int) - Site index at which the local update has taken place.
- move_right (bool) - Whether the next \(i 0\) of the sweep will be right or left of the current one.
- next_B (Array) - The subspace expansion requires to change the tensor on the next site as well. If move_right, it should correspond to engine.psi.get_B(i0+1, form='B'). If not move_right, it should correspond to engine.psi.get_B(i0-1, form='A').

\section*{Returns}
- theta - Local MPS tensor at site \(i 0\) after subspace expansion.
- next_B - MPS tensor at site \(i 0+1\) or \(i 0-1\) (depending on sweep direction) after subspace expansion.
update_amplitude (sweeps)
Update the amplitude, possibly disable the mixer.
Parameters sweeps (int) - The number of performed sweeps, to check if we need to disable the mixer.

Returns mixer - Returns self if we should continue mixing, or None, if the mixer should be disabled.

Return type Mixer I None

\subsection*{17.5.5 TwoSiteMixer}
- full name: tenpy.algorithms.dmrg.TwoSiteMixer
- parent module: tenpy.algorithms.dmrg
- type: class

\section*{Inheritance Diagram}


\section*{Methods}
\begin{tabular}{ll}
\hline TwoSiteMixer.__init__(options) & Initialize self. \\
\hline \begin{tabular}{l} 
TwoSiteMixer.perturb_Svd(engine, theta, i0, \\
\(\ldots\)..)
\end{tabular} & Mix extra terms to theta and perform an SVD. \\
\hline TwoSiteMixer.subspace_expand(engine, & \begin{tabular}{l} 
Expand the MPS subspace, to allow the bond dimension \\
to increase.
\end{tabular} \\
theta,...) & \\
\hline TwoSiteMixer.update_amplitude(sweeps) & Update the amplitude, possibly disable the mixer. \\
\hline & \\
class tenpy.algorithms.dmrg.TwoSiteMixer (options) \\
Bases: tenpy.algorithms.dmrg.SingleSiteMixer \\
Mixer for two-site DMRG.
\end{tabular}

This is the two-site version of the mixer described in [[hubig2015]]. Equivalent to the DensityMatrixMixer, but never construct the full density matrix.
perturb_svd (engine, theta, iO, move_right)
Mix extra terms to theta and perform an SVD.

\section*{Parameters}
- engine (DMRGEngine) - The DMRG engine calling the mixer.
- theta (Array) - The optimized wave function, prepared for svd.
- i0 (int) - Site index; theta lives on i0, i0+1.
- update_LP (bool) - Whether to calculate the next env.LP [i0+1].
- update_RP (bool) - Whether to calculate the next env. RP [i0].

\section*{Returns}
- U(Array) - Left-canonical part of theta. Labels '(vL.p0)', 'vR'.
- S (1D ndarray | 2D Array) - Without mixer just the singluar values of the array; with mixer it might be a general matrix; see comment above.
- VH (Array) - Right-canonical part of theta. Labels 'vL', '(vR.p1)'.
- \(\mathbf{e r r}\) (TruncationError) - The truncation error introduced.
subspace_expand (engine, theta, i0, move_right, next_B)
Expand the MPS subspace, to allow the bond dimension to increase.
This is the subspace expansion following [[hubig2015]].

\section*{Parameters}
- engine (DMRGEngine) - The DMRG engine calling the mixer.
- theta (Array) - Optimized guess for the ground state of the effective local Hamiltonian.
- i0 (int) - Site index at which the local update has taken place.
- move_right (bool) - Whether the next \(i 0\) of the sweep will be right or left of the current one.
- next_B (Array) - The subspace expansion requires to change the tensor on the next site as well. If move_right, it should correspond to engine.psi.get_B(i0+1, form='B'). If not move_right, it should correspond to engine.psi.get_B(i0-1, form='A').

\section*{Returns}
- theta - Local MPS tensor at site \(i 0\) after subspace expansion.
- next_B - MPS tensor at site \(i 0+1\) or \(i 0-1\) (depending on sweep direction) after subspace expansion.
update_amplitude (sweeps)
Update the amplitude, possibly disable the mixer.
Parameters sweeps (int) - The number of performed sweeps, to check if we need to disable the mixer.

Returns mixer - Returns self if we should continue mixing, or None, if the mixer should be disabled.

\section*{Return type Mixer I None}

Functions
\begin{tabular}{ll}
\hline chi_list(chi_max[, dchi, nsweeps]) & Compute a 'ramping-up' chi_list. \\
\hline full_diag_effH(effH, theta_guess[, keep_sector]) & Perform an exact diagonalization of effH. \\
\hline run(psi, model, options) & \begin{tabular}{l} 
Run the DMRG algorithm to find the ground state of the \\
given model.
\end{tabular} \\
\hline
\end{tabular}

\subsection*{17.5.6 chi list}
- full name: tenpy.algorithms.dmrg.chi_list
- parent module: tenpy.algorithms.dmrg
- type: function
tenpy.algorithms.dmrg.chi_list (chi_max, dchi=20, nsweeps=20)
Compute a 'ramping-up' chi_list.
The resulting chi_list allows to increases chi by dchi every nsweeps sweeps up to a given maximal chi_max.

\section*{Parameters}
- chi_max (int) - Final value for the bond dimension.
- dchi (int) - Step size how to increase chi
- nsweeps (int) - Step size for sweeps

Returns chi_list - To be used as chi_list parameter for DMRG, see run (). Keys increase by nsweeps, values by dchi, until a maximum of chi_max is reached.
Return type dict

\subsection*{17.5.7 full_diag_effH}
- full name: tenpy.algorithms.dmrg.full_diag_effH
- parent module: tenpy.algorithms.dmrg
- type: function
tenpy.algorithms.dmrg.full_diag_effH (effH, theta_guess, keep_sector=True)
Perform an exact diagonalization of effH.
This function offers an alternative to lanczos().

\section*{Parameters}
- effin (EffectiveH) - The effective Hamiltonian.
- theta_guess (Array) - Current guess to select the charge sector. Labels as specified by effi.acts_on.

\section*{Module description}

Density Matrix Renormalization Group (DMRG).
Although it was originally not formulated with tensor networks, the DMRG algorithm (invented by Steven White in 1992 [[white1992]]) opened the whole field with its enormous success in finding ground states in 1D.
We implement DMRG in the modern formulation of matrix product states [[schollwoeck2011]], both for finite systems ('finite' or 'segment' boundary conditions) and in the thermodynamic limit ('infinite' b.c.).

The function run () - well - runs one DMRG simulation. Internally, it generates an instance of an Sweep. This class implements the common functionality like defining a sweep, but leaves the details of the contractions to be performed to the derived classes.

Currently, there are two derived classes implementing the contractions: SingleSiteDMRGEngine and TwoSiteDMRGEngine. They differ (as their name implies) in the number of sites which are optimized simultaneously. They should both give the same results (up to rounding errors). However, if started from a product state, SingleSiteDMRGEngine depends critically on the use of a Mixer, while TwoSiteDMRGEngine is in principle more computationally expensive to run and has occasionally displayed some convergence issues.. Which one is preffered in the end is not obvious a priori and might depend on the used model. Just try both of them.

A Mixer should be used initially to avoid that the algorithm gets stuck in local energy minima, and then slowly turned off in the end. For SingleSiteDMRGEngine, using a mixer is crucial, as the one-site algorithm cannot increase the MPS bond dimension by itself.

A generic protocol for approaching a physics question using DMRG is given in /intro/protocol.

\section*{17.6 tdvp}
- full name: tenpy.algorithms.tdvp
- parent module: tenpy.algorithms
- type: module

\section*{Classes}

\begin{tabular}{ll}
\hline Engine(psi, model, options) & Deprecated old name of TDVPEngine. \\
\hline H0_mixed \((\mathrm{Lp}, \mathrm{Rp})\) & Class defining the zero site Hamiltonian for Lanczos. \\
\hline H1_mixed(Lp, Rp, W) & Class defining the one site Hamiltonian for Lanczos. \\
\hline H2_mixed(Lp, Rp, W0, W1) & Class defining the two sites Hamiltonian for Lanczos. \\
\hline TDVPEngine(psi, model, options[, environment]) & Time dependent variational principle algorithm for \\
& MPS. \\
\hline
\end{tabular}

\subsection*{17.6.1 Engine}
- full name: tenpy.algorithms.tdvp.Engine
- parent module: tenpy.algorithms.tdvp
- type: class

\section*{Inheritance Diagram}


\section*{Methods}
\begin{tabular}{ll}
\hline Engine.__init__(psi, model, options) & Initialize self. \\
\hline Engine.get_resume_data() & \begin{tabular}{l} 
Return necessary data to resume a run () interrupted at \\
a checkpoint.
\end{tabular} \\
\hline Engine.resume_run() & Resume a run that was interrupted. \\
\hline Engine.run() & (Real-)time evolution with TDVP. \\
\hline Engine.run_one_site([N_steps]) & Run the TDVP algorithm with the one site algorithm. \\
\hline Engine.run_two_sites([N_steps]) & Run the TDVP algorithm with two sites update. \\
\hline Engine.set_anonymous_svd(U, new_label) & Relabel the svd. \\
\hline Engine.sweep_left_right() & \begin{tabular}{l} 
Performs the sweep left->right of the second order \\
TDVP scheme with one site update.
\end{tabular} \\
\hline Engine.sweep_left_right_two() & \begin{tabular}{l} 
Performs the sweep left->right of the second order \\
\\
TDVP scheme with two sites update.
\end{tabular} \\
\hline Engine.sweep_right_left() & \begin{tabular}{l} 
Performs the sweep right->left of the second order \\
\\
TDVP scheme with one site update.
\end{tabular} \\
\hline Engine.sweep_right_left_two() & \begin{tabular}{l} 
Performs the sweep left->right of the second order \\
\\
TDVP scheme with two sites update.
\end{tabular} \\
\hline Engine.theta_svd_left_right(theta) & Performs the SVD from left to right. \\
\hline Engine.theta_svd_right_left(theta) & Performs the SVD from right to left. \\
\hline Engine.update_s_ho(s, H, dt) & \begin{tabular}{l} 
Update with the zero site Hamiltonian (update of the \\
singular value)
\end{tabular} \\
\hline Engine.update_theta_hl(Lp, Rp, theta, W, dt) & Update with the one site Hamiltonian. \\
\hline Engine.update_theta_h2(Lp, Rp, theta, W0, & Update with the two sites Hamiltonian. \\
\hline W1, dt) & \\
\hline
\end{tabular}

\section*{Class Attributes and Properties}
```

Engine.TDVP_params

```

Engine.verbose
class tenpy.algorithms.tdvp.Engine (psi, model, options)
Bases: tenpy.algorithms.tdvp.TDVPEngine
Deprecated old name of TDVPEngine.
get_resume_data()
Return necessary data to resume a run () interrupted at a checkpoint.
At a checkpoint, you can save psi, model and options along with the data returned by this function. When the simulation aborts, you can resume it using this saved data with:
```

eng = AlgorithmClass(psi, model, options, resume_data=resume_data)
eng.resume_run(resume_data)

```

An algorithm which doesn't support this should override resume_run to raise an Error.
Returns resume_data - Dictionary with necessary data (apart from copies of psi, model, options) that allows to continue the simulation from where we are now.

\section*{Return type dict}
resume_run ()
Resume a run that was interrupted.
In case we saved an intermediate result at a checkpoint, this function allows to resume the run () of the algorithm (after re-initialization with the resume_data). Since most algorithms just have a while loop with break conditions, the default behaviour implemented here is to just call run ().
run()
(Real-)time evolution with TDVP.
run_one_site ( \(N\) _steps \(=\) None)
Run the TDVP algorithm with the one site algorithm.

Warning: Be aware that the bond dimension will not increase!

Parameters N_steps (integer. Number of steps)-
run_two_sites ( \(N\) _steps=None) Run the TDVP algorithm with two sites update.
The bond dimension will increase. Truncation happens at every step of the sweep, according to the parameters set in trunc_params.

Parameters N_steps (integer. Number of steps) -
set_anonymous_svd ( \(U\), new_label)
Relabel the svd.
Parameters U (tenpy.linalg.np_conserved.Array) - the tensor which lacks a leg_label
sweep_left_right()
Performs the sweep left->right of the second order TDVP scheme with one site update.
Evolve from \(0.5 * \mathrm{dt}\).
sweep_left_right_two ()
Performs the sweep left->right of the second order TDVP scheme with two sites update.
Evolve from \(0.5^{*} \mathrm{dt}\)
sweep_right_left()
Performs the sweep right->left of the second order TDVP scheme with one site update.
Evolve from \(0.5 * \mathrm{dt}\)
sweep_right_left_two()
Performs the sweep left->right of the second order TDVP scheme with two sites update.
Evolve from \(0.5 * \mathrm{dt}\)
theta_svd_left_right (theta)
Performs the SVD from left to right.
Parameters theta (tenpy.linalg.np_conserved.Array) - the theta tensor on which the SVD is applied
theta_svd_right_left (theta)
Performs the SVD from right to left.
Parameters theta (tenpy.linalg.np_conserved.Array,) - The theta tensor on which the SVD is applied
update_s_h0 \((s, H, d t)\)
Update with the zero site Hamiltonian (update of the singular value)

\section*{Parameters}
- s(tenpy.linalg.np_conserved.Array) - representing the singular value matrix which is updated
- H (HO_mixed) - zero site Hamiltonian that we need to apply on the singular value matrix
- \(\mathbf{d t}\) (complex number) - time step of the evolution
update_theta_h1 \((L p, R p\), theta, \(W, d t)\)
Update with the one site Hamiltonian.

\section*{Parameters}
- Lp (Array) - tensor representing the left environment
- \(\operatorname{Rp}\) (Array) - tensor representing the right environment
- theta (Array) - the theta tensor which needs to be updated
- W (Array) - MPO which is applied to the 'p' leg of theta
update_theta_h2 ( \(L p, R p\), theta, W0, W1, dt)
Update with the two sites Hamiltonian.

\section*{Parameters}
- Lp (tenpy.Iinalg.np_conserved.Array) - tensor representing the left environment
- Rp (tenpy.linalg.np_conserved.Array) - tensor representing the right environment
- theta (tenpy.linalg.np_conserved.Array) - the theta tensor which needs to be updated
- W (tenpy.Iinalg.np_conserved.Array) - MPO which is applied to the 'p0' leg of theta
- W1 (tenpy.linalg.np_conserved.Array) - MPO which is applied to the 'pl' leg of theta

\subsection*{17.6.2 H0_mixed}
- full name: tenpy.algorithms.tdvp.H0_mixed
- parent module: tenpy.algorithms.tdvp
- type: class

\section*{Inheritance Diagram}
H0_mixed

\section*{Methods}
\begin{tabular}{ll}
\hline H0_mixed.__init__(Lp, Rp) & Initialize self. \\
\hline H0_mixed.matvec(x) & \\
\hline
\end{tabular}
class tenpy.algorithms.tdvp. H0_mixed ( \(L p, R p\) )
Bases: object
Class defining the zero site Hamiltonian for Lanczos.

\section*{Parameters}
- Lp (tenpy.Iinalg.np_conserved.Array) - left part of the environment
- Rp (tenpy.Iinalg.np_conserved.Array) - right part of the environment

Lp
left part of the environment
Type tenpy.linalg.np_conserved.Array
Rp
right part of the environment
Type tenpy.linalg.np_conserved.Array

\subsection*{17.6.3 H1_mixed}
- full name: tenpy.algorithms.tdvp.H1_mixed
- parent module: tenpy.algorithms.tdvp
- type: class

Inheritance Diagram

H1_mixed

\section*{Methods}
\begin{tabular}{ll}
\hline H1_mixed.__init__(Lp, Rp, W) & Initialize self. \\
\hline H1_mixed.matvec(theta) &
\end{tabular}
class tenpy.algorithms.tdvp. H1_mixed ( \(L p, R p, W\) )
Bases: ob ject
Class defining the one site Hamiltonian for Lanczos.

\section*{Parameters}
- Lp (tenpy. Iinalg.np_conserved.Array) - left part of the environment
- Rp (tenpy. Iinalg.np_conserved.Array) - right part of the environment
- M(tenpy. Iinalg.np_conserved.Array) - MPO which is applied to the ' p ' leg of theta

Lp
left part of the environment
Type tenpy.linalg.np_conserved.Array
Rp
right part of the environment
Type tenpy.linalg.np_conserved.Array
w
MPO which is applied to the ' \(\mathrm{p} 0^{\prime}\) leg of theta
Type tenpy.linalg.np_conserved.Array

\subsection*{17.6.4 H2_mixed}
- full name: tenpy.algorithms.tdvp.H2_mixed
- parent module: tenpy.algorithms.tdvp
- type: class

Inheritance Diagram

H2_mixed

\section*{Methods}
\begin{tabular}{ll}
\hline H2_mixed.__init__(Lp, Rp, W0, W1) & Initialize self. \\
\hline H2_mixed.matvec(theta) &
\end{tabular}
class tenpy.algorithms.tdvp. H2_mixed ( \(L p, R p, W 0, W 1\) )
Bases: ob ject
Class defining the two sites Hamiltonian for Lanczos.

\section*{Parameters}
- Lp (tenpy.Iinalg.np_conserved.Array) - left part of the environment
- Rp (tenpy.Iinalg.np_conserved.Array) - right part of the environment
- \(\mathbf{W}\) (tenpy.linalg.np_conserved.Array) - MPO which is applied to the 'p0' leg of theta

Lp
left part of the environment
Type tenpy.linalg.np_conserved.Array
Rp
right part of the environment
Type tenpy.linalg.np_conserved.Array
wo
MPO which is applied to the 'p0' leg of theta
Type tenpy.linalg.np_conserved.Array
W1
MPO which is applied to the 'p1' leg of theta
Type tenpy.linalg.np_conserved.Array

\section*{Module description}

Time Dependant Variational Principle (TDVP) with MPS (finite version only).
The TDVP MPS algorithm was first proposed by [[haegeman2011]]. However the stability of the algorithm was later improved in [[haegeman2016]], that we are following in this implementation. The general idea of the algorithm is to project the quantum time evolution in the manyfold of MPS with a given bond dimension. Compared to e.g. TEBD, the algorithm has several advantages: e.g. it conserves the unitarity of the time evolution and the energy (for the singlesite version), and it is suitable for time evolution of Hamiltonian with arbitrary long range in the form of MPOs. We have implemented the one-site formulation which does not allow for growth of the bond dimension, and the two-site algorithm which does allow the bond dimension to grow - but requires truncation as in the TEBD case.

Todo: This is still a beta version, use with care. The interface might still change.

Todo: long-term: Much of the code is similar as in DMRG. To avoid too much duplicated code, we should have a general way to sweep through an MPS and updated one or two sites, used in both cases.

\section*{17.7 purification}
- full name: tenpy.algorithms.purification
- parent module: tenpy.algorithms
- type: module

\section*{Classes}

\begin{tabular}{ll}
\hline \begin{tabular}{l} 
PurificationApplyMPO(psi, U_MPO, options[, \\
\(\ldots])\).
\end{tabular} & \begin{tabular}{l} 
Variant of VariationalApplyMPO suitable for purifica- \\
tion.
\end{tabular} \\
\hline PurificationTEBD(psi, model, options) & \begin{tabular}{l} 
Time evolving block decimation (TEBD) for purifica- \\
tion MPS.
\end{tabular} \\
\hline PurificationTEBD2(psi, model, options) & \begin{tabular}{l} 
Similar as PurificationTEBD, but perform sweeps in- \\
stead of brickwall.
\end{tabular} \\
\hline PurificationTwoSiteU(env, i0[, combine,...]) & Variant of TwoSiteH suitable for purification. \\
\hline
\end{tabular}

\subsection*{17.7.1 PurificationApplyMPO}
- full name: tenpy.algorithms.purification.PurificationApplyMPO
- parent module: tenpy.algorithms.purification
- type: class

\section*{Inheritance Diagram}


\section*{Methods}
\begin{tabular}{|c|c|}
\hline PurificationApplyMPO.__init__(psi, U_MPO, ...) & Initialize self. \\
\hline PurificationApplyMPO. environment_sweeps(N_sweeps) & Perform \(N\) _sweeps sweeps without optimization to update the environment. \\
\hline PurificationApplyMPO. get_resume_data() & Return necessary data to resume a run () interrupted at a checkpoint. \\
\hline PurificationApplyMPO. get_sweep_schedule() & Define the schedule of the sweep. \\
\hline ```
PurificationApplyMPO.init_env(U_MPO[,
...])
``` & Initialize the environment. \\
\hline PurificationApplyMPO.make_eff_H() & Create new instance of self.EffectiveH at self.i0 and set it to self.eff_ \(H\). \\
\hline PurificationApplyMPO. post_update_local(...) & Algorithm-specific actions to be taken after local update. \\
\hline PurificationApplyMPO.prepare_update() & Prepare everything algorithm-specific to perform a local update. \\
\hline PurificationApplyMPO. reset_stats([resume_data]) & Reset the statistics. \\
\hline PurificationApplyMPO.resume_run() & Resume a run that was interrupted. \\
\hline
\end{tabular}

Table 33 - continued from previous page
\begin{tabular}{ll}
\hline PurificationApplyMPO.run() & Run the compression. \\
\hline PurificationApplyMPO.sweep([optimize]) & One 'sweep' of a sweeper algorithm. \\
\hline PurificationApplyMPO.update_LP(_) & \\
\hline PurificationApplyMPO. update_RP(_) & \\
\hline \begin{tabular}{l} 
PurificationApplyMPO.update_local(_[, \\
optimize])
\end{tabular} & Perform local update. \\
\hline \begin{tabular}{l} 
PurificationApplyMPO. \\
update_new_psi(theta)
\end{tabular} & \begin{tabular}{l} 
Given a new two-site wave function theta, split it and \\
save it in psi.
\end{tabular} \\
\hline
\end{tabular}

\section*{Class Attributes and Properties}

PurificationApplyMPO.engine_params

PurificationApplyMPO.n_optimize the number of sites to be optimized over at once.
PurificationApplyMPO.verbose
class tenpy.algorithms.purification. PurificationApplyMPO (psi, U_MPO, options, resume_data=None)
Bases: tenpy.algorithms.mps_common.VariationalApplyMPO
Variant of VariationalApplyMPO suitable for purification.

\section*{EffectiveH}
alias of tenpy.algorithms.purification.PurificationTwoSiteU
update_local (_, optimize=True)
Perform local update.
This simply contracts the environments and theta from the ket to get an updated theta for the bra self.psi (to be changed in place).
update_new_psi (theta)
Given a new two-site wave function theta, split it and save it in psi.
environment_sweeps ( \(N_{-}\)sweeps)
Perform \(N_{-}\)sweeps sweeps without optimization to update the environment.
Parameters N_sweeps (int) - Number of sweeps to run without optimization
get_resume_data()
Return necessary data to resume a run () interrupted at a checkpoint.
At a checkpoint, you can save psi, model and options along with the data returned by this function. When the simulation aborts, you can resume it using this saved data with:
```

eng = AlgorithmClass(psi, model, options, resume_data=resume_data)
eng.resume_run(resume_data)

```

An algorithm which doesn't support this should override resume_run to raise an Error.
Returns resume_data - Dictionary with necessary data (apart from copies of psi, model, options) that allows to continue the simulation from where we are now.
Return type dict
get_sweep_schedule()
Define the schedule of the sweep.
One 'sweep' is a full sequence from the leftmost site to the right and back. Only those \(L P\) and \(R P\) that can be used later should be updated.

Returns schedule - Schedule for the sweep. Each entry is (i0, move_right, (update_LP, update_RP)), where \(i 0\) is the leftmost of the self.EffectiveH. length sites to be updated in update_local (), move_right indicates whether the next \(i 0\) in the schedule is rigth (True) of the current one, and update_LP, update_RP indicate whether it is necessary to update the \(L P\) and \(R P\). The latter are chosen such that the environment is growing for infinite systems, but we only keep the minimal number of environment tensors in memory.

Return type iterable of (int, bool, (bool, bool))
init_env ( \(U_{-} M P O\), resume_data=None)
Initialize the environment.

\section*{Parameters}
- U_MPO (MPO) - The MPO to be applied to the sate.
- resume_data (dict) - May contain in
make_eff_H()
Create new instance of self.EffectiveH at self.i0 and set it to self.eff_H.
property n_optimize
the number of sites to be optimized over at once.
Indirectly set by the class attribute EffectiveH and it's length. For example, TwoSiteDMRGEngine uses the TwoSiteH and hence has n_optimize=2, while the SingleSiteDMRGEngine has n_optimize=1.
post_update_local (update_data)
Algorithm-specific actions to be taken after local update.
An example would be to collect statistics.
prepare_update()
Prepare everything algorithm-specific to perform a local update.
reset_stats (resume_data=None)
Reset the statistics. Useful if you want to start a new Sweep run.
This method is expected to be overwritten by subclass, and should then define self.update_stats and self.sweep_stats dicts consistent with the statistics generated by the algorithm particular to that subclass.
option Sweep.sweep_0: int
Number of sweeps that have already been performed.
option Sweep.chi_list: None | dict(int -> int)
By default (None) this feature is disabled. A dict allows to gradually increase the chi_max. An entry at_sweep: chi states that starting from sweep at_sweep, the value chi is to be used for trunc_params['chi_max']. For example chi_list=\{0: 50, 20: 100\} uses chi_max=50 for the first 20 sweeps and chi_max=100 afterwards.

Resume a run that was interrupted.

In case we saved an intermediate result at a checkpoint, this function allows to resume the run () of the algorithm (after re-initialization with the resume_data). Since most algorithms just have a while loop with break conditions, the default behaviour implemented here is to just call run ().
run()
Run the compression.
The state psi is compressed in place.
Returns max_trunc_err - The maximal truncation error of a two-site wave function.
Return type TruncationError
sweep (optimize \(=\) True )
One 'sweep' of a sweeper algorithm.
Iteratate over the bond which is optimized, to the right and then back to the left to the starting point. If optimize=False, don't actually diagonalize the effective hamiltonian, but only update the environment.

Parameters optimize (bool, optional) - Whether we actually optimize to find the ground state of the effective Hamiltonian. (If False, just update the environments).

Returns max_trunc_err - Maximal truncation error introduced.
Return type float

\subsection*{17.7.2 PurificationTEBD2}
- full name: tenpy.algorithms.purification.PurificationTEBD2
- parent module: tenpy.algorithms.purification
- type: class

\section*{Inheritance Diagram}


\section*{Methods}
\begin{tabular}{|c|c|}
\hline PurificationTEBD2.__init__(psi, model, options) & Initialize self. \\
\hline ```
PurificationTEBD2.calc_U(order, delta_t[,
...])
``` & see calc_U () \\
\hline PurificationTEBD2.disentangle(theta) & Disentangle theta before splitting with svd. \\
\hline \[
\begin{aligned}
& \text { PurificationTEBD2. } \\
& \text { disentangle_global([pair]) }
\end{aligned}
\] & Try global disentangling by determining the maximally entangled pairs of sites. \\
\hline \begin{tabular}{l}
PurificationTEBD2. \\
disentangle_global_nsite([n])
\end{tabular} & Perform a sweep through the system and disentangle with disentangle_n_site(). \\
\hline PurificationTEBD2. disentangle_n_site(i, n, theta) & Generalization of disentangle () to \(n\) sites. \\
\hline PurificationTEBD2.get_resume_data() & Return necessary data to resume a run () interrupted at a checkpoint. \\
\hline PurificationTEBD2.resume_run() & Resume a run that was interrupted. \\
\hline PurificationTEBD2.run() & Run TEBD real time evolution by \(N_{\text {_ }}\) steps \({ }^{*}\) 'dt. \\
\hline PurificationTEBD2.run_GS() & TEBD algorithm in imaginary time to find the ground state. \\
\hline PurificationTEBD2.run_imaginary(beta) & Run imaginary time evolution to cool down to the given beta. \\
\hline
\end{tabular}

Table 35 - continued from previous page
\begin{tabular}{|c|c|}
\hline \begin{tabular}{l}
PurificationTEBD2. \\
suzuki_trotter_decomposition(...)
\end{tabular} & Returns list of necessary steps for the suzuki trotter decomposition. \\
\hline \begin{tabular}{l}
PurificationTEBD2. \\
suzuki_trotter_time_steps(order)
\end{tabular} & Return time steps of U for the Suzuki Trotter decomposition of desired order. \\
\hline PurificationTEBD2.update(N_steps) & Evolve by N_steps * U_param['dt']. \\
\hline PurificationTEBD2.update_bond(i, U_bond) & Updates the B matrices on a given bond. \\
\hline PurificationTEBD2.update_bond_imag(i, U_bond) & Update a bond with a (possibly non-unitary) U_bond. \\
\hline PurificationTEBD2.update_imag(N_steps) & Perform an update suitable for imaginary time evolution. \\
\hline PurificationTEBD2.update_step(U_idx_dt, odd) & Updates bonds in unit cell. \\
\hline
\end{tabular}

\section*{Class Attributes and Properties}
\begin{tabular}{ll}
\hline PurificationTEBD2.TEBD_params & \\
\hline PurificationTEBD2.disent_iterations & \begin{tabular}{l} 
For each bond the total number of iterations performed \\
in any Disentangler.
\end{tabular} \\
\hline PurificationTEBD2.trunc_err_bonds & truncation error introduced on each non-trivial bond. \\
\hline PurificationTEBD2.verbose &
\end{tabular}
class tenpy.algorithms.purification. PurificationTEBD2 (psi, model, options)
Bases: tenpy.algorithms.purification.PurificationTEBD
Similar as PurificationTEBD, but perform sweeps instead of brickwall.
Instead of the A-B pattern of even/odd bonds used in TEBD, perform sweeps similar as in DMRG for real-time evolution (similar as update_imag () does for imaginary time evolution).
update (N_steps)
Evolve by N_steps * U_param['dt'].
Parameters N_steps (int) - The number of steps for which the whole lattice should be updated.

Returns trunc_err - The error of the represented state which is introduced due to the truncation during this sequence of update steps.

\section*{Return type TruncationError}
update_step ( \(U\) _idx_dt,odd)
Updates bonds in unit cell.
Depending on the choice of \(o d d\), perform a sweep to the left or right, updating once per site with a time step given by U_idx_dt.

\section*{Parameters}
- U_idx_dt (int) - Time step index in self._U, evolve with Us[i] = self. \(\mathrm{U}\left[\mathrm{U} \_i d x \_d t\right][i]\) at bond (i-1,i).
- odd (bool/int) - Indication of whether to update even (odd=False,0) or even (odd=True, 1) sites

Returns trunc_err - The error of the represented state which is introduced due to the truncation during this sequence of update steps.
Return type TruncationError
calc_U (order, delta_t, type_evo='real', E_offset=None)
see calc_U()
property disent_iterations
For each bond the total number of iterations performed in any Disentangler.

\section*{disentangle (theta)}

Disentangle theta before splitting with svd.
For the purification we write \(\rho_{P}=\operatorname{Tr}_{Q}\left|\psi_{P, Q}><\psi_{P, Q}\right|\). Thus, we can actually apply any unitary to the auxiliar \(Q\) space of \(\mid \psi>\) without changing the result.

Note: We have to apply the same unitary to the 'bra' and 'ket' used for expectation values / correlation functions!

The behaviour of this function is set by used_disentangler, which in turn is obtained from get_disentangler (options['disentangle']), see get_disentangler() for details on the syntax.

Parameters theta (Array) - Wave function to disentangle, with legs 'vL', 'vR', 'p0', 'p1', 'q0', 'q1'.

\section*{Returns}
- theta_disentangled (Array) - Disentangled theta; npc.tensordot ( U , theta, axes=[['q0*', 'q1*'], ['q0', 'q1']]).
- \(\mathbf{U}\) (Array) - The unitary used to disentangle theta, with labels ' \(q 0\) ', ' \(q 1\) ', ' \(q 0 *\) ', ' \(q 1 *\) ' . If no unitary was found/applied, it might also be None.

\section*{disentangle_global (pair=None)}

Try global disentangling by determining the maximally entangled pairs of sites.
Caclulate the mutual information (in the auxiliar space) between two sites and determine where it is maximal. Disentangle these two sites with disentangle ()

\section*{disentangle_global_nsite ( \(n=2\) )}

Perform a sweep through the system and disentangle with disentangle_n_site().
Parameters \(\mathrm{n}(i n t)\) - maximal number of sites to disentangle at once.

\section*{disentangle_n_site ( \(i, n\), theta)}

Generalization of disentangle () to \(n\) sites.
Simply group left and right \(n / 2\) physical legs, adjust labels, and apply disentangle () to disentangle the central bond. Recursively proceed to disentangle left and right parts afterwards. Scales (for even \(n\) ) as \(O\left(\chi^{3} d^{n} d^{n / 2}\right)\).
```

get_resume_data()

```

Return necessary data to resume a run () interrupted at a checkpoint.
At a checkpoint, you can save psi, model and options along with the data returned by this function. When the simulation aborts, you can resume it using this saved data with:
```

eng = AlgorithmClass(psi, model, options, resume_data=resume_data)
eng.resume_run(resume_data)

```

An algorithm which doesn't support this should override resume_run to raise an Error.
Returns resume_data - Dictionary with necessary data (apart from copies of psi, model, options) that allows to continue the simulation from where we are now.

Return type dict
resume_run ()
Resume a run that was interrupted.
In case we saved an intermediate result at a checkpoint, this function allows to resume the run () of the algorithm (after re-initialization with the resume_data). Since most algorithms just have a while loop with break conditions, the default behaviour implemented here is to just call run ().
run()
Run TEBD real time evolution by \(N_{-}\)steps \({ }^{\text {** }} d t\).
run_GS ()
TEBD algorithm in imaginary time to find the ground state.

Note: It is almost always more efficient (and hence advisable) to use DMRG. This algorithms can nonetheless be used quite well as a benchmark and for comparison.
option TEBDEngine.delta_tau_list: list
A list of floats: the timesteps to be used. Choosing a large timestep delta_tau introduces large (Trotter) errors, but a too small time step requires a lot of steps to reach exp ( - tau \(H\) ) --> \(|p s i 0><p s i 0|\). Therefore, we start with fairly large time steps for a quick time evolution until convergence, and the gradually decrease the time step.
option TEBDEngine.order: int
Order of the Suzuki-Trotter decomposition.
option TEBDEngine.N_steps: int
Number of steps before measurement can be performed
run_imaginary (beta)
Run imaginary time evolution to cool down to the given beta.
Note that we don't change the norm attribute of the MPS, i.e. normalization is preserved.
Parameters beta (float) - The inverse temperature beta \(=1 / T\), by which we should cool down. We evolve to the closest multiple of options ['dt' ], see also evolved_time.
static suzuki_trotter_decomposition (order, N_steps)
Returns list of necessary steps for the suzuki trotter decomposition.
We split the Hamiltonian as \(H=H_{\text {even }}+H_{o d d}=H[0]+H[1]\). The Suzuki-Trotter decomposition is an approximation \(\exp (t H) \approx \operatorname{prod}_{(j, k) \in S T} \exp (d[j] t H[k])+O\left(t^{\text {order }+1}\right)\).

Parameters order (1, 2, 4, '4_opt') - The desired order of the Suzuki-Trotter decomposition. Order 1 approximation is simply \(e^{A} a^{B}\). Order 2 is the "leapfrog" \(e^{\wedge}\{A / 2\} e^{\wedge} B e^{\wedge}\{A / 2\}\). Order 4 is the fourth-order from [[suzuki1991]] (also referenced in [[schollwoeck2011]]), and '4_opt' gives the optmized version of Equ. (30a) in [[barthel2020]].
Returns ST_decomposition - Indices j, k of the time-steps \(\mathrm{d}=\) suzuki_trotter_time_step (order) and the decomposition of \(H\). They are chosen such that a subsequent application of \(\exp (d[j] t H[k])\) to a given state |psi> yields \(\left(\exp \left(N \_s t e p s t H[k]\right)+O\left(N \_s t e p s t^{\wedge}\{o r d e r+1\}\right)\right) \mid p s i>\).

Return type list of (int, int)
static suzuki_trotter_time_steps (order)
Return time steps of \(U\) for the Suzuki Trotter decomposition of desired order.
See suzuki_trotter_decomposition() for details.
Parameters order (int) - The desired order of the Suzuki-Trotter decomposition.
Returns time_steps - We need \(U=\exp \left(-i \quad H \_\{e v e n / o d d\}\right.\) delta_t * dt) for the \(d t\) returned in this list.

Return type list of float
property trunc_err_bonds
truncation error introduced on each non-trivial bond.
update_bond ( \(i, U \_\)bond)
Updates the B matrices on a given bond.
Function that updates the B matrices, the bond matrix s between and the bond dimension chi for bond i.
This would look something like:
\(\square\)

\section*{Parameters}
- i (int) - Bond index; we update the matrices at sites i-1, \(i\).
- U_bond (Array) - The bond operator which we apply to the wave function. We expect labels 'p0', 'p1', 'p0*', 'p1*' for \(U \_\)bond.
Returns trunc_err - The error of the represented state which is introduced by the truncation during this update step.
Return type TruncationError
update_bond_imag ( \(i, U \_\)bond \()\)
Update a bond with a (possibly non-unitary) \(U \_b o n d\).
Similar as update_bond (); but after the SVD just keep the \(A, S, B\) canonical form. In that way, one can sweep left or right without using old singular values, thus preserving the canonical form during imaginary time evolution.

\section*{Parameters}
- i (int) - Bond index; we update the matrices at sites i-1, \(i\).
- U_bond (Array) - The bond operator which we apply to the wave function. We expect labels 'p0', 'p1', 'p0*', 'p1*'.

Returns trunc_err - The error of the represented state which is introduced by the truncation during this update step.

\section*{Return type TruncationError}
update_imag (N_steps)
Perform an update suitable for imaginary time evolution.

Instead of the even/odd brick structure used for ordinary TEBD, we 'sweep' from left to right and right to left, similar as DMRG. Thanks to that, we are actually able to preserve the canonical form.

Parameters N_steps (int) - The number of steps for which the whole lattice should be updated.
Returns trunc_err - The error of the represented state which is introduced due to the truncation during this sequence of update steps.

\section*{Return type TruncationError}

\subsection*{17.7.3 PurificationTwoSiteU}
- full name: tenpy.algorithms.purification.PurificationTwoSiteU
- parent module: tenpy.algorithms.purification
- type: class

\section*{Inheritance Diagram}


\section*{Methods}
\begin{tabular}{|c|c|}
\hline PurificationTwoSiteU.__init__(env, ...]) & Initialize self. \\
\hline PurificationTwoSiteU.adjoint() & Return the hermitian conjugate of self. \\
\hline PurificationTwoSiteU.combine_Heff() & Combine LP and RP with W to form LHeff and RHeff. \\
\hline PurificationTwoSiteU. combine_theta(theta) & Combine the legs of theta, such that it fits to how we combined the legs of self. \\
\hline
\end{tabular}

Table 37 - continued from previous page
\begin{tabular}{ll}
\hline PurificationTwoSiteU.matvec(theta) & Apply the effective Hamiltonian to theta. \\
\hline PurificationTwoSiteU.to_matrix() & Contract self to a matrix. \\
\hline
\end{tabular}

Class Attributes and Properties
```

PurificationTwoSiteU.acts_on

```
PurificationTwoSiteU.length
class tenpy.algorithms.purification.PurificationTwoSiteU(env, i0, combine=False, move_right=True)
Bases: tenpy.algorithms.mps_common.TwoSiteH
Variant of TwoSiteH suitable for purification.
The MPO gets only applied to the physical legs \(p 0, p 1\), the ancialla legs \(q 0, q 1\) of theta are ignored.
combine_Heff()
Combine LP and RP with W to form LHeff and RHeff.
Combine LP with W0 and RP with W1 to get the effective parts of the Hamiltonian with piped legs.
adjoint()
Return the hermitian conjugate of self.
combine_theta (theta)
Combine the legs of theta, such that it fits to how we combined the legs of self.
Parameters theta (Array) - Wave function with labels 'vL', 'po', 'p1', 'vR'
Returns theta - Wave function with labels 'vL', 'p0', 'p1', 'vR'
Return type Array
matvec (theta)
Apply the effective Hamiltonian to theta.
Parameters theta (Array) - Labels: vL, p0, p1, vR if combine=False, (vL.p0), ( \(\mathrm{p} 1 . \mathrm{vR}\) ) if True
Returns Product of theta and the effective Hamiltonian.
Return type theta Array
to_matrix()
Contract self to a matrix.

\section*{Module description}

Algorithms for using Purification.

\section*{17.8 mpo_evolution}
- full name: tenpy.algorithms.mpo_evolution
- parent module: tenpy.algorithms
- type: module

\section*{Classes}


\section*{Module description}

Time evolution using the WI or WII approximation of the time evolution operator.

\section*{17.9 network_contractor}
- full name: tenpy.algorithms.network_contractor
- parent module: tenpy.algorithms
- type: module

\section*{Functions}
\begin{tabular}{ll}
\hline contract(tensor_list[, tensor_names, ...]) & Contract a network of tensors. \\
\hline ncon(tensor_list, leg_links, sequence) & Implementation of \(\mathrm{ncon} . \mathrm{m}\) for TeNPy Arrays. \\
\hline
\end{tabular}

\subsection*{17.9.1 contract}
- full name: tenpy.algorithms.network_contractor.contract
- parent module: tenpy.algorithms.network_contractor
- type: function
tenpy.algorithms.network_contractor.contract (tensor_list, tensor_names=None, leg_contractions=None, open_legs=None, sequence=None)
Contract a network of tensors.
Based on the MatLab function ncon.m as described in arXiv:1402.0939.

\section*{Parameters}
- tensor_list (list of Array) - The tensors to be contracted.
- leg_contractions (list of [n1, l1, n2, 12])-A list of contraction instructions. An entry of leg_contractions has the form [n1, l1, n2, l2], where n1, n2 are entries of tensor_names and each identify an Array in tensor_list. 11, 12 are leg labels of the corresponding Array. The instruction implies to contract leg 11 of tensor n1 with leg 12 of tensor n 2 .
- open_legs (list of [n1, l1, l]) - A list of instructions for "open" (uncontracted) legs. [n1, l1, l] implies that leg 11 of tensor \(n 1\) is not contracted and is labelled 1 in the result.
- tensor_names (list of str) - A list of names for each tensor, to be used in leg_contractions and open_legs. The default value is list(range(len(tensor_list))), so that the tensor "names" are \(0,1,2, \ldots\)
- sequence (list of int) - The order in which the leg_contractions are to be performed. An entry of network_contractor.outer_product indicates performing an outer product. This corresponds to the zero-in-sequence convention of arXiv:1304.6112

Returns result - The number or tensor resulting from the contraction.
Return type Array I complex

\subsection*{17.9.2 ncon}
- full name: tenpy.algorithms.network_contractor.ncon
- parent module: tenpy.algorithms.network_contractor
- type: function
```

tenpy.algorithms.network_contractor.ncon(tensor_list,leg_links, sequence)
Implementation of ncon.m for TeNPy Arrays.
This function is a python implementation of ncon.m (arXiv:1304.6112) for tenpy Array. contract () is a wrapper that translates from a more python/tenpy input style

```

\section*{Parameters}
- tensor_list (list of :class:'Array')- Tensors to be contracted.
- leg_links (list of list of int) - Each entry of leg_links describes the connectivity of the corresponding tensor in tensor_list. Each entry is a list that has an entry for each leg of the corresponding tensor. Values \(0,1,2, \ldots\) are labels of contracted legs and should appear exactly twice in leg_links. Values \(-1,-2,-3, \ldots\) are labels of uncontracted legs and indicate the final ordering ( -1 is first axis).
- sequence (list of int) - The order in which the contractions are to be performed. An entry of network_contractor.outer_product indicates performing an outer product. This corresponds to the zero-in-sequence convention of arXiv:1304.6112

Returns result - The number or tensor resulting from the contraction.
Return type Array I complex

\section*{Module description}

\section*{Network Contractor.}

A tool to contract a network of multiple tensors.
This is an implementation of 'NCON: A tensor network contractor for MATLAB' by Robert N. C. Pfeifer, Glen Evenbly, Sukhwinder Singh, Guifre Vidal, see arXiv:1402.0939
```

tenpy.algorithms.network_contractor.outer_product = -66666666

```
a constant that represents an outer product in the sequence of ncon

\section*{Todo:}
- implement or wrap netcon.m, a function to find optimal contractionn sequences (arXiv:1304.6112)
- improve helpfulness of Warnings
- _do_trace: trace over all pairs of legs at once. need the corresponding npc function first.

\subsection*{17.10 exact_diag}
- full name: tenpy.algorithms.exact_diag
- parent module: tenpy.algorithms
- type: module

\section*{Classes}

\section*{ExactDiag}
```

ExactDiag(model[, charge_sector, sparse, ...]) (Full) exact diagonalization of the Hamiltonian.

```

\subsection*{17.10.1 ExactDiag}
- full name: tenpy.algorithms.exact_diag.ExactDiag
- parent module: tenpy.algorithms.exact_diag
- type: class

\section*{Inheritance Diagram}

\section*{ExactDiag}

\section*{Methods}
\begin{tabular}{|c|c|}
\hline ExactDiag.__init__(model[, charge_sector, ...]) & Initialize self. \\
\hline ExactDiag.build_full_H_from_bonds() & Calculate self.full_H from self.mpo. \\
\hline ExactDiag.build_full_H_from_mpo() & Calculate self.full_H from self.mpo. \\
\hline ExactDiag.exp_H(dt) & Return U (dt) : = exp (-i H dt). \\
\hline \[
\begin{aligned}
& \text { ExactDiag.from_H_mpo(H_MPO, } \\
& \text { **kwargs) }
\end{aligned}
\] & Wrapper taking directly an MPO instead of a Model. \\
\hline ExactDiag.full_diagonalization(*args, **kwargs) & Full diagonalization to obtain all eigenvalues and eigenvectors. \\
\hline ExactDiag.full_to_mps(psi[, canonical_form]) & Convert a full state (with a single leg) to an MPS. \\
\hline ExactDiag.groundstate([charge_sector]) & Pick the ground state energy and ground state from self.V. \\
\hline ExactDiag.matvec(psi) & Allow to use self as LinearOperator for lanczos. \\
\hline ExactDiag.mps_to_full(mps) & Contract an MPS along the virtual bonds and combine its legs. \\
\hline
\end{tabular}

Table 42 - continued from previous page
ExactDiag.sparse_diag(k, *args, **kwargs) Call speigs().
class tenpy.algorithms.exact_diag.ExactDiag (model, charge_sector=None, sparse=False, max_size=2000000.0)

\section*{Bases: ob ject}
(Full) exact diagonalization of the Hamiltonian.

\section*{Parameters}
- model (MPOmodel I CouplingModel) - The model which is to be diagonalized.
- charge_sector (None I charges) - If not None, project onto the given charge sector.
- sparse (bool) - If True, don't sort/bunch the LegPipe used to combine the physical legs. This results in array blocks with just one entry, requires much more charge data, and is not what np_conserved was designed for, so it's not recommended.
- max_size (int) - The build_H_* functions will do nothing (but emit a warning) if the total size of the Hamiltonian would be larger than this.
model
The model which is to be diagonalized.
Type MPOmodel I CouplingModel
chinfo
The nature of the charge (which is the same for all sites).
Type ChargeInfo
charge_sector
If not None, we project onto the given charge sector.
Type Nonel charges

\section*{max_size}

The build_H_* functions will do nothing (but emit a warning) if the total size of the Hamiltonian would be larger than this.

Type int
full_H
The full Hamiltonian to be diagonalized with legs ' \((\mathrm{p} 0 . \mathrm{p} 1 \ldots)^{\prime}, \quad\), \((\mathrm{p} 0 *, \mathrm{p} 1 * \ldots)\) ' (in that order). None if the build_H_* functions haven't been called yet, or if max_size would have been exceeded.

Type ArraylNone

E
1D array of eigenvalues.
Type ndarray I None
v
Eigenvectors. First leg 'ps' are physical legs, the second leg 'ps*' corresponds to the eigenvalues.
Type ArraylNone
_sites
The sites in the given order.
Type list of Site

\section*{_labels_p}

The labels use for the physical legs; just ['p0', 'p1', ...., 'p\{L-1\}'].
Type list or str

\section*{_labels_pconj}

Just each of _labels_p with an *.
Type list or str
_pipe
The pipe from the single physical legs to the full combined leg.
Type LegPipe
_pipe_conj
Just_pipe.conj().
Type LegPipe

\section*{_mask}

Bool mask, which of the indices of the pipe are in the desired charge_sector.
Type 1D bool ndarray I None
classmethod from_H_mpo (H_MPO, *args, **kwargs)
Wrapper taking directly an MPO instead of a Model.

\section*{Parameters}
- H_MPO (MPO) - The MPO representing the Hamiltonian.
- *args - Further keyword arguments as for the \(\qquad\) init \(\qquad\) of the class.
- **kwargs - Further keyword arguments as for the \(\qquad\) _init \(\qquad\) of the class.
build_full_H_from_mpo ()
Calculate self.full_H from self.mpo.
build_full_H_from_bonds()
Calculate self.full_H from self.mpo.
full_diagonalization(*args, **kwargs)
Full diagonalization to obtain all eigenvalues and eigenvectors.
Arguments are given to eigh.
groundstate (charge_sector=None)
Pick the ground state energy and ground state from self.V.
Parameters charge_sector (None / ID ndarray) - By default (None), consider all charge sectors. Alternatively, give the qtotal which the returned state should have.

\section*{Returns}
- E0 (float) - Ground state energy (possibly in the given sector).
- psi0 (Array) - Ground state (possibly in the given sector).
exp_H (dt)
Return U(dt) : \(=\exp (-i \quad H d t)\).
mps_to_full (mps)
Contract an MPS along the virtual bonds and combine its legs.
Parameters mps (MPS) - The MPS to be contracted.

Returns psi - The MPS contracted along the virtual bonds.
Return type Array
full_to_mps ( \(p\) si, canonical_form='B')
Convert a full state (with a single leg) to an MPS.

\section*{Parameters}
- psi (Array) - The state (with a single leg) which should be splitted into an MPS.
- canonical_from (Array) - The form in which the MPS will be afterwards.

Returns mps - An normalized MPS representation in canonical form.
Return type MPS
matvec ( \(p\) si)
Allow to use self as LinearOperator for lanczos.
Just applies full_H to (the first axis of) the given psi.
sparse_diag ( \(k\), *args, **kwargs)
Call speigs().

\section*{Module description}

Full diagonalization (ED) of the Hamiltonian.
The full diagonalization of a small system is a simple approach to test other algorithms. In case you need the full spectrum, a full diagonalization is often the only way. This module provides functionality to quickly diagonalize the Hamiltonian of a given model. This might be used to obtain the spectrum, the ground state or highly excited states.

Note: Good use of symmetries is crucial to increase the treatable system size. While we can simply use the defined LegCharge of a model, we don't make use of any other symmetries like translation symmetry, \(\mathrm{SU}(2)\) symmetry or inversion symmetries. In other words, this code does not aim to provide state-of-the-art exact diagonalization, but just the ability to diagonalize the defined models for small system sizes without addional extra work.

\section*{LINALG}
- full name: tenpy.linalg
- parent module: tenpy
- type: module

\section*{Module description}

Linear-algebra tools for tensor networks.
Most notably is the module np_conserved, which contains everything needed to make use of charge conservervation in the context of tensor networks.

Relevant contents of charges are imported to np_conserved, so you propably won't need to import charges directly.

Submodules
\begin{tabular}{ll}
\hline np_conserved & \begin{tabular}{l} 
A module to handle charge conservation in tensor net- \\
works.
\end{tabular} \\
\hline charges & Basic definitions of a charge. \\
\hline svd_robust & (More) robust version of singular value decomposition. \\
\hline random_matrix & Provide some random matrix ensembles for numpy. \\
\hline sparse & \begin{tabular}{l} 
Providing support for sparse algorithms (using matrix- \\
vector products only).
\end{tabular} \\
\hline lanczos & Lanczos algorithm for np_conserved arrays. \\
\hline
\end{tabular}

\section*{18.1 np_conserved}
- full name: tenpy.linalg.np_conserved
- parent module: tenpy. linalg
- type: module

\section*{Classes}

\section*{Array}
\begin{tabular}{ll}
\hline Array(legcharges[, dtype, qtotal, labels]) & \begin{tabular}{l} 
A multidimensional array (=tensor) for using charge \\
conservation.
\end{tabular} \\
\hline
\end{tabular}

\subsection*{18.1.1 Array}
- full name: tenpy.linalg.np_conserved.Array
- parent module: tenpy. linalg.np_conserved
- type: class

\section*{Inheritance Diagram}

\section*{Array}

\section*{Methods}
\begin{tabular}{ll}
\hline Array.__init__(legcharges[, dtype, qtotal,...]) & see help(self) \\
\hline Array.add_charge(add_legs[, chinfo, qtotal]) & Add charges. \\
\hline Array.add_leg(leg, i[, axis, label]) & \begin{tabular}{l} 
Add a leg to self, setting the current array as slice for a \\
given index.
\end{tabular} \\
\hline Array.add_trivial_leg([axis, label, qconj]) & Add a trivial leg (with just one entry) to self. \\
\hline Array.as_completely_blocked() & \begin{tabular}{l} 
Gives a version of self which is completely blocked by \\
charges.
\end{tabular} \\
\hline Array.astype(dtype[, copy]) & \begin{tabular}{l} 
Return copy with new dtype, upcasting all blocks in \\
-data.
\end{tabular} \\
\hline Array.binary_blockwise(func, other, *args, & \begin{tabular}{l} 
Roughly return func (self, other), block- \\
wise.
\end{tabular} \\
\hline Array.change_charge(charge, new_qmod[,...]) & Change the qmod of one charge in chinfo. \\
\hline Array.combine_legs(combine_legs[, new_axes, & Reshape: combine multiple legs into multiple pipes. \\
\(\ldots])\). & \\
\hline
\end{tabular}

Table 3 - continued from previous page
\begin{tabular}{|c|c|}
\hline Array.complex_conj() & Return copy which is complex conjugated without conjugating the charge data. \\
\hline Array.conj([complex_conj, inplace]) & Conjugate: complex conjugate data, conjugate charge data. \\
\hline Array.copy([deep]) & Return a (deep or shallow) copy of self. \\
\hline Array.drop_charge([charge, chinfo]) & Drop (one of) the charges. \\
\hline Array.extend(axis, extra) & Increase the dimension of a given axis, filling the values with zeros. \\
\hline Array.from_func(func, legcharges[, dtype, ...]) & Create an Array from a numpy func. \\
\hline ```
Array.from_func_square(func, leg[, dtype,
...])
``` & Create an Array from a (numpy) function. \\
\hline Array.from_hdf5(hdf5_loader, h5gr, subpath) & Load instance from a HDF5 file. \\
\hline Array.from_ndarray(data_flat, legcharges[, ...]) & convert a flat (numpy) ndarray to an Array. \\
\hline Array.from_ndarray_trivial(data_flat[,...]) & convert a flat numpy ndarray to an Array with trivial charge conservation. \\
\hline Array.gauge_total_charge(axis[, newqtotal, ...]) & Changes the total charge by adjusting the charge on a certain leg. \\
\hline Array.get_block(qindices[, insert]) & Return the ndarray in _data representing the block corresponding to qindices. \\
\hline Array.get_leg(label) &  \\
\hline Array.get_leg_index(label) & translate a leg-index or leg-label to a leg-index. \\
\hline Array.get_leg_indices(labels) & Translate a list of leg-indices or leg-labels to leg indices. \\
\hline Array.get_leg_labels() & Return list of the leg labels, with None for anonymous legs. \\
\hline Array.has_label(label) & Check whether a given label exists. \\
\hline Array.iadd_prefactor_other(prefactor, other) & self += prefactor * other for scalar prefactor and Array other. \\
\hline Array.ibinary_blockwise(func, other, *args, ...) & Roughly self = func(self, other), blockwise; in place. \\
\hline Array.iconj([complex_conj]) & Wraper around self.conj() with
inplace=True. \\
\hline Array.idrop_labels([old_labels]) & Remove leg labels from self; in place. \\
\hline Array.iproject(mask, axes) & Applying masks to one or multiple axes; in place. \\
\hline Array.ipurge_zeros([cutoff, norm_order]) & Removes self._data blocks with norm less than cutoff; in place. \\
\hline Array.ireplace_label(old_label, new_label) & Replace the leg label old_label with new_label; in place. \\
\hline Array.ireplace_labels(old_labels, new_labels) & Replace leg label old_labels[i] with new_labels[i]; in place. \\
\hline Array.is_completely_blocked() & Return bool whether all legs are blocked by charge. \\
\hline Array.iscale_axis(s[, axis]) & Scale with varying values along an axis; in place. \\
\hline Array.iscale_prefactor(prefactor) & self *= prefactor for scalar prefactor. \\
\hline Array.iset_leg_labels(labels) & Set labels for the different axes/legs; in place. \\
\hline Array.isort_qdata() & (Lexiographically) sort self._qdata; in place. \\
\hline Array.iswapaxes(axis1, axis2) & Similar as np.swapaxes; in place. \\
\hline Array.itranspose([axes]) & Transpose axes like np.transpose; in place. \\
\hline Array.iunary_blockwise(func,
**kwargs) & Roughly self \(=\mathrm{f}(\mathrm{self})\), block-wise; in place. \\
\hline Array.make_pipe(axes, **kwargs) & Generates a LegPipe for specified axes. \\
\hline Array.matvec(other) & This function is used by the Lanczos algorithm needed for DMRG. \\
\hline
\end{tabular}
continues on next page

Table 3 - continued from previous page
\begin{tabular}{ll}
\hline Array.norm([ord, convert_to_float]) & Norm of flattened data. \\
\hline Array.permute(perm, axis) & Apply a permutation in the indices of an axis. \\
\hline Array.replace_label(old_label, new_label) & \begin{tabular}{l} 
Return a shallow copy with the leg label old_label re- \\
placed by new_label.
\end{tabular} \\
\hline Array.replace_labels(old_labels, new_labels) & \begin{tabular}{l} 
Return a shallow copy with old_labels [i] re- \\
placed by new_labels [i].
\end{tabular} \\
\hline Array.save_hdf5(hdf5_saver, h5gr, subpath) & Export self into a HDF5 file. \\
\hline Array.scale_axis(s[, axis]) & Same as iscale_axis (), but return a (deep) copy. \\
\hline Array.sort_legcharge([sort, bunch]) & Return a copy with one or all legs sorted by charges. \\
\hline Array.sparse_stats() & Returns a string detailing the sparse statistics. \\
\hline Array.split_legs([axes, cutoff]) & \begin{tabular}{l} 
Reshape: opposite of combine_legs: split (some) legs \\
which are LegPipes.
\end{tabular} \\
\hline Array.squeeze([axes]) & \begin{tabular}{l} 
Remove single-dimenisional legs, \\
\\
squeeze ( ).
\end{tabular} \\
\hline Array.take_slice(indices, axes) & \begin{tabular}{l} 
Return a copy of self fixing indices along one or multi- \\
ple axes.
\end{tabular} \\
\hline Array.test_sanity() & Sanity check. \\
\hline Array.to_ndarray() & Convert self to a dense numpy ndarray. \\
\hline Array.transpose([axes]) & Like itranspose (), but on a deep copy. \\
\hline Array.unary_blockwise(func, *args, **kwargs) & Roughly return func (self), block-wise. \\
\hline Array.zeros_like() & \begin{tabular}{l} 
Return a copy of self with only zeros as entries, contain- \\
ing no _data.
\end{tabular} \\
\hline
\end{tabular}

\section*{Class Attributes and Properties}
\begin{tabular}{ll}
\hline Array.labels & \\
\hline Array.ndim & Alias for rank or len(self.shape). \\
\hline Array.size & The number of dtype-objects stored. \\
\hline Array.stored_blocks & The number of (non-zero) blocks stored in_data. \\
\hline
\end{tabular}
class tenpy.linalg.np_conserved.Array(legcharges, dtype=<class 'numpy.float64'>, qto-
Bases: ob ject
A multidimensional array (=tensor) for using charge conservation.
An Array represents a multi-dimensional tensor, together with the charge structure of its legs (for abelian charges). Further information can be found in Charge conservation with np_conserved.

The default __init__() (i.e. Array (...)) does not insert any data, and thus yields an Array 'full' of zeros, equivalent to zeros (). Further, new arrays can be created with one of from_ndarray_trivial(), from_ndarray (), or from_func (), and of course by copying/tensordot/svd etc.

In-place methods are indicated by a name starting with i. (But is_completely_blocked is not inplace...)

\section*{Parameters}
- legcharges (list of LegCharge) - The leg charges for each of the legs. The ChargeInfo is read out from it.
- dtype (type or string) - The data type of the array entries. Defaults to np.float64.
- qtotal (1D array of QTYPE) - The total charge of the array. Defaults to 0 .
- labels (list of \(\{\) str / None \}) - Labels associated to each leg, None for nonnamed labels.
rank
The rank or "number of dimensions", equivalent to len (shape).
Type int
shape
The number of indices for each of the legs.
Type tuple(int)
dtype
The data type of the entries.
Type np.dtype
chinfo
The nature of the charge.
Type ChargeInfo

\section*{qtotal}

The total charge of the tensor.
Type 1D array
legs
The leg charges for each of the legs.
Type list of LegCharge

\section*{_labels}

Labels for the different legs, None for non-labeled legs.
Type list of \{ str I None \}
_data
The actual entries of the tensor.
Type list of arrays
```

_qdata

```

For each of the _data entries the qindices of the different legs.
Type 2D array (len(_data), rank), dtype np.intp
_qdata_sorted
Whether self._qdata is lexsorted. Defaults to True, but must be set to False by algorithms changing _qdata.
Type Bool
test_sanity()
Sanity check.
Raises ValueErrors, if something is wrong.
copy (deep=True)
Return a (deep or shallow) copy of self.
Both deep and shallow copies will share chinfo and the LegCharges in legs.
In contrast to a deep copy, the shallow copy will also share the tensor entries, namely the same instances of _qdata and _data and labels (and other 'immutable' properties like the shape or dtype).

Note: Shallow copies are not recommended unless you know the consequences! See the following examples illustrating some of the pitfalls.

\section*{Examples}

Be very careful when making non-deep copies: In the following example, the original \(a\) is changed if and only if the corresponding block existed in \(a\) before.
```

>>> a = npc.Array.from_ndarray_trivial(np.arange(6.).reshape(2, 3))
>>> print(a.to_ndarray())
[[0. 1. 2.]
[3. 4. 5.]]
>>> b = a.copy(deep=False) \# shallow copy
>>> b[1, 2] = 8.
>>> a[1, 2] \# changed!
8.0

```

Other inplace operations might* have no effect at all (although we don't guarantee that):
```

>>> a *=2 \# has no effect on `b`
>>> b.iconj() \# nor does this change `a

```
save_hdf5 (hdf5_saver, h5gr, subpath)
Export self into a HDF5 file.
This method saves all the data it needs to reconstruct self with from_hdf5 ().
Specifically, it saves chinfo, legs, dtype under these names, qtotal as "total_charge", _data as "blocks", _qdata as :block_inds", the labels in the list-form (as returned by get_leg_labels()). Moreover, it saves rank, shape and _qdata_sorted (under the name "block_inds_sorted") as HDF5 attributes.

\section*{Parameters}
- hdf5_saver (Hdf5Saver) - Instance of the saving engine.
- h5gr (: class `Group ') - HDF5 group which is supposed to represent self.
- subpath (str) - The name of \(h 5 g r\) with a ' / ' in the end.
classmethod from_hdf5 (hdf5_loader, h gr, subpath)
Load instance from a HDF5 file.
This method reconstructs a class instance from the data saved with save_hdf5 ().

\section*{Parameters}
- hdf5_loader (Hdf5Loader) - Instance of the loading engine.
- h5gr (Group) - HDF5 group which is represent the object to be constructed.
- subpath (str) - The name of \(h 5 g r\) with a ' / ' in the end.

Returns obj - Newly generated class instance containing the required data.
Return type cls
classmethod from_ndarray_trivial (data_flat, dtype=None, labels=None) convert a flat numpy ndarray to an Array with trivial charge conservation.

\section*{Parameters}
- data_flat (array_like) - The data to be converted to a Array.
- dtype (np.dtype) - The data type of the array entries. Defaults to dtype of data_flat.
- labels (list of \{str / None \}) - Labels associated to each leg, None for nonnamed labels.

Returns res - An Array with data of data_flat.
Return type Array
classmethod from_ndarray (data_flat, legcharges, dtype=None, qtotal=None, cutoff=None, labels=None, raise_wrong_sector=True)
convert a flat (numpy) ndarray to an Array.

\section*{Parameters}
- data_flat (array_like) - The flat ndarray which should be converted to a npc Array. The shape has to be compatible with legcharges.
- legcharges (list of LegCharge) - The leg charges for each of the legs. The ChargeInfo is read out from it.
- dtype (np.dtype) - The data type of the array entries. Defaults to dtype of data_flat.
- qtotal (None / charges) - The total charge of the new array.
- cutoff (float) - Blocks with np.max (np.abs(block)) > cutoff are considered as zero. Defaults to QCUTOFF.
- labels (list of \{str / None\}) - Labels associated to each leg, None for nonnamed labels.
- raise_wrong_sector (bool) - If True, raise a ValueError in case of non-zero entries (larger than cutoff) in the wrong blocks of data_flat. If False, just raise a warning.
Returns res - An Array with data of data_flat.
Return type Array

\section*{See also:}
detect_qtotal used to detect qtotal if not given.
classmethod from_func (func, legcharges, dtype=None, qtotal=None, func_args=(), func_kwargs \(=\{ \}\), shape_kw=None, labels=None)
Create an Array from a numpy func.
This function creates an array and fills the blocks compatible with the charges using func, where func is a function returning a array_like when given a shape, e.g. one of np.ones or np.random. standard_normal.

\section*{Parameters}
- func (callable) - A function-like object which is called to generate the data blocks. We expect that func returns a flat array of the given shape convertible to dtype. If no shape_kw is given, it is called as func (shape, *func_args, **func_kwargs), otherwise as func(*func_args, `shape_kw`=shape, **func_kwargs). shape is a tuple of int.
- legcharges (list of LegCharge) - The leg charges for each of the legs. The ChargeInfo is read out from it.
- dtype (None / type / string) - The data type of the output entries. Defaults to np.float64. Defaults to None: obtain it from the return value of the function. Note that this argument is not given to func, but rather a type conversion is performed afterwards. You might want to set a dtype in func_kwargs as well.
- qtotal (None / charges) - The total charge of the new array. Defaults to charge 0 .
- func_args (iterable) - Additional arguments given to func.
- func_kwargs (dict) - Additional keyword arguments given to func.
- shape_kw (None / str) - If given, the keyword with which shape is given to func.
- labels (list of \{str / None \}) - Labels associated to each leg, None for nonnamed labels.

Returns res - An Array with blocks filled using func.
Return type Array
classmethod from_func_square (func, leg, dtype=None, func_args=(), func_kwargs=\{\}, shape_kw=None, labels=None)
Create an Array from a (numpy) function.
This function creates an array and fills the blocks compatible with the charges using func, where func is a function returning a array_like when given a shape, e.g. one of np . ones or np.random. standard_normal or the functions defined in random_matrix.

\section*{Parameters}
- func (callable) - A function-like object which is called to generate the data blocks. We expect that func returns a flat array of the given shape convertible to dtype. If no shape_kw is given, it is called like func (shape, *fargs, **fkwargs), otherwise as func (*fargs, `shape_kw`=shape, **fkwargs). shape is a tuple of int.
- leg (LegCharge) - The leg charges for the first leg; the second leg is set to leg. conj(). The ChargeInfo is read out from it.
- dtype (None / type / string) - The data type of the output entries. Defaults to None: obtain it from the return value of the function. Note that this argument is not given to func, but rather a type conversion is performed afterwards. You might want to set a dtype in func_kwargs as well.
- func_args (iterable) - Additional arguments given to func.
- func_kwargs (dict) - Additional keyword arguments given to func.
- shape_kw (None / str) - If given, the keyword with which shape is given to func.
- labels (list of \{str / None \})- Labels associated to each leg, None for nonnamed labels.

Returns res - An Array with blocks filled using func.
Return type Array
zeros_like()
Return a copy of self with only zeros as entries, containing no _data.
property size
The number of dtype-objects stored.
property stored_blocks
The number of (non-zero) blocks stored in _data.
property ndim
Alias for rank or len (self.shape).
get_leg_index (label)
translate a leg-index or leg-label to a leg-index.
Parameters label (int / string) - The leg-index directly or a label (string) set before.
Returns leg_index - The index of the label.
Return type int

\section*{See also:}
get_leg_indices calls get_leg_index for a list of labels.
iset_leg_labels set the labels of different legs.
get_leg_indices (labels)
Translate a list of leg-indices or leg-labels to leg indices.
Parameters labels (iterable of string/int) - The leg-labels (or directly indices) to be translated in leg-indices.

Returns leg_indices - The translated labels.
Return type list of int

\section*{See also:}
get_leg_index used to translate each of the single entries.
iset_leg_labels set the labels of different legs.
iset_leg_labels (labels)
Set labels for the different axes/legs; in place.
Introduction to leg labeling can be found in Charge conservation with np_conserved.
Parameters labels (iterable (strings / None), len=self.rank) - One label for each of the legs. An entry can be None for an anonymous leg.

\section*{See also:}
get_leg translate the labels to indices.
get_leg_labels()
Return list of the leg labels, with None for anonymous legs.
has_label (label)
Check whether a given label exists.
get_leg (label)
Return self.legs[self.get_leg_index(label)].
Convenient function returning the leg corresponding to a leg label/index.
ireplace_label (old_label, new_label)
Replace the leg label old_label with new_label; in place.
replace_label (old_label, new_label)
Return a shallow copy with the leg label old_label replaced by new_label.
ireplace_labels (old_labels, new_labels)
Replace leg labelold_labels[i] with new_labels[i]; in place.
replace_labels (old_labels, new_labels)
Return a shallow copy with old_labels[i] replaced by new_labels[i].
idrop_labels (old_labels=None)
Remove leg labels from self; in place.
Parameters old_labels (list of str/int) - The leg labels/indices for which the label should be removed. By default (None), remove all labels.
```

sparse_stats()

```

Returns a string detailing the sparse statistics.
to_ndarray ()
Convert self to a dense numpy ndarray.
get_block (qindices, insert=False)
Return the ndarray in _dat a representing the block corresponding to qindices.

\section*{Parameters}
- qindices (1D array of np. intp) - The qindices, for which we need to look in _qdata.
- insert (bool) - If True, insert a new (zero) block, if qindices is not existent in self. _data. Otherwise just return None.

Returns block - The block in _dat a corresponding to qindices. If insert \(=\) False and there is not block with qindices, return "'None`.

\section*{Return type ndarray I None}

Raises IndexError - If qindices are incompatible with charge and raise_incomp_q.
take_slice (indices, axes)
Return a copy of self fixing indices along one or multiple axes.
For a rank-4 Array A.take_slice ([i, j], [1,2]) is equivalent to \(A[:, ~ i, ~ j, ~:]\).

\section*{Parameters}
- indices ((iterable of) int) - The (flat) index for each of the legs specified by axes.
- axes ((iterable of) str/int) - Leg labels or indices to specify the legs for which the indices are given.

Returns sliced_self - A copy of self, equivalent to taking slices with indices inserted in axes.
Return type Array
See also:
add_leg opposite action of inserting a new leg.
add_trivial_leg (axis=0, label=None, qconj=1)
Add a trivial leg (with just one entry) to self.

\section*{Parameters}
- axis (int) - The new leg is inserted before index axis.
- label (str I None) - If not None, use it as label for the new leg.
- qconj \(\left(\begin{array}{lll}+1 & / & -1\end{array}\right)\) - The direction of the new leg.

Returns extended - A (possibly) shallow copy of self with an additional leg of ind_len 1 and charge 0 .
Return type Array
add_leg (leg, \(i\), axis=0, label=None)
Add a leg to self, setting the current array as slice for a given index.

\section*{Parameters}
- leg (LegCharge) - The charge data of the leg to be added.
- i (int) - Index within the leg for which the data of self should be set.
- axis (axis) - The new leg is inserted before this current axis.
- label (str I None) - If not None, use it as label for the new leg.

Returns extended - A copy of self with the new leg at axis axis, such that extended. take_slice(i, axis) returns a copy of self.

Return type Array

\section*{See also:}
take_slice opposite action reducing the number of legs.
extend (axis, extra)
Increase the dimension of a given axis, filling the values with zeros.

\section*{Parameters}
- axis (int / str) - The axis (or axis-label) to be extended.
- extra (LegCharge I int) - By what to extend, i.e. the charges to be appended to the leg of axis. An int stands for extending the length of the array by a single new block of that size with zero charges.

Returns extended - A copy of self with the specified axis increased.

\section*{Return type Array}
gauge_total_charge (axis, newqtotal=None, new_qconj=None)
Changes the total charge by adjusting the charge on a certain leg.
The total charge is given by finding a nonzero entry \([i 1, i 2, \ldots]\) and calculating:
```

qtotal = self.chinfo.make_valid(
np.sum([l.get_charge(l.get_qindex(qi) [0])
for i, l in zip([i1,i2,...], self.legs)], axis=0))

```

Thus, the total charge can be changed by redefining (= shifting) the LegCharge of a single given leg. This is exaclty what this function does.

\section*{Parameters}
- axis (int or string) - The new leg (index or label), for which the charge is changed.
- newqtotal (charge values, defaults to 0 ) - The new total charge.
- new_qconj \((\{+1,-1, N o n e\})\) - Whether the new LegCharge points inward (+1) or outward (-1) afterwards. By default (None) use the previous self.legs [leg]. qconj.
Returns copy - A shallow copy of self with copy. qtotal == newqtotal and new copy. legs [leg]. The new leg will be a :class`LegCharge`, even if the old leg was a LegPipe.

Return type Array
add_charge (add_legs, chinfo=None, qtotal=None)
Add charges.

\section*{Parameters}
- add_legs (iterable of LegCharge) - One LegCharge for each axis of self, to be added to the one in legs.
- chargeinfo (ChargeInfo) - The ChargeInfo for all charges; create new if None.
- qtotal (None | charges) - The total charge with respect to add_legs. If None, derive it from non-zero entries of self.

Returns charges_added - A copy of self, where the LegCharges add_legs where added to self.legs. Note that the LegCharges are neither bunched or sorted; you might want to use sort_legcharge().

\section*{Return type Array}
drop_charge \((\) charge \(=\) None, chinfo \(=\) None \()\)
Drop (one of) the charges.

\section*{Parameters}
- charge (int / str) - Number or name of the charge (within chinfo) which is to be dropped. None means dropping all charges.
- chinfo (ChargeInfo) - The ChargeInfo with charge dropped; create a new one if None.

Returns dropped - A copy of self, where the specified charge has been removed. Note that the LegCharges are neither bunched or sorted; you might want to use sort_legcharge ().

\section*{Return type Array}
change_charge (charge, new_qmod, new_name=", chinfo=None)
Change the qmod of one charge in chinfo.

\section*{Parameters}
- charge (int / str) - Number or name of the charge (within chinfo) which is to be changed. None means dropping all charges.
- new_qmod (int) - The new qmod to be set.
- new_name (str) - The new name of the charge.
- chinfo (ChargeInfo) - The ChargeInfo with qmod of charge changed; create a new one if None.
Returns changed - A copy of self, where the qmod of the specified charge has been changed. Note that the LegCharges are neither bunched or sorted; you might want to use sort_legcharge().

Return type Array

\section*{is_completely_blocked()}

Return bool whether all legs are blocked by charge.
sort_legcharge ( sort=True, bunch=True)
Return a copy with one or all legs sorted by charges.
Sort/bunch one or multiple of the LegCharges. Legs which are sorted and bunched are guaranteed to be blocked by charge.

\section*{Parameters}
- sort (True | False | list of \{True, False, perm\}) - A single bool holds for all legs, default=True. Else, sort should contain one entry for each leg, with a bool for sort/don't sort, or a 1D array perm for a given permuation to apply to a leg.
- bunch (True | False | list of \{True, False\})-A single bool holds for all legs, default=True. Whether or not to bunch at each leg, i.e. combine contiguous blocks with equal charges.

\section*{Returns}
- perm (tuple of \(1 D\) arrays) - The permutation applied to each of the legs, such that cp . to_ndarray() = self.to_ndarray()[np.ix_(*perm)].
- result (Array) - A shallow copy of self, with legs sorted/bunched.
```

isort_qdata()

```
(Lexiographically) sort self._qdata; in place.
Lexsort self._qdata and self._data and set self._qdata_sorted = True.
make_pipe (axes, **kwargs)
Generates a LegPipe for specified axes.

\section*{Parameters}
- axes (iterable of str/int) - The leg labels for the axes which should be combined. Order matters!
- **kwargs - Additional keyword arguments given to LegPipe.

Returns pipe - A pipe of the legs specified by axes.
Return type LegPipe
combine_legs (combine_legs, new_axes=None, pipes=None, qconj=None)
Reshape: combine multiple legs into multiple pipes. If necessary, transpose before.

\section*{Parameters}
- combine_legs ((iterable of) iterable of \{str/int\})-Bundles of leg indices or labels, which should be combined into a new output pipes. If multiple pipes should be created, use a list fore each new pipe.
- new_axes (None / (iterable of) int) - The leg-indices, at which the combined legs should appear in the resulting array. Default: for each pipe the position of its first pipe in the original array, (taking into account that some axes are 'removed' by combining). Thus no transposition is perfomed if combine_legs contains only contiguous ranges.
- pipes (None I (iterable of) \{LegPipes I None\}) - Optional: provide one or multiple of the resulting LegPipes to avoid overhead of computing new leg pipes for the same legs multiple times. The LegPipes are conjugated, if that is necessary for compatibility with the legs.
- qconj ( (iterable of) \(\{+1,-1\}\) ) - Specify whether new created pipes point inward or outward. Defaults to +1 . Ignored for given pipes, which are not newly calculated.
Returns reshaped - A copy of self, whith some legs combined into pipes as specified by the arguments.
Return type Array

\section*{See also:}
split_legs inverse reshaping splitting LegPipes.

\section*{Notes}

Labels are inherited from self. New pipe labels are generated as ' (' + '.'.join (*leglabels) + ') '. For these new labels, previously unlabeled legs are replaced by '? \#', where \# is the leg-index in the original tensor self.

\section*{Examples}
```

>>> orig_array = npc.Array.from_ndarray_trivial(np.arange(60).reshape([2, 3, ,
\hookrightarrow2, 1, 5]),
\hookrightarrow'])
>>> c1 = orig_array.combine_legs([1, 2], qconj=-1) \# only single output pipe
>>> c1.get_leg_labels()
['a', '(b.c)', 'd', 'e']
>>> c1.shape
(2, 6, 1, 5)

```

Indices of combine_legs refer to the original array. If transposing is necessary, it is performed automatically:
```

>>> c2 = orig_array.combine_legs([[0, 3], [4, 1]], qconj=[+1, -1]) \# two_
@output pipes
>>> c2.get_leg_labels()
['(a.d)', 'c', '(e.b)']
>>> c2.shape
(2, 2, 15)
>>> c3 = orig_array.combine_legs([['a', 'd'], ['e', 'b']], new_axes=[2, 1],
... pipes=[c2.legs[0], c2.legs[2]])
>>> c3.get_leg_labels()
['c', '(e.b)', '(a.d)']

```
split_legs (axes=None, cutoff=0.0)
Reshape: opposite of combine_legs: split (some) legs which are LegPipes.
Reverts combine_legs () (except a possibly performed transpose). The splited legs are replacing the LegPipes at their position, see the examples below. Labels are split reverting what was done in combine_legs (). '?\#' labels are replaced with None.

\section*{Parameters}
- axes ((iterable of) int/str) - Leg labels or indices determining the axes to split. The corresponding entries in self.legs must be LegPipe instances. Defaults to all legs, which are LegPipe instances.
- cutoff (float) - Splitted data blocks with np.max(np.abs(block)) > cutoff are considered as zero. Defaults to 0 .
Returns reshaped - A copy of self where the specified legs are splitted.
Return type Array

\section*{See also:}
combine_legs this is reversed by split_legs.

\section*{Examples}

Given a rank-5 Array orig_array, you can combine it and split it again:
```

>>> orig_array = npc.Array.from_ndarray_trivial(np.arange(60).reshape([2, 3, b
@2, 1, 5]),
... labels=['a', 'b', 'c', 'd', 'e
\hookrightarrow'])
>>> orig_array.shape
(2, 3, 2, 1, 5)
>>> comb_array = orig_array.combine_legs([['a', 'd'], ['c', 'e']] )
>>> comb_array.get_leg_labels()
['(a.d)', 'b', '(c.e)']
>>> comb_array.shape
(2, 3, 10)
>>> split_array = comb_array.split_legs()
>>> split_array.get_leg_labels()
['a', 'd', 'b', 'c', 'e']
>>> npc.norm(split_array.transpose(orig_array.get_leg_labels()) - orig_array)
0.0

```

\section*{as_completely_blocked()}

Gives a version of self which is completely blocked by charges.
Functions like svd() or eigh() require a complete blocking by charges. This can be achieved by encapsulating each leg which is not completely blocked into a LegPipe (containing only that single leg). The LegPipe will then contain all necessary information to revert the blocking.

\section*{Returns}
- encapsulated_axes (list of int) - The leg indices which have been encapsulated into Pipes.
- blocked_self (Array) - Self (if len (encapsulated_axes) = 0) or a copy of self, which is completely blocked.
squeeze (axes=None)
Remove single-dimenisional legs, like np. squeeze ().
If a squeezed leg has non-zero charge, this charge is added to qtotal.
Parameters axes (None \(/\) (iterable of) \{int/str\}) - Labels or indices of the legs which should be 'squeezed', i.e. the legs removed. The corresponding legs must be trivial, i.e., have ind_len 1.
Returns squeezed - A scalar of self.dtype, if all axes were squeezed. Else a copy of self with reduced rank as specified by axes.
Return type :class:Array I scalar
astype (dtype, copy=True)
Return copy with new dtype, upcasting all blocks in _data.

\section*{Parameters}
- dtype (convertible to a np.dtype) - The new data type. If None, deduce the new dtype as common type of self._data.
- copy ( \(b \circ \circ \mathrm{l})\) - Whether to make a copy of the blocks even if the type didn't change.

Returns copy - Deep copy of self with new dtype.

\section*{Return type Array}
ipurge_zeros (cutoff \(=2.220446049250313 e-15\), norm_order \(=\) None)
Removes self._data blocks with norm less than cutoff; in place.

\section*{Parameters}
- cutoff (float) - Blocks with norm <= cutoff are removed. defaults to QCUTOFF.
- norm_order - A valid ord argument for np.linalg.norm. Default None gives the Frobenius norm/2-norm for matrices/everything else. Note that this differs from other methods, e.g. from_ndarray (), which use the maximum norm.
iproject (mask, axes)
Applying masks to one or multiple axes; in place.
This function is similar as np.compress with boolean arrays For each specified axis, a boolean 1D array mask can be given, which chooses the indices to keep.

Warning: Although it is possible to use an 1D int array as a mask, the order is ignored! If you need to permute an axis, use permute () or sort_legcharge().

\section*{Parameters}
- mask ((list of) 1D array (bool/int)) - For each axis specified by axes a mask, which indices of the axes should be kept. If mask is a bool array, keep the indices where mask is True. If mask is an int array, keep the indices listed in the mask, ignoring the order or multiplicity.
- axes ((list of) int / string)- The \(i\) ith entry in this list specifies the axis for the ` \(i\) th entry of `mask, either as an int, or with a leg label. If axes is just a single int/string, specify just a single mask.

\section*{Returns}
- map_qind (list of \(1 D\) arrays) - The mapping of qindices for each of the specified axes.
- block_masks (list of lists of 1D bool arrays) - block_masks [a] [qind] is a boolen mask which indices to keep in block qindex of axes [a].
permute (perm, axis)
Apply a permutation in the indices of an axis.
Similar as np.take with a 1D array. Roughly equivalent to res [:, ...] = self[perm, ...] for the corresponding axis. Note: This function is quite slow, and usually not needed!

\section*{Parameters}
- perm (array_like \(1 D\) int) - The permutation which should be applied to the leg given by axis.
- axis (str / int) - A leg label or index specifying on which leg to take the permutation.
Returns res - A copy of self with leg axis permuted, such that res[i, ...] = self[perm[i], ...] for i along axis.
Return type Array

\section*{See also:}
sort_legcharge can also be used to perform a general permutation. Preferable, since it is faster for permutations which don't mix charge blocks.

\section*{itranspose (axes=None)}

Transpose axes like np.transpose; in place.
Parameters axes (iterable (intlstring), len rank I None) - The new order of the axes. By default (None), reverse axes.
transpose (axes=None)
Like itranspose (), but on a deep copy.
iswapaxes (axis1, axis2)
Similar as np.swapaxes; in place.
iscale_axis (s,axis=- l)
Scale with varying values along an axis; in place.
Rescale to new_self[i1, ..., i_axis, ...] = s[i_axis] * self[i1, ..., i_axis, ...].

\section*{Parameters}
- s(1D array, len=self.shape[axis]) - The vector with which the axis should be scaled.
- axis (str/int) - The leg label or index for the axis which should be scaled.

\section*{See also:}
iproject can be used to discard indices for which \(s\) is zero.
scale_axis \((s, a x i s=-1)\)
Same as iscale_axis(), but return a (deep) copy.
iunary_blockwise (func, *args, **kwargs)
Roughly self \(=f(s e l f)\), block-wise; in place.
Applies an unary function func to the non-zero blocks in self._data.

Note: Assumes implicitly that func(np.zeros(...), *args, **kwargs) gives 0, since we don't let func act on zero blocks!

\section*{Parameters}
- func (function) - A function acting on flat arrays, returning flat arrays. It is called like new_block \(=\) func (block, *args, **kwargs).
- *args - Additional arguments given to function after the block.
- **kwargs - Keyword arguments given to the function.

\section*{Examples}
```

>>> a = npc.Array.from_ndarray_trivial([1., 2.j])
>>> a.iunary_blockwise(np.conj).to_ndarray() \# same data as a.iconj(), but,
@oesn't charge conjugate.
array([1.-0.j, 0.-2.j])
>>> a.iunary_blockwise(np.real).to_ndarray() \# get real part
array([1., 0.])

```
unary_blockwise (func, *args, **kwargs)
Roughly return func (self), block-wise. Copies.
Same as iunary_blockwise (), but makes a shallow copy first.
iconj(complex_conj=True)
Wraper around self.conj() with inplace=True.
conj (complex_conj=True, inplace=False)
Conjugate: complex conjugate data, conjugate charge data.
Conjugate all legs, set negative qtotal.
Labeling: takes 'a' -> 'a*', 'a*'-> 'a' and ' \(\left(a,\left(b^{*}, c\right)\right)^{\prime}->^{\prime}\left(a^{*},\left(b, c^{*}\right)\right)^{\prime}\)

\section*{Parameters}
- complex_conj (bool) - Whether the data should be complex conjugated.
- inplace (bool) - Whether to apply changes to self, or to return a deep copy.
```

complex_conj()

```

Return copy which is complex conjugated without conjugating the charge data.
norm (ord=None, convert_to_float=True)
Norm of flattened data.
See norm () for details.
ibinary_blockwise (func, other, *args, **kwargs)
Roughly self = func (self, other), block-wise; in place.
Applies a binary function 'block-wise' to the non-zero blocks of self._data and other._data, storing result in place. Assumes that other is an Array as well, with the same shape and compatible legs. If leg labels of other and self are same up to permutations, other gets transposed accordingly before the action.

Note: Assumes implicitly that func(np.zeros(...), np.zeros(...), *args, \(* *\) kwargs ) gives 0 , since we don't let func act on zero blocks!

\section*{Parameters}
- func (function) - Binary function, called as new_block = func (block_self, block_other, *args, **kwargs) for blocks (=Numpy arrays) of equal shape.
- other (Array) - Other Array from which to take blocks. Should have the same leg structure as self.
- *args - Extra arguments given to func.
- **kwargs - Extra keyword arguments given to func.

\section*{Examples}
```

>>> a = npc.Array.from_ndarray_trivial([1., 3.])
>>> b = npc.Array.from_ndarray_trivial([4., 2.])
>>> a.ibinary_blockwise(np.maximum, b).to_ndarray() \# a = max(a, b)
array([4., 3.])
>>> a.ibinary_blockwise(np.add, b).to_ndarray() \# roughly `` a += b`
array([8., 5.])

```
binary_blockwise (func, other, *args, **kwargs)
Roughly return func (self, other), block-wise. Copies.
Same as ibinary_blockwise (), but makes a shallow copy first.
matvec (other)
This function is used by the Lanczos algorithm needed for DMRG.
It is supposed to calculate the matrix - vector - product for a rank-2 matrix self and a rank-1 vector other.
iadd_prefactor_other (prefactor, other)
self += prefactor * other for scalar prefactor and Array other.
Note that we allow the type of self to change if necessary. Moreover, if self and other have the same labels in different order, other gets transposed before the action.
iscale_prefactor (prefactor)
self \(\star=\) prefactor for scalar prefactor.
Note that we allow the type of self to change if necessary.

\section*{Functions}
\begin{tabular}{ll}
\hline concatenate(arrays[, axis, copy]) & \begin{tabular}{l} 
Stack arrays along a given axis, similar as \\
np.concatenate.
\end{tabular} \\
\hline \begin{tabular}{l} 
detect_grid_outer_legcharge(grid, \\
grid_legs)
\end{tabular} & \begin{tabular}{l} 
Derive a LegCharge for a grid used for \\
grid_outer ().
\end{tabular} \\
\hline detect_legcharge(flat_array, chargeinfo, ...) & \begin{tabular}{l} 
Calculate a missing LegCharge by looking for nonzero \\
entries of a flat array.
\end{tabular} \\
\hline detect_qtotal(flat_array, legcharges[, cutoff]) & \begin{tabular}{l} 
Returns the total charge (w.r.t legs) of first non-zero sec- \\
tor found in flat_array.
\end{tabular} \\
\hline diag(s, leg[, dtype, labels]) & Returns a square, diagonal matrix of entries \(s\). \\
\hline eig(a[, sort]) & \begin{tabular}{l} 
Calculate eigenvalues and eigenvectors for a non- \\
hermitian matrix.
\end{tabular} \\
\hline eigh(a[, UPLO, sort]) & \begin{tabular}{l} 
Calculate eigenvalues and eigenvectors for a hermitian \\
matrix.
\end{tabular} \\
\hline eigvals(a[, sort]) & Calculate eigenvalues for a hermitian matrix. \\
\hline eigvalsh(a[, UPLO, sort]) & Calculate eigenvalues for a hermitian matrix. \\
\hline expm(a) & \begin{tabular}{l} 
Use scipy.linalg.expm to calculate the matrix exponen- \\
tial of a square matrix.
\end{tabular} \\
\hline eye_like(a[, axis, labels]) & \begin{tabular}{l} 
Return an identity matrix contractible with the leg axis \\
of the Array.
\end{tabular} \\
\hline
\end{tabular}

Table 5 - continued from previous page
\begin{tabular}{|c|c|}
\hline grid_concat(grid, axes[, copy]) & Given an np.array of npc.Arrays, performs a multidimensional concatentation along 'axes'. \\
\hline grid_outer(grid, grid_legs[, qtotal, ...]) & Given an np.array of npc.Arrays, return the corresponding higher-dimensional Array. \\
\hline inner(a, b[, axes, do_conj]) & Contract all legs in \(a\) and \(b\), return scalar. \\
\hline norm(a[, ord, convert_to_float]) & Norm of flattened data. \\
\hline ones(legcharges[, dtype, qtotal, labels]) & Short-hand for Array. from_func () with function numpy.ones (). \\
\hline outer ( \(\mathrm{a}, \mathrm{b}\) ) & Forms the outer tensor product, equivalent to tensordot (a, b, axes=0). \\
\hline \(\operatorname{pinv}(\mathrm{a}\) [, cutoff]) & Compute the (Moore-Penrose) pseudo-inverse of a matrix. \\
\hline qr(a[, mode, inner_labels, cutoff]) & Q-R decomposition of a matrix. \\
\hline speigs(a, charge_sector, k, *args, **kwargs) & Sparse eigenvalue decomposition \(\mathrm{w}, \mathrm{v}\) of square \(a\) in a given charge sector. \\
\hline \(\operatorname{svd}(\mathrm{a}\) [, full_matrices, compute_uv, cutoff, ...]) & Singualar value decomposition of an Array \(a\). \\
\hline tensordot(a, b[, axes]) & Similar as np.tensordot but for Array. \\
\hline to_iterable_arrays(array_list) & Similar as to_iterable (), but also enclose npc Arrays in a list. \\
\hline trace(a[, leg1, leg2]) & Trace of \(a\), summing over leg 1 and leg2. \\
\hline zeros(legcharges[, dtype, qtotal, labels]) & Create a npc array full of zeros (with no _data). \\
\hline
\end{tabular}

\subsection*{18.1.2 concatenate}
- full name: tenpy.linalg.np_conserved.concatenate
- parent module: tenpy.Iinalg.np_conserved
- type: function
tenpy.linalg.np_conserved.concatenate (arrays, axis=0, copy=True)
Stack arrays along a given axis, similar as np.concatenate.
Stacks the qind of the array, without sorting/blocking. Labels are inherited from the first array only.

\section*{Parameters}
- arrays (iterable of Array) - The arrays to be stacked. They must have the same shape and charge data except on the specified axis.
- axis (int | str) - Leg index or label of the first array. Defines the axis along which the arrays are stacked.
- copy (bool) - Whether to copy the data blocks.

Returns stacked - Concatenation of the given arrays along the specified axis.
Return type Array

\section*{See also:}

Array.sort_legcharge can be used to block by charges along the axis.

\subsection*{18.1.3 detect_grid_outer_legcharge}
- full name: tenpy.linalg.np_conserved.detect_grid_outer_legcharge
- parent module: tenpy.Iinalg.np_conserved
- type: function
tenpy.linalg.np_conserved.detect_grid_outer_legcharge (grid, grid_legs, qtotal=None, qconj=1, bunch=False )
Derive a LegCharge for a grid used for grid_outer ().
Note: The resulting LegCharge is not bunched.

\section*{Parameters}
- grid (array_like of \{Array INone\}) - The grid as it will be given to grid_outer ().
- grid_legs (list of \{LegCharge I None\}) - One LegCharge for each dimension of the grid, except for one entry which is None. This missing entry is to be calculated.
- qtotal (charge) - The desired total charge of the array. Defaults to 0 .

Returns new_grid_legs - A copy of the given grid_legs with the None replaced by a compatible LegCharge. The new LegCharge is neither bunched nor sorted!
Return type list of LegCharge

\section*{See also:}
detect_legcharge similar functionality for a flat numpy array instead of a grid.

\subsection*{18.1.4 detect_legcharge}
- full name: tenpy.linalg.np_conserved.detect_legcharge
- parent module: tenpy.Iinalg.np_conserved
- type: function
tenpy.linalg.np_conserved.detect_legcharge (flat_array, chargeinfo, legcharges, qtotal \(=\) None, qconj=1, cutoff=None)
Calculate a missing LegCharge by looking for nonzero entries of a flat array.

\section*{Parameters}
- flat_array (ndarray) - A flat array, in which we look for non-zero entries.
- chargeinfo (ChargeInfo) - The nature of the charge.
- legcharges (list of LegCharge) - One LegCharge for each dimension of flat_array, except for one entry which is None. This missing entry is to be calculated.
- qconj \((\{+1,-1\})\) - qconj for the new calculated LegCharge.
- qtotal (charges) - Desired total charge of the array. Defaults to zeros.
- cutoff (float) - Blocks with np.max(np.abs(block)) > cutoff are considered as zero. Defaults to QCUTOFF.
Returns new_legcharges - A copy of the given legcharges with the None replaced by a compatible LegCharge. The new legcharge is 'bunched', but not sorted!

Return type list of LegCharge

\section*{See also:}
detect_grid_outer_legcharge similar functionality if the flat array is given by a 'grid'.
detect_qtotal detects the total charge, if all legs are known.

\subsection*{18.1.5 detect_qtotal}
- full name: tenpy.linalg.np_conserved.detect_qtotal
- parent module: tenpy.linalg.np_conserved
- type: function
tenpy.linalg.np_conserved.detect_qtotal (flat_array, legcharges, cutoff=None)
Returns the total charge (w.r.t legs) of first non-zero sector found in flat_array.

\section*{Parameters}
- flat_array (array) - The flat numpy array from which you want to detect the charges.
- legcharges (list of LegCharge) - For each leg the LegCharge.
- cutoff (float) - Blocks with np.max(np.abs(block)) > cutoff are considered as zero. Defaults to QCUTOFF.

Returns qtotal - The total charge fo the first non-zero (i.e. > cutoff) charge block.
Return type charge
See also:
detect_legcharge detects the charges of one missing LegCharge if qtotal is known.
detect_grid_outer_legcharge similar functionality if the flat array is given by a 'grid'.

\subsection*{18.1.6 diag}
- full name: tenpy.linalg.np_conserved.diag
- parent module: tenpy. linalg.np_conserved
- type: function
```

tenpy.linalg.np_conserved.diag(s,leg,dtype=None,labels=None)

```

Returns a square, diagonal matrix of entries \(s\).
The resulting matrix has legs (leg, leg.conj()) and charge 0.

\section*{Parameters}
- s (scalar / ID array) - The entries to put on the diagonal. If scalar, all diagonal entries are the same.
- leg (LegCharge) - The first leg of the resulting matrix.
- dtype (None / type) - The data type to be used for the result. By default, use dtype of \(s\).
- labels (list of \(\{\) str / None \}) - Labels associated to each leg, None for nonnamed labels.

Returns diagonal - A square matrix with diagonal entries \(s\).

Return type Array

\section*{See also:}

Array.scale_axis similar as tensordot (diag(s), ...), but faster.

\subsection*{18.1.7 eig}
- full name: tenpy.linalg.np_conserved.eig
- parent module: tenpy.Iinalg.np_conserved
- type: function
tenpy.linalg.np_conserved.eig ( \(a\), sort=None)
Calculate eigenvalues and eigenvectors for a non-hermitian matrix.
\(\mathrm{W}, \mathrm{V}=\operatorname{eig}(\mathrm{a})\) yields \(a V=V \operatorname{diag}(w)\).

\section*{Parameters}
- a (Array) - The hermitian square matrix to be diagonalized.
- sort ( \(\{\) 'm>', 'm<', '>', '<', None \}) - How the eigenvalues should are sorted within each charge block. Defaults to None, which is same as ' \(<\) '. See argsort () for details.

\section*{Returns}
- W ( \(1 D\) ndarray) - The eigenvalues, sorted within the same charge blocks according to sort.
- V (Array) - Unitary matrix; \(\mathrm{V}[:, ~ i]\) is normalized eigenvector with eigenvalue \(\mathrm{W}[\mathrm{i}]\). The first label is inherited from \(A\), the second label is 'eig'.

\section*{Notes}

Requires the legs to be contractible. If \(a\) is not blocked by charge, a blocked copy is made via a permutation P , \(a^{\prime}=P a P^{-1}=V^{\prime} W^{\prime}\left(V^{\prime}\right)^{\dagger}\). The eigenvectors \(V\) are then obtained by the reverse permutation, \(V=P^{-1} V^{\prime}\) such that \(a=V W V^{\dagger}\).

\subsection*{18.1.8 eigh}
- full name: tenpy.linalg.np_conserved.eigh
- parent module: tenpy. Iinalg.np_conserved
- type: function
```

tenpy.linalg.np_conserved.eigh(a,UPLO='L', sort=None)

```

Calculate eigenvalues and eigenvectors for a hermitian matrix.
\(\mathrm{W}, \mathrm{V}=\operatorname{eigh}(\mathrm{a})\) yields \(a=V \operatorname{diag}(w) V^{\dagger}\). Assumes that a is hermitian, a.conj().transpose() == a.

\section*{Parameters}
- a (Array) - The hermitian square matrix to be diagonalized.
- UPLO ( \(\left\{\right.\) ' \(\left.L^{\prime}, '^{\prime} U^{\prime}\right\}\) ) - Whether to take the lower ('L', default) or upper ('U') triangular part of \(a\).
- sort ( \(\{\) 'm>', 'm<', '>', '<', None \}) - How the eigenvalues should are sorted within each charge block. Defaults to None, which is same as '<'. See argsort () for details.

\section*{Returns}
- W (1D ndarray) - The eigenvalues, sorted within the same charge blocks according to sort.
- V (Array) - Unitary matrix; \(\mathrm{V}[:, ~ i]\) is normalized eigenvector with eigenvalue \(\mathrm{W}[\mathrm{i}]\). The first label is inherited from \(A\), the second label is 'eig'.

\section*{Notes}

Requires the legs to be contractible. If \(a\) is not blocked by charge, a blocked copy is made via a permutation P , \(a^{\prime}=P a P^{-1}=V^{\prime} W^{\prime}\left(V^{\prime}\right)^{\dagger}\). The eigenvectors \(V\) are then obtained by the reverse permutation, \(V=P^{-1} V^{\prime}\) such that \(a=V W V^{\dagger}\).

\subsection*{18.1.9 eigvals}
- full name: tenpy.linalg.np_conserved.eigvals
- parent module: tenpy. linalg.np_conserved
- type: function
```

tenpy.linalg.np_conserved.eigvals(a, sort=None)

```

Calculate eigenvalues for a hermitian matrix.

\section*{Parameters}
- a (Array) - The hermitian square matrix to be diagonalized.
- sort ( \(\{\) ' \(m>\) ', ' \(m<\) ', ' \(>\) ', '<', None \(\}\) ) - How the eigenvalues should are sorted within each charge block. Defaults to None, which is same as ' \(<\) '. See argsort () for details.
Returns W - The eigenvalues, sorted within the same charge blocks according to sort.
Return type 1D ndarray

\section*{Notes}

The eigenvalues are sorted within blocks of the completely blocked legs.

\subsection*{18.1.10 eigvalsh}
- full name: tenpy.linalg.np_conserved.eigvalsh
- parent module: tenpy. linalg.np_conserved
- type: function
```

tenpy.linalg.np_conserved.eigvalsh(a,UPLO='L', sort=None)

```

Calculate eigenvalues for a hermitian matrix.
Assumes that a is hermitian, \(a \cdot\) conj().transpose() \(==a\).
Parameters
- a (Array) - The hermitian square matrix to be diagonalized.
- UPLO ( \(\left\{\right.\) ' \(L\) ', ' \(\left.U^{\prime}\right\}\) ) - Whether to take the lower ('L', default) or upper ('U') triangular part of \(a\).
- sort ( \(\{\) 'm>', 'm<', '>', '<', None\}) - How the eigenvalues should are sorted within each charge block. Defaults to None, which is same as ' \(<\) '. See argsort () for details.
Returns W - The eigenvalues, sorted within the same charge blocks according to sort.
Return type 1D ndarray

\section*{Notes}

The eigenvalues are sorted within blocks of the completely blocked legs.

\subsection*{18.1.11 expm}
- full name: tenpy.linalg.np_conserved.expm
- parent module: tenpy.linalg.np_conserved
- type: function
tenpy.linalg.np_conserved.expm (a)
Use scipy.linalg.expm to calculate the matrix exponential of a square matrix.
Parameters a (Array) - A square matrix to be exponentiated.
Returns exp_a - The matrix exponential expm (a), calculated using scipy.linalg.expm. Same legs/labels as \(a\).
Return type Array

\subsection*{18.1.12 eye_like}
- full name: tenpy.linalg.np_conserved.eye_like
- parent module: tenpy.Iinalg.np_conserved
- type: function
tenpy.linalg.np_conserved.eye_like (a, axis=0, labels=None)
Return an identity matrix contractible with the leg axis of the Array \(a\).

\subsection*{18.1.13 grid_concat}
- full name: tenpy.linalg.np_conserved.grid_concat
- parent module: tenpy.Iinalg.np_conserved
- type: function
tenpy.linalg.np_conserved.grid_concat (grid, axes, copy=True)
Given an np.array of npc.Arrays, performs a multi-dimensional concatentation along 'axes'.
Similar to numpy.block (), but only for uniform blocking.
Stacks the qind of the array, without sorting/blocking.

\section*{Parameters}
- grid (array_like of Array) - The grid of arrays.
- axes (list of int) - The axes along which to concatenate the arrays, same len as the dimension of the grid. Concatenate arrays of the \(i\) 'th axis of the grid along the axis "axes \(\left[i{ }^{\prime}\right.\)
- copy (bool) - Whether the _data blocks are copied.

\section*{Examples}

Assume we have prepared rank 2 Arrays A, B, C, D sharing the legs of equal sizes and looking like this:
```

>>> print(A.to_ndarray())
[[0 1]]
>>> print(B.to_ndarray())
[[[10 11 12 13]]
>>> print(C.to_ndarray())
[[$$
\begin{array}{lll}{20}&{21]}\end{array}
$$]
[22 23]
[24 25]]
>>> print(D.to_ndarray())
[[[30 31 32 33]
[34 35 36 37]
[38 39 40 41]]

```

Then the following grid will result in a \((1+3,2+4)\) shaped array:
```

>>> g = npc.grid_concat([[A, B],
···. [C, D]], axes=[0, 1])
>>> g.shape
(4, 6)
>>> print(g.to_ndarray())
[[[ 0 1 10 11 12 13]
[20 21 30 31 32 33]
[22 23 34 35 36 37]
[[24 25 38 39 40 41]]

```

If A, B, C, D were rank 4 arrays, with the first and last leg as before, and sharing common legs 1 and 2 of dimensions 1,2 , then you would get a rank-4 array:
```

>>> g = grid_concat([[A, B], [C, D]], axes=[0, 3])
>>> g.shape
(4, 1, 2, 6)

```

\section*{See also:}

Array.sort_legcharge can be used to block by charges.

\subsection*{18.1.14 grid_outer}
- full name: tenpy.linalg.np_conserved.grid_outer
- parent module: tenpy.linalg.np_conserved
- type: function
tenpy.linalg.np_conserved.grid_outer (grid, grid_legs, qtotal=None, grid_labels=None) Given an np.array of npc.Arrays, return the corresponding higher-dimensional Array.

\section*{Parameters}
- grid (array_like of \(\{\) Array \(\mid\) None \(\}\) ) - The grid gives the first part of the axes of the resulting array. Entries have to have all the same shape and charge-data, giving the remaining axes. None entries in the grid are interpreted as zeros.
- grid_legs (list of LegCharge) - One LegCharge for each dimension of the grid along the grid.
- qtotal (charge) - The total charge of the Array. By default (None), derive it out from a non-trivial entry of the grid.
- grid_labels (list of \(\{s t r \mid N o n e\}\) ) - One label associated to each of the grid axes. None for non-named labels.

Returns res - An Array with shape grid.shape + nontrivial_grid_entry.shape. Constructed such that res[idx] == grid[idx] for any index idx of the grid the grid entry is not trivial (None).

\section*{Return type Array}

\section*{See also:}
detect_grid_outer_legcharge can calculate one missing LegCharge of the grid.

\section*{Examples}

A typical use-case for this function is the generation of an MPO. Say you have npc.Arrays Splus, Sminus, Sz, Id, each with legs [phys.conj(), phys]. Further, you have to define appropriate LegCharges l_left and \(l_{-}\)right. Then one 'matrix' of the MPO for a nearest neighbour Heisenberg Hamiltonian could look like:
```

>>> s = tenpy.networks.site.SpinHalfSite(conserve='Sz')
>>> Id, Splus, Sminus, Sz = s.Id, s.Sp, s.Sm, s.Sz
>>> J = 1.
>>> leg_wR = npc.LegCharge.from_qflat(s.leg.chinfo,
... [op.qtotal for op in [Id, Splus, Sminus, Sz,
\hookrightarrow Id]],
\cdots.. qconj=-1)
>>> W_mpo = npc.grid_outer([[Id, Splus, Sminus, Sz, None],
... [None, None, None, None, J*0.5*Sminus],
... [None, None, None, None, J*0.5*Splus],
... [None, None, None, None, J*Sz],
... [None, None, None, None, Id]],
... grid_legs=[leg_wR.conj(), leg_wR],
... grid_labels=['wL', 'wR'])
>>> W_mpo.shape
(5, 5, 2, 2)
>>> W_mpo.get_leg_labels()
['wL', 'wR', 'p', 'p*']

```

\subsection*{18.1.15 inner}
- full name: tenpy.linalg.np_conserved.inner
- parent module: tenpy.Iinalg.np_conserved
- type: function
tenpy.linalg.np_conserved.inner ( \(a, b\), axes=None, do_conj=False)
Contract all legs in \(a\) and \(b\), return scalar.

\section*{Parameters}
- a (class:Array) - The arrays for which to calculate the product. Must have same rank, and compatible LegCharges.
- \(\mathbf{b}\) (class:Array) - The arrays for which to calculate the product. Must have same rank, and compatible LegCharges.
- axes ((axes_a, axes_b) | 'range', 'labels') - axes_a and axes_b specifiy the legs of \(a\) and \(b\), respectively, which should be contracted. Legs can be specified with leg labels or indices. We contract leg axes_a[i] of \(a\) with leg axes_b[i] of \(b\). The default axes='range' is equivalent to (range (rank), range (rank)). axes='labels' is equivalent to either (a.get_leg_labels(), a.get_leg_labels()) for do_conj=True, or to (a.get_leg_labels(), conj_labels(a.get_leg_labels())) for do_conj=False. In other words, axes='labels' requires \(a\) and \(b\) to have the same/conjugated labels up to a possible transposition, which is then reverted.
- do_conj (bool) - If False (Default), ignore it. If True, conjugate \(a\) before, i.e., return inner (a.conj(), b, axes).

Returns inner_product - A scalar (of common dtype of \(a\) and \(b\) ) giving the full contraction of \(a\) and \(b\).
Return type dtype

\subsection*{18.1.16 norm}
- full name: tenpy.linalg.np_conserved.norm
- parent module: tenpy.Iinalg.np_conserved
- type: function
tenpy.linalg.np_conserved.norm(a, ord=None, convert_to_float=True)
Norm of flattened data.
Equivalent to np.linalg.norm(a.to_ndarray().flatten(), ord).
In contrast to numpy, we don't distinguish between matrices and vectors, but simply calculate the norm for the flat (block) data. The usual ord-norm is defined as \(\left(\sum_{i}\left|a_{i}\right|^{\text {ord }}\right)^{1 / \text { ord }}\).
\begin{tabular}{|l|l|}
\hline Ord & norm \\
\hline None/'fro' & Frobenius norm (same as 2-norm) \\
\hline np.inf & \(\max (\operatorname{abs}(\mathrm{x}))\) \\
\hline -np.inf & \(\min (\operatorname{abs}(\mathrm{x}))\) \\
\hline 0 & sum \((\mathrm{a} \quad!=0)==\) np. count_nonzero \((\mathrm{x})\) \\
\hline other & ususal ord-norm \\
\hline
\end{tabular}

\section*{Parameters}
- a (Array I np.ndarray) - The array of which the norm should be calculated.
- ord - The order of the norm. See table above.
- convert_to_float - Convert integer to float before calculating the norm, avoiding int overflow.

Returns norm - The norm over the flat data of the array.
Return type float

\subsection*{18.1.17 ones}
- full name: tenpy.linalg.np_conserved.ones
- parent module: tenpy.linalg.np_conserved
- type: function
tenpy.linalg.np_conserved.ones(legcharges, dtype=<class 'numpy.float64'>, qtotal=None, labels=None)
Short-hand for Array.from_func() with function numpy.ones ().

Warning: For non-trivial charges, only blocks with compatible charges are filled with ones!

\subsection*{18.1.18 outer}
- full name: tenpy.linalg.np_conserved.outer
- parent module: tenpy.Iinalg.np_conserved
- type: function
tenpy.linalg.np_conserved.outer ( \(a, b\) )
Forms the outer tensor product, equivalent to tensordot (a, b, axes=0).
Labels are inherited from \(a\) and \(b\). In case of a collision (same label in both \(a\) and \(b\) ), they are both dropped.

\section*{Parameters}
- a (Array) - The arrays for which to form the product.
- \(\mathbf{b}\) (Array) - The arrays for which to form the product.

\section*{Returns}
c -
Array of rank a.rank + b.rank such that (for \(\mathrm{Ra}=\mathrm{a} \cdot \mathrm{rank} ; \mathrm{Rb}=\mathrm{b} \cdot \mathrm{rank}\) ):
```

c[i_1, ..., i_Ra, j_1, ... j_R] = a[i_1, ..., i_Ra] * b[j_1, ..., j_
\hookrightarrowrank_b]

```

\section*{Return type Array}

\subsection*{18.1.19 pinv}
- full name: tenpy.linalg.np_conserved.pinv
- parent module: tenpy.linalg.np_conserved
- type: function
tenpy.linalg.np_conserved.pinv ( \(a\), cutoff=le-15)
Compute the (Moore-Penrose) pseudo-inverse of a matrix.
Equivalent to the following procedure: Perform a SVD, U, S, VH \(=\operatorname{svd}(a\), cutoff=cutoff) with a cutoff \(>0\), calculate \(\mathrm{P}=\mathrm{U} * \operatorname{diag}(1 / \mathrm{S}) * \mathrm{VH}\) (with * denoting tensordot) and return P. conj. transpose().

\section*{Parameters}
- a ((M, N) Array) - Matrix to be pseudo-inverted.
- cuttof (float) - Cutoff for small singular values, as given to svd (). (Note: different convetion than numpy.)
Returns B - The pseudo-inverse of \(a\).
Return type (N, M) Array

\subsection*{18.1.20 qr}
- full name: tenpy.linalg.np_conserved.qr
- parent module: tenpy.linalg.np_conserved
- type: function
```

tenpy.linalg.np_conserved.qr (a,mode='reduced', inner_labels=[None, None], cutoff=None)

```

Q-R decomposition of a matrix.
Decomposition such that \(A==n p c . t e n s o r d o t(q, r\), axes=1) up to numerical rounding errors.

\section*{Parameters}
- a (Array) - A square matrix to be exponentiated, shape (M,N).
- mode ('reduced', 'complete') - 'reduced': return \(q\) and \(r\) with shapes ( \(\mathrm{M}, \mathrm{K}\) ) and \((\mathrm{K}, \mathrm{N})\), where \(\mathrm{K}=\min (\mathrm{M}, \mathrm{N})\) 'complete': return \(q\) with shape (M,M).
- inner_labels ([\{str/None\}, \{str/None\}]) - The first label is used for \(Q\). legs [1], the second for R.legs [0].
- cutoff (None or float) - If not None, discard linearly dependent vectors to given precision, which might reduce \(K\) of the 'reduced' mode even further.

\section*{Returns}
- \(\mathbf{q}\) (Array) - If mode is 'complete', a unitary matrix. For mode 'reduced' such thatOtherwise such that \(q_{j, i}^{*} q_{j, k}=\delta_{i, k}\)
- r (Array) - Upper triangular matrix if both legs of A are sorted by charges; Otherwise a simple transposition (performed when sorting by charges) brings it to upper triangular form.

\subsection*{18.1.21 speigs}
- full name: tenpy.linalg.np_conserved.speigs
- parent module: tenpy.Iinalg.np_conserved
- type: function
tenpy.linalg.np_conserved.speigs ( \(a\), charge_sector, \(k\), *args, **kwargs)
Sparse eigenvalue decomposition \(\mathrm{w}, \mathrm{v}\) of square \(a\) in a given charge sector.
Finds \(k\) right eigenvectors (chosen by kwargs['which']) in a given charge sector, tensordot (A, \(\mathrm{V}[i]\), axes=1) \(=W[i] * V[i]\).

\section*{Parameters}
- a (Array) - A square array with contractible legs and vanishing total charge.
- charge_sector (charges) - ndim charges to select the block.
- \(\mathbf{k}\) (int) - How many eigenvalues/vectors should be calculated. If the block of charge_sector is smaller than \(k, k\) may be reduced accordingly.
- *args - Additional arguments given to scipy.sparse.linalg.eigs.
- **kwargs - Additional keyword arguments given to scipy.sparse.linalg.eigs.

\section*{Returns}
- W (ndarray) - \(k\) (or less) eigenvalues
- V (list of Array) - \(k\) (or less) right eigenvectors of \(A\) with total charge charge_sector. Note that when interpreted as a matrix, this is the transpose of what np.eigs normally gives.

\subsection*{18.1.22 svd}
- full name: tenpy.linalg.np_conserved.svd
- parent module: tenpy.linalg.np_conserved
- type: function
tenpy.linalg.np_conserved.svd (a, full_matrices=False, compute_uv=True, cutoff=None, qtotal_LR=[None, None], inner_labels=[None, None], inner_qconj=1)
Singualar value decomposition of an Array \(a\).
Factorizes \(U, S, V H=\operatorname{svd}(a)\), such that \(a=U * d i a g(S) * V H\) (where \(*\) stands for atensordot() and diag creates an correctly shaped Array with \(S\) on the diagonal). For a non-zero cutoff this holds only approximately.

There is a gauge freedom regarding the charges, see also Array.gauge_total_charge (). We ensure contractibility by setting U.legs[1] = VH.legs[0].conj(). Further, we gauge the LegCharge such that \(U\) and \(V\) have the desired qtotal_LR.

\section*{Parameters}
- a (Array, shape \((\mathrm{M}, \mathrm{N})\) ) - The matrix to be decomposed.
- full_matrices (bool) - If False (default), \(U\) and \(V\) have shapes (M, K) and (K, N ), where \(\mathrm{K}=\operatorname{len}(\mathrm{S})\). If True, \(U\) and \(V\) are full square unitary matrices with shapes ( M , \(\mathrm{M})\) and ( \(\mathrm{N}, \mathrm{N}\) ). Note that the arrays are not directly contractible in that case; diag ( S ) would need to be a rectangluar ( \(\mathrm{M}, \mathrm{N}\) ) matrix.
- compute_uv (bool) - Whether to compute and return \(U\) and \(V\).
- cutoff (None I float) - Keep only singular values which are (strictly) greater than cutoff. (Then the factorization holds only approximately). If None (default), ignored.
- qtotal_LR ([ \{charges/None\}, \{charges/None\}]) - The desired qtotal for \(U\) and \(V H\), respectively. [None, None] (Default) is equivalent to [None, a.qtotal]. A single None entry is replaced the unique charge satisfying the requirement U . qtotal + VH.qtotal \(=\) a.qtotal (modulo qmod).
- inner_labels_LR ([\{str/None\}, \{str/None\}]) - The first label corresponds to U. legs [1], the second to VH. legs [0].
- inner_qconj \((\{+1,-1\})\) - Direction of the charges for the new leg. Default +1 . The new LegCharge is constructed such that VH.legs [0].qconj \(=\) qconj.

\section*{Returns}
- U (Array) - Matrix with left singular vectors as columns. Shape (M, M) or (M, K) depending on full_matrices.
- \(\mathbf{S}\) (1D ndarray) - The singluar values of the array. If no cutoff is given, it has lenght min ( M , N) .
- VH (Array) - Matrix with right singular vectors as rows. Shape (N, N) or (K, N) depending on full_matrices.

\subsection*{18.1.23 tensordot}
- full name: tenpy.linalg.np_conserved.tensordot
- parent module: tenpy.Iinalg.np_conserved
- type: function
tenpy.linalg.np_conserved.tensordot ( \(a, b\), axes=2)
Similar as np.tensordot but for Array.
Builds the tensor product of \(a\) and \(b\) and sums over the specified axes. Does not require complete blocking of the charges.

Labels are inherited from \(a\) and \(b\). In case of a collision (= the same label would be inherited from \(a\) and \(b\) after the contraction), both labels are dropped.

Detailed implementation notes are available in the doc-string of _tensordot_worker().

\section*{Parameters}
- a (Array) - The first and second npc Array for which axes are to be contracted.
- b (Array) - The first and second npc Array for which axes are to be contracted.
- axes ((axes_a, axes_b) I int) - A single integer is equivalent to (range (-axes, \(0)\), range (axes)). Alternatively, axes_a and axes_b specifiy the legs of \(a\) and \(b\), respectively, which should be contracted. Legs can be specified with leg labels or indices. Contract leg axes_a [i] of \(a\) with leg axes_b [i] of \(b\).

Returns a_dot_b - The tensorproduct of \(a\) and \(b\), summed over the specified axes. Returns a scalar in case of a full contraction.

Return type Array

\subsection*{18.1.24 to_iterable_arrays}
- full name: tenpy.linalg.np_conserved.to_iterable_arrays
- parent module: tenpy.linalg.np_conserved
- type: function
tenpy.linalg.np_conserved.to_iterable_arrays (array_list)
Similar as to_iterable (), but also enclose npc Arrays in a list.

\subsection*{18.1.25 trace}
- full name: tenpy.linalg.np_conserved.trace
- parent module: tenpy.Iinalg.np_conserved
- type: function
tenpy.linalg.np_conserved.trace \((a, \operatorname{leg} 1=0, \operatorname{leg} 2=1)\)
Trace of \(a\), summing over leg1 and leg2.
Requires that the contracted legs are contractible (i.e. have opposite charges). Labels are inherited from \(a\).

\section*{Parameters}
- leg1 (str/int) - The leg label or index for the two legs which should be contracted (i.e. summed over).
- leg2 (str/int) - The leg label or index for the two legs which should be contracted (i.e. summed over).
Returns traced - A scalar if a.rank \(==2\), else an Array of rank a.rank - 2. Equivalent to sum([a.take_slice([i, i], [leg1, leg2]) for i in range(a. shape[leg1])]).

Return type Arrayla.dtype

\subsection*{18.1.26 zeros}
- full name: tenpy.linalg.np_conserved.zeros
- parent module: tenpy.Iinalg.np_conserved
- type: function
```

tenpy.linalg.np_conserved.zeros(legcharges, dtype=<class 'numpy.float64'>, qtotal=None, la- bels=None)

```

Create a npc array full of zeros (with no _data).
This is just a wrapper around Array (. . .), detailed documentation can be found in the class doc-string of Array.

\section*{Module description}

A module to handle charge conservation in tensor networks.
A detailed introduction to this module (including notations) can be found in Charge conservation with np_conserved.
This module np_conserved implements a class Array designed to make use of charge conservation in tensor networks. The idea is that the Array class is used in a fashion very similar to the numpy. ndarray, e.g you can call the functions tensordot () or svd () (of this module) on them. The structure of the algorithms (as DMRG) is thus the same as with basic numpy ndarrays.

Internally, an Array saves charge meta data to keep track of blocks which are nonzero. All possible operations (e.g. tensordot, svd, ...) on such arrays preserve the total charge structure. In addition, these operations make use of the charges to figure out which of the blocks it has to use/combine - this is the basis for the speed-up.
```

tenpy.linalg.np_conserved.QCUTOFF = 2.220446049250313e-15

```

A cutoff to ignore machine precision rounding errors when determining charges
tenpy.linalg.np_conserved.QTYPE = <class 'numpy.int64'>
the type used for charges

\subsection*{18.1.27 Overview}

\section*{Classes}
\begin{tabular}{ll}
\hline Array(legcharges[, dtype, qtotal, labels]) & \begin{tabular}{l} 
A multidimensional array (=tensor) for using charge \\
conservation.
\end{tabular} \\
\hline ChargeInfo([mod, names]) & Meta-data about the charge of a tensor. \\
\hline LegCharge(chargeinfo, slices, charges[, qconj]) & Save the charge data associated to a leg of a tensor. \\
\hline LegPipe(legs[, qconj, sort, bunch]) & A LegPipe combines multiple legs of a tensor to one. \\
\hline
\end{tabular}

\section*{Array creation}
\begin{tabular}{ll}
\hline Array.from_ndarray_trivial(data_flat[,...]) & \begin{tabular}{l} 
convert a flat numpy ndarray to an Array with trivial \\
charge conservation.
\end{tabular} \\
\hline Array.from_ndarray(data_flat, legcharges[,...]) & convert a flat (numpy) ndarray to an Array. \\
\hline Array.from_func(func, legcharges[, dtype, ...]) & Create an Array from a numpy func. \\
\hline Array.from_func_square(func, leg[, dtype, & Create an Array from a (numpy) function. \\
\(\ldots])\). & Create a npc array full of zeros (with no _data). \\
\hline zeros(legcharges[, dtype, qtotal, labels]) & \begin{tabular}{l} 
Return an identity matrix contractible with the leg axis \\
of the Array \(a\).
\end{tabular} \\
\hline eye_like(a[, axis, labels]) & Returns a square, diagonal matrix of entries \(s\). \\
\hline diag(s, leg[, dtype, labels]) &
\end{tabular}

\section*{Concatenation}
\begin{tabular}{ll}
\hline concatenate(arrays[, axis, copy]) & \begin{tabular}{l} 
Stack arrays along a given axis, similar as \\
np.concatenate.
\end{tabular} \\
\hline grid_concat(grid, axes[, copy]) & \begin{tabular}{l} 
Given an np.array of npc.Arrays, performs a multi- \\
dimensional concatentation along 'axes'.
\end{tabular} \\
\hline grid_outer(grid, grid_legs[, qtotal, ...]) & \begin{tabular}{l} 
Given an n.array of npc.Arrays, return the correspond- \\
ing higher-dimensional Array.
\end{tabular} \\
\hline
\end{tabular}

\section*{Detecting charges of flat arrays}
\begin{tabular}{ll}
\hline detect_qtotal(flat_array, legcharges[, cutoff]) & \begin{tabular}{l} 
Returns the total charge (w.r.t legs) of first non-zero sec- \\
tor found in lat_array.
\end{tabular} \\
\hline detect_legcharge(flat_array, chargeinfo,...) & \begin{tabular}{l} 
Calculate a missing LegCharge by looking for nonzero \\
entries of a flat array.
\end{tabular} \\
\hline \begin{tabular}{l} 
detect_grid_outer_legcharge(grid, \\
grid_legs)
\end{tabular} & \begin{tabular}{l} 
Derive a LegCharge for a grid used for \\
grid_outer().
\end{tabular} \\
\hline
\end{tabular}

\section*{Contraction of some legs}
\begin{tabular}{ll}
\hline tensordot \((\mathrm{a}, \mathrm{b}[\), axes \(])\) & Similar as np.tensordot but for Array. \\
\hline outer \((\mathrm{a}, \mathrm{b})\) & \begin{tabular}{l} 
Forms the outer tensor product, equivalent to \\
tensordot \((\mathrm{a}, \mathrm{b}\), axes \(=0)\).
\end{tabular} \\
\hline inner \((\mathrm{a}, \mathrm{b}[\), axes, do_conj] \()\) & Contract all legs in \(a\) and \(b\), return scalar. \\
\hline trace \((\mathrm{a}[, \operatorname{leg} 1\), leg2] \()\) & Trace of \(a\), summing over leg1 and leg2. \\
\hline
\end{tabular}

\section*{Linear algebra}
\begin{tabular}{ll}
\hline \(\operatorname{svd}(\mathrm{a}[\), full_matrices, compute_uv, cutoff,...\(])\) & Singualar value decomposition of an Array \(a\). \\
\hline \(\operatorname{pinv}(\mathrm{a}[\), cutoff \(])\) & \begin{tabular}{l} 
Compute the (Moore-Penrose) pseudo-inverse of a ma- \\
trix.
\end{tabular} \\
\hline norm(a[, ord, convert_to_float \(])\) & Norm of flattened data. \\
\hline \(\operatorname{qr}(\mathrm{a}[\), mode, inner_labels, cutoff \(])\) & Q-R decomposition of a matrix. \\
\hline \(\operatorname{expm}(\mathrm{a})\) & \begin{tabular}{l} 
Use scipy.linalg.expm to calculate the matrix exponen- \\
tial of a square matrix.
\end{tabular} \\
\hline
\end{tabular}

\section*{Eigen systems}
\begin{tabular}{ll}
\hline eigh(a[, UPLO, sort \(])\) & \begin{tabular}{l} 
Calculate eigenvalues and eigenvectors for a hermitian \\
matrix.
\end{tabular} \\
\hline eig(a[, sort]) & \begin{tabular}{l} 
Calculate eigenvalues and eigenvectors for a non- \\
hermitian matrix.
\end{tabular} \\
\hline eigvalsh(a[, UPLO, sort \(])\) & Calculate eigenvalues for a hermitian matrix. \\
\hline eigvals(a[, sort \(])\) & Calculate eigenvalues for a hermitian matrix. \\
\hline &
\end{tabular}

Table 12-continued from previous page
\begin{tabular}{ll}
\hline speigs(a, charge_sector, k, *args, \({ }^{* *}\) kwargs) & \begin{tabular}{l} 
Sparse eigenvalue decomposition \(\mathrm{w}, ~ \mathrm{v}\) of square \(a\) in a \\
given charge sector.
\end{tabular} \\
\hline
\end{tabular}

\section*{18.2 charges}
- full name: tenpy.linalg.charges
- parent module: tenpy. linalg
- type: module

\section*{Classes}

Chargelnfo

\begin{tabular}{ll}
\hline Charge Info([mod, names]) & Meta-data about the charge of a tensor. \\
\hline LegCharge(chargeinfo, slices, charges[, qconj]) & Save the charge data associated to a leg of a tensor. \\
\hline LegPipe(legs[, qconj, sort, bunch]) & A LegPipe combines multiple legs of a tensor to one. \\
\hline
\end{tabular}

\subsection*{18.2.1 ChargeInfo}
- full name: tenpy.linalg.charges.ChargeInfo
- parent module: tenpy.linalg.charges
- type: class

\section*{Inheritance Diagram}

> \begin{tabular}{|l|} \hline ChargeInfo \\ \hline \end{tabular}

\section*{Methods}
\begin{tabular}{ll}
\hline ChargeInfo.__init__([mod, names]) & Initialize self. \\
\hline ChargeInfo.add(chinfos) & Create a Charge Info combining multiple charges. \\
\hline ChargeInfo.change(chinfo, charge, new_qmod) & Change the qmod of a given charge. \\
\hline ChargeInfo.check_valid(charges) & \begin{tabular}{l} 
Check, if charges has all entries as expected from \\
self.mod.
\end{tabular} \\
\hline \begin{tabular}{ll} 
ChargeInfo.drop(chinfo[, charge]) & Remove a charge from a ChargeInfo. \\
\hline \begin{tabular}{l} 
ChargeInfo.from_hdf5(hdf5_loader, h5gr, sub- \\
path)
\end{tabular} & Load instance from a HDF5 file. \\
\hline \begin{tabular}{l} 
ChargeInfo.make_valid([charges]) \\
ChargeInfo.save_hdf5(hdf5_saver, h5gr, sub- \\
path)
\end{tabular} & Take charges modulo self.mod. \\
\hline ChargeInfo.test_sanity() & Sanity check, raises ValueErrors, if something is wrong. \\
\hline
\end{tabular} \\
\hline
\end{tabular}

\section*{Class Attributes and Properties}
\begin{tabular}{ll}
\hline ChargeInfo.mod & Modulo how much each of the charges is taken. \\
\hline ChargeInfo. qnumber & The number of charges. \\
class tenpy. linalg. charges. ChargeInfo (mod=[], names=None) \\
Bases: object \\
Meta-data about the charge of a tensor. \\
Saves info about the nature of the charge of a tensor. Provides make_valid() for taking modulo m. \\
(This class is implemented in tenpy. Iinalg.charges but also imported in tenpy.linalg. \\
np_conserved for convenience.)
\end{tabular}

\section*{Parameters}
- mod (iterable of QTYPE) - The len gives the number of charges, qnumber. Each entry is a positive integer, where 1 implies a \(U(1)\) charge and \(N>1\) implies a \(Z_{N}\) symmetry. Defaults to "trivial", i.e., no charge.
- names (list of str) - Descriptive names for the charges. Defaults to ['']*qnumber.

\section*{names}

A descriptive name for each of the charges. May have ' entries.
Type list of strings

\section*{_mask}
mask (mod \(==1\) ), to speed up make_valid in pure python.
Type 1D array bool

\section*{_mod_masked}

Equivalent to self.mod[self._maks_mod1]
Type 1D array QTYPE
_qnumber, _mod
Storage of quimber and mod.

\section*{Notes}

Instances of this class can (should) be shared between different LegCharge and Array's.
save_hdf5 (hdf5_saver, h5gr, subpath)
Export self into a HDF5 file.
This method saves all the data it needs to reconstruct self with from_hdf5 ().
It stores the names under the path "names", and mod as dataset "U1_ZN".

\section*{Parameters}
- hdf5_saver (Hdf5Saver) - Instance of the saving engine.
- h5gr (: class `Group \(\left.{ }^{\prime}\right)\) - HDF5 group which is supposed to represent self.
- subpath (str) - The name of \(h 5 g r\) with a '/' in the end.
classmethod from_hdf5 (hdf5_loader, h5gr, subpath)
Load instance from a HDF5 file.
This method reconstructs a class instance from the data saved with save_hdf5 ().
The "U1_ZN" dataset is mandatory, 'names ' are optional.

\section*{Parameters}
- hdf5_loader (Hdf5Loader) - Instance of the loading engine.
- h5gr (Group) - HDF5 group which is represent the object to be constructed.
- subpath (str) - The name of \(h 5 g r\) with a '/' in the end.

Returns obj - Newly generated class instance containing the required data.
Return type cls
classmethod add (chinfos)
Create a ChargeInfo combining multiple charges.
Parameters chinfos (iterable of ChargeInfo) - ChargeInfo instances to be combined into a single one (in the given order).

Returns chinfo - ChargeInfo combining all the given charges.
Return type ChargeInfo
classmethod drop (chinfo, charge=None)
Remove a charge from a ChargeInfo.

\section*{Parameters}
- chinfo (ChargeInfo) - The ChargeInfo from where to drop/remove a charge.
- charge (int / str) - Number or name of the charge (within chinfo) which is to be dropped. None means dropping all charges.

Returns chinfo - ChargeInfo where the specified charge is dropped.
Return type ChargeInfo
classmethod change (chinfo, charge, new_qmod, new_name=")
Change the qmod of a given charge.

\section*{Parameters}
- chinfo (ChargeInfo) - The ChargeInfo for which qmod of charge should be changed.
- new_qmod (int) - The new qmod to be set.
- new_name (str) - The new name of the charge.

Returns chinfo - ChargeInfo where qmod of the specified charge was changed.
Return type ChargeInfo
test_sanity ()
Sanity check, raises ValueErrors, if something is wrong.
property qnumber
The number of charges.
property mod
Modulo how much each of the charges is taken.
Entries are 1 for a \(U(1)\) charge, and N for a \(Z_{N}\) symmetry.
make_valid (charges=None)
Take charges modulo self.mod.
Parameters charges (array_like or None) - 1D or 2D array of charges, last dimension self.qnumber None defaults to trivial charges np.zeros (qnumber, dtype=QTYPE) .

Returns A copy of charges taken modulo mod, but with \(\mathrm{x} \div 1:=\mathrm{x}\)
Return type charges
check_valid (charges)
Check, if charges has all entries as expected from self.mod.
Parameters charges (2D ndarray QTYPE_t) - Charge values to be checked.
Returns res - True, if all \(0<=\) charges \(<=\) self.mod (wherever self.mod != 1 )
Return type bool

\subsection*{18.2.2 LegCharge}
- full name: tenpy.linalg.charges.LegCharge
- parent module: tenpy.Iinalg.charges
- type: class

\section*{Inheritance Diagram}

> LegCharge

\section*{Methods}
\begin{tabular}{|c|c|}
\hline LegCharge.__init__(chargeinfo, slices, charges) & Initialize self. \\
\hline LegCharge.bunch() & Return a copy with bunched self.charges: form blocks for contiguous equal charges. \\
\hline LegCharge.charge_sectors() & Return unique rows of self.charges. \\
\hline LegCharge.conj() & Return a (shallow) copy with opposite self.qconj. \\
\hline LegCharge.copy() & Return a (shallow) copy of self. \\
\hline LegCharge.extend(extra) & Return a new LegCharge, which extends self with futher charges. \\
\hline LegCharge.flip_charges_qconj() & Return a copy with both negative qconj and charges. \\
\hline LegCharge.from_add_charge(legs[, chargeinfo]) & Add the (independent) charges of two or more legs to get larger qnumber. \\
\hline LegCharge.from_change_charge(leg, charge, ...) & Remove a charge from a LegCharge. \\
\hline \[
\begin{aligned}
& \text { LegCharge.from_drop_charge(leg[, charge, } \\
& \text {...]) }
\end{aligned}
\] & Remove a charge from a LegCharge. \\
\hline LegCharge.from_hdf5(hdf5_loader, h5gr, subpath) & Load instance from a HDF5 file. \\
\hline LegCharge.from_qdict(chargeinfo, qdict[, qconj]) & Create a LegCharge from qdict form. \\
\hline  & Create a LegCharge from qflat form. \\
\hline LegCharge.from_qind(chargeinfo, slices,
charges) & Just a wrapper around self.__init__(), see class docstring for parameters. \\
\hline LegCharge.from_trivial(ind_len[,...]) & Create trivial (qnumber=0) LegCharge for given len of indices ind_len. \\
\hline LegCharge.get_block_sizes() & Return the sizes of the individual blocks. \\
\hline LegCharge.get_charge(qindex) & \[
\begin{aligned}
& \text { Return charge self.charges [qindex] * } \\
& \text { self. qconj for a given qindex. }
\end{aligned}
\] \\
\hline LegCharge.get_qindex(flat_index) & Find qindex containing a flat index. \\
\hline LegCharge.get_qindex_of_charges(charges) & Return the slice selecting the block for given charge values. \\
\hline LegCharge.get_slice(qindex) & Return slice selecting the block for a given qindex. \\
\hline LegCharge.is_blocked() & Returns whether self is blocked, i.e. qindex map 1:1 to charge values. \\
\hline LegCharge.is_bunched() & Checks whether bunch () would change something. \\
\hline LegCharge.is_sorted() & Returns whether self.charges is sorted lexiographically. \\
\hline LegCharge.perm_flat_from_perm_qind(perm & \&indyert a permutation of qind (acting on self) into a flat permutation. \\
\hline LegCharge.perm_qind_from_perm_flat(perm_ & flativert flat permutation into qind permutation. \\
\hline LegCharge.project(mask) & Return copy keeping only the indices specified by mask. \\
\hline LegCharge.save_hdf5(hdf5_saver, h5gr, subpath) & Export self into a HDF5 file. \\
\hline LegCharge.sort([bunch]) & Return a copy of self sorted by charges (but maybe not bunched). \\
\hline LegCharge.test_contractible(other) & Raises a ValueError if charges are incompatible for contraction with other. \\
\hline LegCharge.test_equal(other) & Test if charges are equal including qconj. \\
\hline LegCharge.test_sanity() & Sanity check, raises ValueErrors, if something is wrong. \\
\hline LegCharge.to_qdict() & Return charges in qdict form. \\
\hline
\end{tabular}

Table 16 - continued from previous page
\begin{tabular}{l}
\hline LegCharge.to_qflat() \\
class tenpy.linalg.charges.LegCharge (chargeinfo, slices, charges, qconj=1) \\
Bases: object
\end{tabular}

Save the charge data associated to a leg of a tensor.
This class is more or less a wrapper around a 2D numpy array charges and a 1D array slices. See Charge conservation with \(n p \_\)conserved for more details.
(This class is implemented in tenpy.linalg.charges but also imported in tenpy.linalg. np_conserved for convenience.)

\section*{Parameters}
- chargeinfo (ChargeInfo) - The nature of the charge.
- slices (1D array_like, len (block_number+1)) - A block with 'qindex' qi correspondes to the leg indices in slice (slices [qi], slices [qi+1]).
- charges (2D array_like, shape (block_number, chargeinfo. qnumber)) - charges [qi] gives the charges for a block with 'qindex' qi.
- qconj \((\{+1,-1\})\) - A flag telling whether the charge points inwards \((+1\), default) or outwards (-1).
ind_len
The number of indices for this leg.
Type int
block_number
The number of blocks, i.e., a 'qindex' for this leg is in range (block_number).
chinfo
The nature of the charge. Can be shared between LegCharges.
Type ChargeInfoinstance

\section*{slices}

A block with 'qindex' qi correspondes to the leg indices in slice (self.slices[qi], self. slices[qi+1]). See get_slice().

Type ndarray[np.intp_t,ndim=1] (block_number+1)

\section*{charges}
charges [qi] gives the charges for a block with 'qindex' qi. Note: the sign might be changed by qconj. See also get_charge ().

Type ndarray[QTYPE_t,ndim=1] (block_number, chinfo.qnumber)

\section*{qconj}

A flag telling whether the charge points inwards (+1) or outwards ( -1 ). Whenever charges are added, they should be multiplied with their qconj value.

Type \(\{-1,1\}\)
sorted
Whether the charges are guaranteed to be sorted.
Type bool

\section*{bunched}

Whether the charges are guaranteed to be bunched.

\section*{Type bool}

\section*{Notes}

Instances of this class can be shared between different npc.Array. Thus, functions changing self.slices or self. charges must always make copies. Further they must set sorted and bunched to False (if they might not preserve them).
copy ()
Return a (shallow) copy of self.
save_hdf5 (hdf5_saver, h5gr, subpath)
Export self into a HDF5 file.
This method saves all the data it needs to reconstruct self with from_hdf5().
Checks format for an ouput format key "LegCharge". Possible choices are:
"blocks" (default) Store slices and charges directly as datasets, and block_number, sorted, bunched as further attributes.
"compact" A single array np.hstack([self.slices[:-1], self.slices[1:], self. charges]) as dataset "blockcharges", and block_number, sorted, bunched as further attributes.
"flat" Insufficient (!) to recover the exact blocks; saves only the array returned by to_flat () as dataset 'charges'.

The ind_len, qconj, and the format parameter are saved as group attributes under the same names. chinfo is always saved as subgroup.

\section*{Parameters}
- hdf5_saver (Hdf5Saver) - Instance of the saving engine.
- h5gr (: class`Group`) - HDF5 group which is supposed to represent self.
- subpath (str) - The name of \(h 5 g r\) with a ' /' in the end.
classmethod from_hdf5 (hdf5_loader, h5gr, subpath)
Load instance from a HDF5 file.
This method reconstructs a class instance from the data saved with save_hdf5 ().

\section*{Parameters}
- hdf5_loader (Hdf5Loader) - Instance of the loading engine.
- h5gr (Group) - HDF5 group which is represent the object to be constructed.
- subpath (str) - The name of \(h 5 g r\) with a '/' in the end.

Returns obj - Newly generated class instance containing the required data.
Return type cls
classmethod from_trivial (ind_len, chargeinfo=None, qconj=1)
Create trivial (qnumber=0) LegCharge for given len of indices ind_len.
classmethod from_qflat (chargeinfo, qfat, qconj=1)
Create a LegCharge from qflat form.
Does neither bunch nor sort. We recommend to sort (and bunch) afterwards, if you expect that tensors using the LegCharge have entries at all positions compatible with the charges.

\section*{Parameters}
- chargeinfo (Charge Info) - The nature of the charge.
- qflat (array_like (ind_len, qnumber)) - qnumber charges for each index of the leg on entry.
- qconj ( \(\{-1,1\}\) ) - A flag telling whether the charge points inwards (+1) or outwards (-1).

\section*{See also:}
sort sorts by charges
bunch bunches contiguous blocks of the same charge.
classmethod from_qind (chargeinfo, slices, charges, qconj=1)
Just a wrapper around self. \(\qquad\) nit_( (), see class doc-string for parameters.

\section*{See also:}
sort sorts by charges
bunch bunches contiguous blocks of the same charge.
classmethod from_qdict (chargeinfo, qdict, qconj=1)
Create a LegCharge from qdict form.

\section*{Parameters}
- chargeinfo (Charge Info) - The nature of the charge.
- qdict (dict) - A dictionary mapping a tuple of charges to slices.
classmethod from_add_charge (legs, chargeinfo=None)
Add the (independent) charges of two or more legs to get larger qnumber.

\section*{Parameters}
- legs (iterable of LegCharge) - The legs for which the charges are to be combined/added.
- chargeinfo (ChargeInfo) - The ChargeInfo for all charges; create new if None.

Returns combined - A LegCharge with the charges of both legs. Is neither sorted nor bunched!

\section*{Return type LegCharge}
classmethod from_drop_charge (leg, charge=None, chargeinfo=None)
Remove a charge from a LegCharge.

\section*{Parameters}
- leg (LegCharge) - The leg from which to drop/remove a charge.
- charge (int | str) - Number or name of the charge (within chinfo) which is to be dropped. None means dropping all charges.
- chargeinfo (ChargeInfo) - The ChargeInfo with charge dropped; create new if None.
Returns dropped - A LegCharge with the specified charge dropped. Is neither sorted nor bunched!

\section*{Return type LegCharge}
classmethod from_change_charge (leg, charge, new_qmod, new_name= ", chargeinfo=None)
Remove a charge from a LegCharge.

\section*{Parameters}
- leg (LegCharge) - The leg from which to drop/remove a charge.
- charge (int / str) - Number or name of the charge (within chinfo) for which mod is to be changed.
- new_qmod (int) - The new mod to be set for charge in the ChargeInfo.
- new_name (str) - The new name for charge.
- chargeinfo (ChargeInfo) - The ChargeInfo with charge changed; create new if None.

Returns leg - A LegCharge with the specified charge changed. Is neither sorted nor bunched!
Return type LegCharge
test_sanity ()
Sanity check, raises ValueErrors, if something is wrong.
conj()
Return a (shallow) copy with opposite self.qconj.
Returns conjugated - Shallow copy of self with flipped qconj. test_contractible ()
of self with conjugated will not raise an error.
Return type LegCharge
flip_charges_qconj()
Return a copy with both negative qconj and charges.
Returns conj_charges - (Shallow) copy of self with negative qconj and charges, thus representing the very same charges. test_equal () of self with conj_charges will not raise an error.

\section*{Return type LegCharge}
```

to_qflat()

```

Return charges in qflat form.
to_qdict()
Return charges in qdict form.
Raises ValueError, if not blocked.
is_blocked()
Returns whether self is blocked, i.e. qindex map 1:1 to charge values.
```

is_sorted()

```

Returns whether self.charges is sorted lexiographically.
```

is_bunched()

```

Checks whether bunch () would change something.
test_contractible (other)
Raises a ValueError if charges are incompatible for contraction with other.
Parameters other (LegCharge) - The LegCharge of the other leg condsidered for contraction.
Raises ValueError - If the charges are incompatible for direct contraction.

\section*{Notes}

This function checks that two legs are ready for contraction. This is the case, if all of the following conditions are met:
- the ChargeInfo is equal
- the slices are equal
- the charges are the same up to opposite signs qcon \(j\) :
```

self.charges * self.qconj = - other.charges * other.qconj

```

In general, there could also be a change of the total charge, see Charge conservation with np_conserved This special case is not considered here - instead use gauge_total_charge (), if a change of the charge is desired.

If you are sure that the legs should be contractable, check whether the charges are actually valid or whether self and other are blocked or should be sorted.

\section*{See also:}
```

test_equal self.test_contractible(other) just performs self.
test_equal(other.conj()).

```
test_equal (other)
Test if charges are equal including qconj.
Check that all of the following conditions are met:
- the ChargeInfo is equal
- the slices are equal
- the charges are the same up to the signs qcon \(j\) :
```

self.charges * self.qconj = other.charges * other.qconj

```

\section*{See also:}
```

test_contractible self.test_equal(other) is equivalent to self.

```
        test_contractible(other.conj()).
get_block_sizes()
Return the sizes of the individual blocks.
Returns sizes - The sizes of the individual blocks; sizes[i] = slices[i+1] -
slices[i].
Return type ndarray, shape (block_number,)
get_slice (qindex)
Return slice selecting the block for a given qindex.
get_qindex (flat_index)
Find qindex containing a flat index.
Given a flat index, to find the corresponding entry in an Array, we need to determine the block it is saved in. For example, if slices \(=[[0,3],[3,7],[7,12]]\), the flat index 5 corresponds to the second entry, qindex \(=1\) (since 5 is in [3:7]), and the index within the block would be \(2=5-3\).

Parameters flat_index (int) - A flat index of the leg. Negative index counts from behind.

\section*{Returns}
- qindex (int) - The qindex, i.e. the index of the block containing flat_index.
- index_within_block (int) - The index of flat_index within the block given by qindex.
get_qindex_of_charges (charges)
Return the slice selecting the block for given charge values.
Inverse function of get_charge ().
Parameters charges (1D array_like) - Charge values for which the slice of the block is to be determined.

Returns slice \((\mathbf{i}, \mathbf{j})\) - Slice of the charge values for
Return type slice
:raises ValueError : if the answer is not unique (because self is not blocked).:
get_charge (qindex)
Return charge self.charges [qindex] * self.qconj for a given qindex.
sort (bunch=True)
Return a copy of self sorted by charges (but maybe not bunched).
If bunch=True, the returned copy is completely blocked by charge.
Parameters bunch (bool) - Whether self.bunch is called after sorting. If True, the leg is guaranteed to be fully blocked by charge.

\section*{Returns}
- perm_qind (array (self.block_len, )) - The permutation of the qindices (before bunching) used for the sorting. To obtain the flat permuation such that sorted_array[. .., :] = unsorted_array[..., perm_flat], use perm_flat = unsorted_leg.perm_flat_from_perm_qind (perm_qind)
- sorted_copy (LegCharge) - A shallow copy of self, with new qind sorted (and thus blocked if bunch) by charges.

\section*{See also:}
bunch enlarge blocks for contiguous qind of the same charges.
numpy.take can apply perm_flat to a given axis
tenpy.tools.misc.inverse_permutation returns inverse of a permutation
bunch ()
Return a copy with bunched self.charges: form blocks for contiguous equal charges.

\section*{Returns}
- idx (lD array) - idx [:-1] are the indices of the old qind which are kept, idx [-1] = old_block_number.
- \(\mathbf{~ p ~ ( L e g C h a r g e ) ~ - ~ A ~ n e w ~ L e g C h a r g e ~ w i t h ~ t h e ~ s a m e ~ c h a r g e s ~ a t ~ g i v e n ~ i n d i c e s ~ o f ~ t h e ~ l e g , ~}\) but (possibly) shorter self.charges and self.slices.

\section*{See also:}
sort sorts by charges, thus enforcing complete blocking in combination with bunch.

\section*{project (mask)}

Return copy keeping only the indices specified by mask.
Parameters mask (1D array (bool)) - Whether to keep of the indices.

\section*{Returns}
- map_qind ( \(1 D\) array) - Map of qindices, such that qind_new \(=\) map_qind[qind_old], and map_qind[qind_old] \(=-1\) for qindices projected out.
- block_masks ( \(1 D\) array) - The bool mask for each of the remaining blocks.
- projected_copy (LegCharge) - Copy of self with the qind projected by mask.
extend (extra)
Return a new LegCharge, which extends self with futher charges.
This is needed to formally increase the dimension of an Array.
Parameters extra (LegChargelint) - By what to extend, i.e. the charges to be appended to self. An int stands for extending the length of the array by a single new block of that size and zero charges.

Returns extended_leg - Copy of self extended by the charge blocks of the extra leg.
Return type LegCharge
charge_sectors()
Return unique rows of self.charges.
Returns charges - Rows are the rows of self.charges lexsorted and without duplicates.
Return type array[QTYPE, ndim=2]
perm_flat_from_perm_qind (perm_qind)
Convert a permutation of qind (acting on self) into a flat permutation.
perm_qind_from_perm_flat (perm_flat)
Convert flat permutation into qind permutation.
Parameters perm_flat (1D array) - A permutation acting on self, which doesn't mix the blocks of qind.

Returns perm_qind - The permutation of self.qind described by perm_flat.
Return type 1D array
Raises ValueError - If perm_flat mixes blocks of different qindex.

\subsection*{18.2.3 LegPipe}
- full name: tenpy.linalg.charges.LegPipe
- parent module: tenpy.linalg.charges
- type: class

\section*{Inheritance Diagram}


\section*{Methods}
\begin{tabular}{|c|c|}
\hline LegPipe.__init__(legs[, qconj, sort, bunch]) & Initialize self. \\
\hline LegPipe.bunch(*args, **kwargs) & Convert to LegCharge and call LegCharge. bunch(). \\
\hline LegPipe.charge_sectors() & Return unique rows of self.charges. \\
\hline LegPipe.conj() & Return a shallow copy with opposite self.qconj. \\
\hline LegPipe.copy() & Return a (shallow) copy of self. \\
\hline LegPipe.extend(extra) & Return a new LegCharge, which extends self with futher charges. \\
\hline LegPipe.flip_charges_qconj() & Return a copy with both negative qconj and charges. \\
\hline LegPipe.from_add_charge(legs[, chargeinfo]) & Add the (independent) charges of two or more legs to get larger qnumber. \\
\hline LegPipe.from_change_charge(leg, charge, new_qmod) & Remove a charge from a LegCharge. \\
\hline LegPipe.from_drop_charge(leg[, charge, ...]) & Remove a charge from a LegCharge. \\
\hline LegPipe.from_hdf5(hdf5_loader, h5gr, subpath) & Load instance from a HDF5 file. \\
\hline LegPipe.from_qdict(chargeinfo, qdict[, qconj]) & Create a LegCharge from qdict form. \\
\hline LegPipe.from_qflat(chargeinfo, qflat[, qconj]) & Create a LegCharge from qflat form. \\
\hline LegPipe.from_qind(chargeinfo, slices, charges) & Just a wrapper around self.__init__(), see class docstring for parameters. \\
\hline ```
LegPipe.from_trivial(ind_len[, chargeinfo,
...])
``` & Create trivial (qnumber=0) LegCharge for given len of indices ind_len. \\
\hline LegPipe.get_block_sizes() & Return the sizes of the individual blocks. \\
\hline LegPipe.get_charge(qindex) & Return charge self.charges[qindex] * self. qconj for a given qindex. \\
\hline LegPipe.get_qindex(flat_index) & Find qindex containing a flat index. \\
\hline LegPipe.get_qindex_of_charges(charges) & Return the slice selecting the block for given charge values. \\
\hline LegPipe.get_slice(qindex) & Return slice selecting the block for a given qindex. \\
\hline LegPipe.is_blocked() & Returns whether self is blocked, i.e. qindex map 1:1 to charge values. \\
\hline LegPipe.is_bunched() & Checks whether bunch () would change something. \\
\hline LegPipe.is_sorted() & Returns whether self.charges is sorted lexiographically. \\
\hline
\end{tabular}

Table 17 - continued from previous page
\begin{tabular}{ll}
\hline LegPipe.map_incoming_flat(incoming_indices) & \begin{tabular}{l} 
Map (flat) incoming indices to an index in the outgoing \\
pipe.
\end{tabular} \\
\hline LegPipe.outer_conj() & \begin{tabular}{l} 
Like conj (), but don't change qconj for incoming \\
legs.
\end{tabular} \\
\hline LegPipe.perm_flat_from_perm_qind(perm_qin@lonvert a permutation of qind (acting on self) into a flat \\
permutation.
\end{tabular}
class tenpy.linalg.charges.LegPipe (legs, qconj=1, sort=True, bunch=True)
Bases: tenpy.linalg.charges. LegCharge
A LegPipe combines multiple legs of a tensor to one.
Often, it is necessary to "combine" multiple legs into one: for example to perfom a SVD, the tensor needs to be viewed as a matrix.
This class does exactly this job: it combines multiple LegCharges ('incoming legs') into one 'pipe' (the 'outgoing leg'). The pipe itself is a LegCharge, with indices running from 0 to the product of the individual legs' ind_len, corresponding to all possible combinations of input leg indices.
(This class is implemented in tenpy.linalg.charges but also imported in tenpy.linalg. np_conserved for convenience.)

\section*{Parameters}
- legs (list of LegCharge) - The legs which are to be combined.
- qconj \((\{+1,-1\})\) - A flag telling whether the charge of the resulting pipe points inwards ( +1 , default) or outwards ( -1 ).
- sort (bool) - Whether the outgoing pipe should be sorted. Default True; recommended. Note: calling sort () after initialization converts to a LegCharge.
- bunch (bool) - Whether the outgoing pipe should be bunched. Default True; recommended. Note: calling bunch () after initialization converts to a LegCharge.

\section*{nlegs}

The number of legs.
Type int
legs
The original legs, which were combined in the pipe.
Type tuple of LegCharge

\section*{subshape}
ind_len for each of the incoming legs.
Type tuple of int

\section*{subqshape}
block_number for each of the incoming legs.
Type tuple of int

\section*{q_map}

Shape (block_number, \(3+n l e g s\) ). Rows: [ b_j, b_\{j+1\}, I_s, i_1, ..., i_\{nlegs \}], See Notes below for details.

Type array[np.intp, ndim=2]
q_map_slices
Defined such that the row indices of in range (q_map_slices [I_s], q_map_slices [I_s+1]) have q_map [:, 2] == I_s.

Type array[np.intp, ndim=1]
```

_perm

```

A permutation such that q_map [_perm, 3:] is sorted by i_l.
Type 1D array

\section*{_strides}

Strides for mapping incoming qindices \(i \_l\) to the index of q_map [_perm, : ].
Type 1D array

\section*{Notes}

For np.reshape, taking, for example, \(i, j, \ldots \rightarrow k\) amounted to \(k=s_{1} * i+s_{2} * j+\ldots\) for appropriate strides \(s_{1}, s_{2}\).

In the charged case, however, we want to block \(k\) by charge, so we must implicitly permute as well. This reordering is encoded in \(q_{-} m a p\).

Each qindex combination of the nlegs input legs \(\left(i_{1}, \ldots, i_{n l e g s}\right)\), will end up getting placed in some slice \(a_{j}: a_{j+1}\) of the outgoing pipe. Within this slice, the data is simply reshaped in usual row-major fashion (' C '-order), i.e., with strides \(s_{1}>s_{2}>\ldots\)

It will be a subslice of a new total block labeled by qindex \(I_{s}\). Because many charge combinations fuse to the same total charge, in general there will be many tuples \(\left(i_{1}, \ldots, i_{n l e g s}\right)\) belonging to the same \(I_{s}\). The rows of \(q_{\_}\)map are precisely the collections of \(\left[b_{\_} j, b_{\_}\{j+1\}, ~ I_{\_} s, i \_1, ~ . ~ . ~, ~ i \_\{n l e g s\}\right]\). Here, \(b_{j}: b_{j+1}\) denotes the slice of this qindex combination within the total block \(I_{-}\)s, i.e., \(\mathrm{b}_{-} j=a_{-j}-\) self. slices[I_s].
The rows of \(q_{-}\)map are lex-sorted first by \(I_{-} s\), then the \(i\). Each \(I_{-} s\) will have multiple rows, and the order in which they are stored in q_map is the order the data is stored in the actual tensor, i.e., it might look like
```

[ ...,
[ b_j, b_{j+1}, I_s, i_1, ..., i_{nlegs} ],
[ b_{j+1}, b_{j+2}, I_s, i'_1, ..., i'_{nlegs} ],
[0, b_{j+3}, I_s + 1, i''_1, ..., i''_{nlegs} ],
[ b_{j+3}, b_{j+4}, I_s + 1, i'''_1, ..., i'''__{nlegs}],
...]

```

The charge fusion rule is:
```

self.charges[Qi]*self.qconj == sum([l.charges[qi_l]*l.qconj for l in self.legs]) ,
mod qmod

```

Here the qindex Qi of the pipe corresponds to qindices qi_l on the individual legs.
copy ()
Return a (shallow) copy of self.
save_hdf5 (hdf5_saver, h5gr, subpath)
Export self into a HDF5 file.
This method saves all the data it needs to reconstruct self with \(f r o m \_h d f 5\) ().
In addition to the data saved for the LegCharge, it just saves the legs as subgroup.

\section*{Parameters}
- hdf5_saver (Hdf5Saver) - Instance of the saving engine.
- h5gr (: class`Group ) - HDF5 group which is supposed to represent self.
- subpath (str) - The name of \(h 5 g r\) with a '/' in the end.
classmethod from_hdf5 (hdf5_loader, h5gr, subpath)
Load instance from a HDF5 file.
This method reconstructs a class instance from the data saved with save_hdf5 ().

\section*{Parameters}
- hdf5_loader (Hdf5Loader) - Instance of the loading engine.
- h5gr (Group) - HDF5 group which is represent the object to be constructed.
- subpath (str) - The name of \(h 5 g r\) with a '/' in the end.

Returns obj - Newly generated class instance containing the required data.
Return type cls
```

test_sanity()

```

Sanity check, raises ValueErrors, if something is wrong.

\section*{to_LegCharge ()}

Convert self to a LegCharge, discarding the information how to split the legs.
Usually not needed, but called by functions, which are not implemented for a LegPipe.
conj()
Return a shallow copy with opposite self. qconj.
Returns conjugated - Shallow copy of self with flipped qconj. Whenever we contract two legs, they need to be conjugated to each other. The incoming legs of the pipe are also conjugated.

Return type LegCharge
outer_conj()
Like con \(j\) (), but don't change qcon \(j\) for incoming legs.
sort (*args, **kwargs)
Convert to LegCharge and call LegCharge. sort ().
bunch (*args, **kwargs)
Convert to LegCharge and call LegCharge. bunch ().
project (*args, **kwargs)
Convert self to LegCharge and call LegCharge. project ().
In general, this could be implemented for a LegPipe, but would make split_legs () more complicated, thus we keep it simple. If you really want to project and split afterwards, use the following work-around, which is for example used in exact_diagonalization:
1) Create the full pipe and save it separetely.
2) Convert the Pipe to a Leg \& project the array with it.
3) [... do calculations ...]
4) To split the 'projected pipe' of \(A\), create and empty array \(B\) with the legs of A , but replace the projected leg by the full pipe. Set \(A\) as a slice of \(B\). Finally split the pipe.
map_incoming_flat (incoming_indices)
Map (flat) incoming indices to an index in the outgoing pipe.
Parameters incoming_indices (iterable of int) - One (flat) index on each of the incoming legs.

Returns outgoing_index - The index in the outgoing leg.
Return type int
charge_sectors()
Return unique rows of self.charges.
Returns charges - Rows are the rows of self.charges lexsorted and without duplicates.
Return type array[QTYPE, ndim=2]
extend (extra)
Return a new LegCharge, which extends self with futher charges.
This is needed to formally increase the dimension of an Array.
Parameters extra (LegChargelint) - By what to extend, i.e. the charges to be appended to self. An int stands for extending the length of the array by a single new block of that size and zero charges.

Returns extended_leg - Copy of self extended by the charge blocks of the extra leg.
Return type LegCharge
flip_charges_qconj()
Return a copy with both negative qconj and charges.
Returns conj_charges - (Shallow) copy of self with negative qconj and charges, thus representing the very same charges. test_equal () of self with conj_charges will not raise an error.

\section*{Return type LegCharge}
classmethod from_add_charge (legs, chargeinfo=None)
Add the (independent) charges of two or more legs to get larger qnumber.

\section*{Parameters}
- legs (iterable of LegCharge) - The legs for which the charges are to be combined/added.
- chargeinfo (ChargeInfo) - The ChargeInfo for all charges; create new if None.

Returns combined - A LegCharge with the charges of both legs. Is neither sorted nor bunched!

Return type LegCharge
classmethod from_change_charge (leg, charge, new_qmod, new_name=", chargeinfo=None) Remove a charge from a LegCharge.

\section*{Parameters}
- leg (LegCharge) - The leg from which to drop/remove a charge.
- charge (int / str) - Number or name of the charge (within chinfo) for which mod is to be changed.
- new_qmod (int) - The new mod to be set for charge in the ChargeInfo.
- new_name (str) - The new name for charge.
- chargeinfo (ChargeInfo) - The ChargeInfo with charge changed; create new if None.

Returns leg - A LegCharge with the specified charge changed. Is neither sorted nor bunched!
Return type LegCharge
classmethod from_drop_charge (leg, charge=None, chargeinfo=None)
Remove a charge from a LegCharge.

\section*{Parameters}
- leg (LegCharge) - The leg from which to drop/remove a charge.
- charge (int / str)-Number or name of the charge (within chinfo) which is to be dropped. None means dropping all charges.
- chargeinfo (ChargeInfo) - The ChargeInfo with charge dropped; create new if None.

Returns dropped - A LegCharge with the specified charge dropped. Is neither sorted nor bunched!

Return type LegCharge
classmethod from_qdict (chargeinfo, qdict, qconj=1)
Create a LegCharge from qdict form.

\section*{Parameters}
- chargeinfo (ChargeInfo) - The nature of the charge.
- qdict (dict) - A dictionary mapping a tuple of charges to slices.
classmethod from_qflat (chargeinfo, qflat, qconj=1)
Create a LegCharge from qflat form.
Does neither bunch nor sort. We recommend to sort (and bunch) afterwards, if you expect that tensors using the LegCharge have entries at all positions compatible with the charges.

\section*{Parameters}
- chargeinfo (ChargeInfo) - The nature of the charge.
- qflat (array_like (ind_len, qnumber)) - qnumber charges for each index of the leg on entry.
- qconj \((\{-1,1\})\) - A flag telling whether the charge points inwards \((+1)\) or outwards (-1).

\section*{See also:}
sort sorts by charges
bunch bunches contiguous blocks of the same charge.
classmethod from_qind (chargeinfo, slices, charges, qconj=1)
Just a wrapper around self.__init__(), see class doc-string for parameters.
See also:
sort sorts by charges
bunch bunches contiguous blocks of the same charge.
classmethod from_trivial (ind_len, chargeinfo=None, qconj=1)
Create trivial (qnumber=0) LegCharge for given len of indices ind_len.
get_block_sizes()
Return the sizes of the individual blocks.
Returns sizes - The sizes of the individual blocks; sizes[i] = slices[i+1] slices[i].

Return type ndarray, shape (block_number,)
get_charge (qindex)
Return charge self.charges[qindex] * self.qconj for a given qindex.
get_qindex (flat_index)
Find qindex containing a flat index.
Given a flat index, to find the corresponding entry in an Array, we need to determine the block it is saved in. For example, if slices \(=[[0,3],[3,7],[7,12]]\), the flat index 5 corresponds to the second entry, qindex \(=1\) (since 5 is in [3:7]), and the index within the block would be \(2=5-3\).

Parameters flat_index (int) - A flat index of the leg. Negative index counts from behind.

\section*{Returns}
- qindex (int) - The qindex, i.e. the index of the block containing flat_index.
- index_within_block (int) - The index of flat_index within the block given by qindex.
get_qindex_of_charges (charges)
Return the slice selecting the block for given charge values.
Inverse function of get_charge ().
Parameters charges (1D array_like) - Charge values for which the slice of the block is to be determined.

Returns slice( \(\mathbf{i}, \mathbf{j}\) ) - Slice of the charge values for
Return type slice
:raises ValueError : if the answer is not unique (because self is not blocked).:
```

get_slice(qindex)

```

Return slice selecting the block for a given qindex.
```

is_blocked()

```

Returns whether self is blocked, i.e. qindex map 1:1 to charge values.

\section*{is_bunched()}

Checks whether bunch () would change something.
is_sorted()
Returns whether self.charges is sorted lexiographically.
perm_flat_from_perm_qind (perm_qind)
Convert a permutation of qind (acting on self) into a flat permutation.
perm_qind_from_perm_flat (perm_flat)
Convert flat permutation into qind permutation.
Parameters perm_flat (1D array) - A permutation acting on self, which doesn't mix the blocks of qind.

Returns perm_qind - The permutation of self.qind described by perm_flat.
Return type 1D array
Raises ValueError - If perm_flat mixes blocks of different qindex.
test_contractible (other)
Raises a ValueError if charges are incompatible for contraction with other.
Parameters other (LegCharge) - The LegCharge of the other leg condsidered for contraction.

Raises ValueError - If the charges are incompatible for direct contraction.

\section*{Notes}

This function checks that two legs are ready for contraction. This is the case, if all of the following conditions are met:
- the ChargeInfo is equal
- the slices are equal
- the charges are the same up to opposite signs qcon j:
```

self.charges * self.qconj = - other.charges * other.qconj

```

In general, there could also be a change of the total charge, see Charge conservation with np_conserved This special case is not considered here - instead use gauge_total_charge (), if a change of the charge is desired.
If you are sure that the legs should be contractable, check whether the charges are actually valid or whether self and other are blocked or should be sorted.

\section*{See also:}
test_equal self.test_contractible(other) just performs self. test_equal(other.conj()).
test_equal (other)
Test if charges are equal including qconj.
Check that all of the following conditions are met:
- the ChargeInfo is equal
- the slices are equal
- the charges are the same up to the signs qcon \(j\) :
```

self.charges * self.qconj = other.charges * other.qconj

```

See also:
test_contractible self.test_equal(other) is equivalent to self. test_contractible(other.conj()).
to_qdict ()
Return charges in qdict form.
Raises ValueError, if not blocked.
to_qflat()
Return charges in qflat form.

\section*{Module description}

Basic definitions of a charge.
This module contains implementations for handling the quantum numbers ("charges") of the Array.
In particular, the classes ChargeInfo, LegCharge and LegPipe are implemented here.

Note: The contents of this module are imported in np_conserved, so you usually don't need to import this module in your application.

A detailed introduction to np_conserved can be found in Charge conservation with np_conserved.
In this module, some functions have the python decorator @use_cython. Functions with this decorator are replaced by the ones written in Cython, implemented in the file tenpy/linalg/_npc_helper.pyx. For further details, see the definition of use_cython ().
```

tenpy.linalg.charges.QTYPE = <class 'numpy.int64'>
Numpy data type for the charges.

```

\section*{18.3 svd_robust}
- full name: tenpy.linalg.svd_robust
- parent module: tenpy. linalg
- type: module

\section*{Functions}
\begin{tabular}{ll}
\hline \(\operatorname{svd}(\mathrm{a}[\), full_matrices, compute_uv, ...]) & \begin{tabular}{l} 
Wrapper around scipy.linalg.svd() with gesvd \\
backup plan.
\end{tabular} \\
\hline svd_gesvd(a[, full_matrices, compute_uv, ...]) & \begin{tabular}{l} 
svd with LAPACK's '\#gesvd' (with \# = d/z for \\
float/complex).
\end{tabular} \\
\hline
\end{tabular}

\subsection*{18.3.1 svd}
- full name: tenpy.linalg.svd_robust.svd
- parent module: tenpy. Iinalg.svd_robust
- type: function
tenpy.linalg.svd_robust.svd (a, full_matrices=True, compute_uv=True, overwrite_a=False, check_finite=True, lapack_driver='gesdd', warn=True )
Wrapper around scipy.linalg.svd() with gesvd backup plan.
Tries to avoid raising an LinAlgError by using using the lapack_driver gesvd, if gesdd failed.
Parameters not described below are as in scipy.linalg.svd()

\section*{Parameters}
- overwrite_a (bool) - Ignored (i.e. set to False) if lapack_driver='gesdd'. Otherwise described in scipy.linalg.svd().
- lapack_driver (\{'gesdd', 'gesvd'\}, optional) - Whether to use the more efficient divide-and-conquer approach ('gesdd') or general rectangular approach ('gesvd') to compute the SVD. MATLAB and Octave use the 'gesvd' approach. Default is 'gesdd'. If 'gesdd' fails, 'gesvd' is used as backup.
- warn (bool) - Whether to create a warning when the SVD failed.

Returns U, S, Vh - As described in doc-string of scipy. linalg.svd().
Return type ndarray

\subsection*{18.3.2 svd_gesvd}
- full name: tenpy.linalg.svd_robust.svd_gesvd
- parent module: tenpy.linalg.svd_robust
- type: function
tenpy.linalg.svd_robust.svd_gesvd (a, full_matrices=True, compute_uv=True, check_finite=True )
svd with LAPACK's ‘\#gesvd’ (with \# = d/z for float/complex).
Similar as numpy.linalg.svd (), but use LAPACK 'gesvd' driver. Works only with 2D arrays. Outer part is based on the code of numpy.linalg.svd.

\section*{Parameters}
- a - See numpy.linalg.svd() for details.
- full_matrices - See numpy. linalg.svd () for details.
- compute_uv - See numpy.linalg.svd() for details.
- check_finite - check whether input arrays contain 'NaN' or 'inf'.

Returns U, S, Vh - See numpy. linalg.svd () for details.
Return type ndarray

\section*{Module description}
(More) robust version of singular value decomposition.
We often need to perform an SVD. In general, an SVD is a matrix factorization that is always well defined and should also work for ill-conditioned matrices. But sadly, both numpy.linalg.svd() and scipy.linalg.svd () fail from time to time, raising LinalgError ("SVD did not converge"). The reason is that both of them call the LAPACK function \#gesdd (where \# depends on the data type), which takes an iterative approach that can fail. However, it is usually much faster than the alternative (and robust) \#gesvd.

Our workaround is as follows: we provide a function svd () with call signature as scipy's svd. This function is basically just a wrapper around scipy's svd, i.e., we keep calling the faster dgesdd. But if that fails, we can still use dgesvd as a backup.

Sadly, dgesvd and zgesvd were not included into scipy until version ' 0.18 .0 ' (nor in numpy), which is as the time of this writing the latest stable scipy version. For scipy version newer than ' 0.18 .0 ', we make use of the new keyword 'lapack_driver' for svd, otherwise we (try to) load dgesvd and zgesvd from shared LAPACK libraries.

The tribute for the dgesvd wrapper code goes to 'jgarcke’, originally posted at http://projects.scipy.org/ numpy/ticket/990, which is now hosted at https://github.com/numpy/numpy/issues/1588 He explains a bit more in detail what fails.

The include of dgesvd to scipy was done in https://github.com/scipy/scipy/pull/5994.

\section*{Examples}

The idea is that you just import the \(s v d\) from this module and use it as replacement for np.linalg.svd or scipy. linalg.svd:
```

>>> from tenpy.linalg.svd_robust import svd

```
\(\ggg \mathrm{U}, \mathrm{S}, \mathrm{VT}=\operatorname{svd}([[1 ., 1],.[0 ., 1]]\).

\section*{18.4 random_matrix}
- full name: tenpy.linalg.random_matrix
- parent module: tenpy. Iinalg
- type: module

\section*{Functions}
\begin{tabular}{ll}
\hline\(C O E(\) size \()\) & Circular orthogonal ensemble (COE). \\
\hline\(C R E(\) size \()\) & Circular real ensemble (CRE). \\
\hline\(C U E(\) size \()\) & Circular unitary ensemble (CUE). \\
\hline\(G O E(\) size \()\) & Gaussian orthogonal ensemble (GOE). \\
\hline GUE(size) & Gaussian unitary ensemble (GUE). \\
\hline\(O_{-C l o s e \_1(s i z e[, ~ a]) ~}^{\text {return an random orthogonal matrix 'close' to the Iden- }}\) \\
\hline tity.
\end{tabular}

Table 19 - continued from previous page
\begin{tabular}{ll} 
standard_normal_complex(size \()\) & return \((\mathrm{R}+1 \cdot j * I)\) for independent \(R\) and \(I\) from \\
& np.random.standard_normal.
\end{tabular}

\subsection*{18.4.1 COE}
- full name: tenpy.linalg.random_matrix.COE
- parent module: tenpy.Iinalg.random_matrix
- type: function
```

tenpy.linalg.random_matrix.COE (size)

```

Circular orthogonal ensemble (COE).
Parameters size (tuple) - (n, n), where \(n\) is the dimension of the output matrix.
Returns U - Unitary, symmetric (complex) matrix drawn from the COE (=Haar measure on this space).
Return type ndarray

\subsection*{18.4.2 CRE}
- full name: tenpy.linalg.random_matrix.CRE
- parent module: tenpy.linalg.random_matrix
- type: function
tenpy.linalg.random_matrix.CRE (size)
Circular real ensemble (CRE).
Parameters size (tuple) \(-(n, n)\), where \(n\) is the dimension of the output matrix.
Returns \(\mathbf{U}\) - Orthogonal matrix drawn from the CRE (=Haar measure on \(\mathrm{O}(\mathrm{n})\) ).
Return type ndarray

\subsection*{18.4.3 CUE}
- full name: tenpy.linalg.random_matrix.CUE
- parent module: tenpy.linalg.random_matrix
- type: function
tenpy.linalg.random_matrix.CUE (size)
Circular unitary ensemble (CUE).
Parameters size \((\operatorname{tuple})-(n, n)\), where \(n\) is the dimension of the output matrix.
Returns \(\mathbf{U}\) - Unitary matrix drawn from the CUE (=Haar measure on \(\mathrm{U}(\mathrm{n})\) ).
Return type ndarray

\subsection*{18.4.4 GOE}
- full name: tenpy.linalg.random_matrix.GOE
- parent module: tenpy.Iinalg.random_matrix
- type: function
tenpy.linalg.random_matrix. GOE (size)
Gaussian orthogonal ensemble (GOE).
Parameters size (tuple) \(-(n, n)\), where \(n\) is the dimension of the output matrix.
Returns H - Real, symmetric numpy matrix drawn from the GOE, i.e. \(p(H)=\) \(1 / Z \exp \left(-n / 4 \operatorname{tr}\left(H^{2}\right)\right)\)
Return type ndarray

\subsection*{18.4.5 GUE}
- full name: tenpy.linalg.random_matrix.GUE
- parent module: tenpy.linalg.random_matrix
- type: function
tenpy.linalg.random_matrix. GUE (size)
Gaussian unitary ensemble (GUE).
Parameters size (tuple) - (n, n), where \(n\) is the dimension of the output matrix.
Returns H - Hermitian (complex) numpy matrix drawn from the GUE, i.e. \(p(H)=\) \(1 / Z \exp \left(-n / 4 \operatorname{tr}\left(H^{2}\right)\right)\).
Return type ndarray

\subsection*{18.4.6 O_close_1}
- full name: tenpy.linalg.random_matrix.O_close_1
- parent module: tenpy.Iinalg.random_matrix
- type: function
tenpy.linalg.random_matrix.o_close_1 (size, \(a=0.01\) ) return an random orthogonal matrix 'close' to the Identity.

\section*{Parameters}
- size \((\operatorname{tuple})-(n, n)\), where \(n\) is the dimension of the output matrix.
- a (float) - Parameter determining how close the result is on \(O ; \lim _{a \rightarrow 0}<|O-E|>_{a}=0\) (where \(E\) is the identity).
Returns \(\mathbf{O}\) - Orthogonal matrix close to the identiy (for small \(a\) ).
Return type ndarray

\subsection*{18.4.7 U_close_1}
- full name: tenpy.linalg.random_matrix.U_close_1
- parent module: tenpy.Iinalg.random_matrix
- type: function
tenpy.linalg.random_matrix.U_close_1 (size, \(a=0.01\) )
return an random orthogonal matrix 'close' to the identity.

\section*{Parameters}
- size (tuple) - ( \(n, n\) ), where \(n\) is the dimension of the output matrix.
- a (float) - Parameter determining how close the result is to the identity. \(\lim _{a \rightarrow 0}<\) \(|O-E|>_{a}=0\) (where \(E\) is the identity).

Returns \(\mathbf{U}\) - Unitary matrix close to the identiy (for small \(a\) ). Eigenvalues are chosen i.i.d. as \(\exp (1 . j * a * x)\) with \(x\) uniform in \([-1,1]\).
Return type ndarray

\subsection*{18.4.8 box}
- full name: tenpy.linalg.random_matrix.box
- parent module: tenpy.linalg.random_matrix
- type: function
tenpy.linalg.random_matrix.box (size, \(W=1.0\) )
return random number uniform in (-W, W].

\subsection*{18.4.9 standard_normal_complex}
- full name: tenpy.linalg.random_matrix.standard_normal_complex
- parent module: tenpy.Iinalg.random_matrix
- type: function
```

tenpy.linalg.random_matrix.standard_normal_complex(size)
return (R + 1.j*I) for independent R and I from np.random.standard_normal.

```

\section*{Module description}

Provide some random matrix ensembles for numpy.
The implemented ensembles are:
\begin{tabular}{|l|l|l|l|l|}
\hline ensemble & matrix class drawn from & measure & invariant under & beta \\
\hline GOE & real, symmetric & \(\sim \exp \left(-\mathrm{n} / 4 \operatorname{tr}\left(\mathrm{H}^{\wedge} 2\right)\right)\) & orthogonal O & 1 \\
\hline GUE & hermitian & \(\sim \exp \left(-\mathrm{n} / 2 \operatorname{tr}\left(\mathrm{H}^{\wedge} 2\right)\right)\) & unitary U & 2 \\
\hline CRE & O(n) & Haar & orthogonal O & \(/\) \\
\hline COE & U in U(n) with \(\mathrm{U}=\mathrm{U}^{\wedge} \mathrm{T}\) & Haar & orthogonal O & 1 \\
\hline CUE & \(\mathrm{U}(\mathrm{n})\) & Haar & \(/\) & 2 \\
\hline O_close_1 & O(n) & \(?\) & \(/\) & \(/\) \\
\hline U_close_1 & \(\mathrm{U}(\mathrm{n})\) & \(?\) & \(/\) \\
\hline
\end{tabular}

All functions in this module take a tuple ( \(\mathrm{n}, \mathrm{n}\) ) as first argument, such that we can use the function from_func () to generate a block diagonal Array with the block from the corresponding ensemble, for example:
```

npc.Array.from_func_square(GOE, [leg, leg.conj()])

```

\section*{18.5 sparse}
- full name: tenpy.linalg.sparse
- parent module: tenpy. linalg
- type: module

\section*{Classes}

\begin{tabular}{|c|c|}
\hline FlatHermitianoperator(*args, **kwargs) & Hermitian variant of FlatLinearOperator. \\
\hline FlatLinearOperator(*args, **kwargs) & Square Linear operator acting on numpy arrays based on a matvec acting on npc Arrays. \\
\hline NpcLinearOperator() & Prototype for a Linear Operator acting on Array. \\
\hline NpcLinearOperatorWrapper(orig_operator) & Base class for wrapping around another NpcLinearOperator. \\
\hline OrthogonalNpcLinearOperator(orig_operator, ...) & Replace H -> P H P with the projector P = 1 sum - |0> <ol. \\
\hline ShiftNpcLinearOperator(orig_operator, shift) & ```
Representes original_operator + shift *
identity.
``` \\
\hline SumNpCLinearOperator(orig_operator, ...) & Sum of two linear operators. \\
\hline
\end{tabular}

\subsection*{18.5.1 FlatHermitianOperator}
- full name: tenpy.linalg.sparse.FlatHermitianOperator
- parent module: tenpy. Iinalg.sparse
- type: class

\section*{Inheritance Diagram}


\section*{Methods}
\begin{tabular}{ll}
\hline \begin{tabular}{l} 
FlatHermitianOperator. \\
init__(npc_matvec,...)
\end{tabular} & Initialize this LinearOperator. \\
\hline FlatHermitianOperator.adjoint() & Hermitian adjoint. \\
\hline FlatHermitianOperator.dot(x) & Matrix-matrix or matrix-vector multiplication. \\
\hline \begin{tabular}{l} 
FlatHermitianOperator. \\
eigenvectors(*args,...)
\end{tabular} & Same as FlatLinearOperator(..., hermitian=True). \\
\hline FlatHermitianOperator.flat_to_npc(vec) & \begin{tabular}{l} 
Convert flat numpy vector of selected charge sector into \\
npc Array.
\end{tabular} \\
\hline FlatHermitianOperator. & \begin{tabular}{l} 
Convert flat vector of undetermined charge sectors into \\
flat_to_npc_None_sector(vec)
\end{tabular} \\
\hline FlatHermitianOperator. & \begin{tabular}{l} 
Convert flat vector of all charge sectors into npc Array \\
flat_to_npc_all_sectors(vec)
\end{tabular} \\
\hline FlatHermitianOperator. & Create a FlatLinearOperator from a square Array. \\
from_NpcArray(mat[,...]) & \begin{tabular}{l} 
Create a FlatLinearOperator from a matvec function \\
FlatHermitianOperator.
\end{tabular} \\
acting on multiple legs.
\end{tabular}

Table 21 - continued from previous page
\begin{tabular}{ll}
\hline \begin{tabular}{l} 
FlatHermitianOperator. \\
npc_to_flat_all_sectors(npc_vec)
\end{tabular} & \begin{tabular}{l} 
Convert npc Array with qtotal = self.charge_sector into \\
ndarray.
\end{tabular} \\
\hline FlatHermitianOperator.rmatmat \((\mathrm{X})\) & Adjoint matrix-matrix multiplication. \\
\hline FlatHermitianOperator.rmatvec \((\mathrm{x})\) & Adjoint matrix-vector multiplication. \\
\hline FlatHermitianOperator.transpose () & Transpose this linear operator. \\
\hline
\end{tabular}

\section*{Class Attributes and Properties}
\begin{tabular}{ll}
\hline FlatHermitianOperator. H & Hermitian adjoint. \\
\hline FlatHermitianOperator. \(T\) & Transpose this linear operator. \\
\hline FlatHermitianOperator.charge_sector & Charge sector of the vector which is acted on. \\
\hline FlatHermitianOperator. ndim &
\end{tabular}
class tenpy.linalg.sparse.FlatHermitianOperator(*args, **kwargs)
Bases: tenpy.linalg. sparse.FlatLinearOperator
Hermitian variant of \(F\) latLinearOperator.
Note that we don't check matvec () to return a hermitian result, we only define an adjoint to be self.
property H
Hermitian adjoint.
Returns the Hermitian adjoint of self, aka the Hermitian conjugate or Hermitian transpose. For a complex matrix, the Hermitian adjoint is equal to the conjugate transpose.

Can be abbreviated self.H instead of self.adjoint().
Returns \(\mathbf{A} \_\mathbf{H}\) - Hermitian adjoint of self.
Return type LinearOperator
property \(T\)
Transpose this linear operator.
Returns a LinearOperator that represents the transpose of this one. Can be abbreviated self.T instead of self.transpose().
adjoint()
Hermitian adjoint.
Returns the Hermitian adjoint of self, aka the Hermitian conjugate or Hermitian transpose. For a complex matrix, the Hermitian adjoint is equal to the conjugate transpose.
Can be abbreviated self.H instead of self.adjoint().
Returns \(\mathbf{A} \_\mathbf{H}\) - Hermitian adjoint of self.
Return type LinearOperator
property charge_sector
Charge sector of the vector which is acted on.
\(\operatorname{dot}(x)\)
Matrix-matrix or matrix-vector multiplication.
Parameters \(\mathbf{x}\) (array_like)-1-d or 2-d array, representing a vector or matrix.

Returns Ax-1-d or 2-d array (depending on the shape of x ) that represents the result of applying this linear operator on x .
Return type array
flat_to_npc (vec)
Convert flat numpy vector of selected charge sector into npc Array.
If charge_sector is not None, convert to a 1D npc vector with leg self.leg. Otherwise convert vec, which can be non-zero in all charge sectors, to a npc matrix with an additional 'charge' leg to allow representing the full vector at once.

Parameters vec (1D ndarray) - Numpy vector to be converted. Should have the entries according to self.charge_sector.
Returns npc_vec - Same as vec, but converted into a npc array.
Return type Array
flat_to_npc_None_sector (vec, cutoff=1e-10)
Convert flat vector of undetermined charge sectors into npc Array.
The charge sector to be used is chosen as the block with the maximal norm, not by self.charge_sector (which might be None).

Parameters vec (1D ndarray) - Numpy vector to be converted.
Returns npc_vec - Same as vec, but converted into a npc array.

\section*{Return type Array}
flat_to_npc_all_sectors (vec)
Convert flat vector of all charge sectors into npc Array with extra "charge" leg.
Deprecated since version 0.7.3: This is merged into flat_to_npc () with self.charge_sector = None.

Parameters vec (1D ndarray) - Numpy vector to be converted.
Returns npc_vec - Same as vec, but converted into a npc array.
Return type Array
classmethod from_NpcArray (mat, charge_sector=0)
Create a FlatLinearOperator from a square Array.

\section*{Parameters}
- mat (Array) - A square matrix, with contractable legs.
- charge_sector (None I charges | 0 ) - Selects the charge sector of the vector onto which the Linear operator acts. None stands for all sectors, 0 stands for the zero-charge sector. Defaults to 0 , i.e., assumes the dominant eigenvector is in charge sector 0 .
```

classmethod from_guess_with_pipe (npc_matvec, v0_guess, labels_split=None,
dtype=None)

```

Create a FlatLinearOperator` from a matvec function acting on multiple legs.
This function creates a wrapper matvec function to allow acting on a "vector" with multiple legs. The wrapper combines the legs into a LegPipe before calling the actual matvec function, and splits them again in the end.

\section*{Parameters}
- npc_matvec (function) - Function to calculate the action of the linear operator on an npc vector with the given split labels labels_split. Has to return an npc vector with the same legs.
- v0_guess (Array) - Initial guess/starting vector which can be applied to npc_matvec.
- labels_split (None / list of str) - Labels of v0_guess in the order in which they are to be combined into a LegPipe. None defaults to v0_guess. get_leg_labels().
- dtype (np.dtype / None) - The data type of the arrays. None defaults to dtype of v0_guess (!).

\section*{Returns}
- lin_op (cls) - Instance of the class to be used as linear operator
- guess_flat (np.ndarray) - Numpy vector representing the guess v0_guess.
matmat ( \(X\) )
Matrix-matrix multiplication.
Performs the operation \(y=A * X\) where \(A\) is an \(M x N\) linear operator and \(X\) dense \(N * K\) matrix or ndarray.
Parameters \(\mathbf{X}\) ( \(\{\) matrix, ndarray \(\})\) - An array with shape \((\mathrm{N}, \mathrm{K})\).
Returns \(\mathbf{Y}\) - A matrix or ndarray with shape \((\mathrm{M}, \mathrm{K})\) depending on the type of the X argument.
Return type \{matrix, ndarray\}

Notes
This matmat wraps any user-specified matmat routine or overridden _matmat method to ensure that y has the correct type.
```

matvec(x)

```

Matrix-vector multiplication.
Performs the operation \(y=A^{*} x\) where \(A\) is an \(M x N\) linear operator and \(x\) is a column vector or 1-d array.
Parameters \(\mathbf{x}\) (\{matrix, ndarray \(\})\) - An array with shape ( N, ) or \((\mathrm{N}, 1)\).
Returns \(\mathbf{y}\) - A matrix or ndarray with shape ( M ,) or ( \(\mathrm{M}, 1\) ) depending on the type and shape of the x argument.

Return type \{matrix, ndarray\}

\section*{Notes}

This matvec wraps the user-specified matvec routine or overridden _matvec method to ensure that y has the correct shape and type.
```

npc_to_flat(npc_vec)

```

Convert npc Array into a 1D ndarray, inverse of flat_to_npc ().
Parameters npc_vec (Array) - Npc Array to be converted. If self.charge_sector is not None, this should be a 1D array with that qtotal. If self.charge_sector is not None, it should have an additional "charge" leg, (as returned by flat_to_npc ()).

Returns vec - Same entries as \(n p c_{-} v e c\), but converted into a flat Numpy array.
Return type 1D ndarray
npc_to_flat_all_sectors (npc_vec)
Convert npc Array with qtotal \(=\) self.charge_sector into ndarray.
Deprecated since version 0.7.3: This is merged into npc_to_flat () with self.charge_sector = None.

Parameters npc_vec (Array) - Npc Array to be converted. Should only have entries in self.charge_sector.

Returns vec - Same as \(n p c_{-} v e c\), but converted into a flat Numpy array.
Return type 1D ndarray
rmatmat ( \(X\) )
Adjoint matrix-matrix multiplication.
Performs the operation \(y=A^{\wedge} H^{*} x\) where \(A\) is an \(M x N\) linear operator and \(x\) is a column vector or 1-d array, or 2-d array. The default implementation defers to the adjoint.

Parameters \(\mathbf{X}\) (\{matrix, ndarray\}) - A matrix or 2D array.
Returns \(\mathbf{Y}\) - A matrix or 2D array depending on the type of the input.
Return type \{matrix, ndarray

\section*{Notes}

This rmatmat wraps the user-specified rmatmat routine.
rmatvec \((x)\)
Adjoint matrix-vector multiplication.
Performs the operation \(y=A^{\wedge} H^{*} x\) where \(A\) is an \(M x N\) linear operator and \(x\) is a column vector or 1-d array.

Parameters \(\mathbf{x}\) (\{matrix, ndarray\}) - An array with shape (M,) or (M,1).
Returns \(\mathbf{y}\) - A matrix or ndarray with shape \((\mathrm{N}\), ) or \((\mathrm{N}, 1)\) depending on the type and shape of the x argument.

Return type \{matrix, ndarray

\section*{Notes}

This rmatvec wraps the user-specified rmatvec routine or overridden _rmatvec method to ensure that y has the correct shape and type.

\section*{transpose()}

Transpose this linear operator.
Returns a LinearOperator that represents the transpose of this one. Can be abbreviated self.T instead of self.transpose().
eigenvectors (*args, **kwargs)
Same as FlatLinearOperator(..., hermitian=True).

\subsection*{18.5.2 FlatLinearOperator}
- full name: tenpy.linalg.sparse.FlatLinearOperator
- parent module: tenpy. linalg.sparse
- type: class

\section*{Inheritance Diagram}


\section*{Methods}
\begin{tabular}{|c|c|}
\hline FlatLinearOperator.__init__(npc_matvec, leg, ...) & Initialize this LinearOperator. \\
\hline FlatLinearOperator.adjoint() & Hermitian adjoint. \\
\hline FlatLinearOperator.dot(x) & Matrix-matrix or matrix-vector multiplication. \\
\hline \[
\begin{aligned}
& \text { FlatLinearOperator. } \\
& \text { eigenvectors([num_ev, ...]) }
\end{aligned}
\] & Find (dominant) eigenvector(s) of self using scipy. sparse.linalg.eigs(). \\
\hline FlatLinearOperator.flat_to_npc(vec) & Convert flat numpy vector of selected charge sector into npc Array. \\
\hline \begin{tabular}{l}
FlatLinearOperator. \\
flat_to_npc_None_sector(vec)
\end{tabular} & Convert flat vector of undetermined charge sectors into npc Array. \\
\hline \begin{tabular}{l}
FlatLinearOperator. \\
flat_to_npc_all_sectors(vec)
\end{tabular} & Convert flat vector of all charge sectors into npc Array with extra "charge" leg. \\
\hline FlatLinearOperator.from_NpcArray(mat[,
...] ] & Create a FlatLinearOperator from a square Array. \\
\hline \begin{tabular}{l}
FlatLinearOperator. \\
from_guess_with_pipe(...)
\end{tabular} & Create a FlatLinearOperator` from a matvec function acting on multiple legs. \\
\hline FlatLinearOperator.matmat(X) & Matrix-matrix multiplication. \\
\hline FlatLinearOperator.matvec(x) & Matrix-vector multiplication. \\
\hline FlatLinearOperator.npc_to_flat(npc_vec) & Convert npc Array into a 1D ndarray, inverse of flat_to_npc(). \\
\hline \begin{tabular}{l}
FlatLinearOperator. \\
npc_to_flat_all_sectors(npc_vec)
\end{tabular} & Convert npc Array with qtotal = self.charge_sector into ndarray. \\
\hline FlatLinearOperator.rmatmat (X) & Adjoint matrix-matrix multiplication. \\
\hline FlatLinearOperator.rmatvec(x) & Adjoint matrix-vector multiplication. \\
\hline FlatLinearOperator.transpose() & Transpose this linear operator. \\
\hline
\end{tabular}

\section*{Class Attributes and Properties}
\begin{tabular}{ll}
\hline FlatLinearOperator. H & Hermitian adjoint. \\
\hline FlatLinearOperator. & Transpose this linear operator. \\
\hline FlatLinearOperator.charge_sector & Charge sector of the vector which is acted on. \\
\hline FlatLinearOperator. ndim & \\
\hline
\end{tabular}
```

class tenpy.linalg.sparse.FlatLinearOperator(*args, **kwargs)

```

Bases: scipy.sparse.linalg.interface.LinearOperator
Square Linear operator acting on numpy arrays based on a matvec acting on npc Arrays.
Note that this class represents a square linear operator. In terms of charges, this means it has legs [self.leg. conj(), self.leg] and trivial (zero) qtotal.

\section*{Parameters}
- npc_matvec (function) - Function to calculate the action of the linear operator on an npc vector (with the specified leg). Has to return an npc vector with the same leg.
- leg (LegCharge) - Leg of the vector on which npc_matvec can act on.
- dtype (np.dtype) - The data type of the arrays.
- charge_sector (None I charges I 0) - Selects the charge sector of the vector onto which the Linear operator acts. None stands for all sectors, 0 stands for the zero-charge sector. Defaults to 0, i.e., assumes the dominant eigenvector is in charge sector 0 .
- vec_label (None / str) - Label to be set to the npc vector before acting on it with npc_matvec. Ignored if None.
possible_charge_sectors
Each row corresponds to one possible choice for charge_sector.
Type ndarray[QTYPE, ndim=2]
shape
The dimensions for the selected charge sector.
Type (int, int)

\section*{dtype}

The data type of the arrays.
Type np.dtype
leg
Leg of the vector on which npc_matvec can act on.
Type LegCharge

\section*{vec_label}

Label to be set to the npc vector before acting on it with npc_matvec. Ignored if None.
Type None Istr

\section*{npc_matvec}

Function to calculate the action of the linear operator on an npc vector (with one leg).
Type function

\section*{matvec_count}

The number of times npc_matvec was called.
Type int
_mask
The indices of leg corresponding to the charge_sector to be diagonalized.
Type ndarray[ndim=1, bool]
```

_npc_matvec_multileg

```

Only set if initalized with from_guess_with_pipe (). The npc_matvec function to be wrapped around. Takes the npc Array in multidimensional form and returns it that way.

Type function I None

\section*{_labels_split}

Only set if initalized with from_guess_with_pipe (). Labels of the guess before combining them into a pipe (stored as leg).

Type list of str
classmethod from_NpcArray (mat, charge_sector=0)
Create a FlatLinearOperator from a square Array.

\section*{Parameters}
- mat (Array) - A square matrix, with contractable legs.
- charge_sector (None I charges I 0) - Selects the charge sector of the vector onto which the Linear operator acts. None stands for all sectors, 0 stands for the zero-charge sector. Defaults to 0 , i.e., assumes the dominant eigenvector is in charge sector 0 .
classmethod from_guess_with_pipe (npc_matvec, v0_guess, labels_split=None, dtype \(=\) None )
Create a FlatLinearOperator` from a matvec function acting on multiple legs.
This function creates a wrapper matvec function to allow acting on a "vector" with multiple legs. The wrapper combines the legs into a LegPipe before calling the actual matvec function, and splits them again in the end.

\section*{Parameters}
- npc_matvec (function) - Function to calculate the action of the linear operator on an npc vector with the given split labels labels_split. Has to return an npe vector with the same legs.
- v0_guess (Array) - Initial guess/starting vector which can be applied to npc_matvec.
- labels_split (None | list of str) - Labels of v0_guess in the order in which they are to be combined into a LegPipe. None defaults to v0_guess. get_leg_labels().
- dtype (np.dtype | None) - The data type of the arrays. None defaults to dtype of v0_guess (!).

\section*{Returns}
- lin_op (cls) - Instance of the class to be used as linear operator
- guess_flat (np.ndarray) - Numpy vector representing the guess v0_guess.
```

property charge_sector

```

Charge sector of the vector which is acted on.
flat_to_npc (vec)
Convert flat numpy vector of selected charge sector into npc Array.
If charge_sector is not None, convert to a 1D npc vector with leg self.leg. Otherwise convert vec, which can be non-zero in all charge sectors, to a npc matrix with an additional 'charge' leg to allow representing the full vector at once.

Parameters vec (1D ndarray) - Numpy vector to be converted. Should have the entries according to self.charge_sector.

Returns npc_vec - Same as vec, but converted into a npc array.
Return type Array
npc_to_flat (npc_vec)
Convert npc Array into a 1D ndarray, inverse of flat_to_npc ().
Parameters npc_vec (Array) - Npc Array to be converted. If self.charge_sector is not None, this should be a 1D array with that qtotal. If self.charge_sector is not None, it should have an additional "charge" leg, (as returned by flat_to_npc ()).

Returns vec - Same entries as npc_vec, but converted into a flat Numpy array.
Return type 1D ndarray
flat_to_npc_all_sectors (vec)
Convert flat vector of all charge sectors into npc Array with extra "charge" leg.
Deprecated since version 0.7.3: This is merged into flat_to_npc () with self.charge_sector = None.

Parameters vec (1D ndarray) - Numpy vector to be converted.
Returns npc_vec - Same as vec, but converted into a npc array.
Return type Array
flat_to_npc_None_sector (vec, cutoff=1e-10)
Convert flat vector of undetermined charge sectors into npc Array.
The charge sector to be used is chosen as the block with the maximal norm, not by self.charge_sector (which might be None).

Parameters vec (1D ndarray) - Numpy vector to be converted.
Returns npc_vec - Same as vec, but converted into a npc array.
Return type Array
npc_to_flat_all_sectors (npc_vec)
Convert npc Array with qtotal \(=\) self.charge_sector into ndarray.
Deprecated since version 0.7.3: This is merged into npc_to_flat () with self.charge_sector = None.

Parameters npc_vec (Array) - Npc Array to be converted. Should only have entries in self.charge_sector.
Returns vec - Same as \(n p c \_v e c\), but converted into a flat Numpy array.
Return type 1D ndarray
eigenvectors (num_ev=1, max_num_ev=None, max_tol=1e-12, which='LM', v0=None, \(v 0 \_n p c=\) None, cutoff \(=1 e-12\), hermitian \(=\) False, \(* *\) kwargs \()\)
Find (dominant) eigenvector(s) of self using scipy.sparse.linalg.eigs().
If no charge_sector was selected, we look in all charge sectors.

\section*{Parameters}
- num_ev (int) - Number of eigenvalues/vectors to look for.
- max_num_ev (int) - scipy.sparse.linalg.speigs () somtimes raises a NoConvergenceError for small num_ev, which might be avoided by increasing num_ev. As a work-around, we try it again in the case of an error, just with larger num_ev up to max_num_ev. None defaults to num_ev +2 .
- max_tol (float) - After the first NoConvergenceError we increase the tol argument to that value.
- which (str) - Which eigenvalues to look for, see scipy.sparse.linalg. eigs (). More details also in argsort ().
- v0 (Array) - Initial guess as a "flat" numpy array.
- v0_npc (Array) - Initial guess, to be converted by npc_to_flat ().
- cutoff (float) - Only used if self.charge_sector is None; in that case it determines when entries in a given charge-block are considered nonzero, and what counts as degenerate.
- hermitian (bool) - If False (default), use scipy.sparse.linalg.eigs() If True, assume that self is hermitian and use scipy.sparse.linalg.eigsh().
- **kwargs - Further keyword arguments given to scipy.sparse.linalg. eigsh() or scipy.sparse.linalg.eigs(), respectively.

\section*{Returns}
- eta (1D ndarray) - The eigenvalues, sorted according to which.
- w (list of Array) - The eigenvectors corresponding to eta, as npc.Array with LegPipe.

\section*{property H}

Hermitian adjoint.
Returns the Hermitian adjoint of self, aka the Hermitian conjugate or Hermitian transpose. For a complex matrix, the Hermitian adjoint is equal to the conjugate transpose.

Can be abbreviated self.H instead of self.adjoint().
Returns A_H - Hermitian adjoint of self.
Return type LinearOperator
property \(T\)
Transpose this linear operator.
Returns a LinearOperator that represents the transpose of this one. Can be abbreviated self.T instead of self.transpose().

\section*{adjoint()}

Hermitian adjoint.
Returns the Hermitian adjoint of self, aka the Hermitian conjugate or Hermitian transpose. For a complex matrix, the Hermitian adjoint is equal to the conjugate transpose.

Can be abbreviated self.H instead of self.adjoint().
Returns A_H - Hermitian adjoint of self.
Return type LinearOperator
\(\operatorname{dot}(x)\)
Matrix-matrix or matrix-vector multiplication.
Parameters \(\mathbf{x}\) (array_like)-1-d or 2-d array, representing a vector or matrix.
Returns Ax-1-d or 2-d array (depending on the shape of \(x\) ) that represents the result of applying this linear operator on \(x\).

Return type array
matmat ( \(X\) )
Matrix-matrix multiplication.
Performs the operation \(y=A * X\) where \(A\) is an \(M x N\) linear operator and \(X\) dense \(N^{*} K\) matrix or ndarray.
Parameters \(\mathbf{X}(\{\) matrix, ndarray \(\})\) - An array with shape \((N, K)\).
Returns \(\mathbf{Y}\) - A matrix or ndarray with shape ( \(\mathrm{M}, \mathrm{K}\) ) depending on the type of the X argument.
Return type \{matrix, ndarray

\section*{Notes}

This matmat wraps any user-specified matmat routine or overridden _matmat method to ensure that y has the correct type.
matvec ( \(x\) )
Matrix-vector multiplication.
Performs the operation \(y=A * x\) where \(A\) is an \(M x N\) linear operator and \(x\) is a column vector or 1-d array.
Parameters \(\mathbf{x}(\{\) matrix, ndarray \(\})\) - An array with shape \((\mathrm{N}\), ) or \((\mathrm{N}, 1)\).
Returns \(\mathbf{y}\) - A matrix or ndarray with shape \((\mathrm{M}\), ) or \((\mathrm{M}, 1)\) depending on the type and shape of the x argument.
Return type \{matrix, ndarray

\section*{Notes}

This matvec wraps the user-specified matvec routine or overridden _matvec method to ensure that y has the correct shape and type.
rmatmat ( \(X\) )
Adjoint matrix-matrix multiplication.
Performs the operation \(y=A^{\wedge} H^{*} x\) where \(A\) is an \(M x N\) linear operator and \(x\) is a column vector or 1-d array, or 2-d array. The default implementation defers to the adjoint.

Parameters X (\{matrix, ndarray\}) - A matrix or 2D array.
Returns \(\mathbf{Y}\) - A matrix or 2D array depending on the type of the input.
Return type \(\{\) matrix, ndarray \(\}\)

\section*{Notes}

This rmatmat wraps the user-specified rmatmat routine.
rmatvec \((x)\)
Adjoint matrix-vector multiplication.
Performs the operation \(y=A^{\wedge} H^{*} x\) where \(A\) is an \(M x N\) linear operator and \(x\) is a column vector or 1-d array.

Parameters \(\mathbf{x}(\{\) matrix, \(n d a r r a y\})\) - An array with shape ( \(M\), ) or ( \(M, 1\) ).
Returns \(\mathbf{y}\) - A matrix or ndarray with shape \((\mathrm{N}\),\() or (\mathrm{N}, 1)\) depending on the type and shape of the x argument.
Return type \{matrix, ndarray

\section*{Notes}

This rmatvec wraps the user-specified rmatvec routine or overridden _rmatvec method to ensure that \(y\) has the correct shape and type.

\section*{transpose()}

Transpose this linear operator.
Returns a LinearOperator that represents the transpose of this one. Can be abbreviated self.T instead of self.transpose().

\subsection*{18.5.3 NpcLinearOperator}
- full name: tenpy.linalg.sparse.NpcLinearOperator
- parent module: tenpy.linalg.sparse
- type: class

\section*{Inheritance Diagram}

\footnotetext{
NpcLinearOperator
}

\section*{Methods}
\begin{tabular}{ll}
\hline NpcLinearOperator.__init__() & Initialize self. \\
\hline NpcLinearOperator.adjoint() & Return the hermitian conjugate of self \\
\hline NpcLinearOperator.matvec(vec) & Calculate the action of the operator on a vector vec. \\
\hline NpcLinearOperator.to_matrix() & Contract self to a matrix. \\
\hline
\end{tabular}

\section*{Class Attributes and Properties}

\section*{NpcLinearOperator.acts_on}

\section*{class tenpy.linalg.sparse.NpcLinearOperator}

Bases: ob ject
Prototype for a Linear Operator acting on Array.
Note that an Array implements a matvec function. Thus you can use any (square) npc Array as an NpcLinearOperator.

\section*{dtype}

The data type of its action.
Type np.type
acts_on
Labels of the state on which the operator can act. NB: Class attribute.
Type list of str
matvec (vec)
Calculate the action of the operator on a vector vec.
Note that we don't require vec to be one-dimensional. However, for square operators we require that the result of matvec has the same legs (in the same order) as vec such that they can be added. Note that this excludes a non-trivial qtotal for square operators.
to_matrix()
Contract self to a matrix.
If self represents an operator with very small shape, e.g. because the MPS bond dimension is very small, an algorithm might choose to contract self to a single tensor.

Returns matrix - Contraction of the represented operator.
Return type Array
adjoint()
Return the hermitian conjugate of self
If self is hermitian, subclasses can choose to implement this to define the adjoint operator of self.

\subsection*{18.5.4 NpcLinearOperatorWrapper}
- full name: tenpy.linalg.sparse.NpcLinearOperatorWrapper
- parent module: tenpy.linalg.sparse
- type: class

\section*{Inheritance Diagram}
```

NpcLinearOperatorWrapper

```

\section*{Methods}
\begin{tabular}{ll}
\hline \begin{tabular}{l} 
NpcLinearOperatorWrapper. \\
init__(orig_operator)
\end{tabular} & Initialize self. \\
\hline NpcLinearOperatorWrapper.adjoint() & Return the hermitian conjugate of self. \\
\hline NpcLinearOperatorWrapper.to_matrix() & Contract self to a matrix. \\
\hline NpcLinearOperatorWrapper.unwrapped() & Return to the original NpcLinearOperator. \\
\hline class tenpy.linalg.sparse. NpcLinearOperatorWrapper (orig_operator) \\
Bases: object \\
Base class for wrapping around another NpcLinearOperator. \\
Attributes not explicitly set with self.attribute = value (or by defining methods) default to the attributes of the \\
wrapped orig_operator.
\end{tabular}

Warning: If there are multiple levels of wrapping operators, the order might be critical to get correct results; e.g. OrthogonalNpcLinearOperator needs to be the outer-most wrapper to produce correct results and/or be efficient.

Parameters orig_operator (NpcLinearOperator) - The original operator implementing the matvec.
orig_operator
The original operator implementing the matvec.
Type NpcLinearOperator
unwrapped ()
Return to the original NpcLinearOperator.
If multiple levels of wrapping were used, this returns the most unwrapped one.
```

to_matrix()

```

Contract self to a matrix.
adjoint()
Return the hermitian conjugate of self.
If self is hermitian, subclasses can choose to implement this to define the adjoint operator of self.

\subsection*{18.5.5 OrthogonalNpcLinearOperator}
- full name: tenpy.linalg.sparse.OrthogonalNpcLinearOperator
- parent module: tenpy.linalg.sparse
- type: class

\section*{Inheritance Diagram}


\section*{Methods}
\begin{tabular}{|c|c|}
\hline \[
\begin{aligned}
& \text { OrthogonalNpcLinearOperator. } \\
& \text { _init__(...) }
\end{aligned}
\] & Initialize self. \\
\hline OrthogonalNpcLinearOperator.adjoint() & Return the hermitian conjugate of self. \\
\hline OrthogonalNpcLinearOperator. matvec(vec) & \\
\hline OrthogonalNpcLinearOperator.
to_matrix() & Contract self to a matrix. \\
\hline OrthogonalNpcLinearOperator. unwrapped() & Return to the original NpcLinearOperator. \\
\hline \multicolumn{2}{|l|}{class tenpy.linalg.sparse.OrthogonalNpcLinearOperator (orig_operator,ortho_vecs) Bases: tenpy.linalg.sparse.NpcLinearOperatorWrapper} \\
\hline \multicolumn{2}{|l|}{Replace H -> P H P with the projector P = 1 - sum_0 |0> <o|.} \\
\hline \multicolumn{2}{|l|}{Here, | \(0>\) are the vectors from ortho_vecs.} \\
\hline \multicolumn{2}{|l|}{Parameters} \\
\hline
\end{tabular}
- ortho_vecs (list of Array) - The vectors to orthogonalize against.
to_matrix()
Contract self to a matrix.
adjoint()
Return the hermitian conjugate of self.
If self is hermitian, subclasses can choose to implement this to define the adjoint operator of self.

\section*{unwrapped ()}

Return to the original NpcLinearOperator.
If multiple levels of wrapping were used, this returns the most unwrapped one.

\subsection*{18.5.6 ShiftNpcLinearOperator}
- full name: tenpy.linalg.sparse.ShiftNpcLinearOperator
- parent module: tenpy.linalg.sparse
- type: class

\section*{Inheritance Diagram}


\section*{Methods}
\begin{tabular}{ll}
\hline ShiftNpcLinearOperator.__init__(...) & Initialize self. \\
\hline ShiftNpcLinearOperator.adjoint() & Return the hermitian conjugate of self. \\
\hline ShiftNpcLinearOperator.matvec(vec) & \\
\hline ShiftNpcLinearOperator.to_matrix() & Contract self to a matrix. \\
\hline ShiftNpcLinearOperator.unwrapped() & Return to the original NpcLinearOperator. \\
\hline & \\
class tenpy.linalg.sparse.ShiftNpcLinearOperator (orig_operator, shift) \\
Bases: tenpy.linalg.sparse.NpcLinearOperatorWrapper \\
Representes original_operator + shift * identity. \\
This can be usefule.g. for better Lanczos convergence.
\end{tabular}

\section*{to_matrix()}

Contract self to a matrix.

\section*{adjoint()}

Return the hermitian conjugate of self.
If self is hermitian, subclasses can choose to implement this to define the adjoint operator of self. unwrapped ()

Return to the original NpcLinearOperator.
If multiple levels of wrapping were used, this returns the most unwrapped one.

\subsection*{18.5.7 SumNpcLinearOperator}
- full name: tenpy.linalg.sparse.SumNpcLinearOperator
- parent module: tenpy.linalg.sparse
- type: class

\section*{Inheritance Diagram}


\section*{Methods}
\begin{tabular}{ll}
\hline \begin{tabular}{l} 
SumNpcLinearOperator. \\
init_(orig_operator,...)
\end{tabular} & Initialize self. \\
\hline SumNpcLinearOperator.adjoint() & Return the hermitian conjugate of self. \\
\hline SumNpcLinearOperator.matvec(vec) & \\
\hline SumNpcLinearOperator.to_matrix() & Contract self to a matrix. \\
\hline SumNpcLinearOperator.unwrapped() & Return to the original NpcLinearOperator. \\
\hline & \\
class tenpy.linalg.sparse.SumNpcLinearOperator (orig_operator, other_operator) \\
Bases: tenpy.linalg.sparse.NpcLinearOperatorWrapper \\
Sum of two linear operators. \\
to_matrix() \\
Contract self to a matrix.
\end{tabular}
```

adjoint()

```

Return the hermitian conjugate of self.
If self is hermitian, subclasses can choose to implement this to define the adjoint operator of self.
```

unwrapped()

```

Return to the original NpcLinearOperator.
If multiple levels of wrapping were used, this returns the most unwrapped one.

\section*{Module description}

Providing support for sparse algorithms (using matrix-vector products only).
Some linear algebra algorithms, e.g. Lanczos, do not require the full representations of a linear operator, but only the action on a vector, i.e., a matrix-vector product matvec. Here we define the strucuture of such a general operator, NpcLinearOperator, as it is used in our own implementations of these algorithms (e.g., lanczos). Moreover, the FlatLinearOperator allows to use all the scipy sparse methods by providing functionality to convert flat numpy arrays to and from np_conserved arrays.

\section*{18.6 lanczos}
- full name: tenpy.linalg.lanczos
- parent module: tenpy. Iinalg
- type: module

\section*{Classes}

\begin{tabular}{ll}
\hline LanczosEvolution(H, psi0, options) & Calculate \(\exp (\) delta \(H) \mid p\) si \(0>\) using Lanczos. \\
\hline LanczosGroundState \((H\), psi0, options \([, \ldots])\) & Lanczos algorithm working on npc arrays. \\
\hline
\end{tabular}

\section*{Functions}
\begin{tabular}{ll}
\hline gram_schmidt(vecs[, rcond, verbose]) & \begin{tabular}{l} 
In place Gram-Schmidt Orthogonalization and normal- \\
ization for npc Arrays.
\end{tabular} \\
\hline lanczos(H, psi[, options, orthogonal_to]) & \begin{tabular}{l} 
Simple wrapper calling LanczosGroundState (H, \\
psi, options, orthogonal_to).run()
\end{tabular} \\
\hline lanczos_arpack(H, psi[, options, orthogonal_to]) & \begin{tabular}{l} 
Usescipy.sparse.linalg.eigsh() to find the \\
ground state of \(H\).
\end{tabular} \\
\hline plot_stats(ax, Es) & Plot the convergence of the energies. \\
\hline
\end{tabular}

\subsection*{18.6.1 gram_schmidt}
- full name: tenpy.linalg.lanczos.gram_schmidt
- parent module: tenpy.linalg.lanczos
- type: function
tenpy.linalg.lanczos.gram_schmidt (vecs, rcond=le-14, verbose=None)
In place Gram-Schmidt Orthogonalization and normalization for npc Arrays.

\section*{Parameters}
- vecs (list of Array) - The vectors which should be orthogonalized. All with the same order of the legs. Entries are modified in place. if a norm < rcond, the entry is set to None.
- rcond (float) - Vectors of norm < rcond (after projecting out previous vectors) are discarded.

\section*{Returns}
- vecs (list of Array) - The ortho-normalized vectors (without any None).
- ov (2D Array) - For \(j>=i, o v[j, i]=n p c . i n n e r(v e c s[j], ~ v e c s[i]\), 'range', do_conj=True) (where vecs[j] was orthogonalized to all vecs[k], \(k\) < i).

\subsection*{18.6.2 lanczos}
- full name: tenpy.linalg.lanczos.lanczos
- parent module: tenpy. Iinalg. lanczos
- type: function
```

tenpy.linalg.lanczos.lanczos(H,psi,options={},orthogonal_to=[])
Simple wrapper calling LanczosGroundState(H, psi, options, orthogonal_to).run()

```

Deprecated since version 0.6.0: Going to remove the orthogonal_to argument. Instead, replace H with OrthogonalNpcLinearOperator(H, orthogonal_to) using the OrthogonalNpcLinearOperator.

\section*{Parameters}
- H-See LanczosGroundState.
- psi-See LanczosGroundState.
- options - See LanczosGroundState.
- orthogonal_to-See LanczosGroundState.

Returns See LanczosGroundState.run().
Return type E0, psi0, N

\subsection*{18.6.3 lanczos_arpack}
- full name: tenpy.linalg.lanczos.lanczos_arpack
- parent module: tenpy.linalg.lanczos
- type: function
tenpy.linalg.lanczos.lanczos_arpack (H, psi, options=\{\},orthogonal_to=[])
Use scipy.sparse.linalg.eigsh() to find the ground state of \(H\).
This function has the same call/return structure as \(\operatorname{lanczos}()\), but uses the ARPACK package through the functions speigsh () instead of the custom lanczos implementation in LanczosGroundState. This function is mostly intended for debugging, since it requires to convert the vector from np_conserved Array into a flat numpy array and back during each matvec-operation!

Deprecated since version 0.6.0: Going to remove the orthogonal_to argument. Instead, replace H with OrthogonalNpcLinearOperator(H, orthogonal_to) using the OrthogonalNpcLinearoperator.

\section*{Parameters}
- \(\mathbf{H}\) - See LanczosGroundState. \(H\) and psi should have/use labels.
- psi - See LanczosGroundState. \(H\) and psi should have/use labels.
- options - See LanczosGroundState. \(H\) and psi should have/use labels.
- orthogonal_to-See LanczosGroundState. \(H\) and psi should have/use labels.

\section*{Returns}
- E0 (float) - Ground state energy.
- psi0 (Array) - Ground state vector.

\subsection*{18.6.4 plot_stats}
- full name: tenpy.linalg.lanczos.plot_stats
- parent module: tenpy.linalg.lanczos
- type: function
tenpy.linalg.lanczos.plot_stats (ax, Es)
Plot the convergence of the energies.

\section*{Parameters}
- ax (matplotlib. axes.Axes) - The axes on which we should plot.
- Es(list of ndarray.) - The energies Lanczos.Es.

Module description
Lanczos algorithm for np_conserved arrays.

\section*{MODELS}
- full name: tenpy.models
- parent module: tenpy
- type: module

\section*{Module description}

Definition of the various models.
For an introduction to models see Models.
The module tenpy.models.model contains base classes for models. The module tenpy.models.lattice contains base classes and implementations of lattices. All other modules in this folder contain model classes derived from these base classes.

\section*{Submodules}
\begin{tabular}{ll}
\hline lattice & Classes to define the lattice structure of a model. \\
\hline model & This module contains some base classes for models. \\
\hline
\end{tabular}

\section*{19.1 lattice}
- full name: tenpy.models.lattice
- parent module: tenpy.models
- type: module

\section*{Classes}

\begin{tabular}{ll}
\hline Chain(L, site, **kwargs) & A chain of L equal sites. \\
\hline HelicalLattice(regular_lattice, N_unit_cells) & \begin{tabular}{l} 
Translation invariant version of a tilted, regular 2D lat- \\
tice.
\end{tabular} \\
\hline Honeycomb(Lx, Ly, sites, **kwargs) & A honeycomb lattice. \\
\hline IrregularLattice(regular_lattice[, remove, ...]) & \begin{tabular}{l} 
A variant of a regular lattice, where we might have extra \\
sites or sites missing.
\end{tabular} \\
\hline Kagome(Lx, Ly, sites, **kwargs) & A Kagome lattice. \\
\hline Ladder(L, sites, **kwargs) & A ladder coupling two chains. \\
\hline Lattice(Ls, unit_cell[, order, bc, bc_MPS, ...]) & A general, regular lattice. \\
\hline SimpleLattice(Ls, site, **kwargs) & A lattice with a unit cell consiting of just a single site. \\
\hline Square(Lx, Ly, site, **kwargs) & A square lattice. \\
\hline Triangular(Lx, Ly, site, **kwargs) & A triangular lattice. \\
\hline TrivialLattice(mps_sites, **kwargs) & \begin{tabular}{l} 
Trivial lattice consisting of a single (possibly large) unit \\
cell in 1D.
\end{tabular} \\
\hline
\end{tabular}

\subsection*{19.1.1 Chain}
- full name: tenpy.models.lattice.Chain
- parent module: tenpy.models.lattice
- type: class

\section*{Inheritance Diagram}


\section*{Methods}
\begin{tabular}{|c|c|}
\hline Chain.__init__(L, site, **kwargs) & Initialize self. \\
\hline Chain.count_neighbors([u, key]) & Count e.g. \\
\hline Chain.coupling_shape(dx) & Calculate correct shape of the strengths for a coupling. \\
\hline Chain.distance(u1, u2, dx) & Get the distance for a given coupling between two sites in the lattice. \\
\hline Chain.enlarge_mps_unit_cell([factor]) & Repeat the unit cell for infinite MPS boundary conditions; in place. \\
\hline Chain.find_coupling_pairs([max_dx, cutoff, eps]) & Automatically find coupling pairs grouped by distances. \\
\hline Chain.from_hdf5(hdf5_loader, h5gr, subpath) & Load instance from a HDF5 file. \\
\hline Chain.lat2mps_idx(lat_idx) & Translate lattice indices (x_0, ..., \(x \_\{D-1\}\), u) to MPS index \(i\). \\
\hline Chain.mps2lat_idx(i) & Translate MPS index \(i\) to lattice indices ( \(\mathrm{x} \_0\), ...., x_\{dim-1\}, u). \\
\hline Chain.mps2lat_values(A[, axes, u]) & same as Lattice.mps2lat_values(), but ignore \(u\), setting it to 0 . \\
\hline Chain.mps2lat_values_masked(A[, axes,...]) & Reshape/reorder an array \(A\) to replace an MPS index by lattice indices. \\
\hline Chain.mps_idx_fix_u([u]) & return an index array of MPS indices for which the site within the unit cell is \(u\). \\
\hline Chain.mps_lat_idx_fix_u([u]) & Similar as mps_idx_fix_u(), but return also the corresponding lattice indices. \\
\hline Chain.mps_sites() & Return a list of sites for all MPS indices. \\
\hline Chain.multi_coupling_shape(dx) & Calculate correct shape of the strengths for a multi_coupling. \\
\hline Chain.number_nearest_neighbors([u]) & Deprecated. \\
\hline Chain.number_next_nearest_neighbors([u]) & Deprecated. \\
\hline Chain.ordering(order) & Provide possible orderings of the \(N\) lattice sites. \\
\hline
\end{tabular}

Table 3 - continued from previous page
\begin{tabular}{|c|c|}
\hline Chain.plot_basis(ax[, origin, shade]) & Plot arrows indicating the basis vectors of the lattice. \\
\hline Chain.plot_bc_identified(ax[, direction, ...]) & Mark two sites indified by periodic boundary conditions. \\
\hline Chain.plot_coupling(ax[, coupling, wrap]) & Plot lines connecting nearest neighbors of the lattice. \\
\hline Chain.plot_order(ax[, order, textkwargs]) & Plot a line connecting sites in the specified "order" and text labels enumerating them. \\
\hline Chain.plot_sites(ax[, markers]) & Plot the sites of the lattice with markers. \\
\hline Chain.position(lat_idx) & return 'space' position of one or multiple sites. \\
\hline \(\qquad\) strength]) & Find possible MPS indices for two-site couplings. \\
\hline Chain.possible_multi_couplings(ops[, strength]) & Generalization of possible_couplings() to couplings with more than 2 sites. \\
\hline Chain.save_hdf5(hdf5_saver, h5gr, subpath) & Export self into a HDF5 file. \\
\hline Chain.site(i) & return Site instance corresponding to an MPS index \(i\) \\
\hline Chain.test_sanity() & Sanity check. \\
\hline
\end{tabular}

\section*{Class Attributes and Properties}
\begin{tabular}{ll}
\hline Chain.Lu & \begin{tabular}{l} 
the (expected) number of sites in the \\
len (unit_cell).
\end{tabular} \\
\hline Chain.boundary_conditions & \begin{tabular}{l} 
Human-readable list of boundary conditions from bc \\
and bc_shift.
\end{tabular} \\
\hline Chain.dim & the dimension of the lattice \\
\hline Chain.nearest_neighbors & \\
\hline Chain.next_next_nearest_neighbors & Defines an ordering of the lattice sites, thus mapping the \\
Chain.order lattice to a 1D chain.
\end{tabular}


\section*{Parameters}
- L(int) - The lenght of the chain.
- site (Site) - The local lattice site. The unit_cell of the Lattice is just [site].
- **kwargs - Additional keyword arguments given to the Lattice. pairs are initialize with [next_] next_] nearest_neighbors. positions can be specified as a single vector.

\section*{\(\operatorname{dim}=1\)}
the dimension of the lattice
ordering (order)
Provide possible orderings of the \(N\) lattice sites.
The following orders are defined in this method compared to Lattice.ordering():
\begin{tabular}{|c|c|}
\hline \[
\begin{aligned}
& \mathrm{or}- \\
& \mathrm{der}
\end{aligned}
\] & Resulting order \\
\hline \multicolumn{2}{|l|}{'defaum,t '1, 2, 3, 4, ... , L-1} \\
\hline & ed, \(\mathrm{L}-1,1, \mathrm{~L}-2, \ldots, \mathrm{~L} / / 2\). This order might be usefull if you want to consider a ring with periodic boundary conditions with a finite MPS: It avoids the ultra-long range of the coupling from site 0 to L present in the default order. \\
\hline
\end{tabular}
property boundary_conditions
Human-readable list of boundary conditions from bc and bc_shift.
Returns boundary_conditions - List of "open" or "periodic", one entry for each direction of the lattice.

Return type list of str
count_neighbors ( \(u=0\), key='nearest_neighbors')
Count e.g. the number of nearest neighbors for a site in the bulk.

\section*{Parameters}
- u(int) - Specifies the site in the unit cell, for which we should count the number of neighbors (or whatever key specifies).
- key (str) - Key of pairs to select what to count.

Returns number - Number of nearest neighbors (or whatever key specified) for the \(u\)-th site in the unit cell, somewhere in the bulk of the lattice. Note that it might not be the correct value at the edges of a lattice with open boundary conditions.

\section*{Return type int}

\section*{coupling_shape ( \(d x\) )}

Calculate correct shape of the strengths for a coupling.
Parameters \(\mathbf{d x}\) (tuple of int) - Translation vector in the lattice for a coupling of two operators. Corresponds to \(d x\) argument of tenpy.models.model. CouplingModel. add_multi_coupling().

\section*{Returns}
- coupling_shape (tuple of int) - Len dim. The correct shape for an array specifying the coupling strength. lat_indices has only rows within this shape.
- shift_lat_indices (array) - Translation vector from origin to the lower left corner of box spanned by \(d x\).
distance ( \(u 1, u 2, d x\) )
Get the distance for a given coupling between two sites in the lattice.

The \(u 1, u 2, d x\) parameters are defined in analogy with add_coupling(), i.e., this function calculates the distance between a pair of operators added with add_coupling (using the basis and unit_cell_positions of the lattice).

Warning: This function ignores "wrapping" arround the cylinder in the case of periodic boundary conditions.

\section*{Parameters}
- u1 (int) - Indices within the unit cell; the \(u 1\) and \(u 2\) of add_coupling()
- u2 (int) - Indices within the unit cell; the \(u 1\) and \(u 2\) of add_coupling()
- \(\mathbf{d x}\) (array) - Length dim. The translation in terms of basis vectors for the coupling.

Returns distance - The distance between site at lattice indices [ \(\mathrm{x}, \mathrm{y}, \mathrm{ul}\) ] and [x + \(d x[0], y+d x[1], u 2]\), ignoring any boundary effects.

Return type float
enlarge_mps_unit_cell (factor=2)
Repeat the unit cell for infinite MPS boundary conditions; in place.
Parameters factor (int) - The new number of sites in the MPS unit cell will be increased from \(N\) _sites to factor*N_sites_per_ring. Since MPS unit cells are repeated in the \(x\)-direction in our convetion, the lattice shape goes from ( \(\mathrm{Lx}, \mathrm{Ly}, \ldots, \mathrm{Lu}\) ) to (Lx*factor, Ly, ..., Lu).
find_coupling_pairs (max_dx=3, cutoff=None, eps=1e-10)
Automatically find coupling pairs grouped by distances.
Given the unit_cell_positions and basis, the coupling pairs of nearest_neighbors, next_nearest_neighbors etc at a given distance are basically fixed (although not uniquely, since we take out half of them to avoid double-counting couplings in both directions A_i B_j + B_i A_i). This function iterates through all possible couplings up to a given cutoff distance and then determines the possible pairs at fixed distances (up to round-off errors).

\section*{Parameters}
- max_dx (int) - Maximal index for each index of \(d x\) to iterate over. You need large enough values to include every possible coupling up to the desired distance, but choosing it too large might make this function run for a long time.
- cutoff (float) - Maximal distance (in the units in which basis and unit_cell_positions is given).
- eps (float) - Tolerance up to which to distances are considered the same.

Returns coupling_pairs - Keys are distances of nearest-neighbors, next-nearest-neighbors etc.
Values are [(u1, u2, dx), ...] as in pairs.
Return type dict
classmethod from_hdf5 (hdf5_loader, h5gr, subpath)
Load instance from a HDF5 file.
This method reconstructs a class instance from the data saved with save_hdf5 ().

\section*{Parameters}
- hdf5_loader (Hdf5Loader) - Instance of the loading engine.
- h5gr (Group) - HDF5 group which is represent the object to be constructed.
- subpath (str) - The name of \(h 5 g r\) with a ' /' in the end.

Returns obj - Newly generated class instance containing the required data.
Return type cls
lat2mps_idx (lat_idx)
Translate lattice indices \(\left(\mathrm{x}_{-} 0, \ldots, \mathrm{x}_{-}\{\mathrm{D}-1\}, \mathrm{u}\right)\) to MPS index \(i\).
Parameters lat_idx (array_like [.., dim+1]) - The last dimension corresponds to lattice indices ( \(\left.x \_0, \ldots, x_{-}\{D-1\}, u\right)\). All lattice indices should be positive and smaller than the corresponding entry in self.shape. Exception: for "infinite" \(b c_{-} M P S\), an \(x \_0\) outside indicates shifts accross the boundary.

Returns \(\mathbf{i}\) - MPS index/indices corresponding to lat_idx. Has the same shape as lat_idx without the last dimension.

Return type array_like
mps2lat_idx (i)
Translate MPS index \(i\) to lattice indices ( \(\left.x \_0, \ldots, x_{-}\{\operatorname{dim}-1\}, u\right)\).
Parameters i (int / array_like of int)-MPS index/indices.
Returns lat_idx - First dimensions like \(i\), last dimension has len dim \({ }^{`}+1\) and contains the lattice indices " \(\left(x_{-} 0, \ldots, x_{\_}\{d i m-1\}, u\right)\) ` corresponding to \(i\). For \(i\) accross the MPS unit cell and "infinite" \(b c_{-} M P S\), we shift \(x_{-} 0\) accordingly.

\section*{Return type array}
mps2lat_values \((A\), axes \(=0, u=\) None \()\)
same as Lattice.mps2lat_values(), but ignore \(u\), setting it to 0 .
mps2lat_values_masked \(\left(A\right.\), axes \(=-1, m p s \_i n d s=N o n e\), include_u=None)
Reshape/reorder an array \(A\) to replace an MPS index by lattice indices.
This is a generalization of mps2lat_values () allowing for the case of an arbitrary set of MPS indices present in each axis of \(A\).

\section*{Parameters}
- A (ndarray) - Some values.
- axes ((iterable of) int) - Chooses the axis of \(A\) which should be replaced. If multiple axes are given, you also need to give multiple index arrays as mps_inds.
- mps_inds ((list of) 1D ndarray) - Specifies for each axis in axes, for which MPS indices we have values in the corresponding axis of \(A\). Defaults to [np. arange (A. shape[ax]) for ax in axes]. For indices accross the MPS unit cell and "infinite" \(b c_{-} M P S\), we shift \(x_{-} 0\) accordingly.
- include_u ((list of) bool) - Specifies for each axis in axes, whether the \(u\) index of the lattice should be included into the output array res_A. Defaults to len (self. unit_cell) > 1 .

Returns res_A - Reshaped and reordered copy of A. Such that MPS indices along the specified axes are replaced by lattice indices, i.e., if MPS index \(j\) maps to lattice site ( \(x 0, x 1, x 2\) ), then res_A[..., x0, x1, x2, ...] = A[..., mps_inds[j], ...].
Return type np.ma.MaskedArray
mps_idx_fix_u ( \(u=\) None) return an index array of MPS indices for which the site within the unit cell is \(u\).
If you have multiple sites in your unit-cell, an onsite operator is in general not defined for all sites. This functions returns an index array of the mps indices which belong to sites given by self. unit_cell[u].

Parameters u(None / int) - Selects a site of the unit cell. None (default) means all sites.
Returns mps_idx - MPS indices for which self.site(i) is self.unit_cell[u]. Ordered ascending.

\section*{Return type array}
mps_lat_idx_fix_u ( \(u=\) None)
Similar as mps_idx_fix_u(), but return also the corresponding lattice indices.
Parameters u(None / int) - Selects a site of the unit cell. None (default) means all sites.

\section*{Returns}
- mps_idx (array) - MPS indices \(i\) for which self.site(i) is self. unit_cell[u].
- lat_idx ( \(2 D\) array) - The row \(j\) contains the lattice index (without \(u\) ) corresponding to mps_idx[j].
mps_sites()
Return a list of sites for all MPS indices.
Equivalent to [self.site(i) for i in range(self.N_sites)].
This should be used for sites of 1D tensor networks (MPS, MPO,...).
multi_coupling_shape ( \(d x\) )
Calculate correct shape of the strengths for a multi_coupling.
Parameters \(d \mathbf{x}(2 \mathrm{D}\) array, shape (N_ops, dim)) - dx[i, :] is the translation vector in the lattice for the \(i\)-th operator. Corresponds to the \(d x\) of each operator given in the argument ops of tenpy.models.model.CouplingModel.add_multi_coupling().

\section*{Returns}
- coupling_shape (tuple of int) - Len dim. The correct shape for an array specifying the coupling strength. lat_indices has only rows within this shape.
- shift_lat_indices (array) - Translation vector from origin to the lower left corner of box spanned by \(d x\). (Unlike for coupling_shape () it can also contain entries \(>0\) )
number_nearest_neighbors ( \(u=0\) )
Deprecated.
Deprecated since version 0.5.0: Use count_neighbors () instead.
```

number_next_nearest_neighbors ( }u=0\mathrm{ )

```

Deprecated.
Deprecated since version 0.5.0: Use count_neighbors () instead.

\section*{property order}

Defines an ordering of the lattice sites, thus mapping the lattice to a 1 D chain.
Each row of the array contains the lattice indices for one site, the order of the rows thus specifies a path through the lattice, along which an MPS will wind through through the lattice.
You can visualize the order with plot_order ().
plot_basis (ax, origin \(=(0.0,0.0)\), shade \(=\) None, \({ }^{* *}\) kwargs)
Plot arrows indicating the basis vectors of the lattice.

\section*{Parameters}
- ax (matplotlib. axes.Axes) - The axes on which we should plot.
- **kwargs - Keyword arguments for ax. arrow.
plot_bc_identified (ax, direction=- 1, origin=None, cylinder_axis=False, **kwargs)
Mark two sites indified by periodic boundary conditions.
Works only for lattice with a 2 -dimensional basis.

\section*{Parameters}
- ax (matplotlib. axes.Axes) - The axes on which we should plot.
- direction (int) - The direction of the lattice along which we should mark the idenitified sites. If None, mark it along all directions with periodic boundary conditions.
- cylinder_axis (b○○l) - Whether to plot the cylinder axis as well.
- origin (None / np.ndarray) - The origin starting from where we mark the identified sites. Defaults to the first entry of unit_cell_positions.
- **kwargs - Keyword arguments for the used ax. plot.
plot_coupling ( \(a x\), coupling=None, wrap=False, **kwargs)
Plot lines connecting nearest neighbors of the lattice.

\section*{Parameters}
- \(\mathbf{a x}\) (matplotlib. axes.Axes) - The axes on which we should plot.
- coupling (list of (u1, u2, dx)) - By default (None), use self. pairs['nearest_neighbors']. Specifies the connections to be plotted; iteating over lattice indices ( \(i 0, i 1, \ldots\) ), we plot a connection from the site (i0, i1, ..., u1) to the site (i0+dx[0], i1+dx[1], ..., u2), taking into account the boundary conditions.
- wrap (bool) - If True, plot couplings going around the boundary by directly connecting the sites it connects. This might be hard to see, as this puts lines from one end of the lattice to the other. If False, plot the couplings as dangling lines.
- **kwargs - Further keyword arguments given to ax.plot().
plot_order ( \(a x\), order=None, textkwargs=\{'color': 'r'\}, **kwargs)
Plot a line connecting sites in the specified "order" and text labels enumerating them.

\section*{Parameters}
- ax (matplotlib. axes.Axes) - The axes on which we should plot.
- order (None / 2D array (self.N_sites, self.dim+1))- The order as returned by ordering(); by default (None) use order.
- textkwargs (None I dict) - If not None, we add text labels enumerating the sites in the plot. The dictionary can contain keyword arguments for ax.text ().
- **kwargs - Further keyword arguments given to ax. plot ().
plot_sites (ax, markers=['o', ' \({ }^{\prime}\) ', 's', ' \(p\) ', ' \(h\) ', 'D'], **kwargs)
Plot the sites of the lattice with markers.

\section*{Parameters}
- ax (matplotlib.axes.Axes) - The axes on which we should plot.
- markers (list) - List of values for the keywork marker of ax.plot () to distinguish the different sites in the unit cell, a site \(u\) in the unit cell is plotted with a marker markers[u \% len(markers)].
- **kwargs - Further keyword arguments given to ax.plot ().
position (lat_idx)
return 'space' position of one or multiple sites.
Parameters lat_idx (ndarray, (... , dim+1))-Lattice indices.
Returns pos - The position of the lattice sites specified by lat_idx in real-space.
Return type ndarray, (..., dim)
possible_couplings ( \(u 1, u 2, d x\), strength=None)
Find possible MPS indices for two-site couplings.
For periodic boundary conditions (bc [a] == False) the index x_a is taken modulo Ls [a] and runs through range (Ls [a]). For open boundary conditions, \(x_{-}\)a is limited to \(0<=x_{-}\)< Ls[a] and \(0<=x_{-} a+d x[a]\) < lat. Ls [a].

\section*{Parameters}
- u1 (int) - Indices within the unit cell; the \(u 1\) and \(u 2\) of add_coupling ()
- u2 (int) - Indices within the unit cell; the \(u 1\) and \(u 2\) of add_coupling ()
- \(\mathbf{d x}\) (array) - Length dim. The translation in terms of basis vectors for the coupling.
- strength (array_like | None) - If given, instead of returning lat_indices and coupling_shape directly return the correct strength_12.

\section*{Returns}
- mps1, mps2 (ID array) - For each possible two-site coupling the MPS indices for the ul and \(u 2\).
- strength_vals (lD array) - (Only returend if strength is not None.) Such that for (i, j, s) in zip(mps1, mps2, strength_vals): iterates over all possible couplings with \(s\) being the strength of that coupling. Couplings where strength_vals \(=0\). are filtered out.
- lat_indices ( \(2 D\) int array) - (Only returend if strength is None.) Rows of lat_indices correspond to entries of mpsl and mps2 and contain the lattice indices of the "lower left corner" of the box containing the coupling.
- coupling_shape (tuple of int) - (Only returend if strength is None.) Len dim. The correct shape for an array specifying the coupling strength. lat_indices has only rows within this shape.

\section*{possible_multi_couplings (ops, strength=None)}

Generalization of possible_couplings() to couplings with more than 2 sites.
Parameters ops (list of (opname, \(\mathrm{dx}, \mathrm{u})\) ) - Same as the argument ops of add_multi_coupling().

\section*{Returns}
- mps_ijkl (2D int array) - Each row contains MPS indices \(i, j, k, l, \ldots\). for each of the operators positions. The positions are defined by \(d x(\mathrm{j}, \mathrm{k}, \mathrm{l}, \ldots\) relative to \(i\) ) and boundary coundary conditions of self (how much the box for given \(d x\) can be shifted around without hitting a boundary - these are the different rows).
- strength_vals ( \(1 D\) array) - (Only returend if strength is not None.) Such that for (ijkl, s) in zip(mps_ijkl, strength_vals): iterates over all possible couplings with \(s\) being the strength of that coupling. Couplings where strength_vals \(==0\). are filtered out.
- lat_indices ( \(2 D\) int array) - (Only returend if strength is None.) Rows of lat_indices correspond to rows of mps_ijkl and contain the lattice indices of the "lower left corner" of the box containing the coupling.
- coupling_shape (tuple of int) - (Only returend if strength is None.) Len dim. The correct shape for an array specifying the coupling strength. lat_indices has only rows within this shape.
save_hdf5 (hdf5_saver, h5gr, subpath)
Export self into a HDF5 file.
This method saves all the data it needs to reconstruct self with from_hdf5 ().
Specifically, it saves unit_cell, Ls, unit_cell_positions, basis, boundary_conditions, pairs under their name, bc_MPS as "boundary_conditions_MPS", and order as "order_for_MPS". Moreover, it saves dim and N_sites as HDF5 attributes.

\section*{Parameters}
- hdf5_saver (Hdf5Saver) - Instance of the saving engine.
- h5gr (: class `Group`) - HDF5 group which is supposed to represent self.
- subpath (str) - The name of \(h 5 g r\) with a '/' in the end.
```

site(i)

```
return Site instance corresponding to an MPS index \(i\)
```

test_sanity()

```

Sanity check.
Raises ValueErrors, if something is wrong.

\subsection*{19.1.2 HelicalLattice}
- full name: tenpy.models.lattice.HelicalLattice
- parent module: tenpy.models.lattice
- type: class

\section*{Inheritance Diagram}


\section*{Methods}
\begin{tabular}{ll}
\hline Helicallattice.__init__(regular_lattice,...) & Initialize self. \\
\hline Helicallattice.count_neighbors([u, key]) & Count e.g. \\
\hline Helicallattice.coupling_shape(dx) & Calculate correct shape of the strengths for a coupling. \\
\hline Helicallattice.distance(u1, u2, dx) & \begin{tabular}{l} 
Get the distance for a given coupling between two sites \\
in the lattice.
\end{tabular} \\
\hline Helicallattice.enlarge_mps_unit_cell([facRerpeat the unit cell for infinite MPS boundary condi- \\
tions; in place.
\end{tabular}

Table 5 - continued from previous page
\begin{tabular}{|c|c|}
\hline Helicallattice.plot_coupling(ax[, coupling, ...]) & Plot lines connecting nearest neighbors of the lattice. \\
\hline Helicallattice.plot_order(ax[, order, ...]) & Plot a line connecting sites in the specified "order" and text labels enumerating them. \\
\hline HelicalLattice.plot_sites(ax[, markers]) & Plot the sites of the lattice with markers. \\
\hline HelicalLattice.position(lat_idx) & return 'space' position of one or multiple sites. \\
\hline Helicallattice.possible_couplings(u1, u2, dx) & Find possible MPS indices for two-site couplings. \\
\hline Helicallattice.possible_multi_coupling & (Gpxseralization of possible_couplings () to couplings with more than 2 sites. \\
\hline HelicalLattice.save_hdf5(hdf5_saver, h5gr, ...) & Export self into a HDF5 file. \\
\hline HelicalLattice.site(i) & return Site instance corresponding to an MPS index \(i\) \\
\hline Helicallattice.test_sanity() & Sanity check. \\
\hline
\end{tabular}

\section*{Class Attributes and Properties}
\begin{tabular}{ll}
\hline HelicalLattice.Lu & \begin{tabular}{l} 
the (expected) number of sites in the unit cell, \\
len(unit_cell).
\end{tabular} \\
\hline HelicalLattice.boundary_conditions & \begin{tabular}{l} 
Human-readable list of boundary conditions from bc \\
and bc_shift.
\end{tabular} \\
\hline HelicalLattice.dim & The dimension of the lattice. \\
\hline HelicalLattice.nearest_neighbors & \\
\hline HelicalLattice.next_nearest_neighbors & \\
\hline HelicalLattice.next_next_nearest_neighbors \\
HelicalLattice.order & \begin{tabular}{l} 
Defines an ordering of the lattice sites, thus mapping the \\
lattice to a 1D chain.
\end{tabular} \\
\hline
\end{tabular}
class tenpy.models.lattice. HelicalLattice (regular_lattice, N_unit_cells)
Bases: tenpy.models.lattice.Lattice
Translation invariant version of a tilted, regular 2D lattice.
A 2D lattice on an infinite cylinder becomes translation invariant by a single lattice unit cell if we tilt/shift the boundary conditions around the cylinder such that the unit cell at ( \(x, y=L y-1\) ) is neighbored by \((x+1\), \(\mathrm{y}=0\) ), and the MPS winds as a helix around the cylinder. Let's illustrate this for the Square lattice with a single-site unit cell - for a multi-site unit cell, imagine it being inserted at each of the sites of a Square lattice.

Warning: Some assumptions of the regular lattice like "the number of the sites in the MPS unit cell is product (lat. shape)" no longer hold for this model! Be very careful e.g. for getting the units of the correlation_length() right.

Parameters N_unit_cells (int) - Number of lattice unit cells to include into the MPS unit cell. The total number of sites will be N_unit_cells * len(regular_lattice. unit_cell).

```

save_hdf5 (hdf5_saver, h5gr, subpath)

```

Export self into a HDF5 file.
This method saves all the data it needs to reconstruct self with from_hdf5 ().
Specifically, it saves unit_cell, Ls, unit_cell_positions, basis, boundary_conditions, pairs under their name, bc_MPS as "boundary_conditions_MPS", and order as "order_for_MPS". Moreover, it saves dim and N_sites as HDF5 attributes.

\section*{Parameters}
- hdf5_saver (Hdf5Saver) - Instance of the saving engine.
- h5gr (:class`Group`) - HDF5 group which is supposed to represent self.
- subpath (str) - The name of \(h 5 g r\) with a '/' in the end.
classmethod from_hdf5 (hdf5_loader, h5gr, subpath)
Load instance from a HDF5 file.
This method reconstructs a class instance from the data saved with save_hdf5 ().

\section*{Parameters}
- hdf5_loader (Hdf5Loader) - Instance of the loading engine.
- h5gr (Group) - HDF5 group which is represent the object to be constructed.
- subpath (str) - The name of \(h 5 g r\) with a '/' in the end.

Returns obj - Newly generated class instance containing the required data.
Return type cls
ordering (order)
Provide possible orderings of the lattice sites.
Parameters order - Argument for the Lattice.ordering() of the regular_lattice, or 2D ndarray providing the order of the regular lattice. Note that really the only freedom is the order of the sites in the unit_cell.

Returns order - The order to be used for order, i.e. order with added/removed sites as specified by remove and add.

Return type array, shape ( \(\mathrm{N}, \mathrm{D}+1\) )

\section*{property order}

Defines an ordering of the lattice sites, thus mapping the lattice to a 1 D chain.
Each row of the array contains the lattice indices for one site, the order of the rows thus specifies a path through the lattice, along which an MPS will wind through through the lattice.
You can visualize the order with plot_order().
mps2lat_idx (i)
Translate MPS index \(i\) to lattice indices ( \(\mathrm{x} \_0\), ..., \(\left.\mathrm{x} \_\{\operatorname{dim}-1\}, u\right)\).
Parameters i (int / array_like of int)-MPS index/indices.
Returns lat_idx - First dimensions like \(i\), last dimension has len dim \({ }^{`}+1\) and contains the lattice indices " \(\left(x \_0, \ldots, x_{\_}\{d i m-1\}, u\right)\) ` corresponding to \(i\). For \(i\) accross the MPS unit cell and "infinite" \(b c_{-} M P S\), we shift \(x_{-} 0\) accordingly.

\section*{Return type array}
lat2mps_idx (lat_idx)
Translate lattice indices ( \(\mathrm{x} \_0, \ldots, \mathrm{x}_{-}\{\mathrm{D}-1\}, \mathrm{u}\) ) to MPS index \(i\).

Parameters lat_idx (array_like [.., dim+1]) - The last dimension corresponds to lattice indices ( \(\left.x_{-} 0, \ldots, x_{-}\{D-1\}, u\right)\). All lattice indices should be positive and smaller than the corresponding entry in self.shape. Exception: for "infinite" \(b c_{-} M P S\), an \(x \_0\) outside indicates shifts accross the boundary.
Returns i - MPS index/indices corresponding to lat_idx. Has the same shape as lat_idx without the last dimension.

\section*{Return type array_like}
mps2lat_values (*args, **kwargs)
Not implemented, use mps2lat_values_masked() instead.
mps2lat_values_masked (*args, **kwargs)
Reshape/reorder an array \(A\) to replace an MPS index by lattice indices.
This is a generalization of mps2lat_values () allowing for the case of an arbitrary set of MPS indices present in each axis of \(A\).

\section*{Parameters}
- A (ndarray) - Some values.
- axes ( (iterable of) int) - Chooses the axis of \(A\) which should be replaced. If multiple axes are given, you also need to give multiple index arrays as mps_inds.
- mps_inds ((list of) 1D ndarray) - Specifies for each axis in axes, for which MPS indices we have values in the corresponding axis of A. Defaults to [np. arange (A. shape[ax]) for ax in axes]. For indices accross the MPS unit cell and "infinite" \(b c_{-} M P S\), we shift \(x_{-} 0\) accordingly.
- include_u ((list of) bool) - Specifies for each axis in axes, whether the \(u\) index of the lattice should be included into the output array res_A. Defaults to len (self. unit_cell) > 1 .

Returns res_A - Reshaped and reordered copy of A. Such that MPS indices along the specified axes are replaced by lattice indices, i.e., if MPS index \(j\) maps to lattice site ( \(x 0, x 1, x 2\) ), then res_A[..., x0, x1, x2, ...] = A[..., mps_inds[j], ...].

\section*{Return type np.ma.MaskedArray}
```

enlarge_mps_unit_cell(factor=2)

```

Repeat the unit cell for infinite MPS boundary conditions; in place.
Parameters factor (int) - The new number of sites in the MPS unit cell will be increased from \(N_{\text {_s }}\) sites to factor*N_sites_per_ring. Since MPS unit cells are repeated in the \(x\)-direction in our convetion, the lattice shape goes from ( \(\mathrm{Lx}, \mathrm{Ly}, \ldots, \mathrm{Lu}\) ) to (Lx*factor, Ly, ..., Lu).
possible_couplings ( \(u 1, u 2, d x\), strength=None)
Find possible MPS indices for two-site couplings.
For periodic boundary conditions ( bc [a] == False) the index x _a is taken modulo Ls [a] and runs through range (Ls [a]). For open boundary conditions, \(x_{-}\)a is limited to \(0<=x_{-}<\)Ls [a] and \(0<=x \_a+d x[a]<\) lat.Ls [a].

\section*{Parameters}
- u1 (int) - Indices within the unit cell; the \(u 1\) and \(u 2\) of add_coupling()
- u2 (int) - Indices within the unit cell; the \(u 1\) and \(u 2\) of add_coupling ()
- \(\mathbf{d x}\) (array) - Length dim. The translation in terms of basis vectors for the coupling.
- strength (array_like / None) - If given, instead of returning lat_indices and coupling_shape directly return the correct strength_12.

\section*{Returns}
- mps1, mps2 (1D array) - For each possible two-site coupling the MPS indices for the \(u 1\) and \(u 2\).
- strength_vals (1D array) - (Only returend if strength is not None.) Such that for (i, \(j, ~ s) ~ i n ~ z i p\left(m p s 1, ~ m p s 2, ~ s t r e n g t h \_v a l s\right): ~ i t e r a t e s ~ o v e r ~ a l l ~ p o s s i b l e ~ c o u-~\) plings with \(s\) being the strength of that coupling. Couplings where strength_vals \(==0\). are filtered out.
- lat_indices ( 2 D int array) - (Only returend if strength is None.) Rows of lat_indices correspond to entries of \(m p s l\) and \(m p s 2\) and contain the lattice indices of the "lower left corner" of the box containing the coupling.
- coupling_shape (tuple of int) - (Only returend if strength is None.) Len dim. The correct shape for an array specifying the coupling strength. lat_indices has only rows within this shape.

\section*{possible_multi_couplings (ops, strength=None)}

Generalization of possible_couplings () to couplings with more than 2 sites.
Parameters ops (list of (opname, \(\mathrm{dx}, \mathrm{u})\) ) - Same as the argument ops of add_multi_coupling().

\section*{Returns}
- mps_ijkl ( 2 D int array) - Each row contains MPS indices \(i, j, k, l, \ldots\) for each of the operators positions. The positions are defined by \(d x\) ( \(\mathrm{j}, \mathrm{k}, 1, \ldots\) relative to \(i\) ) and boundary coundary conditions of self (how much the box for given \(d x\) can be shifted around without hitting a boundary - these are the different rows).
- strength_vals ( \(1 D\) array) - (Only returend if strength is not None.) Such that for (ijkl, s) in zip(mps_ijkl, strength_vals): iterates over all possible couplings with \(s\) being the strength of that coupling. Couplings where strength_vals \(==0\). are filtered out.
- lat_indices ( \(2 D\) int array) - (Only returend if strength is None.) Rows of lat_indices correspond to rows of \(m p s \_i j k l\) and contain the lattice indices of the "lower left corner" of the box containing the coupling.
- coupling_shape (tuple of int) - (Only returend if strength is None.) Len dim. The correct shape for an array specifying the coupling strength. lat_indices has only rows within this shape.
plot_coupling (ax, coupling=None, wrap=True, **kwargs)
Plot lines connecting nearest neighbors of the lattice.

\section*{Parameters}
- ax (matplotlib. axes.Axes) - The axes on which we should plot.
- coupling (list of (ul, \(u 2, d x)\) ) - By default (None), use self. pairs['nearest_neighbors']. Specifies the connections to be plotted; iteating over lattice indices ( \(i 0, i l, \ldots\) ), we plot a connection from the site (i0, i1, ..., u1) to the site ( \(i 0+d x[0]\), \(i 1+d x[1], \ldots, u 2\) ), taking into account the boundary conditions.
- wrap (bool) - If True, plot couplings going around the boundary by directly connecting the sites it connects. This might be hard to see, as this puts lines from one end of the lattice to the other. If False, plot the couplings as dangling lines.
- **kwargs - Further keyword arguments given to ax. plot ().
property boundary_conditions
Human-readable list of boundary conditions from bc and bc_shift.
Returns boundary_conditions - List of "open" or "periodic", one entry for each direction of the lattice.

Return type list of str
count_neighbors ( \(u=0\), key='nearest_neighbors')
Count e.g. the number of nearest neighbors for a site in the bulk.

\section*{Parameters}
- u(int) - Specifies the site in the unit cell, for which we should count the number of neighbors (or whatever key specifies).
- key (str) - Key of pairs to select what to count.

Returns number - Number of nearest neighbors (or whatever key specified) for the \(u\)-th site in the unit cell, somewhere in the bulk of the lattice. Note that it might not be the correct value at the edges of a lattice with open boundary conditions.

\section*{Return type int}

\section*{coupling_shape ( \(d x\) )}

Calculate correct shape of the strengths for a coupling.
Parameters \(\mathbf{d x}\) (tuple of int) - Translation vector in the lattice for a coupling of two operators. Corresponds to \(d x\) argument of tenpy.models.model. CouplingModel. add_multi_coupling().

\section*{Returns}
- coupling_shape (tuple of int) - Len dim. The correct shape for an array specifying the coupling strength. lat_indices has only rows within this shape.
- shift_lat_indices (array) - Translation vector from origin to the lower left corner of box spanned by \(d x\).

\section*{property dim}

The dimension of the lattice.
distance ( \(u 1, u 2, d x\) )
Get the distance for a given coupling between two sites in the lattice.
The \(u 1, u 2, d x\) parameters are defined in analogy with add_coupling(), i.e., this function calculates the distance between a pair of operators added with add_coupling (using the basis and unit_cell_positions of the lattice).

Warning: This function ignores "wrapping" arround the cylinder in the case of periodic boundary conditions.

\section*{Parameters}
- u1 (int) - Indices within the unit cell; the \(u 1\) and \(u 2\) of add_coupling()
- u2 (int) - Indices within the unit cell; the \(u 1\) and \(u 2\) of add_coupling()
- \(\mathbf{d} \mathbf{x}\) (array) - Length dim. The translation in terms of basis vectors for the coupling.

Returns distance - The distance between site at lattice indices [ \(\mathrm{x}, \mathrm{y}, \mathrm{u} 1\) ] and [ \(\mathrm{x}+\) \(d x[0], y+d x[1], u 2]\), ignoring any boundary effects.
Return type float
find_coupling_pairs (max_dx=3, cutoff=None, eps=1e-10)
Automatically find coupling pairs grouped by distances.
Given the unit_cell_positions and basis, the coupling pairs of nearest_neighbors, next_nearest_neighbors etc at a given distance are basically fixed (although not uniquely, since we take out half of them to avoid double-counting couplings in both directions A_i B_j + B_i A_i). This function iterates through all possible couplings up to a given cutoff distance and then determines the possible pairs at fixed distances (up to round-off errors).

\section*{Parameters}
- max_dx (int) - Maximal index for each index of \(d x\) to iterate over. You need large enough values to include every possible coupling up to the desired distance, but choosing it too large might make this function run for a long time.
- cutoff (float) - Maximal distance (in the units in which basis and unit_cell_positions is given).
- eps (float) - Tolerance up to which to distances are considered the same.

Returns coupling_pairs - Keys are distances of nearest-neighbors, next-nearest-neighbors etc.
Values are [(u1, u2, dx), ...] as in pairs.
Return type dict
mps_idx_fix_u(u=None)
return an index array of MPS indices for which the site within the unit cell is \(u\).
If you have multiple sites in your unit-cell, an onsite operator is in general not defined for all sites. This functions returns an index array of the mps indices which belong to sites given by self. unit_cell[u].

Parameters u(None / int) - Selects a site of the unit cell. None (default) means all sites.
Returns mps_idx - MPS indices for which self.site(i) is self.unit_cell[u]. Ordered ascending.
Return type array
mps_lat_idx_fix_u (u=None)
Similar as mps_idx_fix_u(), but return also the corresponding lattice indices.
Parameters u(None / int) - Selects a site of the unit cell. None (default) means all sites.

\section*{Returns}
- mps_idx (array) - MPS indices \(i\) for which self.site(i) is self. unit_cell[u].
- lat_idx (2D array) - The row \(j\) contains the lattice index (without \(u\) ) corresponding to mps_idx[j].
mps_sites()
Return a list of sites for all MPS indices.
Equivalent to [self.site(i) for i in range(self.N_sites)].
This should be used for sites of 1D tensor networks (MPS, MPO,...).
multi_coupling_shape ( \(d x\) )
Calculate correct shape of the strengths for a multi_coupling.
Parameters dx (2D array, shape (N_ops, dim)) - dx[i, :] is the translation vector in the lattice for the \(i\)-th operator. Corresponds to the \(d x\) of each operator given in the argument ops of tenpy.models.model.CouplingModel.add_multi_coupling().

\section*{Returns}
- coupling_shape (tuple of int) - Len dim. The correct shape for an array specifying the coupling strength. lat_indices has only rows within this shape.
- shift_lat_indices (array) - Translation vector from origin to the lower left corner of box spanned by \(d x\). (Unlike for coupling_shape () it can also contain entries \(>0\) )
number_nearest_neighbors ( \(u=0\) )
Deprecated.
Deprecated since version 0.5.0: Use count_neighbors () instead.
number_next_nearest_neighbors ( \(u=0\) )
Deprecated.
Deprecated since version 0.5.0: Use count_neighbors () instead.
plot_basis \((a x\), origin \(=(0.0,0.0)\), shade \(=\) None, \(* * k w a r g s)\)
Plot arrows indicating the basis vectors of the lattice.

\section*{Parameters}
- ax (matplotlib.axes.Axes) - The axes on which we should plot.
- **kwargs - Keyword arguments for ax. arrow.
plot_bc_identified (ax, direction=- 1, origin=None, cylinder_axis=False, **kwargs)
Mark two sites indified by periodic boundary conditions.
Works only for lattice with a 2 -dimensional basis.

\section*{Parameters}
- ax (matplotlib.axes.Axes) - The axes on which we should plot.
- direction (int) - The direction of the lattice along which we should mark the idenitified sites. If None, mark it along all directions with periodic boundary conditions.
- cylinder_axis (bool) - Whether to plot the cylinder axis as well.
- origin (None | np.ndarray) - The origin starting from where we mark the identified sites. Defaults to the first entry of unit_cell_positions.
- **kwargs - Keyword arguments for the used ax. plot.
plot_order (ax, order=None, textkwargs=\{'color': 'r'\}, **kwargs)
Plot a line connecting sites in the specified "order" and text labels enumerating them.

\section*{Parameters}
- ax (matplotlib.axes.Axes) - The axes on which we should plot.
- order (None / 2D array (self.N_sites, self.dim+1))- The order as returned by ordering (); by default (None) use order.
- textkwargs (None I dict) - If not None, we add text labels enumerating the sites in the plot. The dictionary can contain keyword arguments for ax.text ().
- **kwargs - Further keyword arguments given to ax.plot ().
plot_sites (ax, markers=['o', ' \({ }^{\prime}\) ', 's', ' \(\left.p^{\prime}, h^{\prime} h^{\prime}, D^{\prime}\right]\), **kwargs)
Plot the sites of the lattice with markers.

\section*{Parameters}
- ax (matplotlib. axes.Axes) - The axes on which we should plot.
- markers (Iist) - List of values for the keywork marker of ax.plot () to distinguish the different sites in the unit cell, a site \(u\) in the unit cell is plotted with a marker markers[u \% len(markers)].
- **kwargs - Further keyword arguments given to ax.plot().
position(lat_idx)
return 'space' position of one or multiple sites.
Parameters lat_idx (ndarray, (... , dim+1))-Lattice indices.
Returns pos - The position of the lattice sites specified by lat_idx in real-space.
Return type ndarray, (..., dim)
site ( \(i\) )
return Site instance corresponding to an MPS index \(i\)
```

test_sanity()

```

Sanity check.
Raises ValueErrors, if something is wrong.

\subsection*{19.1.3 Honeycomb}
- full name: tenpy.models.lattice.Honeycomb
- parent module: tenpy.models.lattice
- type: class

\section*{Inheritance Diagram}


\section*{Methods}
\begin{tabular}{|c|c|}
\hline Honeycomb.__init__(Lx, Ly, sites, **kwargs) & Initialize self. \\
\hline Honeycomb.count_neighbors([u, key]) & Count e.g. \\
\hline Honeycomb.coupling_shape(dx) & Calculate correct shape of the strengths for a coupling. \\
\hline Honeycomb.distance(u1, u2, dx) & Get the distance for a given coupling between two sites in the lattice. \\
\hline Honeycomb.enlarge_mps_unit_cell([factor]) & Repeat the unit cell for infinite MPS boundary conditions; in place. \\
\hline Honeycomb.find_coupling_pairs([max_dx, ...]) & Automatically find coupling pairs grouped by distances. \\
\hline Honeycomb.from_hdf5(hdf5_loader, h5gr, subpath) & Load instance from a HDF5 file. \\
\hline Honeycomb.lat2mps_idx(lat_idx) & Translate lattice indices \(\left(x \_0, \ldots, x_{-}\{D-1\}\right.\), u) to MPS index \(i\). \\
\hline Honeycomb.mps2lat_idx(i) & Translate MPS index \(i\) to lattice indices ( \(x \_0\), ..., x_\{dim-1\}, u). \\
\hline Honeycomb.mps2lat_values(A[, axes, u]) & Reshape/reorder \(A\) to replace an MPS index by lattice indices. \\
\hline ```
Honeycomb.mps2lat_values_masked(A[,
axes,...])
``` & Reshape/reorder an array \(A\) to replace an MPS index by lattice indices. \\
\hline Honeycomb.mps_idx_fix_u([u]) & return an index array of MPS indices for which the site within the unit cell is \(u\). \\
\hline Honeycomb.mps_lat_idx_fix_u([u]) & Similar as mps_idx_fix_u(), but return also the corresponding lattice indices. \\
\hline Honeycomb.mps_sites() & Return a list of sites for all MPS indices. \\
\hline Honeycomb.multi_coupling_shape(dx) & Calculate correct shape of the strengths for a multi_coupling. \\
\hline Honeycomb.number_nearest_neighbors([u]) & Deprecated. \\
\hline Honeycomb.number_next_nearest_neighbor & (Dep)recated. \\
\hline Honeycomb.ordering(order) & Provide possible orderings of the \(N\) lattice sites. \\
\hline Honeycomb.plot_basis(ax[, origin, shade]) & Plot arrows indicating the basis vectors of the lattice. \\
\hline Honeycomb.plot_bc_identified(ax[,..]) & Mark two sites indified by periodic boundary conditions. \\
\hline Honeycomb.plot_coupling(ax[, coupling, wrap]) & Plot lines connecting nearest neighbors of the lattice. \\
\hline Honeycomb.plot_order(ax[, order, textkwargs]) & Plot a line connecting sites in the specified "order" and text labels enumerating them. \\
\hline Honeycomb.plot_sites(ax[, markers]) & Plot the sites of the lattice with markers. \\
\hline Honeycomb.position(lat_idx) & return 'space' position of one or multiple sites. \\
\hline ```
Honeycomb.possible_couplings(u1, u2, dx[,
...])
``` & Find possible MPS indices for two-site couplings. \\
\hline Honeycomb.possible_multi_couplings(ops[, ...]) & Generalization of possible_couplings() to couplings with more than 2 sites. \\
\hline Honeycomb.save_hdf5(hdf5_saver, h5gr, subpath) & Export self into a HDF5 file. \\
\hline Honeycomb.site(i) & return Site instance corresponding to an MPS index \(i\) \\
\hline Honeycomb.test_sanity() & Sanity check. \\
\hline
\end{tabular}

\section*{Class Attributes and Properties}
\begin{tabular}{ll}
\hline Honeycomb. Lu & \begin{tabular}{l} 
the (expected) number of sites in the unit cell, \\
len (unit_cell).
\end{tabular} \\
\hline Honeycomb.boundary_conditions & \begin{tabular}{l} 
Human-readable list of boundary conditions from bc \\
and bc_shift.
\end{tabular} \\
\hline Honeycomb.dim & the dimension of the lattice \\
\hline Honeycomb.fifth_nearest_neighbors & \\
\hline Honeycomb.fourth_nearest_neighbors & \\
\hline Honeycomb. next_nearest_neighbors & \\
\hline Honeycomb. next_next_nearest_neighbors & Defines an ordering of the lattice sites, thus mapping the \\
\hline
\end{tabular}
class tenpy.models.lattice. Honeycomb (Lx, Ly, sites, **kwargs)
Bases: tenpy.models.lattice.Lattice
A honeycomb lattice.

\section*{Parameters}
- Lx (int) - The length in each direction.
- Ly (int) - The length in each direction.
- sites ((list of) Site) - The two local lattice sites making the unit_cell of the Lattice. If only a single Site is given, it is used for both sites.
- **kwargs - Additional keyword arguments given to the Lattice. basis, pos and pairs are set accordingly. For the Honeycomb lattice 'fourth_nearest_neighbors', 'fifth_nearest_neighbors' are set in pairs.
\(\operatorname{dim}=2\)
the dimension of the lattice
\(\mathrm{Lu}=2\)
the (expected) number of sites in the unit cell, len (unit_cell).
ordering (order)
Provide possible orderings of the \(N\) lattice sites.
The following orders are defined in this method compared to Lattice.ordering():
\begin{tabular}{|l|l|l|}
\hline order & equivalent priority & equivalent snake_winding \\
\hline 'default' & \((0,2,1)\) & (False, False, False) \\
\hline 'snake' & \((0,2,1)\) & (False, True, False) \\
\hline
\end{tabular}
property boundary_conditions
Human-readable list of boundary conditions from bc and bc_shift.



Returns boundary_conditions - List of "open" or "periodic", one entry for each direction of the lattice.

Return type list of str
count_neighbors ( \(u=0\), key='nearest_neighbors')
Count e.g. the number of nearest neighbors for a site in the bulk.

\section*{Parameters}
- u (int) - Specifies the site in the unit cell, for which we should count the number of neighbors (or whatever key specifies).
- key (str) - Key of pairs to select what to count.

Returns number - Number of nearest neighbors (or whatever key specified) for the \(u\)-th site in the unit cell, somewhere in the bulk of the lattice. Note that it might not be the correct value at the edges of a lattice with open boundary conditions.

\section*{Return type int}

\section*{coupling_shape ( \(d x\) )}

Calculate correct shape of the strengths for a coupling.
Parameters \(\mathbf{d x}\) (tuple of int) - Translation vector in the lattice for a coupling of two operators. Corresponds to \(d x\) argument of tenpy.models.model. CouplingModel. add_multi_coupling().

\section*{Returns}
- coupling_shape (tuple of int) - Len dim. The correct shape for an array specifying the coupling strength. lat_indices has only rows within this shape.
- shift_lat_indices (array) - Translation vector from origin to the lower left corner of box spanned by \(d x\).
distance ( \(u 1, u 2, d x\) )
Get the distance for a given coupling between two sites in the lattice.
The \(u 1, u 2, d x\) parameters are defined in analogy with add_coupling(), i.e., this function calculates the distance between a pair of operators added with add_coupling (using the basis and unit_cell_positions of the lattice).

Warning: This function ignores "wrapping" arround the cylinder in the case of periodic boundary conditions.

\section*{Parameters}
- u1 (int) - Indices within the unit cell; the \(u 1\) and \(u 2\) of add_coupling ()
- u2 (int) - Indices within the unit cell; the \(u 1\) and \(u 2\) of add_coupling()
- \(\mathbf{d x}\) (array) - Length dim. The translation in terms of basis vectors for the coupling.

Returns distance - The distance between site at lattice indices [x, y, ul] and [x + \(d x[0], y+d x[1], u 2]\), ignoring any boundary effects.

\section*{Return type float}
```

enlarge_mps_unit_cell(factor=2)

```

Repeat the unit cell for infinite MPS boundary conditions; in place.

Parameters factor (int) - The new number of sites in the MPS unit cell will be increased from \(N\) _sites to factor \(* N\) _sites_per_ring. Since MPS unit cells are repeated in the \(x\)-direction in our convetion, the lattice shape goes from ( \(\mathrm{Lx}, \mathrm{Ly}, \ldots, \mathrm{Lu}\) ) to (Lx*factor, Ly, ..., Lu).
find_coupling_pairs (max_dx=3, cutoff=None, eps=1e-10)
Automatically find coupling pairs grouped by distances.
Given the unit_cell_positions and basis, the coupling pairs of nearest_neighbors, next_nearest_neighbors etc at a given distance are basically fixed (although not uniquely, since we take out half of them to avoid double-counting couplings in both directions A_i B_j + B_i A_i). This function iterates through all possible couplings up to a given cutoff distance and then determines the possible pairs at fixed distances (up to round-off errors).

\section*{Parameters}
- max_dx (int) - Maximal index for each index of \(d x\) to iterate over. You need large enough values to include every possible coupling up to the desired distance, but choosing it too large might make this function run for a long time.
- cutoff (float) - Maximal distance (in the units in which basis and unit_cell_positions is given).
- eps (float) - Tolerance up to which to distances are considered the same.

Returns coupling_pairs - Keys are distances of nearest-neighbors, next-nearest-neighbors etc.
Values are [(u1, u2, dx), ...] as in pairs.
Return type dict
classmethod from_hdf5 (hdf5_loader, h5gr, subpath)
Load instance from a HDF5 file.
This method reconstructs a class instance from the data saved with save_hdf5 ().

\section*{Parameters}
- hdf5_loader (Hdf5Loader) - Instance of the loading engine.
- h5gr (Group) - HDF5 group which is represent the object to be constructed.
- subpath (str) - The name of \(h 5 g r\) with a '/' in the end.

Returns obj - Newly generated class instance containing the required data.
Return type cls
lat2mps_idx (lat_idx)
Translate lattice indices ( \(\mathrm{x} \_0, \ldots, \mathrm{x}_{-}\{\mathrm{D}-1\}, \mathrm{u}\) ) to MPS index \(i\).
Parameters lat_idx (array_like [.., dim+1]) - The last dimension corresponds to lattice indices \(\left(x_{-} 0, \ldots, x_{-}\{D-1\}, u\right)\). All lattice indices should be positive and smaller than the corresponding entry in self.shape. Exception: for "infinite" \(b c_{-} M P S\), an \(x \_0\) outside indicates shifts accross the boundary.

Returns i - MPS index/indices corresponding to lat_idx. Has the same shape as lat_idx without the last dimension.
Return type array_like
mps2lat_idx(i)
Translate MPS index \(i\) to lattice indices ( \(\mathrm{x} \_0\), ..., \(\left.\mathrm{x}_{-}\{\operatorname{dim}-1\}, u\right)\).
Parameters i(int | array_like of int)-MPS index/indices.

Returns lat_idx - First dimensions like \(i\), last dimension has len dim +1 and contains the lattice indices \({ }^{`}\left(x_{-} 0, \ldots, x_{\_}\{d i m-1\}, u\right)\) ` corresponding to \(i\). For \(i\) accross the MPS unit cell and "infinite" \(b c_{-} M P S\), we shift \(x_{-} 0\) accordingly.

\section*{Return type array}
mps2lat_values \((A\), axes \(=0, u=\) None \()\)
Reshape/reorder \(A\) to replace an MPS index by lattice indices.

\section*{Parameters}
- A (ndarray) - Some values. Must have A.shape [axes] = self.N_sites if \(u\) is None, or A.shape[axes] = self. N_cells if \(u\) is an int.
- axes ( (iterable of) int) - chooses the axis which should be replaced.
- u (None I int) - Optionally choose a subset of MPS indices present in the axes of \(A\), namely the indices corresponding to self.unit_cell[u], as returned by mps_idx_fix_u(). The resulting array will not have the additional dimension(s) of u.

Returns res_A - Reshaped and reordered verions of A. Such that MPS indices along the specified axes are replaced by lattice indices, i.e., if MPS index \(j\) maps to lattice site ( \(x 0, x 1, x 2\) ), then res_A[..., \(x 0, x 1, x 2, \ldots]=A[\ldots, j, \ldots]\).

Return type ndarray

\section*{Examples}

Say you measure expection values of an onsite term for an MPS, which gives you an 1D array \(A\), where \(A[i]\) is the expectation value of the site given by self.mps2lat_idx(i). Then this function gives you the expectation values ordered by the lattice:
```

>>> print(lat.shape, A.shape)
(10, 3, 2) (60,)
>>> A_res = lat.mps2lat_values(A)
>>> A_res.shape
(10, 3, 2)
>>> A_res[tuple(lat.mps2lat_idx(5))] == A[5]
True

```

If you have a correlation function C [i, j], it gets just slightly more complicated:
```

>>> print(lat.shape, C.shape)
(10, 3, 2) (60, 60)
>>> lat.mps2lat_values(C, axes=[0, 1]).shape
(10, 3, 2, 10, 3, 2)

```

If the unit cell consists of different physical sites, an onsite operator might be defined only on one of the sites in the unit cell. Then you can use mps_idx_fix_u() to get the indices of sites it is defined on, measure the operator on these sites, and use the argument \(u\) of this function.
```

>>> u = 0
>>> idx_subset = lat.mps_idx_fix_u(u)
>>> A_u = A[idx_subset]
>>> A_u_res = lat.mps2lat_values(A_u, u=u)
>>> A_u_res.shape
(10, 3)
>>> np.all(A_res[:, :, u] == A_u_res[:, :])
True

```
mps2lat_values_masked ( \(A\), axes=- 1, mps_inds=None, include_u=None)
Reshape/reorder an array \(A\) to replace an MPS index by lattice indices.
This is a generalization of mps2lat_values () allowing for the case of an arbitrary set of MPS indices present in each axis of \(A\).

\section*{Parameters}
- A (ndarray) - Some values.
- axes ((iterable of) int) - Chooses the axis of \(A\) which should be replaced. If multiple axes are given, you also need to give multiple index arrays as mps_inds.
- mps_inds ((Iist of) 1D ndarray) - Specifies for each axis in axes, for which MPS indices we have values in the corresponding axis of \(A\). Defaults to [np. arange (A. shape[ax]) for ax in axes]. For indices accross the MPS unit cell and "infinite" \(b c_{-} M P S\), we shift \(x \_0\) accordingly.
- include_u ((list of) bool) - Specifies for each axis in axes, whether the \(u\) index of the lattice should be included into the output array res_A. Defaults to len (self. unit_cell) > 1 .

Returns res_A - Reshaped and reordered copy of A. Such that MPS indices along the specified axes are replaced by lattice indices, i.e., if MPS index \(j\) maps to lattice site ( \(x 0, x 1, x 2\) ), then

Return type np.ma.MaskedArray
mps_idx_fix_u (u=None)
return an index array of MPS indices for which the site within the unit cell is \(u\).
If you have multiple sites in your unit-cell, an onsite operator is in general not defined for all sites. This functions returns an index array of the mps indices which belong to sites given by self. unit_cell[u].

Parameters \(\mathbf{u}\) (None / int) - Selects a site of the unit cell. None (default) means all sites.
Returns mps_idx - MPS indices for which self.site(i) is self.unit_cell[u]. Ordered ascending.

\section*{Return type array}
mps_lat_idx_fix_u ( \(u=\) None)
Similar as mps_idx_fix_u(), but return also the corresponding lattice indices.
Parameters \(\mathbf{u}\) (None / int) - Selects a site of the unit cell. None (default) means all sites.

\section*{Returns}
- mps_idx (array) - MPS indices \(i\) for which self.site(i) is self. unit_cell[u].
- lat_idx ( \(2 D\) array) - The row \(j\) contains the lattice index (without \(u\) ) corresponding to mps_idx[j].
mps_sites()
Return a list of sites for all MPS indices.
Equivalent to [self.site(i) for i in range(self.N_sites)].
This should be used for sites of 1D tensor networks (MPS, MPO,...).
multi_coupling_shape ( \(d x\) )
Calculate correct shape of the strengths for a multi_coupling.
Parameters dx (2D array, shape (N_ops, dim)) - dx[i, :] is the translation vector in the lattice for the \(i\)-th operator. Corresponds to the \(d x\) of each operator given in the argument ops of tenpy.models.model.CouplingModel.add_multi_coupling().

\section*{Returns}
- coupling_shape (tuple of int) - Len dim. The correct shape for an array specifying the coupling strength. lat_indices has only rows within this shape.
- shift_lat_indices (array) - Translation vector from origin to the lower left corner of box spanned by \(d x\). (Unlike for coupling_shape () it can also contain entries \(>0\) )
number_nearest_neighbors ( \(u=0\) )
Deprecated.
Deprecated since version 0.5.0: Use count_neighbors () instead.
number_next_nearest_neighbors \((u=0)\)
Deprecated.
Deprecated since version 0.5.0: Use count_neighbors () instead.
property order
Defines an ordering of the lattice sites, thus mapping the lattice to a 1D chain.
Each row of the array contains the lattice indices for one site, the order of the rows thus specifies a path through the lattice, along which an MPS will wind through through the lattice.
You can visualize the order with plot_order ().
plot_basis (ax, origin=(0.0, 0.0), shade=None, **kwargs)
Plot arrows indicating the basis vectors of the lattice.

\section*{Parameters}
- ax (matplotlib.axes.Axes) - The axes on which we should plot.
- **kwargs - Keyword arguments for ax. arrow.
plot_bc_identified (ax, direction=- 1, origin=None, cylinder_axis=False, **kwargs)
Mark two sites indified by periodic boundary conditions.
Works only for lattice with a 2 -dimensional basis.

\section*{Parameters}
- ax (matplotlib.axes. Axes) - The axes on which we should plot.
- direction (int) - The direction of the lattice along which we should mark the idenitified sites. If None, mark it along all directions with periodic boundary conditions.
- cylinder_axis (bool) - Whether to plot the cylinder axis as well.
- origin (None | np.ndarray) - The origin starting from where we mark the identified sites. Defaults to the first entry of unit_cell_positions.
- **kwargs - Keyword arguments for the used ax.plot.
plot_coupling (ax, coupling=None, wrap=False, **kwargs)
Plot lines connecting nearest neighbors of the lattice.

\section*{Parameters}
- ax (matplotlib. axes.Axes) - The axes on which we should plot.
- coupling (list of (u1, \(u 2, d x)\) ) - By default (None), use self. pairs['nearest_neighbors']. Specifies the connections to be plotted; iteating over lattice indices ( \(i 0, i 1, \ldots\) ), we plot a connection from the site (i0, i1, ..., u1) to the site (i0 \(+\mathrm{dx}[0]\), \(i 1+d x[1], \ldots, u 2\) ), taking into account the boundary conditions.
- wrap (bool) - If True, plot couplings going around the boundary by directly connecting the sites it connects. This might be hard to see, as this puts lines from one end of the lattice to the other. If False, plot the couplings as dangling lines.
- **kwargs - Further keyword arguments given to ax.plot ().
plot_order (ax, order=None, textkwargs=\{'color': 'r'\}, **kwargs)
Plot a line connecting sites in the specified "order" and text labels enumerating them.

\section*{Parameters}
- ax (matplotlib. axes.Axes) - The axes on which we should plot.
- order (None / 2D array (self.N_sites, self.dim+1))- The order as returned by ordering (); by default (None) use order.
- textkwargs (None I dict) - If not None, we add text labels enumerating the sites in the plot. The dictionary can contain keyword arguments for ax.text ().
- **kwargs - Further keyword arguments given to ax.plot ().
plot_sites (ax, markers=['o', '^', 's', 'p', 'h', 'D'], **kwargs)
Plot the sites of the lattice with markers.

\section*{Parameters}
- ax (matplotlib. axes.Axes) - The axes on which we should plot.
- markers (list) - List of values for the keywork marker of ax.plot () to distinguish the different sites in the unit cell, a site \(u\) in the unit cell is plotted with a marker markers [u \% len(markers)].
- **kwargs - Further keyword arguments given to ax.plot().
position (lat_idx)
return 'space' position of one or multiple sites.
Parameters lat_idx (ndarray, (... , dim+1))-Lattice indices.
Returns pos - The position of the lattice sites specified by lat_idx in real-space.
Return type ndarray, (..., dim)
possible_couplings ( \(u 1, u 2, d x\), strength=None)
Find possible MPS indices for two-site couplings.
For periodic boundary conditions ( \(\mathrm{bc}[\mathrm{a}]==\mathrm{False}\) ) the index x _a is taken modulo Ls [a] and runs through range (Ls [a]). For open boundary conditions, x_a is limited to \(0<=x \_a<\) Ls [a] and \(0<=x \_a+d x[a]<\) lat.Ls[a].

\section*{Parameters}
- u1 (int) - Indices within the unit cell; the \(u 1\) and \(u 2\) of add_coupling ()
- u2 (int) - Indices within the unit cell; the \(u 1\) and \(u 2\) of add_coupling()
- \(\mathbf{d x}\) (array) - Length dim. The translation in terms of basis vectors for the coupling.
- strength (array_like / None) - If given, instead of returning lat_indices and coupling_shape directly return the correct strength_12.

\section*{Returns}
- mps1, mps2 (ID array) - For each possible two-site coupling the MPS indices for the \(u\) l and \(u 2\).
- strength_vals (1D array) - (Only returend if strength is not None.) Such that for (i, j, s) in zip(mps1, mps2, strength_vals): iterates over all possible couplings with \(s\) being the strength of that coupling. Couplings where strength_vals \(==0\). are filtered out.
- lat_indices ( \(2 D\) int array) - (Only returend if strength is None.) Rows of lat_indices correspond to entries of mpsl and mps2 and contain the lattice indices of the "lower left corner" of the box containing the coupling.
- coupling_shape (tuple of int) - (Only returend if strength is None.) Len dim. The correct shape for an array specifying the coupling strength. lat_indices has only rows within this shape.
possible_multi_couplings (ops, strength=None)
Generalization of possible_couplings() to couplings with more than 2 sites.
Parameters ops (list of (opname, \(\mathrm{dx}, \mathrm{u})\) ) - Same as the argument ops of add_multi_coupling().

\section*{Returns}
- mps_ijkl (2D int array) - Each row contains MPS indices \(i, j, k, l, \ldots\). for each of the operators positions. The positions are defined by \(d x(\mathrm{j}, \mathrm{k}, \mathrm{l}, \ldots\) relative to \(i\) ) and boundary coundary conditions of self (how much the box for given \(d x\) can be shifted around without hitting a boundary - these are the different rows).
- strength_vals (1D array) - (Only returend if strength is not None.) Such that for (ijkl, s) in zip(mps_ijkl, strength_vals): iterates over all possible couplings with \(s\) being the strength of that coupling. Couplings where strength_vals \(=0\). are filtered out.
- lat_indices ( 2 D int array) - (Only returend if strength is None.) Rows of lat_indices correspond to rows of \(m p s \_i j k l\) and contain the lattice indices of the "lower left corner" of the box containing the coupling.
- coupling_shape (tuple of int) - (Only returend if strength is None.) Len dim. The correct shape for an array specifying the coupling strength. lat_indices has only rows within this shape.
save_hdf5 (hdf5_saver, h5gr, subpath)
Export self into a HDF5 file.
This method saves all the data it needs to reconstruct self with from_hdf5().
Specifically, it saves unit_cell, Ls, unit_cell_positions, basis, boundary_conditions, pairs under their name, bc_MPS as "boundary_conditions_MPS", and order as "order_for_MPS". Moreover, it saves dim and N_sites as HDF5 attributes.

\section*{Parameters}
- hdf5_saver (Hdf5Saver) - Instance of the saving engine.
- h5gr (: class `Group ') - HDF5 group which is supposed to represent self.
- subpath (str) - The name of \(h 5 g r\) with a ' / ' in the end.
site (i)
return Site instance corresponding to an MPS index \(i\)
```

test_sanity()

```

Sanity check.
Raises ValueErrors, if something is wrong.

\subsection*{19.1.4 IrregularLattice}
- full name: tenpy.models.lattice.IrregularLattice
- parent module: tenpy.models.lattice
- type: class

\section*{Inheritance Diagram}


\section*{Methods}
\begin{tabular}{|c|c|}
\hline IrregularLattice. \(\qquad\) init__(regular_lattice[, ...]) & Initialize self. \\
\hline IrregularLattice.count_neighbors([u, key]) & Count e.g. \\
\hline IrregularLattice.coupling_shape(dx) & Calculate correct shape of the strengths for a coupling. \\
\hline IrregularLattice.distance(u1, u2, dx) & Get the distance for a given coupling between two sites in the lattice. \\
\hline \[
\begin{aligned}
& \text { IrregularLattice. } \\
& \text { enlarge_mps_unit_cell([factor]) }
\end{aligned}
\] & Repeat the unit cell for infinite MPS boundary conditions; in place. \\
\hline \[
\begin{aligned}
& \text { IrregularLattice. } \\
& \text { find_coupling_pairs([...]) }
\end{aligned}
\] & Automatically find coupling pairs grouped by distances. \\
\hline IrregularLattice.from_hdf5(hdf5_loader, ...) & Load instance from a HDF5 file. \\
\hline IrregularLattice.lat2mps_idx(lat_idx) & Translate lattice indices (x_0, ..., \(x_{-}\{D-1\}\), u) to MPS index \(i\). \\
\hline IrregularLattice.mps2lat_idx(i) & Translate MPS index \(i\) to lattice indices ( \(x_{-} 0, \ldots\), x_\{dim-1\}, u). \\
\hline \[
\begin{aligned}
& \text { IrregularLattice.mps2lat_values(A[, } \\
& \text { axes, u]) }
\end{aligned}
\] & Reshape/reorder \(A\) to replace an MPS index by lattice indices. \\
\hline
\end{tabular}

Table 9 - continued from previous page
\begin{tabular}{|c|c|}
\hline \begin{tabular}{l}
IrregularLattice. \\
mps2lat_values_masked(A[,...])
\end{tabular} & Reshape/reorder an array \(A\) to replace an MPS index by lattice indices. \\
\hline IrregularLattice.mps_idx_fix_u([u]) & return an index array of MPS indices for which the site within the unit cell is \(u\). \\
\hline \[
\begin{aligned}
& \text { IrregularLattice. } \\
& \text { mps_lat_idx_fix_u([u]) }
\end{aligned}
\] & Similar as mps_idx_fix_u(), but return also the corresponding lattice indices. \\
\hline Irregularlattice.mps_sites() & Return a list of sites for all MPS indices. \\
\hline \[
\begin{aligned}
& \text { Irregularlattice. } \\
& \text { multi_coupling_shape(dx) }
\end{aligned}
\] & Calculate correct shape of the strengths for a multi_coupling. \\
\hline \begin{tabular}{l}
Irregularlattice. \\
number_nearest_neighbors([u])
\end{tabular} & Deprecated. \\
\hline Irregularlattice. number_next_nearest_neighbors([u]) & Deprecated. \\
\hline Irregularlattice.ordering(order) & Provide possible orderings of the lattice sites. \\
\hline IrregularLattice.plot_basis(ax[, origin, shade]) & Plot arrows indicating the basis vectors of the lattice. \\
\hline \[
\begin{aligned}
& \text { IrregularLattice. } \\
& \text { plot_bc_identified(ax[,...]) }
\end{aligned}
\] & Mark two sites indified by periodic boundary conditions. \\
\hline Irregularlattice.plot_coupling(ax[,..]) & Plot lines connecting nearest neighbors of the lattice. \\
\hline IrregularLattice.plot_order(ax[, order, ...]) & Plot a line connecting sites in the specified "order" and text labels enumerating them. \\
\hline IrregularLattice.plot_sites(ax[, markers]) & Plot the sites of the lattice with markers. \\
\hline IrregularLattice.position(lat_idx) & return 'space' position of one or multiple sites. \\
\hline ```
IrregularLattice.
possible_couplings(u1,u2, dx)
``` & Find possible MPS indices for two-site couplings. \\
\hline Irregularlattice. possible_multi_couplings(ops) & Generalization of possible_couplings() to couplings with more than 2 sites. \\
\hline IrregularLattice.save_hdf5(hdf5_saver, h5gr, ...) & Export self into a HDF5 file. \\
\hline IrregularLattice.site(i) & return Site instance corresponding to an MPS index \(i\) \\
\hline IrregularLattice.test_sanity() & Sanity check. \\
\hline
\end{tabular}

\section*{Class Attributes and Properties}
\begin{tabular}{ll}
\hline IrregularLattice.Lu & \begin{tabular}{l} 
the (expected) number of sites in the unit cell, \\
len(unit_cell).
\end{tabular} \\
\hline IrregularLattice.boundary_conditions & \begin{tabular}{l} 
Human-readable list of boundary conditions from bc \\
andbc_shift.
\end{tabular} \\
\hline IrregularLattice.dim & The dimension of the lattice. \\
\hline IrregularLattice.nearest_neighbors & \\
\hline \begin{tabular}{l} 
IrregularLattice. \\
next_nearest_neighbors
\end{tabular} & \begin{tabular}{l} 
IrregularLattice. \\
next_next_nearest_neighbors an ordering of the lattice sites, thus mapping the \\
lattice to a 1D chain.
\end{tabular} \\
\hline IrregularLattice.order &
\end{tabular}
class tenpy.models.lattice.IrregularLattice (regular_lattice, remove=None, add=None, add_unit_cell=[], add_positions=None)

Bases: tenpy.models.lattice.Lattice
A variant of a regular lattice, where we might have extra sites or sites missing.

Note: The lattice defines only the geometry of the sites, not the couplings; you can have position-dependent couplings/onsite terms despite having a regular lattice.

By adjusting the order and a few private attributes and methods, we can make functions like possible_couplings () work with a more "irregular" lattice structure, where some of the sites are missing and other sites added instead.

\section*{Parameters}
- regular_lattice (Lattice) - The lattice this is based on.
- remove ( 2 D array / None) - Each row is a lattice index (x_0, ..., \(\left.x_{-}\{\operatorname{dim}-1\}, u\right)\) of a site to be removed. If None, don't remove something.
- add (Tuple[2D array, list] / None) - Each row of the 2D array is a lattice index \(\left(x_{-} 0, \ldots, x_{-}\{\operatorname{dim}-1\}, u\right)\) specifiying where a site is to be added; \(u\) is the index of the site within the final unit_cell of the irregular lattice. For each row of the 2D array, there is one entry in the list specifying where the site is inserted in the MPS; the values are compared to the MPS indices of the regular lattice and sorted into it, so " 2.5 " goes between what was site 2 and 3 in the regular lattice. An entry None indicates that the site should be inserted after the lattice site ( \(x \_0, \ldots, x_{-}\{\operatorname{dim}-1\},-1\) ) of the regular_lattice.
- add_unit_cell (list of Site) - Extra sites to be added to the unit cell.
- add_positions (iterable of ID arrays) - For each extra site in add_unit_cell the position within the unit cell. Defaults to np.zeros ((len(add_unit_cell), dim)).
regular_lattice
The lattice this is based on.
Type Lattice
remove, add
See above. Used in ordering () only.
Type 2D array I None

\section*{Examples}

Let's imagine that we have two different sites; for concreteness we can thing of a fermion site, which we represent with ' F ', and a spin site ' S '. If you want to preserve charges, take a look at set_common_charges () for the proper way to initialize the sites.

You could now imagine that to have fermion chain with spins on the "bonds". In the periodic/infinite case, you would simply define
```

>>> lat = Lattice([2], unit_cell=['F', 'S'], bc='periodic', bc_MPS='infinite')
>>> lat.mps_sites()
['F', 'S', 'F', 'S']

```

For a finite system, you don't want to terminate with a spin on the right, so you need to remove the very last site by specifying the lattice index \([L-1,1]\) of that site:
```

>>> L = 4
>>> reg_lat = Lattice([L], unit_cell=['F', 'S'], bc='open', bc_MPS='finite')
>>> irr_lat = IrregularLattice(reg_lat, remove=[[L - 1, 1]])
>>> irr_lat.mps_sites()
['F', 'S', 'F', 'S', 'E', 'S', 'F']

```

Another simple example would be to add a spin in the center of a fermion chain. In that case, we add another site to the unit cell and specify the lattice index as \([(L-1) / / 2,1]\) (where the 1 is the index of ' \(S\) ' in the unit cell [' \(\mathrm{F}^{\prime}\), ' \(\mathrm{S}^{\prime}\) ] of the irregular lattice). The None for the MPS index is equivalent to ( \(\mathrm{L}-1\) )/2 in this case.
```

>>> reg_lat = Lattice([L], unit_cell=['F'])
>>> irr_lat = IrregularLattice(reg_lat, add=([[(L - 1)//2, 1]], [None]),
... add_unit_cell=['S'])
>>> irr_lat.mps_sites()
['F', 'F', 'S', 'F', 'F']

```
save_hdf5 (hdf5_saver, h5gr, subpath)
Export self into a HDF5 file.
This method saves all the data it needs to reconstruct self with from_hdf5 ().
Specifically, it saves unit_cell, Ls, unit_cell_positions, basis, boundary_conditions, pairs under their name, bc_MPS as "boundary_conditions_MPS", and order as "order_for_MPS". Moreover, it saves dim and N_sites as HDF5 attributes.

\section*{Parameters}
- hdf5_saver (Hdf5Saver) - Instance of the saving engine.
- h5gr (: class `Group`) - HDF5 group which is supposed to represent self.
- subpath (str) - The name of \(h 5 g r\) with a ' /' in the end.
classmethod from_hdf5 (hdf5_loader, h5gr, subpath)
Load instance from a HDF5 file.
This method reconstructs a class instance from the data saved with save_hdf5 ().

\section*{Parameters}
- hdf5_loader (Hdf5Loader) - Instance of the loading engine.
- h5gr (Group) - HDF5 group which is represent the object to be constructed.
- subpath (str) - The name of \(h 5 g r\) with a ' / ' in the end.

Returns obj - Newly generated class instance containing the required data.
Return type cls

\section*{ordering (order)}

Provide possible orderings of the lattice sites.
Parameters order - Argument for the Lattice.ordering() of the regular_lattice, or 2D ndarray providing the order of the regular lattice.

Returns order - The order to be used for order, i.e. order with added/removed sites as specified by remove and add.
Return type array, shape ( \(\mathrm{N}, \mathrm{D}+1\) )

\section*{property order}

Defines an ordering of the lattice sites, thus mapping the lattice to a 1 D chain.
Each row of the array contains the lattice indices for one site, the order of the rows thus specifies a path through the lattice, along which an MPS will wind through through the lattice.
You can visualize the order with plot_order ().
mps_idx_fix_u (u=None)
return an index array of MPS indices for which the site within the unit cell is \(u\).
If you have multiple sites in your unit-cell, an onsite operator is in general not defined for all sites. This functions returns an index array of the mps indices which belong to sites given by self. unit_cell[u].

Parameters u(None / int) - Selects a site of the unit cell. None (default) means all sites.
Returns mps_idx - MPS indices for which self.site(i) is self.unit_cell[u]. Ordered ascending.

\section*{Return type array}
property boundary_conditions
Human-readable list of boundary conditions from bc and bc_shift.
Returns boundary_conditions - List of "open" or "periodic", one entry for each direction of the lattice.

\section*{Return type list of str}
count_neighbors ( \(u=0\), key='nearest_neighbors')
Count e.g. the number of nearest neighbors for a site in the bulk.

\section*{Parameters}
- u(int) - Specifies the site in the unit cell, for which we should count the number of neighbors (or whatever key specifies).
- key (str) - Key of pairs to select what to count.

Returns number - Number of nearest neighbors (or whatever key specified) for the \(u\)-th site in the unit cell, somewhere in the bulk of the lattice. Note that it might not be the correct value at the edges of a lattice with open boundary conditions.

\section*{Return type int}
coupling_shape ( \(d x\) )
Calculate correct shape of the strengths for a coupling.
Parameters \(\mathbf{d x}\) (tuple of int) - Translation vector in the lattice for a coupling of two operators. Corresponds to \(d x\) argument of tenpy.models.model. CouplingModel. add_multi_coupling().

\section*{Returns}
- coupling_shape (tuple of int) - Len dim. The correct shape for an array specifying the coupling strength. lat_indices has only rows within this shape.
- shift_lat_indices (array) - Translation vector from origin to the lower left corner of box spanned by \(d x\).
property dim
The dimension of the lattice.
distance ( \(u 1, u 2, d x\) )
Get the distance for a given coupling between two sites in the lattice.
The \(u 1, u 2, d x\) parameters are defined in analogy with add_coupling(), i.e., this function calculates the distance between a pair of operators added with add_coupling (using the basis and unit_cell_positions of the lattice).

Warning: This function ignores "wrapping" arround the cylinder in the case of periodic boundary conditions.

\section*{Parameters}
- u1 (int) - Indices within the unit cell; the \(u 1\) and \(u 2\) of add_coupling()
- u2 (int) - Indices within the unit cell; the \(u 1\) and \(u 2\) of add_coupling()
- \(\mathbf{d x}\) (array) - Length dim. The translation in terms of basis vectors for the coupling.

Returns distance - The distance between site at lattice indices [ \(\mathrm{x}, \mathrm{y}, \mathrm{ul}\) ] and [ \(\mathrm{x}+\) \(d x[0], y+d x[1], u 2]\), ignoring any boundary effects.
Return type float
enlarge_mps_unit_cell(factor=2)
Repeat the unit cell for infinite MPS boundary conditions; in place.
Parameters factor (int) - The new number of sites in the MPS unit cell will be increased from \(N\) _sites to factor \(* N\) _sites_per_ring. Since MPS unit cells are repeated in the \(x\)-direction in our convetion, the lattice shape goes from ( \(L x, L y, \ldots, L u\) ) to (Lx*factor, Ly, ..., Lu).
find_coupling_pairs (max_dx=3, cutoff=None, eps=le-10)
Automatically find coupling pairs grouped by distances.
Given the unit_cell_positions and basis, the coupling pairs of nearest_neighbors, next_nearest_neighbors etc at a given distance are basically fixed (although not uniquely, since we take out half of them to avoid double-counting couplings in both directions A_i B_j + B_i A_i). This function iterates through all possible couplings up to a given cutoff distance and then determines the possible pairs at fixed distances (up to round-off errors).

\section*{Parameters}
- max_dx (int) - Maximal index for each index of \(d x\) to iterate over. You need large enough values to include every possible coupling up to the desired distance, but choosing it too large might make this function run for a long time.
- cutoff (float) - Maximal distance (in the units in which basis and unit_cell_positions is given).
- eps (float) - Tolerance up to which to distances are considered the same.

Returns coupling_pairs - Keys are distances of nearest-neighbors, next-nearest-neighbors etc. Values are \([(u 1, u 2, d x), \ldots]\) as in pairs.

Return type dict

\section*{lat2mps_idx (lat_idx)}

Translate lattice indices ( \(\mathrm{x} \_0, \ldots, \mathrm{x} \_\{\mathrm{D}-1\}, \mathrm{u}\) ) to MPS index \(i\).
Parameters lat_idx (array_like [.., dim+1]) - The last dimension corresponds to lattice indices \(\left(x_{-} 0, \ldots, x_{-}\{D-1\}, u\right)\). All lattice indices should be positive and
smaller than the corresponding entry in self.shape. Exception: for "infinite" \(b c_{-} M P S\), an \(x \_0\) outside indicates shifts accross the boundary.
Returns i-MPS index/indices corresponding to lat_idx. Has the same shape as lat_idx without the last dimension.

Return type array_like
mps2lat_idx (i)
Translate MPS index \(i\) to lattice indices ( \(\left.\mathrm{x} \_0, \ldots, \mathrm{x}_{-}\{\operatorname{dim}-1\}, u\right)\).
Parameters i (int | array_like of int)-MPS index/indices.
Returns lat_idx - First dimensions like \(i\), last dimension has len dim \({ }^{`}+1\) and contains the lattice indices " \(\left(x_{-} 0, \ldots, x_{\_}\{d i m-1\}, u\right)\) ' corresponding to \(i\). For \(i\) accross the MPS unit cell and "infinite" \(b c_{-} M P S\), we shift \(x_{-} 0\) accordingly.

\section*{Return type array}
mps2lat_values \((A\), axes \(=0, u=\) None \()\)
Reshape/reorder \(A\) to replace an MPS index by lattice indices.

\section*{Parameters}
- A (ndarray) - Some values. Must have A.shape [axes] = self. N_sites if \(u\) is None, or A.shape[axes] = self.N_cells if \(u\) is an int.
- axes ( (iterable of) int) - chooses the axis which should be replaced.
- u (None | int) - Optionally choose a subset of MPS indices present in the axes of \(A\), namely the indices corresponding to self.unit_cell[u], as returned by \(m p s \_i d x \_f i x \_u()\). The resulting array will not have the additional dimension(s) of \(u\).

Returns res_A - Reshaped and reordered verions of A. Such that MPS indices along the specified axes are replaced by lattice indices, i.e., if MPS index \(j\) maps to lattice site ( \(x 0, x 1, x 2\) ), then res_A[..., \(x 0, x 1, x 2, \ldots]=A[\ldots, j, \ldots]\).

Return type ndarray

\section*{Examples}

Say you measure expection values of an onsite term for an MPS, which gives you an 1D array \(A\), where \(A[i]\) is the expectation value of the site given by self.mps2lat_idx(i). Then this function gives you the expectation values ordered by the lattice:
```

>>> print(lat.shape, A.shape)
(10, 3, 2) (60,)
>>> A_res = lat.mps2lat_values(A)
>>> A_res.shape
(10, 3, 2)
>>> A_res[tuple(lat.mps2lat_idx(5))] == A[5]
True

```

If you have a correlation function C [i, j], it gets just slightly more complicated:
```

>>> print(lat.shape, C.shape)
(10, 3, 2) (60, 60)
>>> lat.mps2lat_values(C, axes=[0, 1]).shape
(10, 3, 2, 10, 3, 2)

```

If the unit cell consists of different physical sites, an onsite operator might be defined only on one of the sites in the unit cell. Then you can use mps_idx_fix_u() to get the indices of sites it is defined on, measure the operator on these sites, and use the argument \(u\) of this function.
```

>>> u = 0
>>> idx_subset = lat.mps_idx_fix_u(u)
>>> A__u = A[idx_subset]
>>> A_u_res = lat.mps2lat_values(A_u, u=u)
>>> A_u_res.shape
(10, 3)
>>> np.all(A_res[:, :, u] == A_u_res[:, :])
True

```
mps2lat_values_masked (A, axes=- \(1, m p s \_i n d s=N o n e\), include_u=None)
Reshape/reorder an array \(A\) to replace an MPS index by lattice indices.
This is a generalization of mps2lat_values () allowing for the case of an arbitrary set of MPS indices present in each axis of \(A\).

\section*{Parameters}
- A (ndarray) - Some values.
- axes ( (iterable of) int) - Chooses the axis of \(A\) which should be replaced. If multiple axes are given, you also need to give multiple index arrays as mps_inds.
- mps_inds ((list of) ID ndarray) - Specifies for each axis in axes, for which MPS indices we have values in the corresponding axis of \(A\). Defaults to [np. arange (A. shape[ax]) for ax in axes]. For indices accross the MPS unit cell and "infinite" \(b c_{-} M P S\), we shift \(x \_0\) accordingly.
- include_u ((list of) bool) - Specifies for each axis in axes, whether the \(u\) index of the lattice should be included into the output array res_A. Defaults to len (self. unit_cell) > 1 .

Returns res_A - Reshaped and reordered copy of A. Such that MPS indices along the specified axes are replaced by lattice indices, i.e., if MPS index \(j\) maps to lattice site \((x 0, x 1, x 2)\), then res_A[..., \(x 0, x 1, x 2, \ldots]=A\left[. . ., m p s \_i n d s[j], \ldots\right]\).

\section*{Return type np.ma.MaskedArray}
mps_lat_idx_fix_u ( \(u=\) None)
Similar as mps_idx_fix_u(), but return also the corresponding lattice indices.
Parameters u(None / int)-Selects a site of the unit cell. None (default) means all sites.

\section*{Returns}
- mps_idx (array) - MPS indices \(i\) for which self.site(i) is self. unit_cell[u].
- lat_idx (2D array) - The row \(j\) contains the lattice index (without \(u\) ) corresponding to mps_idx[j].
mps_sites()
Return a list of sites for all MPS indices.
Equivalent to [self.site(i) for i in range(self.N_sites)].
This should be used for sites of 1D tensor networks (MPS, MPO,...).
multi_coupling_shape ( \(d x\) )
Calculate correct shape of the strengths for a multi_coupling.
Parameters dx (2D array, shape (N_ops, dim)) - dx [i, :] is the translation vector in the lattice for the \(i\)-th operator. Corresponds to the \(d x\) of each operator given in the argument ops of tenpy.models.model. CouplingModel.add_multi_coupling().

\section*{Returns}
- coupling_shape (tuple of int) - Len dim. The correct shape for an array specifying the coupling strength. lat_indices has only rows within this shape.
- shift_lat_indices (array) - Translation vector from origin to the lower left corner of box spanned by \(d x\). (Unlike for coupling_shape () it can also contain entries \(>0\) )
number_nearest_neighbors ( \(u=0\) )
Deprecated.
Deprecated since version 0.5.0: Use count_neighbors () instead.
number_next_nearest_neighbors ( \(u=0\) )
Deprecated.
Deprecated since version 0.5.0: Use count_neighbors () instead.
plot_basis (ax, origin=(0.0, 0.0), shade=None, **kwargs)
Plot arrows indicating the basis vectors of the lattice.

\section*{Parameters}
- ax (matplotlib.axes.Axes) - The axes on which we should plot.
- **kwargs - Keyword arguments for ax. arrow.
plot_bc_identified (ax, direction=- 1, origin=None, cylinder_axis=False, **kwargs)
Mark two sites indified by periodic boundary conditions.
Works only for lattice with a 2 -dimensional basis.

\section*{Parameters}
- ax (matplotlib.axes.Axes) - The axes on which we should plot.
- direction (int) - The direction of the lattice along which we should mark the idenitified sites. If None, mark it along all directions with periodic boundary conditions.
- cylinder_axis (bool) - Whether to plot the cylinder axis as well.
- origin (None / np.ndarray) - The origin starting from where we mark the identified sites. Defaults to the first entry of unit_cell_positions.
- **kwargs - Keyword arguments for the used ax.plot.
plot_coupling (ax, coupling=None, wrap=False, **kwargs)
Plot lines connecting nearest neighbors of the lattice.

\section*{Parameters}
- ax (matplotlib.axes.Axes) - The axes on which we should plot.
- coupling (list of (u1, u2, dx)) - By default (None), use self. pairs['nearest_neighbors']. Specifies the connections to be plotted; iteating over lattice indices \((i 0, i l, \ldots\) ), we plot a connection from the site (i0, i1, ..., u1) to the site ( \(\mathrm{i} 0+\mathrm{dx}[0]\), \(i 1+\mathrm{dx}[1], \ldots, \mathrm{u}\) ), taking into account the boundary conditions.
- wrap (bool) - If True, plot couplings going around the boundary by directly connecting the sites it connects. This might be hard to see, as this puts lines from one end of the lattice to the other. If False, plot the couplings as dangling lines.
- **kwargs - Further keyword arguments given to ax.plot ().
plot_order (ax, order=None, textkwargs=\{'color': 'r'\}, **kwargs)
Plot a line connecting sites in the specified "order" and text labels enumerating them.

\section*{Parameters}
- ax (matplotlib. axes.Axes) - The axes on which we should plot.
- order (None / 2D array (self.N_sites, self.dim+1))- The order as returned by ordering(); by default (None) use order.
- textkwargs (None I dict) - If not None, we add text labels enumerating the sites in the plot. The dictionary can contain keyword arguments for ax.text ().
- **kwargs - Further keyword arguments given to ax. plot ().
plot_sites (ax, markers=['o', ' \({ }^{\prime}\) ', 's', ' \(p^{\prime}\) ' ' \(h^{\prime}\) ', D'], **kwargs)
Plot the sites of the lattice with markers.

\section*{Parameters}
- ax (matplotlib. axes.Axes) - The axes on which we should plot.
- markers (list) - List of values for the keywork marker of ax.plot () to distinguish the different sites in the unit cell, a site \(u\) in the unit cell is plotted with a marker markers[u \% len(markers)].
- **kwargs - Further keyword arguments given to ax.plot ().

\section*{position (lat_idx)}
return 'space' position of one or multiple sites.
Parameters lat_idx (ndarray, (. . , dim+1))-Lattice indices.
Returns pos - The position of the lattice sites specified by lat_idx in real-space.
Return type ndarray, (..., dim)
possible_couplings ( \(u 1, u 2, d x\), strength=None)
Find possible MPS indices for two-site couplings.
For periodic boundary conditions ( \(\mathrm{bc}[\mathrm{a}]==\mathrm{False}\) ) the index x _a is taken modulo Ls [a] and runs through range (Ls [a]). For open boundary conditions, x_a is limited to \(0<=\) x_a \(<\) Ls [a] and \(0<=x \_a+d x[a]<\) lat.Ls [a].

\section*{Parameters}
- u1 (int) - Indices within the unit cell; the \(u 1\) and \(u 2\) of add_coupling()
- u2 (int) - Indices within the unit cell; the \(u 1\) and \(u 2\) of add_coupling()
- \(\mathbf{d x}(\operatorname{array})\) - Length dim. The translation in terms of basis vectors for the coupling.
- strength (array_like / None) - If given, instead of returning lat_indices and coupling_shape directly return the correct strength_12.

\section*{Returns}
- mps1, mps2 ( \(1 D\) array) - For each possible two-site coupling the MPS indices for the \(u 1\) and \(u 2\).
- strength_vals ( \(1 D\) array) - (Only returend if strength is not None.) Such that for (i, \(j, ~ s) ~ i n ~ z i p\left(m p s 1, ~ m p s 2, ~ s t r e n g t h \_v a l s\right): ~ i t e r a t e s ~ o v e r ~ a l l ~ p o s s i b l e ~ c o u-~\) plings with \(s\) being the strength of that coupling. Couplings where strength_vals \(=0\). are filtered out.
- lat_indices ( \(2 D\) int array) - (Only returend if strength is None.) Rows of lat_indices correspond to entries of mpsl and mps 2 and contain the lattice indices of the "lower left corner" of the box containing the coupling.
- coupling_shape (tuple of int) - (Only returend if strength is None.) Len dim. The correct shape for an array specifying the coupling strength. lat_indices has only rows within this shape.
possible_multi_couplings (ops, strength=None)
Generalization of possible_couplings() to couplings with more than 2 sites.
Parameters ops (list of (opname, \(\mathrm{dx}, \mathrm{u})\) ) - Same as the argument ops of add_multi_coupling().

\section*{Returns}
- mps_ijkl ( \(2 D\) int array) - Each row contains MPS indices \(i, j, k, l, \ldots\) ` for each of the operators positions. The positions are defined by \(d x\) ( \(\mathrm{j}, \mathrm{k}, 1, \ldots\) relative to \(i\) ) and boundary coundary conditions of self (how much the box for given \(d x\) can be shifted around without hitting a boundary - these are the different rows).
- strength_vals ( \(1 D\) array) - (Only returend if strength is not None.) Such that for (ijkl, s) in zip(mps_ijkl, strength_vals): iterates over all possible couplings with \(s\) being the strength of that coupling. Couplings where strength_vals \(==0\). are filtered out.
- lat_indices ( \(2 D\) int array) - (Only returend if strength is None.) Rows of lat_indices correspond to rows of mps_ijkl and contain the lattice indices of the "lower left corner" of the box containing the coupling.
- coupling_shape (tuple of int) - (Only returend if strength is None.) Len dim. The correct shape for an array specifying the coupling strength. lat_indices has only rows within this shape.
```

site(i)

```
return Site instance corresponding to an MPS index \(i\)
```

test_sanity()

```

Sanity check.
Raises ValueErrors, if something is wrong.

\subsection*{19.1.5 Kagome}
- full name: tenpy.models.lattice.Kagome
- parent module: tenpy.models.lattice
- type: class

\section*{Inheritance Diagram}


\section*{Methods}
\begin{tabular}{|c|c|}
\hline Kagome.__init__(Lx, Ly, sites, **kwargs) & Initialize self. \\
\hline Kagome.count_neighbors([u, key]) & Count e.g. \\
\hline Kagome.coupling_shape(dx) & Calculate correct shape of the strengths for a coupling. \\
\hline Kagome.distance(u1, u2, dx) & Get the distance for a given coupling between two sites in the lattice. \\
\hline Kagome.enlarge_mps_unit_cell([factor]) & Repeat the unit cell for infinite MPS boundary conditions; in place. \\
\hline Kagome.find_coupling_pairs([max_dx, cutoff, eps]) & Automatically find coupling pairs grouped by distances. \\
\hline Kagome.from_hdf5(hdf5_loader, h5gr, subpath) & Load instance from a HDF5 file. \\
\hline Kagome.lat2mps_idx(lat_idx) & Translate lattice indices \(\left(x \_0, \ldots, x_{-}\{D-1\}\right.\), u) to MPS index \(i\). \\
\hline Kagome.mps2lat_idx(i) & Translate MPS index \(i\) to lattice indices ( \(\mathrm{x} \_0\), \(\ldots\), x_\{dim-1\}, u). \\
\hline Kagome.mps2lat_values(A[, axes, u]) & Reshape/reorder \(A\) to replace an MPS index by lattice indices. \\
\hline Kagome.mps2lat_values_masked(A[, axes, ...]) & Reshape/reorder an array \(A\) to replace an MPS index by lattice indices. \\
\hline Kagome.mps_idx_fix_u([u]) & return an index array of MPS indices for which the site within the unit cell is \(u\). \\
\hline Kagome.mps_lat_idx_fix_u([u]) & Similar as mps_idx_fix_u(), but return also the corresponding lattice indices. \\
\hline Kagome.mps_sites() & Return a list of sites for all MPS indices. \\
\hline Kagome.multi_coupling_shape(dx) & Calculate correct shape of the strengths for a multi_coupling. \\
\hline Kagome.number_nearest_neighbors([u]) & Deprecated. \\
\hline Kagome.number_next_nearest_neighbors([u]) & Deprecated. \\
\hline Kagome.ordering(order) & Provide possible orderings of the \(N\) lattice sites. \\
\hline Kagome.plot_basis(ax[, origin, shade]) & Plot arrows indicating the basis vectors of the lattice. \\
\hline ```
Kagome.plot_bc_identified(ax[, direction,
...])
``` & Mark two sites indified by periodic boundary conditions. \\
\hline Kagome.plot_coupling(ax[, coupling, wrap]) & Plot lines connecting nearest neighbors of the lattice. \\
\hline
\end{tabular}

Table 11 - continued from previous page
\begin{tabular}{|c|c|}
\hline Kagome.plot_order(ax[, order, textkwargs]) & Plot a line connecting sites in the specified "order" and text labels enumerating them. \\
\hline Kagome.plot_sites(ax[, markers]) & Plot the sites of the lattice with markers. \\
\hline Kagome.position(lat_idx) & return 'space' position of one or multiple sites. \\
\hline Kagome.possible_couplings(u1, u2, dx[, strength]) & Find possible MPS indices for two-site couplings. \\
\hline Kagome.possible_multi_couplings(ops[, strength]) & Generalization of possible_couplings() to couplings with more than 2 sites. \\
\hline Kagome.save_hdf5(hdf5_saver, h5gr, subpath) & Export self into a HDF5 file. \\
\hline Kagome.site(i) & return Site instance corresponding to an MPS index \(i\) \\
\hline Kagome.test_sanity() & Sanity check. \\
\hline
\end{tabular}

\section*{Class Attributes and Properties}
\begin{tabular}{ll}
\hline Kagome. Lu & \begin{tabular}{l} 
the (expected) number of sites in the unit cell, \\
len (unit_cell).
\end{tabular} \\
\hline Kagome.boundary_conditions & \begin{tabular}{l} 
Human-readable list of boundary conditions from bc \\
and bc_shift.
\end{tabular} \\
\hline Kagome.dim & the dimension of the lattice \\
\hline Kagome. nearest_neighbors & \\
\hline Kagome.next_nearest_neighbors & \begin{tabular}{l} 
Defines an ordering of the lattice sites, thus mapping the \\
lattice to a 1D chain.
\end{tabular} \\
\hline Kagome.next_next_nearest_neighbors &
\end{tabular}
class tenpy.models.lattice. Kagome (Lx,Ly, sites, **kwargs)
Bases: tenpy.models.lattice.Lattice
A Kagome lattice.

\section*{Parameters}
- \(\mathbf{L x}(i n t)\) - The length in each direction.
- Ly (int) - The length in each direction.
- sites ((list of) Site) - The two local lattice sites making the unit_cell of the Lattice. If only a single Site is given, it is used for both sites.
- **kwargs - Additional keyword arguments given to the Lattice. basis, pos and pairs are set accordingly.
property boundary_conditions
Human-readable list of boundary conditions from bc and bc_shift.
Returns boundary_conditions - List of "open" or "periodic", one entry for each direction of the lattice.

Return type list of str
count_neighbors ( \(u=0\), key='nearest_neighbors')
Count e.g. the number of nearest neighbors for a site in the bulk.


\section*{Parameters}
- u (int) - Specifies the site in the unit cell, for which we should count the number of neighbors (or whatever key specifies).
- key (str) - Key of pairs to select what to count.

Returns number - Number of nearest neighbors (or whatever key specified) for the \(u\)-th site in the unit cell, somewhere in the bulk of the lattice. Note that it might not be the correct value at the edges of a lattice with open boundary conditions.

\section*{Return type int}
coupling_shape ( \(d x\) )
Calculate correct shape of the strengths for a coupling.
Parameters \(\mathbf{d x}\) (tuple of int) - Translation vector in the lattice for a coupling of two operators. Corresponds to \(d x\) argument of tenpy.models.model. CouplingModel. add_multi_coupling().

\section*{Returns}
- coupling_shape (tuple of int) - Len dim. The correct shape for an array specifying the coupling strength. lat_indices has only rows within this shape.
- shift_lat_indices (array) - Translation vector from origin to the lower left corner of box spanned by \(d x\).
distance ( \(u 1, u 2, d x\) )
Get the distance for a given coupling between two sites in the lattice.
The \(u 1, u 2, d x\) parameters are defined in analogy with add_coupling(), i.e., this function calculates the distance between a pair of operators added with add_coupling (using the basis and unit_cell_positions of the lattice).

Warning: This function ignores "wrapping" arround the cylinder in the case of periodic boundary conditions.

\section*{Parameters}
- u1 (int) - Indices within the unit cell; the \(u 1\) and \(u 2\) of add_coupling()
- u2 (int) - Indices within the unit cell; the \(u 1\) and \(u 2\) of add_coupling()
- \(\mathbf{d x}\) (array) - Length dim. The translation in terms of basis vectors for the coupling.

Returns distance - The distance between site at lattice indices [ \(\mathrm{x}, \mathrm{y}, \mathrm{u} 1]\) and \([\mathrm{x}+\) \(d x[0], y+d x[1], u 2]\), ignoring any boundary effects.

Return type float

\section*{enlarge_mps_unit_cell (factor=2)}

Repeat the unit cell for infinite MPS boundary conditions; in place.
Parameters factor (int) - The new number of sites in the MPS unit cell will be increased from \(N\) _sites to factor*N_sites_per_ring. Since MPS unit cells are repeated in the \(x\)-direction in our convetion, the lattice shape goes from ( \(L x, L y, \ldots, L u\) ) to (Lx*factor, Ly, ..., Lu).
find_coupling_pairs (max_dx=3, cutoff=None, eps=1e-10)
Automatically find coupling pairs grouped by distances.
Given the unit_cell_positions and basis, the coupling pairs of nearest_neighbors, next_nearest_neighbors etc at a given distance are basically fixed (although not uniquely, since we take out half of them to avoid double-counting couplings in both directions A_i B_j + B_i A_i). This function iterates through all possible couplings up to a given cutoff distance and then determines the possible pairs at fixed distances (up to round-off errors).

\section*{Parameters}
- max_dx (int) - Maximal index for each index of \(d x\) to iterate over. You need large enough values to include every possible coupling up to the desired distance, but choosing it too large might make this function run for a long time.
- cutoff (float) - Maximal distance (in the units in which basis and unit_cell_positions is given).
- eps (float) - Tolerance up to which to distances are considered the same.

Returns coupling_pairs - Keys are distances of nearest-neighbors, next-nearest-neighbors etc.
Values are [(u1, u2, dx), ...] as in pairs.
Return type dict
classmethod from_hdf5 (hdf5_loader, h5gr, subpath)
Load instance from a HDF5 file.
This method reconstructs a class instance from the data saved with save_hdf5 ().

\section*{Parameters}
- hdf5_loader (Hdf5Loader) - Instance of the loading engine.
- h5gr (Group) - HDF5 group which is represent the object to be constructed.
- subpath (str) - The name of \(h 5 g r\) with a '/' in the end.

Returns obj - Newly generated class instance containing the required data.

Return type cls
lat2mps_idx (lat_idx)
Translate lattice indices ( \(\mathrm{x} \_0, \ldots, \mathrm{x}_{-}\{\mathrm{D}-1\}, \mathrm{u}\) ) to MPS index \(i\).
Parameters lat_idx (array_like [.., dim+1]) - The last dimension corresponds to lattice indices \(\left(x_{-} 0, \ldots, x_{-}\{D-1\}, u\right)\). All lattice indices should be positive and smaller than the corresponding entry in self.shape. Exception: for "infinite" \(b c_{-} M P S\), an \(x \_0\) outside indicates shifts accross the boundary.

Returns \(\mathbf{i}\) - MPS index/indices corresponding to lat_idx. Has the same shape as lat_idx without the last dimension.

Return type array_like
mps2lat_idx (i)
Translate MPS index \(i\) to lattice indices ( \(\left.\mathrm{x} \_0, \ldots, \mathrm{x}_{-}\{\operatorname{dim}-1\}, u\right)\).
Parameters i (int / array_like of int)-MPS index/indices.
Returns lat_idx - First dimensions like \(i\), last dimension has len dim \({ }^{`}+1\) and contains the lattice indices " \(\left(x \_0, \ldots, x_{\_}\{d i m-1\}, u\right)\) ` corresponding to \(i\). For \(i\) accross the MPS unit cell and "infinite" \(b c_{-} M P S\), we shift \(x_{-} 0\) accordingly.

\section*{Return type array}
mps2lat_values \((A\), axes \(=0, u=\) None)
Reshape/reorder \(A\) to replace an MPS index by lattice indices.

\section*{Parameters}
- A (ndarray) - Some values. Must have A.shape[axes] = self.N_sites if \(u\) is None, or A. shape [axes] = self.N_cells if \(u\) is an int.
- axes ( (iterable of) int) - chooses the axis which should be replaced.
- u (None I int) - Optionally choose a subset of MPS indices present in the axes of \(A\), namely the indices corresponding to self.unit_cell[u], as returned by mps_idx_fix_u(). The resulting array will not have the additional dimension(s) of \(u\).

Returns res_A - Reshaped and reordered verions of A. Such that MPS indices along the specified axes are replaced by lattice indices, i.e., if MPS index \(j\) maps to lattice site ( \(x 0, x 1, x 2\) ), then res_A[..., \(x 0, x 1, x 2, \ldots]=A[\ldots, j, \ldots]\).
Return type ndarray

\section*{Examples}

Say you measure expection values of an onsite term for an MPS, which gives you an 1D array \(A\), where \(A[i]\) is the expectation value of the site given by self.mps2lat_idx(i). Then this function gives you the expectation values ordered by the lattice:
```

>>> print(lat.shape, A.shape)
(10, 3, 2) (60,)
>>> A_res = lat.mps2lat_values(A)
>>> A_res.shape
(10, 3, 2)
>>> A_res[tuple(lat.mps2lat_idx(5))] == A[5]
True

```

If you have a correlation function C [i, j], it gets just slightly more complicated:
```

>>> print(lat.shape, C.shape)
(10, 3, 2) (60, 60)
>>> lat.mps2lat_values(C, axes=[0, 1]).shape
(10, 3, 2, 10, 3, 2)

```

If the unit cell consists of different physical sites, an onsite operator might be defined only on one of the sites in the unit cell. Then you can use mps_idx_fix_u() to get the indices of sites it is defined on, measure the operator on these sites, and use the argument \(u\) of this function.
```

>>> u = 0
>>> idx_subset = lat.mps_idx_fix_u(u)
>>> A_u = A[idx_subset]
>>> A_u__res = lat.mps2lat_values(A_u, u=u)
>>> A_u_res.shape
(10, 3)
>>> np.all(A_res[:, :, u] == A_u_res[:, :])
True

```
mps2lat_values_masked ( \(A\), axes=- 1, mps_inds=None, include_u=None)
Reshape/reorder an array \(A\) to replace an MPS index by lattice indices.
This is a generalization of mps2lat_values () allowing for the case of an arbitrary set of MPS indices present in each axis of \(A\).

\section*{Parameters}
- A (ndarray) - Some values.
- axes ( (iterable of) int) - Chooses the axis of \(A\) which should be replaced. If multiple axes are given, you also need to give multiple index arrays as mps_inds.
- mps_inds ((list of) 1D ndarray) - Specifies for each axis in axes, for which MPS indices we have values in the corresponding axis of A. Defaults to [np. arange (A. shape[ax]) for ax in axes]. For indices accross the MPS unit cell and "infinite" \(b c_{-} M P S\), we shift \(x \_0\) accordingly.
- include_u ((list of) bool) - Specifies for each axis in axes, whether the \(u\) index of the lattice should be included into the output array res_A. Defaults to len (self. unit_cell) > 1 .

Returns res_A - Reshaped and reordered copy of A. Such that MPS indices along the specified axes are replaced by lattice indices, i.e., if MPS index \(j\) maps to lattice site ( \(x 0, x 1, x 2\) ), then res_A[..., x0, x1, x2, ...] = A[..., mps_inds[j], ...].

\section*{Return type np.ma.MaskedArray}
mps_idx_fix_u(u=None)
return an index array of MPS indices for which the site within the unit cell is \(u\).
If you have multiple sites in your unit-cell, an onsite operator is in general not defined for all sites. This functions returns an index array of the mps indices which belong to sites given by self. unit_cell[u].

Parameters \(\mathbf{u}\) (None / int) - Selects a site of the unit cell. None (default) means all sites.
Returns mps_idx - MPS indices for which self.site(i) is self.unit_cell[u]. Ordered ascending.

\section*{Return type array}
mps_lat_idx_fix_u (u=None)
Similar as mps_idx_fix_u(), but return also the corresponding lattice indices.
Parameters \(\mathbf{u}\) (None / int) - Selects a site of the unit cell. None (default) means all sites.

\section*{Returns}
- mps_idx (array) - MPS indices \(i\) for which self.site(i) is self. unit_cell[u].
- lat_idx ( \(2 D\) array) - The row \(j\) contains the lattice index (without \(u\) ) corresponding to mps_idx[j].
mps_sites()
Return a list of sites for all MPS indices.
Equivalent to [self.site(i) for i in range(self.N_sites)].
This should be used for sites of 1D tensor networks (MPS, MPO,...).
multi_coupling_shape ( \(d x\) )
Calculate correct shape of the strengths for a multi_coupling.
Parameters \(d \mathbf{x}(2 \mathrm{D}\) array, shape (N_ops, dim)) - dx[i, :] is the translation vector in the lattice for the \(i\)-th operator. Corresponds to the \(d x\) of each operator given in the argument ops of tenpy.models.model.CouplingModel.add_multi_coupling().

\section*{Returns}
- coupling_shape (tuple of int) - Len dim. The correct shape for an array specifying the coupling strength. lat_indices has only rows within this shape.
- shift_lat_indices (array) - Translation vector from origin to the lower left corner of box spanned by \(d x\). (Unlike for coupling_shape () it can also contain entries \(>0\) )
number_nearest_neighbors ( \(u=0\) )
Deprecated.
Deprecated since version 0.5.0: Use count_neighbors () instead.
number_next_nearest_neighbors ( \(u=0\) )
Deprecated.
Deprecated since version 0.5.0: Use count_neighbors () instead.

\section*{property order}

Defines an ordering of the lattice sites, thus mapping the lattice to a 1 D chain.
Each row of the array contains the lattice indices for one site, the order of the rows thus specifies a path through the lattice, along which an MPS will wind through through the lattice.
You can visualize the order with plot_order ().
ordering (order)
Provide possible orderings of the \(N\) lattice sites.
This function can be overwritten by derived lattices to define additional orderings. The following orders are defined in this method:
\begin{tabular}{|l|l|l|}
\hline order & equivalent priority & equivalent snake_winding \\
\hline 'Cstyle' & \((0,1, \ldots, \operatorname{dim}-1, \operatorname{dim})\) & (False, \(\ldots\), False, False) \\
\hline 'default' & & \\
\hline 'snake' & \((0,1, \ldots, \operatorname{dim}-1, \operatorname{dim})\) & (True, \(\ldots\), True, True) \\
\hline 'snakeCstyle' & \\
\hline 'Fstyle' & (dim-1, \(\ldots, 1,0, \operatorname{dim})\) & (False, \(\ldots\), False, False) \\
\hline 'snakeFstyle' & (dim-1, \(, \ldots, 1,0, \operatorname{dim})\) & (False,.., False, False) \\
\hline
\end{tabular}


Note: For lattices with a non-trivial unit cell (e.g. Honeycomb, Kagome), the grouped order might be more appropriate, see get_order_grouped()

Parameters order (str | ('standard', snake_winding, priority) | ('grouped', groups, ...)) - Specifies the desired ordering using one of the strings of the above tables. Alternatively, an ordering is specified by a tuple with first entry specifying a function, 'standard' for get_order() and 'grouped' for get_order_grouped(), and other arguments in the tuple as specified in the documentation of these functions.

Returns order - the order to be used for order.
Return type array, shape (N, D+1), dtype np.intp

\section*{See also:}
get_order generates the order from equivalent priority and snake_winding.
get_order_grouped variant of get_order.
plot_order visualizes the resulting order.
plot_basis (ax, origin=(0.0, 0.0), shade \(=\) None, \({ }^{* *}\) kwargs )
Plot arrows indicating the basis vectors of the lattice.

\section*{Parameters}
- ax (matplotlib. axes.Axes) - The axes on which we should plot.
- **kwargs - Keyword arguments for ax. arrow.
plot_bc_identified (ax, direction=- 1, origin=None, cylinder_axis=False, **kwargs)
Mark two sites indified by periodic boundary conditions.
Works only for lattice with a 2-dimensional basis.

\section*{Parameters}
- ax (matplotlib. axes.Axes) - The axes on which we should plot.
- direction (int) - The direction of the lattice along which we should mark the idenitified sites. If None, mark it along all directions with periodic boundary conditions.
- cylinder_axis (b○○l) - Whether to plot the cylinder axis as well.
- origin (None / np.ndarray) - The origin starting from where we mark the identified sites. Defaults to the first entry of unit_cell_positions.
- **kwargs - Keyword arguments for the used ax. plot.
plot_coupling (ax, coupling=None, wrap=False, **kwargs)
Plot lines connecting nearest neighbors of the lattice.

\section*{Parameters}
- ax (matplotlib. axes.Axes) - The axes on which we should plot.
- coupling (list of (u1, \(u 2, d x)\) ) - By default (None), use self. pairs['nearest_neighbors']. Specifies the connections to be plotted; iteating over lattice indices ( \(i 0, i 1, \ldots\) ), we plot a connection from the site (i0, i1, ..., u1) to the site (i0 \(+d x[0]\), \(i 1+d x[1], \ldots, u 2\) ), taking into account the boundary conditions.
- wrap (b○○l) - If True, plot couplings going around the boundary by directly connecting the sites it connects. This might be hard to see, as this puts lines from one end of the lattice to the other. If Fal se, plot the couplings as dangling lines.
- **kwargs - Further keyword arguments given to ax.plot().
plot_order (ax, order=None, textkwargs=\{'color': 'r'\}, **kwargs)
Plot a line connecting sites in the specified "order" and text labels enumerating them.

\section*{Parameters}
- ax (matplotlib. axes.Axes) - The axes on which we should plot.
- order (None / 2D array (self.N_sites, self.dim+1))- The order as returned by ordering(); by default (None) use order.
- textkwargs (None I dict) - If not None, we add text labels enumerating the sites in the plot. The dictionary can contain keyword arguments for ax.text ().
- **kwargs - Further keyword arguments given to ax.plot ().
plot_sites (ax, markers=['o', ' \({ }^{\prime}\) ', 's', ' \({ }^{\prime}\) ', 'h', 'D'], **kwargs)
Plot the sites of the lattice with markers.

\section*{Parameters}
- ax (matplotlib. axes.Axes) - The axes on which we should plot.
- markers (list) - List of values for the keywork marker of ax. plot () to distinguish the different sites in the unit cell, a site \(u\) in the unit cell is plotted with a marker markers[u \% len(markers)].
- **kwargs - Further keyword arguments given to ax. plot ().
position (lat_idx)
return 'space' position of one or multiple sites.
Parameters lat_idx (ndarray, (. . . , dim+1)) - Lattice indices.
Returns pos - The position of the lattice sites specified by lat_idx in real-space.
Return type ndarray, (..., dim)
possible_couplings ( \(u 1, u 2, d x\), strength=None)
Find possible MPS indices for two-site couplings.
For periodic boundary conditions ( \(\mathrm{bc}[\mathrm{a}]==\mathrm{False}\) ) the index x _a is taken modulo Ls [a] and runs through range (Ls [a]). For open boundary conditions, x_a is limited to \(0<=x \_a<\) Ls [a] and \(0<=x \_a+d x[a]<\) lat.Ls[a].

\section*{Parameters}
- u1 (int) - Indices within the unit cell; the \(u 1\) and \(u 2\) of add_coupling ()
- u2 (int) - Indices within the unit cell; the \(u 1\) and \(u 2\) of add_coupling()
- \(\mathbf{d x}(\operatorname{array})\) - Length dim. The translation in terms of basis vectors for the coupling.
- strength (array_like / None) - If given, instead of returning lat_indices and coupling_shape directly return the correct strength_12.

\section*{Returns}
- mps1, mps2 (lD array) - For each possible two-site coupling the MPS indices for the \(u 1\) and \(u 2\).
- strength_vals (1D array) - (Only returend if strength is not None.) Such that for (i, \(j, ~ s) ~ i n ~ z i p\left(m p s 1, ~ m p s 2, ~ s t r e n g t h \_v a l s\right): ~ i t e r a t e s ~ o v e r ~ a l l ~ p o s s i b l e ~ c o u-~\) plings with \(s\) being the strength of that coupling. Couplings where strength_vals \(==0\). are filtered out.
- lat_indices ( \(2 D\) int array) - (Only returend if strength is None.) Rows of lat_indices correspond to entries of \(m p s 1\) and \(m p s 2\) and contain the lattice indices of the "lower left corner" of the box containing the coupling.
- coupling_shape (tuple of int) - (Only returend if strength is None.) Len dim. The correct shape for an array specifying the coupling strength. lat_indices has only rows within this shape.
possible_multi_couplings (ops, strength=None)
Generalization of possible_couplings () to couplings with more than 2 sites.

Parameters ops (list of (opname, \(\mathrm{dx}, \mathrm{u})\) ) - Same as the argument ops of add_multi_coupling().

\section*{Returns}
- mps_ijkl (2D int array) - Each row contains MPS indices \(i, j, k, l, \ldots\). for each of the operators positions. The positions are defined by \(d x(\mathrm{j}, \mathrm{k}, 1, \ldots\) relative to \(i\) ) and boundary coundary conditions of self (how much the box for given \(d x\) can be shifted around without hitting a boundary - these are the different rows).
- strength_vals ( \(1 D\) array) - (Only returend if strength is not None.) Such that for (ijkl, s) in zip(mps_ijkl, strength_vals): iterates over all possible couplings with \(s\) being the strength of that coupling. Couplings where strength_vals \(==0\). are filtered out.
- lat_indices ( \(2 D\) int array) - (Only returend if strength is None.) Rows of lat_indices correspond to rows of \(m p s \_i j k l\) and contain the lattice indices of the "lower left corner" of the box containing the coupling.
- coupling_shape (tuple of int) - (Only returend if strength is None.) Len dim. The correct shape for an array specifying the coupling strength. lat_indices has only rows within this shape.
```

save_hdf5 (hdf5_saver, h5gr, subpath)

```

Export self into a HDF5 file.
This method saves all the data it needs to reconstruct self with from_hdf5 ().
Specifically, it saves unit_cell, Ls, unit_cell_positions, basis, boundary_conditions, pairs under their name, bc_MPS as "boundary_conditions_MPS", and order as "order_for_MPS". Moreover, it saves dim and N_sites as HDF5 attributes.

\section*{Parameters}
- hdf5_saver (Hdf5Saver) - Instance of the saving engine.
- h5gr (: class`Group ) - HDF5 group which is supposed to represent self.
- subpath (str) - The name of \(h 5 g r\) with a '/' in the end.
```

site(i)

```
return Site instance corresponding to an MPS index \(i\)
```

test_sanity()

```

Sanity check.
Raises ValueErrors, if something is wrong.

\section*{\(\operatorname{dim}=2\)}
the dimension of the lattice
\(\mathrm{Lu}=3\)
the (expected) number of sites in the unit cell, len (unit_cell).

\subsection*{19.1.6 Ladder}
- full name: tenpy.models.lattice.Ladder
- parent module: tenpy.models.lattice
- type: class

\section*{Inheritance Diagram}


\section*{Methods}
\begin{tabular}{|c|c|}
\hline Ladder.__init__(L, sites, **kwargs) & Initialize self. \\
\hline Ladder.count_neighbors([u, key]) & Count e.g. \\
\hline Ladder.coupling_shape(dx) & Calculate correct shape of the strengths for a coupling. \\
\hline Ladder.distance(u1, u2, dx) & Get the distance for a given coupling between two sites in the lattice. \\
\hline Ladder.enlarge_mps_unit_cell([factor]) & Repeat the unit cell for infinite MPS boundary conditions; in place. \\
\hline ```
Ladder.find_coupling_pairs([max_dx, cut-
off, eps])
``` & Automatically find coupling pairs grouped by distances. \\
\hline Ladder.from_hdf5(hdf5_loader, h5gr, subpath) & Load instance from a HDF5 file. \\
\hline Ladder.lat2mps_idx(lat_idx) & Translate lattice indices ( \(x\) _ \(0, \ldots, x_{-}\{D-1\}\), u) to MPS index \(i\). \\
\hline Ladder.mps2lat_idx(i) & Translate MPS index \(i\) to lattice indices ( \(\mathrm{x} \_0\), ...., x_\{dim-1\}, u). \\
\hline Ladder.mps2lat_values(A[, axes, u]) & Reshape/reorder \(A\) to replace an MPS index by lattice indices. \\
\hline Ladder.mps2lat_values_masked(A[, axes, ...]) & Reshape/reorder an array \(A\) to replace an MPS index by lattice indices. \\
\hline Ladder.mps_idx_fix_u([u]) & return an index array of MPS indices for which the site within the unit cell is \(u\). \\
\hline Ladder.mps_lat_idx_fix_u([u]) & Similar as mps_idx_fix_u(), but return also the corresponding lattice indices. \\
\hline Ladder.mps_sites() & Return a list of sites for all MPS indices. \\
\hline Ladder.multi_coupling_shape(dx) & Calculate correct shape of the strengths for a multi_coupling. \\
\hline
\end{tabular}

Table 13 - continued from previous page
\begin{tabular}{|c|c|}
\hline Ladder.number_nearest_neighbors([u]) & Deprecated. \\
\hline Ladder.number_next_nearest_neighbors([u]) & )Deprecated. \\
\hline Ladder.ordering(order) & Provide possible orderings of the \(N\) lattice sites. \\
\hline Ladder.plot_basis(ax[, origin, shade]) & Plot arrows indicating the basis vectors of the lattice. \\
\hline Ladder.plot_bc_identified(ax[, direction, ...]) & Mark two sites indified by periodic boundary conditions. \\
\hline Ladder.plot_coupling(ax[, coupling, wrap]) & Plot lines connecting nearest neighbors of the lattice. \\
\hline Ladder.plot_order(ax[, order, textkwargs]) & Plot a line connecting sites in the specified "order" and text labels enumerating them. \\
\hline Ladder.plot_sites(ax[, markers]) & Plot the sites of the lattice with markers. \\
\hline Ladder.position(lat_idx) & return 'space' position of one or multiple sites. \\
\hline ```
Ladder.possible_couplings(u1, u2, dx[,
strength])
``` & Find possible MPS indices for two-site couplings. \\
\hline Ladder.possible_multi_couplings(ops[, strength]) & Generalization of possible_couplings() to couplings with more than 2 sites. \\
\hline Ladder.save_hdf5(hdf5_saver, h5gr, subpath) & Export self into a HDF5 file. \\
\hline Ladder.site(i) & return Site instance corresponding to an MPS index \(i\) \\
\hline Ladder.test_sanity() & Sanity check. \\
\hline
\end{tabular}

\section*{Class Attributes and Properties}
\begin{tabular}{ll}
\hline Ladder.Lu & \begin{tabular}{l} 
the (expected) number of sites in the unit cell, \\
len_(unit_cell).
\end{tabular} \\
\hline Ladder.boundary_conditions & \begin{tabular}{l} 
Human-readable list of boundary conditions from bc \\
and bc_shift.
\end{tabular} \\
\hline Ladder.dim & the dimension of the lattice \\
\hline Ladder.nearest_neighbors & \begin{tabular}{l} 
Defines an ordering of the lattice sites, thus mapping the \\
lattice to a 1D chain.
\end{tabular} \\
\hline Ladder.next_nearest_neighbors & \\
\hline Ladder.next_next_nearest_neighbors & \\
class tenpy.models.lattice. Ladder \((L\), sites, **kwargs) \\
Bases: tenpy.models.lattice.Lattice
\end{tabular}

A ladder coupling two chains.


\section*{Parameters}
- L (int) - The length of each chain, we have \(2 * \mathrm{~L}\) sites in total.
- sites ((list of) Site) - The two local lattice sites making the unit_cell of the Lattice. If only a single Site is given, it is used for both chains.
- **kwargs - Additional keyword arguments given to the Lattice. basis, pos and pairs are set accordingly.

\section*{\(\mathrm{Lu}=2\)}
the (expected) number of sites in the unit cell, len (unit_cell).
\(\operatorname{dim}=1\)
the dimension of the lattice
property boundary_conditions
Human-readable list of boundary conditions from bc and bc_shift.
Returns boundary_conditions - List of "open" or "periodic", one entry for each direction of the lattice.
Return type list of str
count_neighbors ( \(u=0\), key='nearest_neighbors')
Count e.g. the number of nearest neighbors for a site in the bulk.

\section*{Parameters}
- u (int) - Specifies the site in the unit cell, for which we should count the number of neighbors (or whatever key specifies).
- key (str) - Key of pairs to select what to count.

Returns number - Number of nearest neighbors (or whatever key specified) for the \(u\)-th site in the unit cell, somewhere in the bulk of the lattice. Note that it might not be the correct value at the edges of a lattice with open boundary conditions.

\section*{Return type int}

\section*{coupling_shape ( \(d x\) )}

Calculate correct shape of the strengths for a coupling.
Parameters \(\mathbf{d x}\) (tuple of int) - Translation vector in the lattice for a coupling of two operators. Corresponds to \(d x\) argument of tenpy.models.model. CouplingModel. add_multi_coupling().

\section*{Returns}
- coupling_shape (tuple of int) - Len dim. The correct shape for an array specifying the coupling strength. lat_indices has only rows within this shape.
- shift_lat_indices (array) - Translation vector from origin to the lower left corner of box spanned by \(d x\).
distance ( \(u 1, u 2, d x\) )
Get the distance for a given coupling between two sites in the lattice.
The \(u 1, u 2, d x\) parameters are defined in analogy with add_coupling(), i.e., this function calculates the distance between a pair of operators added with add_coupling (using the basis and unit_cell_positions of the lattice).

\footnotetext{
Warning: This function ignores "wrapping" arround the cylinder in the case of periodic boundary conditions.
}

\section*{Parameters}
- u1 (int) - Indices within the unit cell; the \(u 1\) and \(u 2\) of add_coupling ()
- u2 (int) - Indices within the unit cell; the \(u 1\) and \(u 2\) of add_coupling ()
- \(\mathbf{d x}\) (array) - Length dim. The translation in terms of basis vectors for the coupling.

Returns distance - The distance between site at lattice indices [x, y, u1] and [x + \(d x[0], y+d x[1], u 2]\), ignoring any boundary effects.

\section*{Return type float}
```

enlarge_mps_unit_cell (factor=2)

```

Repeat the unit cell for infinite MPS boundary conditions; in place.
Parameters factor (int) - The new number of sites in the MPS unit cell will be increased from \(N_{-}\)sites to factor*N_sites_per_ring. Since MPS unit cells are repeated in the \(x\)-direction in our convetion, the lattice shape goes from ( \(\mathrm{Lx}, \mathrm{Ly}, \ldots, \mathrm{Lu}\) ) to (Lx*factor, Ly, ..., Lu).
find_coupling_pairs ( max_dx=3, cutoff=None, eps=le-10)
Automatically find coupling pairs grouped by distances.
Given the unit_cell_positions and basis, the coupling pairs of nearest_neighbors, next_nearest_neighbors etc at a given distance are basically fixed (although not uniquely, since we take out half of them to avoid double-counting couplings in both directions A_i B_j + B_i A_i). This function iterates through all possible couplings up to a given cutoff distance and then determines the possible pairs at fixed distances (up to round-off errors).

\section*{Parameters}
- max_dx (int) - Maximal index for each index of \(d x\) to iterate over. You need large enough values to include every possible coupling up to the desired distance, but choosing it too large might make this function run for a long time.
- cutoff (float) - Maximal distance (in the units in which basis and unit_cell_positions is given).
- eps (float) - Tolerance up to which to distances are considered the same.

Returns coupling_pairs - Keys are distances of nearest-neighbors, next-nearest-neighbors etc.
Values are [ (u1, u2, dx), ...] as in pairs.
Return type dict
classmethod from_hdf5 (hdf5_loader, h5gr, subpath)
Load instance from a HDF5 file.
This method reconstructs a class instance from the data saved with save_hdf5().

\section*{Parameters}
- hdf5_loader (Hdf5Loader) - Instance of the loading engine.
- h5gr (Group) - HDF5 group which is represent the object to be constructed.
- subpath (str) - The name of \(h 5 g r\) with a ' / ' in the end.

Returns obj - Newly generated class instance containing the required data.
Return type cls
lat2mps_idx (lat_idx)
Translate lattice indices ( \(\left.x_{-} 0, \ldots, x_{-}\{D-1\}, u\right)\) to MPS index \(i\).

Parameters lat_idx (array_like [.., dim+1]) - The last dimension corresponds to lattice indices ( \(\left.x \_0, \ldots, x_{-}\{D-1\}, u\right)\). All lattice indices should be positive and smaller than the corresponding entry in self.shape. Exception: for "infinite" \(b c_{-} M P S\), an \(x \_0\) outside indicates shifts accross the boundary.
Returns i - MPS index/indices corresponding to lat_idx. Has the same shape as lat_idx without the last dimension.

Return type array_like

\section*{mps2lat_idx(i)}

Translate MPS index \(i\) to lattice indices ( \(\mathrm{x} \_0\), ..., \(\left.\mathrm{x} \_\{\operatorname{dim}-1\}, u\right)\).
Parameters i (int / array_like of int)-MPS index/indices.
Returns lat_idx - First dimensions like \(i\), last dimension has len dim \({ }^{`}+1\) and contains the lattice indices \({ }^{`}\left(x \_0, \ldots, x_{-}\{\operatorname{dim}-1\}, u\right)\) corresponding to \(i\). For \(i\) accross the MPS unit cell and "infinite" \(b c_{-} M P S\), we shift \(x \_0\) accordingly.

\section*{Return type array}
mps2lat_values ( \(A\), axes \(=0, u=\) None)
Reshape/reorder \(A\) to replace an MPS index by lattice indices.

\section*{Parameters}
- A (ndarray) - Some values. Must have A.shape[axes] = self.N_sites if \(u\) is None, or A. shape[axes] = self. N_cells if \(u\) is an int.
- axes ((iterable of) int) - chooses the axis which should be replaced.
- u (None I int) - Optionally choose a subset of MPS indices present in the axes of \(A\), namely the indices corresponding to self.unit_cell[u], as returned by mps_idx_fix_u(). The resulting array will not have the additional dimension(s) of \(u\).

Returns res_A - Reshaped and reordered verions of A. Such that MPS indices along the specified axes are replaced by lattice indices, i.e., if MPS index \(j\) maps to lattice site ( \(x 0, x 1, x 2\) ), then res_A[..., \(x 0, x 1, x 2, \ldots]=A[\ldots, j, \ldots]\).

\section*{Return type ndarray}

\section*{Examples}

Say you measure expection values of an onsite term for an MPS, which gives you an 1D array \(A\), where \(A[i]\) is the expectation value of the site given by self.mps2lat_idx(i). Then this function gives you the expectation values ordered by the lattice:
```

>>> print(lat.shape, A.shape)
(10, 3, 2) (60,)
>>> A_res = lat.mps2lat_values(A)
>>> A_res.shape
(10, 3, 2)
>>> A_res[tuple(lat.mps2lat_idx(5))] == A[5]
True

```

If you have a correlation function C [i, j], it gets just slightly more complicated:
```

>>> print(lat.shape, C.shape)
(10, 3, 2) (60, 60)

```
```

>>> lat.mps2lat_values(C, axes=[0, 1]).shape
(10, 3, 2, 10, 3, 2)

```

If the unit cell consists of different physical sites, an onsite operator might be defined only on one of the sites in the unit cell. Then you can use mps_idx_fix_u() to get the indices of sites it is defined on, measure the operator on these sites, and use the argument \(u\) of this function.
```

>>> u = 0
>>> idx_subset = lat.mps_idx_fix_u(u)
>>> A_u = A[idx_subset]
>>> A_u_res = lat.mps2lat_values(A_u, u=u)
>>> A_u_res.shape
(10, 3)
>>> np.all(A_res[:, :, u] == A_u_res[:, :])
True

```
mps2lat_values_masked ( \(A\), axes=- \(1, m p s \_i n d s=N o n e\), include_u=None)
Reshape/reorder an array \(A\) to replace an MPS index by lattice indices.
This is a generalization of mps2lat_values () allowing for the case of an arbitrary set of MPS indices present in each axis of \(A\).

\section*{Parameters}
- A (ndarray) - Some values.
- axes ((iterable of) int) - Chooses the axis of \(A\) which should be replaced. If multiple axes are given, you also need to give multiple index arrays as mps_inds.
- mps_inds ((list of) 1D ndarray) - Specifies for each axis in axes, for which MPS indices we have values in the corresponding axis of \(A\). Defaults to [np. arange (A. shape[ax]) for ax in axes]. For indices accross the MPS unit cell and "infinite" \(b c_{-} M P S\), we shift \(x \_0\) accordingly.
- include_u ((list of) bool) - Specifies for each axis in axes, whether the \(u\) index of the lattice should be included into the output array res_A. Defaults to len (self. unit_cell) > 1 .

Returns res_A - Reshaped and reordered copy of A. Such that MPS indices along the specified axes are replaced by lattice indices, i.e., if MPS index \(j\) maps to lattice site ( \(x 0, x 1, x 2\) ), then


\section*{Return type np.ma.MaskedArray}
mps_idx_fix_u ( \(u=\) None)
return an index array of MPS indices for which the site within the unit cell is \(u\).
If you have multiple sites in your unit-cell, an onsite operator is in general not defined for all sites. This functions returns an index array of the mps indices which belong to sites given by self. unit_cell[u].

Parameters \(\mathbf{u}\) (None / int) - Selects a site of the unit cell. None (default) means all sites.
Returns mps_idx - MPS indices for which self.site(i) is self.unit_cell[u]. Ordered ascending.

Return type array
mps_lat_idx_fix_u ( \(u=\) None)
Similar as mps_idx_fix_u(), but return also the corresponding lattice indices.

Parameters \(\mathbf{u}\) (None / int) - Selects a site of the unit cell. None (default) means all sites.

\section*{Returns}
- mps_idx (array) - MPS indices \(i\) for which self.site(i) is self. unit_cell[u].
- lat_idx (2D array) - The row \(j\) contains the lattice index (without \(u\) ) corresponding to mps_idx[j].

\section*{mps_sites()}

Return a list of sites for all MPS indices.
Equivalent to [self.site(i) for i in range(self.N_sites)].
This should be used for sites of 1D tensor networks (MPS, MPO,...).

\section*{multi_coupling_shape ( \(d x\) )}

Calculate correct shape of the strengths for a multi_coupling.
Parameters \(d \mathbf{x}\) (2D array, shape (N_ops, dim)) - dx[i, : ] is the translation vector in the lattice for the \(i\)-th operator. Corresponds to the \(d x\) of each operator given in the argument ops of tenpy.models.model.CouplingModel.add_multi_coupling().

\section*{Returns}
- coupling_shape (tuple of int) - Len dim. The correct shape for an array specifying the coupling strength. lat_indices has only rows within this shape.
- shift_lat_indices (array) - Translation vector from origin to the lower left corner of box spanned by \(d x\). (Unlike for coupling_shape () it can also contain entries \(>0\) )

\section*{number_nearest_neighbors ( \(u=0\) )}

Deprecated.
Deprecated since version 0.5.0: Use count_neighbors() instead.
```

number_next_nearest_neighbors ( }u=0\mathrm{ )

```

Deprecated.
Deprecated since version 0.5.0: Use count_neighbors () instead.

\section*{property order}

Defines an ordering of the lattice sites, thus mapping the lattice to a 1 D chain.
Each row of the array contains the lattice indices for one site, the order of the rows thus specifies a path through the lattice, along which an MPS will wind through through the lattice.
You can visualize the order with plot_order ().

\section*{ordering (order)}

Provide possible orderings of the \(N\) lattice sites.
This function can be overwritten by derived lattices to define additional orderings. The following orders are defined in this method:
\begin{tabular}{|l|l|l|}
\hline order & equivalent priority & equivalent snake_winding \\
\hline 'Cstyle' & \((0,1, \ldots\), dim-1, dim) & (False, \(\ldots\), False, False) \\
\hline 'default' & & \\
\hline 'snake' & \((0,1, \ldots, \operatorname{dim}-1, \operatorname{dim})\) & (True, \(\ldots\), True, True) \\
\hline 'snakeCstyle' & \\
\hline 'Fstyle' & \((\operatorname{dim}-1, \ldots, 1,0, \operatorname{dim})\) & (False, \(\ldots\), False, False) \\
\hline 'snakeFstyle' & \((\operatorname{dim}-1, \ldots, 1,0, \operatorname{dim})\) & (False,.., False, False) \\
\hline
\end{tabular}


Note: For lattices with a non-trivial unit cell (e.g. Honeycomb, Kagome), the grouped order might be more appropriate, see get_order_grouped ()

Parameters order (str | ('standard', snake_winding, priority) | ('grouped', groups, ...)) - Specifies the desired ordering using one of the strings of the above tables. Alternatively, an ordering is specified by a tuple with first entry specifying a function, 'standard' for get_order() and 'grouped' for get_order_grouped(), and other arguments in the tuple as specified in the documentation of these functions.
Returns order - the order to be used for order.
Return type array, shape ( \(\mathrm{N}, \mathrm{D}+1\) ), dtype np.intp

\section*{See also:}
get_order generates the order from equivalent priority and snake_winding.
get_order_grouped variant of get_order.
plot_order visualizes the resulting order.
plot_basis (ax, origin \(=(0.0,0.0)\), shade=None, **kwargs)
Plot arrows indicating the basis vectors of the lattice.

\section*{Parameters}
- ax (matplotlib.axes.Axes) - The axes on which we should plot.
- **kwargs - Keyword arguments for ax. arrow.
plot_bc_identified (ax, direction=- 1 , origin=None, cylinder_axis=False, **kwargs)
Mark two sites indified by periodic boundary conditions.
Works only for lattice with a 2 -dimensional basis.

\section*{Parameters}
- ax (matplotlib.axes.Axes) - The axes on which we should plot.
- direction (int) - The direction of the lattice along which we should mark the idenitified sites. If None, mark it along all directions with periodic boundary conditions.
- cylinder_axis (bool) - Whether to plot the cylinder axis as well.
- origin (None / np.ndarray) - The origin starting from where we mark the identified sites. Defaults to the first entry of unit_cell_positions.
- **kwargs - Keyword arguments for the used ax.plot.
plot_coupling (ax, coupling=None, wrap=False, **kwargs)
Plot lines connecting nearest neighbors of the lattice.

\section*{Parameters}
- ax (matplotlib.axes.Axes) - The axes on which we should plot.
- coupling (list of (u1, u2, dx)) - By default (None), use self. pairs['nearest_neighbors']. Specifies the connections to be plotted; iteating over lattice indices ( \(i 0, i 1, \ldots\) ), we plot a connection from the site (i0, i1, ...,
u1) to the site ( \(i 0+d x[0]\), \(i 1+d x[1], \ldots, u 2\) ), taking into account the boundary conditions.
- wrap (bool) - If True, plot couplings going around the boundary by directly connecting the sites it connects. This might be hard to see, as this puts lines from one end of the lattice to the other. If False, plot the couplings as dangling lines.
- **kwargs - Further keyword arguments given to ax.plot ().
plot_order (ax, order=None, textkwargs=\{'color': 'r'\}, **kwargs)
Plot a line connecting sites in the specified "order" and text labels enumerating them.

\section*{Parameters}
- ax (matplotlib. axes.Axes) - The axes on which we should plot.
- order (None / 2D array (self.N_sites, self.dim+1))- The order as returned by ordering (); by default (None) use order.
- textkwargs (None I dict) - If not None, we add text labels enumerating the sites in the plot. The dictionary can contain keyword arguments for ax.text ().
- **kwargs - Further keyword arguments given to ax.plot ().
```

plot_sites (ax, markers=['o', '^', 's', 'p', 'h', 'D'], **kwargs)

```

Plot the sites of the lattice with markers.

\section*{Parameters}
- ax (matplotlib. axes.Axes) - The axes on which we should plot.
- markers (list) - List of values for the keywork marker of ax.plot () to distinguish the different sites in the unit cell, a site \(u\) in the unit cell is plotted with a marker markers[u \% len(markers)].
- **kwargs - Further keyword arguments given to ax. plot ().
```

position(lat_idx)

```
return 'space' position of one or multiple sites.
Parameters lat_idx (ndarray, (. . , dim+1)) - Lattice indices.
Returns pos - The position of the lattice sites specified by lat_idx in real-space.
Return type ndarray, (..., dim)
possible_couplings ( \(u 1, u 2, d x\), strength=None)
Find possible MPS indices for two-site couplings.
For periodic boundary conditions ( \(\mathrm{bc}[\mathrm{a}]==\mathrm{False}\) ) the index x _a is taken modulo Ls [a] and runs through range (Ls [a]). For open boundary conditions, x_a is limited to \(0<=x \_a<L s[a]\) and \(0<=x \_a+d x[a]<\) lat.Ls[a].

\section*{Parameters}
- u1 (int) - Indices within the unit cell; the \(u 1\) and \(u 2\) of add_coupling ()
- u2 (int) - Indices within the unit cell; the \(u 1\) and \(u 2\) of add_coupling()
- \(\mathbf{d x}\) (array) - Length dim. The translation in terms of basis vectors for the coupling.
- strength (array_like / None) - If given, instead of returning lat_indices and coupling_shape directly return the correct strength_12.

\section*{Returns}
- mps1, mps2 (1D array) - For each possible two-site coupling the MPS indices for the \(u 1\) and \(u 2\).
- strength_vals (1D array) - (Only returend if strength is not None.) Such that for (i, j, s) in zip(mps1, mps2, strength_vals) : iterates over all possible couplings with \(s\) being the strength of that coupling. Couplings where strength_vals \(==0\). are filtered out.
- lat_indices ( \(2 D\) int array) - (Only returend if strength is None.) Rows of lat_indices correspond to entries of \(m p s l\) and \(m p s 2\) and contain the lattice indices of the "lower left corner" of the box containing the coupling.
- coupling_shape (tuple of int) - (Only returend if strength is None.) Len dim. The correct shape for an array specifying the coupling strength. lat_indices has only rows within this shape.
possible_multi_couplings (ops, strength=None)
Generalization of possible_couplings() to couplings with more than 2 sites.
Parameters ops (list of (opname, \(\mathrm{dx}, \mathrm{u})\) ) - Same as the argument ops of add_multi_coupling().

\section*{Returns}
- mps_ijkl (2D int array) - Each row contains MPS indices \(i, j, k, l, \ldots\). for each of the operators positions. The positions are defined by \(d x(\mathrm{j}, \mathrm{k}, \mathrm{l}, \ldots\) relative to \(i\) ) and boundary coundary conditions of self (how much the box for given \(d x\) can be shifted around without hitting a boundary - these are the different rows).
- strength_vals ( \(1 D\) array) - (Only returend if strength is not None.) Such that for (ijkl, s) in zip(mps_ijkl, strength_vals): iterates over all possible couplings with \(s\) being the strength of that coupling. Couplings where strength_vals \(=0\). are filtered out.
- lat_indices ( 2 D int array) - (Only returend if strength is None.) Rows of lat_indices correspond to rows of mps_ijkl and contain the lattice indices of the "lower left corner" of the box containing the coupling.
- coupling_shape (tuple of int) - (Only returend if strength is None.) Len dim. The correct shape for an array specifying the coupling strength. lat_indices has only rows within this shape.
save_hdf5 (hdf5_saver, h5gr, subpath)
Export self into a HDF5 file.
This method saves all the data it needs to reconstruct self with from_hdf5 ().
Specifically, it saves unit_cell, Ls, unit_cell_positions, basis, boundary_conditions, pairs under their name, bc_MPS as "boundary_conditions_MPS", and order as "order_for_MPS". Moreover, it saves dim and N_sites as HDF5 attributes.

\section*{Parameters}
- hdf5_saver (Hdf5Saver) - Instance of the saving engine.
- h5gr (:class `Group \({ }^{\text {) }}\) - HDF5 group which is supposed to represent self.
- subpath (str) - The name of \(h 5 g r\) with a ' / ' in the end.
site (i)
return Site instance corresponding to an MPS index \(i\)
test_sanity()
Sanity check.
Raises ValueErrors, if something is wrong.

\subsection*{19.1.7 Lattice}
- full name: tenpy.models.lattice.Lattice
- parent module: tenpy.models.lattice
- type: class

\section*{Inheritance Diagram}

\section*{Lattice}

\section*{Methods}
\begin{tabular}{|c|c|}
\hline Lattice.__init__(Ls, unit_cell[, order, bc, ...]) & Initialize self. \\
\hline Lattice.count_neighbors([u, key]) & Count e.g. \\
\hline Lattice.coupling_shape(dx) & Calculate correct shape of the strengths for a coupling. \\
\hline Lattice.distance(u1, u2, dx) & Get the distance for a given coupling between two sites in the lattice. \\
\hline Lattice.enlarge_mps_unit_cell([factor]) & Repeat the unit cell for infinite MPS boundary conditions; in place. \\
\hline ```
Lattice.find_coupling_pairs([max_dx,
...])
``` & Automatically find coupling pairs grouped by distances. \\
\hline Lattice.from_hdf5(hdf5_loader, h5gr, subpath) & Load instance from a HDF5 file. \\
\hline Lattice.lat2mps_idx(lat_idx) & Translate lattice indices \(\left(x \_0, \ldots, x_{-}\{D-1\}\right.\), u) to MPS index \(i\). \\
\hline Lattice.mps2lat_idx(i) & Translate MPS index \(i\) to lattice indices ( \(x \_0\), ...., x_\{dim-1\}, u). \\
\hline Lattice.mps2lat_values(A[, axes, u]) & Reshape/reorder \(A\) to replace an MPS index by lattice indices. \\
\hline Lattice.mps2lat_values_masked(A[, axes, ...]) & Reshape/reorder an array \(A\) to replace an MPS index by lattice indices. \\
\hline Lattice.mps_idx_fix_u([u]) & return an index array of MPS indices for which the site within the unit cell is \(u\). \\
\hline Lattice.mps_lat_idx_fix_u([u]) & Similar as mps_idx_fix_u(), but return also the corresponding lattice indices. \\
\hline Lattice.mps_sites() & Return a list of sites for all MPS indices. \\
\hline
\end{tabular}

Table 15 - continued from previous page
\begin{tabular}{|c|c|}
\hline Lattice.multi_coupling_shape(dx) & Calculate correct shape of the strengths for a multi_coupling. \\
\hline Lattice.number_nearest_neighbors([u]) & Deprecated. \\
\hline \multicolumn{2}{|l|}{Lattice.number_next_nearest_neighbors([uD) precated.} \\
\hline Lattice.ordering(order) & Provide possible orderings of the \(N\) lattice sites. \\
\hline Lattice.plot_basis(ax[, origin, shade]) & Plot arrows indicating the basis vectors of the lattice. \\
\hline Lattice.plot_bc_identified(ax[, direction, ...]) & Mark two sites indified by periodic boundary conditions. \\
\hline Lattice.plot_coupling(ax[, coupling, wrap]) & Plot lines connecting nearest neighbors of the lattice. \\
\hline Lattice.plot_order(ax[, order, textkwargs]) & Plot a line connecting sites in the specified "order" and text labels enumerating them. \\
\hline Lattice.plot_sites(ax[, markers]) & Plot the sites of the lattice with markers. \\
\hline Lattice.position(lat_idx) & return 'space' position of one or multiple sites. \\
\hline ```
Lattice.possible_couplings(u1, u2, dx[,
...])
``` & Find possible MPS indices for two-site couplings. \\
\hline Lattice.possible_multi_couplings(ops[, strength]) & Generalization of possible_couplings() to couplings with more than 2 sites. \\
\hline Lattice.save_hdf5(hdf5_saver, h5gr, subpath) & Export self into a HDF5 file. \\
\hline Lattice.site(i) & return Site instance corresponding to an MPS index \(i\) \\
\hline Lattice.test_sanity() & Sanity check. \\
\hline
\end{tabular}

\section*{Class Attributes and Properties}
\begin{tabular}{ll}
\hline Lattice.Lu & \begin{tabular}{l} 
the (expected) number of sites in the unit cell, \\
len (unit_cell).
\end{tabular} \\
\hline Lattice.boundary_conditions & \begin{tabular}{l} 
Human-readable list of boundary conditions from bc \\
and bc_shift.
\end{tabular} \\
\hline Lattice.dim & the dimension of the lattice \\
\hline Lattice.nearest_neighbors & \\
\hline Lattice.next_nearest_neighbors & \begin{tabular}{l} 
Defines an ordering of the lattice sites, thus mapping the \\
lattice to a 1D chain.
\end{tabular} \\
\hline Lattice.next_next_nearest_neighbors &
\end{tabular}
class tenpy.models.lattice.Lattice(Ls, unit_cell, order='default', bc='open', \(b c \_M P S=\) 'finite', basis=None, positions=None, nearest_neighbors=None, next_nearest_neighbors=None, next_next_nearest_neighbors=None, pairs=None)
Bases: ob ject
A general, regular lattice.
The lattice consists of a unit cell which is repeated in dim different directions. A site of the lattice is thus identified by lattice indices \(\left(x \_0, \ldots, x_{\_}\{\operatorname{dim}-1\}, u\right)\), where \(0<=x_{\_} l<L s[l]\) pick the position of the unit cell in the lattice and \(0<=u<\) len (unit_cell) picks the site within the unit cell. The site is located in 'space' at sum_l x_l*basis[l] + unit_cell_positions[u] (see position()). (Note that the position in space is only used for plotting, not for defining the couplings.)
In addition to the pure geometry, this class also defines an order of all sites. This order maps the lattice to
a finite 1D chain and defines the geometry of MPSs and MPOs. The MPS index \(i\) corresponds thus to the lattice sites given by ( \(\left.x \_0, \ldots, x_{-}\{\operatorname{dim}-1\}, u\right)=\) tuple(self.order[i]). Infinite boundary conditions of the MPS repeat in the first spatial direction of the lattice, i.e., if the site at ( \(x \_0, x \_1\), . . ., \(\left.x_{-}\{\operatorname{dim}-1\}, u\right)\) has MPS index \(i\), the site at at ( \(x_{-} 0+\operatorname{Ls}[0], x_{-} 1, \ldots, x_{-}\{d i m-1\}\), \(u\) ) corresponds to MPS index i + N_sites. Use mps2lat_idx () and lat2mps_idx () for conversion of indices. The function mps2lat_values() performs the necessary reshaping and re-ordering from arrays indexed in MPS form to arrays indexed in lattice form.

Deprecated since version 0.5.0: The parameters and attributes nearest_neighbors, next_nearest_neighbors and next_next_nearest_neighbors are deprecated. Instead, we use a dictionary pairs with those names as keys and the corresponding values as specified before.

\section*{Parameters}
- Ls (list of int) - the length in each direction
- unit_cell (list of Site) - The sites making up a unit cell of the lattice. If you want to specify it only after initialization, use None entries in the list.
- order (str l ('standard', snake_winding, priority) l ('grouped', groups, ...))-A string or tuple specifying the order, given to ordering().
- bc ((iterable of) \{'open' | 'periodic' / int \})-Boundary conditions in each direction of the lattice. A single string holds for all directions. An integer shift means that we have periodic boundary conditions along this direction, but shift/tilt by -shift*lattice.basis[0] (~cylinder axis for bc_MPS='infinite') when going around the boundary along this direction.
- bc_MPS ('finite' / 'segment' / 'infinite')-Boundary conditions for an MPS/MPO living on the ordered lattice. If the system is 'infinite', the infinite direction is always along the first basis vector (justifying the definition of \(N_{-}\)rings and N_sites_per_ring).
- basis (iterable of 1D arrays) - For each direction one translation vectors shifting the unit cell. Defaults to the standard ONB np. eye (dim).
- positions (iterable of \(1 D\) arrays) - For each site of the unit cell the position within the unit cell. Defaults to np.zeros ((len (unit_cell), dim)).
- nearest_neighbors (None I list of (u1, u2, dx)) - Deprecated. Specify as pairs['nearest_neighbors'] instead.
- next_nearest_neighbors (None l list of (u1, u2, dx)) - Deprecated. Specify as pairs['next_nearest_neighbors'] instead.
- next_next_nearest_neighbors (None l list of (u1, u2, dx)) - Deprecated. Specify as pairs['next_next_nearest_neighbors'] instead.
- pairs (dict) - Of the form \{'nearest_neighbors': [(u1, u2, dx), ...], ...\}. Typical keys are 'nearest_neighbors', 'next_nearest_neighbors'. For each of them, it specifies a list of tuples ( \(u 1, \mathrm{u} 2, \mathrm{dx}\) ) which can be used as parameters for add_coupling() to generate couplings over each pair of ,e.g., ' nearest_neighbors'. Note that this adds couplings for each pair only in one direction!

Ls
the length in each direction.
Type tuple of int
shape
the 'shape' of the lattice, same as Ls + (len (unit_cell), )

Type tuple of int
N_cells
the number of unit cells in the lattice, np.prod (self.Ls).
Type int
N_sites
the number of sites in the lattice, np.prod (self. shape).
Type int
N_sites_per_ring
Defined as N_sites / Ls [0], for an infinite system the number of cites per "ring".
Type int

\section*{N_rings}

Alias for Ls [0], for an infinite system the number of "rings" in the unit cell.
Type int

\section*{unit_cell}
the sites making up a unit cell of the lattice.
Type list of Site
bc
Boundary conditions of the couplings in each direction of the lattice, translated into a bool array with the global bc_choices.

Type bool ndarray

\section*{bc_shift}

The shift in x -direction when going around periodic boundaries in other directions.
Type None I ndarray(int)
bc_MPS
Boundary conditions for an MPS/MPO living on the ordered lattice. If the system is 'infinite', the infinite direction is always along the first basis vector (justifying the definition of \(N\) _rings and N_sites_per_ring).

Type 'finite'। 'segment' I 'infinite'

\section*{basis}
translation vectors shifting the unit cell. The row \(i\) gives the vector shifting in direction \(i\).
Type ndarray (dim, Dim)
unit_cell_positions
for each site in the unit cell a vector giving its position within the unit cell.
Type ndarray, shape (len(unit_cell), Dim)
pairs
See above.
Type dict
_order
The place where order is stored.
Type ndarray (N_sites, dim+1)
```

_strides

```
necessary for lat2mps_idx().
Type ndarray (dim, )
_perm
permutation needed to make \(\operatorname{order}\) lexsorted,_perm \(=n p\).lexsort(_order.T).
Type ndarray ( N, )
```

_mps2lat_vals_idx

```
index array for reshape/reordering in mps2lat_vals()
Type ndarray shape
_mps_fix_u
for each site of the unit cell an index array selecting the mps indices of that site.
Type tuple of ndarray (N_cells, ) np.intp
_mps2lat_vals_idx_fix_u
similar as _mps2lat_vals_idx, but for a fixed \(u\) picking a site from the unit cell.
Type tuple of ndarray of shape \(L s\)
Lu = None
the (expected) number of sites in the unit cell, len (unit_cell).
test_sanity()
Sanity check.
Raises ValueErrors, if something is wrong.
save_hdf5 (hdf5_saver, h5gr, subpath)
Export self into a HDF5 file.
This method saves all the data it needs to reconstruct self with from_hof5 ().
Specifically, it saves unit_cell, Ls, unit_cell_positions, basis, boundary_conditions, pairs under their name, bc_MPS as "boundary_conditions_MPS
", and order as "order_for_MPS". Moreover, it saves dim and N_sites as HDF5 attributes.

\section*{Parameters}
- hdf5_saver (Hdf5Saver) - Instance of the saving engine.
- h5gr (: class `Group`) - HDF5 group which is supposed to represent self.
- subpath (str) - The name of \(h 5 g r\) with a ' /' in the end.
classmethod from_hdf5 (hdf5_loader, h5gr, subpath)
Load instance from a HDF5 file.
This method reconstructs a class instance from the data saved with save_hdf5 ().

\section*{Parameters}
- hdf5_loader (Hdf5Loader) - Instance of the loading engine.
- h5gr (Group) - HDF5 group which is represent the object to be constructed.
- subpath (str) - The name of \(h 5 g r\) with a '/' in the end.

Returns obj - Newly generated class instance containing the required data.
Return type cls

\section*{property dim}
the dimension of the lattice
property order
Defines an ordering of the lattice sites, thus mapping the lattice to a 1D chain.
Each row of the array contains the lattice indices for one site, the order of the rows thus specifies a path through the lattice, along which an MPS will wind through through the lattice.

You can visualize the order with plot_order ().

\section*{ordering (order)}

Provide possible orderings of the \(N\) lattice sites.
This function can be overwritten by derived lattices to define additional orderings. The following orders are defined in this method:
\begin{tabular}{|c|c|c|}
\hline order & equivalent priority & equivalent snake_winding \\
\hline 'Cstyle' & ( \(0,1, \ldots\), dim-1, dim) & (False, ..., False, False) \\
\hline 'default' & & \\
\hline 'snake' & ( \(0,1, \ldots\), dim-1, dim) & (True, ..., True, True) \\
\hline 'snakeCstyle' & & \\
\hline 'Fstyle' & (dim-1, .., 1, 0, dim) & (False, ..., False, False) \\
\hline 'snakeFstyle' & \((\operatorname{dim}-1, \ldots, 1,0, \operatorname{dim})\) & (False, ..., False, False) \\
\hline
\end{tabular}


Note: For lattices with a non-trivial unit cell (e.g. Honeycomb, Kagome), the grouped order might be more appropriate, see get_order_grouped ()

Parameters order (str | ('standard', snake_winding, priority) | ('grouped', groups, ...)) - Specifies the desired ordering using one of the strings of the above tables. Alternatively, an ordering is specified by a tuple with first entry specifying a function, 'standard' for get_order() and 'grouped' for get_order_grouped(), and other arguments in the tuple as specified in the documentation of these functions.

Returns order - the order to be used for order.
Return type array, shape ( \(\mathrm{N}, \mathrm{D}+1\) ), dtype np.intp

\section*{See also:}
get_order generates the order from equivalent priority and snake_winding.
get_order_grouped variant of get_order.
plot_order visualizes the resulting order.

\section*{property boundary_conditions}

Human-readable list of boundary conditions from bc and bc_shift.
Returns boundary_conditions - List of "open" or "periodic", one entry for each direction of the lattice.
Return type list of str
enlarge_mps_unit_cell (factor=2)
Repeat the unit cell for infinite MPS boundary conditions; in place.
Parameters factor (int) - The new number of sites in the MPS unit cell will be increased from \(N\) _sites to factor*N_sites_per_ring. Since MPS unit cells are repeated in the \(x\)-direction in our convetion, the lattice shape goes from ( \(\mathrm{Lx}, \mathrm{Ly}, \ldots, \mathrm{Lu}\) ) to (Lx*factor, Ly, ..., Lu).
position (lat_idx)
return 'space' position of one or multiple sites.
Parameters lat_idx (ndarray, (. . . , dim+1)) - Lattice indices.
Returns pos - The position of the lattice sites specified by lat_idx in real-space.
Return type ndarray, (..., dim)
site (i)
return Site instance corresponding to an MPS index \(i\)
mps_sites()
Return a list of sites for all MPS indices.
Equivalent to [self.site(i) for i in range(self.N_sites)].
This should be used for sites of 1D tensor networks (MPS, MPO,....).
mps2lat_idx (i)
Translate MPS index \(i\) to lattice indices ( \(\mathrm{x} \_0, \ldots, \mathrm{x}_{-}\{\)dim-1\}, \(u\) ).
Parameters i (int | array_like of int)-MPS index/indices.

Returns lat_idx - First dimensions like \(i\), last dimension has len dim +1 and contains the lattice indices " \(\left(x \_0, \ldots, x_{\_}\{d i m-1\}, u\right)\) ` corresponding to \(i\). For \(i\) accross the MPS unit cell and "infinite" \(b c_{-} M P S\), we shift \(x_{-} 0\) accordingly.

\section*{Return type array}
lat2mps_idx (lat_idx)
Translate lattice indices \(\left(\mathrm{x}_{-} 0, \ldots, \mathrm{x}_{-}\{\mathrm{D}-1\}, \mathrm{u}\right)\) to MPS index \(i\).
Parameters lat_idx (array_like [.., dim+1]) - The last dimension corresponds to lattice indices ( \(\left.x \_0, \ldots, x_{-}\{D-1\}, u\right)\). All lattice indices should be positive and smaller than the corresponding entry in self.shape. Exception: for "infinite" \(b c_{-} M P S\), an \(x \_0\) outside indicates shifts accross the boundary.

Returns \(\mathbf{i}\) - MPS index/indices corresponding to lat_idx. Has the same shape as lat_idx without the last dimension.

Return type array_like
mps_idx_fix_u(u=None)
return an index array of MPS indices for which the site within the unit cell is \(u\).
If you have multiple sites in your unit-cell, an onsite operator is in general not defined for all sites. This functions returns an index array of the mps indices which belong to sites given by self. unit_cell[u].

Parameters \(\mathbf{u}\) (None / int) - Selects a site of the unit cell. None (default) means all sites.
Returns mps_idx - MPS indices for which self.site(i) is self.unit_cell[u]. Ordered ascending.
Return type array
mps_lat_idx_fix_u (u=None)
Similar as mps_idx_fix_u(), but return also the corresponding lattice indices.
Parameters u(None / int)-Selects a site of the unit cell. None (default) means all sites.

\section*{Returns}
- mps_idx (array) - MPS indices \(i\) for which self.site(i) is self. unit_cell[u].
- lat_idx (2D array) - The row \(j\) contains the lattice index (without \(u\) ) corresponding to mps_idx[j].
mps2lat_values \((A\), axes \(=0, u=\) None \()\)
Reshape/reorder \(A\) to replace an MPS index by lattice indices.

\section*{Parameters}
- A (ndarray) - Some values. Must have A.shape [axes] = self.N_sites if \(u\) is None, or A.shape[axes] = self.N_cells if \(u\) is an int.
- axes ( (iterable of) int) - chooses the axis which should be replaced.
- u (None | int) - Optionally choose a subset of MPS indices present in the axes of \(A\), namely the indices corresponding to self.unit_cell[u], as returned by \(m p s \_i d x \_f i x \_u()\). The resulting array will not have the additional dimension(s) of \(u\).
Returns res_A - Reshaped and reordered verions of A. Such that MPS indices along the specified axes are replaced by lattice indices, i.e., if MPS index \(j\) maps to lattice site ( \(x 0, x 1, x 2\) ), then res_A [..., \(x 0, x 1, x 2, \ldots]=A[\ldots, j, \ldots]\).

Return type ndarray

\section*{Examples}

Say you measure expection values of an onsite term for an MPS, which gives you an 1D array \(A\), where \(A[i]\) is the expectation value of the site given by self.mps2lat_idx(i). Then this function gives you the expectation values ordered by the lattice:
```

>>> print(lat.shape, A.shape)
(10, 3, 2) (60,)
>>> A_res = lat.mps2lat_values(A)
>>> A_res.shape
(10, 3, 2)
>>> A_res[tuple(lat.mps2lat_idx(5))] == A[5]
True

```

If you have a correlation function \(C[i, j]\), it gets just slightly more complicated:
```

>>> print(lat.shape, C.shape)
(10, 3, 2) (60, 60)
>>> lat.mps2lat_values(C, axes=[0, 1]).shape
(10, 3, 2, 10, 3, 2)

```

If the unit cell consists of different physical sites, an onsite operator might be defined only on one of the sites in the unit cell. Then you can use mps_idx_fix_u() to get the indices of sites it is defined on, measure the operator on these sites, and use the argument \(u\) of this function.
```

>>> u = 0
>>> idx_subset = lat.mps_idx_fix_u(u)
>>> A_u = A[idx_subset]
>>> A_u_res = lat.mps2lat_values(A_u, u=u)
>>> A_u_res.shape
(10, 3)
>>> np.all(A_res[:, :, u] == A_u_res[:, :])
True

```
mps2lat_values_masked ( \(A\), axes=- 1, mps_inds=None, include_u=None) Reshape/reorder an array \(A\) to replace an MPS index by lattice indices.
This is a generalization of mps2lat_values () allowing for the case of an arbitrary set of MPS indices present in each axis of \(A\).

\section*{Parameters}
- A (ndarray) - Some values.
- axes ((iterable of) int) - Chooses the axis of \(A\) which should be replaced. If multiple axes are given, you also need to give multiple index arrays as mps_inds.
- mps_inds ((list of) 1D ndarray) - Specifies for each axis in axes, for which MPS indices we have values in the corresponding axis of \(A\). Defaults to [np. arange (A. shape[ax]) for ax in axes]. For indices accross the MPS unit cell and "infinite" \(b c_{-} M P S\), we shift \(x \_0\) accordingly.
- include_u ((list of) bool) - Specifies for each axis in axes, whether the \(u\) index of the lattice should be included into the output array res_A. Defaults to len (self. unit_cell) > 1 .

Returns res_A - Reshaped and reordered copy of A. Such that MPS indices along the specified axes are replaced by lattice indices, i.e., if MPS index \(j\) maps to lattice site ( \(x 0, x 1, x 2\) ), then res_A[..., x0, x1, x2, ...] = A[..., mps_inds[j], ...].
Return type np.ma.MaskedArray
count_neighbors ( \(u=0\), key='nearest_neighbors')
Count e.g. the number of nearest neighbors for a site in the bulk.

\section*{Parameters}
- u (int) - Specifies the site in the unit cell, for which we should count the number of neighbors (or whatever key specifies).
- key (str) - Key of pairs to select what to count.

Returns number - Number of nearest neighbors (or whatever key specified) for the \(u\)-th site in the unit cell, somewhere in the bulk of the lattice. Note that it might not be the correct value at the edges of a lattice with open boundary conditions.

\section*{Return type int}
number_nearest_neighbors ( \(u=0\) )
Deprecated.
Deprecated since version 0.5.0: Use count_neighbors () instead.
number_next_nearest_neighbors ( \(u=0\) )
Deprecated.
Deprecated since version 0.5.0: Use count_neighbors () instead.
distance ( \(u 1, u 2, d x\) )
Get the distance for a given coupling between two sites in the lattice.
The \(u 1, u 2, d x\) parameters are defined in analogy with add_coupling(), i.e., this function calculates the distance between a pair of operators added with add_coupling (using the basis and unit_cell_positions of the lattice).

Warning: This function ignores "wrapping" arround the cylinder in the case of periodic boundary conditions.

\section*{Parameters}
- u1 (int) - Indices within the unit cell; the \(u 1\) and \(u 2\) of add_coupling()
- u2 (int) - Indices within the unit cell; the \(u 1\) and \(u 2\) of add_coupling ()
- \(\mathbf{d x}\) (array) - Length dim. The translation in terms of basis vectors for the coupling.

Returns distance - The distance between site at lattice indices [ \(\mathrm{x}, \mathrm{y}, \mathrm{ul}\) ] and [ \(\mathrm{x}+\) \(d x[0], y+d x[1], u 2]\), ignoring any boundary effects.
Return type float
find_coupling_pairs (max_dx=3, cutoff=None, eps=1e-10)
Automatically find coupling pairs grouped by distances.
Given the unit_cell_positions and basis, the coupling pairs of nearest_neighbors, next_nearest_neighbors etc at a given distance are basically fixed (although not uniquely, since we take out
half of them to avoid double-counting couplings in both directions A_i B_j + B_i A_i). This function iterates through all possible couplings up to a given cutoff distance and then determines the possible pairs at fixed distances (up to round-off errors).

\section*{Parameters}
- max_dx (int) - Maximal index for each index of \(d x\) to iterate over. You need large enough values to include every possible coupling up to the desired distance, but choosing it too large might make this function run for a long time.
- cutoff (float) - Maximal distance (in the units in which basis and unit_cell_positions is given).
- eps (float) - Tolerance up to which to distances are considered the same.

Returns coupling_pairs - Keys are distances of nearest-neighbors, next-nearest-neighbors etc. Values are [(u1, u2, dx), ...] as in pairs.

Return type dict
coupling_shape ( \(d x\) )
Calculate correct shape of the strengths for a coupling.
Parameters \(\mathbf{d x}\) (tuple of int) - Translation vector in the lattice for a coupling of two operators. Corresponds to \(d x\) argument of tenpy.models.model. CouplingModel. add_multi_coupling().

\section*{Returns}
- coupling_shape (tuple of int) - Len dim. The correct shape for an array specifying the coupling strength. lat_indices has only rows within this shape.
- shift_lat_indices (array) - Translation vector from origin to the lower left corner of box spanned by \(d x\).
possible_couplings ( \(u 1, u 2, d x\), strength=None)
Find possible MPS indices for two-site couplings.
For periodic boundary conditions ( \(\mathrm{bc}[\mathrm{a}]==\mathrm{False}\) ) the index x _a is taken modulo Ls [a] and runs through range (Ls [a]). For open boundary conditions, \(x\) _a is limited to \(0<=x_{\text {_ }}<\) Ls [a] and \(0<=x \_a+d x[a]<\) lat.Ls [a].

\section*{Parameters}
- u1 (int) - Indices within the unit cell; the \(u 1\) and \(u 2\) of add_coupling()
- u2 (int) - Indices within the unit cell; the \(u 1\) and \(u 2\) of add_coupling ()
- \(\mathbf{d} \mathbf{x}\) (array) - Length dim. The translation in terms of basis vectors for the coupling.
- strength (array_like / None) - If given, instead of returning lat_indices and coupling_shape directly return the correct strength_12.

\section*{Returns}
- mps1, mps2 (1D array) - For each possible two-site coupling the MPS indices for the \(u 1\) and \(u 2\).
- strength_vals (lD array) - (Only returend if strength is not None.) Such that for (i, j, s) in zip (mps1, mps2, strength_vals): iterates over all possible couplings with \(s\) being the strength of that coupling. Couplings where strength_vals \(==0\). are filtered out.
- lat_indices ( \(2 D\) int array) - (Only returend if strength is None.) Rows of lat_indices correspond to entries of \(m p s l\) and \(m p s 2\) and contain the lattice indices of the "lower left corner" of the box containing the coupling.
- coupling_shape (tuple of int) - (Only returend if strength is None.) Len dim. The correct shape for an array specifying the coupling strength. lat_indices has only rows within this shape.
multi_coupling_shape ( \(d x\) )
Calculate correct shape of the strengths for a multi_coupling.
Parameters \(d \mathbf{x}\) (2D array, shape (N_ops, dim)) - dx[i, :] is the translation vector in the lattice for the \(i\)-th operator. Corresponds to the \(d x\) of each operator given in the argument ops of tenpy.models.model.CouplingModel.add_multi_coupling().

\section*{Returns}
- coupling_shape (tuple of int) - Len dim. The correct shape for an array specifying the coupling strength. lat_indices has only rows within this shape.
- shift_lat_indices (array) - Translation vector from origin to the lower left corner of box spanned by \(d x\). (Unlike for coupling_shape () it can also contain entries \(>0\) )
possible_multi_couplings (ops, strength=None)
Generalization of possible_couplings () to couplings with more than 2 sites.
Parameters ops (list of (opname, \(\mathrm{dx}, \mathrm{u})\) ) - Same as the argument ops of add_multi_coupling().

\section*{Returns}
- mps_ijkl (2D int array) - Each row contains MPS indices \(i, j, k, l, \ldots\). for each of the operators positions. The positions are defined by \(d x(\mathrm{j}, \mathrm{k}, 1, \ldots\) relative to \(i\) ) and boundary coundary conditions of self (how much the box for given \(d x\) can be shifted around without hitting a boundary - these are the different rows).
- strength_vals ( \(1 D\) array) - (Only returend if strength is not None.) Such that for (ijkl, s) in zip(mps_ijkl, strength_vals): iterates over all possible couplings with \(s\) being the strength of that coupling. Couplings where strength_vals \(==0\). are filtered out.
- lat_indices ( \(2 D\) int array) - (Only returend if strength is None.) Rows of lat_indices correspond to rows of \(m p s \_i j k l\) and contain the lattice indices of the "lower left corner" of the box containing the coupling.
- coupling_shape (tuple of int) - (Only returend if strength is None.) Len dim. The correct shape for an array specifying the coupling strength. lat_indices has only rows within this shape.
```

plot_sites (ax,markers=['o', '^', 's', 'p', 'h', 'D'], **kwargs)

```

Plot the sites of the lattice with markers.

\section*{Parameters}
- ax (matplotlib.axes.Axes) - The axes on which we should plot.
- markers (list) - List of values for the keywork marker of ax.plot () to distinguish the different sites in the unit cell, a site \(u\) in the unit cell is plotted with a marker markers[u \% len(markers)].
- **kwargs - Further keyword arguments given to ax.plot ().
plot_order (ax, order=None, textkwargs=\{'color': 'r'\}, **kwargs)
Plot a line connecting sites in the specified "order" and text labels enumerating them.

\section*{Parameters}
- ax (matplotlib. axes.Axes) - The axes on which we should plot.
- order (None / 2D array (self.N_sites, self.dim+1))- The order as returned by ordering(); by default (None) use order.
- textkwargs (None I dict) - If not None, we add text labels enumerating the sites in the plot. The dictionary can contain keyword arguments for ax.text ().
- **kwargs - Further keyword arguments given to ax.plot ().
plot_coupling (ax, coupling=None, wrap=False, **kwargs)
Plot lines connecting nearest neighbors of the lattice.

\section*{Parameters}
- ax (matplotlib. axes.Axes) - The axes on which we should plot.
- coupling (list of (ul, \(u 2, d x)\) ) - By default (None), use self. pairs['nearest_neighbors']. Specifies the connections to be plotted; iteating over lattice indices ( \(i 0, i 1, \ldots\) ), we plot a connection from the site (i0, i1, ..., u1) to the site ( \(i 0+d x[0]\), \(i 1+d x[1], \ldots, u 2\) ), taking into account the boundary conditions.
- wrap (bool) - If True, plot couplings going around the boundary by directly connecting the sites it connects. This might be hard to see, as this puts lines from one end of the lattice to the other. If False, plot the couplings as dangling lines.
- **kwargs - Further keyword arguments given to ax.plot ().
plot_basis (ax, origin=(0.0, 0.0), shade \(=\) None, \(* *\) kwargs \()\)
Plot arrows indicating the basis vectors of the lattice.

\section*{Parameters}
- ax (matplotlib. axes.Axes) - The axes on which we should plot.
- **kwargs - Keyword arguments for ax. arrow.
plot_bc_identified (ax, direction=- 1, origin=None, cylinder_axis=False, **kwargs)
Mark two sites indified by periodic boundary conditions.
Works only for lattice with a 2-dimensional basis.

\section*{Parameters}
- ax (matplotlib. axes.Axes) - The axes on which we should plot.
- direction (int) - The direction of the lattice along which we should mark the idenitified sites. If None, mark it along all directions with periodic boundary conditions.
- cylinder_axis (bool) - Whether to plot the cylinder axis as well.
- origin (None / np.ndarray) - The origin starting from where we mark the identified sites. Defaults to the first entry of unit_cell_positions.
- **kwargs - Keyword arguments for the used ax.plot.

\subsection*{19.1.8 SimpleLattice}
- full name: tenpy.models.lattice.SimpleLattice
- parent module: tenpy.models.lattice
- type: class

\section*{Inheritance Diagram}


\section*{Methods}
\begin{tabular}{ll}
\hline SimpleLattice.__init__(Ls, site, ***kwargs) & Initialize self. \\
\hline SimpleLattice.count_neighbors([u, key]) & Count e.g. \\
\hline SimpleLattice.coupling_shape(dx) & Calculate correct shape of the strengths for a coupling. \\
\hline SimpleLattice.distance(u1, u2, dx) & \begin{tabular}{l} 
Get the distance for a given coupling between two sites \\
in the lattice.
\end{tabular} \\
\hline SimpleLattice.enlarge_mps_unit_cell([factdRłpeat the unit cell for infinite MPS boundary condi- \\
tions; in place.
\end{tabular}

Table 17-continued from previous page
\begin{tabular}{|c|c|}
\hline SimpleLattice.multi_coupling_shape(dx) & Calculate correct shape of the strengths for multi_coupling. \\
\hline \multicolumn{2}{|l|}{SimpleLattice.number_nearest_neighbors([D])precated.} \\
\hline \multicolumn{2}{|l|}{SimpleLattice.number_next_nearest_neighDepre(dat)d.} \\
\hline SimpleLattice.ordering(order) & Provide possible orderings of the \(N\) lattice sites. \\
\hline  & Plot arrows indicating the basis vectors of the lattice. \\
\hline ```
SimpleLattice.plot_bc_identified(ax[,
...])
``` & Mark two sites indified by periodic boundary conditions. \\
\hline SimpleLattice.plot_coupling(ax[, coupling, wrap]) & Plot lines connecting nearest neighbors of the lattice. \\
\hline SimpleLattice.plot_order(ax[, order, textkwargs]) & Plot a line connecting sites in the specified "order" and text labels enumerating them. \\
\hline SimpleLattice.plot_sites(ax[, markers]) & Plot the sites of the lattice with markers. \\
\hline SimpleLattice.position(lat_idx) & return 'space' position of one or multiple sites. \\
\hline SimpleLattice.possible_couplings(u1, u2, dx) & Find possible MPS indices for two-site couplings. \\
\hline \multicolumn{2}{|l|}{\[
\begin{gathered}
\text { SimpleLattice.possible_multi_couplings(opsheralization of possible_couplings () to cou- } \\
\text { plings with more than } 2 \text { sites. }
\end{gathered}
\]} \\
\hline SimpleLattice.save_hdf5(hdf5_saver, h5gr, ...) & Export self into a HDF5 file. \\
\hline SimpleLattice.site(i) & return Site instance corresponding to an MPS index \(i\) \\
\hline SimpleLattice.test_sanity() & Sanity check. \\
\hline
\end{tabular}

\section*{Class Attributes and Properties}
\begin{tabular}{ll}
\hline SimpleLattice.Lu & \begin{tabular}{l} 
the (expected) number of sites in the unit cell, \\
len_(unit_cell).
\end{tabular} \\
\hline SimpleLattice.boundary_conditions & \begin{tabular}{l} 
Human-readable list of boundary conditions from bc \\
and bc_shift.
\end{tabular} \\
\hline SimpleLattice.dim & The dimension of the lattice. \\
\hline SimpleLattice.nearest_neighbors & \\
\hline SimpleLattice.next_nearest_neighbors & \begin{tabular}{l} 
Defines an ordering of the lattice sites, thus mapping the \\
lattice to a 1D chain.
\end{tabular} \\
\hline SimpleLattice.next_next_nearest_neighbors
\end{tabular}
class tenpy.models.lattice.SimpleLattice (Ls, site, **kwargs)
Bases: tenpy.models.lattice.Lattice
A lattice with a unit cell consiting of just a single site.
In many cases, the unit cell consists just of a single site, such that the the last entry of \(u\) of an 'lattice index' can only be 0 . From the point of internal algorithms, we handle this class like a Lattice - in that way we don't need to distinguish special cases in the algorithms.

Yet, from the point of a tenpy user, for example if you measure an expectation value on each site in a SimpleLattice, you expect to get an ndarray of dimensions self.Ls, not self.shape. To avoid that problem, SimpleLattice overwrites just the meaning of \(u=N o n e\) in mps2lat_values () to be the same as \(u=0\).

\section*{Parameters}
- Ls (list of int) - the length in each direction
- site (Site) - the lattice site. The unit_cell of the Lattice is just [site].
- **kwargs - Additional keyword arguments given to the Lattice. If order is specified in the form ('standard', snake_windingi, priority), the snake_winding and priority should only be specified for the spatial directions. Similarly, positions can be specified as a single vector.
\(\mathrm{Lu}=1\)
the (expected) number of sites in the unit cell, len (unit_cell).
mps2lat_values \((A\), axes \(=0, u=\) None)
same as Lattice.mps2lat_values(), but ignore \(u\), setting it to 0 .
property boundary_conditions
Human-readable list of boundary conditions from bc and bc_shift.
Returns boundary_conditions - List of "open" or "periodic", one entry for each direction of the lattice.

Return type list of str
count_neighbors ( \(u=0\), key='nearest_neighbors')
Count e.g. the number of nearest neighbors for a site in the bulk.

\section*{Parameters}
- u (int) - Specifies the site in the unit cell, for which we should count the number of neighbors (or whatever key specifies).
- key (str) - Key of pairs to select what to count.

Returns number - Number of nearest neighbors (or whatever key specified) for the \(u\)-th site in the unit cell, somewhere in the bulk of the lattice. Note that it might not be the correct value at the edges of a lattice with open boundary conditions.

\section*{Return type int}
coupling_shape ( \(d x\) )
Calculate correct shape of the strengths for a coupling.
Parameters \(\mathbf{d x}\) (tuple of int) - Translation vector in the lattice for a coupling of two operators. Corresponds to \(d x\) argument of tenpy.models.model. CouplingModel. add_multi_coupling().

\section*{Returns}
- coupling_shape (tuple of int) - Len dim. The correct shape for an array specifying the coupling strength. lat_indices has only rows within this shape.
- shift_lat_indices (array) - Translation vector from origin to the lower left corner of box spanned by \(d x\).
property dim
The dimension of the lattice.
distance ( \(u 1, u 2, d x\) )
Get the distance for a given coupling between two sites in the lattice.
The \(u 1, u 2, d x\) parameters are defined in analogy with add_coupling(), i.e., this function calculates the distance between a pair of operators added with add_coupling (using the basis and unit_cell_positions of the lattice).

Warning: This function ignores "wrapping" arround the cylinder in the case of periodic boundary conditions.

\section*{Parameters}
- u1 (int) - Indices within the unit cell; the \(u 1\) and \(u 2\) of add_coupling ()
- u2 (int) - Indices within the unit cell; the \(u 1\) and \(u 2\) of add_coupling()
- \(\mathbf{d x}\) (array) - Length dim. The translation in terms of basis vectors for the coupling.

Returns distance - The distance between site at lattice indices [ \(\mathrm{x}, \mathrm{y}, \mathrm{ul}\) ] and [ \(\mathrm{x}+\) \(d x[0], y+d x[1], u 2]\), ignoring any boundary effects.

Return type float

\section*{enlarge_mps_unit_cell (factor=2)}

Repeat the unit cell for infinite MPS boundary conditions; in place.
Parameters factor (int) - The new number of sites in the MPS unit cell will be increased from \(N\) _sites to factor*N_sites_per_ring. Since MPS unit cells are repeated in the \(x\)-direction in our convetion, the lattice shape goes from ( \(L x, L y, \ldots, L u\) ) to (Lx*factor, Ly, ..., Lu).
find_coupling_pairs (max_dx=3, cutoff=None, eps=1e-10)
Automatically find coupling pairs grouped by distances.
Given the unit_cell_positions and basis, the coupling pairs of nearest_neighbors, next_nearest_neighbors etc at a given distance are basically fixed (although not uniquely, since we take out half of them to avoid double-counting couplings in both directions A_i B_j + B_i A_i). This function iterates through all possible couplings up to a given cutoff distance and then determines the possible pairs at fixed distances (up to round-off errors).

\section*{Parameters}
- max_dx (int) - Maximal index for each index of \(d x\) to iterate over. You need large enough values to include every possible coupling up to the desired distance, but choosing it too large might make this function run for a long time.
- cutoff (float) - Maximal distance (in the units in which basis and unit_cell_positions is given).
- eps (float) - Tolerance up to which to distances are considered the same.

Returns coupling_pairs - Keys are distances of nearest-neighbors, next-nearest-neighbors etc.
Values are [(u1, u2, dx), ...] as in pairs.
Return type dict
classmethod from_hdf5 (hdf5_loader, h5gr, subpath)
Load instance from a HDF5 file.
This method reconstructs a class instance from the data saved with save_hdf5 ().

\section*{Parameters}
- hdf5_loader (Hdf5Loader) - Instance of the loading engine.
- h5gr (Group) - HDF5 group which is represent the object to be constructed.
- subpath (str) - The name of \(h 5 g r\) with a ' /' in the end.

Returns obj - Newly generated class instance containing the required data.

Return type cls
lat2mps_idx (lat_idx)
Translate lattice indices ( \(\mathrm{x} \_0, \ldots, \mathrm{x}_{-}\{\mathrm{D}-1\}, \mathrm{u}\) ) to MPS index \(i\).
Parameters lat_idx (array_like [.., dim+1]) - The last dimension corresponds to lattice indices \(\left(x_{-} 0, \ldots, x_{-}\{D-1\}, u\right)\). All lattice indices should be positive and smaller than the corresponding entry in self.shape. Exception: for "infinite" \(b c_{-} M P S\), an \(x \_0\) outside indicates shifts accross the boundary.

Returns i - MPS index/indices corresponding to lat_idx. Has the same shape as lat_idx without the last dimension.

Return type array_like
mps2lat_idx (i)
Translate MPS index \(i\) to lattice indices ( \(\left.\mathrm{x} \_0, \ldots, \mathrm{x}_{-}\{\operatorname{dim}-1\}, u\right)\).
Parameters i (int / array_like of int)-MPS index/indices.
Returns lat_idx - First dimensions like \(i\), last dimension has len dim \({ }^{`}+1\) and contains the lattice indices " \(\left(x \_0, \ldots, x_{\_}\{d i m-1\}, u\right)\) ` corresponding to \(i\). For \(i\) accross the MPS unit cell and "infinite" \(b c_{-} M P S\), we shift \(x_{-} 0\) accordingly.

\section*{Return type array}
mps2lat_values_masked ( \(A\), axes=- 1, mps_inds=None, include_u=None)
Reshape/reorder an array \(A\) to replace an MPS index by lattice indices.
This is a generalization of mps2lat_values () allowing for the case of an arbitrary set of MPS indices present in each axis of \(A\).

\section*{Parameters}
- A (ndarray) - Some values.
- axes ( (iterable of) int) - Chooses the axis of \(A\) which should be replaced. If multiple axes are given, you also need to give multiple index arrays as mps_inds.
- mps_inds ((list of) 1D ndarray) - Specifies for each axis in axes, for which MPS indices we have values in the corresponding axis of \(A\). Defaults to [np. arange (A. shape[ax]) for ax in axes]. For indices accross the MPS unit cell and "infinite" \(b c_{-} M P S\), we shift \(x_{-} 0\) accordingly.
- include_u ((list of) bool) - Specifies for each axis in axes, whether the \(u\) index of the lattice should be included into the output array res_A. Defaults to len (self. unit_cell) > 1 .

Returns res_A - Reshaped and reordered copy of A. Such that MPS indices along the specified axes are replaced by lattice indices, i.e., if MPS index \(j\) maps to lattice site ( \(x 0, x 1, x 2\) ), then res_A[..., \(x 0, x 1, x 2, \ldots]=A\left[. . ., m p s \_i n d s[j], \ldots\right]\).

\section*{Return type np.ma.MaskedArray}
mps_idx_fix_u (u=None)
return an index array of MPS indices for which the site within the unit cell is \(u\).
If you have multiple sites in your unit-cell, an onsite operator is in general not defined for all sites. This functions returns an index array of the mps indices which belong to sites given by self. unit_cell[u].

Parameters \(\mathbf{u}\) (None / int) - Selects a site of the unit cell. None (default) means all sites.

Returns mps_idx - MPS indices for which self.site(i) is self.unit_cell[u].
Ordered ascending.
Return type array
mps_lat_idx_fix_u (u=None)
Similar as mps_idx_fix_u(), but return also the corresponding lattice indices.
Parameters u(None / int) - Selects a site of the unit cell. None (default) means all sites.

\section*{Returns}
- mps_idx (array) - MPS indices \(i\) for which self.site(i) is self. unit_cell[u].
- lat_idx (2D array) - The row \(j\) contains the lattice index (without \(u\) ) corresponding to mps_idx[j].
mps_sites()
Return a list of sites for all MPS indices.
Equivalent to [self.site(i) for i in range(self.N_sites)].
This should be used for sites of 1D tensor networks (MPS, MPO,...).
multi_coupling_shape ( \(d x\) )
Calculate correct shape of the strengths for a multi_coupling.
Parameters \(d \mathbf{x}\) (2D array, shape (N_ops, dim)) - dx[i, :] is the translation vector in the lattice for the \(i\)-th operator. Corresponds to the \(d x\) of each operator given in the argument ops of tenpy.models.model.CouplingModel.add_multi_coupling().

\section*{Returns}
- coupling_shape (tuple of int) - Len dim. The correct shape for an array specifying the coupling strength. lat_indices has only rows within this shape.
- shift_lat_indices (array) - Translation vector from origin to the lower left corner of box spanned by \(d x\). (Unlike for coupling_shape () it can also contain entries \(>0\) )
number_nearest_neighbors ( \(u=0\) )
Deprecated.
Deprecated since version 0.5.0: Use count_neighbors () instead.
number_next_nearest_neighbors ( \(u=0\) )
Deprecated.
Deprecated since version 0.5.0: Use count_neighbors () instead.
property order
Defines an ordering of the lattice sites, thus mapping the lattice to a 1 D chain.
Each row of the array contains the lattice indices for one site, the order of the rows thus specifies a path through the lattice, along which an MPS will wind through through the lattice.

You can visualize the order with plot_order ().

\section*{ordering (order)}

Provide possible orderings of the \(N\) lattice sites.
This function can be overwritten by derived lattices to define additional orderings. The following orders are defined in this method:
\begin{tabular}{|l|l|l|}
\hline order & equivalent priority & equivalent snake_winding \\
\hline 'Cstyle' & \((0,1, \ldots, \operatorname{dim}-1, \operatorname{dim})\) & (False, \(\ldots\), False, False) \\
\hline 'default' & & \\
\hline 'snake' & \((0,1, \ldots, \operatorname{dim}-1, \operatorname{dim})\) & (True, \(\ldots\), True, True) \\
\hline 'snakeCstyle' & \\
\hline 'Fstyle' & (dim-1, \(\ldots, 1,0, \operatorname{dim})\) & (False, \(\ldots\), False, False) \\
\hline 'snakeFstyle' & (dim-1, \(, \ldots, 1,0, \operatorname{dim})\) & (False,.., False, False) \\
\hline
\end{tabular}


Note: For lattices with a non-trivial unit cell (e.g. Honeycomb, Kagome), the grouped order might be more appropriate, see get_order_grouped()

Parameters order (str | ('standard', snake_winding, priority) | ('grouped', groups, ...)) - Specifies the desired ordering using one of the strings of the above tables. Alternatively, an ordering is specified by a tuple with first entry specifying a function, 'standard' for get_order() and 'grouped' for get_order_grouped(), and other arguments in the tuple as specified in the documentation of these functions.

Returns order - the order to be used for order.
Return type array, shape (N, D+1), dtype np.intp

\section*{See also:}
get_order generates the order from equivalent priority and snake_winding.
get_order_grouped variant of get_order.
plot_order visualizes the resulting order.
plot_basis (ax, origin=(0.0, 0.0), shade \(=\) None, \(* *\) kwargs \()\)
Plot arrows indicating the basis vectors of the lattice.

\section*{Parameters}
- ax (matplotlib. axes.Axes) - The axes on which we should plot.
- **kwargs - Keyword arguments for ax. arrow.
plot_bc_identified (ax, direction=- 1, origin=None, cylinder_axis=False, **kwargs)
Mark two sites indified by periodic boundary conditions.
Works only for lattice with a 2-dimensional basis.

\section*{Parameters}
- ax (matplotlib. axes.Axes) - The axes on which we should plot.
- direction (int) - The direction of the lattice along which we should mark the idenitified sites. If None, mark it along all directions with periodic boundary conditions.
- cylinder_axis (b○○l) - Whether to plot the cylinder axis as well.
- origin (None / np.ndarray) - The origin starting from where we mark the identified sites. Defaults to the first entry of unit_cell_positions.
- **kwargs - Keyword arguments for the used ax. plot.
plot_coupling (ax, coupling=None, wrap=False, **kwargs)
Plot lines connecting nearest neighbors of the lattice.

\section*{Parameters}
- ax (matplotlib. axes.Axes) - The axes on which we should plot.
- coupling (list of (u1, \(u 2, d x)\) ) - By default (None), use self. pairs['nearest_neighbors']. Specifies the connections to be plotted; iteating over lattice indices ( \(i 0, i 1, \ldots\) ), we plot a connection from the site (i0, i1, ..., u1) to the site ( \(i 0+d x[0]\), \(i 1+d x[1], \ldots, u 2\) ), taking into account the boundary conditions.
- wrap (b○○l) - If True, plot couplings going around the boundary by directly connecting the sites it connects. This might be hard to see, as this puts lines from one end of the lattice to the other. If Fal se, plot the couplings as dangling lines.
- **kwargs - Further keyword arguments given to ax.plot().
plot_order (ax, order=None, textkwargs=\{'color': 'r'\}, **kwargs)
Plot a line connecting sites in the specified "order" and text labels enumerating them.

\section*{Parameters}
- ax (matplotlib. axes.Axes) - The axes on which we should plot.
- order (None / 2D array (self.N_sites, self.dim+1))- The order as returned by ordering(); by default (None) use order.
- textkwargs (None I dict) - If not None, we add text labels enumerating the sites in the plot. The dictionary can contain keyword arguments for ax.text ().
- **kwargs - Further keyword arguments given to ax.plot ().
plot_sites (ax, markers=['o', ' \({ }^{\prime}\) ', 's', ' \({ }^{\prime}\) ', 'h', 'D'], **kwargs)
Plot the sites of the lattice with markers.

\section*{Parameters}
- ax (matplotlib. axes.Axes) - The axes on which we should plot.
- markers (list) - List of values for the keywork marker of ax. plot () to distinguish the different sites in the unit cell, a site \(u\) in the unit cell is plotted with a marker markers[u \% len(markers)].
- **kwargs - Further keyword arguments given to ax. plot ().
position (lat_idx)
return 'space' position of one or multiple sites.
Parameters lat_idx (ndarray, (. . . , dim+1)) - Lattice indices.
Returns pos - The position of the lattice sites specified by lat_idx in real-space.
Return type ndarray, (..., dim)
possible_couplings ( \(u 1, u 2, d x\), strength=None)
Find possible MPS indices for two-site couplings.
For periodic boundary conditions ( \(\mathrm{bc}[\mathrm{a}]==\mathrm{False}\) ) the index x _a is taken modulo Ls [a] and runs through range (Ls [a]). For open boundary conditions, x_a is limited to \(0<=x \_a<\) Ls [a] and \(0<=x \_a+d x[a]<\) lat.Ls[a].

\section*{Parameters}
- u1 (int) - Indices within the unit cell; the \(u 1\) and \(u 2\) of add_coupling ()
- u2 (int) - Indices within the unit cell; the \(u 1\) and \(u 2\) of add_coupling()
- \(\mathbf{d x}(\operatorname{array})\) - Length dim. The translation in terms of basis vectors for the coupling.
- strength (array_like / None) - If given, instead of returning lat_indices and coupling_shape directly return the correct strength_12.

\section*{Returns}
- mps1, mps2 (lD array) - For each possible two-site coupling the MPS indices for the \(u 1\) and \(u 2\).
- strength_vals ( \(1 D\) array) - (Only returend if strength is not None.) Such that for (i, \(j, ~ s) ~ i n ~ z i p\left(m p s 1, ~ m p s 2, ~ s t r e n g t h \_v a l s\right): ~ i t e r a t e s ~ o v e r ~ a l l ~ p o s s i b l e ~ c o u-~\) plings with \(s\) being the strength of that coupling. Couplings where strength_vals \(==0\). are filtered out.
- lat_indices ( \(2 D\) int array) - (Only returend if strength is None.) Rows of lat_indices correspond to entries of \(m p s 1\) and \(m p s 2\) and contain the lattice indices of the "lower left corner" of the box containing the coupling.
- coupling_shape (tuple of int) - (Only returend if strength is None.) Len dim. The correct shape for an array specifying the coupling strength. lat_indices has only rows within this shape.
possible_multi_couplings (ops, strength=None)
Generalization of possible_couplings () to couplings with more than 2 sites.

Parameters ops (list of (opname, \(\mathrm{dx}, \mathrm{u})\) ) - Same as the argument ops of add_multi_coupling().

\section*{Returns}
- mps_ijkl (2D int array) - Each row contains MPS indices \(i, j, k, l, \ldots\). for each of the operators positions. The positions are defined by \(d x(\mathrm{j}, \mathrm{k}, 1, \ldots\) relative to \(i\) ) and boundary coundary conditions of self (how much the box for given \(d x\) can be shifted around without hitting a boundary - these are the different rows).
- strength_vals ( \(1 D\) array) - (Only returend if strength is not None.) Such that for (ijkl, s) in zip(mps_ijkl, strength_vals): iterates over all possible couplings with \(s\) being the strength of that coupling. Couplings where strength_vals \(==0\). are filtered out.
- lat_indices ( \(2 D\) int array) - (Only returend if strength is None.) Rows of lat_indices correspond to rows of \(m p s \_i j k l\) and contain the lattice indices of the "lower left corner" of the box containing the coupling.
- coupling_shape (tuple of int) - (Only returend if strength is None.) Len dim. The correct shape for an array specifying the coupling strength. lat_indices has only rows within this shape.
save_hdf5 (hdf5_saver, h5gr, subpath)
Export self into a HDF5 file.
This method saves all the data it needs to reconstruct self with from_hdf5 ().
Specifically, it saves unit_cell, Ls, unit_cell_positions, basis, boundary_conditions, pairs under their name, bc_MPS as "boundary_conditions_MPS", and order as "order_for_MPS". Moreover, it saves dim and N_sites as HDF5 attributes.

\section*{Parameters}
- hdf5_saver (Hdf5Saver) - Instance of the saving engine.
- h5gr (: class`Group ) - HDF5 group which is supposed to represent self.
- subpath (str) - The name of \(h 5 g r\) with a '/' in the end.
site ( \(i\) )
return Site instance corresponding to an MPS index \(i\)
```

test_sanity()

```

Sanity check.
Raises ValueErrors, if something is wrong.

\subsection*{19.1.9 Square}
- full name: tenpy.models.lattice.Square
- parent module: tenpy.models.lattice
- type: class

\section*{Inheritance Diagram}


\section*{Methods}
\begin{tabular}{|c|c|}
\hline Square.__init__(Lx, Ly, site, **kwargs) & Initialize self. \\
\hline Square.count_neighbors([u, key]) & Count e.g. \\
\hline Square.coupling_shape(dx) & Calculate correct shape of the strengths for a coupling. \\
\hline Square.distance(u1, u2, dx) & Get the distance for a given coupling between two sites in the lattice. \\
\hline Square.enlarge_mps_unit_cell([factor]) & Repeat the unit cell for infinite MPS boundary conditions; in place. \\
\hline Square.find_coupling_pairs([max_dx, cutoff, eps]) & Automatically find coupling pairs grouped by distances. \\
\hline Square.from_hdf5(hdf5_loader, h5gr, subpath) & Load instance from a HDF5 file. \\
\hline Square.lat2mps_idx(lat_idx) & Translate lattice indices (x_0, ..., x_\{D-1\}, u) to MPS index \(i\). \\
\hline Square.mps2lat_idx(i) & Translate MPS index \(i\) to lattice indices ( \(x_{-} 0, \ldots\), x_\{dim-1\}, u). \\
\hline Square.mps2lat_values(A[, axes, u]) & same as Lattice.mps2lat_values(), but ignore \(u\), setting it to 0 . \\
\hline Square.mps2lat_values_masked(A[, axes, ...]) & Reshape/reorder an array \(A\) to replace an MPS index by lattice indices. \\
\hline Square.mps_idx_fix_u([u]) & return an index array of MPS indices for which the site within the unit cell is \(u\). \\
\hline Square.mps_lat_idx_fix_u([u]) & Similar as mps_idx_fix_u(), but return also the corresponding lattice indices. \\
\hline Square.mps_sites() & Return a list of sites for all MPS indices. \\
\hline Square.multi_coupling_shape(dx) & Calculate correct shape of the strengths for a multi_coupling. \\
\hline Square.number_nearest_neighbors([u]) & Deprecated. \\
\hline Square.number_next_nearest_neighbors(lu] & Deprecated. \\
\hline Square.ordering(order) & Provide possible orderings of the \(N\) lattice sites. \\
\hline
\end{tabular}

Table 19 - continued from previous page
\begin{tabular}{|c|c|}
\hline Square.plot_basis(ax[, origin, shade]) & Plot arrows indicating the basis vectors of the lattice. \\
\hline Square.plot_bc_identified(ax[, direction, ...]) & Mark two sites indified by periodic boundary conditions. \\
\hline Square.plot_coupling(ax[, coupling, wrap]) & Plot lines connecting nearest neighbors of the lattice. \\
\hline Square.plot_order(ax[, order, textkwargs]) & Plot a line connecting sites in the specified "order" and text labels enumerating them. \\
\hline Square.plot_sites(ax[ & Plot the sites of the lattice with markers. \\
\hline Square.position(lat_idx) & return 'space' position of one or multiple sites. \\
\hline ```
Square.possible_couplings(u1, u2, dx[,
strength])
``` & Find possible MPS indices for two-site couplings. \\
\hline Square.possible_multi_couplings(ops[, strength]) & Generalization of possible_couplings () to couplings with more than 2 sites. \\
\hline Square.save_hdf5(hdf5_saver, h5gr, subpath) & Export self into a HDF5 file. \\
\hline Square.site(i) & return Site instance corresponding to an MPS index \(i\) \\
\hline Square.test_sanity() & Sanity check. \\
\hline
\end{tabular}

\section*{Class Attributes and Properties}
\begin{tabular}{ll}
\hline Square.Lu & \begin{tabular}{l} 
the (expected) number of sites in the unit cell, \\
len_unit_cell).
\end{tabular} \\
\hline Square.boundary_conditions & \begin{tabular}{l} 
Human-readable list of boundary conditions from bc \\
and bc_shift.
\end{tabular} \\
\hline Square.dim & the dimension of the lattice \\
\hline Square.nearest_neighbors & \\
\hline Square.next_nearest_neighbors & \begin{tabular}{l} 
Defines an ordering of the lattice sites, thus mapping the \\
lattice to a 1D chain.
\end{tabular} \\
\hline Square.next_next_nearest_neighbors &
\end{tabular}
class tenpy.models.lattice. Square (Lx, Ly, site, **kwargs)
Bases: tenpy.models.lattice.SimpleLattice
A square lattice.

\section*{Parameters}
- Lx (int) - The length in each direction.
- Ly (int) - The length in each direction.
- site (Site) - The local lattice site. The unit_cell of the Lattice is just [site].
- **kwargs - Additional keyword arguments given to the Lattice. pairs are set accordingly. If order is specified in the form ('standard', snake_winding, priority), the snake_winding and priority should only be specified for the spatial directions. Similarly, positions can be specified as a single vector.
\(\operatorname{dim}=2\)
the dimension of the lattice


\section*{property boundary_conditions}

Human-readable list of boundary conditions from bc and bc_shift.
Returns boundary_conditions - List of "open" or "periodic", one entry for each direction of the lattice.

Return type list of str
count_neighbors ( \(u=0\), key='nearest_neighbors')
Count e.g. the number of nearest neighbors for a site in the bulk.

\section*{Parameters}
- \(\mathbf{u}\) (int) - Specifies the site in the unit cell, for which we should count the number of neighbors (or whatever key specifies).
- key (str) - Key of pairs to select what to count.

Returns number - Number of nearest neighbors (or whatever key specified) for the \(u\)-th site in the unit cell, somewhere in the bulk of the lattice. Note that it might not be the correct value at the edges of a lattice with open boundary conditions.

\section*{Return type int}

\section*{coupling_shape ( \(d x\) )}

Calculate correct shape of the strengths for a coupling.
Parameters \(\mathbf{d x}\) (tuple of int) - Translation vector in the lattice for a coupling of two operators. Corresponds to \(d x\) argument of tenpy.models.model. CouplingModel. add_multi_coupling().

\section*{Returns}
- coupling_shape (tuple of int) - Len dim. The correct shape for an array specifying the coupling strength. lat_indices has only rows within this shape.
- shift_lat_indices (array) - Translation vector from origin to the lower left corner of box spanned by \(d x\).
distance ( \(u 1, u 2, d x\) )
Get the distance for a given coupling between two sites in the lattice.
The \(u 1, u 2, d x\) parameters are defined in analogy with add_coupling(), i.e., this function calculates the distance between a pair of operators added with add_coupling (using the basis and unit_cell_positions of the lattice).

Warning: This function ignores "wrapping" arround the cylinder in the case of periodic boundary conditions.

\section*{Parameters}
- u1 (int) - Indices within the unit cell; the \(u 1\) and \(u 2\) of add_coupling()
- u2 (int) - Indices within the unit cell; the \(u 1\) and \(u 2\) of add_coupling ()
- \(\mathbf{d x}\) (array) - Length dim. The translation in terms of basis vectors for the coupling.

Returns distance - The distance between site at lattice indices [ \(\mathrm{x}, \mathrm{y}, \mathrm{ul}\) ] and [ \(\mathrm{x}+\) \(d x[0], y+d x[1], u 2]\), ignoring any boundary effects.

\section*{Return type float}
```

enlarge_mps_unit_cell(factor=2)

```

Repeat the unit cell for infinite MPS boundary conditions; in place.
Parameters factor (int) - The new number of sites in the MPS unit cell will be increased from \(N\) _sites to factor \(* N\) _sites_per_ring. Since MPS unit cells are repeated in the \(x\)-direction in our convetion, the lattice shape goes from ( \(\mathrm{Lx}, \mathrm{Ly}, \ldots, \mathrm{Lu}\) ) to (Lx*factor, Ly, ..., Lu).
find_coupling_pairs (max_dx=3, cutoff=None, eps=le-10)
Automatically find coupling pairs grouped by distances.
Given the unit_cell_positions and basis, the coupling pairs of nearest_neighbors, next_nearest_neighbors etc at a given distance are basically fixed (although not uniquely, since we take out half of them to avoid double-counting couplings in both directions A_i B_j + B_i A_i). This function iterates through all possible couplings up to a given cutoff distance and then determines the possible pairs at fixed distances (up to round-off errors).

\section*{Parameters}
- max_dx (int) - Maximal index for each index of \(d x\) to iterate over. You need large enough values to include every possible coupling up to the desired distance, but choosing it too large might make this function run for a long time.
- cutoff (float) - Maximal distance (in the units in which basis and unit_cell_positions is given).
- eps (float) - Tolerance up to which to distances are considered the same.

Returns coupling_pairs - Keys are distances of nearest-neighbors, next-nearest-neighbors etc.
Values are [ (u1, u2, dx), ...] as in pairs.
Return type dict
classmethod from_hdf5 (hdf5_loader, h5gr, subpath)
Load instance from a HDF5 file.
This method reconstructs a class instance from the data saved with save_hdf5().

\section*{Parameters}
- hdf5_loader (Hdf5Loader) - Instance of the loading engine.
- h5gr (Group) - HDF5 group which is represent the object to be constructed.
- subpath (str) - The name of \(h 5 g r\) with a '/' in the end.

Returns obj - Newly generated class instance containing the required data.
Return type cls
lat2mps_idx (lat_idx)
Translate lattice indices ( \(\mathrm{x} \_0, \ldots, \mathrm{x}_{-}\{\mathrm{D}-1\}, \mathrm{u}\) ) to MPS index \(i\).
Parameters lat_idx (array_like [.., dim+1]) - The last dimension corresponds to lattice indices ( \(\left.x \_0, \ldots, x_{-}\{D-1\}, u\right)\). All lattice indices should be positive and smaller than the corresponding entry in self.shape. Exception: for "infinite" \(b c_{-} M P S\), an \(x \_0\) outside indicates shifts accross the boundary.

Returns i-MPS index/indices corresponding to lat_idx. Has the same shape as lat_idx without the last dimension.

Return type array_like
mps2lat_idx (i)
Translate MPS index \(i\) to lattice indices ( \(\left.x \_0, \ldots, x_{-}\{\operatorname{dim}-1\}, u\right)\).

Parameters i (int / array_like of int)-MPS index/indices.
Returns lat_idx - First dimensions like \(i\), last dimension has len dim \({ }^{`}+1\) and contains the lattice indices " \(\left(x_{-} 0, \ldots, x_{\_}\{d i m-1\}, u\right)\) ` corresponding to \(i\). For \(i\) accross the MPS unit cell and "infinite" \(b c_{-} M P S\), we shift \(x \_0\) accordingly.

\section*{Return type array}
mps2lat_values \((A\), axes \(=0, u=\) None)
same as Lattice.mps2lat_values(), but ignore \(u\), setting it to 0 .
mps2lat_values_masked (A, axes=- \(1, m p s \_i n d s=N o n e\), include_u=None)
Reshape/reorder an array \(A\) to replace an MPS index by lattice indices.
This is a generalization of mps2lat_values () allowing for the case of an arbitrary set of MPS indices present in each axis of \(A\).

\section*{Parameters}
- A (ndarray) - Some values.
- axes ( (iterable of) int) - Chooses the axis of \(A\) which should be replaced. If multiple axes are given, you also need to give multiple index arrays as mps_inds.
- mps_inds ((list of) 1D ndarray) - Specifies for each axis in axes, for which MPS indices we have values in the corresponding axis of A. Defaults to [np. arange (A. shape[ax]) for ax in axes]. For indices accross the MPS unit cell and "infinite" \(b c_{-} M P S\), we shift \(x \_0\) accordingly.
- include_u ((list of) bool) - Specifies for each axis in axes, whether the \(u\) index of the lattice should be included into the output array res_A. Defaults to len (self. unit_cell) > 1 .
Returns res_A - Reshaped and reordered copy of A. Such that MPS indices along the specified axes are replaced by lattice indices, i.e., if MPS index \(j\) maps to lattice site ( \(x 0, x 1, x 2\) ), then res_A[..., \(x 0, x 1, x 2, \ldots]=A\left[. . ., m p s \_i n d s[j], \ldots\right]\).

\section*{Return type np.ma.MaskedArray}
mps_idx_fix_u (u=None)
return an index array of MPS indices for which the site within the unit cell is \(u\).
If you have multiple sites in your unit-cell, an onsite operator is in general not defined for all sites. This functions returns an index array of the mps indices which belong to sites given by self. unit_cell[u].

Parameters \(\mathbf{u}\) (None / int) - Selects a site of the unit cell. None (default) means all sites.
Returns mps_idx - MPS indices for which self.site(i) is self.unit_cell[u]. Ordered ascending.

\section*{Return type array}
mps_lat_idx_fix_u ( \(u=\) None)
Similar as mps_idx_fix_u(), but return also the corresponding lattice indices.
Parameters \(\mathbf{u}\) (None / int) - Selects a site of the unit cell. None (default) means all sites.

\section*{Returns}
- mps_idx (array) - MPS indices \(i\) for which self.site(i) is self. unit_cell[u].
- lat_idx (2D array) - The row \(j\) contains the lattice index (without \(u\) ) corresponding to mps_idx[j].
mps_sites()
Return a list of sites for all MPS indices.
Equivalent to [self.site(i) for i in range(self.N_sites)].
This should be used for sites of 1D tensor networks (MPS, MPO,...).
multi_coupling_shape ( \(d x\) )
Calculate correct shape of the strengths for a multi_coupling.
Parameters \(d \mathbf{x}(2 \mathrm{D}\) array, shape (N_ops, dim)) - dx[i, :] is the translation vector in the lattice for the \(i\)-th operator. Corresponds to the \(d x\) of each operator given in the argument ops of tenpy.models.model.CouplingModel.add_multi_coupling().

\section*{Returns}
- coupling_shape (tuple of int) - Len dim. The correct shape for an array specifying the coupling strength. lat_indices has only rows within this shape.
- shift_lat_indices (array) - Translation vector from origin to the lower left corner of box spanned by \(d x\). (Unlike for coupling_shape () it can also contain entries \(>0\) )
number_nearest_neighbors ( \(u=0\) )
Deprecated.
Deprecated since version 0.5.0: Use count_neighbors () instead.
number_next_nearest_neighbors ( \(u=0\) )
Deprecated.
Deprecated since version 0.5.0: Use count_neighbors () instead.
property order
Defines an ordering of the lattice sites, thus mapping the lattice to a 1 D chain.
Each row of the array contains the lattice indices for one site, the order of the rows thus specifies a path through the lattice, along which an MPS will wind through through the lattice.

You can visualize the order with plot_order ().

\section*{ordering (order)}

Provide possible orderings of the \(N\) lattice sites.
This function can be overwritten by derived lattices to define additional orderings. The following orders are defined in this method:
\begin{tabular}{|c|c|c|}
\hline order & equivalent priority & equivalent snake_winding \\
\hline 'Cstyle' & ( \(0,1, \ldots\), dim-1, dim) & (False, ..., False, False) \\
\hline 'default' & & \\
\hline 'snake' & ( \(0,1, \ldots\), dim-1, dim) & (True, ..., True, True) \\
\hline 'snakeCstyle' & & \\
\hline 'Fstyle' & (dim-1, .., 1, 0, dim) & (False, ..., False, False) \\
\hline 'snakeFstyle' & \((\operatorname{dim}-1, \ldots, 1,0, \operatorname{dim})\) & (False, ...., False, False) \\
\hline
\end{tabular}

Note: For lattices with a non-trivial unit cell (e.g. Honeycomb, Kagome), the grouped order might be more appropriate, see get_order_grouped()

Parameters order (str | ('standard', snake_winding, priority) |
('grouped', groups, ...)) - Specifies the desired ordering using one of the

strings of the above tables. Alternatively, an ordering is specified by a tuple with first entry specifying a function, 'standard' for get_order() and 'grouped' for get_order_grouped(), and other arguments in the tuple as specified in the documentation of these functions.

Returns order - the order to be used for order.
Return type array, shape (N, D+1), dtype np.intp

\section*{See also:}
get_order generates the order from equivalent priority and snake_winding.
get_order_grouped variant of get_order.
plot_order visualizes the resulting order.
plot_basis \((a x\), origin \(=(0.0,0.0)\), shade \(=\) None, \(* *\) kwargs \()\)
Plot arrows indicating the basis vectors of the lattice.

\section*{Parameters}
- ax (matplotlib. axes.Axes) - The axes on which we should plot.
- **kwargs - Keyword arguments for ax. arrow.
plot_bc_identified (ax, direction=- 1, origin=None, cylinder_axis=False, **kwargs)
Mark two sites indified by periodic boundary conditions.
Works only for lattice with a 2-dimensional basis.

\section*{Parameters}
- ax (matplotlib. axes.Axes) - The axes on which we should plot.
- direction (int) - The direction of the lattice along which we should mark the idenitified sites. If None, mark it along all directions with periodic boundary conditions.
- cylinder_axis (bool) - Whether to plot the cylinder axis as well.
- origin (None / np.ndarray) - The origin starting from where we mark the identified sites. Defaults to the first entry of unit_cell_positions.
- **kwargs - Keyword arguments for the used ax.plot.
plot_coupling (ax, coupling=None, wrap=False, **kwargs)
Plot lines connecting nearest neighbors of the lattice.

\section*{Parameters}
- ax (matplotlib.axes.Axes) - The axes on which we should plot.
- coupling (list of (u1, u2, dx)) - By default (None), use self. pairs['nearest_neighbors']. Specifies the connections to be plotted; iteating over lattice indices ( \(i 0, i 1, \ldots\) ), we plot a connection from the site (i0, i1, ..., u1) to the site ( \(i 0+d x[0]\), \(i 1+d x[1], \ldots, u 2\) ), taking into account the boundary conditions.
- wrap (bool) - If True, plot couplings going around the boundary by directly connecting the sites it connects. This might be hard to see, as this puts lines from one end of the lattice to the other. If False, plot the couplings as dangling lines.
- **kwargs - Further keyword arguments given to ax. plot ().
```

plot_order (ax, order=None, textkwargs=\{'color': 'r'\}, **kwargs)

```

Plot a line connecting sites in the specified "order" and text labels enumerating them.

\section*{Parameters}
- ax (matplotlib. axes.Axes) - The axes on which we should plot.
- order (None / 2D array (self.N_sites, self.dim+1))- The order as returned by ordering(); by default (None) use order.
- textkwargs (None I dict) - If not None, we add text labels enumerating the sites in the plot. The dictionary can contain keyword arguments for ax.text ().
- **kwargs - Further keyword arguments given to ax.plot ().
plot_sites (ax, markers=['o', '^', 's', 'p', 'h', 'D'], **kwargs)
Plot the sites of the lattice with markers.

\section*{Parameters}
- ax (matplotlib. axes.Axes) - The axes on which we should plot.
- markers (list) - List of values for the keywork marker of ax.plot () to distinguish the different sites in the unit cell, a site \(u\) in the unit cell is plotted with a marker markers[u \% len(markers)].
- **kwargs - Further keyword arguments given to ax. plot ().
position(lat_idx)
return 'space' position of one or multiple sites.
Parameters lat_idx (ndarray, (. . , dim+1)) - Lattice indices.
Returns pos - The position of the lattice sites specified by lat_idx in real-space.
Return type ndarray, (..., dim)
possible_couplings ( \(u 1, u 2, d x\), strength=None)
Find possible MPS indices for two-site couplings.
For periodic boundary conditions ( \(\mathrm{bc}[\mathrm{a}]==\mathrm{False}\) ) the index x _a is taken modulo Ls [a] and runs through range (Ls [a]). For open boundary conditions, x_a is limited to \(0<=x \_a<\) Ls [a] and \(0<=x \_a+d x[a]<\) lat.Ls [a].

\section*{Parameters}
- u1 (int) - Indices within the unit cell; the \(u 1\) and \(u 2\) of add_coupling()
- u2 (int) - Indices within the unit cell; the \(u 1\) and \(u 2\) of add_coupling()
- \(\mathbf{d x}\) (array) - Length dim. The translation in terms of basis vectors for the coupling.
- strength (array_like / None) - If given, instead of returning lat_indices and coupling_shape directly return the correct strength_12.

\section*{Returns}
- mps1, mps2 (1D array) - For each possible two-site coupling the MPS indices for the \(u 1\) and \(u 2\).
- strength_vals (1D array) - (Only returend if strength is not None.) Such that for (i, \(j, ~ s) ~ i n ~ z i p\left(m p s 1, ~ m p s 2, ~ s t r e n g t h \_v a l s\right): ~ i t e r a t e s ~ o v e r ~ a l l ~ p o s s i b l e ~ c o u-~\) plings with \(s\) being the strength of that coupling. Couplings where strength_vals \(=0\). are filtered out.
- lat_indices ( \(2 D\) int array) - (Only returend if strength is None.) Rows of lat_indices correspond to entries of \(m p s l\) and \(m p s 2\) and contain the lattice indices of the "lower left corner" of the box containing the coupling.
- coupling_shape (tuple of int) - (Only returend if strength is None.) Len dim. The correct shape for an array specifying the coupling strength. lat_indices has only rows within this shape.

\section*{possible_multi_couplings (ops, strength=None)}

Generalization of possible_couplings () to couplings with more than 2 sites.
Parameters ops (list of (opname, \(\mathrm{dx}, \mathrm{u})\) ) - Same as the argument ops of add_multi_coupling().

\section*{Returns}
- mps_ijkl (2D int array) - Each row contains MPS indices \(i, j, k, l, \ldots\). for each of the operators positions. The positions are defined by \(d x\) ( \(\mathrm{j}, \mathrm{k}, \mathrm{l}, \ldots\) relative to \(i\) ) and boundary coundary conditions of self (how much the box for given \(d x\) can be shifted around without hitting a boundary - these are the different rows).
- strength_vals ( \(1 D\) array) - (Only returend if strength is not None.) Such that for (ijkl, s) in zip(mps_ijkl, strength_vals): iterates over all possible couplings with \(s\) being the strength of that coupling. Couplings where strength_vals \(==0\). are filtered out.
- lat_indices ( 2 D int array) - (Only returend if strength is None.) Rows of lat_indices correspond to rows of \(m p s \_i j k l\) and contain the lattice indices of the "lower left corner" of the box containing the coupling.
- coupling_shape (tuple of int) - (Only returend if strength is None.) Len dim. The correct shape for an array specifying the coupling strength. lat_indices has only rows within this shape.
save_hdf5 (hdf5_saver, h5gr, subpath)
Export self into a HDF5 file.
This method saves all the data it needs to reconstruct self with from_hdf5 ().
Specifically, it saves unit_cell, Ls, unit_cell_positions, basis, boundary_conditions, pairs under their name, bc_MPS as "boundary_conditions_MPS", and order as "order_for_MPS". Moreover, it saves dim and N_sites as HDF5 attributes.

\section*{Parameters}
- hdf5_saver (Hdf5Saver) - Instance of the saving engine.
- h5gr (: class `Group \()\) - HDF5 group which is supposed to represent self.
- subpath (str) - The name of \(h 5 g r\) with a ' /' in the end.
```

site(i)

```
return Site instance corresponding to an MPS index \(i\)
```

test_sanity()

```

Sanity check.
Raises ValueErrors, if something is wrong.

\subsection*{19.1.10 Triangular}
- full name: tenpy.models.lattice.Triangular
- parent module: tenpy.models.lattice
- type: class

\section*{Inheritance Diagram}


\section*{Methods}
\begin{tabular}{|c|c|}
\hline Triangular.__init__(Lx, Ly, site, **kwargs) & Initialize self. \\
\hline Triangular.count_neighbors([u, key]) & Count e.g. \\
\hline Triangular.coupling_shape(dx) & Calculate correct shape of the strengths for a coupling. \\
\hline Triangular.distance(u1, u2, dx) & Get the distance for a given coupling between two sites in the lattice. \\
\hline Triangular.enlarge_mps_unit_cell([factor]) & Repeat the unit cell for infinite MPS boundary conditions; in place. \\
\hline ```
Triangular.find_coupling_pairs([max_dx,
...])
``` & Automatically find coupling pairs grouped by distances. \\
\hline Triangular.from_hdf5(hdf5_loader, h5gr, subpath) & Load instance from a HDF5 file. \\
\hline Triangular.lat2mps_idx(lat_idx) & Translate lattice indices \(\left(x \_0, \ldots, x_{-}\{D-1\}\right.\), u) to MPS index \(i\). \\
\hline Triangular.mps2lat_idx(i) & Translate MPS index \(i\) to lattice indices (x_0, ..., \(\left.x_{-}\{d i m-1\}, u\right)\). \\
\hline Triangular.mps2lat_values(A[, axes, u]) & same as Lattice.mps2lat_values(), but ignore \(u\), setting it to 0 . \\
\hline Triangular.mps2lat_values_masked(A[, axes,...]) & Reshape/reorder an array \(A\) to replace an MPS index by lattice indices. \\
\hline Triangular.mps_idx_fix_u([u]) & return an index array of MPS indices for which the site within the unit cell is \(u\). \\
\hline
\end{tabular}

Table 21 - continued from previous page
\begin{tabular}{|c|c|}
\hline Triangular.mps_lat_idx_fix_u([u]) & Similar as mps_idx_fix_u(), but return also the corresponding lattice indices. \\
\hline Triangular.mps_sites() & Return a list of sites for all MPS indices. \\
\hline Triangular.multi_coupling_shape(dx) & Calculate correct shape of the strengths for a multi_coupling. \\
\hline Triangular.number_nearest_neighbors([u]) & Deprecated. \\
\hline Triangular.number_next_near &  \\
\hline Triangular.ordering(order) & Provide possible orderings of the \(N\) lattice sites. \\
\hline Triangular.plot_basis(ax[, origin, shade]) & Plot arrows indicating the basis vectors of the lattice. \\
\hline Triangular.plot_bc_identified(ax[,..]) & Mark two sites indified by periodic boundary conditions. \\
\hline Triangular.plot_coupling(ax[, coupling, wrap]) & Plot lines connecting nearest neighbors of the lattice. \\
\hline Triangular.plot_order(ax[, order, textkwargs]) & Plot a line connecting sites in the specified "order" and text labels enumerating them. \\
\hline Triangular.plot_sites(ax[, markers]) & Plot the sites of the lattice with markers. \\
\hline Triangular.position(lat_idx) & return 'space' position of one or multiple sites. \\
\hline Triangular.possible_couplings(u1, u2, dx[, ...]) & Find possible MPS indices for two-site couplings. \\
\hline Triangular.possible_multi_couplings(ops
...]) & Generalization of possible_couplings () to couplings with more than 2 sites. \\
\hline Triangular.save_hdf5(hdf5_saver, h5gr, subpath) & Export self into a HDF5 file. \\
\hline Triangular.site(i) & return Site instance corresponding to an MPS index \(i\) \\
\hline Triangular.test_sanity() & Sanity check. \\
\hline
\end{tabular}

\section*{Class Attributes and Properties}
\begin{tabular}{ll}
\hline Triangular.Lu & \begin{tabular}{l} 
the (expected) number of sites in the unit cell, \\
len (unit_cell).
\end{tabular} \\
\hline Triangular.boundary_conditions & \begin{tabular}{l} 
Human-readable list of boundary conditions from bc \\
and bc_shift.
\end{tabular} \\
\hline Triangular.dim & the dimension of the lattice \\
\hline Triangular.nearest_neighbors & \\
\hline Triangular.next_nearest_neighbors & \begin{tabular}{l} 
Defines an ordering of the lattice sites, thus mapping the \\
Triangular.next_next_nearest_neighbors \\
Triangular.order
\end{tabular} \\
\begin{tabular}{ll} 
class tenpy.models.lattice. Triangular (Lx, Ly, site, **kwargs) \\
Bases: tenpy.models.Iattice.SimpleLattice \\
A triangularlattice.
\end{tabular}
\end{tabular}

\section*{Parameters}
- \(\mathbf{L x}(i n t)\) - The length in each direction.
- Ly (int) - The length in each direction.

- site (Site) - The local lattice site. The unit_cell of the Lattice is just [site].
- **kwargs - Additional keyword arguments given to the Lattice. pairs are set accordingly. If order is specified in the form ('standard', snake_windingi, priority), the snake_winding and priority should only be specified for the spatial directions. Similarly, positions can be specified as a single vector.
```

dim = 2

```
the dimension of the lattice

\section*{property boundary_conditions}

Human-readable list of boundary conditions from bc and bc_shift.
Returns boundary_conditions - List of "open" or "periodic", one entry for each direction of the lattice.
Return type list of str
count_neighbors ( \(u=0\), key='nearest_neighbors')
Count e.g. the number of nearest neighbors for a site in the bulk.

\section*{Parameters}
- \(\mathbf{u}\) (int) - Specifies the site in the unit cell, for which we should count the number of neighbors (or whatever key specifies).
- key (str) - Key of pairs to select what to count.

Returns number - Number of nearest neighbors (or whatever key specified) for the \(u\)-th site in the unit cell, somewhere in the bulk of the lattice. Note that it might not be the correct value at the edges of a lattice with open boundary conditions.

\section*{Return type int}

\section*{coupling_shape ( \(d x\) )}

Calculate correct shape of the strengths for a coupling.
Parameters \(\mathbf{d x}\) (tuple of int) - Translation vector in the lattice for a coupling of two operators. Corresponds to \(d x\) argument of tenpy.models.model. CouplingModel. add_multi_coupling().

\section*{Returns}
- coupling_shape (tuple of int) - Len dim. The correct shape for an array specifying the coupling strength. lat_indices has only rows within this shape.
- shift_lat_indices (array) - Translation vector from origin to the lower left corner of box spanned by \(d x\).
distance ( \(u 1, u 2, d x\) )
Get the distance for a given coupling between two sites in the lattice.
The \(u 1, u 2, d x\) parameters are defined in analogy with add_coupling(), i.e., this function calculates the distance between a pair of operators added with add_coupling (using the basis and unit_cell_positions of the lattice).

Warning: This function ignores "wrapping" arround the cylinder in the case of periodic boundary conditions.

\section*{Parameters}
- u1 (int) - Indices within the unit cell; the \(u 1\) and \(u 2\) of add_coupling ()
- u2 (int) - Indices within the unit cell; the \(u 1\) and \(u 2\) of add_coupling()
- \(\mathbf{d x}\) (array) - Length dim. The translation in terms of basis vectors for the coupling.

Returns distance - The distance between site at lattice indices [x, y, ul] and [x + \(d x[0], y+d x[1], u 2]\), ignoring any boundary effects.

Return type float

\section*{enlarge_mps_unit_cell (factor=2)}

Repeat the unit cell for infinite MPS boundary conditions; in place.
Parameters factor (int) - The new number of sites in the MPS unit cell will be increased from \(N\) _sites to factor*N_sites_per_ring. Since MPS unit cells are repeated in the \(x\)-direction in our convetion, the lattice shape goes from ( \(L x, L y, \ldots, L u\) ) to (Lx*factor, Ly, ..., Lu).
find_coupling_pairs (max_dx=3, cutoff=None, eps=le-10)
Automatically find coupling pairs grouped by distances.
Given the unit_cell_positions and basis, the coupling pairs of nearest_neighbors, next_nearest_neighbors etc at a given distance are basically fixed (although not uniquely, since we take out half of them to avoid double-counting couplings in both directions A_i B_j + B_i A_i). This function iterates through all possible couplings up to a given cutoff distance and then determines the possible pairs at fixed distances (up to round-off errors).

\section*{Parameters}
- max_dx (int) - Maximal index for each index of \(d x\) to iterate over. You need large enough values to include every possible coupling up to the desired distance, but choosing it too large might make this function run for a long time.
- cutoff (float) - Maximal distance (in the units in which basis and unit_cell_positions is given).
- eps (float) - Tolerance up to which to distances are considered the same.

Returns coupling_pairs - Keys are distances of nearest-neighbors, next-nearest-neighbors etc.
Values are [(u1, u2, dx), ...] as in pairs.
Return type dict
classmethod from_hdf5 (hdf5_loader, h5gr, subpath)
Load instance from a HDF5 file.
This method reconstructs a class instance from the data saved with save_hdf5 ().

\section*{Parameters}
- hdf5_loader (Hdf5Loader) - Instance of the loading engine.
- h5gr (Group) - HDF5 group which is represent the object to be constructed.
- subpath (str) - The name of \(h 5 g r\) with a ' /' in the end.

Returns obj - Newly generated class instance containing the required data.
Return type cls
lat2mps_idx (lat_idx)
Translate lattice indices ( \(\mathrm{x} \_0, \ldots, \mathrm{x} \_\{\mathrm{D}-1\}, \mathrm{u}\) ) to MPS index \(i\).
Parameters lat_idx (array_like [.., dim+1]) - The last dimension corresponds to lattice indices \(\left(x_{-} 0, \ldots, x_{-}\{D-1\}, u\right)\). All lattice indices should be positive and
smaller than the corresponding entry in self.shape. Exception: for "infinite" \(b c_{-} M P S\), an \(x \_0\) outside indicates shifts accross the boundary.
Returns i - MPS index/indices corresponding to lat_idx. Has the same shape as lat_idx without the last dimension.

Return type array_like
mps2lat_idx (i)
Translate MPS index \(i\) to lattice indices ( \(\left.\mathrm{x} \_0, \ldots, \mathrm{x}_{-}\{\operatorname{dim}-1\}, u\right)\).
Parameters i (int | array_like of int)-MPS index/indices.
Returns lat_idx - First dimensions like \(i\), last dimension has len dim \({ }^{`}+1\) and contains the lattice indices \({ }^{`}\left(x_{-} 0, \ldots, x_{\_}\{d i m-1\}, u\right)\) ' corresponding to \(i\). For \(i\) accross the MPS unit cell and "infinite" \(b c_{-} M P S\), we shift \(x_{-} 0\) accordingly.

\section*{Return type array}
mps2lat_values \((A\), axes \(=0, u=\) None \()\)
same as Lattice.mps2lat_values(), but ignore \(u\), setting it to 0 .
mps2lat_values_masked ( \(A\), axes=- \(1, m p s \_i n d s=N o n e\), include_u=None)
Reshape/reorder an array \(A\) to replace an MPS index by lattice indices.
This is a generalization of mps2lat_values () allowing for the case of an arbitrary set of MPS indices present in each axis of \(A\).

\section*{Parameters}
- A (ndarray) - Some values.
- axes ( (iterable of) int) - Chooses the axis of \(A\) which should be replaced. If multiple axes are given, you also need to give multiple index arrays as mps_inds.
- mps_inds ((list of) 1D ndarray) - Specifies for each axis in axes, for which MPS indices we have values in the corresponding axis of \(A\). Defaults to [np. arange (A. shape[ax]) for ax in axes]. For indices accross the MPS unit cell and "infinite" \(b c_{-} M P S\), we shift \(x_{-} 0\) accordingly.
- include_u ((list of) bool) - Specifies for each axis in axes, whether the \(u\) index of the lattice should be included into the output array res_A. Defaults to len (self. unit_cell) > 1.
Returns res_A - Reshaped and reordered copy of A. Such that MPS indices along the specified axes are replaced by lattice indices, i.e., if MPS index \(j\) maps to lattice site \((x 0, x 1, x 2)\), then res_A[..., x0, x1, x2, ...] =A[..., mps_inds[j], ...].
Return type np.ma.MaskedArray
mps_idx_fix_u (u=None)
return an index array of MPS indices for which the site within the unit cell is \(u\).
If you have multiple sites in your unit-cell, an onsite operator is in general not defined for all sites. This functions returns an index array of the mps indices which belong to sites given by self. unit_cell[u].

Parameters \(\mathbf{u}\) (None / int) - Selects a site of the unit cell. None (default) means all sites.
Returns mps_idx - MPS indices for which self.site(i) is self.unit_cell[u]. Ordered ascending.
Return type array
mps_lat_idx_fix_u (u=None)
Similar as mps_idx_fix_u(), but return also the corresponding lattice indices.
Parameters \(\mathbf{u}\) (None / int) - Selects a site of the unit cell. None (default) means all sites.

\section*{Returns}
- mps_idx (array) - MPS indices \(i\) for which self.site(i) is self. unit_cell[u].
- lat_idx ( \(2 D\) array) - The row \(j\) contains the lattice index (without \(u\) ) corresponding to mps_idx[j].
mps_sites()
Return a list of sites for all MPS indices.
Equivalent to [self.site(i) for i in range(self.N_sites)].
This should be used for sites of 1D tensor networks (MPS, MPO,...).
multi_coupling_shape ( \(d x\) )
Calculate correct shape of the strengths for a multi_coupling.
Parameters \(d \mathbf{x}(2 \mathrm{D}\) array, shape (N_ops, dim)) - dx[i, :] is the translation vector in the lattice for the \(i\)-th operator. Corresponds to the \(d x\) of each operator given in the argument ops of tenpy.models.model.CouplingModel.add_multi_coupling().

\section*{Returns}
- coupling_shape (tuple of int) - Len dim. The correct shape for an array specifying the coupling strength. lat_indices has only rows within this shape.
- shift_lat_indices (array) - Translation vector from origin to the lower left corner of box spanned by \(d x\). (Unlike for coupling_shape () it can also contain entries \(>0\) )
number_nearest_neighbors ( \(u=0\) )
Deprecated.
Deprecated since version 0.5.0: Use count_neighbors () instead.
number_next_nearest_neighbors ( \(u=0\) )
Deprecated.
Deprecated since version 0.5.0: Use count_neighbors () instead.

\section*{property order}

Defines an ordering of the lattice sites, thus mapping the lattice to a 1 D chain.
Each row of the array contains the lattice indices for one site, the order of the rows thus specifies a path through the lattice, along which an MPS will wind through through the lattice.
You can visualize the order with plot_order ().
ordering (order)
Provide possible orderings of the \(N\) lattice sites.
This function can be overwritten by derived lattices to define additional orderings. The following orders are defined in this method:
\begin{tabular}{|l|l|l|}
\hline order & equivalent priority & equivalent snake_winding \\
\hline 'Cstyle' & \((0,1, \ldots, \operatorname{dim}-1, \operatorname{dim})\) & (False, \(\ldots\), False, False) \\
\hline 'default' & & \\
\hline 'snake' & \((0,1, \ldots, \operatorname{dim}-1, \operatorname{dim})\) & (True, \(\ldots\), True, True) \\
\hline 'snakeCstyle' & \\
\hline 'Fstyle' & (dim-1, \(\ldots, 1,0, \operatorname{dim})\) & (False, \(\ldots\), False, False) \\
\hline 'snakeFstyle' & (dim-1, \(, \ldots, 1,0, \operatorname{dim})\) & (False,.., False, False) \\
\hline
\end{tabular}


Note: For lattices with a non-trivial unit cell (e.g. Honeycomb, Kagome), the grouped order might be more appropriate, see get_order_grouped()

Parameters order (str | ('standard', snake_winding, priority) | ('grouped', groups, ...)) - Specifies the desired ordering using one of the strings of the above tables. Alternatively, an ordering is specified by a tuple with first entry specifying a function, 'standard' for get_order() and 'grouped' for get_order_grouped(), and other arguments in the tuple as specified in the documentation of these functions.

Returns order - the order to be used for order.
Return type array, shape (N, D+1), dtype np.intp

\section*{See also:}
get_order generates the order from equivalent priority and snake_winding.
get_order_grouped variant of get_order.
plot_order visualizes the resulting order.
plot_basis (ax, origin=(0.0, 0.0), shade \(=\) None, \({ }^{* *}\) kwargs )
Plot arrows indicating the basis vectors of the lattice.

\section*{Parameters}
- ax (matplotlib. axes.Axes) - The axes on which we should plot.
- **kwargs - Keyword arguments for ax. arrow.
plot_bc_identified (ax, direction=- 1, origin=None, cylinder_axis=False, **kwargs)
Mark two sites indified by periodic boundary conditions.
Works only for lattice with a 2-dimensional basis.

\section*{Parameters}
- ax (matplotlib. axes.Axes) - The axes on which we should plot.
- direction (int) - The direction of the lattice along which we should mark the idenitified sites. If None, mark it along all directions with periodic boundary conditions.
- cylinder_axis (b○○l) - Whether to plot the cylinder axis as well.
- origin (None / np.ndarray) - The origin starting from where we mark the identified sites. Defaults to the first entry of unit_cell_positions.
- **kwargs - Keyword arguments for the used ax. plot.
plot_coupling (ax, coupling=None, wrap=False, **kwargs)
Plot lines connecting nearest neighbors of the lattice.

\section*{Parameters}
- ax (matplotlib. axes.Axes) - The axes on which we should plot.
- coupling (list of (u1, \(u 2, d x)\) ) - By default (None), use self. pairs['nearest_neighbors']. Specifies the connections to be plotted; iteating over lattice indices ( \(i 0, i 1, \ldots\) ), we plot a connection from the site (i0, i1, ..., u1) to the site ( \(i 0+d x[0]\), \(i 1+d x[1], \ldots, u 2\) ), taking into account the boundary conditions.
- wrap (b○○l) - If True, plot couplings going around the boundary by directly connecting the sites it connects. This might be hard to see, as this puts lines from one end of the lattice to the other. If Fal se, plot the couplings as dangling lines.
- **kwargs - Further keyword arguments given to ax.plot().
plot_order (ax, order=None, textkwargs=\{'color': 'r'\}, **kwargs)
Plot a line connecting sites in the specified "order" and text labels enumerating them.

\section*{Parameters}
- ax (matplotlib. axes.Axes) - The axes on which we should plot.
- order (None / 2D array (self.N_sites, self.dim+1))- The order as returned by ordering(); by default (None) use order.
- textkwargs (None I dict) - If not None, we add text labels enumerating the sites in the plot. The dictionary can contain keyword arguments for ax.text ().
- **kwargs - Further keyword arguments given to ax.plot ().
plot_sites (ax, markers=['o', ' \({ }^{\prime}\) ', 's', ' \({ }^{\prime}\) ', 'h', 'D'], **kwargs)
Plot the sites of the lattice with markers.

\section*{Parameters}
- ax (matplotlib. axes.Axes) - The axes on which we should plot.
- markers (list) - List of values for the keywork marker of ax. plot () to distinguish the different sites in the unit cell, a site \(u\) in the unit cell is plotted with a marker markers[u \% len(markers)].
- **kwargs - Further keyword arguments given to ax. plot ().
position (lat_idx)
return 'space' position of one or multiple sites.
Parameters lat_idx (ndarray, (. . . , dim+1)) - Lattice indices.
Returns pos - The position of the lattice sites specified by lat_idx in real-space.
Return type ndarray, (..., dim)
possible_couplings ( \(u 1, u 2, d x\), strength=None)
Find possible MPS indices for two-site couplings.
For periodic boundary conditions ( \(\mathrm{bc}[\mathrm{a}]==\mathrm{False}\) ) the index x _a is taken modulo Ls [a] and runs through range (Ls [a]). For open boundary conditions, x_a is limited to \(0<=x \_a<\) Ls [a] and \(0<=x \_a+d x[a]<\) lat.Ls[a].

\section*{Parameters}
- u1 (int) - Indices within the unit cell; the \(u 1\) and \(u 2\) of add_coupling ()
- u2 (int) - Indices within the unit cell; the \(u 1\) and \(u 2\) of add_coupling()
- \(\mathbf{d x}(\operatorname{array})\) - Length dim. The translation in terms of basis vectors for the coupling.
- strength (array_like / None) - If given, instead of returning lat_indices and coupling_shape directly return the correct strength_12.

\section*{Returns}
- mps1, mps2 (lD array) - For each possible two-site coupling the MPS indices for the \(u 1\) and \(u 2\).
- strength_vals (1D array) - (Only returend if strength is not None.) Such that for (i, \(j, ~ s) ~ i n ~ z i p\left(m p s 1, ~ m p s 2, ~ s t r e n g t h \_v a l s\right): ~ i t e r a t e s ~ o v e r ~ a l l ~ p o s s i b l e ~ c o u-~\) plings with \(s\) being the strength of that coupling. Couplings where strength_vals \(==0\). are filtered out.
- lat_indices ( \(2 D\) int array) - (Only returend if strength is None.) Rows of lat_indices correspond to entries of \(m p s 1\) and \(m p s 2\) and contain the lattice indices of the "lower left corner" of the box containing the coupling.
- coupling_shape (tuple of int) - (Only returend if strength is None.) Len dim. The correct shape for an array specifying the coupling strength. lat_indices has only rows within this shape.
possible_multi_couplings (ops, strength=None)
Generalization of possible_couplings () to couplings with more than 2 sites.

Parameters ops (list of (opname, \(\mathrm{dx}, \mathrm{u})\) ) - Same as the argument ops of add_multi_coupling().

\section*{Returns}
- mps_ijkl (2D int array) - Each row contains MPS indices \(i, j, k, l, \ldots\). for each of the operators positions. The positions are defined by \(d x(\mathrm{j}, \mathrm{k}, 1, \ldots\) relative to \(i\) ) and boundary coundary conditions of self (how much the box for given \(d x\) can be shifted around without hitting a boundary - these are the different rows).
- strength_vals ( \(1 D\) array) - (Only returend if strength is not None.) Such that for (ijkl, s) in zip(mps_ijkl, strength_vals): iterates over all possible couplings with \(s\) being the strength of that coupling. Couplings where strength_vals \(==0\). are filtered out.
- lat_indices ( \(2 D\) int array) - (Only returend if strength is None.) Rows of lat_indices correspond to rows of \(m p s \_i j k l\) and contain the lattice indices of the "lower left corner" of the box containing the coupling.
- coupling_shape (tuple of int) - (Only returend if strength is None.) Len dim. The correct shape for an array specifying the coupling strength. lat_indices has only rows within this shape.
```

save_hdf5 (hdf5_saver, h5gr, subpath)

```

Export self into a HDF5 file.
This method saves all the data it needs to reconstruct self with from_hdf5 ().
Specifically, it saves unit_cell, Ls, unit_cell_positions, basis, boundary_conditions, pairs under their name, bc_MPS as "boundary_conditions_MPS", and order as "order_for_MPS". Moreover, it saves dim and N_sites as HDF5 attributes.

\section*{Parameters}
- hdf5_saver (Hdf5Saver) - Instance of the saving engine.
- h5gr (: class`Group ) - HDF5 group which is supposed to represent self.
- subpath (str) - The name of \(h 5 g r\) with a '/' in the end.
```

site(i)

```
return Site instance corresponding to an MPS index \(i\)
```

test_sanity()

```

Sanity check.
Raises ValueErrors, if something is wrong.

\subsection*{19.1.11 TrivialLattice}
- full name: tenpy.models.lattice.TrivialLattice
- parent module: tenpy.models.lattice
- type: class

\section*{Inheritance Diagram}


\section*{Methods}
\begin{tabular}{ll}
\hline \begin{tabular}{l} 
TrivialLattice.__init__(mps_sites, \\
**kargs)
\end{tabular} & Initialize self. \\
\hline Triviallattice.count_neighbors([u, key]) & Count e.g. \\
\hline Triviallattice.coupling_shape(dx) & Calculate correct shape of the strengths for a coupling. \\
\hline Triviallattice.distance(ul, u2, dx) & \begin{tabular}{l} 
Get the distance for a given coupling between two sites \\
in the lattice.
\end{tabular} \\
\hline Triviallattice.enlarge_mps_unit_cell([facRorpeat the unit cell for infinite MPS boundary condi- \\
tions; in place.
\end{tabular}

Table 23 - continued from previous page
\begin{tabular}{|c|c|}
\hline ```
TrivialLattice.plot_bc_identified(ax[,
...])
``` & Mark two sites indified by periodic boundary conditions. \\
\hline Triviallattice.plot_coupling(ax[, coupling, ...]) & Plot lines connecting nearest neighbors of the lattice. \\
\hline TrivialLattice.plot_order(ax[,order,...]) & Plot a line connecting sites in the specified "order" and text labels enumerating them. \\
\hline TrivialLattice.plot_sites(ax[, markers]) & Plot the sites of the lattice with markers. \\
\hline TrivialLattice.position(lat_idx) & return 'space' position of one or multiple sites. \\
\hline ```
TrivialLattice.possible_couplings(u1,
u2, dx)
``` & Find possible MPS indices for two-site couplings. \\
\hline Triviallattice.possible_multi_coupling & (Gpaseralization of possible_couplings () to couplings with more than 2 sites. \\
\hline Triviallattice.save_hdf5(hdf5_saver, h5gr, ...) & Export self into a HDF5 file. \\
\hline TrivialLattice.site(i) & return Site instance corresponding to an MPS index \(i\) \\
\hline TrivialLattice.test_sanity() & Sanity check. \\
\hline
\end{tabular}

\section*{Class Attributes and Properties}
\begin{tabular}{|c|c|}
\hline TrivialLattice.Lu & the (expected) number of sites in the unit cell, len (unit_cell). \\
\hline TrivialLattice.boundary_conditions & Human-readable list of boundary conditions from bc and bc_shift. \\
\hline TrivialLattice.dim & The dimension of the lattice. \\
\hline \multicolumn{2}{|l|}{TrivialLattice.nearest_neighbors} \\
\hline \multicolumn{2}{|l|}{TrivialLattice.next_nearest_neighbors} \\
\hline \multicolumn{2}{|l|}{TrivialLattice.next_next_nearest_neighbors} \\
\hline TrivialLattice.order & Defines an ordering of the lattice sites, thus mapping the lattice to a 1D chain. \\
\hline \multicolumn{2}{|l|}{class tenpy.models.lattice.TrivialLattice (mps_sites, **kwargs)} \\
\hline \multicolumn{2}{|l|}{Bases: tenpy.models.lattice.Lattice} \\
\hline \multicolumn{2}{|l|}{Trivial lattice consisting of a single (possibly large) unit cell in 1D.} \\
\hline \multicolumn{2}{|l|}{This is usefull if you need a valid Lattice with given mps_sites () and don't care about the actual geometry, e.g, because you don't intend to use the CouplingModel.} \\
\hline
\end{tabular}

\section*{Parameters}
- mps_sites (list of Site) - The sites making up a unit cell of the lattice.
- **kwargs - Further keyword arguments given to Lattice.

\section*{property boundary_conditions}

Human-readable list of boundary conditions from bc and bc_shift.
Returns boundary_conditions - List of "open" or "periodic", one entry for each direction of the lattice.
Return type list of str
```

count_neighbors ( }u=0,\mathrm{ key='nearest_neighbors')

```

Count e.g. the number of nearest neighbors for a site in the bulk.

\section*{Parameters}
- u (int) - Specifies the site in the unit cell, for which we should count the number of neighbors (or whatever key specifies).
- key (str) - Key of pairs to select what to count.

Returns number - Number of nearest neighbors (or whatever key specified) for the \(u\)-th site in the unit cell, somewhere in the bulk of the lattice. Note that it might not be the correct value at the edges of a lattice with open boundary conditions.

\section*{Return type int}

\section*{coupling_shape ( \(d x\) )}

Calculate correct shape of the strengths for a coupling.
Parameters \(\mathbf{d x}\) (tuple of int) - Translation vector in the lattice for a coupling of two operators. Corresponds to \(d x\) argument of tenpy.models.model. CouplingModel. add_multi_coupling().

\section*{Returns}
- coupling_shape (tuple of int) - Len dim. The correct shape for an array specifying the coupling strength. lat_indices has only rows within this shape.
- shift_lat_indices (array) - Translation vector from origin to the lower left corner of box spanned by \(d x\).

\section*{property dim}

The dimension of the lattice.
distance ( \(u 1, u 2, d x\) )
Get the distance for a given coupling between two sites in the lattice.
The \(u 1, u 2, d x\) parameters are defined in analogy with add_coupling(), i.e., this function calculates the distance between a pair of operators added with add_coupling (using the basis and unit_cell_positions of the lattice).

Warning: This function ignores "wrapping" arround the cylinder in the case of periodic boundary conditions.

\section*{Parameters}
- u1 (int) - Indices within the unit cell; the \(u 1\) and \(u 2\) of add_coupling()
- u2 (int) - Indices within the unit cell; the \(u 1\) and \(u 2\) of add_coupling ()
- \(\mathbf{d} \mathbf{x}\) (array) - Length dim. The translation in terms of basis vectors for the coupling.

Returns distance - The distance between site at lattice indices [ \(\mathrm{x}, \mathrm{y}, \mathrm{u} 1]\) and \([\mathrm{x}+\) \(d x[0], y+d x[1], u 2]\), ignoring any boundary effects.

Return type float
```

enlarge_mps_unit_cell(factor=2)

```

Repeat the unit cell for infinite MPS boundary conditions; in place.
Parameters factor (int) - The new number of sites in the MPS unit cell will be increased from \(N\) _sites to factor*N_sites_per_ring. Since MPS unit cells are repeated in
the \(x\)-direction in our convetion, the lattice shape goes from ( \(L x, L y, \ldots, L u\) ) to (Lx*factor, Ly, ..., Lu).
find_coupling_pairs (max_dx=3, cutoff=None, eps=1e-10)
Automatically find coupling pairs grouped by distances.
Given the unit_cell_positions and basis, the coupling pairs of nearest_neighbors, next_nearest_neighbors etc at a given distance are basically fixed (although not uniquely, since we take out half of them to avoid double-counting couplings in both directions A_i B_j + B_i A_i). This function iterates through all possible couplings up to a given cutoff distance and then determines the possible pairs at fixed distances (up to round-off errors).

\section*{Parameters}
- max_dx (int) - Maximal index for each index of \(d x\) to iterate over. You need large enough values to include every possible coupling up to the desired distance, but choosing it too large might make this function run for a long time.
- cutoff (float) - Maximal distance (in the units in which basis and unit_cell_positions is given).
- eps (float) - Tolerance up to which to distances are considered the same.

Returns coupling_pairs - Keys are distances of nearest-neighbors, next-nearest-neighbors etc. Values are [ (u1, u2, dx), ...] as in pairs.
Return type dict
classmethod from_hdf5 (hdf5_loader, h5gr, subpath)
Load instance from a HDF5 file.
This method reconstructs a class instance from the data saved with save_hdf5 ().

\section*{Parameters}
- hdf5_loader (Hdf5Loader) - Instance of the loading engine.
- h5gr (Group) - HDF5 group which is represent the object to be constructed.
- subpath (str) - The name of \(h 5 g r\) with a ' /' in the end.

Returns obj - Newly generated class instance containing the required data.
Return type cls
lat2mps_idx (lat_idx)
Translate lattice indices ( \(\mathrm{x} \_0, \ldots, \mathrm{x}_{-}\{\mathrm{D}-1\}, \mathrm{u}\) ) to MPS index \(i\).
Parameters lat_idx (array_like [.., dim+1]) - The last dimension corresponds to lattice indices \(\left(x_{-} 0, \ldots, x_{-}\{D-1\}, u\right)\). All lattice indices should be positive and smaller than the corresponding entry in self.shape. Exception: for "infinite" \(b c_{-} M P S\), an \(x \_0\) outside indicates shifts accross the boundary.

Returns i - MPS index/indices corresponding to lat_idx. Has the same shape as lat_idx without the last dimension.

Return type array_like
mps2lat_idx(i)
Translate MPS index \(i\) to lattice indices ( \(\left.x_{-} 0, \ldots, x_{-}\{d i m-1\}, u\right)\).
Parameters i (int / array_like of int)-MPS index/indices.

Returns lat_idx - First dimensions like \(i\), last dimension has len \(\operatorname{dim}^{`}+1\) and contains the lattice indices " \(\left(x \_0, \ldots, x_{\_}\{d i m-1\}, u\right)\) ` corresponding to \(i\). For \(i\) accross the MPS unit cell and "infinite" \(b c_{-} M P S\), we shift \(x_{-} 0\) accordingly.

\section*{Return type array}
mps2lat_values ( \(A\), axes \(=0, u=\) None)
Reshape/reorder \(A\) to replace an MPS index by lattice indices.

\section*{Parameters}
- A (ndarray) - Some values. Must have A.shape[axes] = self.N_sites if \(u\) is None, or A.shape[axes] = self.N_cells if \(u\) is an int.
- axes ( (iterable of) int) - chooses the axis which should be replaced.
- u (None I int) - Optionally choose a subset of MPS indices present in the axes of \(A\), namely the indices corresponding to self.unit_cell[u], as returned by mps_idx_fix_u(). The resulting array will not have the additional dimension(s) of \(u\).

Returns res_A - Reshaped and reordered verions of A. Such that MPS indices along the specified axes are replaced by lattice indices, i.e., if MPS index \(j\) maps to lattice site ( \(x 0, x 1, x 2\) ), then res_A[..., \(x 0, x 1, x 2, \ldots]=A[\ldots, j, \ldots]\).

Return type ndarray

\section*{Examples}

Say you measure expection values of an onsite term for an MPS, which gives you an 1D array \(A\), where \(A[i]\) is the expectation value of the site given by self.mps2lat_idx(i). Then this function gives you the expectation values ordered by the lattice:
```

>>> print(lat.shape, A.shape)
(10, 3, 2) (60,)
>>> A_res = lat.mps2lat_values(A)
>>> A_res.shape
(10, 3, 2)
>>> A_res[tuple(lat.mps2lat_idx(5))] == A[5]
True

```

If you have a correlation function C [i, j], it gets just slightly more complicated:
```

>>> print(lat.shape, C.shape)
(10, 3, 2) (60, 60)
>>> lat.mps2lat_values(C, axes=[0, 1]).shape
(10, 3, 2, 10, 3, 2)

```

If the unit cell consists of different physical sites, an onsite operator might be defined only on one of the sites in the unit cell. Then you can use mps_idx_fix_u() to get the indices of sites it is defined on, measure the operator on these sites, and use the argument \(u\) of this function.
```

>>> u = 0
>>> idx_subset = lat.mps_idx_fix_u(u)
>>> A_u = A[idx_subset]
>>> A_u_res = lat.mps2lat_values(A_u, u=u)
>>> A_u_res.shape
(10, 3)
>>> np.all(A_res[:, :, u] == A_u_res[:, :])
True

```
mps2lat_values_masked (A, axes=- \(1, m p s \_i n d s=N o n e\), include_u=None)
Reshape/reorder an array \(A\) to replace an MPS index by lattice indices.
This is a generalization of mps2lat_values () allowing for the case of an arbitrary set of MPS indices present in each axis of \(A\).

\section*{Parameters}
- A (ndarray) - Some values.
- axes ((iterable of) int) - Chooses the axis of \(A\) which should be replaced. If multiple axes are given, you also need to give multiple index arrays as mps_inds.
- mps_inds ((list of) 1D ndarray) - Specifies for each axis in axes, for which MPS indices we have values in the corresponding axis of \(A\). Defaults to [np. arange (A. shape[ax]) for ax in axes]. For indices accross the MPS unit cell and "infinite" \(b c_{-} M P S\), we shift \(x \_0\) accordingly.
- include_u ((list of) bool) - Specifies for each axis in axes, whether the \(u\) index of the lattice should be included into the output array res_A. Defaults to len (self. unit_cell) > 1 .

Returns res_A - Reshaped and reordered copy of A. Such that MPS indices along the specified axes are replaced by lattice indices, i.e., if MPS index \(j\) maps to lattice site ( \(x 0, x 1, x 2\) ), then res_A[..., \(x 0, x 1, x 2, \ldots]=A\left[. . ., ~ m p s \_i n d s[j], \ldots\right]\).

Return type np.ma.MaskedArray
mps_idx_fix_u (u=None)
return an index array of MPS indices for which the site within the unit cell is \(u\).
If you have multiple sites in your unit-cell, an onsite operator is in general not defined for all sites. This functions returns an index array of the mps indices which belong to sites given by self. unit_cell[u].

Parameters u(None / int) - Selects a site of the unit cell. None (default) means all sites.
Returns mps_idx - MPS indices for which self.site(i) is self.unit_cell[u]. Ordered ascending.

\section*{Return type array}
mps_lat_idx_fix_u (u=None)
Similar as mps_idx_fix_u(), but return also the corresponding lattice indices.
Parameters \(\mathbf{u}\) (None / int) - Selects a site of the unit cell. None (default) means all sites.

\section*{Returns}
- mps_idx (array) - MPS indices \(i\) for which self.site(i) is self. unit_cell[u].
- lat_idx (2D array) - The row \(j\) contains the lattice index (without \(u\) ) corresponding to mps_idx[j].
mps_sites ()
Return a list of sites for all MPS indices.
Equivalent to [self.site(i) for i in range(self.N_sites)].
This should be used for sites of 1D tensor networks (MPS, MPO,...).
multi_coupling_shape ( \(d x\) )
Calculate correct shape of the strengths for a multi_coupling.
Parameters \(d \mathbf{x}(2 D\) array, shape (N_ops, dim)) \(-d x[i, \quad:]\) is the translation vector in the lattice for the \(i\)-th operator. Corresponds to the \(d x\) of each operator given in the argument ops of tenpy.models.model.CouplingModel.add_multi_coupling().

\section*{Returns}
- coupling_shape (tuple of int) - Len dim. The correct shape for an array specifying the coupling strength. lat_indices has only rows within this shape.
- shift_lat_indices (array) - Translation vector from origin to the lower left corner of box spanned by \(d x\). (Unlike for coupling_shape () it can also contain entries \(>0\) )
number_nearest_neighbors ( \(u=0\) )
Deprecated.
Deprecated since version 0.5.0: Use count_neighbors () instead.
number_next_nearest_neighbors ( \(u=0\) )
Deprecated.
Deprecated since version 0.5.0: Use count_neighbors () instead.

\section*{property order}

Defines an ordering of the lattice sites, thus mapping the lattice to a 1 D chain.
Each row of the array contains the lattice indices for one site, the order of the rows thus specifies a path through the lattice, along which an MPS will wind through through the lattice.
You can visualize the order with plot_order ().
ordering (order)
Provide possible orderings of the \(N\) lattice sites.
This function can be overwritten by derived lattices to define additional orderings. The following orders are defined in this method:
\begin{tabular}{|c|c|c|}
\hline order & equivalent priority & equivalent snake_winding \\
\hline 'Cstyle' & (0, 1, .., dim-1, dim) & (False, ..., False, False) \\
\hline 'default' & & \\
\hline 'snake' & (0, 1, .., dim-1, dim) & (True, . . . True, True) \\
\hline 'snakeCstyle' & & \\
\hline 'Fstyle' & (dim-1, .., 1, 0, dim) & (False, ..., False, False) \\
\hline 'snakeFstyle' & (dim-1, .., 1, 0, dim) & (False, ..., False, False) \\
\hline
\end{tabular}

Note: For lattices with a non-trivial unit cell (e.g. Honeycomb, Kagome), the grouped order might be more appropriate, see get_order_grouped()

Parameters order (str | ('standard', snake_winding, priority) | ('grouped', groups, ...)) - Specifies the desired ordering using one of the strings of the above tables. Alternatively, an ordering is specified by a tuple with first entry specifying a function, 'standard' for get_order() and 'grouped' for get_order_grouped(), and other arguments in the tuple as specified in the documentation of these functions.
Returns order - the order to be used for order.


Return type array, shape (N, D+1), dtype np.intp

\section*{See also:}
get_order generates the order from equivalent priority and snake_winding.
get_order_grouped variant of get_order.
plot_order visualizes the resulting order.
plot_basis \(\left(a x\right.\), origin \(=(0.0,0.0)\), shade \(=\) None, \({ }^{* * * w a r g s) ~}\)
Plot arrows indicating the basis vectors of the lattice.

\section*{Parameters}
- ax (matplotlib.axes.Axes) - The axes on which we should plot.
- **kwargs - Keyword arguments for ax. arrow.
plot_bc_identified (ax, direction=- 1, origin=None, cylinder_axis=False, **kwargs)
Mark two sites indified by periodic boundary conditions.
Works only for lattice with a 2-dimensional basis.

\section*{Parameters}
- ax (matplotlib.axes.Axes) - The axes on which we should plot.
- direction (int) - The direction of the lattice along which we should mark the idenitified sites. If None, mark it along all directions with periodic boundary conditions.
- cylinder_axis (b○○l) - Whether to plot the cylinder axis as well.
- origin (None / np.ndarray) - The origin starting from where we mark the identified sites. Defaults to the first entry of unit_cell_positions.
- **kwargs - Keyword arguments for the used ax. plot.
plot_coupling (ax, coupling=None, wrap=False, **kwargs)
Plot lines connecting nearest neighbors of the lattice.

\section*{Parameters}
- ax (matplotlib. axes.Axes) - The axes on which we should plot.
- coupling (list of (u1, u2, dx)) - By default (None), use self. pairs['nearest_neighbors']. Specifies the connections to be plotted; iteating over lattice indices ( \(i 0, i 1, \ldots\) ), we plot a connection from the site (i0, i1, ..., u1) to the site ( \(i 0+d x[0]\), \(i 1+d x[1], \ldots, u 2\) ), taking into account the boundary conditions.
- wrap (bool) - If True, plot couplings going around the boundary by directly connecting the sites it connects. This might be hard to see, as this puts lines from one end of the lattice to the other. If False, plot the couplings as dangling lines.
- **kwargs - Further keyword arguments given to ax.plot ().
plot_order (ax, order=None, textkwargs=\{'color': 'r'\}, **kwargs)
Plot a line connecting sites in the specified "order" and text labels enumerating them.

\section*{Parameters}
- ax (matplotlib. axes.Axes) - The axes on which we should plot.
- order (None / 2D array (self.N_sites, self.dim+1))- The order as returned by ordering(); by default (None) use order.
- textkwargs (None I dict) - If not None, we add text labels enumerating the sites in the plot. The dictionary can contain keyword arguments for ax.text ().
- **kwargs - Further keyword arguments given to ax. plot ().
plot_sites (ax, markers=['o', ' \({ }^{\prime}\) ', 's', ' \(p^{\prime}\) ' ' \(h^{\prime}\) ', \(\left.D^{\prime}\right]\), **kwargs)
Plot the sites of the lattice with markers.

\section*{Parameters}
- ax (matplotlib. axes.Axes) - The axes on which we should plot.
- markers (list) - List of values for the keywork marker of ax.plot () to distinguish the different sites in the unit cell, a site \(u\) in the unit cell is plotted with a marker markers [u \% len(markers)].
- **kwargs - Further keyword arguments given to ax. plot ().
position(lat_idx)
return 'space' position of one or multiple sites.
Parameters lat_idx (ndarray, (. . , dim+1)) - Lattice indices.
Returns pos - The position of the lattice sites specified by lat_idx in real-space.
Return type ndarray, (..., dim)
possible_couplings ( \(u 1, u 2, d x\), strength=None)
Find possible MPS indices for two-site couplings.
For periodic boundary conditions ( \(\mathrm{bc}[\mathrm{a}]==\) False ) the index x _a is taken modulo Ls [a] and runs through range (Ls [a]). For open boundary conditions, \(x_{-}\)a is limited to \(0<=x_{-}<\)Ls [a] and \(0<=x \_a+d x[a]<\) lat.Ls[a].

\section*{Parameters}
- u1 (int) - Indices within the unit cell; the \(u 1\) and \(u 2\) of add_coupling ()
- u2 (int) - Indices within the unit cell; the \(u 1\) and \(u 2\) of add_coupling ()
- \(\mathbf{d x}\) (array) - Length dim. The translation in terms of basis vectors for the coupling.
- strength (array_like / None) - If given, instead of returning lat_indices and coupling_shape directly return the correct strength_12.

\section*{Returns}
- mps1, mps2 (1D array) - For each possible two-site coupling the MPS indices for the \(u 1\) and \(u 2\).
- strength_vals (1D array) - (Only returend if strength is not None.) Such that for (i, \(j, ~ s) ~ i n ~ z i p\left(m p s 1, ~ m p s 2, ~ s t r e n g t h \_v a l s\right): ~ i t e r a t e s ~ o v e r ~ a l l ~ p o s s i b l e ~ c o u-~\) plings with \(s\) being the strength of that coupling. Couplings where strength_vals \(==0\). are filtered out.
- lat_indices ( \(2 D\) int array) - (Only returend if strength is None.) Rows of lat_indices correspond to entries of \(m p s l\) and \(m p s 2\) and contain the lattice indices of the "lower left corner" of the box containing the coupling.
- coupling_shape (tuple of int) - (Only returend if strength is None.) Len dim. The correct shape for an array specifying the coupling strength. lat_indices has only rows within this shape.
possible_multi_couplings (ops, strength=None)
Generalization of possible_couplings () to couplings with more than 2 sites.
Parameters ops (list of (opname, \(\mathrm{dx}, \mathrm{u})\) ) - Same as the argument ops of add_multi_coupling().

\section*{Returns}
- mps_ijkl (2D int array) - Each row contains MPS indices \(i, j, k, l, \ldots\). . for each of the operators positions. The positions are defined by \(d x(\mathrm{j}, \mathrm{k}, \mathrm{l}, \ldots\) relative to \(i\) ) and boundary coundary conditions of self (how much the box for given \(d x\) can be shifted around without hitting a boundary - these are the different rows).
- strength_vals ( \(1 D\) array) - (Only returend if strength is not None.) Such that for (ijkl, s) in zip(mps_ijkl, strength_vals): iterates over all possible couplings with \(s\) being the strength of that coupling. Couplings where strength_vals \(==0\). are filtered out.
- lat_indices ( 2 D int array) - (Only returend if strength is None.) Rows of lat_indices correspond to rows of mps_ijkl and contain the lattice indices of the "lower left corner" of the box containing the coupling.
- coupling_shape (tuple of int) - (Only returend if strength is None.) Len dim. The correct shape for an array specifying the coupling strength. lat_indices has only rows within this shape.
```

save_hdf5 (hdf5_saver, h5gr, subpath)

```

Export self into a HDF5 file.
This method saves all the data it needs to reconstruct self with \(f r o m \_h d f 5\) ().
Specifically, it saves unit_cell, Ls, unit_cell_positions, basis, boundary_conditions, pairs under their name, bc_MPS as "boundary_conditions_MPS", and order as "order_for_MPS". Moreover, it saves dim and N_sites as HDF5 attributes.

\section*{Parameters}
- hdf5_saver (Hdf5Saver) - Instance of the saving engine.
- h5gr (: class `Group ') - HDF5 group which is supposed to represent self.
- subpath (str) - The name of \(h 5 g r\) with a ' / ' in the end.
```

site(i)

```
return Site instance corresponding to an MPS index \(i\)
test_sanity ()
Sanity check.
Raises ValueErrors, if something is wrong.

\section*{Functions}
\begin{tabular}{ll}
\hline get_lattice(lattice_name) & \begin{tabular}{l} 
Given the name of a Lattice class, get the lattice class \\
itself.
\end{tabular} \\
\hline get_order(shape, snake_winding[, priority]) & \begin{tabular}{l} 
Built the Lattice.order in (Snake-) C-Style for a \\
given lattice shape.
\end{tabular} \\
\hline get_order_grouped(shape, groups[, priority]) & \begin{tabular}{l} 
Variant of get_order(), grouping some sites of the \\
unit cell.
\end{tabular} \\
\hline
\end{tabular}

\subsection*{19.1.12 get_lattice}
- full name: tenpy.models.lattice.get_lattice
- parent module: tenpy.models.lattice
- type: function
tenpy.models.lattice.get_lattice(lattice_name)
Given the name of a Lattice class, get the lattice class itself.
Parameters lattice_name (str) - Name of a Lattice class defined in the module lattice, for example "Chain", "Square", "Honeycomb", ....
Returns LatticeClass - The lattice class (type, not instance) specified by lattice_name.

\section*{Return type Lattice}

\subsection*{19.1.13 get_order}
- full name: tenpy.models.lattice.get_order
- parent module: tenpy.models.lattice
- type: function
tenpy.models.lattice.get_order (shape, snake_winding, priority=None)
Built the Lattice. order in (Snake-) C-Style for a given lattice shape.

Note: In this doc-string, the word 'direction' referst to a physical direction of the lattice or the index \(u\) of the unit cell as an "artificial direction".

\section*{Parameters}
- shape (tuple of int) - The shape of the lattice, i.e., the length in each direction.
- snake_winding (tuple of bool) - For each direction one bool, whether we should wind as a "snake" (True) in that direction (i.e., going forth and back) or simply repeat ascending (False)
- priority (None I tuple of float) - If None (default), use C-Style ordering. Otherwise, this defines the priority along which direction to wind first; the direction with the highest priority increases fastest. For example, "C-Style" order is enforced by priority=(0, 1, 2, ...), and Fortrans F-style order is enforced by priority=(dim, dim-1, ..., 1, 0)
- group (None I tuple of tuple) - If None (default), ignore it. Otherwise, it specifies that we group the fastests changing dimension

Returns order - An order of the sites for Lattice. order in the specified ordering.
Return type ndarray (np.prod(shape), len(shape))

\section*{See also:}

Lattice. ordering method in Lattice to obtain the order from parameters.
Lattice.plot_order visualizes the resulting order in a Lattice.
get_order_grouped a variant grouping sites of the unit cell.

\subsection*{19.1.14 get_order_grouped}
- full name: tenpy.models.lattice.get_order_grouped
- parent module: tenpy.models.lattice
- type: function
tenpy.models.lattice.get_order_grouped (shape, groups, priority=None)
Variant of get_order (), grouping some sites of the unit cell.
This function is usefull for lattices with a unit cell of more than 2 sites (e.g. Kagome). For 2D lattices with a unit cell, the ordering goes first within a group, then along \(y\), then the next group (for the same \(x\)-value), again along y , and finally along x when all groups are done.

As an example, consider the Kagome lattice.


Note: In this doc-string, the word 'direction' referst to a physical direction of the lattice or the index \(u\) of the unit cell as an "artificial direction".

\section*{Parameters}
- shape (tuple of int) - The shape of the lattice, i.e., the length in each direction.
- groups (tuple of tuple of int) - A partition and reordering of range (shape [-1]) into smaller groups. The ordering goes first within a group, then along the last spatial dimensions, then changing between different groups and finally in Cstyle order along the remaining spatial dimensions.
- priority (None | tuple of ints) - By default (None), use C-style order for everything except the unit cell, as shown above. If a tuple, it should have length len ( shape) and specifies which order to go first, similarly as in get_order (). To group sites in the unit cell, you should make the last entry of priority the largest. However, you can also choose to group along another direction - in that case groups should be a partitioning of range (shape (argmax (priority))). Try and plot it, if you need it!

Returns order - An order of the sites for Lattice. order in the specified ordering.
Return type ndarray (np.prod(shape), len(shape))

\section*{See also:}

Lattice.ordering() method in Lattice to obtain the order from parameters.
Lattice.plot_order() visualizes the resulting order in a Lattice.

\section*{Module description}

Classes to define the lattice structure of a model.
The base class Lattice defines the general structure of a lattice, you can subclass this to define you own lattice. The SimpleLattice is a slight simplification for lattices with a single-site unit cell. Further, we have some predefined lattices, namely Chain, Ladder in 1D and Square, Triangular, Honeycomb, and Kagome in 2D.

The IrregularLattice provides a way to remove or add sites to an existing, regular lattice.
See also the Models and Details on the lattice geometry.

\section*{19.2 model}
- full name: tenpy.models.model
- parent module: tenpy.models
- type: module

\section*{Classes}

\begin{tabular}{ll}
\hline CouplingMPOModel(model_params) & \begin{tabular}{l} 
Combination of the CouplingModel and \\
MPOModel.
\end{tabular} \\
\hline CouplingModel(lattice[, bc_coupling, ...]) & \begin{tabular}{l} 
Base class for a general model of a Hamiltonian consist- \\
ing of two-site couplings.
\end{tabular} \\
\hline MPOModel(lattice, H_MPO) & \begin{tabular}{l} 
Base class for a model with an MPO representation of \\
the Hamiltonian.
\end{tabular} \\
\hline Model(lattice) & Base class for all models. \\
\hline MultiCouplingModel(lattice[, bc_coupling, ...]) & \begin{tabular}{l} 
Deprecated class which was a generalization of the Cou- \\
plingModel.
\end{tabular} \\
\hline NearestNeighborModel(lattice, H_bond) & \begin{tabular}{l} 
Base class for a model of nearest neigbor interactions \\
w.r.t.
\end{tabular} \\
\hline
\end{tabular}

\subsection*{19.2.1 CouplingModel}
- full name: tenpy.models.model.CouplingModel
- parent module: tenpy.models.model
- type: class

\section*{Inheritance Diagram}


\section*{Methods}
\begin{tabular}{|c|c|}
\hline CouplingModel.__init__(lattice[,...]) In & Initialize self. \\
\hline  & Add twosite coupling terms to the Hamiltonian, summing over lattice sites. \\
\hline CouplingModel.add_coupling_term(strength, A i, ...) & Add a two-site coupling term on given MPS sites. \\
\hline CouplingModel.add_exponentially_decayind & Add anjexpométially decaying long-range coupling. \\
\hline CouplingModel.add_local_term(strength, term) & Add a single term to self. \\
\hline CouplingModel.add_multi_coupling(strength, Ad ops) & Add multi-site coupling terms to the Hamiltonian, summing over lattice sites. \\
\hline ```
CouplingModel.add_multi_coupling_term(..A
...])
``` & Add a general M-site coupling term on given MPS sites. \\
\hline CouplingModel.add_onsite(strength, u, op- A name) & Add onsite terms to onsite_terms. \\
\hline CouplingModel.add_onsite_term(strength, i, op) & Add an onsite term on a given MPS site. \\
\hline CouplingModel.all_coupling_terms() S & Sum of all coupling_terms. \\
\hline CouplingModel.all_onsite_terms() S & Sum of all onsite_terms. \\
\hline CouplingModel.calc_H_MPO([tol_zero]) C & Calculate MPO representation of the Hamiltonian. \\
\hline CouplingModel.calc_H_bond([tol_zero]) & calculate H_bond from coupling_terms and onsite_terms. \\
\hline CouplingModel.calc_H_onsite([tol_zero]) C & Calculate H_onsite from self.onsite_terms. \\
\hline CouplingModel.coupling_strength_add_extA & Add an(external flux to the coupling strength. \\
\hline CouplingModel.enlarge_mps_unit_cell([factdR & dR)peat the unit cell for infinite MPS boundary conditions; in place. \\
\hline CouplingModel.from_hdf5(hdf5_loader, h5gr, L ...) & Load instance from a HDF5 file. \\
\hline
\end{tabular}

Table 27 - continued from previous page
CouplingModel.group_sites([n, Modify self in place to group sites.
grouped_sites])
CouplingModel.save_hdf5(hdf5_saver, h5gr, Export self into a HDF5 file.
...)
CouplingModel.test_sanity () Sanity check, raises ValueErrors, if something is wrong.

\section*{Class Attributes and Properties}
CouplingModel.logger class attribute.
class tenpy.models.model.CouplingModel (lattice, ex-
Bases: tenpy.models.model.Model plicit_plus_hc=False)

Base class for a general model of a Hamiltonian consisting of two-site couplings.
In this class, the terms of the Hamiltonian are specified explicitly as OnsiteTerms or CouplingTerms.
Deprecated since version 0.4.0: bc_coupling will be removed in 1.0.0. To specify the full geometry in the lattice, use the \(b c\) parameter of the Lattice.

\section*{Parameters}
- lattice (Lattice) - The lattice defining the geometry and the local Hilbert space(s).
- bc_coupling ((iterable of) \{'open' | 'periodic' | int\}) - Boundary conditions of the couplings in each direction of the lattice. Defines how the couplings are added in add_coupling (). A single string holds for all directions. An integer shift means that we have periodic boundary conditions along this direction, but shift/tilt by -shift*lattice.basis[0] (~cylinder axis for bc_MPS='infinite') when going around the boundary along this direction.
- explicit_plus_hc (bool) - If True, the Hermitian conjugate of the MPO is computed at runtime, rather than saved in the MPO.

\section*{onsite_terms}

The OnsiteTerms ordered by category.
Type \{'category': OnsiteTerms\}
coupling_terms
The CouplingTerms ordered by category. In case we've added terms with more than 2 operators, e.g. with add_multi_coupling(), the values of the dictionary may also be MultiCouplingTerms.

Type \{'category': CouplingTerms \(\}\)
exp_decaying_terms
Collection of coupling terms with exponentially decaying long-range interactions. Filled by add_exponentially_decaying_coupling().

Type ExponentiallyDecayingTerms
explicit_plus_hc
If True, self represents the terms in onsite_terms, coupling_terms and exp_decaying_terms plus their hermitian conjugate added. The flag will be carried on to the MPO, which will have a reduced bond dimension if self.add_coupling (..., plus_hc=True) was used. Note that add_onsite(), add_coupling(), add_multi_coupling() and
add_exponentially_decaying_coupling() respect this flag, ensuring that the represented Hamiltonian is indepentent of the explicit_plus_hc flag.

Type bool
test_sanity()
Sanity check, raises ValueErrors, if something is wrong.
add_local_term (strength, term, category=None, plus_hc=False)
Add a single term to self.
The repesented term is strength times the product of the operators given in terms. Each operator is specified by the name and the site it acts on; the latter given by a lattice index, see Lattice.

Depending on the length of term, it can add an onsite term or a coupling term to onsite_terms or coupling_terms, respectively.

\section*{Parameters}
- strength (float/complex) - The prefactor of the term.
- term (list of (str, array_like))-List of tuples (opname, lat_idx) where opname is a string describing the operator acting on the site given by the lattice index lat_idx. Here, lat_idx is for example \([x, y, u]\) for a 2D lattice, with \(u\) being the index within the unit cell.
- category - Descriptive name used as key for onsite_terms or coupling_terms.
- plus_hc (b○○l) - If True, the hermitian conjugate of the terms is added automatically.
add_onsite (strength, u, opname, category=None, plus_hc=False)
Add onsite terms to onsite_terms.
Adds \(\sum_{\vec{x}}\) strength \([\vec{x}] * O P\) to the represented Hamiltonian, where the operator OP=lat. unit_cell[u].get_op (opname) acts on the site given by a lattice index (x_0, ..., \(\left.x_{-}\{\operatorname{dim}-1\}, u\right)\),

The necessary terms are just added to onsite_terms; doesn't rebuild the MPO.

\section*{Parameters}
- strength (scalar / array) - Prefactor of the onsite term. May vary spatially. If an array of smaller size is provided, it gets tiled to the required shape.
- u(int)-Picks a Site lat.unit_cell[u] out of the unit cell.
- opname (str) - valid operator name of an onsite operator in lat. unit_cell[u].
- category (str) - Descriptive name used as key for onsite_terms. Defaults to opname.
- plus_hc (bool) - If True, the hermitian conjugate of the terms is added automatically.

\section*{See also:}
add_coupling Add a terms acting on two sites.
add_onsite_term Add a single term without summing over vecx.
add_onsite_term (strength, \(i, o p\), category=None, plus_hc=False)
Add an onsite term on a given MPS site.
Wrapper for self.onsite_terms[category].add_onsite_term(....).

\section*{Parameters}
- strength (float) - The strength of the term.
- i (int) - The MPS index of the site on which the operator acts. We require \(0<=i<\) L.
- op (str) - Name of the involved operator.
- category (str) - Descriptive name used as key for onsite_terms. Defaults to op.
- plus_hc (bool) - If True, the hermitian conjugate of the term is added automatically.
all_onsite_terms()
Sum of all onsite_terms.
add_coupling (strength, u1, op1, u2, op2, dx, op_string=None, str_on_first=True, raise_op2_left=False, category=None, plus_hc=False)
Add twosite coupling terms to the Hamiltonian, summing over lattice sites.
Represents couplings of the form \(\sum_{x_{0}, \ldots, x_{d i m-1}} \operatorname{strength}[\operatorname{shift}(\vec{x})] * O P 0 * O P 1\), where OP 0 := lat.unit_cell[u0].get_op(op0) acts on the site (x_0, ..., x_\{dim-1\}, u1), and OP1 := lat.unit_cell[u1].get_op(op1) acts on the site (x_0+dx[0], ..., \(x_{-}\{\operatorname{dim}-1\}+d x\left[\right.\) dim-1], u1). Possible combinations \(x_{-} 0, \ldots, x_{-}\{d i m-1\}\) are determined from the boundary conditions in possible_couplings ().
The coupling strength may vary spatially if the given strength is a numpy array. The correct shape of this array is the coupling_shape returned by tenpy.models.lattice.coupling_shape() and depends on the boundary conditions. The shift (...) depends on \(d x\), and is chosen such that the first entry strength \([0,0, \ldots]\) of strength is the prefactor for the first possible coupling fitting into the lattice if you imagine open boundary conditions.

The necessary terms are just added to coupling_terms; this function does not rebuild the MPO.
Deprecated since version 0.4.0: The arguments str_on_first and raise_op2_left will be removed in version 1.0.0.

\section*{Parameters}
- strength (scalar | array) - Prefactor of the coupling. May vary spatially (see above). If an array of smaller size is provided, it gets tiled to the required shape.
- u1 (int) - Picks the site lat. unit_cell [u1] for OP1.
- op1 (str) - Valid operator name of an onsite operator in lat.unit_cell[u1] for OP1.
- u2 (int) - Picks the site lat.unit_cell [u2] for OP2.
- op2 (str) - Valid operator name of an onsite operator in lat.unit_cell[u2] for OP2.
- \(\mathbf{d x}\) (iterable of int) - Translation vector (of the unit cell) between OP1 and OP2. For a 1D lattice, a single int is also fine.
- op_string (str | None) - Name of an operator to be used between the OP1 and OP2 sites. Typical use case is the phase for a Jordan-Wigner transformation. The operator should be defined on all sites in the unit cell. If None, auto-determine whether a JordanWigner string is needed, using op_needs_JW ().
- str_on_first (bool) - Whether the provided op_string should also act on the first site. This option should be chosen as True for Jordan-Wigner strings. When handling Jordan-Wigner strings we need to extend the op_string to also act on the 'left', first site (in the sense of the MPS ordering of the sites given by the lattice). In this case, there is a
well-defined ordering of the operators in the physical sense (i.e. which of opl or op2 acts first on a given state). We follow the convention that op2 acts first (in the physical sense), independent of the MPS ordering. Deprecated.
- raise_op2_left (bool) - Raise an error when op2 appears left of opl (in the sense of the MPS ordering given by the lattice). Deprecated.
- category (str) - Descriptive name used as key for coupling_terms. Defaults to a string of the form " \(\{o p 1\} \_i \quad\{o p 2\} \_j "\).
- plus_hc (bool) - If True, the hermitian conjugate of the terms is added automatically.

\section*{Examples}

When initializing a model, you can add a term \(J \sum_{<i, j>} S_{i}^{z} S_{j}^{z}\) on all nearest-neighbor bonds of the lattice like this:
```

>>> J = 1. \# the strength
>>> for ul, u2, dx in self.lat.pairs['nearest_neighbors']:
... self.add_coupling(J, u1, 'Sz', u2, 'Sz', dx)

```

The strength can be an array, which gets tiled to the correct shape. For example, in a 1D Chain with an even number of sites and periodic (or infinite) boundary conditions, you can add alternating strong and weak couplings with a line like:
```

>>> self.add_coupling([1.5, 1.], u1, 'Sz', u2, 'Sz', dx)

```

Make sure to use the plus_hc argument if necessary, e.g. for hoppings:
```

>>> t = 1. \# hopping strength
>>> for u1, u2, dx in self.lat.pairs['nearest_neighbors']:
... self.add_coupling(t, u1, 'Cd', u2, 'C', dx, plus_hc=True)

```

Alternatively, you can add the hermitian conjugate terms explictly. The correct way is to complex conjugate the strength, take the hermitian conjugate of the operators and swap the order (including a swap \(u 1<->u 2\) ), and use the opposite direction -dx , i.e. the h.c. of add_coupling ( \(t, \mathrm{u} 1\), 'A', u2, 'B', dx) is add_coupling(np.conj(t), u2, hc('B'), u1, hc('A'), -dx), where \(h c\) takes the hermitian conjugate of the operator names, see get_hc_op_name (). For spin-less fermions (FermionSite), this would be
```

>>> for u1, u2, dx in self.lat.pairs['nearest_neighbors']:
... self.add_coupling(t, u1, 'Cd', u2, 'C', dx)
... self.add_coupling(np.conj(t), u2, 'Cd', u1, 'C', -dx) \# h.c.

```

With spin-full fermions (SpinHalfFermions), it could be:
```

>>> for ul, u2, dx in self.lat.pairs['nearest_neighbors']:
... self.add_coupling(t, u1, 'Cdu', u2, 'Cd', dx) \# Cdagger_up C_down
... self.add_coupling(np.conj(t), u2, 'Cdd', ul, 'Cu', -dx) \# h.c.,
\hookrightarrowCdagger_down C_up

```

Note that the Jordan-Wigner strings for the fermions are added automatically!

\section*{See also:}
add_onsite Add terms acting on one site only.
add_multi_coupling_term for terms on more than two sites.
add_coupling_term Add a single term without summing over \(\vec{x}\).
add_coupling_term (strength, \(i, j, o p_{-} i, o p_{-} j, o p_{-}\)string \(=' I d^{\prime}\), category=None, plus_hc=False)
Add a two-site coupling term on given MPS sites.
Wrapper forself.coupling_terms[category].add_coupling_term(...).

Warning: This function does not handle Jordan-Wigner strings! You might want to use add_local_term() instead.

\section*{Parameters}
- strength (float) - The strength of the coupling term.
- i (int) - The MPS indices of the two sites on which the operator acts. We require 0 \(<=i<N \_s i t e s\) and \(i<j\), i.e., op_i acts "left" of \(o p_{-} j\). If \(j>=N_{-}\)sites, it indicates couplings between unit cells of an infinite MPS.
- j (int) - The MPS indices of the two sites on which the operator acts. We require 0 \(<=i<N \_s i t e s\) and \(i<j\), i.e., op_i acts "left" of \(o p_{-} j\). If \(\mathrm{j}>=\mathrm{N}_{-}\)sites, it indicates couplings between unit cells of an infinite MPS.
- op1 (str) - Names of the involved operators.
- op2 (str) - Names of the involved operators.
- op_string (str) - The operator to be inserted between \(i\) and \(j\).
- category (str) - Descriptive name used as key for coupling_terms. Defaults to a string of the form " \(\{o p 1\} \_i \quad\{o p 2\} \_j "\).
- plus_hc (bool) - If True, the hermitian conjugate of the term is added automatically.

\section*{all_coupling_terms()}

Sum of all coupling_terms.
add_multi_coupling (strength, ops, _deprecate_l='DEPRECATED', _deprecate_2 = 'DEPRECATED', op_string=None, category=None, plus_hc=False)
Add multi-site coupling terms to the Hamiltonian, summing over lattice sites.
Represents couplings of the form \(\operatorname{sum}_{\vec{x}} \operatorname{strength}[\operatorname{shift}(\vec{x})] * O P_{0} * O P_{1} * \ldots * O P_{M-1}\), involving \(M\) operators. Here, \(O P_{m}\) stands for the operator defined by the \(m\)-th tuple (opname, \(\mathrm{dx}, \mathrm{u}\) ) given in the argument ops, which determines the position \(\vec{x}+\overrightarrow{d x}\) and unit-cell index \(u\) of the site it acts on; the actual operator is given by self.lat.unit_cell[u].get_op(opname).
The coupling strength may vary spatially if the given strength is a numpy array. The correct shape of this array is the coupling_shape returned by tenpy.models.lattice. possible_multi_couplings() and depends on the boundary conditions. The shift (...) depends on the \(d x\) entries of ops and is chosen such that the first entry st rength \([0,0, \ldots]\) of strength is the prefactor for the first possible coupling fitting into the lattice if you imagine open boundary conditions.

The necessary terms are just added to coupling_terms; this function does not rebuild the MPO.
Deprecated since version 0.6.0: We switched from the three arguments \(u 0\), op0 and other_op with other_ops=[(u1, op1, dx1), (op2, u2, dx2), ...] to a single, equivalent argment ops which should now read ops=[(op0, dx0, u0), (op1, dx1, u1), (op2, dx2, u2), . ..], where \(\mathrm{dx} 0=[0] *\) self.lat.dim. Note the changed order inside the tuples!

\section*{Parameters}
- strength (scalar / array) - Prefactor of the coupling. May vary spatially, and is tiled to the required shape.
- ops (list of (opname, \(d x, u\) )) - Each tuple determines one operator of the coupling, see the description above. opname (str) is the name of the operator, \(d x\) (list of length lat.dim) is a translation vector, and \(u\) (int) is the index of lat.unit_cell on which the operator acts. The first entry of ops corresponds to \(O P_{0}\) and acts last in the physical sense.
- op_string (str / None) - If a string is given, we use this as the name of an operator to be used inbetween the operators, excluding the sites on which any operators act. This operator should be defined on all sites in the unit cell.

If None, auto-determine whether a Jordan-Wigner string is needed (using op_needs_JW ()) for each of the segments inbetween the operators and also on the sites of the left operators.
- category (str) - Descriptive name used as key for coupling_terms. Defaults to a string of the form " \(\{0 \mathrm{p} 0\}\) _i \(\left\{o t h e r \_o p s[0]\right\} \_j\) \{other_ops[1]\}_k ...".
- plus_hc (bool) - If True, the hermitian conjugate of the terms is added automatically.

\section*{Examples}

A call to add_coupling() with arguments add_coupling(strength, u1, 'A', u2, ' B ', dx ) is equivalent to the following:
```

>> dx_0 = [0] * self.lat.dim \# = [0] for a ID lattice, [0, 0] in 2D
>>> self.add_multi_coupling(strength, [('A', dx_0, u1), ('B', dx, u2)])

```

To explicitly add the hermitian conjugate (instead of simply using plus_hc = True), you need to take the complex conjugate of the strength, reverse the order of the operators and take the hermitian conjugates of the individual operator names (indicated by the hc (...), see get_hc_op_name ()):
\(\ggg\) self.add_multi_coupling(np.conj(strength), [(hc('B'), dx, u2), (hc('A'), \(\quad\), \(\left.\left.\hookrightarrow d x \_0, \quad u 1\right)\right]\) )

\section*{See also:}
add_onsite Add terms acting on one site only.
add_coupling Add terms acting on two sites.
add_multi_coupling_term Add a single term, not summing over the possible \(\vec{x}\).
add_multi_coupling_term (strength, ijkl,ops_ijkl,op_string, category=None, plus_hc=False)
Add a general M-site coupling term on given MPS sites.
Wrapper for self.coupling_terms[category].add_multi_coupling_term(...).

Warning: This function does not handle Jordan-Wigner strings! You might want to use add_local_term() instead.

\section*{Parameters}
- strength (float) - The strength of the coupling term.
- ijkl (list of int) - The MPS indices of the sites on which the operators acts. With \(i, j, k, \ldots=i j k l\), we require that they are ordered ascending, \(i<j<\mathrm{k}<\ldots\) and that \(0<=\) i \(<\) N_sites. Inidces \(>=\) N_sites indicate couplings between different unit cells of an infinite MPS.
- ops_ijkl (list of str) - Names of the involved operators on sites \(i, j, k, \ldots\).
- op_string (list of str) - Names of the operator to be inserted between the operators, e.g., op_string[0] is inserted between \(i\) and \(j\).
- category (str) - Descriptive name used as key for coupling_terms. Defaults to a string of the form " \(\{\mathrm{op} 0\} \_i \quad\{o p 1\} \_j \quad\{o p 2\} \_k \quad .\). ".
- plus_hc (bool) - If True, the hermitian conjugate of the term is added automatically.
add_exponentially_decaying_coupling (strength, lambda_, op_i,op_j, subsites=None, op_string \(=\) None, plus_hc=False)
Add an exponentially decaying long-range coupling.
\[
\text { strength } \sum_{i<j} \lambda^{|i-j|} A_{\text {subsites }[i]} B_{\text {subsites }[j]}
\]

Where the operator \(A\) is given by \(o p_{-} i\), and \(B\) is given by \(o p_{-} j\). Note that the sum over \(i, j\) is long-range, for infinite systems going beyond the MPS unit cell. Moreover, note that the distance in the exponent is the distance within subsites.

\section*{Parameters}
- strength (float) - Overall prefactor.
- lambda (float) - Decay-rate
- op_i (string) - Names for the operators.
- op_j (string) - Names for the operators.
- subsites (None / 1D array) - Selects a subset of sites within the MPS unit cell on which the operators act. Needs to be sorted. None selects all sites.
- op_string (None / str) - The operator to be inserted between \(A\) and \(B\); If None, this function checks whether a fermionic "JW" string is needed for the given operators; in this case the right \(o p_{-} j\) acts first.
- plus_hc (bool) - If True, the hermitian conjugate of the term is added automatically.

\section*{Examples}

At least for simple enough 1D chains (or ladders), you can use fit_with_sum_of_exp () to approximate a long-range function with a few sum of exponentials and then add them with this function.
```

>>> def decay(x):
... return np.exp (-0.1*x) / x**2
>>> from tenpy.tools.fit import fit_with_sum_of_exp, sum_of_exp
>>> n_exp = 5
>>> fit_range = 50
>>> lam, pref = fit_with_sum_of_exp(decay, n_exp, fit_range)
>>> x = np.arange(1, fit_range + 1)
>>> print('error in fit: {0:.3e}'.format(np.sum(np.abs(decay(x) - sum_of_
\leftrightarrowexp(lam, pref, x)))))
error in fit: 1.073e-04
>>> for pr, la in zip(pref, lam):
... self.add_exponentially_decaying_coupling(pr, la, 'N', 'N')

```
calc_H_onsite (tol_zero=le-15)
Calculate H_onsite from self.onsite_terms.
Deprecated since version 0.4.0: This function will be removed in 1.0.0. Replace calls to this function by self.all_onsite_terms().remove_zeros(tol_zero).to_Arrays(self. lat.mps_sites()). You might also want to take explicit_plus_hc into account.

Parameters tol_zero (float) - prefactors with abs (strength) < tol_zero are considered to be zero.

\section*{Returns}

\section*{- H_onsite (list of npc.Array)}
- onsite terms of the Hamiltonian. If explicit_plus_hc is True, - Hermitian conjugates of the onsite terms will be included.
calc_H_bond (tol_zero \(=1 e-15\) )
calculate \(H_{-}\)bond from coupling_terms and onsite_terms.
Parameters tol_zero (float) - prefactors with abs (strength) < tol_zero are considered to be zero.

Returns H_bond - Bond terms as required by the constructor of NearestNeighborModel. Legs are ['p0', 'p0*', 'p1', 'p1*']
Return type list of Array
:raises ValueError : if the Hamiltonian contains longer-range terms.:
calc_H_MPO (tol_zero \(=1 e-15\) )
Calculate MPO representation of the Hamiltonian.
Uses onsite_terms and coupling_terms to build an MPOGraph (and then an MPO).
Parameters tol_zero (float) - Prefactors with abs (strength) < tol_zero are considered to be zero.

Returns H_MPO - MPO representation of the Hamiltonian.
Return type MPO
coupling_strength_add_ext_flux (strength, \(d x\), phase)
Add an external flux to the coupling strength.
When performing DMRG on a "cylinder" geometry, it might be useful to put an "external flux" through the cylinder. This means that a particle hopping around the cylinder should pick up a phase given by the external flux [[resta1998]]. This is also called "twisted boundary conditions" in literature. This function adds a complex phase to the strength array on some bonds, such that particles hopping in positive direction around the cylinder pick up \(\exp (+i\) phase \()\).

Warning: For the sign of phase it is important that you consistently use the creation operator as opl and the annihilation operator as op2 in add_coupling().

\section*{Parameters}
- strength (scalar / array) - The strength to be used in add_coupling(), when no external flux would be present.
- dx (iterable of int) - Translation vector (of the unit cell) between opl and op2 in add_coupling().
- phase (iterable of float) - The phase of the external flux for hopping in each direction of the lattice. E.g., if you want flux through the cylinder on which you have an infinite MPS, you should give phase \(=[0\), phi] souch that particles pick up a phase phi when hopping around the cylinder.

Returns strength - The strength array to be used as strength in add_coupling() with the given \(d x\).

Return type complex array

\section*{Examples}

Let's say you have an infinite MPS on a cylinder, and want to add nearest-neighbor hopping of fermions with the FermionSite. The cylinder axis is the \(x\)-direction of the lattice, so to put a flux through the cylinder, you want particles hopping around the cylinder to pick up a phase phi given by the external flux.
```

>>> strength = 1. \# hopping strength without external flux
>>> phi = np.pi/4 \# determines the external flux strength
>>> for u1, u2, dx in self.lat.pairs['nearest_neighbors']:
... strength_with_flux = self.coupling_strength_add_ext_flux(strength, dx,
@ [0, phi])
... self.add_coupling(strength_with_flux, u1, 'Cd', u2, 'C', dx)
... self.add_coupling(np.conj(strength_with_flux), u2, 'cd', u1, 'C', -dx)

```
enlarge_mps_unit_cell (factor=2)
Repeat the unit cell for infinite MPS boundary conditions; in place.
This has to be done after finishing initialization and can not be reverted.
Parameters factor (int) - The new number of sites in the MPS unit cell will be increased from \(N_{\text {_s }}\) sites to factor*N_sites_per_ring. Since MPS unit cells are repeated in the \(x\)-direction in our convetion, the lattice shape goes from ( \(\mathrm{Lx}, \mathrm{Ly}, \ldots, \mathrm{Lu}\) ) to (Lx*factor, Ly, ..., Lu).
classmethod from_hdf5 (hdf5_loader, h5gr, subpath)
Load instance from a HDF5 file.
This method reconstructs a class instance from the data saved with save_hdf5().

\section*{Parameters}
- hdf5_loader (Hdf5Loader) - Instance of the loading engine.
- h5gr (Group) - HDF5 group which is represent the object to be constructed.
- subpath (str) - The name of \(h 5 g r\) with a '/' in the end.

Returns obj - Newly generated class instance containing the required data.

\section*{Return type cls}
group_sites ( \(n=2\), grouped_sites=None)
Modify self in place to group sites.
Group each \(n\) sites together using the GroupedSite. This might allow to do TEBD with a Trotter decomposition, or help the convergence of DMRG (in case of too long range interactions).
This has to be done after finishing initialization and can not be reverted.

\section*{Parameters}
- \(\mathbf{n}(i n t)\) - Number of sites to be grouped together.
- grouped_sites (None I list of GroupedSite) - The sites grouped together.

Returns grouped_sites - The sites grouped together.
Return type list of GroupedSite
save_hdf5 (hdf5_saver, h5gr, subpath)
Export self into a HDF5 file.
This method saves all the data it needs to reconstruct self with from_hdf5 ().
This implementation saves the content of \(\qquad\) dict \(\qquad\) with save_dict_content (), storing the format under the attribute 'format'.

\section*{Parameters}
- hdf5_saver (Hdf5Saver) - Instance of the saving engine.
- h5gr (: class `Group \(\left.{ }^{\prime}\right)\) - HDF5 group which is supposed to represent self.
- subpath (str) - The name of \(h 5 g r\) with a ' / ' in the end.

\subsection*{19.2.2 MPOModel}
- full name: tenpy.models.model.MPOModel
- parent module: tenpy.models.model
- type: class

\section*{Inheritance Diagram}


\section*{Methods}
\begin{tabular}{ll}
\hline MPOModel.__init__(lattice, H_MPO) & Initialize self. \\
\hline MPOModel.calc_H_bond_from_MPO([tol_zero]) & \begin{tabular}{l} 
Calculate the bond Hamiltonian from the MPO Hamil- \\
tonian.
\end{tabular} \\
\hline MPOModel.enlarge_mps_unit_cell([factor]) & \begin{tabular}{l} 
Repeat the unit cell for infinite MPS boundary condi- \\
tions; in place.
\end{tabular} \\
\hline MPOModel.from_hdf5(hdf5_loader, h5gr, subpath) & Load instance from a HDF5 file. \\
\hline MPOModel.group_sites([n, grouped_sites]) & Modify self in place to group sites. \\
\hline MPOModel.save_hdf5(hdf5_saver, h5gr, subpath) & Export self into a HDF5 file. \\
\hline MPOModel.test_sanity () &
\end{tabular}

\section*{Class Attributes and Properties}
MPOModel.logger class attribute.
class tenpy.models.model. MPOModel (lattice, \(H_{-} M P O\) )
Bases: tenpy.models.model.Model
Base class for a model with an MPO representation of the Hamiltonian.
In this class, the Hamiltonian gets represented by an MPO. Thus, instances of this class are suitable for MPObased algorithms like DMRG dmrg and MPO time evolution.

Parameters H_MPO (MPO) - The Hamiltonian rewritten as an MPO.
H_MPO
MPO representation of the Hamiltonian. If the explicit_plus_hc flag of the MPO is True, the represented Hamiltonian is H_MPO + hermitian_cojugate (H_MPO).

Type tenpy.networks.mpo.MPO
enlarge_mps_unit_cell (factor=2)
Repeat the unit cell for infinite MPS boundary conditions; in place.
This has to be done after finishing initialization and can not be reverted.
Parameters factor (int) - The new number of sites in the MPS unit cell will be increased from \(N\) _sites to factor*N_sites_per_ring. Since MPS unit cells are repeated in the \(x\)-direction in our convetion, the lattice shape goes from ( \(\mathrm{Lx}, \mathrm{Ly}, \ldots, \mathrm{Lu}\) ) to (Lx*factor, Ly, ..., Lu).
group_sites ( \(n=2\), grouped_sites \(=\) None)
Modify self in place to group sites.
Group each \(n\) sites together using the GroupedSite. This might allow to do TEBD with a Trotter decomposition, or help the convergence of DMRG (in case of too long range interactions).

This has to be done after finishing initialization and can not be reverted.

\section*{Parameters}
- \(\mathbf{n}(i n t)\) - Number of sites to be grouped together.
- grouped_sites (None l list of GroupedSite) - The sites grouped together.

Returns grouped_sites - The sites grouped together.

Return type list of GroupedSite
calc_H_bond_from_MPO (tol_zero=le-15)
Calculate the bond Hamiltonian from the MPO Hamiltonian.
Parameters tol_zero (float) - Arrays with norm <tol_zero are considered to be zero.
Returns H_bond - Bond terms as required by the constructor of NearestNeighborModel. Legs are ['p0', 'p0*', 'p1', 'p1*']

Return type list of Array
:raises ValueError : if the Hamiltonian contains longer-range terms.:
classmethod from_hdf5 (hdf5_loader, h5gr, subpath)
Load instance from a HDF5 file.
This method reconstructs a class instance from the data saved with save_hdf5 ().

\section*{Parameters}
- hdf5_loader (Hdf5Loader) - Instance of the loading engine.
- h5gr (Group) - HDF5 group which is represent the object to be constructed.
- subpath (str) - The name of \(h 5 g r\) with a '/' in the end.

Returns obj - Newly generated class instance containing the required data.
Return type cls
save_hdf5 (hdf5_saver, h5gr, subpath)
Export self into a HDF5 file.
This method saves all the data it needs to reconstruct self with from_hdf5 ().
This implementation saves the content of __ dict__ with save_dict_content (), storing the format under the attribute 'format'.

\section*{Parameters}
- hdf5_saver (Hdf5Saver) - Instance of the saving engine.
- h5gr (: class `Group`) - HDF5 group which is supposed to represent self.
- subpath (str) - The name of \(h 5 g r\) with a '/' in the end.

\subsection*{19.2.3 Model}
- full name: tenpy.models.model.Model
- parent module: tenpy.models.model
- type: class

\section*{Inheritance Diagram}


\section*{Methods}
\begin{tabular}{ll}
\hline Model.__init__(lattice) & Initialize self. \\
\hline Model.enlarge_mps_unit_cell([factor]) & \begin{tabular}{l} 
Repeat the unit cell for infinite MPS boundary condi- \\
tions; in place.
\end{tabular} \\
\hline Model.from_hdf5(hdf5_loader, h5gr, subpath) & Load instance from a HDF5 file. \\
\hline Model.group_sites([n, grouped_sites]) & Modify self in place to group sites. \\
\hline Model.save_hdf5(hdf5_saver, h5gr, subpath) & Export self into a HDF5 file. \\
\hline
\end{tabular}

\section*{Class Attributes and Properties}
Model.logger class attribute.
class tenpy.models.model.Model (lattice)
Bases: tenpy.tools.hdf5_io.Hdf5Exportable
Base class for all models.
The common base to all models is the underlying Hilbert space and geometry, specified by a Lattice.
Parameters lattice (Lattice) - The lattice defining the geometry and the local Hilbert space(s).
lat
The lattice defining the geometry and the local Hilbert space(s).
Type Lattice
dtype
The data type of the Hamiltonian
Type dtype
logger \(=\) <Logger tenpy.models.model.Model (WARNING)> class attribute.

Type logging.Logger
Type An instance of a logger; see Logging and terminal output. NB
enlarge_mps_unit_cell (factor=2)
Repeat the unit cell for infinite MPS boundary conditions; in place.
This has to be done after finishing initialization and can not be reverted.
Parameters factor (int) - The new number of sites in the MPS unit cell will be increased from \(N\) _sites to factor*N_sites_per_ring. Since MPS unit cells are repeated in the \(x\)-direction in our convetion, the lattice shape goes from ( \(\mathrm{Lx}, \mathrm{Ly}, \ldots, \mathrm{Lu}\) ) to (Lx*factor, Ly, ..., Lu).
group_sites ( \(n=2\), grouped_sites \(=\) None)
Modify self in place to group sites.
Group each \(n\) sites together using the GroupedSite. This might allow to do TEBD with a Trotter decomposition, or help the convergence of DMRG (in case of too long range interactions).
This has to be done after finishing initialization and can not be reverted.

\section*{Parameters}
- \(\mathbf{n}(i n t)\) - Number of sites to be grouped together.
- grouped_sites (None l list of GroupedSite) - The sites grouped together.

Returns grouped_sites - The sites grouped together.
Return type list of GroupedSite
classmethod from_hdf5 (hdf5_loader, h5gr, subpath)
Load instance from a HDF5 file.
This method reconstructs a class instance from the data saved with save_hdf5 ().

\section*{Parameters}
- hdf5_loader (Hdf5Loader) - Instance of the loading engine.
- h5gr (Group) - HDF5 group which is represent the object to be constructed.
- subpath (str) - The name of \(h 5 g r\) with a '/' in the end.

Returns obj - Newly generated class instance containing the required data.
Return type cls
save_hdf5 (hdf5_saver, h5gr, subpath)
Export self into a HDF5 file.
This method saves all the data it needs to reconstruct self with \(f r o m \_h d f 5\) ().
This implementation saves the content of __dict__ with save_dict_content (), storing the format under the attribute 'format'.

\section*{Parameters}
- hdf5_saver (Hdf5Saver) - Instance of the saving engine.
- h5gr (: class `Group`) - HDF5 group which is supposed to represent self.
- subpath (str) - The name of \(h 5 g r\) with a '/' in the end.

\subsection*{19.2.4 MultiCouplingModel}
- full name: tenpy.models.model.MultiCouplingModel
- parent module: tenpy.models.model
- type: class

\section*{Inheritance Diagram}


\section*{Methods}
\begin{tabular}{|c|c|}
\hline MultiCouplingModel.__init__(lattice[,...]) & Initialize self. \\
\hline MultiCouplingModel. add_coupling(strength, ...) & Add twosite coupling terms to the Hamiltonian, summing over lattice sites. \\
\hline \begin{tabular}{l}
MultiCouplingModel. \\
add_coupling_term(...[,...])
\end{tabular} & Add a two-site coupling term on given MPS sites. \\
\hline \begin{tabular}{l}
MultiCouplingModel. \\
add_exponentially_decaying_coupling(...)
\end{tabular} & Add an exponentially decaying long-range coupling. \\
\hline \begin{tabular}{l}
MultiCouplingModel. \\
add_local_term(strength, term)
\end{tabular} & Add a single term to self. \\
\hline \begin{tabular}{l}
MultiCouplingModel. \\
add_multi_coupling(...[,...])
\end{tabular} & Add multi-site coupling terms to the Hamiltonian, summing over lattice sites. \\
\hline \begin{tabular}{l}
MultiCouplingModel. \\
add_multi_coupling_term(...)
\end{tabular} & Add a general M-site coupling term on given MPS sites. \\
\hline MultiCouplingModel.add_onsite(strength, u, ...) & Add onsite terms to onsite_terms. \\
\hline
\end{tabular}

Table 33 - continued from previous page
\begin{tabular}{|c|c|}
\hline MultiCouplingModel. add_onsite_term(strength, ...) & Add an onsite term on a given MPS site. \\
\hline MultiCouplingModel. all_coupling_terms() & Sum of all coupling_terms. \\
\hline MultiCouplingModel.all_onsite_terms() & Sum of all onsite_terms. \\
\hline MultiCouplingModel.calc_H_MPO([tol_zero]) & Calculate MPO representation of the Hamiltonian. \\
\hline MultiCouplingModel. calc_H_bond([tol_zero]) & calculate H_bond from coupling_terms and onsite_terms. \\
\hline MultiCouplingModel. calc_H_onsite([tol_zero]) & Calculate H_onsite from self.onsite_terms. \\
\hline MultiCouplingModel. coupling_strength_add_ext_flux(...) & Add an external flux to the coupling strength. \\
\hline \begin{tabular}{l}
MultiCouplingModel. \\
enlarge_mps_unit_cell([...])
\end{tabular} & Repeat the unit cell for infinite MPS boundary conditions; in place. \\
\hline MultiCouplingModel.from_hdf5(hdf5_loader, ...) & Load instance from a HDF5 file. \\
\hline MultiCouplingModel.group_sites([n,...]) & Modify self in place to group sites. \\
\hline MultiCouplingModel.save_hdf5(hdf5_saver, ...) & Export self into a HDF5 file. \\
\hline MultiCouplingModel.test_sanity() & Sanity check, raises ValueErrors, if something is wrong. \\
\hline
\end{tabular}

\section*{Class Attributes and Properties}
\begin{tabular}{l} 
MultiCouplingModel.logger class attribute. \\
\hline \\
class tenpy.models.model.MultiCouplingModel(lattice, \(\quad\) plicit_plus_hc=False)
\end{tabular}

Bases: tenpy.models.model. CouplingModel
Deprecated class which was a generalization of the CouplingModel.
Deprecated since version 0.7.2: In earlier versions of TeNPy, this class contained the methods add_multi_coupling() and add_multi_coupling_term(). However, since we introduced the MultiCouplingTerms, this separation within the Model class is no longer necessary. We hence merged the MultiCouplingModel with the CouplingModel.
add_coupling \(\left(\right.\) strength, \(\quad u 1, \quad o p 1, \quad u 2, \quad o p 2, \quad d x, \quad o p \_\)string \(=N o n e, \quad\) str_on_first=True, raise_op2_left=False, category=None, plus_hc=False)
Add twosite coupling terms to the Hamiltonian, summing over lattice sites.
Represents couplings of the form \(\sum_{x_{0}, \ldots, x_{\text {dim-1 }}} \operatorname{strength}[\operatorname{shift}(\vec{x})] * O P 0 * O P 1\), where OP 0 : = lat.unit_cell[u0].get_op (op0) acts on the site (x_0, ..., x_\{dim-1\}, u1), and OP1 := lat.unit_cell[u1].get_op(op1) acts on the site (x_0+dx[0], ..., \(x_{-}\{\operatorname{dim}-1\}+d x[\operatorname{dim}-1]\), \(\left.u 1\right)\). Possible combinations \(x_{-} 0, \ldots, x_{-}\{d i m-1\}\) are determined from the boundary conditions in possible_couplings().

The coupling strength may vary spatially if the given strength is a numpy array. The correct shape of this array is the coupling_shape returned by tenpy.models.lattice.coupling_shape () and depends on the boundary conditions. The shift (...) depends on \(d x\), and is chosen such that the first entry strength [0, 0, ...] of strength is the prefactor for the first possible coupling fitting into the lattice if you imagine open boundary conditions.

The necessary terms are just added to coupling_terms; this function does not rebuild the MPO.

Deprecated since version 0.4.0: The arguments str_on_first and raise_op2_left will be removed in version 1.0.0.

\section*{Parameters}
- strength (scalar / array) - Prefactor of the coupling. May vary spatially (see above). If an array of smaller size is provided, it gets tiled to the required shape.
- u1 (int) - Picks the site lat. unit_cell[u1] for OP1.
- op1 (str) - Valid operator name of an onsite operator in lat.unit_cell[u1] for OP1.
- u2 (int) - Picks the site lat.unit_cell[u2] for OP2.
- op2 (str) - Valid operator name of an onsite operator in lat.unit_cell[u2] for OP2.
- \(\mathbf{d x}\) (iterable of int) - Translation vector (of the unit cell) between OP1 and OP2. For a 1D lattice, a single int is also fine.
- op_string (str / None) - Name of an operator to be used between the OP1 and OP2 sites. Typical use case is the phase for a Jordan-Wigner transformation. The operator should be defined on all sites in the unit cell. If None, auto-determine whether a JordanWigner string is needed, using op_needs_JW ().
- str_on_first (bool) - Whether the provided op_string should also act on the first site. This option should be chosen as True for Jordan-Wigner strings. When handling Jordan-Wigner strings we need to extend the op_string to also act on the 'left', first site (in the sense of the MPS ordering of the sites given by the lattice). In this case, there is a well-defined ordering of the operators in the physical sense (i.e. which of opl or op2 acts first on a given state). We follow the convention that op2 acts first (in the physical sense), independent of the MPS ordering. Deprecated.
- raise_op2_left (bool) - Raise an error when op2 appears left of opl (in the sense of the MPS ordering given by the lattice). Deprecated.
- category (str) - Descriptive name used as key for coupling_terms. Defaults to a string of the form " \(\{o p 1\} \_i \quad\{o p 2\} \_j "\).
- plus_hc (bool) - If True, the hermitian conjugate of the terms is added automatically.

\section*{Examples}

When initializing a model, you can add a term \(J \sum_{<i, j>} S_{i}^{z} S_{j}^{z}\) on all nearest-neighbor bonds of the lattice like this:
```

>>> J = 1. \# the strength
>>> for u1, u2, dx in self.lat.pairs['nearest_neighbors']:
... self.add_coupling(J, u1, 'Sz', u2, 'Sz', dx)

```

The strength can be an array, which gets tiled to the correct shape. For example, in a 1D Chain with an even number of sites and periodic (or infinite) boundary conditions, you can add alternating strong and weak couplings with a line like:
```

>>> self.add_coupling([1.5, 1.], u1, 'Sz', u2, 'Sz', dx)

```

Make sure to use the plus_hc argument if necessary, e.g. for hoppings:
```

>>> t = 1. \# hopping strength
>>> for u1, u2, dx in self.lat.pairs['nearest_neighbors']:
... self.add_coupling(t, u1, 'Cd', u2, 'C', dx, plus_hc=True)

```

Alternatively, you can add the hermitian conjugate terms explictly. The correct way is to complex conjugate the strength, take the hermitian conjugate of the operators and swap the order (including a swap \(u 1<->u 2\) ), and use the opposite direction \(-d x\), i.e. the h.c. of add_coupling ( \(t, u 1\), 'A', u2, 'B', dx) is add_coupling(np.conj(t), u2, hc('B'), u1, hc('A'), -dx), where \(h c\) takes the hermitian conjugate of the operator names, see get_hc_op_name (). For spin-less fermions (FermionSite), this would be
```

>>> for u1, u2, dx in self.lat.pairs['nearest_neighbors']:
... self.add_coupling(t, u1, 'Cd', u2, 'C', dx)
... self.add_coupling(np.conj(t), u2, 'Cd', u1, 'C', -dx) \# h.c.

```

With spin-full fermions (SpinHalfFermions), it could be:
```

>>> for u1, u2, dx in self.lat.pairs['nearest_neighbors']:
... self.add_coupling(t, u1, 'Cdu', u2, 'Cd', dx) \# Cdagger_up C_down
... self.add_coupling(np.conj(t), u2, 'Cdd', u1, 'Cu', -dx) \# h.c.u
\hookrightarrowCdagger_down C_up

```

Note that the Jordan-Wigner strings for the fermions are added automatically!

\section*{See also:}
add_onsite Add terms acting on one site only.
add_multi_coupling_term for terms on more than two sites.
add_coupling_term Add a single term without summing over \(\vec{x}\).
add_coupling_term (strength, \(i, j, o p_{-} i, o p_{-} j, o p_{-}\)string \(=' I d^{\prime}\), category=None, plus_hc=False)
Add a two-site coupling term on given MPS sites.
Wrapper for self.coupling_terms[category].add_coupling_term(...).

Warning: This function does not handle Jordan-Wigner strings! You might want to use add_local_term() instead.

\section*{Parameters}
- strength (float) - The strength of the coupling term.
- i (int) - The MPS indices of the two sites on which the operator acts. We require 0 \(<=i<N \_s i t e s\) and \(i<j\), i.e., op_i acts "left" of \(o p_{-} j\). If \(\mathrm{j}>=\mathrm{N}_{-}\)sites, it indicates couplings between unit cells of an infinite MPS.
- j \((i n t)\) - The MPS indices of the two sites on which the operator acts. We require 0 \(<=i<N_{\_} s i t e s\) and \(i<j\), i.e., op_i acts "left" of \(o p_{-} j\). If \(j>=N_{-}\)sites, it indicates couplings between unit cells of an infinite MPS.
- op1 (str) - Names of the involved operators.
- op2 (str) - Names of the involved operators.
- op_string (str) - The operator to be inserted between \(i\) and \(j\).
- category (str) - Descriptive name used as key for coupling_terms. Defaults to a string of the form " \(\{0 \mathrm{p} 1\} \_i \quad\{o p 2\} \_j "\).
- plus_hc (bool) - If True, the hermitian conjugate of the term is added automatically.
add_exponentially_decaying_coupling (strength, lambda_, op_i,op_j, subsites=None, \(o p_{-}\)string \(=\)None, \(p l u s \_h c=\) False \()\)
Add an exponentially decaying long-range coupling.
\[
\text { strength } \sum_{i<j} \lambda^{|i-j|} A_{\text {subsites }[i]} B_{\text {subsites }[j]}
\]

Where the operator \(A\) is given by \(o p_{-} i\), and \(B\) is given by \(o p_{-} j\). Note that the sum over \(i, j\) is long-range, for infinite systems going beyond the MPS unit cell. Moreover, note that the distance in the exponent is the distance within subsites.

\section*{Parameters}
- strength (float) - Overall prefactor.
- lambda (float) - Decay-rate
- op_i (string) - Names for the operators.
- op_j (string) - Names for the operators.
- subsites (None / 1D array) - Selects a subset of sites within the MPS unit cell on which the operators act. Needs to be sorted. None selects all sites.
- op_string (None / str) - The operator to be inserted between \(A\) and \(B\); If None, this function checks whether a fermionic "JW" string is needed for the given operators; in this case the right \(o p_{-} j\) acts first.
- plus_hc (bool) - If True, the hermitian conjugate of the term is added automatically.

\section*{Examples}

At least for simple enough 1D chains (or ladders), you can use fit_with_sum_of_exp () to approximate a long-range function with a few sum of exponentials and then add them with this function.
```

>>> def decay(x):
return np.exp(-0.1*x) / x**2
>>> from tenpy.tools.fit import fit_with_sum_of_exp, sum_of_exp
>>> n_exp = 5
>>> fit_range = 50
>>> lam, pref = fit_with_sum_of_exp(decay, n_exp, fit_range)
>>> x = np.arange(1, fit_range + 1)
>>> print('error in fit: {0:.3e}'.format(np.sum(np.abs(decay(x) - sum_of_
@exp(lam, pref, x)))))
error in fit: 1.073e-04
>>> for pr, la in zip(pref, lam):
self.add_exponentially_decaying_coupling(pr, la, 'N', 'N')

```
add_local_term (strength, term, category=None, plus_hc=False)
Add a single term to self.
The repesented term is strength times the product of the operators given in terms. Each operator is specified by the name and the site it acts on; the latter given by a lattice index, see Lattice.

Depending on the length of term, it can add an onsite term or a coupling term to onsite_terms or coupling_terms, respectively.

\section*{Parameters}
- strength (float/complex) - The prefactor of the term.
- term (list of (str, array_like)) - List of tuples (opname, lat_idx) where opname is a string describing the operator acting on the site given by the lattice index lat_idx. Here, lat_idx is for example \([x, y, u]\) for a 2D lattice, with \(u\) being the index within the unit cell.
- category - Descriptive name used as key for onsite_terms or coupling_terms.
- plus_hc (bool) - If True, the hermitian conjugate of the terms is added automatically.
add_multi_coupling (strength, ops, _deprecate_l='DEPRECATED', _depre-
cate_2='DEPRECATED', op_string=None, category=None, plus_hc=False)
Add multi-site coupling terms to the Hamiltonian, summing over lattice sites.
Represents couplings of the form \(\operatorname{sum}_{\vec{x}} \operatorname{strength}[\operatorname{shift}(\vec{x})] * O P_{0} * O P_{1} * \ldots * O P_{M-1}\), involving \(M\) operators. Here, \(O P_{m}\) stands for the operator defined by the \(m\)-th tuple (opname, dx, u) given in the argument ops, which determines the position \(\vec{x}+\overrightarrow{d x}\) and unit-cell index \(u\) of the site it acts on; the actual operator is given by self.lat.unit_cell[u].get_op(opname).

The coupling strength may vary spatially if the given strength is a numpy array. The correct shape of this array is the coupling_shape returned by tenpy.models.lattice. possible_multi_couplings() and depends on the boundary conditions. The shift(...) depends on the \(d x\) entries of ops and is chosen such that the first entry st rength \([0,0, \ldots]\) of strength is the prefactor for the first possible coupling fitting into the lattice if you imagine open boundary conditions.

The necessary terms are just added to coupling_terms; this function does not rebuild the MPO.
Deprecated since version 0.6.0: We switched from the three arguments \(u 0\), op0 and other_op with other_ops=[(u1, op1, dx1), (op2, u2, dx2), ...] to a single, equivalent argment ops which should now read ops=[(op0, dx0, u0), (op1, dx1, u1), (op2, dx2, u2), . ..], where \(d x 0=[0] *\) self.lat.dim. Note the changed order inside the tuples!

\section*{Parameters}
- strength (scalar | array) - Prefactor of the coupling. May vary spatially, and is tiled to the required shape.
- ops (list of (opname, dx, u)) - Each tuple determines one operator of the coupling, see the description above. opname (str) is the name of the operator, \(d x\) (list of length lat.dim) is a translation vector, and \(u\) (int) is the index of lat.unit_cell on which the operator acts. The first entry of ops corresponds to \(O P_{0}\) and acts last in the physical sense.
- op_string (str | None) - If a string is given, we use this as the name of an operator to be used inbetween the operators, excluding the sites on which any operators act. This operator should be defined on all sites in the unit cell.
If None, auto-determine whether a Jordan-Wigner string is needed (using op_needs_JW ()) for each of the segments inbetween the operators and also on the sites of the left operators.
- category (str) - Descriptive name used as key for coupling_terms. Defaults to a string of the form " \(\{0 \mathrm{p} 0\}\) _i \(\{\) other_ops [0]\}_j \{other_ops [1]\}_k ...".
- plus_hc (bool) - If True, the hermitian conjugate of the terms is added automatically.

\section*{Examples}

A call to add_coupling() with arguments add_coupling(strength, u1, 'A', u2, ' \(\mathrm{B}^{\prime}, \mathrm{dx}\) ) is equivalent to the following:
```

>>> dx_0 = [0] * self.lat.dim \# = [0] for a ID lattice, [0, 0] in 2D
>>> self.add_multi_coupling(strength, [('A', dx_0, ul), ('B', dx, u2)])

```

To explicitly add the hermitian conjugate (instead of simply using plus_hc = True), you need to take the complex conjugate of the strength, reverse the order of the operators and take the hermitian conjugates of the individual operator names (indicated by the hc (...), see get_hc_op_name ()):
```

>>> self.add_multi_coupling(np.conj(strength), [(hc('B'), dx, u2), (hc('A'), -
\hookrightarrowdx_0, u1)])

```

\section*{See also:}
add_onsite Add terms acting on one site only.
add_coupling Add terms acting on two sites.
add_multi_coupling_term Add a single term, not summing over the possible \(\vec{x}\).
add_multi_coupling_term (strength, ijkl,ops_ijkl,op_string, category=None, plus_hc=False)
Add a general M-site coupling term on given MPS sites.
Wrapper for self.coupling_terms[category].add_multi_coupling_term(...).

Warning: This function does not handle Jordan-Wigner strings! You might want to use add_local_term() instead.

\section*{Parameters}
- strength (float) - The strength of the coupling term.
- ijkl (list of int)- The MPS indices of the sites on which the operators acts. With \(i, j, k, \ldots=i j k l\), we require that they are ordered ascending, \(i<j<k<\ldots\) and that \(0<=i<N \_s i t e s\). Inidces \(>=\) N_sites indicate couplings between different unit cells of an infinite MPS.
- ops_ijkl(list of str) - Names of the involved operators on sites \(i, j, k, \ldots\).
- op_string (list of str) - Names of the operator to be inserted between the operators, e.g., op_string[0] is inserted between \(i\) and \(j\).
- category (str) - Descriptive name used as key for coupling_terms. Defaults to a string of the form " \(\{\mathrm{op} 0\} \_i \quad\{o p 1\} \_j \quad\{o p 2\} \_k \quad .\). ".
- plus_hc (bool) - If True, the hermitian conjugate of the term is added automatically.
add_onsite (strength, u, opname, category=None, plus_hc=False)
Add onsite terms to onsite_terms.
Adds \(\sum_{\vec{x}} \operatorname{strength}[\vec{x}] * O P\) to the represented Hamiltonian, where the operator \(\mathrm{OP}=1\) at. unit_cell[u].get_op (opname) acts on the site given by a lattice index (x_0, ..., \(\left.x_{-}\{\operatorname{dim}-1\}, u\right)\),

The necessary terms are just added to onsite_terms; doesn't rebuild the MPO.

\section*{Parameters}
- strength (scalar / array) - Prefactor of the onsite term. May vary spatially. If an array of smaller size is provided, it gets tiled to the required shape.
- u(int) - Picks a Site lat.unit_cell [u] out of the unit cell.
- opname (str) - valid operator name of an onsite operator in lat. unit_cell[u].
- category (str) - Descriptive name used as key for onsite_terms. Defaults to opname.
- plus_hc (bool) - If True, the hermitian conjugate of the terms is added automatically.

\section*{See also:}
add_coupling Add a terms acting on two sites.
add_onsite_term Add a single term without summing over vecx.
add_onsite_term (strength, i, op, category=None, plus_hc=False)
Add an onsite term on a given MPS site.
Wrapper for self.onsite_terms[category].add_onsite_term(...).

\section*{Parameters}
- strength (float) - The strength of the term.
- \(\mathbf{i}(i n t)\) - The MPS index of the site on which the operator acts. We require \(0<=i<\) L.
- op (str) - Name of the involved operator.
- category (str) - Descriptive name used as key for onsite_terms. Defaults to op.
- plus_hc (bool) - If True, the hermitian conjugate of the term is added automatically.
all_coupling_terms()
Sum of all coupling_terms.
all_onsite_terms()
Sum of all onsite_terms.
calc_H_MPO (tol_zero \(=1 e-15\) )
Calculate MPO representation of the Hamiltonian.
Uses onsite_terms and coupling_terms to build an MPOGraph (and then an MPO).
Parameters tol_zero (float) - Prefactors with abs(strength) < tol_zero are considered to be zero.

Returns H_MPO - MPO representation of the Hamiltonian.
Return type MPO
calc_H_bond (tol_zero=le-15)
calculate \(H\) _bond from coupling_terms and onsite_terms.
Parameters tol_zero (float) - prefactors with abs (strength) < tol_zero are considered to be zero.

Returns H_bond - Bond terms as required by the constructor of NearestNeighborModel. Legs are ['p0', 'p0*', 'p1', 'p1*']
Return type list of Array
:raises ValueError : if the Hamiltonian contains longer-range terms.:
```

calc_H_onsite(tol_zero=le-15)

```

Calculate H_onsite from self.onsite_terms.
Deprecated since version 0.4.0: This function will be removed in 1.0.0. Replace calls to this function by self.all_onsite_terms().remove_zeros(tol_zero).to_Arrays (self. lat.mps_sites()). You might also want to take explicit_plus_hc into account.

Parameters tol_zero (float) - prefactors with abs (strength) < tol_zero are considered to be zero.

\section*{Returns}
- H_onsite (list of npc.Array)
- onsite terms of the Hamiltonian. If explicit_plus_hc is True, - Hermitian conjugates of the onsite terms will be included.
coupling_strength_add_ext_flux (strength, \(d x\), phase)
Add an external flux to the coupling strength.
When performing DMRG on a "cylinder" geometry, it might be useful to put an "external flux" through the cylinder. This means that a particle hopping around the cylinder should pick up a phase given by the external flux [[resta1998]]. This is also called "twisted boundary conditions" in literature. This function adds a complex phase to the strength array on some bonds, such that particles hopping in positive direction around the cylinder pick up \(\exp (+i\) phase \()\).

Warning: For the sign of phase it is important that you consistently use the creation operator as opl and the annihilation operator as op2 in add_coupling().

\section*{Parameters}
- strength (scalar / array) - The strength to be used in add_coupling(), when no external flux would be present.
- \(\mathbf{d x}\) (iterable of int) - Translation vector (of the unit cell) between opl and op2 in add_coupling().
- phase (iterable of float) - The phase of the external flux for hopping in each direction of the lattice. E.g., if you want flux through the cylinder on which you have an infinite MPS, you should give phase \(=[0\), phi] souch that particles pick up a phase phi when hopping around the cylinder.
Returns strength - The strength array to be used as strength in add_coupling() with the given \(d x\).
Return type complex array

\section*{Examples}

Let's say you have an infinite MPS on a cylinder, and want to add nearest-neighbor hopping of fermions with the FermionSite. The cylinder axis is the \(x\)-direction of the lattice, so to put a flux through the cylinder, you want particles hopping around the cylinder to pick up a phase phi given by the external flux.
```

>>> strength = 1. \# hopping strength without external flux
>>> phi = np.pi/4 \# determines the external flux strength
>> for ul, u2, dx in self.lat.pairs['nearest_neighbors']:
... strength_with_flux = self.coupling_strength__add_ext_flux(strength, dx,
\hookrightarrow[0, phi])

```
(continued from previous page)
... self.add_coupling(strength_with_flux, u1, 'Cd', u2, 'C', dx) self.add_coupling(np.conj(strength_with_flux), u2, 'Cd', u1, 'C', -dx)
enlarge_mps_unit_cell(factor=2)
Repeat the unit cell for infinite MPS boundary conditions; in place.
This has to be done after finishing initialization and can not be reverted.
Parameters factor (int) - The new number of sites in the MPS unit cell will be increased from \(N\) _sites to factor*N_sites_per_ring. Since MPS unit cells are repeated in the \(x\)-direction in our convetion, the lattice shape goes from ( \(L x, L y, \ldots, L u\) ) to (Lx*factor, Ly, ..., Lu).
classmethod from_hdf5 (hdf5_loader, h5gr, subpath)
Load instance from a HDF5 file.
This method reconstructs a class instance from the data saved with save_hdf5 ().

\section*{Parameters}
- hdf5_loader (Hdf5Loader) - Instance of the loading engine.
- h5gr (Group) - HDF5 group which is represent the object to be constructed.
- subpath (str) - The name of \(h 5 g r\) with a '/' in the end.

Returns obj - Newly generated class instance containing the required data.
Return type cls
group_sites ( \(n=2\), grouped_sites \(=\) None)
Modify self in place to group sites.
Group each \(n\) sites together using the GroupedSite. This might allow to do TEBD with a Trotter decomposition, or help the convergence of DMRG (in case of too long range interactions).
This has to be done after finishing initialization and can not be reverted.

\section*{Parameters}
- \(\mathrm{n}(\mathrm{int})\) - Number of sites to be grouped together.
- grouped_sites (None l list of GroupedSite) - The sites grouped together.

Returns grouped_sites - The sites grouped together.
Return type list of GroupedSite
save_hdf5 (hdf5_saver, h5gr, subpath)
Export self into a HDF5 file.
This method saves all the data it needs to reconstruct self with from_hdf5 ().
This implementation saves the content of \(\qquad\) dict \(\qquad\) with save_dict_content (), storing the format under the attribute 'format'.

\section*{Parameters}
- hdf5_saver (Hdf5Saver) - Instance of the saving engine.
- h5gr (: class `Group`) - HDF5 group which is supposed to represent self.
- subpath (str) - The name of \(h 5 g r\) with a ' /' in the end.
```

test_sanity()

```

Sanity check, raises ValueErrors, if something is wrong.

\subsection*{19.2.5 NearestNeighborModel}
- full name: tenpy.models.model.NearestNeighborModel
- parent module: tenpy.models.model
- type: class

\section*{Inheritance Diagram}


\section*{Methods}
\begin{tabular}{|c|c|}
\hline NearestNeighborModel.__init__(lattice, H_bond) & Initialize self. \\
\hline \begin{tabular}{l}
NearestNeighborModel. \\
bond_energies(psi)
\end{tabular} & Calculate bond energies <psilH_bondlpsi>. \\
\hline \begin{tabular}{l}
NearestNeighborModel. \\
calc_H_MPO_from_bond([...])
\end{tabular} & Calculate the MPO Hamiltonian from the bond Hamiltonian. \\
\hline \begin{tabular}{l}
NearestNeighborModel. \\
enlarge_mps_unit_cell([...])
\end{tabular} & Repeat the unit cell for infinite MPS boundary conditions; in place. \\
\hline NearestNeighborModel. from_MPOMOdel(mpo_model) & Initialize a NearestNeighborModel from a model class defining an MPO. \\
\hline \begin{tabular}{l}
NearestNeighborModel. \\
from_hdf5(hdf5_loader, ...)
\end{tabular} & Load instance from a HDF5 file. \\
\hline NearestNeighborModel.group_sites([n, ...]) & Modify self in place to group sites. \\
\hline NearestNeighborModel. save_hdf5(hdf5_saver, ...) & Export self into a HDF5 file. \\
\hline
\end{tabular}

Table 35 - continued from previous page
NearestNeighborModel.test_sanity()
NearestNeighborModel.
Return a NearestNeighborModel with same lattice, but trivial_like_NNModel() trivial \((\mathrm{H}=0)\) bonds.

\section*{Class Attributes and Properties}
NearestNeighborModel.logger class attribute.
class tenpy.models.model. NearestNeighborModel (lattice, \(H\) _bond)
Bases: tenpy.models.model.Model
Base class for a model of nearest neigbor interactions w.r.t. the MPS index.
In this class, the Hamiltonian \(H=\sum_{i} H_{i, i+1}\) is represented by "bond terms" \(H_{i, i+1}\) acting only on two neighboring sites \(i\) and \(i+1\), where \(i\) is an integer. Instances of this class are suitable for tebd.

Note that the "nearest-neighbor" in the name refers to the MPS index, not the lattice. In short, this works only for 1-dimensional (1D) nearest-neighbor models: A 2D lattice is internally mapped to a 1D MPS "snake", and even a nearest-neighbor coupling in 2D becomes long-range in the MPS chain.

\section*{Parameters}
- lattice (tenpy.model.lattice.Lattice) - The lattice defining the geometry and the local Hilbert space(s).
- H_bond (list of \{Array I None\}) - The Hamiltonian rewritten as sum_i H_bond [i] for MPS indices i. H_bond [i] acts on sites (i-1, i); we require len (H_bond) == lat. N_sites. Legs of each H_bond[i] are ['p0', 'p0*', 'p1', 'p1*'].

\section*{H_bond}

The Hamiltonian rewritten as sum_i H_bond [i] for MPS indices i. H_bond [i] acts on sites (i-1, i), None represents 0 . Legs of each H_bond[i] are ['p0', 'p0*', 'p1', 'p1*']. H_bond is not affected by the explicit_plus_hc flag of a CouplingModel.

Type list of \{Array I None \}
classmethod from_MPOModel (mpo_model)
Initialize a NearestNeighborModel from a model class defining an MPO.
This is especially usefull in combination with MPOModel.group_sites().
Parameters mpo_model (MPOMOdel) - A model instance implementing the MPO. Does not need to be a NearestNeighborModel, but should only have nearest-neighbor couplings.

\section*{Examples}

The SpinChainNNN2 has next-nearest-neighbor couplings and thus only implements an MPO:
```

>>> from tenpy.models.spins_nnn import SpinChainNNN2
>>> nnn_chain = SpinChainNNN2({'L': 20})
>>> print(isinstance(nnn_chain, NearestNeighborModel))
False
>>> print("range before grouping:", nnn_chain.H_MPO.max_range)
range before grouping: 2

```

By grouping each two neighboring sites, we can bring it down to nearest neighbors.
```

>>> grouped_sites = nnn_chain.group_sites(2)
>>> print("range after grouping:", nnn_chain.H_MPO.max_range)
range after grouping: 1

```

Yet, TEBD will not yet work, as the model doesn't define \(H_{-}\)bond. However, we can initialize a NearestNeighborModel from the MPO:
```

>>> nnn_chain_for_tebd = NearestNeighborModel.from_MPOModel(nnn_chain)
>>> isinstance(nnn_chain_for_tebd, NearestNeighborModel)
True

```

\section*{trivial_like_NNModel()}

Return a NearestNeighborModel with same lattice, but trivial ( \(\mathrm{H}=0\) ) bonds.
bond_energies ( \(p s i\) )
Calculate bond energies <psilH_bondlpsi>.
Parameters psi (MPS) - The MPS for which the bond energies should be calculated.
Returns E_bond - List of bond energies: for finite bc, E_Bond [i] is the energy of bond i, i+1. (i.e. we omit bond 0 between sites L-1 and 0); for infinite bc E_bond[i] is the energy of bond \(i-1\), \(i\).
Return type 1D ndarray
enlarge_mps_unit_cell (factor=2)
Repeat the unit cell for infinite MPS boundary conditions; in place.
This has to be done after finishing initialization and can not be reverted.
Parameters factor (int) - The new number of sites in the MPS unit cell will be increased from \(N\) _sites to factor \(* N\) _sites_per_ring. Since MPS unit cells are repeated in the \(x\)-direction in our convetion, the lattice shape goes from ( \(L x, L y, \ldots, L u\) ) to (Lx*factor, Ly, ..., Lu).
group_sites ( \(n=2\), grouped_sites=None)
Modify self in place to group sites.
Group each \(n\) sites together using the GroupedSite. This might allow to do TEBD with a Trotter decomposition, or help the convergence of DMRG (in case of too long range interactions).
This has to be done after finishing initialization and can not be reverted.

\section*{Parameters}
- \(\mathbf{n}(i n t)\) - Number of sites to be grouped together.
- grouped_sites (None l list of GroupedSite) - The sites grouped together.

Returns grouped_sites - The sites grouped together.
Return type list of GroupedSite
calc_H_MPO_from_bond (tol_zero=1e-15)
Calculate the MPO Hamiltonian from the bond Hamiltonian.
Parameters tol_zero (float) - Arrays with norm <tol_zero are considered to be zero.
Returns H_MPO - MPO representation of the Hamiltonian.
Return type MPO
classmethod from_hdf5 (hdf5_loader, h5gr, subpath)
Load instance from a HDF5 file.
This method reconstructs a class instance from the data saved with save_hdf5 ().

\section*{Parameters}
- hdf5_loader (Hdf5Loader) - Instance of the loading engine.
- h5gr (Group) - HDF5 group which is represent the object to be constructed.
- subpath (str) - The name of \(h 5 g r\) with a ' / ' in the end.

Returns obj - Newly generated class instance containing the required data.
Return type cls
save_hdf5 (hdf5_saver, h5gr, subpath)
Export self into a HDF5 file.
This method saves all the data it needs to reconstruct self with from_hdf5 ().
This implementation saves the content of __ dict__ with save_dict_content (), storing the format under the attribute 'format'.

\section*{Parameters}
- hdf5_saver (Hdf5Saver) - Instance of the saving engine.
- h5gr (: class`Group`) - HDF5 group which is supposed to represent self.
- subpath (str) - The name of \(h 5 g r\) with a '/' in the end.

\section*{Module description}

This module contains some base classes for models.
A 'model' is supposed to represent a Hamiltonian in a generalized way. The Lattice specifies the geometry and underlying Hilbert space, and is thus common to all models. It is needed to intialize the common base class Model of all models.

Different algorithms require different representations of the Hamiltonian. For example for DMRG, the Hamiltonian needs to be given as an MPO, while TEBD needs the Hamiltonian to be represented by 'nearest neighbor' bond terms. This module contains the base classes defining these possible representations, namley the MPOMOdel and NearestNeighborModel.

A particular model like the XXZChain should then yet another class derived from these classes. In it's __init__, it needs to explicitly call the MPOModel.__init__(self, lattice, H_MPO), providing an MPO representation of H, and also the NearestNeighborModel.__init__(self, lattice, H_bond), providing a representation of H by bond terms \(H_{-}\)bond.

The CouplingModel is the attempt to generalize the representation of \(H\) by explicitly specifying the couplings in a general way, and providing functionality for converting them into \(H_{-} M P O\) and \(H_{-}\)bond. This allows to quickly generate new model classes for a very broad class of Hamiltonians.

The CouplingMPOModel aims at structuring the initialization for most models and is used as base class in (most of) the predefined models in TeNPy.

See also the introduction in Models.

\section*{Specific models}
\begin{tabular}{ll}
\hline tf_ising & \begin{tabular}{l} 
Prototypical example of a quantum model: the trans- \\
verse field Ising model.
\end{tabular} \\
\hline xxz_chain & Prototypical example of a 1D quantum model: the spin- \\
& \(1 / 2\) XXZ chain. \\
\hline spins & Nearest-neighbour spin-S models. \\
\hline spins_nnn & Next-Nearest-neighbour spin-S models. \\
\hline fermions_spinless & Spinless fermions with hopping and interaction. \\
\hline hubbard & Bosonic and fermionic Hubbard models. \\
\hline hofstadter & Cold atomic (Harper-)Hofstadter model on a strip or \\
& cylinder. \\
\hline haldane & Bosonic and fermionic Haldane models. \\
\hline toric_code & Kitaev's exactly solvable toric code model. \\
\hline
\end{tabular}

\section*{19.3 tf_ising}
- full name: tenpy.models.tf_ising
- parent module: tenpy.models
- type: module

\section*{Classes}

\begin{tabular}{ll}
\hline TFIChain(model_params) & The TFIModel on a Chain, suitable for TEBD. \\
\hline TFIModel(model_params) & Transverse field Ising model on a general lattice. \\
\hline
\end{tabular}

\subsection*{19.3.1 TFIChain}
- full name: tenpy.models.tf_ising.TFIChain
- parent module: tenpy.models.tf_ising
- type: class

\section*{Inheritance Diagram}


\section*{Methods}
\begin{tabular}{|c|c|}
\hline TFIChain.__init__(model_params) & Initialize self. \\
\hline TFIChain.add_coupling(strength, u1, op1, u2, ...) & Add twosite coupling terms to the Hamiltonian, summing over lattice sites. \\
\hline ```
TFIChain.add_coupling_term(strength, i, j,
...)
``` & Add a two-site coupling term on given MPS sites. \\
\hline TFIChain.add_exponentially_decaying_c & Add ang(expønentially decaying long-range coupling. \\
\hline ```
TFIChain.add_local_term(strength, term[,
...])
``` & Add a single term to self. \\
\hline TFIChain.add_multi_coupling(strength, ops[, ...]) & Add multi-site coupling terms to the Hamiltonian, summing over lattice sites. \\
\hline TFIChain.add_multi_coupling_term(strength, ...) & Add a general M-site coupling term on given MPS sites. \\
\hline TFIChain.add_onsite(strength, u, opname[,...]) & Add onsite terms to onsite_terms. \\
\hline
\end{tabular}

Table 39 - continued from previous page
\begin{tabular}{|c|c|}
\hline TFIChain.add_onsite_term(strength, i, op[, ...]) & Add an onsite term on a given MPS site. \\
\hline TFIChain.all_coupling_terms() & Sum of all coupling_terms. \\
\hline TFIChain.all_onsite_terms() & Sum of all onsite_terms. \\
\hline TFIChain.bond_energies(psi) & Calculate bond energies <psilH_bondlpsi>. \\
\hline TFIChain.calc_H_MPO([tol_zero]) & Calculate MPO representation of the Hamiltonian. \\
\hline TFIChain.calc_H_MPO_from_bond([tol_zero]) & Calculate the MPO Hamiltonian from the bond Hamiltonian. \\
\hline TFIChain.calc_H_bond([tol_zero]) & calculate H_bond from coupling_terms and onsite_terms. \\
\hline TFIChain.calc_H_bond_from_MPO([tol_zero]) & Calculate the bond Hamiltonian from the MPO Hamiltonian. \\
\hline TFIChain.calc_H_onsite([tol_zero]) & Calculate H_onsite from self.onsite_terms. \\
\hline TFIChain.coupling_strength_add_ext & Add) an external flux to the coupling strength. \\
\hline TFIChain.enlarge_mps_unit_cell([factor]) & Repeat the unit cell for infinite MPS boundary conditions; in place. \\
\hline TFIChain.from_MPOModel(mpo_model) & Initialize a NearestNeighborModel from a model class defining an MPO. \\
\hline TFIChain.from_hdf5(hdf5_loader, h5gr, subp & Load instance from a HDF5 file. \\
\hline TFIChain.group_sites([n, grouped_sites]) & Modify self in place to group sites. \\
\hline TFIChain.init_H_from_terms() & Initialize \(H_{-} M P O\) (and \(H_{-}\)bond) from the terms of the CouplingModel. \\
\hline TFIChain.init_lattice(model_params) & Initialize a lattice for the given model parameters. \\
\hline TFIChain.init_sites(model_params) & Define the local Hilbert space and operators; needs to be implemented in subclasses. \\
\hline TFIChain.init_terms(model_params) & Add the onsite and coupling terms to the model; subclasses should implement this. \\
\hline TFIChain.save_hdf5(hdf5_saver, h5gr, subpath) & Export self into a HDF5 file. \\
\hline TFIChain.test_sanity() & Sanity check, raises ValueErrors, if something is wrong. \\
\hline TFIChain.trivial_like_NNModel() & Return a NearestNeighborModel with same lattice, but trivial \((\mathrm{H}=0)\) bonds. \\
\hline
\end{tabular}

\section*{Class Attributes and Properties}
\begin{tabular}{ll}
\hline TFIChain.force_default_lattice & If True, init_lattice () asserts that the initialized \\
lattice is (a subclass of) default_lattice
\end{tabular}
class tenpy.models.tf_ising.TFIChain(model_params)
Bases: tenpy.models.tf_ising.TFIModel, tenpy.models.model.
NearestNeighborModel
The TFIModel on a Chain, suitable for TEBD.
See the TFIModel for the documentation of parameters.
```

    default_lattice
        alias of tenpy.models.lattice.Chain
    ```
add_coupling (strength, u1, op1, u2, op2, dx, op_string=None, str_on_first=True, raise_op2_left=False, category=None, plus_hc=False)
Add twosite coupling terms to the Hamiltonian, summing over lattice sites.
Represents couplings of the form \(\sum_{x_{0}, \ldots, x_{\text {dim-1 }}} \operatorname{strength}[\operatorname{shift}(\vec{x})] * O P 0 * O P 1\), where OP 0 := lat.unit_cell[u0].get_op(op0) acts on the site (x_0, ..., x_\{dim-1\}, u1), and OP1 := lat.unit_cell[u1].get_op(op1) acts on the site (x_0+dx[0], ..., \(x_{\_}\{d i m-1\}+d x[\operatorname{dim}-1]\), \(\left.u 1\right)\). Possible combinations \(x_{-} 0, \ldots, x_{-}\{d i m-1\}\) are determined from the boundary conditions in possible_couplings ().

The coupling strength may vary spatially if the given strength is a numpy array. The correct shape of this array is the coupling_shape returned by tenpy.models.lattice.coupling_shape () and depends on the boundary conditions. The shift (...) depends on \(d x\), and is chosen such that the first entry strength [0, 0, ...] of strength is the prefactor for the first possible coupling fitting into the lattice if you imagine open boundary conditions.

The necessary terms are just added to coupling_terms; this function does not rebuild the MPO.
Deprecated since version 0.4.0: The arguments str_on_first and raise_op2_left will be removed in version 1.0.0.

\section*{Parameters}
- strength (scalar / array) - Prefactor of the coupling. May vary spatially (see above). If an array of smaller size is provided, it gets tiled to the required shape.
- u1 (int) - Picks the site lat. unit_cell[u1] for OP1.
- op1 (str) - Valid operator name of an onsite operator in lat.unit_cell[u1] for OP1.
- u2 (int) - Picks the site lat. unit_cell [u2] for OP2.
- op2 (str) - Valid operator name of an onsite operator in lat.unit_cell[u2] for OP2.
- \(\mathbf{d x}\) (iterable of int) - Translation vector (of the unit cell) between OP1 and OP2. For a 1D lattice, a single int is also fine.
- op_string (str / None) - Name of an operator to be used between the OP1 and OP2 sites. Typical use case is the phase for a Jordan-Wigner transformation. The operator should be defined on all sites in the unit cell. If None, auto-determine whether a JordanWigner string is needed, using op_needs_JW ().
- str_on_first (bool) - Whether the provided op_string should also act on the first site. This option should be chosen as True for Jordan-Wigner strings. When handling Jordan-Wigner strings we need to extend the op_string to also act on the 'left', first site (in the sense of the MPS ordering of the sites given by the lattice). In this case, there is a well-defined ordering of the operators in the physical sense (i.e. which of opl or op2 acts first on a given state). We follow the convention that op2 acts first (in the physical sense), independent of the MPS ordering. Deprecated.
- raise_op2_left (bool) - Raise an error when op2 appears left of opl (in the sense of the MPS ordering given by the lattice). Deprecated.
- category (str) - Descriptive name used as key for coupling_terms. Defaults to a string of the form " \(\{o p 1\} \_i \quad\{o p 2\} \_j "\).
- plus_hc (bool) - If True, the hermitian conjugate of the terms is added automatically.

\section*{Examples}

When initializing a model, you can add a term \(J \sum_{<i, j>} S_{i}^{z} S_{j}^{z}\) on all nearest-neighbor bonds of the lattice like this:
```

>>> J = 1. \# the strength
>>> for u1, u2, dx in self.lat.pairs['nearest_neighbors']:
... self.add_coupling(J, u1, 'Sz', u2, 'Sz', dx)

```

The strength can be an array, which gets tiled to the correct shape. For example, in a 1D Chain with an even number of sites and periodic (or infinite) boundary conditions, you can add alternating strong and weak couplings with a line like:
```

>>> self.add_coupling([1.5, 1.], u1, 'Sz', u2, 'Sz', dx)

```

Make sure to use the plus_hc argument if necessary, e.g. for hoppings:
```

>>> t = 1. \# hopping strength
>>> for u1, u2, dx in self.lat.pairs['nearest_neighbors']:
... self.add_coupling(t, ul, 'Cd', u2, 'C', dx, plus_hc=True)

```

Alternatively, you can add the hermitian conjugate terms explictly. The correct way is to complex conjugate the strength, take the hermitian conjugate of the operators and swap the order (including a swap \(u 1<->u 2\) ), and use the opposite direction \(-d x\), i.e. the h.c. of add_coupling ( \(t, \mathrm{u} 1\), 'A', u2, 'B', dx) is add_coupling(np.conj(t), u2, hc('B'), u1, hc('A'), -dx), where \(h c\) takes the hermitian conjugate of the operator names, see get_hc_op_name (). For spin-less fermions (FermionSite), this would be
```

>>> for ul, u2, dx in self.lat.pairs['nearest_neighbors']:
... self.add_coupling(t, u1, 'Cd', u2, 'C', dx)
... self.add_coupling(np.conj(t), u2, 'Cd', u1, 'C', -dx) \# h.c.

```

With spin-full fermions (SpinHalfFermions), it could be:
```

>>> for u1, u2, dx in self.lat.pairs['nearest_neighbors']:
... self.add_coupling(t, u1, 'Cdu', u2, 'Cd', dx) \# Cdagger_up C_down
... self.add_coupling(np.conj(t), u2, 'Cdd', u1, 'Cu', -dx) \# h.c.u
Cdagger_down C_up

```

Note that the Jordan-Wigner strings for the fermions are added automatically!

\section*{See also:}
add_onsite Add terms acting on one site only.
add_multi_coupling_term for terms on more than two sites.
add_coupling_term Add a single term without summing over \(\vec{x}\).
add_coupling_term (strength, \(i, j, o p_{-} i, o p \_j\), op_string='Id', category=None, plus_hc=False)
Add a two-site coupling term on given MPS sites.
Wrapper for self.coupling_terms[category].add_coupling_term(...).

Warning: This function does not handle Jordan-Wigner strings! You might want to use add_local_term() instead.

\section*{Parameters}
- strength (float) - The strength of the coupling term.
- i (int) - The MPS indices of the two sites on which the operator acts. We require 0 <= i < N_sites and i < j, i.e., op_i acts "left" of op_j. If j >= N_sites, it indicates couplings between unit cells of an infinite MPS.
- \(\mathbf{j}\) (int) - The MPS indices of the two sites on which the operator acts. We require 0 <= i < N_sites and i < j, i.e., op_i acts "left" of \(o p_{-} j\). If j\(\rangle=\mathrm{N} \_\)sites, it indicates couplings between unit cells of an infinite MPS.
- op1 (str) - Names of the involved operators.
- op2 (str) - Names of the involved operators.
- op_string (str) - The operator to be inserted between \(i\) and \(j\).
- category (str) - Descriptive name used as key for coupling_terms. Defaults to a string of the form " \(\{\mathrm{op} 1\} \_\)i \(\{o p 2\} \_j\) ".
- plus_hc (bool) - If True, the hermitian conjugate of the term is added automatically.
add_exponentially_decaying_coupling (strength, lambda_, op_i, op_j, subsites=None, \(o p \_\)string \(=\)None, \(p l u s \_h c=\) False)
Add an exponentially decaying long-range coupling.
\[
\text { strength } \sum_{i<j} \lambda^{|i-j|} A_{\text {subsites }[i]} B_{\text {subsites }[j]}
\]

Where the operator \(A\) is given by op_i, and \(B\) is given by op_j. Note that the sum over \(\mathrm{i}, \mathrm{j}\) is long-range, for infinite systems going beyond the MPS unit cell. Moreover, note that the distance in the exponent is the distance within subsites.

\section*{Parameters}
- strength (float) - Overall prefactor.
- lambda (float) - Decay-rate
- op_i (string) - Names for the operators.
- op_j (string) - Names for the operators.
- subsites (None | ID array) - Selects a subset of sites within the MPS unit cell on which the operators act. Needs to be sorted. None selects all sites.
- op_string (None | str) - The operator to be inserted between \(A\) and \(B\); If None, this function checks whether a fermionic "JW" string is needed for the given operators; in this case the right \(o p_{-} j\) acts first.
- plus_hc (bool) - If True, the hermitian conjugate of the term is added automatically.

\section*{Examples}

At least for simple enough 1D chains (or ladders), you can use fit_with_sum_of_exp () to approximate a long-range function with a few sum of exponentials and then add them with this function.
```

>>> def decay(x):
... return np.exp (-0.1*x) / x**2
>>> from tenpy.tools.fit import fit_with_sum_of_exp, sum_of_exp
>>> n_exp = 5
>>> fit_range = 50
>>> lam, pref = fit_with_sum_of_exp(decay, n_exp, fit_range)

```
(continued from previous page)
```

>>> x = np.arange(1, fit_range + 1)
>>> print('error in fit: {0:.3e}'.format(np.sum(np.abs(decay(x) - sum_of_
\hookrightarrowexp(lam, pref, x)))))
error in fit: 1.073e-04
>>> for pr, la in zip(pref, lam):
self.add_exponentially_decaying_coupling(pr, la, 'N', 'N')

```
add_local_term (strength, term, category=None, plus_hc=False)
Add a single term to self.
The repesented term is strength times the product of the operators given in terms. Each operator is specified by the name and the site it acts on; the latter given by a lattice index, see Lattice.

Depending on the length of term, it can add an onsite term or a coupling term to onsite_terms or coupling_terms, respectively.

\section*{Parameters}
- strength (float/complex) - The prefactor of the term.
- term (list of (str, array_like)) - List of tuples (opname, lat_idx) where opname is a string describing the operator acting on the site given by the lattice index lat_idx. Here, lat_idx is for example \([x, y, u]\) for a 2D lattice, with \(u\) being the index within the unit cell.
- category - Descriptive name used as key for onsite_terms or coupling_terms.
- plus_hc (bool) - If True, the hermitian conjugate of the terms is added automatically.
add_multi_coupling (strength, ops, _deprecate_l='DEPRECATED', _deprecate_2 = 'DEPRECATED', op_string=None, category=None, plus_hc=False)
Add multi-site coupling terms to the Hamiltonian, summing over lattice sites.
Represents couplings of the form \(\operatorname{sum}_{\vec{x}} \operatorname{strength}[\operatorname{shift}(\vec{x})] * O P_{0} * O P_{1} * \ldots * O P_{M-1}\), involving \(M\) operators. Here, \(O P_{m}\) stands for the operator defined by the \(m\)-th tuple (opname, \(\mathrm{dx}, \mathrm{u}\) ) given in the argument ops, which determines the position \(\vec{x}+\overrightarrow{d x}\) and unit-cell index \(u\) of the site it acts on; the actual operator is given by self.lat.unit_cell[u].get_op(opname).

The coupling strength may vary spatially if the given strength is a numpy array. The correct shape of this array is the coupling_shape returned by tenpy.models.lattice. possible_multi_couplings() and depends on the boundary conditions. The shift(...) depends on the \(d x\) entries of ops and is chosen such that the first entry strength [0, 0, ...] of strength is the prefactor for the first possible coupling fitting into the lattice if you imagine open boundary conditions.

The necessary terms are just added to coupling_terms; this function does not rebuild the MPO.
Deprecated since version 0.6.0: We switched from the three arguments \(u 0\), op0 and other_op with other_ops=[(u1, op1, dx1), (op2, u2, dx2), ...] to a single, equivalent argment ops which should now read ops=[(op0, dx0, u0), (op1, dx1, u1), (op2, dx2, u2), . ..], where \(\mathrm{dx} 0=[0] *\) self.lat.dim. Note the changed order inside the tuples!

\section*{Parameters}
- strength (scalar / array) - Prefactor of the coupling. May vary spatially, and is tiled to the required shape.
- ops (list of (opname, \(d x, u\) )) - Each tuple determines one operator of the coupling, see the description above. opname (str) is the name of the operator, \(d x\) (list of length
lat.dim) is a translation vector, and \(u\) (int) is the index of lat.unit_cell on which the operator acts. The first entry of ops corresponds to \(O P_{0}\) and acts last in the physical sense.
- op_string (str / None) - If a string is given, we use this as the name of an operator to be used inbetween the operators, excluding the sites on which any operators act. This operator should be defined on all sites in the unit cell.

If None, auto-determine whether a Jordan-Wigner string is needed (using op_needs_JW ()) for each of the segments inbetween the operators and also on the sites of the left operators.
- category (str) - Descriptive name used as key for coupling_terms. Defaults to a string of the form " \(\{0 \mathrm{p} 0\}\) _i \(\{0\) her_ops [0]\}_j \(\{0\) other_ops [1]\}_k ...".
- plus_hc (bool) - If True, the hermitian conjugate of the terms is added automatically.

\section*{Examples}

A call to add_coupling() with arguments add_coupling(strength, u1, 'A', u2, ' \(\mathrm{B}^{\prime}, \mathrm{dx}\) ) is equivalent to the following:
```

>>> dx_0 = [0] * self.lat.dim \# = [0] for a ID lattice, [0, 0] in 2D
>>> self.add_multi_coupling(strength, [('A', dx_0, u1), ('B', dx, u2)])

```

To explicitly add the hermitian conjugate (instead of simply using plus_hc = True), you need to take the complex conjugate of the strength, reverse the order of the operators and take the hermitian conjugates of the individual operator names (indicated by the hc (...), see get_hc_op_name ()):
```

>>> self.add_multi_coupling(np.conj(strength), [(hc('B'), dx, u2), (hc('A'),
\hookrightarrowdx_0, u1)])

```

\section*{See also:}
add_onsite Add terms acting on one site only.
add_coupling Add terms acting on two sites.
add_multi_coupling_term Add a single term, not summing over the possible \(\vec{x}\).
add_multi_coupling_term (strength, ijkl,ops_ijkl,op_string, category=None, plus_hc=False)
Add a general M-site coupling term on given MPS sites.
Wrapper for self.coupling_terms[category].add_multi_coupling_term(....).

\section*{Warning: This function does not handle Jordan-Wigner strings! You might want to use add_local_term() instead.}

\section*{Parameters}
- strength (float) - The strength of the coupling term.
- ijkl (list of int) - The MPS indices of the sites on which the operators acts. With \(i, j, k, \ldots=i j k l\), we require that they are ordered ascending, \(i<j<k<\ldots\) and that \(0<=i<N \_s i t e s\). Inidces \(>=\) N_sites indicate couplings between different unit cells of an infinite MPS.
- ops_ijkl(list of str) - Names of the involved operators on sites \(i, j, k, \ldots\).
- op_string (list of str) - Names of the operator to be inserted between the operators, e.g., op_string[0] is inserted between \(i\) and \(j\).
- category (str) - Descriptive name used as key for coupling_terms. Defaults to a string of the form " \(\{\mathrm{op} 0\} \_i \quad\{o p 1\} \_j \quad\{o p 2\} \_k \ldots\). . "
- plus_hc (bool) - If True, the hermitian conjugate of the term is added automatically.
add_onsite (strength, \(u\), opname, category=None, plus_hc=False)
Add onsite terms to onsite_terms.
Adds \(\sum_{\vec{x}}\) strength \([\vec{x}] * O P\) to the represented Hamiltonian, where the operator \(\mathrm{OP}=1\) at. unit_cell[u].get_op(opname) acts on the site given by a lattice index (x_0, ..., x_\{dim-1\}, u),

The necessary terms are just added to onsite_terms; doesn't rebuild the MPO.

\section*{Parameters}
- strength (scalar / array) - Prefactor of the onsite term. May vary spatially. If an array of smaller size is provided, it gets tiled to the required shape.
- u(int) - Picks a Site lat.unit_cell[u] out of the unit cell.
- opname (str) - valid operator name of an onsite operator in lat.unit_cell [u].
- category (str) - Descriptive name used as key for onsite_terms. Defaults to opname.
- plus_hc (bool) - If True, the hermitian conjugate of the terms is added automatically.

\section*{See also:}
add_coupling Add a terms acting on two sites.
add_onsite_term Add a single term without summing over vecx.
add_onsite_term (strength, \(i\), op, category=None, plus_hc=False)
Add an onsite term on a given MPS site.
Wrapper for self.onsite_terms[category].add_onsite_term(....).

\section*{Parameters}
- strength (float) - The strength of the term.
- i(int) - The MPS index of the site on which the operator acts. We require \(0<=i<\) L.
- op (str) - Name of the involved operator.
- category (str) - Descriptive name used as key for onsite_terms. Defaults to op.
- plus_hc (bool) - If True, the hermitian conjugate of the term is added automatically.
all_coupling_terms()
Sum of all coupling_terms.
all_onsite_terms()
Sum of all onsite_terms.

\section*{bond_energies ( \(p s i\) )}

Calculate bond energies <psilH_bondlpsi>.
Parameters psi (MPS) - The MPS for which the bond energies should be calculated.

Returns E_bond - List of bond energies: for finite bc, E_Bond [i] is the energy of bond i, i+1. (i.e. we omit bond 0 between sites L-1 and 0 ); for infinite bc E_bond[i] is the energy of bond \(i-1, i\).
Return type 1D ndarray
calc_H_MPO (tol_zero \(=1 e-15\) )
Calculate MPO representation of the Hamiltonian.
Uses onsite_terms and coupling_terms to build an MPOGraph (and then an MPO).
Parameters tol_zero (float) - Prefactors with abs (strength) < tol_zero are considered to be zero.

Returns H_MPO - MPO representation of the Hamiltonian.
Return type MPO
calc_H_MPO_from_bond (tol_zero=le-15)
Calculate the MPO Hamiltonian from the bond Hamiltonian.
Parameters tol_zero \((f l o a t)\) - Arrays with norm <tol_zero are considered to be zero.
Returns H_MPO - MPO representation of the Hamiltonian.
Return type MPO
calc_H_bond (tol_zero \(=1 e-15\) )
calculate \(H\) _bond from coupling_terms and onsite_terms.
Parameters tol_zero (float) - prefactors with abs(strength) < tol_zero are considered to be zero.

Returns H_bond - Bond terms as required by the constructor of NearestNeighborModel. Legs are ['p0', 'p0*', 'p1', 'p1*']
Return type list of Array
:raises ValueError : if the Hamiltonian contains longer-range terms.:
calc_H_bond_from_MPO (tol_zero=le-15)
Calculate the bond Hamiltonian from the MPO Hamiltonian.
Parameters tol_zero (float) - Arrays with norm <tol_zero are considered to be zero.
Returns H_bond - Bond terms as required by the constructor of NearestNeighborModel. Legs are ['p0', 'p0*', 'p1', 'p1*']
Return type list of Array
:raises ValueError : if the Hamiltonian contains longer-range terms.:
calc_H_onsite (tol_zero=le-15)
Calculate H_onsite from self.onsite_terms.
Deprecated since version 0.4.0: This function will be removed in 1.0.0. Replace calls to this function by self.all_onsite_terms().remove_zeros(tol_zero).to_Arrays(self. lat.mps_sites()). You might also want to take explicit_plus_hc into account.

Parameters tol_zero (float) - prefactors with abs (strength) < tol_zero are considered to be zero.

\section*{Returns}
- H_onsite (list of npc.Array)
- onsite terms of the Hamiltonian. If explicit_plus_hc is True, - Hermitian conjugates of the onsite terms will be included.
coupling_strength_add_ext_flux (strength, \(d x\), phase)
Add an external flux to the coupling strength.
When performing DMRG on a "cylinder" geometry, it might be useful to put an "external flux" through the cylinder. This means that a particle hopping around the cylinder should pick up a phase given by the external flux [[resta1998]]. This is also called "twisted boundary conditions" in literature. This function adds a complex phase to the strength array on some bonds, such that particles hopping in positive direction around the cylinder pick up \(\exp (+i\) phase \()\).

Warning: For the sign of phase it is important that you consistently use the creation operator as opl and the annihilation operator as op2 in add_coupling().

\section*{Parameters}
- strength (scalar / array) - The strength to be used in add_coupling(), when no external flux would be present.
- \(\mathbf{d} \mathbf{x}\) (iterable of int) - Translation vector (of the unit cell) between opl and op2 in add_coupling().
- phase (iterable of float) - The phase of the external flux for hopping in each direction of the lattice. E.g., if you want flux through the cylinder on which you have an infinite MPS, you should give phase \(=[0\), phi] souch that particles pick up a phase phi when hopping around the cylinder.
Returns strength - The strength array to be used as strength in add_coupling() with the given \(d x\).
Return type complex array

\section*{Examples}

Let's say you have an infinite MPS on a cylinder, and want to add nearest-neighbor hopping of fermions with the FermionSite. The cylinder axis is the \(x\)-direction of the lattice, so to put a flux through the cylinder, you want particles hopping around the cylinder to pick up a phase phi given by the external flux.
```

>>> strength = 1. \# hopping strength without external flux
>>> phi = np.pi/4 \# determines the external flux strength
>>> for u1, u2, dx in self.lat.pairs['nearest_neighbors']:
... strength_with_flux = self.coupling_strength_add_ext_flux(strength, dx,
@ [0, phi])
... self.add_coupling(strength_with_flux, u1, 'Cd', u2, 'C', dx)
... self.add_coupling(np.conj(strength_with_flux), u2, 'cd', u1, 'C', -dx)

```
enlarge_mps_unit_cell (factor=2)
Repeat the unit cell for infinite MPS boundary conditions; in place.
This has to be done after finishing initialization and can not be reverted.
Parameters factor (int) - The new number of sites in the MPS unit cell will be increased from \(N\) _sites to factor*N_sites_per_ring. Since MPS unit cells are repeated in the \(x\)-direction in our convetion, the lattice shape goes from ( \(L x, L y, \ldots, L u\) ) to (Lx*factor, Ly, ..., Lu).

\section*{classmethod from_MPOModel (mpo_model)}

Initialize a NearestNeighborModel from a model class defining an MPO.
This is especially usefull in combination with MPOModel.group_sites ().
Parameters mpo_model (MPOModel) - A model instance implementing the MPO. Does not need to be a NearestNeighborModel, but should only have nearest-neighbor couplings.

\section*{Examples}

The SpinChainNNN2 has next-nearest-neighbor couplings and thus only implements an MPO:
```

>>> from tenpy.models.spins_nnn import SpinChainNNN2
>>> nnn_chain = SpinChainNNN2({'L': 20})
>>> print(isinstance(nnn_chain, NearestNeighborModel))
False
>>> print("range before grouping:", nnn_chain.H_MPO.max_range)
range before grouping: 2

```

By grouping each two neighboring sites, we can bring it down to nearest neighbors.
```

>>> grouped_sites = nnn_chain.group_sites(2)
>>> print("range after grouping:", nnn_chain.H_MPO.max_range)
range after grouping: 1

```

Yet, TEBD will not yet work, as the model doesn't define \(H_{\_}\)bond. However, we can initialize a NearestNeighborModel from the MPO:
```

>>> nnn_chain_for_tebd = NearestNeighborModel.from_MPOModel(nnn_chain)
>>> isinstance(nnn_chain_for_tebd, NearestNeighborModel)
True

```
classmethod from_hdf5 (hdf5_loader, h5gr, subpath)
Load instance from a HDF5 file.
This method reconstructs a class instance from the data saved with save_hdf5 ().

\section*{Parameters}
- hdf5_loader (Hdf5Loader) - Instance of the loading engine.
- h5gr (Group) - HDF5 group which is represent the object to be constructed.
- subpath (str) - The name of \(h 5 g r\) with a '/' in the end.

Returns obj - Newly generated class instance containing the required data.

\section*{Return type cls}
group_sites ( \(n=2\), grouped_sites=None)
Modify self in place to group sites.
Group each \(n\) sites together using the GroupedSite. This might allow to do TEBD with a Trotter decomposition, or help the convergence of DMRG (in case of too long range interactions).
This has to be done after finishing initialization and can not be reverted.

\section*{Parameters}
- \(\mathbf{n}(i n t)\) - Number of sites to be grouped together.
- grouped_sites (None l list of GroupedSite) - The sites grouped together.

Returns grouped_sites - The sites grouped together.
Return type list of GroupedSite

\section*{init_H_from_terms()}

Initialize \(H_{-} M P O\) (and \(H \_\)bond) from the terms of the CouplingModel.
This function is called automatically during CouplingMPOModel.__init__
If you use one of the add_* methods of the CouplingModel after initialization, you will need to call init_H_from_terms in the end by yourself, in order to update the \(H_{-} M P O\) (and possibly \(H_{-} b o n d\) ) representations. (You should get a warning about this... The way to avoid it is to initialize all the terms in init_terms by defining your own model, as outlined in Models.
init_lattice (model_params)
Initialize a lattice for the given model parameters.
This function reads out the model parameter lattice. This can be a full Lattice instance, in which case it is just returned without further action. Alternatively, the lattice parameter can be a string giving the name of one of the predefined lattices, which then gets initialized. Depending on the dimensionality of the lattice, this requires different model parameters.

Parameters model_params (dict) - The model parameters given to \(\qquad\) init \(\qquad\) _.

Returns lat - An initialized lattice.
Return type Lattice

\section*{Options}
option CouplingMPOModel.lattice: str | Lattice
The name of a lattice pre-defined in TeNPy to be initialized. Alternatively, directly a subclass of Lattice instead of the name. Alternatively, a (possibly self-defined) Lattice instance. In the latter case, no further parameters are read out.
option CouplingMPOModel.bc_MPS: str Boundary conditions for the MPS.
option CouplingMPOModel.order: str
The order of sites within the lattice for non-trivial lattices, e.g, 'default', 'snake', see ordering (). Only used if lattice is a string.
option CouplingMPOModel.L: int
The length in x-direction; only read out for 1D lattices. For an infinite system the length of the unit cell.
option CouplingMPOModel.Lx: int
option CouplingMPOModel.Ly: int
The length in x - and y-direction; only read out for 2D lattices. For "infinite" \(b c_{-}\)MPS, the system is infinite in x-direction and \(L x\) is the number of "rings" in the infinite MPS unit cell, while \(L y\) gives the circumference around the cylinder or width of th the rung for a ladder (depending on \(b c_{-} y\) ).
option CouplingMPOModel.bc_y: str
"cylinder" | "ladder"; only read out for 2D lattices. The boundary conditions in y-direction.
option CouplingMPOModel.bc_x: str
"open" | "periodic". Can be used to force "periodic" boundaries for the lattice, i.e., for the couplings in the Hamiltonian, even if the MPS is finite. Defaults to "open" for bc_MPS="finite " and "periodic" for bc_MPS="infinite. If you are not aware of the consequences, you should probably not use "periodic" boundary conditions. (The MPS is still "open", so this will introduce long-range couplings between the first and last sites of the MPS!)

\section*{init_sites (model_params)}

Define the local Hilbert space and operators; needs to be implemented in subclasses.
This function gets called by init_lattice () to get the Site for the lattice unit cell.

Note: Initializing the sites requires to define the conserved quantum numbers. All pre-defined sites accept conserve=None to disable using quantum numbers. Many models in TeNPy read out the conserve model parameter, which can be set to "best" to indicate the optimal parameters.

If you need to initialize more than one site, the function tenpy.networks.site. set_common_charges () should be helpful.

Parameters model_params (dict) - The model parameters given to \(\qquad\) init \(\qquad\)
Returns sites - The local sites of the lattice, defining the local basis states and operators.
Return type (tuple of) Site
init_terms (model_params)
Add the onsite and coupling terms to the model; subclasses should implement this.
save_hdf5 (hdf5_saver, h5gr, subpath)
Export self into a HDF5 file.
This method saves all the data it needs to reconstruct self with from_hdf5 ().
This implementation saves the content of __dict __ with save_dict_content (), storing the format under the attribute ' format'.

\section*{Parameters}
- hdf5_saver (Hdf5Saver) - Instance of the saving engine.
- h5gr (: class `Group \(\left.{ }^{\prime}\right)\) - HDF5 group which is supposed to represent self.
- subpath (str) - The name of \(h 5 g r\) with a ' / ' in the end.
```

test_sanity()

```

Sanity check, raises ValueErrors, if something is wrong.
trivial_like_NNModel()
Return a NearestNeighborModel with same lattice, but trivial ( \(\mathrm{H}=0\) ) bonds.

\section*{Module description}

Prototypical example of a quantum model: the transverse field Ising model.
Like the XXZChain, the transverse field ising chain TFIChain is contained in the more general SpinChain; the idea is more to serve as a pedagogical example for a 'model'.
We choose the field along z to allow to conserve the parity, if desired.

\section*{19.4 xxz_chain}
- full name: tenpy.models.xxz_chain
- parent module: tenpy.models
- type: module

\section*{Classes}

\begin{tabular}{ll}
\hline XXZChain(model_params) & Spin-1/2 XXZ chain with Sz conservation. \\
\hline XXZChain2(model_params) & \begin{tabular}{l} 
Another implementation of the Spin-1/2 XXZ chain \\
with \(S z\) conservation.
\end{tabular} \\
\hline
\end{tabular}

\subsection*{19.4.1 XXZChain2}
- full name: tenpy.models.xxz_chain.XXZChain2
- parent module: tenpy.models.xxz_chain
- type: class

\section*{Inheritance Diagram}


\section*{Methods}
\begin{tabular}{|c|c|}
\hline XXZChain2.__init__(model_params) & Initialize self. \\
\hline XXZChain2.add_coupling(strength, u1,op1,...) & Add twosite coupling terms to the Hamiltonian, summing over lattice sites. \\
\hline XXZChain2.add_coupling_term(strength, i, j, ...) & Add a two-site coupling term on given MPS sites. \\
\hline \multicolumn{2}{|l|}{XXZChain2.add_exponentially_decaying_coAdd án ¢xponentially decaying long-range coupling.} \\
\hline XXZChain2.add_local_term(strength, term[, ...]) & Add a single term to self. \\
\hline XXZChain2.add_multi_coupling(strength, ops) & Add multi-site coupling terms to the Hamiltonian, summing over lattice sites. \\
\hline \multicolumn{2}{|l|}{XXZChain2.add_multi_coupling_term(strengthAdd a general M-site coupling term on given MPS sites. ...)} \\
\hline ```
XXZChain2.add_onsite(strength, u, opname[,
...])
``` & Add onsite terms to onsite_terms. \\
\hline \multicolumn{2}{|l|}{xxZChain2.add_onsite_term(strength, i, op[, Add an onsite term on a given MPS site. ...])} \\
\hline XXZChain2.all_coupling_terms() & Sum of all coupling_terms. \\
\hline XXZChain2.all_onsite_terms() & Sum of all onsite_terms. \\
\hline XXZChain2.bond_energies(psi) & Calculate bond energies <psilH_bondlpsi>. \\
\hline & continues on next page \\
\hline
\end{tabular}

Table 42 - continued from previous page
\begin{tabular}{|c|c|}
\hline XXZChain2.calc_H_MPO([tol_zero]) & Calculate MPO representation of the Hamiltonian. \\
\hline XXZChain2.calc_H_MPO_from_bond([tol_zero]) & Calculate the MPO Hamiltonian from the bond Hamiltonian. \\
\hline XXZChain2.calc_H_bond([tol_zero]) & calculate \(H_{\text {_bond }}\) from coupling_terms and onsite_terms. \\
\hline XXZChain2.calc_H_bond_from_MPO([tol_zero]) & Calculate the bond Hamiltonian from the MPO Hamiltonian. \\
\hline XXZChain2.calc_H_onsite([tol_zero]) & Calculate H_onsite from self.onsite_terms. \\
\hline XXZChain2.coupling_strength_add_ext_fl & \(u\) Add.ann external flux to the coupling strength. \\
\hline XXZChain2.enlarge_mps_unit_cell([factor]) & Repeat the unit cell for infinite MPS boundary conditions; in place. \\
\hline XXZChain2.from_MPOModel(mpo_model) & Initialize a NearestNeighborModel from a model class defining an MPO. \\
\hline XXZChain2.from_hdf5(hdf5_loader, h5gr, subpath) & Load instance from a HDF5 file. \\
\hline XXZChain2.group_sites([n, grouped_sites]) & Modify self in place to group sites. \\
\hline XXZChain2.init_H_from_terms() & Initialize \(H_{-} M P O\) (and \(H \_b o n d\) ) from the terms of the CouplingModel. \\
\hline XXZChain2.init_lattice(model_params) & Initialize a lattice for the given model parameters. \\
\hline XXZChain2.init_sites(model_params) & Define the local Hilbert space and operators; needs to be implemented in subclasses. \\
\hline XXZChain2.init_terms(model_params) & Add the onsite and coupling terms to the model; subclasses should implement this. \\
\hline XXZChain2.save_hdf5(hdf5_saver, h5gr, subpath) & Export self into a HDF5 file. \\
\hline xxzChain2.test_sanity() & Sanity check, raises ValueErrors, if something is wrong. \\
\hline XXZChain2.trivial_like_NNModel() & Return a NearestNeighborModel with same lattice, but trivial ( \(\mathrm{H}=0\) ) bonds. \\
\hline
\end{tabular}

\section*{Class Attributes and Properties}
\begin{tabular}{ll}
\hline XXZChain2.default_lattice & \begin{tabular}{l} 
The default lattice class or class name to be used in \\
init_lattice().
\end{tabular} \\
\hline XXZChain2.force_default_lattice & \begin{tabular}{l} 
If True, init_lattice() asserts that the initialized \\
lattice is (a subclass of) default_lattice
\end{tabular} \\
\hline XXZChain2.logger & class attribute. \\
\hline XXZChain2.verbose &
\end{tabular}
class tenpy.models.xxz_chain. XxzChain2 (model_params)
Bases: tenpy.models.model.CouplingMPOModel, tenpy.models.model.
NearestNeighborModel
Another implementation of the Spin-1/2 XXZ chain with Sz conservation.
This implementation takes the same parameters as the XXZChain, but is implemented based on the CouplingMPOModel.

Parameters model_params (dictlConfig) - See XXZChain
init_sites (model_params)
Define the local Hilbert space and operators; needs to be implemented in subclasses.

This function gets called by init_lattice () to get the Site for the lattice unit cell.

Note: Initializing the sites requires to define the conserved quantum numbers. All pre-defined sites accept conserve=None to disable using quantum numbers. Many models in TeNPy read out the conserve model parameter, which can be set to "best" to indicate the optimal parameters.

If you need to initialize more than one site, the function tenpy.networks.site. set_common_charges () should be helpful.

Parameters model_params (dict) - The model parameters given to \(\qquad\)
\(\qquad\)
Returns sites - The local sites of the lattice, defining the local basis states and operators.
Return type (tuple of) Site
init_terms (model_params)
Add the onsite and coupling terms to the model; subclasses should implement this.
add_coupling (strength, u1, op1, u2, op2, dx, op_string=None, str_on_first=True, raise_op2_left=False, category=None, plus_hc=False)
Add twosite coupling terms to the Hamiltonian, summing over lattice sites.
Represents couplings of the form \(\sum_{x_{0}, \ldots, x_{d i m-1}} \operatorname{strength}[\operatorname{shift}(\vec{x})] * O P 0 * O P 1\), where opo := lat.unit_cell[u0].get_op(op0) acts on the site (x_0, ..., \(\left.x_{-}\{d i m-1\}, u 1\right)\), and OP1 := lat.unit_cell[u1].get_op(op1) acts on the site (x_0+dx[0], ..., \(x_{-}\{\operatorname{dim}-1\}+d x[\operatorname{dim}-1]\), u1). Possible combinations \(x_{-} 0, \ldots, x_{-}\{d i m-1\}\) are determined from the boundary conditions in possible_couplings ().
The coupling strength may vary spatially if the given strength is a numpy array. The correct shape of this array is the coupling_shape returned by tenpy.models.lattice.coupling_shape() and depends on the boundary conditions. The shift (...) depends on \(d x\), and is chosen such that the first entry strength \([0,0, \ldots]\) of strength is the prefactor for the first possible coupling fitting into the lattice if you imagine open boundary conditions.

The necessary terms are just added to coupling_terms; this function does not rebuild the MPO.
Deprecated since version 0.4.0: The arguments str_on_first and raise_op2_left will be removed in version 1.0.0.

\section*{Parameters}
- strength (scalar | array) - Prefactor of the coupling. May vary spatially (see above). If an array of smaller size is provided, it gets tiled to the required shape.
- u1 (int) - Picks the site lat. unit_cell [u1] for OP1.
- op1 (str) - Valid operator name of an onsite operator in lat.unit_cell[u1] for OP1.
- u2 (int) - Picks the site lat. unit_cell[u2] for OP2.
- op2 (str) - Valid operator name of an onsite operator in lat.unit_cell[u2] for OP2.
- \(\mathbf{d x}\) (iterable of int) - Translation vector (of the unit cell) between OP1 and OP2. For a 1D lattice, a single int is also fine.
- op_string (str | None) - Name of an operator to be used between the OP1 and OP2 sites. Typical use case is the phase for a Jordan-Wigner transformation. The operator should be defined on all sites in the unit cell. If None, auto-determine whether a JordanWigner string is needed, using op_needs_JW ().
- str_on_first (bool) - Whether the provided op_string should also act on the first site. This option should be chosen as True for Jordan-Wigner strings. When handling Jordan-Wigner strings we need to extend the op_string to also act on the 'left', first site (in the sense of the MPS ordering of the sites given by the lattice). In this case, there is a well-defined ordering of the operators in the physical sense (i.e. which of opl or op2 acts first on a given state). We follow the convention that \(o p 2\) acts first (in the physical sense), independent of the MPS ordering. Deprecated.
- raise_op2_left (bool) - Raise an error when op2 appears left of opl (in the sense of the MPS ordering given by the lattice). Deprecated.
- category (str) - Descriptive name used as key for coupling_terms. Defaults to a string of the form " \(\{0 \mathrm{p} 1\} \_i \quad\{o p 2\} \_j\) ".
- plus_hc (bool) - If True, the hermitian conjugate of the terms is added automatically.

\section*{Examples}

When initializing a model, you can add a term \(J \sum_{<i, j>} S_{i}^{z} S_{j}^{z}\) on all nearest-neighbor bonds of the lattice like this:
```

>>> J = 1. \# the strength
>>> for u1, u2, dx in self.lat.pairs['nearest_neighbors']:
... self.add_coupling(J, u1, 'Sz', u2, 'Sz', dx)

```

The strength can be an array, which gets tiled to the correct shape. For example, in a 1D Chain with an even number of sites and periodic (or infinite) boundary conditions, you can add alternating strong and weak couplings with a line like:
```

>>> self.add_coupling([1.5, 1.], u1, 'Sz', u2, 'Sz', dx)

```

Make sure to use the plus_hc argument if necessary, e.g. for hoppings:
```

>>> t = 1. \# hopping strength
>>> for u1, u2, dx in self.lat.pairs['nearest_neighbors']:
self.add_coupling(t, u1, 'Cd', u2, 'C', dx, plus_hc=True)

```

Alternatively, you can add the hermitian conjugate terms explictly. The correct way is to complex conjugate the strength, take the hermitian conjugate of the operators and swap the order (including a swap \(u 1<->u 2\) ), and use the opposite direction -dx , i.e. the h.c. of add_coupling ( \(t, \mathrm{u} 1\), 'A', u2, 'B', dx) is add_coupling(np.conj(t), u2, hc('B'), u1, hc('A'), -dx), where \(h c\) takes the hermitian conjugate of the operator names, see \(g e t \_h c \_o p \_n a m e()\). For spin-less fermions (FermionSite), this would be
```

>>> for u1, u2, dx in self.lat.pairs['nearest_neighbors']:
... self.add_coupling(t, u1, 'Cd', u2, 'C', dx)
... self.add_coupling(np.conj(t), u2, 'Cd', ul, 'C', -dx) \# h.c.

```

With spin-full fermions (SpinHalfFermions), it could be:
```

>>> for u1, u2, dx in self.lat.pairs['nearest_neighbors']:
... self.add_coupling(t, u1, 'Cdu', u2, 'Cd', dx) \# Cdagger_up C_down
... self.add_coupling(np.conj(t), u2, 'Cdd', u1, 'Cu', -dx) \# h.c.ь
Cdagger_down C_up

```

Note that the Jordan-Wigner strings for the fermions are added automatically!

\section*{See also:}
add_onsite Add terms acting on one site only.
add_multi_coupling_term for terms on more than two sites.
add_coupling_term Add a single term without summing over \(\vec{x}\).
add_coupling_term (strength, \(i, j, o p_{-} i, o p_{-} j, o p_{-}\)string \(=' I d^{\prime}\), category=None, plus_hc=False)
Add a two-site coupling term on given MPS sites.
Wrapper for self.coupling_terms[category].add_coupling_term(...).

\section*{Warning: This function does not handle Jordan-Wigner strings! You might want to use add_local_term() instead.}

\section*{Parameters}
- strength (float) - The strength of the coupling term.
- i (int) - The MPS indices of the two sites on which the operator acts. We require 0 \(<=i<N \_s i t e s\) and \(i<j\), i.e., op_i acts "left" of \(o p_{-} j\). If \(j>=N_{-}\)sites, it indicates couplings between unit cells of an infinite MPS.
- j (int) - The MPS indices of the two sites on which the operator acts. We require 0 \(<=i<N \_s i t e s\) and \(i<j\), i.e., op_i acts "left" of \(o p_{-} j\). If \(\mathrm{j}>=\mathrm{N}_{-}\)sites, it indicates couplings between unit cells of an infinite MPS.
- op1 (str) - Names of the involved operators.
- op2 (str) - Names of the involved operators.
- op_string (str) - The operator to be inserted between \(i\) and \(j\).
- category (str) - Descriptive name used as key for coupling_terms. Defaults to a string of the form " \(\{0 \mathrm{p} 1\} \_i \quad\{o p 2\} \_j\) ".
- plus_hc (bool) - If True, the hermitian conjugate of the term is added automatically.
add_exponentially_decaying_coupling (strength, lambda_, op_i, op_j, subsites=None, op_string \(=\) None, plus_hc \(^{\text {F False }}\) )
Add an exponentially decaying long-range coupling.
\[
\text { strength } \sum_{i<j} \lambda^{|i-j|} A_{\text {subsites }[i]} B_{\text {subsites }[j]}
\]

Where the operator \(A\) is given by \(o p_{-} i\), and \(B\) is given by \(o p_{-} j\). Note that the sum over \(i, j\) is long-range, for infinite systems going beyond the MPS unit cell. Moreover, note that the distance in the exponent is the distance within subsites.

\section*{Parameters}
- strength (float) - Overall prefactor.
- lambda (float) - Decay-rate
- op_i (string) - Names for the operators.
- op_j (string) - Names for the operators.
- subsites (None / 1D array) - Selects a subset of sites within the MPS unit cell on which the operators act. Needs to be sorted. None selects all sites.
- op_string (None / str) - The operator to be inserted between \(A\) and \(B\); If None, this function checks whether a fermionic "JW" string is needed for the given operators; in this case the right \(o p_{-} j\) acts first.
- plus_hc (bool) - If True, the hermitian conjugate of the term is added automatically.

\section*{Examples}

At least for simple enough 1D chains (or ladders), you can use fit_with_sum_of_exp () to approximate a long-range function with a few sum of exponentials and then add them with this function.
```

>>> def decay(x):
return np.exp(-0.1*x) / x**2
>>> from tenpy.tools.fit import fit_with_sum_of_exp, sum_of_exp
>>> n_exp = 5
>>> fit_range = 50
>>> lam, pref = fit_with_sum_of_exp(decay, n_exp, fit_range)
>>> x = np.arange(1, fit_range + 1)
>>> print('error in fit: {0:.3e}'.format(np.sum(np.abs(decay(x) - sum_of__
@exp(lam, pref, x)))))
error in fit: 1.073e-04
>>> for pr, la in zip(pref, lam):
... self.add_exponentially_decaying_coupling(pr, la, 'N', 'N')

```
add_local_term (strength, term, category=None, plus_hc=False)
Add a single term to self.
The repesented term is strength times the product of the operators given in terms. Each operator is specified by the name and the site it acts on; the latter given by a lattice index, see Lattice.

Depending on the length of term, it can add an onsite term or a coupling term to onsite_terms or coupling_terms, respectively.

\section*{Parameters}
- strength (float/complex) - The prefactor of the term.
- term (list of (str, array_like))-List of tuples (opname, lat_idx) where opname is a string describing the operator acting on the site given by the lattice index lat_idx. Here, lat_idx is for example \([x, y, u]\) for a 2D lattice, with \(u\) being the index within the unit cell.
- category - Descriptive name used as key for onsite_terms or coupling_terms.
- plus_hc (bool) - If True, the hermitian conjugate of the terms is added automatically.
add_multi_coupling (strength, ops, _deprecate_l='DEPRECATED', _depre-
cate_2 = 'DEPRECATED', op_string=None, category=None, plus_hc=False)
Add multi-site coupling terms to the Hamiltonian, summing over lattice sites.
Represents couplings of the form \(\operatorname{sum}_{\vec{x}} \operatorname{strength}[\operatorname{shift}(\vec{x})] * O P_{0} * O P_{1} * \ldots * O P_{M-1}\), involving \(M\) operators. Here, \(O P_{m}\) stands for the operator defined by the \(m\)-th tuple (opname, \(\mathrm{dx}, \mathrm{u}\) ) given in the argument ops, which determines the position \(\vec{x}+\overrightarrow{d x}\) and unit-cell index \(u\) of the site it acts on; the actual operator is given by self.lat.unit_cell[u].get_op(opname).

The coupling strength may vary spatially if the given strength is a numpy array. The correct shape of this array is the coupling_shape returned by tenpy.models.lattice. possible_multi_couplings() and depends on the boundary conditions. The shift(...) depends on the \(d x\) entries of ops and is chosen such that the first entry strength \([0,0, \ldots]\) of strength
is the prefactor for the first possible coupling fitting into the lattice if you imagine open boundary conditions.

The necessary terms are just added to coupling_terms; this function does not rebuild the MPO.
Deprecated since version 0.6.0: We switched from the three arguments \(u 0\), op0 and other_op with other_ops=[(u1, op1, dx1), (op2, u2, dx2), ...] to a single, equivalent argment ops which should now read ops \(=[(\mathrm{op} 0, \mathrm{dx} 0, \mathrm{u} 0)\), (op1, dx1, u1), (op2, dx2, u2), . ..], where \(d x 0=[0] *\) self.lat.dim. Note the changed order inside the tuples!

\section*{Parameters}
- strength (scalar / array) - Prefactor of the coupling. May vary spatially, and is tiled to the required shape.
- ops (list of (opname, \(d x, u\) ) ) - Each tuple determines one operator of the coupling, see the description above. opname (str) is the name of the operator, \(d x\) (list of length lat.dim) is a translation vector, and \(u\) (int) is the index of lat.unit_cell on which the operator acts. The first entry of ops corresponds to \(O P_{0}\) and acts last in the physical sense.
- op_string (str / None) - If a string is given, we use this as the name of an operator to be used inbetween the operators, excluding the sites on which any operators act. This operator should be defined on all sites in the unit cell.

If None, auto-determine whether a Jordan-Wigner string is needed (using op_needs_JW ()) for each of the segments inbetween the operators and also on the sites of the left operators.
- category (str) - Descriptive name used as key for coupling_terms. Defaults to a string of the form " \(\{0 \mathrm{p} 0\}\) _i \(\left\{o t h e r \_o p s[0]\right\} \_j\) \{other_ops[1]\}_k ...".
- plus_hc (bool) - If True, the hermitian conjugate of the terms is added automatically.

\section*{Examples}

A call to add_coupling() with arguments add_coupling(strength, u1, 'A', u2, ' \(B\) ', \(d x\) ) is equivalent to the following:
```

>>>dx_0 = [0] * self.lat.dim \# = [0] for a ID lattice, [0, 0] in 2D
>> self.add_multi_coupling(strength, [('A', dx_0, u1), ('B', dx, u2)])

```

To explicitly add the hermitian conjugate (instead of simply using plus_hc = True), you need to take the complex conjugate of the strength, reverse the order of the operators and take the hermitian conjugates of the individual operator names (indicated by the hc (...), see get_hc_op_name ()):
```

>>> self.add_multi_coupling(np.conj(strength), [(hc('B'), dx, u2), (hc('A'), r

```
\(\left.\left.\rightarrow d x \_0, u 1\right)\right]\) )

\section*{See also:}
add_onsite Add terms acting on one site only.
add_coupling Add terms acting on two sites.
add_multi_coupling_term Add a single term, not summing over the possible \(\vec{x}\).
add_multi_coupling_term (strength, ijkl, ops_ijkl,op_string, category=None, plus_hc=False)
Add a general M-site coupling term on given MPS sites.
Wrapper for self.coupling_terms[category].add_multi_coupling_term(...).

> Warning: This function does not handle Jordan-Wigner strings! You might want to use add_local_term() instead.

\section*{Parameters}
- strength (float) - The strength of the coupling term.
- ijkl (list of int) - The MPS indices of the sites on which the operators acts. With \(i, j, k, \ldots=i j k l\), we require that they are ordered ascending, \(i<j<k<\ldots\) and that \(0<=i<N \_s i t e s\). Inidces \(>=\) N_sites indicate couplings between different unit cells of an infinite MPS.
- ops_ijkl (list of str) - Names of the involved operators on sites \(i, j, k, \ldots\).
- op_string (list of str) - Names of the operator to be inserted between the operators, e.g., op_string[0] is inserted between \(i\) and \(j\).
- category (str) - Descriptive name used as key for coupling_terms. Defaults to a string of the form " \(\{o p 0\} \_i \quad\{o p 1\} \_j \quad\{o p 2\} \_k \quad . . . "\).
- plus_hc (bool) - If True, the hermitian conjugate of the term is added automatically.
add_onsite (strength, u, opname, category=None, plus_hc=False)
Add onsite terms to onsite_terms.
Adds \(\sum_{\vec{x}}\) strength \([\vec{x}] * O P\) to the represented Hamiltonian, where the operator \(O P=1\) at. unit_cell[u].get_op (opname) acts on the site given by a lattice index (x_0, ..., \(\left.x_{-}\{\operatorname{dim}-1\}, u\right)\),
The necessary terms are just added to onsite_terms; doesn't rebuild the MPO.

\section*{Parameters}
- strength (scalar / array) - Prefactor of the onsite term. May vary spatially. If an array of smaller size is provided, it gets tiled to the required shape.
- u(int) - Picks a Site lat.unit_cell[u] out of the unit cell.
- opname (str) - valid operator name of an onsite operator in lat.unit_cell [u].
- category (str) - Descriptive name used as key for onsite_terms. Defaults to opname.
- plus_hc (bool) - If True, the hermitian conjugate of the terms is added automatically.

\section*{See also:}
add_coupling Add a terms acting on two sites.
add_onsite_term Add a single term without summing over vecx.
add_onsite_term (strength, \(i\), op, category=None, plus_hc=False)
Add an onsite term on a given MPS site.
Wrapper for self.onsite_terms[category].add_onsite_term(...).

\section*{Parameters}
- strength (float) - The strength of the term.
- i (int) - The MPS index of the site on which the operator acts. We require \(0<=i<\) L.
- op (str) - Name of the involved operator.
- category (str) - Descriptive name used as key for onsite_terms. Defaults to op.
- plus_hc (bool) - If True, the hermitian conjugate of the term is added automatically.
all_coupling_terms()
Sum of all coupling_terms.
all_onsite_terms()
Sum of all onsite_terms.
bond_energies (psi)
Calculate bond energies <psilH_bondlpsi>.
Parameters psi (MPS) - The MPS for which the bond energies should be calculated.
Returns E_bond - List of bond energies: for finite bc, E_Bond [i] is the energy of bond i, i+1. (i.e. we omit bond 0 between sites L-1 and 0); for infinite bc E_bond[i] is the energy of bond \(i-1, i\).
Return type 1D ndarray
calc_H_MPO (tol_zero \(=1 e-15\) )
Calculate MPO representation of the Hamiltonian.
Uses onsite_terms and coupling_terms to build an MPOGraph (and then an MPO).
Parameters tol_zero (float) - Prefactors with abs(strength) < tol_zero are considered to be zero.
Returns H_MPO - MPO representation of the Hamiltonian.
Return type MPO
calc_H_MPO_from_bond (tol_zero=le-15)
Calculate the MPO Hamiltonian from the bond Hamiltonian.
Parameters tol_zero (float) - Arrays with norm <tol_zero are considered to be zero.
Returns H_MPO - MPO representation of the Hamiltonian.
Return type MPO
calc_H_bond (tol_zero=le-15)
calculate \(H \_b o n d\) from coupling_terms and onsite_terms.
Parameters tol_zero (float) - prefactors with abs(strength) < tol_zero are considered to be zero.
Returns H_bond - Bond terms as required by the constructor of NearestNeighborModel. Legs are ['p0', 'p0*', 'p1', 'p1*']
Return type list of Array
:raises ValueError : if the Hamiltonian contains longer-range terms.:
calc_H_bond_from_MPO (tol_zero=le-15)
Calculate the bond Hamiltonian from the MPO Hamiltonian.
Parameters tol_zero (float) - Arrays with norm < tol_zero are considered to be zero.
Returns H_bond - Bond terms as required by the constructor of NearestNeighborModel. Legs are ['p0', 'p0*', 'p1', 'p1*']
Return type list of Array
:raises ValueError : if the Hamiltonian contains longer-range terms.:
calc_H_onsite (tol_zero=le-15)
Calculate H_onsite from self.onsite_terms.
Deprecated since version 0.4.0: This function will be removed in 1.0.0. Replace calls to this function by self.all_onsite_terms().remove_zeros(tol_zero).to_Arrays(self. lat.mps_sites()). You might also want to take explicit_plus_hc into account.

Parameters tol_zero (float) - prefactors with abs (strength) < tol_zero are considered to be zero.

\section*{Returns}

\section*{- H_onsite (list of npc.Array)}
- onsite terms of the Hamiltonian. If explicit_plus_hc is True, - Hermitian conjugates of the onsite terms will be included.
coupling_strength_add_ext_flux (strength, \(d x\), phase)
Add an external flux to the coupling strength.
When performing DMRG on a "cylinder" geometry, it might be useful to put an "external flux" through the cylinder. This means that a particle hopping around the cylinder should pick up a phase given by the external flux [[resta1998]]. This is also called "twisted boundary conditions" in literature. This function adds a complex phase to the strength array on some bonds, such that particles hopping in positive direction around the cylinder pick up \(\exp (+i\) phase \()\).

Warning: For the sign of phase it is important that you consistently use the creation operator as opl and the annihilation operator as op2 in add_coupling().

\section*{Parameters}
- strength (scalar / array) - The strength to be used in add_coupling(), when no external flux would be present.
- \(\mathbf{d x}\) (iterable of int) - Translation vector (of the unit cell) between opl and op2 in add_coupling().
- phase (iterable of float) - The phase of the external flux for hopping in each direction of the lattice. E.g., if you want flux through the cylinder on which you have an infinite MPS, you should give phase \(=[0\), phi] souch that particles pick up a phase phi when hopping around the cylinder.
Returns strength - The strength array to be used as strength in add_coupling() with the given \(d x\).

\section*{Return type complex array}

\section*{Examples}

Let's say you have an infinite MPS on a cylinder, and want to add nearest-neighbor hopping of fermions with the FermionSite. The cylinder axis is the \(x\)-direction of the lattice, so to put a flux through the cylinder, you want particles hopping around the cylinder to pick up a phase phi given by the external flux.
```

>>> strength = 1. \# hopping strength without external flux
>>> phi = np.pi/4 \# determines the external flux strength
>>> for u1, u2, dx in self.lat.pairs['nearest_neighbors']:

```
```

... strength_with_flux = self.coupling_strength_add_ext_flux(strength, dx,
@ [0, phi])
self.add_coupling(strength_with_flux, u1, 'Cd', u2, 'C', dx)
... self.add_coupling(np.conj(strength_with_flux), u2, 'Cd', u1, 'C', -dx)

```
enlarge_mps_unit_cell (factor=2)
Repeat the unit cell for infinite MPS boundary conditions; in place.
This has to be done after finishing initialization and can not be reverted.
Parameters factor (int) - The new number of sites in the MPS unit cell will be increased from \(N\) _sites to factor*N_sites_per_ring. Since MPS unit cells are repeated in the \(x\)-direction in our convetion, the lattice shape goes from ( \(L x, L y, \ldots, L u\) ) to (Lx*factor, Ly, ..., Lu).
classmethod from_MPOModel (mpo_model)
Initialize a NearestNeighborModel from a model class defining an MPO.
This is especially usefull in combination with MPOModel.group_sites ().
Parameters mpo_model (MPOModel) - A model instance implementing the MPO. Does not need to be a NearestNeighborModel, but should only have nearest-neighbor couplings.

\section*{Examples}

The SpinChainNNN2 has next-nearest-neighbor couplings and thus only implements an MPO:
```

>>> from tenpy.models.spins_nnn import SpinChainNNN2
>>> nnn_chain = SpinChainNNN2({'L': 20})
>>> print(isinstance(nnn_chain, NearestNeighborModel))
False
>>> print("range before grouping:", nnn_chain.H_MPO.max_range)
range before grouping: 2

```

By grouping each two neighboring sites, we can bring it down to nearest neighbors.
```

>>> grouped_sites = nnn_chain.group_sites(2)
>>> print("range after grouping:", nnn_chain.H_MPO.max_range)
range after grouping: 1

```

Yet, TEBD will not yet work, as the model doesn't define \(H_{\_}\)bond. However, we can initialize a NearestNeighborModel from the MPO:
```

>>> nnn_chain_for_tebd = NearestNeighborModel.from_MPOModel(nnn_chain)
>>> isinstance(nnn_chain_for_tebd, NearestNeighborModel)
True

```
classmethod from_hdf5 (hdf5_loader, h5gr, subpath)
Load instance from a HDF5 file.
This method reconstructs a class instance from the data saved with save_hdf5 ().

\section*{Parameters}
- hdf5_loader (Hdf5Loader) - Instance of the loading engine.
- h5gr (Group) - HDF5 group which is represent the object to be constructed.
- subpath (str) - The name of \(h 5 g r\) with a ' /' in the end.

Returns obj - Newly generated class instance containing the required data.
Return type cls
group_sites ( \(n=2\), grouped_sites=None)
Modify self in place to group sites.
Group each \(n\) sites together using the GroupedSite. This might allow to do TEBD with a Trotter decomposition, or help the convergence of DMRG (in case of too long range interactions).

This has to be done after finishing initialization and can not be reverted.

\section*{Parameters}
- \(\mathbf{n}(i n t)\) - Number of sites to be grouped together.
- grouped_sites (None l list of GroupedSite) - The sites grouped together.

Returns grouped_sites - The sites grouped together.
Return type list of GroupedSite

\section*{init_H_from_terms()}

Initialize \(H_{-} M P O\) (and \(H_{-}\)bond) from the terms of the CouplingModel.
This function is called automatically during CouplingMPOModel.__init__.
If you use one of the add_* methods of the CouplingModel after initialization, you will need to call init_H_from_terms in the end by yourself, in order to update the \(H_{-} M P O\) (and possibly \(H_{-} b o n d\) ) representations. (You should get a warning about this... The way to avoid it is to initialize all the terms in init_terms by defining your own model, as outlined in Models.

\section*{init_lattice (model_params)}

Initialize a lattice for the given model parameters.
This function reads out the model parameter lattice. This can be a full Lattice instance, in which case it is just returned without further action. Alternatively, the lattice parameter can be a string giving the name of one of the predefined lattices, which then gets initialized. Depending on the dimensionality of the lattice, this requires different model parameters.

Parameters model_params (dict) - The model parameters given to __init__
Returns lat - An initialized lattice.
Return type Lattice

\section*{Options}
option CouplingMPOModel.lattice: str | Lattice
The name of a lattice pre-defined in TeNPy to be initialized. Alternatively, directly a subclass of Lattice instead of the name. Alternatively, a (possibly self-defined) Lattice instance. In the latter case, no further parameters are read out.
option CouplingMPOModel.bc_MPS: str Boundary conditions for the MPS.
option CouplingMPOModel.order: str
The order of sites within the lattice for non-trivial lattices, e.g, 'default', 'snake', see ordering(). Only used if lattice is a string.
option CouplingMPOModel.L: int
The length in x-direction; only read out for 1D lattices. For an infinite system the length of the unit cell.
option CouplingMPOModel.Lx: int
option CouplingMPOModel.Ly: int
The length in x - and y -direction; only read out for 2D lattices. For "infinite" \(b c_{-} M P S\), the system is infinite in x-direction and \(L x\) is the number of "rings" in the infinite MPS unit cell, while \(L y\) gives the circumference around the cylinder or width of th the rung for a ladder (depending on \(b c_{-} y\) ).
option CouplingMPOModel.bc_y: str
"cylinder" | "ladder"; only read out for 2D lattices. The boundary conditions in y-direction.
option CouplingMPOModel.bc_x: str
"open" | "periodic". Can be used to force "periodic" boundaries for the lattice, i.e., for the couplings in the Hamiltonian, even if the MPS is finite. Defaults to "open" for bc_MPS="finite " and "periodic" for bc_MPS="infinite. If you are not aware of the consequences, you should probably not use "periodic" boundary conditions. (The MPS is still "open", so this will introduce long-range couplings between the first and last sites of the MPS!)
save_hdf5 (hdf5_saver, h5gr, subpath)
Export self into a HDF5 file.
This method saves all the data it needs to reconstruct self with \(f r o m \_h d f 5\) ().
This implementation saves the content of __dict__ with save_dict_content (), storing the format under the attribute 'format'.

\section*{Parameters}
- hdf5_saver (Hdf5Saver) - Instance of the saving engine.
- h5gr (: class `Group ) - HDF5 group which is supposed to represent self.
- subpath (str) - The name of \(h 5 g r\) with a '/' in the end.

\section*{test_sanity ()}

Sanity check, raises ValueErrors, if something is wrong.
trivial_like_NNModel()
Return a NearestNeighborModel with same lattice, but trivial \((\mathrm{H}=0)\) bonds.

\section*{Module description}

Prototypical example of a 1D quantum model: the spin-1/2 XXZ chain.
The XXZ chain is contained in the more general SpinChain; the idea of this module is more to serve as a pedagogical example for a model.

\section*{19.5 spins}
- full name: tenpy.models.spins
- parent module: tenpy.models
- type: module

\section*{Classes}

\begin{tabular}{ll}
\hline SpinChain(model_params) & The SpinModel on a Chain, suitable for TEBD. \\
\hline SpinModel(model_params) & Spin-S sites coupled by nearest neighbour interactions. \\
\hline
\end{tabular}

\subsection*{19.5.1 SpinChain}
- full name: tenpy.models.spins.SpinChain
- parent module: tenpy.models.spins
- type: class

\section*{Inheritance Diagram}


\section*{Methods}
\begin{tabular}{|c|c|}
\hline Spinchain.__init__(model_params) & Initialize self. \\
\hline SpinChain.add_coupling(strength, u1, op1,... & Add twosite coupling terms to the Hamiltonian, summing over lattice sites. \\
\hline ```
SpinChain.add_coupling_term(strength, i, j,
...)
``` & dd a two-site coupling term on given MPS sites. \\
\hline \multicolumn{2}{|l|}{SpinChain.add_exponentially_decaying_coAdd án ¢xponentially decaying long-range coupling.} \\
\hline SpinChain.add_local_term(strength, term[, ...]) & Add a single term to self. \\
\hline SpinChain.add_multi_coupling(strength, ops) & Add multi-site coupling terms to the Hamiltonian, summing over lattice sites. \\
\hline \multicolumn{2}{|l|}{SpinChain.add_multi_coupling_term(strengthAdd a general M-site coupling term on given MPS sites. ...)} \\
\hline SpinChain.add_onsite(strength, u, opname[, ...]) & Add onsite terms to onsite_terms. \\
\hline
\end{tabular}

Table 45 - continued from previous page
\begin{tabular}{|c|c|}
\hline SpinChain.add_onsite_term(strength, i, op[,
...]) & Add an onsite term on a given MPS site. \\
\hline SpinChain.all_coupling_terms() & Sum of all coupling_terms. \\
\hline SpinChain.all_onsite_terms() & Sum of all onsite_terms. \\
\hline SpinChain.bond_energies(psi) & Calculate bond energies <psilH_bondlpsi>. \\
\hline SpinChain.calc_H_MPO([tol_zero]) & Calculate MPO representation of the Hamiltonian. \\
\hline SpinChain.calc_H_MPO_from_bond([tol_zero]) & Calculate the MPO Hamiltonian from the bond Hamiltonian. \\
\hline SpinChain.calc_H_bond([tol_zero]) & calculate \(H_{\text {_bond }}\) from coupling_terms and onsite_terms. \\
\hline SpinChain.calc_H_bond_from_MPO([tol_zero]) & Calculate the bond Hamiltonian from the MPO Hamiltonian. \\
\hline SpinChain.calc_H_onsite([tol_zero]) & Calculate H_onsite from self.onsite_terms. \\
\hline SpinChain.coupling_strength_add_ext_flut & \(u\) A(dd.a)n external flux to the coupling strength. \\
\hline SpinChain.enlarge_mps_unit_cell([factor]) & Repeat the unit cell for infinite MPS boundary conditions; in place. \\
\hline SpinChain.from_MPOModel(mpo_model) & Initialize a NearestNeighborModel from a model class defining an MPO. \\
\hline SpinChain.from_hdf5(hdf5_loader, h5gr, subpath) & Load instance from a HDF5 file. \\
\hline SpinChain.group_sites([n, grouped_sites]) & Modify self in place to group sites. \\
\hline SpinChain.init_H_from_terms() & Initialize \(H_{-} M P O\) (and \(H \_b o n d\) ) from the terms of the CouplingModel. \\
\hline SpinChain.init_lattice(model_params) & Initialize a lattice for the given model parameters. \\
\hline SpinChain.init_sites(model_params) & Define the local Hilbert space and operators; needs to be implemented in subclasses. \\
\hline SpinChain.init_terms(model_params) & Add the onsite and coupling terms to the model; subclasses should implement this. \\
\hline SpinChain.save_hdf5(hdf5_saver, h5gr, subpath) & Export self into a HDF5 file. \\
\hline SpinChain.test_sanity() & Sanity check, raises ValueErrors, if something is wrong. \\
\hline SpinChain.trivial_like_NNModel() & Return a NearestNeighborModel with same lattice, but trivial ( \(\mathrm{H}=0\) ) bonds. \\
\hline
\end{tabular}

\section*{Class Attributes and Properties}
\begin{tabular}{ll}
\hline SpinChain.force_default_lattice & \begin{tabular}{l} 
If True, init_lattice () asserts that the initialized \\
lattice is (a subclass of) default_lattice
\end{tabular} \\
\hline SpinChain.logger & class attribute. \\
\hline SpinChain.verbose &
\end{tabular}
class tenpy.models.spins.SpinChain (model_params)
Bases: tenpy.models.spins.SpinModel, tenpy.models.model.NearestNeighborModel
The SpinModel on a Chain, suitable for TEBD.
See the SpinModel for the documentation of parameters.

\section*{default_lattice}
alias of tenpy.models.lattice.Chain
add_coupling (strength, u1, op1, u2, op2, dx, op_string=None, str_on_first=True, raise_op2_left=False, category=None, plus_hc=False)
Add twosite coupling terms to the Hamiltonian, summing over lattice sites.
Represents couplings of the form \(\sum_{x_{0}, \ldots, x_{\text {dim-1 }}} \operatorname{strength}[\operatorname{shift}(\vec{x})] * O P 0 * O P 1\), where OP 0 := lat.unit_cell[u0].get_op(op0) acts on the site (x_0, ..., x_\{dim-1\}, u1), and OP1 := lat.unit_cell[u1].get_op(op1) acts on the site (x_0+dx[0], ..., \(x_{\_}\{d i m-1\}+d x[\operatorname{dim}-1]\), \(\left.u 1\right)\). Possible combinations \(x_{-} 0, \ldots, x_{-}\{d i m-1\}\) are determined from the boundary conditions in possible_couplings ().

The coupling strength may vary spatially if the given strength is a numpy array. The correct shape of this array is the coupling_shape returned by tenpy.models.lattice.coupling_shape () and depends on the boundary conditions. The shift (...) depends on \(d x\), and is chosen such that the first entry strength [0, 0, ...] of strength is the prefactor for the first possible coupling fitting into the lattice if you imagine open boundary conditions.

The necessary terms are just added to coupling_terms; this function does not rebuild the MPO.
Deprecated since version 0.4.0: The arguments str_on_first and raise_op2_left will be removed in version 1.0.0.

\section*{Parameters}
- strength (scalar / array) - Prefactor of the coupling. May vary spatially (see above). If an array of smaller size is provided, it gets tiled to the required shape.
- u1 (int) - Picks the site lat. unit_cell[u1] for OP1.
- op1 (str) - Valid operator name of an onsite operator in lat. unit_cell[u1] for OP1.
- u2 (int) - Picks the site lat. unit_cell [u2] for OP2.
- op2 (str) - Valid operator name of an onsite operator in lat.unit_cell[u2] for OP2.
- \(\mathbf{d x}\) (iterable of int) - Translation vector (of the unit cell) between OP1 and OP2. For a 1D lattice, a single int is also fine.
- op_string (str / None) - Name of an operator to be used between the OP1 and OP2 sites. Typical use case is the phase for a Jordan-Wigner transformation. The operator should be defined on all sites in the unit cell. If None, auto-determine whether a JordanWigner string is needed, using op_needs_JW().
- str_on_first (bool) - Whether the provided op_string should also act on the first site. This option should be chosen as True for Jordan-Wigner strings. When handling Jordan-Wigner strings we need to extend the op_string to also act on the 'left', first site (in the sense of the MPS ordering of the sites given by the lattice). In this case, there is a well-defined ordering of the operators in the physical sense (i.e. which of opl or op2 acts first on a given state). We follow the convention that op2 acts first (in the physical sense), independent of the MPS ordering. Deprecated.
- raise_op2_left (bool) - Raise an error when op2 appears left of opl (in the sense of the MPS ordering given by the lattice). Deprecated.
- category (str) - Descriptive name used as key for coupling_terms. Defaults to a string of the form " \(\{o p 1\} \_i \quad\{o p 2\} \_j "\).
- plus_hc (bool) - If True, the hermitian conjugate of the terms is added automatically.

\section*{Examples}

When initializing a model, you can add a term \(J \sum_{<i, j>} S_{i}^{z} S_{j}^{z}\) on all nearest-neighbor bonds of the lattice like this:
```

>>> J = 1. \# the strength
>>> for u1, u2, dx in self.lat.pairs['nearest_neighbors']:
... self.add_coupling(J, u1, 'Sz', u2, 'Sz', dx)

```

The strength can be an array, which gets tiled to the correct shape. For example, in a 1D Chain with an even number of sites and periodic (or infinite) boundary conditions, you can add alternating strong and weak couplings with a line like:
```

>>> self.add_coupling([1.5, 1.], u1, 'Sz', u2, 'Sz', dx)

```

Make sure to use the plus_hc argument if necessary, e.g. for hoppings:
```

>>> t = 1. \# hopping strength
>>> for u1, u2, dx in self.lat.pairs['nearest_neighbors']:
... self.add_coupling(t, ul, 'Cd', u2, 'C', dx, plus_hc=True)

```

Alternatively, you can add the hermitian conjugate terms explictly. The correct way is to complex conjugate the strength, take the hermitian conjugate of the operators and swap the order (including a swap \(u 1<->u 2\) ), and use the opposite direction \(-d x\), i.e. the h.c. of add_coupling ( \(t, \mathrm{u} 1\), 'A', u2, 'B', dx) is add_coupling(np.conj(t), u2, hc('B'), u1, hc('A'), -dx), where \(h c\) takes the hermitian conjugate of the operator names, see get_hc_op_name (). For spin-less fermions (FermionSite), this would be
```

>>> for ul, u2, dx in self.lat.pairs['nearest_neighbors']:
... self.add_coupling(t, u1, 'Cd', u2, 'C', dx)
... self.add_coupling(np.conj(t), u2, 'Cd', u1, 'C', -dx) \# h.c.

```

With spin-full fermions (SpinHalfFermions), it could be:
```

>>> for u1, u2, dx in self.lat.pairs['nearest_neighbors']:
... self.add_coupling(t, u1, 'Cdu', u2, 'Cd', dx) \# Cdagger_up C_down
... self.add_coupling(np.conj(t), u2, 'Cdd', u1, 'Cu', -dx) \# h.c.u
Cdagger_down C_up

```

Note that the Jordan-Wigner strings for the fermions are added automatically!

\section*{See also:}
add_onsite Add terms acting on one site only.
add_multi_coupling_term for terms on more than two sites.
add_coupling_term Add a single term without summing over \(\vec{x}\).
add_coupling_term (strength, \(i, j, o p_{-} i, o p \_j\), op_string='Id', category=None, plus_hc=False)
Add a two-site coupling term on given MPS sites.
Wrapper for self.coupling_terms[category].add_coupling_term(...).

Warning: This function does not handle Jordan-Wigner strings! You might want to use add_local_term() instead.

\section*{Parameters}
- strength (float) - The strength of the coupling term.
- i (int) - The MPS indices of the two sites on which the operator acts. We require 0 <= i < N_sites and i < j, i.e., op_i acts "left" of op_j. If j >= N_sites, it indicates couplings between unit cells of an infinite MPS.
- \(\mathbf{j}\) (int) - The MPS indices of the two sites on which the operator acts. We require 0 <= i < N_sites and i < j, i.e., op_i acts "left" of \(o p_{-} j\). If j\(\rangle=\mathrm{N} \_\)sites, it indicates couplings between unit cells of an infinite MPS.
- op1 (str) - Names of the involved operators.
- op2 (str) - Names of the involved operators.
- op_string (str) - The operator to be inserted between \(i\) and \(j\).
- category (str) - Descriptive name used as key for coupling_terms. Defaults to a string of the form " \(\{\mathrm{op} 1\} \_\)i \(\{o p 2\} \_j\) ".
- plus_hc (bool) - If True, the hermitian conjugate of the term is added automatically.
add_exponentially_decaying_coupling (strength, lambda_, op_i, op_j, subsites=None, \(o p \_\)string \(=\)None, \(p l u s \_h c=\) False)
Add an exponentially decaying long-range coupling.
\[
\text { strength } \sum_{i<j} \lambda^{|i-j|} A_{\text {subsites }[i]} B_{\text {subsites }[j]}
\]

Where the operator \(A\) is given by op_i, and \(B\) is given by op_j. Note that the sum over \(\mathrm{i}, \mathrm{j}\) is long-range, for infinite systems going beyond the MPS unit cell. Moreover, note that the distance in the exponent is the distance within subsites.

\section*{Parameters}
- strength (float) - Overall prefactor.
- lambda (float) - Decay-rate
- op_i (string) - Names for the operators.
- op_j (string) - Names for the operators.
- subsites (None | ID array) - Selects a subset of sites within the MPS unit cell on which the operators act. Needs to be sorted. None selects all sites.
- op_string (None | str) - The operator to be inserted between \(A\) and \(B\); If None, this function checks whether a fermionic "JW" string is needed for the given operators; in this case the right \(o p_{-} j\) acts first.
- plus_hc (bool) - If True, the hermitian conjugate of the term is added automatically.

\section*{Examples}

At least for simple enough 1D chains (or ladders), you can use fit_with_sum_of_exp () to approximate a long-range function with a few sum of exponentials and then add them with this function.
```

>>> def decay(x):
... return np.exp (-0.1*x) / x**2
>>> from tenpy.tools.fit import fit_with_sum_of_exp, sum_of__exp
>>> n_exp = 5
>>> fit_range = 50
>>> lam, pref = fit_with_sum_of_exp(decay, n_exp, fit_range)

```
(continued from previous page)
```

>>> x = np.arange(1, fit_range + 1)
>>> print('error in fit: {0:.3e}'.format(np.sum(np.abs(decay(x) - sum_of_
\hookrightarrowexp(lam, pref, x)))))
error in fit: 1.073e-04
>>> for pr, la in zip(pref, lam):
self.add_exponentially_decaying_coupling(pr, la, 'N', 'N')

```
add_local_term (strength, term, category=None, plus_hc=False)
Add a single term to self.
The repesented term is strength times the product of the operators given in terms. Each operator is specified by the name and the site it acts on; the latter given by a lattice index, see Lattice.

Depending on the length of term, it can add an onsite term or a coupling term to onsite_terms or coupling_terms, respectively.

\section*{Parameters}
- strength (float/complex) - The prefactor of the term.
- term (list of (str, array_like)) - List of tuples (opname, lat_idx) where opname is a string describing the operator acting on the site given by the lattice index lat_idx. Here, lat_idx is for example \([x, y, u]\) for a 2D lattice, with \(u\) being the index within the unit cell.
- category - Descriptive name used as key for onsite_terms or coupling_terms.
- plus_hc (bool) - If True, the hermitian conjugate of the terms is added automatically.
add_multi_coupling (strength, ops, _deprecate_l='DEPRECATED', _deprecate_2 = 'DEPRECATED', op_string=None, category=None, plus_hc=False)
Add multi-site coupling terms to the Hamiltonian, summing over lattice sites.
Represents couplings of the form \(\operatorname{sum}_{\vec{x}} \operatorname{strength}[\operatorname{shift}(\vec{x})] * O P_{0} * O P_{1} * \ldots * O P_{M-1}\), involving \(M\) operators. Here, \(O P_{m}\) stands for the operator defined by the \(m\)-th tuple (opname, \(\mathrm{dx}, \mathrm{u}\) ) given in the argument ops, which determines the position \(\vec{x}+\overrightarrow{d x}\) and unit-cell index \(u\) of the site it acts on; the actual operator is given by self.lat.unit_cell[u].get_op(opname).

The coupling strength may vary spatially if the given strength is a numpy array. The correct shape of this array is the coupling_shape returned by tenpy.models.lattice. possible_multi_couplings() and depends on the boundary conditions. The shift(...) depends on the \(d x\) entries of ops and is chosen such that the first entry strength [0, 0, ...] of strength is the prefactor for the first possible coupling fitting into the lattice if you imagine open boundary conditions.

The necessary terms are just added to coupling_terms; this function does not rebuild the MPO.
Deprecated since version 0.6.0: We switched from the three arguments \(u 0\), op0 and other_op with other_ops=[(u1, op1, dx1), (op2, u2, dx2), ...] to a single, equivalent argment ops which should now read ops=[(op0, dx0, u0), (op1, dx1, u1), (op2, dx2, u2), . ..], where \(\mathrm{dx} 0=[0] *\) self.lat.dim. Note the changed order inside the tuples!

\section*{Parameters}
- strength (scalar / array) - Prefactor of the coupling. May vary spatially, and is tiled to the required shape.
- ops (list of (opname, \(d x, u\) )) - Each tuple determines one operator of the coupling, see the description above. opname (str) is the name of the operator, \(d x\) (list of length
lat.dim) is a translation vector, and \(u\) (int) is the index of lat.unit_cell on which the operator acts. The first entry of ops corresponds to \(O P_{0}\) and acts last in the physical sense.
- op_string (str / None) - If a string is given, we use this as the name of an operator to be used inbetween the operators, excluding the sites on which any operators act. This operator should be defined on all sites in the unit cell.

If None, auto-determine whether a Jordan-Wigner string is needed (using op_needs_JW ()) for each of the segments inbetween the operators and also on the sites of the left operators.
- category (str) - Descriptive name used as key for coupling_terms. Defaults to a string of the form " \(\{0 \mathrm{p} 0\}\) _i \(\{0\) her_ops [0]\}_j \(\{0\) other_ops [1]\}_k ...".
- plus_hc (bool) - If True, the hermitian conjugate of the terms is added automatically.

\section*{Examples}

A call to add_coupling() with arguments add_coupling(strength, u1, 'A', u2, ' \(\mathrm{B}^{\prime}, \mathrm{dx}\) ) is equivalent to the following:
```

>>> dx_0 = [0] * self.lat.dim \# = [0] for a ID lattice, [0, 0] in 2D
>>> self.add_multi_coupling(strength, [('A', dx_0, u1), ('B', dx, u2)])

```

To explicitly add the hermitian conjugate (instead of simply using plus_hc = True), you need to take the complex conjugate of the strength, reverse the order of the operators and take the hermitian conjugates of the individual operator names (indicated by the hc (...), see get_hc_op_name ()):
```

>>> self.add_multi_coupling(np.conj(strength), [(hc('B'), dx, u2), (hc('A'),
\hookrightarrowdx_0, u1)])

```

\section*{See also:}
add_onsite Add terms acting on one site only.
add_coupling Add terms acting on two sites.
add_multi_coupling_term Add a single term, not summing over the possible \(\vec{x}\).
add_multi_coupling_term (strength, ijkl,ops_ijkl,op_string, category=None, plus_hc=False)
Add a general M-site coupling term on given MPS sites.
Wrapper for self.coupling_terms[category].add_multi_coupling_term(....).

\section*{Warning: This function does not handle Jordan-Wigner strings! You might want to use add_local_term() instead.}

\section*{Parameters}
- strength (float) - The strength of the coupling term.
- ijkl (list of int) - The MPS indices of the sites on which the operators acts. With \(i, j, k, \ldots=i j k l\), we require that they are ordered ascending, \(i<j<k<\ldots\) and that \(0<=i<N \_s i t e s\). Inidces \(>=\) N_sites indicate couplings between different unit cells of an infinite MPS.
- ops_ijkl(list of str) - Names of the involved operators on sites \(i, j, k, \ldots\).
- op_string (list of str) - Names of the operator to be inserted between the operators, e.g., op_string[0] is inserted between \(i\) and \(j\).
- category (str) - Descriptive name used as key for coupling_terms. Defaults to a string of the form " \(\{\mathrm{op} 0\} \_i \quad\{o p 1\} \_j \quad\{o p 2\} \_k \ldots\). . "
- plus_hc (bool) - If True, the hermitian conjugate of the term is added automatically.
add_onsite (strength, \(u\), opname, category=None, plus_hc=False)
Add onsite terms to onsite_terms.
Adds \(\sum_{\vec{x}}\) strength \([\vec{x}] * O P\) to the represented Hamiltonian, where the operator \(\mathrm{OP}=1\) at. unit_cell[u].get_op(opname) acts on the site given by a lattice index (x_0, ..., x_\{dim-1\}, u),

The necessary terms are just added to onsite_terms; doesn't rebuild the MPO.

\section*{Parameters}
- strength (scalar / array) - Prefactor of the onsite term. May vary spatially. If an array of smaller size is provided, it gets tiled to the required shape.
- u(int) - Picks a Site lat.unit_cell[u] out of the unit cell.
- opname (str) - valid operator name of an onsite operator in lat.unit_cell [u].
- category (str) - Descriptive name used as key for onsite_terms. Defaults to opname.
- plus_hc (bool) - If True, the hermitian conjugate of the terms is added automatically.

\section*{See also:}
add_coupling Add a terms acting on two sites.
add_onsite_term Add a single term without summing over vecx.
add_onsite_term (strength, \(i\), op, category=None, plus_hc=False)
Add an onsite term on a given MPS site.
Wrapper for self.onsite_terms[category].add_onsite_term(....).

\section*{Parameters}
- strength (float) - The strength of the term.
- i(int) - The MPS index of the site on which the operator acts. We require \(0<=i<\) L.
- op (str) - Name of the involved operator.
- category (str) - Descriptive name used as key for onsite_terms. Defaults to op.
- plus_hc (bool) - If True, the hermitian conjugate of the term is added automatically.
all_coupling_terms()
Sum of all coupling_terms.
all_onsite_terms()
Sum of all onsite_terms.

\section*{bond_energies ( \(p s i\) )}

Calculate bond energies <psilH_bondlpsi>.
Parameters psi (MPS) - The MPS for which the bond energies should be calculated.

Returns E_bond - List of bond energies: for finite bc, E_Bond [i] is the energy of bond i, i+1. (i.e. we omit bond 0 between sites L-1 and 0 ); for infinite bc E_bond[i] is the energy of bond \(i-1, i\).
Return type 1D ndarray
calc_H_MPO (tol_zero \(=1 e-15\) )
Calculate MPO representation of the Hamiltonian.
Uses onsite_terms and coupling_terms to build an MPOGraph (and then an MPO).
Parameters tol_zero (float) - Prefactors with abs (strength) < tol_zero are considered to be zero.

Returns H_MPO - MPO representation of the Hamiltonian.
Return type MPO
calc_H_MPO_from_bond (tol_zero=le-15)
Calculate the MPO Hamiltonian from the bond Hamiltonian.
Parameters tol_zero \((f l o a t)\) - Arrays with norm <tol_zero are considered to be zero.
Returns H_MPO - MPO representation of the Hamiltonian.
Return type MPO
calc_H_bond (tol_zero \(=1 e-15\) )
calculate \(H\) _bond from coupling_terms and onsite_terms.
Parameters tol_zero (float) - prefactors with abs(strength) < tol_zero are considered to be zero.

Returns H_bond - Bond terms as required by the constructor of NearestNeighborModel. Legs are ['p0', 'p0*', 'p1', 'p1*']
Return type list of Array
:raises ValueError : if the Hamiltonian contains longer-range terms.:
calc_H_bond_from_MPO (tol_zero=le-15)
Calculate the bond Hamiltonian from the MPO Hamiltonian.
Parameters tol_zero (float) - Arrays with norm <tol_zero are considered to be zero.
Returns H_bond - Bond terms as required by the constructor of NearestNeighborModel. Legs are ['p0', 'p0*', 'p1', 'p1*']
Return type list of Array
:raises ValueError : if the Hamiltonian contains longer-range terms.:
calc_H_onsite (tol_zero=le-15)
Calculate H_onsite from self.onsite_terms.
Deprecated since version 0.4.0: This function will be removed in 1.0.0. Replace calls to this function by self.all_onsite_terms().remove_zeros(tol_zero).to_Arrays(self. lat.mps_sites()). You might also want to take explicit_plus_hc into account.

Parameters tol_zero (float) - prefactors with abs (strength) < tol_zero are considered to be zero.

\section*{Returns}
- H_onsite (list of npc.Array)
- onsite terms of the Hamiltonian. If explicit_plus_hc is True, - Hermitian conjugates of the onsite terms will be included.
coupling_strength_add_ext_flux (strength, \(d x\), phase)
Add an external flux to the coupling strength.
When performing DMRG on a "cylinder" geometry, it might be useful to put an "external flux" through the cylinder. This means that a particle hopping around the cylinder should pick up a phase given by the external flux [[resta1998]]. This is also called "twisted boundary conditions" in literature. This function adds a complex phase to the strength array on some bonds, such that particles hopping in positive direction around the cylinder pick up \(\exp (+i\) phase \()\).

Warning: For the sign of phase it is important that you consistently use the creation operator as opl and the annihilation operator as op2 in add_coupling().

\section*{Parameters}
- strength (scalar / array) - The strength to be used in add_coupling(), when no external flux would be present.
- \(\mathbf{d} \mathbf{x}\) (iterable of int) - Translation vector (of the unit cell) between opl and op2 in add_coupling().
- phase (iterable of float) - The phase of the external flux for hopping in each direction of the lattice. E.g., if you want flux through the cylinder on which you have an infinite MPS, you should give phase \(=[0\), phi] souch that particles pick up a phase phi when hopping around the cylinder.
Returns strength - The strength array to be used as strength in add_coupling() with the given \(d x\).
Return type complex array

\section*{Examples}

Let's say you have an infinite MPS on a cylinder, and want to add nearest-neighbor hopping of fermions with the FermionSite. The cylinder axis is the \(x\)-direction of the lattice, so to put a flux through the cylinder, you want particles hopping around the cylinder to pick up a phase phi given by the external flux.
```

>>> strength = 1. \# hopping strength without external flux
>>> phi = np.pi/4 \# determines the external flux strength
>>> for u1, u2, dx in self.lat.pairs['nearest_neighbors']:
... strength_with_flux = self.coupling_strength_add_ext_flux(strength, dx,
@ [0, phi])
... self.add_coupling(strength_with_flux, u1, 'Cd', u2, 'C', dx)
... self.add_coupling(np.conj(strength_with_flux), u2, 'cd', u1, 'C', -dx)

```
enlarge_mps_unit_cell (factor=2)
Repeat the unit cell for infinite MPS boundary conditions; in place.
This has to be done after finishing initialization and can not be reverted.
Parameters factor (int) - The new number of sites in the MPS unit cell will be increased from \(N\) _sites to factor*N_sites_per_ring. Since MPS unit cells are repeated in the \(x\)-direction in our convetion, the lattice shape goes from ( \(L x, L y, \ldots, L u\) ) to (Lx*factor, Ly, ..., Lu).

\section*{classmethod from_MPOModel (mpo_model)}

Initialize a NearestNeighborModel from a model class defining an MPO.
This is especially usefull in combination with MPOModel.group_sites ().
Parameters mpo_model (MPOModel) - A model instance implementing the MPO. Does not need to be a NearestNeighborModel, but should only have nearest-neighbor couplings.

\section*{Examples}

The SpinChainNNN2 has next-nearest-neighbor couplings and thus only implements an MPO:
```

>>> from tenpy.models.spins_nnn import SpinChainNNN2
>>> nnn_chain = SpinChainNNN2({'L': 20})
>>> print(isinstance(nnn_chain, NearestNeighborModel))
False
>>> print("range before grouping:", nnn_chain.H_MPO.max_range)
range before grouping: 2

```

By grouping each two neighboring sites, we can bring it down to nearest neighbors.
```

>>> grouped_sites = nnn_chain.group_sites(2)
>>> print("range after grouping:", nnn_chain.H_MPO.max_range)
range after grouping: 1

```

Yet, TEBD will not yet work, as the model doesn't define \(H_{\_}\)bond. However, we can initialize a NearestNeighborModel from the MPO:
```

>>> nnn_chain_for_tebd = NearestNeighborModel.from_MPOModel(nnn_chain)
>>> isinstance(nnn_chain_for_tebd, NearestNeighborModel)
True

```
classmethod from_hdf5 (hdf5_loader, h5gr, subpath)
Load instance from a HDF5 file.
This method reconstructs a class instance from the data saved with save_hdf5 ().

\section*{Parameters}
- hdf5_loader (Hdf5Loader) - Instance of the loading engine.
- h5gr (Group) - HDF5 group which is represent the object to be constructed.
- subpath (str) - The name of \(h 5 g r\) with a '/' in the end.

Returns obj - Newly generated class instance containing the required data.

\section*{Return type cls}
group_sites ( \(n=2\), grouped_sites=None)
Modify self in place to group sites.
Group each \(n\) sites together using the GroupedSite. This might allow to do TEBD with a Trotter decomposition, or help the convergence of DMRG (in case of too long range interactions).
This has to be done after finishing initialization and can not be reverted.

\section*{Parameters}
- \(\mathbf{n}(i n t)\) - Number of sites to be grouped together.
- grouped_sites (None l list of GroupedSite) - The sites grouped together.

Returns grouped_sites - The sites grouped together.
Return type list of GroupedSite

\section*{init_H_from_terms()}

Initialize \(H_{-} M P O\) (and \(H_{-}\)bond) from the terms of the CouplingModel.
This function is called automatically during CouplingMPOModel.__init__
If you use one of the add_* methods of the CouplingModel after initialization, you will need to call init_H_from_terms in the end by yourself, in order to update the \(H_{-} M P O\) (and possibly \(H_{-} b o n d\) ) representations. (You should get a warning about this... The way to avoid it is to initialize all the terms in init_terms by defining your own model, as outlined in Models.
init_lattice (model_params)
Initialize a lattice for the given model parameters.
This function reads out the model parameter lattice. This can be a full Lattice instance, in which case it is just returned without further action. Alternatively, the lattice parameter can be a string giving the name of one of the predefined lattices, which then gets initialized. Depending on the dimensionality of the lattice, this requires different model parameters.

Parameters model_params (dict) - The model parameters given to \(\qquad\) init \(\qquad\) _.

Returns lat - An initialized lattice.
Return type Lattice

\section*{Options}
option CouplingMPOModel.lattice: str | Lattice
The name of a lattice pre-defined in TeNPy to be initialized. Alternatively, directly a subclass of Lattice instead of the name. Alternatively, a (possibly self-defined) Lattice instance. In the latter case, no further parameters are read out.
option CouplingMPOModel.bc_MPS: str Boundary conditions for the MPS.
option CouplingMPOModel.order: str
The order of sites within the lattice for non-trivial lattices, e.g, 'default', 'snake', see ordering (). Only used if lattice is a string.
option CouplingMPOModel.L: int
The length in x-direction; only read out for 1D lattices. For an infinite system the length of the unit cell.
option CouplingMPOModel.Lx: int
option CouplingMPOModel.Ly: int
The length in x - and y-direction; only read out for 2D lattices. For "infinite" \(b c_{-}\)MPS, the system is infinite in x-direction and \(L x\) is the number of "rings" in the infinite MPS unit cell, while \(L y\) gives the circumference around the cylinder or width of th the rung for a ladder (depending on \(b c_{-} y\) ).
option CouplingMPOModel.bc_y: str
"cylinder" | "ladder"; only read out for 2D lattices. The boundary conditions in y-direction.
option CouplingMPOModel.bc_x: str
"open" | "periodic". Can be used to force "periodic" boundaries for the lattice, i.e., for the couplings in the Hamiltonian, even if the MPS is finite. Defaults to "open" for bc_MPS="finite " and "periodic" for bc_MPS="infinite. If you are not aware of the consequences, you should probably not use "periodic" boundary conditions. (The MPS is still "open", so this will introduce long-range couplings between the first and last sites of the MPS!)
```

init_sites (model_params)

```

Define the local Hilbert space and operators; needs to be implemented in subclasses.
This function gets called by init_lattice () to get the Site for the lattice unit cell.

Note: Initializing the sites requires to define the conserved quantum numbers. All pre-defined sites accept conserve=None to disable using quantum numbers. Many models in TeNPy read out the conserve model parameter, which can be set to "best" to indicate the optimal parameters.

If you need to initialize more than one site, the function tenpy.networks.site. set_common_charges () should be helpful.

Parameters model_params (dict) - The model parameters given to \(\qquad\) init \(\qquad\)
Returns sites - The local sites of the lattice, defining the local basis states and operators.
Return type (tuple of) Site
init_terms (model_params)
Add the onsite and coupling terms to the model; subclasses should implement this.
save_hdf5 (hdf5_saver, h5gr, subpath)
Export self into a HDF5 file.
This method saves all the data it needs to reconstruct self with from_hdf5 ().
This implementation saves the content of __dict__ with save_dict_content (), storing the format under the attribute 'format'.

\section*{Parameters}
- hdf5_saver (Hdf5Saver) - Instance of the saving engine.
- h5gr (: class `Group ') - HDF5 group which is supposed to represent self.
- subpath (str) - The name of \(h 5 g r\) with a ' /' in the end.
```

test_sanity()

```

Sanity check, raises ValueErrors, if something is wrong.
trivial_like_NNModel()
Return a NearestNeighborModel with same lattice, but trivial ( \(\mathrm{H}=0\) ) bonds.

\section*{Module description}

Nearest-neighbour spin-S models.
Uniform lattice of spin-S sites, coupled by nearest-neighbour interactions.

\section*{19.6 spins_nnn}
- full name: tenpy.models.spins_nnn
- parent module: tenpy.models
- type: module

\section*{Classes}

\begin{tabular}{ll}
\hline SpinChainNNN(model_params) & \begin{tabular}{l} 
Spin-S sites coupled by (next-)nearest neighbour inter- \\
actions on a GroupedSite.
\end{tabular} \\
\hline SpinChainNNN2(model_params) & \begin{tabular}{l} 
Spin-S sites coupled by next-nearest neighbour interac- \\
tions.
\end{tabular} \\
\hline
\end{tabular}

\section*{Module description}

Next-Nearest-neighbour spin-S models.
Uniform lattice of spin-S sites, coupled by next-nearest-neighbour interactions. We have two variants implementing the same hamiltonian. The SpinChainNNN uses the GroupedSite to keep it a NearestNeighborModel suitable for TEBD, while the SpinChainNNN2 just involves longer-range couplings in the MPO. The latter is preferable for pure DMRG calculations and avoids having to add each of the short range couplings twice for the grouped sites.

Note that you can also get a NearestNeighborModel for TEBD from the latter by using group_sites () and from_MPOModel(). An example for such a case is given in the file examples/c_tebd.py.

\section*{19.7 fermions_spinless}
- full name: tenpy.models.fermions_spinless
- parent module: tenpy.models
- type: module

\section*{Classes}

\begin{tabular}{ll}
\hline FermionChain(model_params) & The FermionModel on a Chain, suitable for TEBD. \\
\hline FermionModel(model_params) & Spinless fermions with particle number conservation. \\
\hline
\end{tabular}

\subsection*{19.7.1 FermionChain}
- full name: tenpy.models.fermions_spinless.FermionChain
- parent module: tenpy.models.fermions_spinless
- type: class

\section*{Inheritance Diagram}


\section*{Methods}
\begin{tabular}{ll}
\hline FermionChain.__init__(model_params) & Initialize self. \\
\hline \begin{tabular}{l} 
FermionChain.add_coupling(strength, \\
op1,...)
\end{tabular} & u1, \\
\hline FermionChain.add_coupling_term(strength, & Add a two-site coupling terms to the Hamiltonian, sum- \\
ming over lattice sites term on given MPS sites. \\
i,...) & \\
\hline FermionChain.add_exponentially_decayingAddap exponentially decaying long-range coupling. \\
\hline
\end{tabular}
continues on next page

Table 49 - continued from previous page
\begin{tabular}{ll}
\hline \begin{tabular}{l} 
FermionChain.add_local_term(strength, \\
term)
\end{tabular} & Add a single term to self. \\
\hline \begin{tabular}{l} 
FermionChain.add_multi_coupling(strength, \\
ops)
\end{tabular} & \begin{tabular}{l} 
Add multi-site coupling terms to the Hamiltonian, sum- \\
ming over lattice sites.
\end{tabular} \\
\hline FermionChain.add_multi_coupling_term(... [Add a general M-site coupling term on given MPS sites. \\
\(\ldots\).\(] )\) & \\
\hline \begin{tabular}{ll} 
FermionChain.add_onsite(strength, \\
name)
\end{tabular} & op-
\end{tabular}

\section*{Class Attributes and Properties}
\begin{tabular}{ll}
\hline FermionChain.force_default_lattice & \begin{tabular}{l} 
If True, init_lattice() asserts that the initialized \\
lattice is (a subclass of) default_lattice
\end{tabular} \\
\hline FermionChain.logger & class attribute. \\
\hline FermionChain.verbose \\
\hline
\end{tabular}

The FermionModel on a Chain, suitable for TEBD.
See the FermionModel for the documentation of parameters.

\section*{default_lattice}
alias of tenpy.models.lattice.Chain
add_coupling (strength, u1, op1, u2, op2, dx, op_string=None, str_on_first=True, raise_op2_left=False, category=None, plus_hc=False)
Add twosite coupling terms to the Hamiltonian, summing over lattice sites.
Represents couplings of the form \(\sum_{x_{0}, \ldots, x_{d i m-1}} \operatorname{strength}[\operatorname{shift}(\vec{x})] * O P 0 * O P 1\), where OP 0 : = lat.unit_cell[u0].get_op(op0) acts on the site (x_0, ..., x_\{dim-1\}, u1), and OP1 := lat.unit_cell[ul].get_op(op1) acts on the site (x_0+dx[0], ..., \(\left.x_{\_}\{\operatorname{dim}-1\}+d x[\operatorname{dim}-1], u 1\right)\). Possible combinations \(x \_0, \ldots, x_{-}\{d i m-1\}\) are determined from the boundary conditions in possible_couplings ().

The coupling strength may vary spatially if the given strength is a numpy array. The correct shape of this array is the coupling_shape returned by tenpy.models.lattice.coupling_shape () and depends on the boundary conditions. The shift (. . .) depends on \(d x\), and is chosen such that the first entry strength \([0,0, \ldots]\) of strength is the prefactor for the first possible coupling fitting into the lattice if you imagine open boundary conditions.

The necessary terms are just added to coupling_terms; this function does not rebuild the MPO.
Deprecated since version 0.4.0: The arguments str_on_first and raise_op2_left will be removed in version 1.0.0.

\section*{Parameters}
- strength (scalar / array) - Prefactor of the coupling. May vary spatially (see above). If an array of smaller size is provided, it gets tiled to the required shape.
- u1 (int) - Picks the site lat. unit_cell[u1] for OP1.
- op1 (str) - Valid operator name of an onsite operator in lat.unit_cell[u1] for OP1.
- u2 (int) - Picks the site lat.unit_cell[u2] for OP2.
- op2 (str) - Valid operator name of an onsite operator in lat.unit_cell[u2] for OP2.
- \(\mathbf{d x}\) (iterable of int) - Translation vector (of the unit cell) between OP1 and OP2. For a 1D lattice, a single int is also fine.
- op_string (str / None) - Name of an operator to be used between the OP1 and OP2 sites. Typical use case is the phase for a Jordan-Wigner transformation. The operator
should be defined on all sites in the unit cell. If None, auto-determine whether a JordanWigner string is needed, using op_needs_JW().
- str_on_first (bool) - Whether the provided op_string should also act on the first site. This option should be chosen as True for Jordan-Wigner strings. When handling Jordan-Wigner strings we need to extend the op_string to also act on the 'left', first site (in the sense of the MPS ordering of the sites given by the lattice). In this case, there is a well-defined ordering of the operators in the physical sense (i.e. which of opl or op2 acts first on a given state). We follow the convention that op2 acts first (in the physical sense), independent of the MPS ordering. Deprecated.
- raise_op2_left (bool) - Raise an error when op2 appears left of opl (in the sense of the MPS ordering given by the lattice). Deprecated.
- category (str) - Descriptive name used as key for coupling_terms. Defaults to a string of the form " \(\{o p 1\} \_i \quad\{o p 2\} \_j "\).
- plus_hc (bool) - If True, the hermitian conjugate of the terms is added automatically.

\section*{Examples}

When initializing a model, you can add a term \(J \sum_{<i, j>} S_{i}^{z} S_{j}^{z}\) on all nearest-neighbor bonds of the lattice like this:
```

>>> J = 1. \# the strength
>>> for u1, u2, dx in self.lat.pairs['nearest_neighbors']:
... self.add_coupling(J, u1, 'Sz', u2, 'Sz', dx)

```

The strength can be an array, which gets tiled to the correct shape. For example, in a 1D Chain with an even number of sites and periodic (or infinite) boundary conditions, you can add alternating strong and weak couplings with a line like:
```

>>> self.add_coupling([1.5, 1.], u1, 'Sz', u2, 'Sz', dx)

```

Make sure to use the plus_hc argument if necessary, e.g. for hoppings:
```

>>> t = 1. \# hopping strength
>>> for ul, u2, dx in self.lat.pairs['nearest_neighbors']:
... self.add_coupling(t, u1, 'Cd', u2, 'C', dx, plus_hc=True)

```

Alternatively, you can add the hermitian conjugate terms explictly. The correct way is to complex conjugate the strength, take the hermitian conjugate of the operators and swap the order (including a swap \(u l<->u 2\) ), and use the opposite direction \(-d x\), i.e. the h.c. of add_coupling ( \(t, \mathrm{u} 1\), 'A', u2, 'B', dx) is add_coupling(np.conj(t), u2, hc('B'), u1, hc('A'), -dx), where \(h c\) takes the hermitian conjugate of the operator names, see get_hc_op_name (). For spin-less fermions (FermionSite), this would be
```

>>> for u1, u2, dx in self.lat.pairs['nearest_neighbors']:
... self.add_coupling(t, u1, 'Cd', u2, 'C', dx)
... self.add_coupling(np.conj(t), u2, 'Cd', ul, 'C', -dx) \# h.c.

```

With spin-full fermions (SpinHalfFermions), it could be:
```

>>> for u1, u2, dx in self.lat.pairs['nearest_neighbors']:
self.add_coupling(t, u1, 'Cdu', u2, 'Cd', dx) \# Cdagger_up C_down
self.add_coupling(np.conj(t), u2, 'Cdd', u1, 'Cu', -dx) \# h.c.v
->Cdagger_down C_up

```

Note that the Jordan-Wigner strings for the fermions are added automatically!

\section*{See also:}
add_onsite Add terms acting on one site only.
add_multi_coupling_term for terms on more than two sites.
add_coupling_term Add a single term without summing over \(\vec{x}\).
add_coupling_term (strength, \(i, j, o p_{-} i, o p \_j, o p_{-}\)string \(=' I d^{\prime}\), category \(=\) None, plus_hc=False)
Add a two-site coupling term on given MPS sites.
Wrapper for self.coupling_terms[category].add_coupling_term(...).

Warning: This function does not handle Jordan-Wigner strings! You might want to use add_local_term() instead.

\section*{Parameters}
- strength (float) - The strength of the coupling term.
- \(\mathbf{i}\) (int) - The MPS indices of the two sites on which the operator acts. We require 0 \(<=i<N \_s i t e s\) and \(i<j\), i.e., \(o p_{-} i\) acts "left" of \(o p_{-} j\). If \(\mathrm{j}>=\mathrm{N}_{-}\)sites, it indicates couplings between unit cells of an infinite MPS.
- j (int) - The MPS indices of the two sites on which the operator acts. We require 0 \(<=i<N \_s i t e s\) and \(i<j\), i.e., op_i acts "left" of \(o p_{-} j\). If \(\mathrm{j}>=\mathrm{N}_{-}\)sites, it indicates couplings between unit cells of an infinite MPS.
- op1 (str) - Names of the involved operators.
- op2 (str) - Names of the involved operators.
- op_string (str) - The operator to be inserted between \(i\) and \(j\).
- category (str) - Descriptive name used as key for coupling_terms. Defaults to a string of the form " \(\{0 \mathrm{p} 1\} \_i \quad\{o p 2\} \_j\) ".
- plus_hc (bool) - If True, the hermitian conjugate of the term is added automatically.
add_exponentially_decaying_coupling (strength, lambda_, op_i,op_j, subsites=None, \(o p_{-}\)string \(=\)None, \(p l u s_{-} h c=\) False \()\)
Add an exponentially decaying long-range coupling.
\[
\text { strength } \sum_{i<j} \lambda^{|i-j|} A_{\text {subsites }[i]} B_{\text {subsites }[j]}
\]

Where the operator \(A\) is given by \(o p_{-} i\), and \(B\) is given by \(o p_{-} j\). Note that the sum over \(i, j\) is long-range, for infinite systems going beyond the MPS unit cell. Moreover, note that the distance in the exponent is the distance within subsites.

\section*{Parameters}
- strength (float) - Overall prefactor.
- lambda (float) - Decay-rate
- op_i (string) - Names for the operators.
- op_j (string) - Names for the operators.
- subsites (None / 1D array) - Selects a subset of sites within the MPS unit cell on which the operators act. Needs to be sorted. None selects all sites.
- op_string (None / str) - The operator to be inserted between \(A\) and \(B\); If None, this function checks whether a fermionic "JW" string is needed for the given operators; in this case the right \(o p_{-} j\) acts first.
- plus_hc (bool) - If True, the hermitian conjugate of the term is added automatically.

\section*{Examples}

At least for simple enough 1D chains (or ladders), you can use fit_with_sum_of_exp () to approximate a long-range function with a few sum of exponentials and then add them with this function.
```

>>> def decay(x):
... return np.exp(-0.1*x) / x**2
>>> from tenpy.tools.fit import fit_with_sum_of_exp, sum_of_exp
>>> n_exp = 5
>>> fit_range = 50
>>> lam, pref = fit_with_sum_of_exp(decay, n_exp, fit_range)
>>> x = np.arange(1, fit_range + 1)
>>> print('error in fit: {0:.3e}'.format(np.sum(np.abs(decay(x) - sum_of__
@exp(lam, pref, x)))))
error in fit: 1.073e-04
>>> for pr, la in zip(pref, lam):
... self.add_exponentially_decaying_coupling(pr, la, 'N', 'N')

```
add_local_term (strength, term, category=None, plus_hc=False)
Add a single term to self.
The repesented term is strength times the product of the operators given in terms. Each operator is specified by the name and the site it acts on; the latter given by a lattice index, see Lattice.

Depending on the length of term, it can add an onsite term or a coupling term to onsite_terms or coupling_terms, respectively.

\section*{Parameters}
- strength (float/complex) - The prefactor of the term.
- term (list of (str, array_like))- List of tuples (opname, lat_idx) where opname is a string describing the operator acting on the site given by the lattice index lat_idx. Here, lat_idx is for example \([x, y, u]\) for a 2D lattice, with \(u\) being the index within the unit cell.
- category - Descriptive name used as key for onsite_terms or coupling_terms.
- plus_hc (bool) - If True, the hermitian conjugate of the terms is added automatically.
add_multi_coupling (strength, ops, _deprecate_l='DEPRECATED', _deprecate_2 = 'DEPRECATED', op_string=None, category=None, plus_hc=False)
Add multi-site coupling terms to the Hamiltonian, summing over lattice sites.
Represents couplings of the form \(\operatorname{sum}_{\vec{x}} \operatorname{strength}[\operatorname{shift}(\vec{x})] * O P_{0} * O P_{1} * \ldots * O P_{M-1}\), involving \(M\) operators. Here, \(O P_{m}\) stands for the operator defined by the \(m\)-th tuple (opname, \(\mathrm{dx}, \mathrm{u}\) ) given in the argument ops, which determines the position \(\vec{x}+\overrightarrow{d x}\) and unit-cell index \(u\) of the site it acts on; the actual operator is given by self.lat.unit_cell[u].get_op(opname).

The coupling strength may vary spatially if the given strength is a numpy array. The correct shape of this array is the coupling_shape returned by tenpy.models.lattice. possible_multi_couplings() and depends on the boundary conditions. The shift(...) depends on the \(d x\) entries of ops and is chosen such that the first entry strength [0, 0, ...] of strength is the prefactor for the first possible coupling fitting into the lattice if you imagine open boundary conditions.

The necessary terms are just added to coupling_terms; this function does not rebuild the MPO.
Deprecated since version 0.6.0: We switched from the three arguments \(u 0\), op0 and other_op with other_ops=[(u1, op1, dx1), (op2, u2, dx2), ...] to a single, equivalent argment ops which should now read ops \(=[(o p 0, d x 0, u 0),(o p 1, d x 1, u 1),(o p 2, d x 2, u 2)\), ..], where \(\mathrm{dx} 0=[0] *\) self.lat.dim. Note the changed order inside the tuples!

\section*{Parameters}
- strength (scalar / array) - Prefactor of the coupling. May vary spatially, and is tiled to the required shape.
- ops (list of (opname, \(d x, u\) ) ) - Each tuple determines one operator of the coupling, see the description above. opname (str) is the name of the operator, \(d x\) (list of length lat.dim) is a translation vector, and \(u\) (int) is the index of lat.unit_cell on which the operator acts. The first entry of ops corresponds to \(O P_{0}\) and acts last in the physical sense.
- op_string (str / None) - If a string is given, we use this as the name of an operator to be used inbetween the operators, excluding the sites on which any operators act. This operator should be defined on all sites in the unit cell.

If None, auto-determine whether a Jordan-Wigner string is needed (using op_needs_JW ()) for each of the segments inbetween the operators and also on the sites of the left operators.
- category (str) - Descriptive name used as key for coupling_terms. Defaults to a

- plus_hc (bool) - If True, the hermitian conjugate of the terms is added automatically.

\section*{Examples}

A call to add_coupling() with arguments add_coupling(strength, u1, 'A', u2, ' B ', dx ) is equivalent to the following:
```

>>> dx_0 = [0] * self.lat.dim \# = [0] for a 1D lattice, [0, 0] in 2D
>>> self.add_multi_coupling(strength, [('A', dx_0, u1), ('B', dx, u2)])

```

To explicitly add the hermitian conjugate (instead of simply using plus_hc = True), you need to take the complex conjugate of the strength, reverse the order of the operators and take the hermitian conjugates of the individual operator names (indicated by the hc (. . .) , see get_hc_op_name ()):
```

>>> self.add_multi_coupling(np.conj(strength), [(hc('B'), dx, u2), (hc('A'), -
\hookrightarrowdx_0, ul)])

```

\section*{See also:}
add_onsite Add terms acting on one site only.
add_coupling Add terms acting on two sites.
add_multi_coupling_term Add a single term, not summing over the possible \(\vec{x}\).
add_multi_coupling_term (strength, ijkl,ops_ijkl,op_string, category=None, plus_hc=False)
Add a general M-site coupling term on given MPS sites.
Wrapper for self.coupling_terms[category].add_multi_coupling_term(...).

Warning: This function does not handle Jordan-Wigner strings! You might want to use add_local_term() instead.

\section*{Parameters}
- strength (float) - The strength of the coupling term.
- ijkl (list of int) - The MPS indices of the sites on which the operators acts. With \(i, j, k, \ldots=i j k l\), we require that they are ordered ascending, \(i<j<k<\ldots\) and that \(0<=i<N \_\)sites. Inidces \(>=\)N_sites indicate couplings between different unit cells of an infinite MPS.
- ops_ijkl (list of str) - Names of the involved operators on sites \(i, j, k, \ldots\).
- op_string (list of str) - Names of the operator to be inserted between the operators, e.g., op_string[0] is inserted between \(i\) and \(j\).
- category (str) - Descriptive name used as key for coupling_terms. Defaults to a string of the form " \(\{\mathrm{op} 0\} \_i \quad\{o p 1\} \_j \quad\{o p 2\} \_k \ldots\). . ".
- plus_hc (bool) - If True, the hermitian conjugate of the term is added automatically.
add_onsite (strength, \(u\), opname, category=None, plus_hc=False)
Add onsite terms to onsite_terms.
Adds \(\sum_{\vec{x}} \operatorname{strength}[\vec{x}] * O P\) to the represented Hamiltonian, where the operator \(\mathrm{OP}=1\) at. unit_cell[u].get_op (opname) acts on the site given by a lattice index (x_0, ..., x_\{dim-1\}, u),

The necessary terms are just added to onsite_terms; doesn't rebuild the MPO.

\section*{Parameters}
- strength (scalar / array) - Prefactor of the onsite term. May vary spatially. If an array of smaller size is provided, it gets tiled to the required shape.
- u (int) - Picks a Site lat.unit_cell[u] out of the unit cell.
- opname (str) - valid operator name of an onsite operator in lat.unit_cell [u].
- category (str) - Descriptive name used as key for onsite_terms. Defaults to opname.
- plus_hc (bool) - If True, the hermitian conjugate of the terms is added automatically.

\section*{See also:}
add_coupling Add a terms acting on two sites.
add_onsite_term Add a single term without summing over vecx.
add_onsite_term (strength, \(i, o p\), category=None, plus_hc=False)
Add an onsite term on a given MPS site.
Wrapper for self.onsite_terms[category].add_onsite_term(....).

\section*{Parameters}
- strength (float) - The strength of the term.
- \(\mathbf{i}(i n t)\) - The MPS index of the site on which the operator acts. We require \(0<=i<\) L.
- op (str) - Name of the involved operator.
- category (str) - Descriptive name used as key for onsite_terms. Defaults to op.
- plus_hc (bool) - If True, the hermitian conjugate of the term is added automatically.
all_coupling_terms()
Sum of all coupling_terms.
all_onsite_terms()
Sum of all onsite_terms.
bond_energies ( \(p s i\) )
Calculate bond energies <psilH_bondlpsi>.
Parameters psi (MPS) - The MPS for which the bond energies should be calculated.
Returns E_bond - List of bond energies: for finite bc, E_Bond [i] is the energy of bond i, i+1. (i.e. we omit bond 0 between sites L-1 and 0); for infinite bc E_bond[i] is the energy of bond \(i-1\), \(i\).

Return type 1D ndarray
calc_H_MPO (tol_zero \(=1 e-15\) )
Calculate MPO representation of the Hamiltonian.
Uses onsite_terms and coupling_terms to build an MPOGraph (and then an MPO).
Parameters tol_zero (float) - Prefactors with abs (strength) < tol_zero are considered to be zero.

Returns H_MPO - MPO representation of the Hamiltonian.
Return type MPO
calc_H_MPO_from_bond (tol_zero=1e-15)
Calculate the MPO Hamiltonian from the bond Hamiltonian.
Parameters tol_zero (float) - Arrays with norm < tol_zero are considered to be zero.
Returns H_MPO - MPO representation of the Hamiltonian.
Return type MPO
calc_H_bond (tol_zero \(=1 e-15\) )
calculate \(H\) _bond from coupling_terms and onsite_terms.
Parameters tol_zero (float) - prefactors with abs (strength) < tol_zero are considered to be zero.

Returns H_bond - Bond terms as required by the constructor of NearestNeighborModel. Legs are ['p0', 'p0*', 'p1', 'p1*']

Return type list of Array
:raises ValueError : if the Hamiltonian contains longer-range terms.:
calc_H_bond_from_MPO (tol_zero=le-15)
Calculate the bond Hamiltonian from the MPO Hamiltonian.
Parameters tol_zero \((f l o a t)\) - Arrays with norm <tol_zero are considered to be zero.

Returns H_bond - Bond terms as required by the constructor of NearestNeighborModel.
Legs are ['p0', 'p0*', 'p1', 'p1*']
Return type list of Array
:raises ValueError : if the Hamiltonian contains longer-range terms.:
calc_H_onsite (tol_zero=le-15)
Calculate H_onsite from self.onsite_terms.
Deprecated since version 0.4.0: This function will be removed in 1.0.0. Replace calls to this function by self.all_onsite_terms().remove_zeros(tol_zero).to_Arrays (self. lat.mps_sites()). You might also want to take explicit_plus_hc into account.

Parameters tol_zero (float) - prefactors with abs(strength) < tol_zero are considered to be zero.

\section*{Returns}
- H_onsite (list of npc.Array)
- onsite terms of the Hamiltonian. If explicit_plus_hc is True, - Hermitian conjugates of the onsite terms will be included.
coupling_strength_add_ext_flux (strength, \(d x\), phase)
Add an external flux to the coupling strength.
When performing DMRG on a "cylinder" geometry, it might be useful to put an "external flux" through the cylinder. This means that a particle hopping around the cylinder should pick up a phase given by the external flux [[resta1998]]. This is also called "twisted boundary conditions" in literature. This function adds a complex phase to the strength array on some bonds, such that particles hopping in positive direction around the cylinder pick up \(\exp (+i\) phase \()\).

Warning: For the sign of phase it is important that you consistently use the creation operator as opl and the annihilation operator as op2 in add_coupling().

\section*{Parameters}
- strength (scalar / array) - The strength to be used in add_coupling(), when no external flux would be present.
- dx (iterable of int) - Translation vector (of the unit cell) between opl and op2 in add_coupling().
- phase (iterable of float) - The phase of the external flux for hopping in each direction of the lattice. E.g., if you want flux through the cylinder on which you have an infinite MPS, you should give phase \(=[0\), phi] souch that particles pick up a phase phi when hopping around the cylinder.

Returns strength - The strength array to be used as strength in add_coupling() with the given \(d x\).

Return type complex array

\section*{Examples}

Let's say you have an infinite MPS on a cylinder, and want to add nearest-neighbor hopping of fermions with the Fermionsite. The cylinder axis is the \(x\)-direction of the lattice, so to put a flux through the cylinder, you want particles hopping around the cylinder to pick up a phase phi given by the external flux.
```

>>> strength = 1. \# hopping strength without external flux
>>> phi = np.pi/4 \# determines the external flux strength
>>> for u1, u2, dx in self.lat.pairs['nearest_neighbors']:
... strength_with_flux = self.coupling_strength_add_ext_flux(strength, dx,
\hookrightarrow [0, phi])
... self.add_coupling(strength_with_flux, u1, 'Cd', u2, 'C', dx)
... self.add_coupling(np.conj(strength_with_flux), u2, 'cd', u1, 'C', -dx)

```
enlarge_mps_unit_cell(factor=2)
Repeat the unit cell for infinite MPS boundary conditions; in place.
This has to be done after finishing initialization and can not be reverted.
Parameters factor (int) - The new number of sites in the MPS unit cell will be increased from \(N_{\text {_s }}\) sites to factor*N_sites_per_ring. Since MPS unit cells are repeated in the \(x\)-direction in our convetion, the lattice shape goes from ( \(\mathrm{Lx}, \mathrm{Ly}, \ldots, \mathrm{Lu}\) ) to (Lx*factor, Ly, ..., Lu).
classmethod from_MPOModel (mpo_model)
Initialize a NearestNeighborModel from a model class defining an MPO.
This is especially usefull in combination with MPOModel.group_sites().
Parameters mpo_model (MPOModel) - A model instance implementing the MPO. Does not need to be a NearestNeighborModel, but should only have nearest-neighbor couplings.

\section*{Examples}

The SpinChainNNN2 has next-nearest-neighbor couplings and thus only implements an MPO:
```

>>> from tenpy.models.spins_nnn import SpinChainNNN2
>>> nnn_chain = SpinChainNNN2({'L': 20})
>>> print(isinstance(nnn_chain, NearestNeighborModel))
False
>>> print("range before grouping:", nnn_chain.H_MPO.max_range)
range before grouping: 2

```

By grouping each two neighboring sites, we can bring it down to nearest neighbors.
```

>>> grouped_sites = nnn_chain.group_sites(2)
>>> print("range after grouping:", nnn_chain.H_MPO.max_range)
range after grouping: 1

```

Yet, TEBD will not yet work, as the model doesn't define \(H \_b o n d\). However, we can initialize a NearestNeighborModel from the MPO:
```

>>> nnn_chain_for_tebd = NearestNeighborModel.from_MPOModel(nnn_chain)
>>> isinstance(nnn_chain_for_tebd, NearestNeighborModel)
True

```
classmethod from_hdf5 (hdf5_loader, h5gr, subpath)
Load instance from a HDF5 file.
This method reconstructs a class instance from the data saved with save_hdf5 ().

\section*{Parameters}
- hdf5_loader (Hdf5Loader) - Instance of the loading engine.
- h5gr (Group) - HDF5 group which is represent the object to be constructed.
- subpath (str) - The name of \(h 5 g r\) with a ' /' in the end.

Returns obj - Newly generated class instance containing the required data.

\section*{Return type cls}
group_sites ( \(n=2\), grouped_sites=None)
Modify self in place to group sites.
Group each \(n\) sites together using the GroupedSite. This might allow to do TEBD with a Trotter decomposition, or help the convergence of DMRG (in case of too long range interactions).

This has to be done after finishing initialization and can not be reverted.

\section*{Parameters}
- \(\mathbf{n}(i n t)\) - Number of sites to be grouped together.
- grouped_sites (None l list of GroupedSite) - The sites grouped together.

Returns grouped_sites - The sites grouped together.
Return type list of GroupedSite
init_H_from_terms()
Initialize \(H_{-} M P O\) (and \(H \_\)bond) from the terms of the CouplingModel.
This function is called automatically during CouplingMPOModel.__init__.
If you use one of the add_* methods of the CouplingModel after initialization, you will need to call init_H_from_terms in the end by yourself, in order to update the \(H_{-} M P O\) (and possibly \(H_{-} b o n d\) ) representations. (You should get a warning about this... The way to avoid it is to initialize all the terms in init_terms by defining your own model, as outlined in Models.

\section*{init_lattice (model_params)}

Initialize a lattice for the given model parameters.
This function reads out the model parameter lattice. This can be a full Lattice instance, in which case it is just returned without further action. Alternatively, the lattice parameter can be a string giving the name of one of the predefined lattices, which then gets initialized. Depending on the dimensionality of the lattice, this requires different model parameters.

Parameters model_params (dict) - The model parameters given to \(\qquad\) init \(\qquad\)
Returns lat - An initialized lattice.
Return type Lattice

\section*{Options}
option CouplingMPOModel.lattice: str | Lattice The name of a lattice pre-defined in TeNPy to be initialized. Alternatively, directly a subclass of Lattice instead of the name. Alternatively, a (possibly self-defined) Lattice instance. In the latter case, no further parameters are read out.
option CouplingMPOModel.bc_MPS: str Boundary conditions for the MPS.
option CouplingMPOModel.order: str
The order of sites within the lattice for non-trivial lattices, e.g, 'default', 'snake', see ordering (). Only used if lattice is a string.
option CouplingMPOModel.L: int
The length in x-direction; only read out for 1D lattices. For an infinite system the length of the unit cell.
option CouplingMPOModel.Lx: int
option CouplingMPOModel.Ly: int
The length in x - and y-direction; only read out for 2D lattices. For "infinite" \(b c_{-} M P S\), the system is infinite in x-direction and \(L x\) is the number of "rings" in the infinite MPS unit cell, while \(L y\) gives the circumference around the cylinder or width of th the rung for a ladder (depending on \(b c_{-} y\) ).
option CouplingMPOModel.bc_y: str
"cylinder" | "ladder"; only read out for 2D lattices. The boundary conditions in y-direction.
option CouplingMPOModel.bc_x: str
"open" | "periodic". Can be used to force "periodic" boundaries for the lattice, i.e., for the couplings in the Hamiltonian, even if the MPS is finite. Defaults to "open" for bc_MPS="finite " and "periodic" for bc_MPS="infinite. If you are not aware of the consequences, you should probably not use "periodic" boundary conditions. (The MPS is still "open", so this will introduce long-range couplings between the first and last sites of the MPS!)

\section*{init_sites (model_params)}

Define the local Hilbert space and operators; needs to be implemented in subclasses.
This function gets called by init_lattice () to get the Site for the lattice unit cell.

Note: Initializing the sites requires to define the conserved quantum numbers. All pre-defined sites accept conserve=None to disable using quantum numbers. Many models in TeNPy read out the conserve model parameter, which can be set to "best" to indicate the optimal parameters.

If you need to initialize more than one site, the function tenpy.networks.site. set_common_charges () should be helpful.

Parameters model_params (dict) - The model parameters given to \(\qquad\)
\(\qquad\)
Returns sites - The local sites of the lattice, defining the local basis states and operators.
Return type (tuple of) Site
init_terms (model_params)
Add the onsite and coupling terms to the model; subclasses should implement this.
save_hdf5 (hdf5_saver, h5gr, subpath)
Export self into a HDF5 file.
This method saves all the data it needs to reconstruct self with from_hdf5 ().
This implementation saves the content of ___dict__ with save_dict_content (), storing the format under the attribute 'format'.

\section*{Parameters}
- hdf5_saver (Hdf5Saver) - Instance of the saving engine.
- h5gr (: class `Group`) - HDF5 group which is supposed to represent self.
- subpath (str) - The name of \(h 5 g r\) with a '/' in the end.

\section*{test_sanity()}

Sanity check, raises ValueErrors, if something is wrong.

\section*{trivial_like_NNModel()}

Return a NearestNeighborModel with same lattice, but trivial ( \(\mathrm{H}=0\) ) bonds.

\section*{Module description}

Spinless fermions with hopping and interaction.

Todo: add further terms (e.g. \(\mathrm{c}^{\wedge}\) dagger \(\mathrm{c}^{\wedge}\) dagger + h.c. \()\) to the Hamiltonian.

\section*{19.8 hubbard}
- full name: tenpy.models.hubbard
- parent module: tenpy.models
- type: module

\section*{Classes}

\begin{tabular}{ll}
\hline BoseHubbardChain(model_params) & \begin{tabular}{l} 
The BoseHubbardModel on a Chain, suitable for \\
TEBD.
\end{tabular} \\
\hline BoseHubbardModel(model_params) & Spinless Bose-Hubbard model. \\
\hline FermiHubbardChain(model_params) & The FermiHubbardModel on a Chain, suitable for \\
& TEBD. \\
\hline FermiHubbardModel(model_params) & Spin-1/2 Fermi-Hubbard model. \\
\hline
\end{tabular}

\subsection*{19.8.1 BoseHubbardChain}
- full name: tenpy.models.hubbard.BoseHubbardChain
- parent module: tenpy.models.hubbard
- type: class

\section*{Inheritance Diagram}


\section*{Methods}
\begin{tabular}{|c|c|}
\hline BoseHubbardChain.__init__(model_params) & Initialize self. \\
\hline BoseHubbardChain.add_coupling(strength, u1,...) & Add twosite coupling terms to the Hamiltonian, summing over lattice sites. \\
\hline \begin{tabular}{l}
BoseHubbardChain. \\
add_coupling_term(strength,...)
\end{tabular} & Add a two-site coupling term on given MPS sites. \\
\hline \begin{tabular}{l}
BoseHubbardChain. \\
add_exponentially_decaying_coupling(...)
\end{tabular} & Add an exponentially decaying long-range coupling. \\
\hline BoseHubbardChain. add_local_term(strength, term) & Add a single term to self. \\
\hline \[
\begin{aligned}
& \text { BoseHubbardChain. } \\
& \text { add_multi_coupling(...[,...]) }
\end{aligned}
\] & Add multi-site coupling terms to the Hamiltonian, summing over lattice sites. \\
\hline BoseHubbardChain. add_multi_coupling_term(...) & Add a general M-site coupling term on given MPS sites. \\
\hline BoseHubbardChain.add_onsite(strength, u, opname) & Add onsite terms to onsite_terms. \\
\hline BoseHubbardChain. add_onsite_term(strength, i, op) & Add an onsite term on a given MPS site. \\
\hline BoseHubbardChain.all_coupling_terms() & Sum of all coupling_terms. \\
\hline BoseHubbardChain.all_onsite_terms() & Sum of all onsite_terms. \\
\hline BoseHubbardChain.bond_energies(psi) & Calculate bond energies <psilH_bondlpsi>. \\
\hline BoseHubbardChain.calc_H_MPO([tol_zero]) & Calculate MPO representation of the Hamiltonian. \\
\hline \[
\begin{aligned}
& \text { BoseHubbardChain. } \\
& \text { calc_H_MPO_from_bond([tol_zero]) }
\end{aligned}
\] & Calculate the MPO Hamiltonian from the bond Hamiltonian. \\
\hline BoseHubbardChain.calc_H_bond([tol_zero]) & calculate H_bond from coupling_terms and onsite_terms. \\
\hline \[
\begin{aligned}
& \text { BoseHubbardChain. } \\
& \text { calc_H_bond_from_MPO([tol_zero]) }
\end{aligned}
\] & Calculate the bond Hamiltonian from the MPO Hamiltonian. \\
\hline BoseHubbardChain. calc_H_onsite([tol_zero]) & Calculate H_onsite from self.onsite_terms. \\
\hline BoseHubbardChain. coupling_strength_add_ext_flux(...) & Add an external flux to the coupling strength. \\
\hline \begin{tabular}{l}
BoseHubbardChain. \\
enlarge_mps_unit_cell([factor])
\end{tabular} & Repeat the unit cell for infinite MPS boundary conditions; in place. \\
\hline \begin{tabular}{l}
BoseHubbardChain. \\
from_MPOModel(mpo_model)
\end{tabular} & Initialize a NearestNeighborModel from a model class defining an MPO. \\
\hline BoseHubbardChain.from_hdf5(hdf5_loader, ...) & Load instance from a HDF5 file. \\
\hline BoseHubbardChain.group_sites([n, grouped_sites]) & Modify self in place to group sites. \\
\hline BoseHubbardChain.init_H_from_terms() & Initialize \(H_{-} M P O\) (and \(H \_b o n d\) ) from the terms of the CouplingModel. \\
\hline \begin{tabular}{l}
BoseHubbardChain. \\
init_lattice(model_params)
\end{tabular} & Initialize a lattice for the given model parameters. \\
\hline BoseHubbardChain. init_sites(model_params) & Define the local Hilbert space and operators; needs to be implemented in subclasses. \\
\hline \[
\begin{aligned}
& \text { BoseHubbardChain. } \\
& \text { init_terms(model_params) }
\end{aligned}
\] & Add the onsite and coupling terms to the model; subclasses should implement this. \\
\hline BoseHubbardChain.save_hdf5(hdf5_saver, h5gr, ...) & Export self into a HDF5 file. \\
\hline
\end{tabular}

Table 52 - continued from previous page
\begin{tabular}{ll}
\hline BoseHubbardChain.test_sanity() & Sanity check, raises ValueErrors, if something is wrong. \\
\hline BoseHubbardChain. & Return a NearestNeighborModel with same lattice, but \\
trivial_like_NNModel) & trivial \((\mathrm{H}=0)\) bonds. \\
\hline
\end{tabular}

\section*{Class Attributes and Properties}
\begin{tabular}{ll}
\hline BoseHubbardChain.default_lattice & \begin{tabular}{l} 
The default lattice class or class name to be used in \\
init_lattice().
\end{tabular} \\
\hline BoseHubbardChain. & If True, init_lattice() asserts that the initialized \\
force_default_lattice & lattice is (a subclass of)default_lattice
\end{tabular}
class tenpy.models.hubbard.BoseHubbardChain (model_params)
Bases: tenpy.models.hubbard.BoseHubbardModel, tenpy.models.model. NearestNeighborModel

The BoseHubbardModel on a Chain, suitable for TEBD.
See the BoseHubbardModel for the documentation of parameters.
add_coupling (strength, u1, op1, u2, op2, dx, op_string=None, str_on_first=True, raise_op2_left=False, category=None, plus_hc=False)
Add twosite coupling terms to the Hamiltonian, summing over lattice sites.
Represents couplings of the form \(\sum_{x_{0}, \ldots, x_{\text {dim-1 }}} \operatorname{strength}[\operatorname{shift}(\vec{x})] * O P 0 * O P 1\), where OPO \(:=\) lat.unit_cell[u0].get_op (op0) acts on the site (x_0, ..., x_\{dim-1\}, u1), and OP1 := lat.unit_cell[u1].get_op(op1) acts on the site (x_0+dx[0], ..., \(\left.x_{\_}\{\operatorname{dim}-1\}+d x[\operatorname{dim}-1], u 1\right)\). Possible combinations \(x_{-} 0, \ldots, x_{-}\{d i m-1\}\) are determined from the boundary conditions in possible_couplings ().

The coupling strength may vary spatially if the given strength is a numpy array. The correct shape of this array is the coupling_shape returned by tenpy.models.lattice.coupling_shape () and depends on the boundary conditions. The shift (...) depends on \(d x\), and is chosen such that the first entry strength \([0,0, \ldots]\) of strength is the prefactor for the first possible coupling fitting into the lattice if you imagine open boundary conditions.
The necessary terms are just added to coupling_terms; this function does not rebuild the MPO.
Deprecated since version 0.4.0: The arguments str_on_first and raise_op2_left will be removed in version 1.0.0.

\section*{Parameters}
- strength (scalar / array) - Prefactor of the coupling. May vary spatially (see above). If an array of smaller size is provided, it gets tiled to the required shape.
- u1 (int) - Picks the site lat. unit_cell[u1] for OP1.
- op1 (str) - Valid operator name of an onsite operator in lat.unit_cell[u1] for OP1.
- u2 (int) - Picks the site lat.unit_cell[u2] for OP2.
- op2 (str) - Valid operator name of an onsite operator in lat.unit_cell[u2] for OP2.
- dx (iterable of int) - Translation vector (of the unit cell) between OP1 and OP2. For a 1D lattice, a single int is also fine.
- op_string (str / None) - Name of an operator to be used between the OP1 and OP2 sites. Typical use case is the phase for a Jordan-Wigner transformation. The operator should be defined on all sites in the unit cell. If None, auto-determine whether a JordanWigner string is needed, using op_needs_JW ().
- str_on_first (bool) - Whether the provided op_string should also act on the first site. This option should be chosen as True for Jordan-Wigner strings. When handling Jordan-Wigner strings we need to extend the op_string to also act on the 'left', first site (in the sense of the MPS ordering of the sites given by the lattice). In this case, there is a well-defined ordering of the operators in the physical sense (i.e. which of op1 or op2 acts first on a given state). We follow the convention that op2 acts first (in the physical sense), independent of the MPS ordering. Deprecated.
- raise_op2_left (bool) - Raise an error when op2 appears left of opl (in the sense of the MPS ordering given by the lattice). Deprecated.
- category (str) - Descriptive name used as key for coupling_terms. Defaults to a string of the form " \(\{o p 1\} \_i \quad\{o p 2\} \_j "\).
- plus_hc (bool) - If True, the hermitian conjugate of the terms is added automatically.

\section*{Examples}

When initializing a model, you can add a term \(J \sum_{<i, j>} S_{i}^{z} S_{j}^{z}\) on all nearest-neighbor bonds of the lattice like this:
```

>>> J = 1. \# the strength
>>> for ul, u2, dx in self.lat.pairs['nearest_neighbors']:
... self.add_coupling(J, u1, 'Sz', u2, 'Sz', dx)

```

The strength can be an array, which gets tiled to the correct shape. For example, in a 1D Chain with an even number of sites and periodic (or infinite) boundary conditions, you can add alternating strong and weak couplings with a line like:
```

>>> self.add_coupling([1.5, 1.], u1, 'Sz', u2, 'Sz', dx)

```

Make sure to use the plus_hc argument if necessary, e.g. for hoppings:
```

>>> t = 1. \# hopping strength
>>> for u1, u2, dx in self.lat.pairs['nearest_neighbors']:
... self.add_coupling(t, ul, 'Cd', u2, 'C', dx, plus_hc=True)

```

Alternatively, you can add the hermitian conjugate terms explictly. The correct way is to complex conjugate the strength, take the hermitian conjugate of the operators and swap the order (including a swap \(u 1<->u 2\) ), and use the opposite direction \(-d x\), i.e. the h.c. of add_coupling ( \(t, \mathrm{u} 1\), 'A', u2, 'B', dx) is add_coupling(np.conj(t), u2, hc('B'), u1, hc('A'), -dx), where \(h c\) takes the hermitian conjugate of the operator names, see get_hc_op_name (). For spin-less fermions (FermionSite), this would be
```

>>> for u1, u2, dx in self.lat.pairs['nearest_neighbors']:
... self.add_coupling(t, u1, 'Cd', u2, 'C', dx)
... self.add_coupling(np.conj(t), u2, 'Cd', ul, 'C', -dx) \# h.c.

```

With spin-full fermions (SpinHalfFermions), it could be:
```

>>> for u1, u2, dx in self.lat.pairs['nearest_neighbors']:
... self.add_coupling(t, u1, 'Cdu', u2, 'Cd', dx) \# Cdagger_up C_down
... self.add_coupling(np.conj(t), u2, 'Cdd', u1, 'Cu', -dx) \# h.c.u
\hookrightarrowCdagger_down C_up

```

Note that the Jordan-Wigner strings for the fermions are added automatically!

\section*{See also:}
add_onsite Add terms acting on one site only.
add_multi_coupling_term for terms on more than two sites.
add_coupling_term Add a single term without summing over \(\vec{x}\).
add_coupling_term (strength, \(i, j, o p_{-} i, o p \_j, o p_{-}\)string \(=' I d^{\prime}\), category \(=\) None, plus_hc=False)
Add a two-site coupling term on given MPS sites.
Wrapper for self.coupling_terms[category].add_coupling_term(...).

\section*{Warning: This function does not handle Jordan-Wigner strings! You might want to use} add_local_term() instead.

\section*{Parameters}
- strength (float) - The strength of the coupling term.
- i (int) - The MPS indices of the two sites on which the operator acts. We require 0 \(<=i<N \_s i t e s\) and \(i<j\), i.e., op_i acts "left" of \(o p_{-} j\). If \(\mathrm{j}>=\mathrm{N}_{-}\)sites, it indicates couplings between unit cells of an infinite MPS.
- j (int) - The MPS indices of the two sites on which the operator acts. We require 0 \(<=i<N \_s i t e s\) and \(i<j\), i.e., op_i acts "left" of \(o p_{-} j\). If \(\mathrm{j}>=\mathrm{N}_{-}\)sites, it indicates couplings between unit cells of an infinite MPS.
- op1 (str) - Names of the involved operators.
- op2 (str) - Names of the involved operators.
- op_string (str) - The operator to be inserted between \(i\) and \(j\).
- category (str) - Descriptive name used as key for coupling_terms. Defaults to a string of the form " \(\{o p 1\} \_i \quad\{o p 2\} \_j "\).
- plus_hc (bool) - If True, the hermitian conjugate of the term is added automatically.
add_exponentially_decaying_coupling (strength, lambda_, op_i,op_j, subsites=None, \(o p_{-}\)string \(=\)None, \(p l u s \_h c=\) False \()\)
Add an exponentially decaying long-range coupling.
\[
\text { strength } \sum_{i<j} \lambda^{|i-j|} A_{\text {subsites }[i]} B_{\text {subsites }[j]}
\]

Where the operator \(A\) is given by \(o p_{-} i\), and \(B\) is given by \(o p_{-} j\). Note that the sum over \(i, j\) is long-range, for infinite systems going beyond the MPS unit cell. Moreover, note that the distance in the exponent is the distance within subsites.

Parameters
- strength (float) - Overall prefactor.
- lambda (float) - Decay-rate
- op_i (string) - Names for the operators.
- op_j (string) - Names for the operators.
- subsites (None / 1D array) - Selects a subset of sites within the MPS unit cell on which the operators act. Needs to be sorted. None selects all sites.
- op_string (None / str) - The operator to be inserted between \(A\) and \(B\); If None, this function checks whether a fermionic "JW" string is needed for the given operators; in this case the right \(o p_{-} j\) acts first.
- plus_hc (bool) - If True, the hermitian conjugate of the term is added automatically.

\section*{Examples}

At least for simple enough 1D chains (or ladders), you can use fit_with_sum_of_exp () to approximate a long-range function with a few sum of exponentials and then add them with this function.
```

>>> def decay(x):
... return np.exp(-0.1*x) / x**2
>>> from tenpy.tools.fit import fit_with_sum_of_exp, sum_of_exp
>>> n_exp = 5
>>> fit_range = 50
>>> lam, pref = fit_with_sum_of_exp(decay, n_exp, fit_range)
>>> x = np.arange(1, fit_range + 1)
>>> print('error in fit: {0:.3e}'.format(np.sum(np.abs(decay(x) - sum_of_
\hookrightarrowexp(lam, pref, x)))))
error in fit: 1.073e-04
>>> for pr, la in zip(pref, lam):
... self.add_exponentially_decaying_coupling(pr, la, 'N', 'N')

```
add_local_term (strength, term, category=None, plus_hc=False)
Add a single term to self.
The repesented term is strength times the product of the operators given in terms. Each operator is specified by the name and the site it acts on; the latter given by a lattice index, see Lattice.

Depending on the length of term, it can add an onsite term or a coupling term to onsite_terms or coupling_terms, respectively.

\section*{Parameters}
- strength (float/complex) - The prefactor of the term.
- term (list of (str, array_like)) - List of tuples (opname, lat_idx) where opname is a string describing the operator acting on the site given by the lattice index lat_idx. Here, lat_idx is for example \([x, y, u]\) for a 2D lattice, with \(u\) being the index within the unit cell.
- category - Descriptive name used as key for onsite_terms or coupling_terms.
- plus_hc (bool) - If True, the hermitian conjugate of the terms is added automatically.
add_multi_coupling (strength, ops, _deprecate_l='DEPRECATED', _deprecate_2 = 'DEPRECATED', op_string=None, category=None, plus_hc=False)
Add multi-site coupling terms to the Hamiltonian, summing over lattice sites.

Represents couplings of the form \(\operatorname{sum}_{\vec{x}} \operatorname{strength}[\operatorname{shift}(\vec{x})] * O P_{0} * O P_{1} * \ldots * O P_{M-1}\), involving \(M\) operators. Here, \(O P_{m}\) stands for the operator defined by the \(m\)-th tuple (opname, \(\mathrm{dx}, \mathrm{u}\) ) given in the argument ops, which determines the position \(\vec{x}+\overrightarrow{d x}\) and unit-cell index \(u\) of the site it acts on; the actual operator is given by self.lat.unit_cell[u].get_op(opname).
The coupling strength may vary spatially if the given strength is a numpy array. The correct shape of this array is the coupling_shape returned by tenpy.models.lattice. possible_multi_couplings() and depends on the boundary conditions. The shift (...) depends on the \(d x\) entries of ops and is chosen such that the first entry strength [0, 0, ...] of strength is the prefactor for the first possible coupling fitting into the lattice if you imagine open boundary conditions.

The necessary terms are just added to coupling_terms; this function does not rebuild the MPO.
Deprecated since version 0.6.0: We switched from the three arguments \(u 0\), op0 and other_op with other_ops=[(u1, op1, dx1), (op2, u2, dx2), ...] to a single, equivalent argment ops which should now read ops \(=[(\mathrm{op} 0, \mathrm{dx} 0, \mathrm{u} 0)\), (op1, \(\mathrm{dx} 1, \mathrm{u} 1)\), (op2, \(\mathrm{dx} 2, \mathrm{u} 2)\), . ..], where \(\mathrm{dx} 0=[0] *\) self.lat.dim. Note the changed order inside the tuples!

\section*{Parameters}
- strength (scalar / array) - Prefactor of the coupling. May vary spatially, and is tiled to the required shape.
- ops (list of (opname, \(d x, u)\) ) - Each tuple determines one operator of the coupling, see the description above. opname (str) is the name of the operator, \(d x\) (list of length lat.dim) is a translation vector, and \(u\) (int) is the index of lat.unit_cell on which the operator acts. The first entry of ops corresponds to \(O P_{0}\) and acts last in the physical sense.
- op_string (str / None) - If a string is given, we use this as the name of an operator to be used inbetween the operators, excluding the sites on which any operators act. This operator should be defined on all sites in the unit cell.

If None, auto-determine whether a Jordan-Wigner string is needed (using op_needs_JW ()) for each of the segments inbetween the operators and also on the sites of the left operators.
- category (str) - Descriptive name used as key for coupling_terms. Defaults to a string of the form " \(\{0 \mathrm{p} 0\}\) _i \(\left\{o t h e r \_o p s[0]\right\} \_j\) \{other_ops [1]\}_k ...".
- plus_hc (bool) - If True, the hermitian conjugate of the terms is added automatically.

\section*{Examples}

A call to add_coupling() with arguments add_coupling(strength, u1, 'A', u2, ' \(B^{\prime}, d x\) ) is equivalent to the following:
```

>>> dx_0 = [0] * self.lat.dim \# = [0] for a 1D lattice, [0, 0] in 2D
>>> self.add_multi_coupling(strength, [('A', dx_0, u1), ('B', dx, u2)])

```

To explicitly add the hermitian conjugate (instead of simply using plus_hc = True), you need to take the complex conjugate of the strength, reverse the order of the operators and take the hermitian conjugates of the individual operator names (indicated by the hc (...), see get_hc_op_name ()):
```

>>> self.add_multi_coupling(np.conj(strength), [(hc('B'), dx, u2), (hc('A'), b
@dx_0, u1)])

```

\section*{See also:}
add_onsite Add terms acting on one site only.
add_coupling Add terms acting on two sites.
add_multi_coupling_term Add a single term, not summing over the possible \(\vec{x}\).
add_multi_coupling_term (strength, ijkl,ops_ijkl,op_string, category=None, plus_hc=False)
Add a general M-site coupling term on given MPS sites.
Wrapper for self.coupling_terms[category].add_multi_coupling_term(....).

\section*{Warning: This function does not handle Jordan-Wigner strings! You might want to use add_local_term() instead.}

\section*{Parameters}
- strength (float) - The strength of the coupling term.
- ijkl (list of int) - The MPS indices of the sites on which the operators acts. With \(i, j, k, \ldots=i j k l\), we require that they are ordered ascending, \(i<j<k<\ldots\) and that \(0<=i<N \_s i t e s\). Inidces \(>=\) N_sites indicate couplings between different unit cells of an infinite MPS.
- ops_ijkl (list of str) - Names of the involved operators on sites \(i, j, k, \ldots\).
- op_string (list of str) - Names of the operator to be inserted between the operators, e.g., op_string[0] is inserted between \(i\) and \(j\).
- category (str) - Descriptive name used as key for coupling_terms. Defaults to a string of the form " \(\{\mathrm{op} 0\} \_\)i \(\{o p 1\} \_j \quad\{o p 2\} \_k \ldots\). .
- plus_hc (bool) - If True, the hermitian conjugate of the term is added automatically.
add_onsite (strength, \(u\), opname, category=None, plus_hc=False)
Add onsite terms to onsite_terms.
Adds \(\sum_{\vec{x}}\) strength \([\vec{x}] * O P\) to the represented Hamiltonian, where the operator \(O P=1\) at. unit_cell[u].get_op (opname) acts on the site given by a lattice index (x_0, ..., \(\left.x_{-}\{\operatorname{dim}-1\}, u\right)\),
The necessary terms are just added to onsite_terms; doesn't rebuild the MPO.

\section*{Parameters}
- strength (scalar / array) - Prefactor of the onsite term. May vary spatially. If an array of smaller size is provided, it gets tiled to the required shape.
- u(int) - Picks a Site lat.unit_cell [u] out of the unit cell.
- opname (str) - valid operator name of an onsite operator in lat. unit_cell[u].
- category (str) - Descriptive name used as key for onsite_terms. Defaults to opname.
- plus_hc (bool) - If True, the hermitian conjugate of the terms is added automatically.

\section*{See also:}
add_coupling Add a terms acting on two sites.
add_onsite_term Add a single term without summing over vecx.
add_onsite_term (strength, \(i\), op, category=None, plus_hc=False)
Add an onsite term on a given MPS site.
Wrapper for self.onsite_terms[category].add_onsite_term(....).

\section*{Parameters}
- strength (float) - The strength of the term.
- i (int) - The MPS index of the site on which the operator acts. We require \(0<=i<\) L.
- op (str) - Name of the involved operator.
- category (str) - Descriptive name used as key for onsite_terms. Defaults to op.
- plus_hc (bool) - If True, the hermitian conjugate of the term is added automatically.
all_coupling_terms()
Sum of all coupling_terms.
all_onsite_terms()
Sum of all onsite_terms.
bond_energies ( \(p s i\) )
Calculate bond energies <psilH_bondlpsi>.
Parameters psi (MPS) - The MPS for which the bond energies should be calculated.
Returns E_bond - List of bond energies: for finite bc, E_Bond[i] is the energy of bond i, i+1. (i.e. we omit bond 0 between sites L-1 and 0 ); for infinite bc E_bond[i] is the energy of bond \(i-1, i\).

Return type 1D ndarray
calc_H_MPO (tol_zero \(=1 e-15\) )
Calculate MPO representation of the Hamiltonian.
Uses onsite_terms and coupling_terms to build an MPOGraph (and then an MPO).
Parameters tol_zero (float) - Prefactors with abs (strength) < tol_zero are considered to be zero.

Returns H_MPO - MPO representation of the Hamiltonian.
Return type MPO
calc_H_MPO_from_bond (tol_zero=1e-15)
Calculate the MPO Hamiltonian from the bond Hamiltonian.
Parameters tol_zero (float) - Arrays with norm <tol_zero are considered to be zero.
Returns H_MPO - MPO representation of the Hamiltonian.
Return type MPO
calc_H_bond (tol_zero=1e-15)
calculate \(H\) _bond from coupling_terms and onsite_terms.
Parameters tol_zero (float) - prefactors with abs(strength) < tol_zero are considered to be zero.

Returns H_bond - Bond terms as required by the constructor of NearestNeighborModel. Legs are ['p0', 'p0*', 'p1', 'p1*']

Return type list of Array
:raises ValueError : if the Hamiltonian contains longer-range terms.:
calc_H_bond_from_MPO (tol_zero=le-15)
Calculate the bond Hamiltonian from the MPO Hamiltonian.
Parameters tol_zero (float) - Arrays with norm <tol_zero are considered to be zero.
Returns H_bond - Bond terms as required by the constructor of NearestNeighborModel. Legs are ['p0', 'p0*', 'p1', 'p1*']

Return type list of Array
:raises ValueError : if the Hamiltonian contains longer-range terms.:
calc_H_onsite (tol_zero=1e-15)
Calculate H_onsite from self.onsite_terms.
Deprecated since version 0.4.0: This function will be removed in 1.0.0. Replace calls to this function by self.all_onsite_terms().remove_zeros(tol_zero).to_Arrays(self. lat.mps_sites()). You might also want to take explicit_plus_hc into account.

Parameters tol_zero (float) - prefactors with abs (strength) < tol_zero are considered to be zero.

\section*{Returns}
- H_onsite (list of npc.Array)
- onsite terms of the Hamiltonian. If explicit_plus_hc is True, - Hermitian conjugates of the onsite terms will be included.
coupling_strength_add_ext_flux (strength, \(d x\), phase)
Add an external flux to the coupling strength.
When performing DMRG on a "cylinder" geometry, it might be useful to put an "external flux" through the cylinder. This means that a particle hopping around the cylinder should pick up a phase given by the external flux [[resta1998]]. This is also called "twisted boundary conditions" in literature. This function adds a complex phase to the strength array on some bonds, such that particles hopping in positive direction around the cylinder pick up \(\exp (+i\) phase \()\).

Warning: For the sign of phase it is important that you consistently use the creation operator as opl and the annihilation operator as op2 in add_coupling().

\section*{Parameters}
- strength (scalar / array) - The strength to be used in add_coupling(), when no external flux would be present.
- \(\mathbf{d x}\) (iterable of int) - Translation vector (of the unit cell) between opl and op2 in add_coupling().
- phase (iterable of float) - The phase of the external flux for hopping in each direction of the lattice. E.g., if you want flux through the cylinder on which you have an infinite MPS, you should give phase=[0, phi] souch that particles pick up a phase phi when hopping around the cylinder.

Returns strength - The strength array to be used as strength in add_coupling() with the given \(d x\).

Return type complex array

\section*{Examples}

Let's say you have an infinite MPS on a cylinder, and want to add nearest-neighbor hopping of fermions with the Fermionsite. The cylinder axis is the \(x\)-direction of the lattice, so to put a flux through the cylinder, you want particles hopping around the cylinder to pick up a phase phi given by the external flux.
```

>>> strength = 1. \# hopping strength without external flux
>>> phi = np.pi/4 \# determines the external flux strength
>>> for ul, u2, dx in self.lat.pairs['nearest_neighbors']:
... strength_with_flux = self.coupling_strength_add_ext_flux(strength, dx,
\hookrightarrow [0, phi])
... self.add_coupling(strength_with_flux, u1, 'Cd', u2, 'C', dx)
... self.add_coupling(np.conj(strength_with_flux), u2, 'Cd', ul, 'C', -dx)

```
enlarge_mps_unit_cell (factor=2)
Repeat the unit cell for infinite MPS boundary conditions; in place.
This has to be done after finishing initialization and can not be reverted.
Parameters factor (int) - The new number of sites in the MPS unit cell will be increased from \(N\) _sites to factor*N_sites_per_ring. Since MPS unit cells are repeated in the \(x\)-direction in our convetion, the lattice shape goes from ( \(\mathrm{Lx}, \mathrm{Ly}, \ldots, \mathrm{Lu}\) ) to (Lx*factor, Ly, ..., Lu).

\section*{classmethod from_MPOModel (mpo_model)}

Initialize a NearestNeighborModel from a model class defining an MPO.
This is especially usefull in combination with MPOModel.group_sites ().
Parameters mpo_model (MPOModel) - A model instance implementing the MPO. Does not need to be a NearestNe ighborModel, but should only have nearest-neighbor couplings.

\section*{Examples}

The SpinChainNNN2 has next-nearest-neighbor couplings and thus only implements an MPO:
```

>>> from tenpy.models.spins_nnn import SpinChainNNN2
>>> nnn__chain = SpinChainNNN2 ({'L': 20})
>>> print(isinstance(nnn_chain, NearestNeighborModel))
False
>>> print("range before grouping:", nnn_chain.H_MPO.max_range)
range before grouping: 2

```

By grouping each two neighboring sites, we can bring it down to nearest neighbors.
```

>>> grouped_sites = nnn_chain.group_sites(2)
>>> print("range after grouping:", nnn_chain.H_MPO.max_range)
range after grouping: 1

```

Yet, TEBD will not yet work, as the model doesn't define \(H_{-}\)bond. However, we can initialize a NearestNeighborModel from the MPO:
```

>>> nnn_chain_for_tebd = NearestNeighborModel.from_MPOModel(nnn_chain)
>>> isinstance(nnn_chain_for_tebd, NearestNeighborModel)
True

```
classmethod from_hdf5 (hdf5_loader, h5gr, subpath)
Load instance from a HDF5 file.
This method reconstructs a class instance from the data saved with save_hdf5 ().

\section*{Parameters}
- hdf5_loader (Hdf5Loader) - Instance of the loading engine.
- h5gr (Group) - HDF5 group which is represent the object to be constructed.
- subpath (str) - The name of \(h 5 g r\) with a '/' in the end.

Returns obj - Newly generated class instance containing the required data.
Return type cls
group_sites ( \(n=2\), grouped_sites=None)
Modify self in place to group sites.
Group each \(n\) sites together using the GroupedSite. This might allow to do TEBD with a Trotter decomposition, or help the convergence of DMRG (in case of too long range interactions).
This has to be done after finishing initialization and can not be reverted.

\section*{Parameters}
- \(\mathbf{n}(i n t)\) - Number of sites to be grouped together.
- grouped_sites (None l list of GroupedSite) - The sites grouped together.

Returns grouped_sites - The sites grouped together.
Return type list of GroupedSite

\section*{init_H_from_terms()}

Initialize \(H_{-} M P O\) (and \(H_{-}\)bond) from the terms of the CouplingModel.
This function is called automatically during CouplingMPOModel.__init__.
\(\qquad\)
If you use one of the add_* methods of the CouplingModel after initialization, you will need to call init_H_from_terms in the end by yourself, in order to update the \(H_{-} M P O\) (and possibly \(H_{-} b o n d\) ) representations. (You should get a warning about this... The way to avoid it is to initialize all the terms in init_terms by defining your own model, as outlined in Models.

\section*{init_lattice (model_params)}

Initialize a lattice for the given model parameters.
This function reads out the model parameter lattice. This can be a full Lattice instance, in which case it is just returned without further action. Alternatively, the lattice parameter can be a string giving the name of one of the predefined lattices, which then gets initialized. Depending on the dimensionality of the lattice, this requires different model parameters.

Parameters model_params (dict) - The model parameters given to \(\qquad\) init \(\qquad\)
Returns lat - An initialized lattice.
Return type Lattice

\section*{Options}
option CouplingMPOModel.lattice: str | Lattice
The name of a lattice pre-defined in TeNPy to be initialized. Alternatively, directly a subclass of Lattice instead of the name. Alternatively, a (possibly self-defined) Lattice instance. In the latter case, no further parameters are read out.
option CouplingMPOModel.bc_MPS: str
Boundary conditions for the MPS.
option CouplingMPOModel.order: str
The order of sites within the lattice for non-trivial lattices, e.g, 'default', 'snake', see ordering (). Only used if lattice is a string.
option CouplingMPOModel.L: int
The length in x-direction; only read out for 1D lattices. For an infinite system the length of the unit cell.
option CouplingMPOModel.Lx: int
option CouplingMPOModel.Ly: int
The length in x - and y-direction; only read out for 2D lattices. For "infinite" \(b c_{-} M P S\), the system is infinite in x-direction and \(L x\) is the number of "rings" in the infinite MPS unit cell, while \(L y\) gives the circumference around the cylinder or width of th the rung for a ladder (depending on \(b c_{-} y\) ).
option CouplingMPOModel.bc_y: str
"cylinder" | "ladder"; only read out for 2D lattices. The boundary conditions in y-direction.
option CouplingMPOModel.bc_x: str
"open" | "periodic". Can be used to force "periodic" boundaries for the lattice, i.e., for the couplings in the Hamiltonian, even if the MPS is finite. Defaults to "open" for bc_MPS="finite " and "periodic" for bc_MPS="infinite. If you are not aware of the consequences, you should probably not use "periodic" boundary conditions. (The MPS is still "open", so this will introduce long-range couplings between the first and last sites of the MPS!)

\section*{init_sites (model_params)}

Define the local Hilbert space and operators; needs to be implemented in subclasses.
This function gets called by init_lattice () to get the Site for the lattice unit cell.

Note: Initializing the sites requires to define the conserved quantum numbers. All pre-defined sites accept conserve=None to disable using quantum numbers. Many models in TeNPy read out the conserve model parameter, which can be set to "best " to indicate the optimal parameters.

If you need to initialize more than one site, the function tenpy.networks.site. set_common_charges () should be helpful.

Parameters model_params (dict) - The model parameters given to \(\qquad\) __.

Returns sites - The local sites of the lattice, defining the local basis states and operators.
Return type (tuple of) Site
init_terms (model_params)
Add the onsite and coupling terms to the model; subclasses should implement this.
save_hdf5 (hdf5_saver, h5gr, subpath)
Export self into a HDF5 file.
This method saves all the data it needs to reconstruct self with from_hdf5 ().

This implementation saves the content of \(\qquad\) dict \(\qquad\) with save_dict_content (), storing the format under the attribute 'format'.

\section*{Parameters}
- hdf5_saver (Hdf5Saver) - Instance of the saving engine.
- h5gr (: class `Group`) - HDF5 group which is supposed to represent self.
- subpath (str) - The name of \(h 5 g r\) with a ' /' in the end.
```

test_sanity()

```

Sanity check, raises ValueErrors, if something is wrong.
trivial_like_NNModel()
Return a NearestNeighborModel with same lattice, but trivial ( \(\mathrm{H}=0\) ) bonds.

\subsection*{19.8.2 FermiHubbardChain}
- full name: tenpy.models.hubbard.FermiHubbardChain
- parent module: tenpy.models.hubbard
- type: class

\section*{Inheritance Diagram}


\section*{Methods}
\begin{tabular}{|c|c|}
\hline FermiHubbardChain.__init__(model_params) & Initialize self. \\
\hline FermiHubbardChain.add_coupling(strength, u1,...) & Add twosite coupling terms to the Hamiltonian, summing over lattice sites. \\
\hline \begin{tabular}{l}
FermiHubbardChain. \\
add_coupling_term(...[,...])
\end{tabular} & Add a two-site coupling term on given MPS sites. \\
\hline \begin{tabular}{l}
FermiHubbardChain. \\
add_exponentially_decaying_coupling(...)
\end{tabular} & Add an exponentially decaying long-range coupling. \\
\hline FermiHubbardChain. add_local_term(strength, term) & Add a single term to self. \\
\hline \[
\begin{aligned}
& \text { FermiHubbardChain. } \\
& \text { add_multi_coupling(...[, ...]) }
\end{aligned}
\] & Add multi-site coupling terms to the Hamiltonian, summing over lattice sites. \\
\hline \begin{tabular}{l}
FermiHubbardChain. \\
add_multi_coupling_term(...)
\end{tabular} & Add a general M-site coupling term on given MPS sites. \\
\hline
\end{tabular}

Table 54 - continued from previous page
FermiHubbardChain.add_onsite(strength, u, Add onsite terms to onsite_terms. opname)
\begin{tabular}{|c|c|}
\hline \begin{tabular}{l}
FermiHubbardChain. \\
add_onsite_term(strength, ...)
\end{tabular} & Add an onsite term on a given MPS site. \\
\hline \begin{tabular}{l}
FermiHubbardChain. \\
all_coupling_terms()
\end{tabular} & Sum of all coupling_terms. \\
\hline FermiHubbardChain.all_onsite_terms() & Sum of all onsite_terms. \\
\hline FermiHubbardChain.bond_energies(psi) & Calculate bond energies <psilH_bondlpsi>. \\
\hline FermiHubbardChain.calc_H_MPO([tol_zero]) & Calculate MPO representation of the Hamiltonian. \\
\hline \begin{tabular}{l}
FermiHubbardChain. \\
calc_H_MPO_from_bond([...])
\end{tabular} & Calculate the MPO Hamiltonian from the bond Hamiltonian. \\
\hline FermiHubbardChain.calc_H_bond([tol_zero]) & calculate H_bond from coupling_terms and onsite_terms. \\
\hline FermiHubbardChain. calc_H_bond_from_MPO([...]) & Calculate the bond Hamiltonian from the MPO Hamiltonian. \\
\hline FermiHubbardChain. calc_H_onsite([tol_zero]) & Calculate H_onsite from self.onsite_terms. \\
\hline FermiHubbardChain. coupling_strength_add_ext_flux(...) & Add an external flux to the coupling strength. \\
\hline \begin{tabular}{l}
FermiHubbardChain. \\
enlarge_mps_unit_cell([factor])
\end{tabular} & Repeat the unit cell for infinite MPS boundary conditions; in place. \\
\hline FermiHubbardChain. from_MPOModel(mpo_model) & Initialize a NearestNeighborModel from a model class defining an MPO. \\
\hline FermiHubbardChain.from_hdf5(hdf5_loader, ...) & Load instance from a HDF5 file. \\
\hline ```
FermiHubbardChain.group_sites([n,
grouped_sites])
``` & Modify self in place to group sites. \\
\hline FermiHubbardChain.init_H_from_terms() & Initialize \(H_{-} M P O\) (and \(H_{-}\)bond) from the terms of the CouplingModel. \\
\hline FermiHubbardChain. init_lattice(model_params) & Initialize a lattice for the given model parameters. \\
\hline FermiHubbardChain. init_sites(model_params) & Define the local Hilbert space and operators; needs to be implemented in subclasses. \\
\hline \begin{tabular}{l}
FermiHubbardChain. \\
init_terms(model_params)
\end{tabular} & Add the onsite and coupling terms to the model; subclasses should implement this. \\
\hline FermiHubbardChain.save_hdf5(hdf5_saver, ...) & Export self into a HDF5 file. \\
\hline FermiHubbardChain.test_sanity() & Sanity check, raises ValueErrors, if something is wrong. \\
\hline FermiHubbardChain. trivial_like_NNModel() & Return a NearestNeighborModel with same lattice, but trivial \((\mathrm{H}=0)\) bonds. \\
\hline
\end{tabular}

\section*{Class Attributes and Properties}
\begin{tabular}{ll}
\hline \begin{tabular}{l} 
FermiHubbardChain. \\
force_default_lattice
\end{tabular} & \begin{tabular}{l} 
If True, init_lattice() asserts that the initialized \\
lattice is (a subclass of)default_lattice
\end{tabular} \\
\hline FermiHubbardChain.logger & class attribute. \\
\hline FermiHubbardChain. verbose &
\end{tabular}

FermiHubbardChain.verbose
class tenpy.models.hubbard.FermiHubbardChain (model_params)

Bases: tenpy.models.hubbard.FermiHubbardModel, tenpy.models.model. NearestNeighborModel

The FermiHubbardModel on a Chain, suitable for TEBD.
See the FermiHubbardModel for the documentation of parameters.

\section*{default_lattice}
alias of tenpy.models.lattice. Chain
add_coupling (strength, u1, op1, u2, op2, dx, op_string=None, str_on_first=True, raise_op2_left=False, category=None, plus_hc=False)
Add twosite coupling terms to the Hamiltonian, summing over lattice sites.
Represents couplings of the form \(\sum_{x_{0}, \ldots, x_{\text {dim-1 }}}\) strength \([\operatorname{shift}(\vec{x})] * O P 0 * O P 1\), where OP 0 : = lat.unit_cell[u0].get_op(op0) acts on the site (x_0, ..., x_\{dim-1\}, u1), and OP1 := lat.unit_cell[u1].get_op(op1) acts on the site (x_0+dx[0], ..., \(x_{-}\{\operatorname{dim}-1\}+d x[\operatorname{dim}-1]\), u1). Possible combinations \(x_{\_} 0, \ldots, x_{-}\{d i m-1\}\) are determined from the boundary conditions in possible_couplings ().

The coupling strength may vary spatially if the given strength is a numpy array. The correct shape of this array is the coupling_shape returned by tenpy.models.lattice.coupling_shape () and depends on the boundary conditions. The shift (. . .) depends on \(d x\), and is chosen such that the first entry strength \([0,0, \ldots]\) of strength is the prefactor for the first possible coupling fitting into the lattice if you imagine open boundary conditions.
The necessary terms are just added to coupling_terms; this function does not rebuild the MPO.
Deprecated since version 0.4.0: The arguments str_on_first and raise_op2_left will be removed in version 1.0.0.

\section*{Parameters}
- strength (scalar / array) - Prefactor of the coupling. May vary spatially (see above). If an array of smaller size is provided, it gets tiled to the required shape.
- u1 (int) - Picks the site lat.unit_cell[u1] for OP1.
- op1 (str) - Valid operator name of an onsite operator in lat.unit_cell[ul] for OP1.
- u2 (int) - Picks the site lat. unit_cell[u2] for OP2.
- op2 (str) - Valid operator name of an onsite operator in lat.unit_cell[u2] for OP2.
- dx (iterable of int) - Translation vector (of the unit cell) between OP1 and OP2. For a 1D lattice, a single int is also fine.
- op_string (str / None) - Name of an operator to be used between the OP1 and OP2 sites. Typical use case is the phase for a Jordan-Wigner transformation. The operator should be defined on all sites in the unit cell. If None, auto-determine whether a JordanWigner string is needed, using op_needs_JW ().
- str_on_first (bool) - Whether the provided op_string should also act on the first site. This option should be chosen as True for Jordan-Wigner strings. When handling Jordan-Wigner strings we need to extend the op_string to also act on the 'left', first site (in the sense of the MPS ordering of the sites given by the lattice). In this case, there is a well-defined ordering of the operators in the physical sense (i.e. which of opl or op2 acts first on a given state). We follow the convention that op2 acts first (in the physical sense), independent of the MPS ordering. Deprecated.
- raise_op2_left (bool) - Raise an error when op2 appears left of opl (in the sense of the MPS ordering given by the lattice). Deprecated.
- category (str) - Descriptive name used as key for coupling_terms. Defaults to a string of the form " \(\{0 \mathrm{p} 1\} \_i \quad\{o p 2\} \_j "\).
- plus_hc (bo○l) - If True, the hermitian conjugate of the terms is added automatically.

\section*{Examples}

When initializing a model, you can add a term \(J \sum_{<i, j>} S_{i}^{z} S_{j}^{z}\) on all nearest-neighbor bonds of the lattice like this:
```

>>> J = 1. \# the strength
>>> for ul, u2, dx in self.lat.pairs['nearest_neighbors']:
self.add_coupling(J, u1, 'Sz', u2, 'Sz', dx)

```

The strength can be an array, which gets tiled to the correct shape. For example, in a 1D Chain with an even number of sites and periodic (or infinite) boundary conditions, you can add alternating strong and weak couplings with a line like:
```

>>> self.add_coupling([1.5, 1.], u1, 'Sz', u2, 'Sz', dx)

```

Make sure to use the plus_hc argument if necessary, e.g. for hoppings:
```

>>> t = 1. \# hopping strength
>>> for ul, u2, dx in self.lat.pairs['nearest_neighbors']:
... self.add_coupling(t, u1, 'Cd', u2, 'C', dx, plus_hc=True)

```

Alternatively, you can add the hermitian conjugate terms explictly. The correct way is to complex conjugate the strength, take the hermitian conjugate of the operators and swap the order (including a swap \(u 1<->u 2\) ), and use the opposite direction -dx , i.e. the h.c. of add_coupling (t, u1, 'A', u2, 'B', dx) is add_coupling(np.conj(t), u2, hc('B'), u1, hc('A'), -dx), where \(h c\) takes the hermitian conjugate of the operator names, see \(g e t \_h c \_o p \_n a m e()\). For spin-less fermions (FermionSite), this would be
```

>>> for u1, u2, dx in self.lat.pairs['nearest_neighbors']:
... self.add_coupling(t, u1, 'Cd', u2, 'C', dx)
... self.add_coupling(np.conj(t), u2, 'Cd', u1, 'C', -dx) \# h.c.

```

With spin-full fermions (SpinHalfFermions), it could be:
```

>>> for u1, u2, dx in self.lat.pairs['nearest_neighbors']:
... self.add_coupling(t, u1, 'Cdu', u2, 'Cd', dx) \# Cdagger_up C_down
... self.add_coupling(np.conj(t), u2, 'Cdd', u1, 'Cu', -dx) \# h.c.s
\hookrightarrowCdagger_down C_up

```

Note that the Jordan-Wigner strings for the fermions are added automatically!

\section*{See also:}
add_onsite Add terms acting on one site only.
add_multi_coupling_term for terms on more than two sites.
add_coupling_term Add a single term without summing over \(\vec{x}\).
add_coupling_term (strength, \(i, j, o p_{-} i, o p \_j\), op_string='Id', category=None, plus_hc=False)
Add a two-site coupling term on given MPS sites.
Wrapper for self.coupling_terms[category].add_coupling_term(...).

Warning: This function does not handle Jordan-Wigner strings! You might want to use add_local_term() instead.

\section*{Parameters}
- strength (float) - The strength of the coupling term.
- i (int) - The MPS indices of the two sites on which the operator acts. We require 0 \(<=i<N \_s i t e s\) and \(i<j\), i.e., op_i acts "left" of \(o p_{-} j\). If \(\mathrm{j}>=\mathrm{N}_{-}\)sites, it indicates couplings between unit cells of an infinite MPS.
- j (int) - The MPS indices of the two sites on which the operator acts. We require 0 <= i < N_sites and \(i<j\), i.e., op_i acts "left" of \(o p_{-} j\). If \(j>=N \_\)sites, it indicates couplings between unit cells of an infinite MPS.
- op1 (str) - Names of the involved operators.
- op2 (str) - Names of the involved operators.
- op_string (str) - The operator to be inserted between \(i\) and \(j\).
- category (str) - Descriptive name used as key for coupling_terms. Defaults to a string of the form " \(\{o p 1\} \_i \quad\{o p 2\} \_j "\).
- plus_hc (bool) - If True, the hermitian conjugate of the term is added automatically.
add_exponentially_decaying_coupling (strength, lambda_, op_i,op_j, subsites=None, op_string \(=\) None, plus_hc=False)
Add an exponentially decaying long-range coupling.
\[
\text { strength } \sum_{i<j} \lambda^{|i-j|} A_{\text {subsites }[i]} B_{\text {subsites }[j]}
\]

Where the operator \(A\) is given by \(o p_{-} i\), and \(B\) is given by \(o p_{-} j\). Note that the sum over \(i, j\) is long-range, for infinite systems going beyond the MPS unit cell. Moreover, note that the distance in the exponent is the distance within subsites.

\section*{Parameters}
- strength (float) - Overall prefactor.
- lambda (float) - Decay-rate
- op_i (string) - Names for the operators.
- op_j (string) - Names for the operators.
- subsites (None / ID array) - Selects a subset of sites within the MPS unit cell on which the operators act. Needs to be sorted. None selects all sites.
- op_string (None / str) - The operator to be inserted between \(A\) and \(B\); If None, this function checks whether a fermionic "JW" string is needed for the given operators; in this case the right \(o p_{-} j\) acts first.
- plus_hc (bool) - If True, the hermitian conjugate of the term is added automatically.

\section*{Examples}

At least for simple enough 1D chains (or ladders), you can use fit_with_sum_of_exp () to approximate a long-range function with a few sum of exponentials and then add them with this function.
```

>>> def decay (x):
... return np.exp (-0.1*x) / x**2
>>> from tenpy.tools.fit import fit_with_sum_of_exp, sum_of__exp
>>> n__exp = 5
>>> fit_range = 50
>>> lam, pref = fit__with__sum_of_exp(decay, n_exp, fit_range)
>>> x = np.arange(1, fit_range + 1)
>> print('error in fit: {0:.3e}'.format(np.sum(np.abs(decay(x) - sum_of_
\hookrightarrowexp(lam, pref, x)))))
error in fit: 1.073e-04
>>> for pr, la in zip(pref, lam):
... self.add_exponentially_decaying_coupling(pr, la, 'N', 'N')

```
add_local_term (strength, term, category=None, plus_hc=False)
Add a single term to self.
The repesented term is strength times the product of the operators given in terms. Each operator is specified by the name and the site it acts on; the latter given by a lattice index, see Lattice.
Depending on the length of term, it can add an onsite term or a coupling term to onsite_terms or coupling_terms, respectively.

\section*{Parameters}
- strength (float/complex) - The prefactor of the term.
- term (list of (str, array_like)) - List of tuples (opname, lat_idx) where opname is a string describing the operator acting on the site given by the lattice index lat_idx. Here, lat_idx is for example \([x, y, u]\) for a 2D lattice, with \(u\) being the index within the unit cell.
- category - Descriptive name used as key for onsite_terms or coupling_terms.
- plus_hc (bool) - If True, the hermitian conjugate of the terms is added automatically.
add_multi_coupling (strength, ops, _deprecate_l='DEPRECATED', _deprecate_2='DEPRECATED', op_string=None, category=None, plus_hc=False ) Add multi-site coupling terms to the Hamiltonian, summing over lattice sites.
Represents couplings of the form \(\operatorname{sum}_{\vec{x}} \operatorname{strength}[\operatorname{shift}(\vec{x})] * O P_{0} * O P_{1} * \ldots * O P_{M-1}\), involving \(M\) operators. Here, \(O P_{m}\) stands for the operator defined by the \(m\)-th tuple (opname, \(\mathrm{dx}, \mathrm{u}\) ) given in the argument ops, which determines the position \(\vec{x}+\overrightarrow{d x}\) and unit-cell index \(u\) of the site it acts on; the actual operator is given by self.lat.unit_cell[u].get_op(opname).

The coupling strength may vary spatially if the given strength is a numpy array. The correct shape of this array is the coupling_shape returned by tenpy.models.lattice. possible_multi_couplings() and depends on the boundary conditions. The shift (...) depends on the \(d x\) entries of ops and is chosen such that the first entry strength [0, 0, ...] of strength is the prefactor for the first possible coupling fitting into the lattice if you imagine open boundary conditions.
The necessary terms are just added to coupling_terms; this function does not rebuild the MPO.

Deprecated since version 0.6.0: We switched from the three arguments \(u 0\), op0 and other_op with other_ops=[(u1, op1, dx1), (op2, u2, dx2), ...] to a single, equivalent argment ops which should now read ops \(=[(\mathrm{op} 0, \mathrm{dx} 0, \mathrm{u} 0)\), (op1, dx1, u1), (op2, dx2, u2), . ..], where \(d x 0=[0] * s e l f . l a t . d i m\). Note the changed order inside the tuples!

\section*{Parameters}
- strength (scalar / array) - Prefactor of the coupling. May vary spatially, and is tiled to the required shape.
- ops (list of (opname, \(d x, u)\) ) - Each tuple determines one operator of the coupling, see the description above. opname (str) is the name of the operator, \(d x\) (list of length lat.dim) is a translation vector, and \(u\) (int) is the index of lat.unit_cell on which the operator acts. The first entry of ops corresponds to \(O P_{0}\) and acts last in the physical sense.
- op_string (str / None) - If a string is given, we use this as the name of an operator to be used inbetween the operators, excluding the sites on which any operators act. This operator should be defined on all sites in the unit cell.
If None, auto-determine whether a Jordan-Wigner string is needed (using op_needs_JW ()) for each of the segments inbetween the operators and also on the sites of the left operators.
- category (str) - Descriptive name used as key for coupling_terms. Defaults to a string of the form " \(\{0 \mathrm{p} 0\}\) _i \(\{\) other_ops [0]\}_j \(\{\) other_ops [1] \}_k ...".
- plus_hc (bool) - If True, the hermitian conjugate of the terms is added automatically.

\section*{Examples}

A call to add_coupling() with arguments add_coupling(strength, u1, 'A', u2, ' \(B\) ', \(d x\) ) is equivalent to the following:
```

>>> dx_0 = [0] * self.lat.dim \# = [0] for a 1D lattice, [0, 0] in 2D
>>> self.add_multi_coupling(strength, [('A', dx_0, u1), ('B', dx, u2)])

```

To explicitly add the hermitian conjugate (instead of simply using plus_hc = True), you need to take the complex conjugate of the strength, reverse the order of the operators and take the hermitian conjugates of the individual operator names (indicated by the hc (...), see get_hc_op_name ()):
```

>>> self.add_multi_coupling(np.conj(strength), [(hc('B'), dx, u2), (hc('A'),
->dx_0, u1)])

```

\section*{See also:}
add_onsite Add terms acting on one site only.
add_coupling Add terms acting on two sites.
add_multi_coupling_term Add a single term, not summing over the possible \(\vec{x}\).
add_multi_coupling_term (strength, ijkl,ops_ijkl,op_string, category=None, plus_hc=False)
Add a general M-site coupling term on given MPS sites.
Wrapper for self.coupling_terms[category].add_multi_coupling_term(...).
```

Warning: This function does not handle Jordan-Wigner strings! You might want to use
add_local_term() instead.

```

\section*{Parameters}
- strength (float) - The strength of the coupling term.
- ijkl (list of int) - The MPS indices of the sites on which the operators acts. With \(i, j, k, \ldots=i j k l\), we require that they are ordered ascending, \(i<j<k<\ldots\) and that \(0<=\) i \(<\) N_sites. Inidces \(>=\) N_sites indicate couplings between different unit cells of an infinite MPS.
- ops_ijkl (list of str) - Names of the involved operators on sites \(i, j, k, \ldots\).
- op_string (list of str) - Names of the operator to be inserted between the operators, e.g., op_string[0] is inserted between \(i\) and \(j\).
- category (str) - Descriptive name used as key for coupling_terms. Defaults to a string of the form " \(\{\mathrm{op} 0\} \_i \quad\{o p 1\} \_j \quad\{o p 2\} \_k \quad .\). ".
- plus_hc (bool) - If True, the hermitian conjugate of the term is added automatically.
add_onsite (strength, \(u\), opname, category=None, plus_hc=False)
Add onsite terms to onsite_terms.
Adds \(\sum_{\vec{x}} \operatorname{strength}[\vec{x}] * O P\) to the represented Hamiltonian, where the operator \(\mathrm{OP}=1\) at. unit_cell[u].get_op(opname) acts on the site given by a lattice index (x_0, ..., \(\left.x_{-}\{\operatorname{dim}-1\}, u\right)\),
The necessary terms are just added to onsite_terms; doesn't rebuild the MPO.

\section*{Parameters}
- strength (scalar / array) - Prefactor of the onsite term. May vary spatially. If an array of smaller size is provided, it gets tiled to the required shape.
- u(int) - Picks a Site lat.unit_cell[u] out of the unit cell.
- opname (str) - valid operator name of an onsite operator in lat.unit_cell [u].
- category (str) - Descriptive name used as key for onsite_terms. Defaults to opname.
- plus_hc (bool) - If True, the hermitian conjugate of the terms is added automatically.

\section*{See also:}
add_coupling Add a terms acting on two sites.
add_onsite_term Add a single term without summing over vecx.
add_onsite_term (strength, \(i\), op, category=None, plus_hc=False)
Add an onsite term on a given MPS site.
Wrapper for self.onsite_terms[category].add_onsite_term(....).

\section*{Parameters}
- strength (float) - The strength of the term.
- i (int) - The MPS index of the site on which the operator acts. We require \(0<=i<\) L.
- op (str) - Name of the involved operator.
- category (str) - Descriptive name used as key for onsite_terms. Defaults to op.
- plus_hc (bool) - If True, the hermitian conjugate of the term is added automatically.
all_coupling_terms()
Sum of all coupling_terms.
```

all_onsite_terms()

```

Sum of all onsite_terms.
bond_energies \((p s i)\)
Calculate bond energies <psilH_bondlpsi>.
Parameters psi (MPS) - The MPS for which the bond energies should be calculated.
Returns E_bond - List of bond energies: for finite bc, E_Bond[i] is the energy of bond i, i+1. (i.e. we omit bond 0 between sites L-1 and 0); for infinite bc E_bond[i] is the energy of bond i-1, i.

Return type 1D ndarray
calc_H_MPO (tol_zero \(=1 e-15\) )
Calculate MPO representation of the Hamiltonian.
Uses onsite_terms and coupling_terms to build an MPOGraph (and then an MPO).
Parameters tol_zero (float) - Prefactors with abs (strength) < tol_zero are considered to be zero.
Returns H_MPO - MPO representation of the Hamiltonian.
Return type MPO
calc_H_MPO_from_bond (tol_zero=1e-15)
Calculate the MPO Hamiltonian from the bond Hamiltonian.
Parameters tol_zero (float) - Arrays with norm < tol_zero are considered to be zero.
Returns H_MPO - MPO representation of the Hamiltonian.
Return type MPO
calc_H_bond (tol_zero \(=1 e-15\) )
calculate \(H \_b o n d\) from coupling_terms and onsite_terms.
Parameters tol_zero (float) - prefactors with abs(strength) < tol_zero are considered to be zero.

Returns H_bond - Bond terms as required by the constructor of NearestNeighborModel. Legs are ['p0', 'p0*', 'p1', 'p1*']

Return type list of Array
:raises ValueError : if the Hamiltonian contains longer-range terms.:
calc_H_bond_from_MPO (tol_zero=le-15)
Calculate the bond Hamiltonian from the MPO Hamiltonian.
Parameters tol_zero (float) - Arrays with norm < tol_zero are considered to be zero.
Returns H_bond - Bond terms as required by the constructor of NearestNeighborModel. Legs are ['p0', 'p0*', 'p1', 'p1*']

Return type list of Array
:raises ValueError : if the Hamiltonian contains longer-range terms.:
calc_H_onsite (tol_zero=le-15)
Calculate H_onsite from self.onsite_terms.
Deprecated since version 0.4.0: This function will be removed in 1.0.0. Replace calls to this function by self.all_onsite_terms().remove_zeros(tol_zero).to_Arrays (self. lat.mps_sites()). You might also want to take explicit_plus_hc into account.

Parameters tol_zero (float) - prefactors with abs (strength) < tol_zero are considered to be zero.

\section*{Returns}

\section*{- H_onsite (list of npc.Array)}
- onsite terms of the Hamiltonian. If explicit_plus_hc is True, - Hermitian conjugates of the onsite terms will be included.
coupling_strength_add_ext_flux (strength, \(d x\), phase)
Add an external flux to the coupling strength.
When performing DMRG on a "cylinder" geometry, it might be useful to put an "external flux" through the cylinder. This means that a particle hopping around the cylinder should pick up a phase given by the external flux [[resta1998]]. This is also called "twisted boundary conditions" in literature. This function adds a complex phase to the strength array on some bonds, such that particles hopping in positive direction around the cylinder pick up \(\exp (+i\) phase \()\).

Warning: For the sign of phase it is important that you consistently use the creation operator as opl and the annihilation operator as op2 in add_coupling().

\section*{Parameters}
- strength (scalar / array) - The strength to be used in add_coupling(), when no external flux would be present.
- dx (iterable of int) - Translation vector (of the unit cell) between opl and op2 in add_coupling().
- phase (iterable of float) - The phase of the external flux for hopping in each direction of the lattice. E.g., if you want flux through the cylinder on which you have an infinite MPS, you should give phase \(=[0\), phi] souch that particles pick up a phase phi when hopping around the cylinder.
Returns strength - The strength array to be used as strength in add_coupling() with the given \(d x\).

\section*{Return type complex array}

\section*{Examples}

Let's say you have an infinite MPS on a cylinder, and want to add nearest-neighbor hopping of fermions with the FermionSite. The cylinder axis is the \(x\)-direction of the lattice, so to put a flux through the cylinder, you want particles hopping around the cylinder to pick up a phase phi given by the external flux.
```

>>> strength = 1. \# hopping strength without external flux
>>> phi = np.pi/4 \# determines the external flux strength
>>> for u1, u2, dx in self.lat.pairs['nearest_neighbors']:

```
(continued from previous page)
```

... strength_with_flux = self.coupling_strength_add_ext_flux(strength, dx,
\hookrightarrow [0, phi])
self.add_coupling(strength_with_flux, u1, 'Cd', u2, 'C', dx)
... self.add_coupling(np.conj(strength_with_flux), u2, 'Cd', u1, 'C', -dx)

```
enlarge_mps_unit_cell (factor=2)
Repeat the unit cell for infinite MPS boundary conditions; in place.
This has to be done after finishing initialization and can not be reverted.
Parameters factor (int) - The new number of sites in the MPS unit cell will be increased from \(N\) _sites to factor*N_sites_per_ring. Since MPS unit cells are repeated in the \(x\)-direction in our convetion, the lattice shape goes from ( \(L x, L y, \ldots, L u\) ) to (Lx*factor, Ly, ..., Lu).
classmethod from_MPOModel (mpo_model)
Initialize a NearestNeighborModel from a model class defining an MPO.
This is especially usefull in combination with MPOModel.group_sites().
Parameters mpo_model (MPOModel) - A model instance implementing the MPO. Does not need to be a NearestNeighborModel, but should only have nearest-neighbor couplings.

\section*{Examples}

The SpinChainNNN2 has next-nearest-neighbor couplings and thus only implements an MPO:
```

>>> from tenpy.models.spins_nnn import SpinChainNNN2
>>> nnn_chain = SpinChainNNN2({'L': 20})
>>> print(isinstance(nnn_chain, NearestNeighborModel))
False
>>> print("range before grouping:", nnn_chain.H_MPO.max_range)
range before grouping: 2

```

By grouping each two neighboring sites, we can bring it down to nearest neighbors.
```

>>> grouped_sites = nnn_chain.group_sites(2)
>>> print("range after grouping:", nnn_chain.H_MPO.max_range)
range after grouping: 1

```

Yet, TEBD will not yet work, as the model doesn't define \(H_{\_}\)bond. However, we can initialize a NearestNeighborModel from the MPO:
```

>>> nnn_chain_for_tebd = NearestNeighborModel.from_MPOModel(nnn_chain)
>>> isinstance(nnn_chain_for_tebd, NearestNeighborModel)
True

```
classmethod from_hdf5 (hdf5_loader, h5gr, subpath)
Load instance from a HDF5 file.
This method reconstructs a class instance from the data saved with save_hdf5 ().

\section*{Parameters}
- hdf5_loader (Hdf5Loader) - Instance of the loading engine.
- h5gr (Group) - HDF5 group which is represent the object to be constructed.
- subpath (str) - The name of \(h 5 g r\) with a ' /' in the end.

Returns obj - Newly generated class instance containing the required data.
Return type cls
group_sites ( \(n=2\), grouped_sites=None)
Modify self in place to group sites.
Group each \(n\) sites together using the GroupedSite. This might allow to do TEBD with a Trotter decomposition, or help the convergence of DMRG (in case of too long range interactions).

This has to be done after finishing initialization and can not be reverted.

\section*{Parameters}
- \(\mathbf{n}(i n t)\) - Number of sites to be grouped together.
- grouped_sites (None l list of GroupedSite) - The sites grouped together.

Returns grouped_sites - The sites grouped together.
Return type list of GroupedSite

\section*{init_H_from_terms()}

Initialize \(H_{-} M P O\) (and \(H \_\)bond) from the terms of the CouplingModel.
This function is called automatically during CouplingMPOModel.__init__.
If you use one of the add_* methods of the CouplingModel after initialization, you will need to call init_H_from_terms in the end by yourself, in order to update the \(H_{-} M P O\) (and possibly \(H_{-} b o n d\) ) representations. (You should get a warning about this... The way to avoid it is to initialize all the terms in init_terms by defining your own model, as outlined in Models.

\section*{init_lattice (model_params)}

Initialize a lattice for the given model parameters.
This function reads out the model parameter lattice. This can be a full Lattice instance, in which case it is just returned without further action. Alternatively, the lattice parameter can be a string giving the name of one of the predefined lattices, which then gets initialized. Depending on the dimensionality of the lattice, this requires different model parameters.

Parameters model_params (dict) - The model parameters given to __init__
Returns lat - An initialized lattice.
Return type Lattice

\section*{Options}
option CouplingMPOModel.lattice: str | Lattice
The name of a lattice pre-defined in TeNPy to be initialized. Alternatively, directly a subclass of Lattice instead of the name. Alternatively, a (possibly self-defined) Lattice instance. In the latter case, no further parameters are read out.
option CouplingMPOModel.bc_MPS: str Boundary conditions for the MPS.
option CouplingMPOModel.order: str
The order of sites within the lattice for non-trivial lattices, e.g, 'default', 'snake', see ordering (). Only used if lattice is a string.
option CouplingMPOModel.L: int
The length in x-direction; only read out for 1D lattices. For an infinite system the length of the unit cell.
option CouplingMPOModel.Lx: int
option CouplingMPOModel.Ly: int
The length in x - and y-direction; only read out for 2D lattices. For "infinite" \(b c_{-} M P S\), the system is infinite in x-direction and \(L x\) is the number of "rings" in the infinite MPS unit cell, while \(L y\) gives the circumference around the cylinder or width of th the rung for a ladder (depending on \(b c_{-} y\) ).
option CouplingMPOModel.bc_y: str
"cylinder" | "ladder"; only read out for 2D lattices. The boundary conditions in y-direction.
option CouplingMPOModel.bc_x: str
"open" | "periodic". Can be used to force "periodic" boundaries for the lattice, i.e., for the couplings in the Hamiltonian, even if the MPS is finite. Defaults to "open" for bc_MPS="finite " and "periodic" for bc_MPS="infinite. If you are not aware of the consequences, you should probably not use "periodic" boundary conditions. (The MPS is still "open", so this will introduce long-range couplings between the first and last sites of the MPS!)

\section*{init_sites (model_params)}

Define the local Hilbert space and operators; needs to be implemented in subclasses.
This function gets called by init_lattice () to get the Site for the lattice unit cell.

Note: Initializing the sites requires to define the conserved quantum numbers. All pre-defined sites accept conserve=None to disable using quantum numbers. Many models in TeNPy read out the conserve model parameter, which can be set to "best" to indicate the optimal parameters.

If you need to initialize more than one site, the function tenpy.networks.site. set_common_charges () should be helpful.

Parameters model_params (dict) - The model parameters given to \(\qquad\)
\(\qquad\)
Returns sites - The local sites of the lattice, defining the local basis states and operators.
Return type (tuple of) Site
init_terms (model_params)
Add the onsite and coupling terms to the model; subclasses should implement this.
save_hdf5 (hdf5_saver, h5gr, subpath)
Export self into a HDF5 file.
This method saves all the data it needs to reconstruct self with from_hdf5 ().
This implementation saves the content of ___dict__ with save_dict_content (), storing the format under the attribute 'format'.

\section*{Parameters}
- hdf5_saver (Hdf5Saver) - Instance of the saving engine.
- h5gr (: class `Group`) - HDF5 group which is supposed to represent self.
- subpath (str) - The name of \(h 5 g r\) with a '/' in the end.

\section*{test_sanity()}

Sanity check, raises ValueErrors, if something is wrong.

\section*{trivial_like_NNModel()}

Return a NearestNeighborModel with same lattice, but trivial ( \(\mathrm{H}=0\) ) bonds.

\section*{Module description}

Bosonic and fermionic Hubbard models.

\section*{19.9 hofstadter}
- full name: tenpy.models.hofstadter
- parent module: tenpy.models
- type: module

\section*{Classes}

\begin{tabular}{ll}
\hline HofstadterBosons(model_params) & Bosons on a square lattice with magnetic flux. \\
\hline HofstadterFermions(model_params) & Fermions on a square lattice with magnetic flux. \\
\hline
\end{tabular}

\section*{Functions}

\subsection*{19.9.1 gauge_hopping}
- full name: tenpy.models.hofstadter.gauge_hopping
- parent module: tenpy.models.hofstadter
- type: function
tenpy.models.hofstadter.gauge_hopping (model_params)
Compute hopping amplitudes for the Hofstadter models based on a gauge choice.
In the Hofstadter model, the magnetic field enters as an Aharonov-Bohm phase. This phase is dependent on a choice of gauge, which simultaneously defines a 'magnetic unit cell' (MUC).

The magnetic unit cell is the smallest set of lattice plaquettes that encloses an integer number of flux quanta. It can be user-defined by setting \(m x\) and my, but for common gauge choices is computed based on the flux density.

\section*{The gauge choices are:}
- 'landau_x': Landau gauge along the \(x\)-axis. The magnetic unit cell will have shape :math \((\) mathtt \(\{\mathrm{mx}\}, 1)\). For flux densities \(p / q, m x\) will default to q. Example: at a flux density \(1 / 3\), the magnetic unit cell will have shape \((3,1)\), so it encloses exactly 1 flux quantum.
- 'landau_y': Landau gauge along the y-axis. The magnetic unit cell will have shape :math ( 1 , mathtt \(\{m y\})^{\prime}\). For flux densities :math`p/q`, my will default to q. Example: at a flux density \(3 / 7\), the magnetic unit cell will have shape \((1,7)\), so it encloses axactly 3 flux quanta.
- 'symmetric': symmetric gauge. The magnetic unit cell will have shape ( \(\mathrm{mx}, \mathrm{my}\) ), with \(m x=m y\). For flux densities \(p / q, m x\) and \(m y\) will default to \(q\) Example: at a flux density 4/9, the magnetic unit cell will have shape \((9,9)\).

\section*{Parameters}
- gauge ('landau_x' / 'landau_y' / 'symmetric') - Choice of the gauge, see table above.
- mx (int / None) - Dimensions of the magnetic unit cell in terms of lattice sites. None defaults to the minimal choice compatible with gauge and phi_pq.
- my (int / None) - Dimensions of the magnetic unit cell in terms of lattice sites. None defaults to the minimal choice compatible with gauge and phi_pq.
- Jx (float) - 'Bare' hopping amplitudes (without phase). Without any flux we have hop_x = -Jx and hop_y = -Jy.
- Jy (float) - 'Bare' hopping amplitudes (without phase). Without any flux we have hop_x \(=-J x\) and hop_y \(=-J y\).
- phi_pq (tuple (int, int))-Magnetic flux as a fraction \(\mathrm{p} / \mathrm{q}\), defined as (p, q)

Returns hop_x, hop_y - Hopping amplitudes to be used as prefactors for \(c_{x, y}^{\dagger} c_{x+1, y}\) (hop_x) and \(c_{x, y}^{\dagger} c_{x, y+1}\left(h o p_{-} x\right)\), respectively, with the necessary phases for the gauge.
Return type float I array

\section*{Module description}

Cold atomic (Harper-)Hofstadter model on a strip or cylinder.
Todo: WARNING: These models are still under development and not yet tested for correctness. Use at your own risk! Replicate known results to confirm models work correctly. Long term: implement different lattices. Long term: implement variable hopping strengths Jx , Jy.

\subsection*{19.10 haldane}
- full name: tenpy.models.haldane
- parent module: tenpy.models
- type: module

\section*{Classes}

\begin{tabular}{ll}
\hline BosonicHaldaneModel(model_params) & Hardcore bosonic Haldane model. \\
\hline FermionicHaldaneModel(model_params) & Spinless fermionic Haldane model. \\
\hline
\end{tabular}

\section*{Module description}

Bosonic and fermionic Haldane models.

\subsection*{19.11 toric_code}
- full name: tenpy.models.toric_code
- parent module: tenpy.models
- type: module

\section*{Classes}

\begin{tabular}{ll}
\hline DualSquare(Lx, Ly, sites, \({ }^{* *}\) kwargs) & The dual lattice of the square lattice (again square). \\
\hline ToricCode(model_params) & Toric code model. \\
\hline
\end{tabular}

\subsection*{19.11.1 DualSquare}
- full name: tenpy.models.toric_code.DualSquare
- parent module: tenpy.models.toric_code
- type: class

\section*{Inheritance Diagram}


\section*{Methods}
\begin{tabular}{|c|c|}
\hline DualSquare.__init__(Lx, Ly, sites, **kwargs) & Initialize self. \\
\hline DualSquare.count_neighbors([u, key]) & Count e.g. \\
\hline DualSquare.coupling_shape(dx) & Calculate correct shape of the strengths for a coupling. \\
\hline DualSquare.distance(u1, u2, dx) & Get the distance for a given coupling between two sites in the lattice. \\
\hline DualSquare.enlarge_mps_unit_cell([factor]) & Repeat the unit cell for infinite MPS boundary conditions; in place. \\
\hline ```
DualSquare.find_coupling_pairs([max_dx,
...])
``` & Automatically find coupling pairs grouped by distances. \\
\hline DualSquare.from_hdf5(hdf5_loader, h5gr, subpath) & Load instance from a HDF5 file. \\
\hline DualSquare.lat2mps_idx(lat_idx) & Translate lattice indices (x_0, ..., \(x_{-}\{D-1\}\), u) to MPS index \(i\). \\
\hline DualSquare.mps2lat_idx(i) & Translate MPS index \(i\) to lattice indices ( \(x \_0, \ldots\), x_\{dim-1\}, u). \\
\hline DualSquare.mps2lat_values(A[, axes, u]) & Reshape/reorder \(A\) to replace an MPS index by lattice indices. \\
\hline DualSquare.mps2lat_values_masked(A[, axes,...]) & Reshape/reorder an array \(A\) to replace an MPS index by lattice indices. \\
\hline DualSquare.mps_idx_fix_u([u]) & return an index array of MPS indices for which the site within the unit cell is \(u\). \\
\hline DualSquare.mps_lat_idx_fix_u([u]) & Similar as mps_idx_fix_u(), but return also the corresponding lattice indices. \\
\hline DualSquare.mps_sites() & Return a list of sites for all MPS indices. \\
\hline
\end{tabular}

Table 60 - continued from previous page
\begin{tabular}{|c|c|}
\hline DualSquare.multi_coupling_shape(dx) & Calculate correct shape of the strengths for a multi_coupling. \\
\hline DualSquare.number_nearest_neighbors([u]) & Deprecated. \\
\hline DualSquare.number_next_nearest_neighb & B(dpuld cated. \\
\hline DualSquare.ordering(order) & Provide possible orderings of the \(N\) lattice sites. \\
\hline DualSquare.plot_basis(ax[, origin, shade]) & Plot arrows indicating the basis vectors of the lattice. \\
\hline DualSquare.plot_bc_identified(ax[,..]) & Mark two sites indified by periodic boundary conditions. \\
\hline DualSquare.plot_coupling(ax[, coupling, wrap]) & Plot lines connecting nearest neighbors of the lattice. \\
\hline DualSquare.plot_order(ax[, order, textkwargs]) & Plot a line connecting sites in the specified "order" and text labels enumerating them. \\
\hline DualSquare.plot_sites(ax[, markers]) & Plot the sites of the lattice with markers. \\
\hline DualSquare.position(lat_idx) & return 'space' position of one or multiple sites. \\
\hline ```
DualSquare.possible_couplings(u1,u2, dx[,
...])
``` & Find possible MPS indices for two-site couplings. \\
\hline DualSquare.possible_multi_couplings(ops[, ...]) & Generalization of possible_couplings () to couplings with more than 2 sites. \\
\hline DualSquare.save_hdf5(hdf5_saver, h5gr, subpath) & Export self into a HDF5 file. \\
\hline DualSquare.site(i) & return Site instance corresponding to an MPS index \(i\) \\
\hline DualSquare.test_sanity() & Sanity check. \\
\hline
\end{tabular}

\section*{Class Attributes and Properties}
\begin{tabular}{ll}
\hline DualSquare.Lu & \begin{tabular}{l} 
the (expected) number of sites in the unit cell, \\
len_(unit_cell).
\end{tabular} \\
\hline DualSquare.boundary_conditions & \begin{tabular}{l} 
Human-readable list of boundary conditions from bc \\
and bc_shift.
\end{tabular} \\
\hline DualSquare.dim & the dimension of the lattice \\
\hline DualSquare.nearest_neighbors & \\
\hline DualSquare.next_nearest_neighbors & \begin{tabular}{l} 
Defines an ordering of the lattice sites, thus mapping the \\
lattice to a 1D chain.
\end{tabular} \\
\hline DualSquare.next_next_nearest_neighbors &
\end{tabular}
class tenpy.models.toric_code.DualSquare (Lx, Ly, sites, **kwargs)
Bases: tenpy.models.lattice.Lattice
The dual lattice of the square lattice (again square).
The sites in this lattice correspond to the vertical and horizontal (nearest neighbor) bonds of a common Square lattice with the same dimensions \(L x, L y\).

\section*{Parameters}
- \(\mathbf{L x}(i n t)\) - Dimensions of the original lattice. This lattice has \(2 * L x * L y\) sites.
- Ly (int) - Dimensions of the original lattice. This lattice has \(2 * L x * L y\) sites.

- sites (Site) - The sites for the horizontal (first entry) and vertical (second entry) bonds.
- **kwargs - Additional keyword arguments given to the Lattice. basis, pos and pairs are set accordingly.

\section*{\(\operatorname{dim}=2\)}
the dimension of the lattice

\section*{ordering (order)}

Provide possible orderings of the \(N\) lattice sites.
The following orders are defined in this method compared to tenpy.models.lattice. Lattice. ordering():
\begin{tabular}{|l|l|l|}
\hline Order & equivalent priority & equivalent snake_winding \\
\hline 'default' & \((0,2,1)\) & (False, False, False) \\
\hline
\end{tabular}
property boundary_conditions
Human-readable list of boundary conditions from bc and bc_shift.
Returns boundary_conditions - List of "open" or "periodic", one entry for each direction of the lattice.

Return type list of str
count_neighbors ( \(u=0\), key='nearest_neighbors')
Count e.g. the number of nearest neighbors for a site in the bulk.

\section*{Parameters}
- u (int) - Specifies the site in the unit cell, for which we should count the number of neighbors (or whatever key specifies).
- key (str) - Key of pairs to select what to count.

Returns number - Number of nearest neighbors (or whatever key specified) for the \(u\)-th site in the unit cell, somewhere in the bulk of the lattice. Note that it might not be the correct value at the edges of a lattice with open boundary conditions.

\section*{Return type int}

\section*{coupling_shape ( \(d x\) )}

Calculate correct shape of the strengths for a coupling.
Parameters \(\mathbf{d x}\) (tuple of int) - Translation vector in the lattice for a coupling of two operators. Corresponds to \(d x\) argument of tenpy.models.model. CouplingModel. add_multi_coupling().

\section*{Returns}
- coupling_shape (tuple of int) - Len dim. The correct shape for an array specifying the coupling strength. lat_indices has only rows within this shape.
- shift_lat_indices (array) - Translation vector from origin to the lower left corner of box spanned by \(d x\).
distance ( \(u 1, u 2, d x\) )
Get the distance for a given coupling between two sites in the lattice.
The \(u 1, u 2, d x\) parameters are defined in analogy with add_coupling(), i.e., this function calculates the distance between a pair of operators added with add_coupling (using the basis and unit_cell_positions of the lattice).

Warning: This function ignores "wrapping" arround the cylinder in the case of periodic boundary conditions.

\section*{Parameters}
- u1 (int) - Indices within the unit cell; the \(u 1\) and \(u 2\) of add_coupling()
- u2 (int) - Indices within the unit cell; the \(u 1\) and \(u 2\) of add_coupling()
- \(\mathbf{d x}\) (array) - Length dim. The translation in terms of basis vectors for the coupling.

Returns distance - The distance between site at lattice indices [ \(\mathrm{x}, \mathrm{y}, \mathrm{u} 1]\) and \([\mathrm{x}+\) \(d x[0], y+d x[1], u 2]\), ignoring any boundary effects.

Return type float

\section*{enlarge_mps_unit_cell (factor=2)}

Repeat the unit cell for infinite MPS boundary conditions; in place.
Parameters factor (int) - The new number of sites in the MPS unit cell will be increased from \(N\) _sites to factor*N_sites_per_ring. Since MPS unit cells are repeated in the \(x\)-direction in our convetion, the lattice shape goes from ( \(L x, L y, \ldots, L u\) ) to (Lx*factor, Ly, ..., Lu).
find_coupling_pairs (max_dx=3, cutoff=None, eps=1e-10)
Automatically find coupling pairs grouped by distances.
Given the unit_cell_positions and basis, the coupling pairs of nearest_neighbors, next_nearest_neighbors etc at a given distance are basically fixed (although not uniquely, since we take out half of them to avoid double-counting couplings in both directions A_i B_j + B_i A_i). This function iterates through all possible couplings up to a given cutoff distance and then determines the possible pairs at fixed distances (up to round-off errors).

\section*{Parameters}
- max_dx (int) - Maximal index for each index of \(d x\) to iterate over. You need large enough values to include every possible coupling up to the desired distance, but choosing it too large might make this function run for a long time.
- cutoff (float) - Maximal distance (in the units in which basis and unit_cell_positions is given).
- eps (float) - Tolerance up to which to distances are considered the same.

Returns coupling_pairs - Keys are distances of nearest-neighbors, next-nearest-neighbors etc.
Values are [(u1, u2, dx), ...] as in pairs.
Return type dict
classmethod from_hdf5 (hdf5_loader, h5gr, subpath)
Load instance from a HDF5 file.
This method reconstructs a class instance from the data saved with save_hdf5 ().

\section*{Parameters}
- hdf5_loader (Hdf5Loader) - Instance of the loading engine.
- h5gr (Group) - HDF5 group which is represent the object to be constructed.
- subpath (str) - The name of \(h 5 g r\) with a ' /' in the end.

Returns obj - Newly generated class instance containing the required data.

Return type cls
lat2mps_idx (lat_idx)
Translate lattice indices ( \(\mathrm{x} \_0, \ldots, \mathrm{x} \_\{\mathrm{D}-1\}, \mathrm{u}\) ) to MPS index \(i\).
Parameters lat_idx (array_like [.., dim+1]) - The last dimension corresponds to lattice indices \(\left(x_{-} 0, \ldots, x_{-}\{D-1\}, u\right)\). All lattice indices should be positive and smaller than the corresponding entry in self.shape. Exception: for "infinite" \(b c_{-} M P S\), an \(x \_0\) outside indicates shifts accross the boundary.

Returns \(\mathbf{i}\) - MPS index/indices corresponding to lat_idx. Has the same shape as lat_idx without the last dimension.

Return type array_like
mps2lat_idx (i)
Translate MPS index \(i\) to lattice indices ( \(\left.\mathrm{x} \_0, \ldots, \mathrm{x}_{-}\{\operatorname{dim}-1\}, u\right)\).
Parameters i (int / array_like of int)-MPS index/indices.
Returns lat_idx - First dimensions like \(i\), last dimension has len dim \({ }^{`}+1\) and contains the lattice indices " \(\left(x \_0, \ldots, x_{\_}\{d i m-1\}, u\right)\) ` corresponding to \(i\). For \(i\) accross the MPS unit cell and "infinite" \(b c_{-} M P S\), we shift \(x_{-} 0\) accordingly.

\section*{Return type array}
mps2lat_values \((A\), axes \(=0, u=\) None)
Reshape/reorder \(A\) to replace an MPS index by lattice indices.

\section*{Parameters}
- A (ndarray) - Some values. Must have A.shape[axes] = self.N_sites if \(u\) is None, or A.shape[axes] = self.N_cells if \(u\) is an int.
- axes ( (iterable of) int) - chooses the axis which should be replaced.
- u (None I int) - Optionally choose a subset of MPS indices present in the axes of \(A\), namely the indices corresponding to self.unit_cell[u], as returned by mps_idx_fix_u(). The resulting array will not have the additional dimension(s) of \(u\).

Returns res_A - Reshaped and reordered verions of A. Such that MPS indices along the specified axes are replaced by lattice indices, i.e., if MPS index \(j\) maps to lattice site ( \(x 0, x 1, x 2\) ), then res_A[..., \(x 0, x 1, x 2, \ldots]=A[\ldots, j, \ldots]\).
Return type ndarray

\section*{Examples}

Say you measure expection values of an onsite term for an MPS, which gives you an 1D array \(A\), where \(A[i]\) is the expectation value of the site given by self.mps2lat_idx(i). Then this function gives you the expectation values ordered by the lattice:
```

>>> print(lat.shape, A.shape)
(10, 3, 2) (60,)
>>> A_res = lat.mps2lat_values(A)
>>> A_res.shape
(10, 3, 2)
>>> A_res[tuple(lat.mps2lat_idx(5))] == A[5]
True

```

If you have a correlation function C [i, j], it gets just slightly more complicated:
```

>>> print(lat.shape, C.shape)
(10, 3, 2) (60, 60)
>>> lat.mps2lat_values(C, axes=[0, 1]).shape
(10, 3, 2, 10, 3, 2)

```

If the unit cell consists of different physical sites, an onsite operator might be defined only on one of the sites in the unit cell. Then you can use mps_idx_fix_u() to get the indices of sites it is defined on, measure the operator on these sites, and use the argument \(u\) of this function.
```

>>> u = 0
>>> idx_subset = lat.mps_idx_fix_u(u)
>>> A_u = A[idx_subset]
>>> A_u__res = lat.mps2lat_values(A_u, u=u)
>>> A_u_res.shape
(10, 3)
>>> np.all(A_res[:, :, u] == A_u_res[:, :])
True

```
mps2lat_values_masked (A, axes=- \(1, m p s \_i n d s=N o n e\), include_u=None)
Reshape/reorder an array \(A\) to replace an MPS index by lattice indices.
This is a generalization of mps2lat_values () allowing for the case of an arbitrary set of MPS indices present in each axis of \(A\).

\section*{Parameters}
- A (ndarray) - Some values.
- axes ( (iterable of) int) - Chooses the axis of \(A\) which should be replaced. If multiple axes are given, you also need to give multiple index arrays as mps_inds.
- mps_inds ((list of) 1D ndarray) - Specifies for each axis in axes, for which MPS indices we have values in the corresponding axis of A. Defaults to [np. arange (A. shape[ax]) for ax in axes]. For indices accross the MPS unit cell and "infinite" \(b c_{-} M P S\), we shift \(x \_0\) accordingly.
- include_u ((list of) bool) - Specifies for each axis in axes, whether the \(u\) index of the lattice should be included into the output array res_A. Defaults to len (self. unit_cell) > 1.
Returns res_A - Reshaped and reordered copy of A. Such that MPS indices along the specified axes are replaced by lattice indices, i.e., if MPS index \(j\) maps to lattice site ( \(x 0, x 1, x 2\) ), then res_A[..., x0, x1, x2, ...] = A[..., mps_inds[j], ...].

\section*{Return type np.ma.MaskedArray}
mps_idx_fix_u(u=None)
return an index array of MPS indices for which the site within the unit cell is \(u\).
If you have multiple sites in your unit-cell, an onsite operator is in general not defined for all sites. This functions returns an index array of the mps indices which belong to sites given by self. unit_cell[u].

Parameters u(None / int) - Selects a site of the unit cell. None (default) means all sites.
Returns mps_idx - MPS indices for which self.site(i) is self.unit_cell[u]. Ordered ascending.

\section*{Return type array}
mps_lat_idx_fix_u (u=None)
Similar as mps_idx_fix_u(), but return also the corresponding lattice indices.
Parameters u(None / int) - Selects a site of the unit cell. None (default) means all sites.

\section*{Returns}
- mps_idx (array) - MPS indices \(i\) for which self.site(i) is self. unit_cell[u].
- lat_idx ( \(2 D\) array) - The row \(j\) contains the lattice index (without \(u\) ) corresponding to mps_idx[j].
mps_sites()
Return a list of sites for all MPS indices.
Equivalent to [self.site(i) for i in range(self.N_sites)].
This should be used for sites of 1D tensor networks (MPS, MPO,...).
multi_coupling_shape ( \(d x\) )
Calculate correct shape of the strengths for a multi_coupling.
Parameters \(d \mathbf{x}(2 \mathrm{D}\) array, shape (N_ops, dim)) - dx[i, :] is the translation vector in the lattice for the \(i\)-th operator. Corresponds to the \(d x\) of each operator given in the argument ops of tenpy.models.model.CouplingModel.add_multi_coupling().

\section*{Returns}
- coupling_shape (tuple of int) - Len dim. The correct shape for an array specifying the coupling strength. lat_indices has only rows within this shape.
- shift_lat_indices (array) - Translation vector from origin to the lower left corner of box spanned by \(d x\). (Unlike for coupling_shape () it can also contain entries \(>0\) )
number_nearest_neighbors ( \(u=0\) )
Deprecated.
Deprecated since version 0.5.0: Use count_neighbors () instead.
```

number_next_nearest_neighbors ( }u=0\mathrm{ )

```

Deprecated.
Deprecated since version 0.5.0: Use count_neighbors () instead.

\section*{property order}

Defines an ordering of the lattice sites, thus mapping the lattice to a 1 D chain.
Each row of the array contains the lattice indices for one site, the order of the rows thus specifies a path through the lattice, along which an MPS will wind through through the lattice.
You can visualize the order with plot_order ().
plot_basis (ax, origin \(=(0.0,0.0)\), shade \(=\) None, \(* * k w a r g s)\)
Plot arrows indicating the basis vectors of the lattice.

\section*{Parameters}
- ax (matplotlib. axes.Axes) - The axes on which we should plot.
- **kwargs - Keyword arguments for ax. arrow.
plot_bc_identified (ax, direction=- 1, origin=None, cylinder_axis=False, **kwargs)
Mark two sites indified by periodic boundary conditions.
Works only for lattice with a 2-dimensional basis.

\section*{Parameters}
- ax (matplotlib.axes.Axes) - The axes on which we should plot.
- direction (int) - The direction of the lattice along which we should mark the idenitified sites. If None, mark it along all directions with periodic boundary conditions.
- cylinder_axis (bool) - Whether to plot the cylinder axis as well.
- origin (None | np.ndarray) - The origin starting from where we mark the identified sites. Defaults to the first entry of unit_cell_positions.
- **kwargs - Keyword arguments for the used ax.plot.
plot_coupling (ax, coupling=None, wrap=False, **kwargs)
Plot lines connecting nearest neighbors of the lattice.

\section*{Parameters}
- \(\mathbf{a x}\) (matplotlib.axes. Axes) - The axes on which we should plot.
- coupling (list of (u1, u2, dx)) - By default (None), use self. pairs['nearest_neighbors']. Specifies the connections to be plotted; iteating over lattice indices ( \(i 0, i 1, \ldots\) ), we plot a connection from the site (i0, i1, ..., u1) to the site ( \(\mathrm{i} 0+\mathrm{dx}[0]\), \(i 1+\mathrm{dx}[1], \ldots, \mathrm{u}\) ), taking into account the boundary conditions.
- wrap (bool) - If True, plot couplings going around the boundary by directly connecting the sites it connects. This might be hard to see, as this puts lines from one end of the lattice to the other. If False, plot the couplings as dangling lines.
- **kwargs - Further keyword arguments given to ax.plot ().
plot_order (ax, order=None, textkwargs=\{'color': 'r'\}, **kwargs)
Plot a line connecting sites in the specified "order" and text labels enumerating them.

\section*{Parameters}
- ax (matplotlib.axes.Axes) - The axes on which we should plot.
- order (None | 2D array (self.N_sites, self.dim+1))- The order as returned by ordering (); by default (None) use order.
- textkwargs (None I dict) - If not None, we add text labels enumerating the sites in the plot. The dictionary can contain keyword arguments for ax.text ().
- **kwargs - Further keyword arguments given to ax.plot ().
plot_sites (ax, markers=['o', ^', 's', 'p', 'h', 'D'], **kwargs)
Plot the sites of the lattice with markers.

\section*{Parameters}
- ax (matplotlib.axes.Axes) - The axes on which we should plot.
- markers (list) - List of values for the keywork marker of ax.plot () to distinguish the different sites in the unit cell, a site \(u\) in the unit cell is plotted with a marker markers [u \% len(markers)].
- **kwargs - Further keyword arguments given to ax.plot ().
position(lat_idx)
return 'space' position of one or multiple sites.
Parameters lat_idx (ndarray, (... , dim+1))- Lattice indices.

Returns pos - The position of the lattice sites specified by lat_idx in real-space.
Return type ndarray, (..., dim)
possible_couplings ( \(u 1, u 2, d x\), strength=None) Find possible MPS indices for two-site couplings.
For periodic boundary conditions ( \(\mathrm{bc}[\mathrm{a}]==\mathrm{False}\) ) the index x _a is taken modulo Ls [a] and runs through range (Ls [a]). For open boundary conditions, \(x_{-}\)a is limited to \(0<=x_{\text {_ }}<\) Ls [a] and \(0<=x_{\text {_ }}+d x[a]<\) lat.Ls [a].

\section*{Parameters}
- u1 (int) - Indices within the unit cell; the \(u 1\) and \(u 2\) of add_coupling()
- u2 (int) - Indices within the unit cell; the \(u 1\) and \(u 2\) of add_coupling()
- \(\mathbf{d x}\) (array) - Length dim. The translation in terms of basis vectors for the coupling.
- strength (array_like / None) - If given, instead of returning lat_indices and coupling_shape directly return the correct strength_12.

\section*{Returns}
- mps1, mps2 (1D array) - For each possible two-site coupling the MPS indices for the \(u 1\) and \(u 2\).
- strength_vals (1D array) - (Only returend if strength is not None.) Such that for (i, \(j, ~ s) ~ i n ~ z i p\left(m p s 1, ~ m p s 2, ~ s t r e n g t h \_v a l s\right): ~ i t e r a t e s ~ o v e r ~ a l l ~ p o s s i b l e ~ c o u-~\) plings with \(s\) being the strength of that coupling. Couplings where strength_vals \(==0\). are filtered out.
- lat_indices ( \(2 D\) int array) - (Only returend if strength is None.) Rows of lat_indices correspond to entries of mpsl and mps2 and contain the lattice indices of the "lower left corner" of the box containing the coupling.
- coupling_shape (tuple of int) - (Only returend if strength is None.) Len dim. The correct shape for an array specifying the coupling strength. lat_indices has only rows within this shape.
```

possible_multi_couplings(ops, strength=None)

```

Generalization of possible_couplings () to couplings with more than 2 sites.
Parameters ops (list of (opname, \(\mathrm{dx}, \mathrm{u})\) ) - Same as the argument ops of add_multi_coupling().

\section*{Returns}
- mps_ijkl ( \(2 D\) int array) - Each row contains MPS indices \(i, j, k, l, \ldots\). for each of the operators positions. The positions are defined by \(d x\) ( \(\mathrm{j}, \mathrm{k}, 1, \ldots\) relative to \(i\) ) and boundary coundary conditions of self (how much the box for given \(d x\) can be shifted around without hitting a boundary - these are the different rows).
- strength_vals ( \(1 D\) array) - (Only returend if strength is not None.) Such that for (ijkl, s) in zip(mps_ijkl, strength_vals): iterates over all possible couplings with \(s\) being the strength of that coupling. Couplings where strength_vals \(==0\). are filtered out.
- lat_indices ( 2 D int array) - (Only returend if strength is None.) Rows of lat_indices correspond to rows of \(m p s \_i j k l\) and contain the lattice indices of the "lower left corner" of the box containing the coupling.
- coupling_shape (tuple of int) - (Only returend if strength is None.) Len dim. The correct shape for an array specifying the coupling strength. lat_indices has only rows within this shape.
```

save_hdf5 (hdf5_saver, h5gr, subpath)
Export self into a HDF5 file.
This method saves all the data it needs to reconstruct self with from_hdf5 ().
Specifically, it saves unit_cell, Ls, unit_cell_positions, basis,
boundary_conditions, pairs under their name, bc_MPS as "boundary_conditions_MPS",
and order as "order_for_MPS". Moreover, it saves dim and N_sites as HDF5 attributes.

```

\section*{Parameters}
- hdf5_saver (Hdf5Saver) - Instance of the saving engine.
- h5gr (: class `Group`) - HDF5 group which is supposed to represent self.
- subpath (str) - The name of \(h 5 g r\) with a '/' in the end.
site ( \(i\) )
return Site instance corresponding to an MPS index \(i\)
```

test_sanity()

```

Sanity check.
Raises ValueErrors, if something is wrong.

\section*{Module description}

Kitaev's exactly solvable toric code model.
As we put the model on a cylinder, the name "toric code" is a bit misleading, but it is the established name for this model...

\section*{NETWORKS}
- full name: tenpy.networks
- parent module: tenpy
- type: module

\section*{Module description}

Definitions of tensor networks like MPS and MPO.
Here, 'tensor network' refers just to the (parital) contraction of tensors. For example an MPS represents the contraction along the 'virtual' legs/bonds of its \(B\).

\section*{Submodules}
\begin{tabular}{ll}
\hline site & \begin{tabular}{l} 
Defines a class describing the local physical Hilbert \\
space.
\end{tabular} \\
\hline mps & \begin{tabular}{l} 
This module contains a base class for a Matrix Product \\
State (MPS).
\end{tabular} \\
\hline mpo & Matrix product operator (MPO). \\
\hline terms & \begin{tabular}{l} 
Classes to store a collection of operator names and sites \\
they act on, together with prefactors.
\end{tabular} \\
\hline purification_mps & \begin{tabular}{l} 
This module contains an MPS class representing an den- \\
sity matrix by purification.
\end{tabular} \\
\hline
\end{tabular}

\section*{20.1 site}
- full name: tenpy.networks.site
- parent module: tenpy. networks
- type: module

\section*{Classes}

\begin{tabular}{ll}
\hline BosonSite([Nmax, conserve, filling]) & Create a Site for up to Nmax bosons. \\
\hline FermionSite([conserve, filling]) & Create a Site for spin-less fermions. \\
\hline GroupedSite(sites[, labels, charges]) & Group two or more Site into a larger one. \\
\hline Site(leg[, state_labels]) & \begin{tabular}{l} 
Collects necessary information about a single local site \\
of a lattice.
\end{tabular} \\
\hline \begin{tabular}{l} 
SpinHalfFermionSite([cons_N, cons_Sz, \\
ing] \()\)
\end{tabular} & fill- \\
\hline Create a Site for spinful (spin-1/2) fermions. \\
\hline SpinSite([S, conserve] \()\) & Spin-1/2 site. \\
\hline
\end{tabular}

\subsection*{20.1.1 BosonSite}
- full name: tenpy.networks.site.BosonSite
- parent module: tenpy. networks.site
- type: class

\section*{Inheritance Diagram}


\section*{Methods}
\begin{tabular}{ll}
\hline BosonSite.__init__([Nmax, conserve, filling]) & Initialize self. \\
\hline BosonSite.add_op(name, op[, need_JW, hc]) & Add one on-site operators. \\
\hline \begin{tabular}{l} 
BosonSite.change_charge([new_leg_charge, \\
\(\ldots])\).
\end{tabular} & Change the charges of the site (in place). \\
\hline \begin{tabular}{l} 
BosonSite.from_hdf5(hdf5_loader, h5gr, sub- \\
path)
\end{tabular} & Load instance from a HDF5 file. \\
\hline BosonSite.get_hc_op_name(name) & Return the hermitian conjugate of a given operator. \\
\hline BosonSite.get_op(name) & Return operator of given name. \\
\hline BosonSite.multiply_op_names(names) & Multiply operator names together. \\
\hline BosonSite.multiply_operators(operators) & \begin{tabular}{l} 
Multiply local operators (possibly given by their names) \\
together.
\end{tabular} \\
\hline BosonSite.op_needs_JW(name) & \begin{tabular}{l} 
Whether an (composite) onsite operator is fermionic and \\
needs a Jordan-Wigner string.
\end{tabular} \\
\hline BosonSite.remove_op(name) & Remove an added operator. \\
\hline BosonSite.rename_op(old_name, new_name) & Rename an added operator. \\
\hline BosonSite.save_hdf5(hdf5_saver, h5gr, sub- & Export self into a HDF5 file. \\
path) & Return index of a basis state from its label. \\
\hline BosonSite.state_index(label) & Same as state_index (), but for multiple labels. \\
\hline BosonSite.state_indices(labels) & Sanity check, raises ValueErrors, if something is wrong. \\
\hline BosonSite.test_sanity() & Check whether 'name' labels a valid onsite-operator. \\
\hline BosonSite.valid_opname(name) &
\end{tabular}

\section*{Class Attributes and Properties}
\begin{tabular}{ll}
\hline BosonSite.dim & Dimension of the local Hilbert space. \\
\hline BosonSite.onsite_ops & Dictionary of on-site operators for iteration. \\
\hline
\end{tabular}
class tenpy.networks.site. BosonSite ( Nmax \(=1\), conserve \(={ }^{\prime} N^{\prime}\), filling=0.0)
Bases: tenpy.networks.site.Site
Create a Site for up to Nmax bosons.
Local states are vac, 1, 2, ... , Nc. (Exception: for parity conservation, we sort as vac, 2, 4, ..., 1, 3, 5, ....)
\begin{tabular}{|l|l|}
\hline operator & description \\
\hline Id, JW & Identity \(\nVdash\) \\
\hline B & Annihilation operator \(b\) \\
\hline Bd & Creation operator \(b^{\dagger}\) \\
\hline N & Number operator \(n=b^{\dagger} b\) \\
\hline NN & \(n^{2}\) \\
\hline dN & \(\delta n:=n-\) filling \\
\hline dNdN & \((\delta n)^{2}\) \\
\hline P & Parity \(I d-2(n \bmod 2)\). \\
\hline
\end{tabular}
\begin{tabular}{|l|l|l|}
\hline conserve & qmod & excluded onsite operators \\
\hline 'N' & {\([1]\)} & - \\
\hline 'parity' & {\([2]\)} & - \\
\hline None & {[]} & - \\
\hline
\end{tabular}

\section*{Parameters}
- Nmax (int) - Cutoff defining the maximum number of bosons per site. The default Nmax \(=1\) describes hard-core bosons.
- conserve (str) - Defines what is conserved, see table above.
- filling (float) - Average filling. Used to define dN .

\section*{conserve}

Defines what is conserved, see table above.
Type str
filling
Average filling. Used to define \(d N\).
Type float
add_op (name, op, need_JW=False, hc=None)
Add one on-site operators.

\section*{Parameters}
- name (str) - A valid python variable name, used to label the operator. The name under which \(o p\) is added as attribute to self.
- op (np.ndarray I Array) - A matrix acting on the local hilbert space representing the local operator. Dense numpy arrays are automatically converted to Array. LegCharges have to be [leg, leg.conj()]. We set labels 'p', 'p*'.
- need_JW (bool) - Whether the operator needs a Jordan-Wigner string. If True, add name to need_JW_string.
- hc (None / False / str) - The name for the hermitian conjugate operator, to be used for hc_ops. By default (None), try to auto-determine it. If False, disable adding antries to hc_ops.
change_charge (new_leg_charge=None, permute=None)
Change the charges of the site (in place).

\section*{Parameters}
- new_leg_charge (LegCharge I None) - The new charges to be used. If None, use trivial charges.
- permute (ndarray / None) - The permuation applied to the physical leg, which gets used to adjust state_labels and perm. If you sorted the previous leg with perm_qind, new_leg_charge \(=\) leg. sort(), use old_leg. perm_flat_from_perm_qind(perm_qind). Ignored if None.

\section*{property dim}

Dimension of the local Hilbert space.
classmethod from_hdf5 (hdf5_loader, h5gr, subpath)
Load instance from a HDF5 file.
This method reconstructs a class instance from the data saved with save_hdf5().

\section*{Parameters}
- hdf5_loader (Hdf5Loader) - Instance of the loading engine.
- h5gr (Group) - HDF5 group which is represent the object to be constructed.
- subpath (str) - The name of \(h 5 g r\) with a '/' in the end.

Returns obj - Newly generated class instance containing the required data.
Return type cls
get_hc_op_name (name)
Return the hermitian conjugate of a given operator.
Parameters name (str) - The name of the operator to be returned. Multiple operators separated by whitespace are interpreted as an operator product, exactly as get_op () does.
Returns hc_op_name - Operator name for the hermi such that get_op () of
Return type str
get_op (name)
Return operator of given name.
Parameters name (str) - The name of the operator to be returned. In case of multiple operator names separated by whitespace, we multiply them together to a single on-site operator (with the one on the right acting first).
Returns op - The operator given by name, with labels 'p', 'p*'. If name already was an npc Array, it's directly returned.

Return type np_conserved
multiply_op_names (names)
Multiply operator names together.
Join the operator names in names such that get_op returns the product of the corresponding operators.
Parameters names (list of str) - List of valid operator labels.
Returns combined_opname - A valid operator name Operatorname representing the product of operators in names.

Return type str
multiply_operators (operators)
Multiply local operators (possibly given by their names) together.
Parameters operators (list of \(\{\operatorname{str|Array}\}\) ) - List of valid operator names (to be translated with get_op ()) or directly on-site operators in the form of npc arrays with 'p', 'p*' label. The operators are multiplied left-to-right.

Returns combined_operator - The product of the given operators in a left-to-right multiplication following the usual mathematical convention. For example, if operators=['Sz', 'Sp', 'Sx'], the final operator is equivalent to site.get_op('Sz Sp Sx'), with the ' \(S x\) ' operator acting first on any physical state.

\section*{Return type Array}
property onsite_ops
Dictionary of on-site operators for iteration.
Single operators are accessible as attributes.
```

op_needs_JW(name)

```

Whether an (composite) onsite operator is fermionic and needs a Jordan-Wigner string.
Parameters name (str) - The name of the operator, as in get_op ().
Returns needs_JW - Whether the operator needs a Jordan-Wigner string, judging from need_JW_string.

\section*{Return type bool}
```

remove_op (name)

```

Remove an added operator.
Parameters name (str) - The name of the operator to be removed.
rename_op (old_name, new_name)
Rename an added operator.

\section*{Parameters}
- old_name (str) - The old name of the operator.
- new_name (str) - The new name of the operator.
save_hdf5 (hdf5_saver, h5gr, subpath)
Export self into a HDF5 file.
This method saves all the data it needs to reconstruct self with from_hdf5 ().
This implementation saves the content of __dict__ with save_dict_content (), storing the format under the attribute 'format'.

\section*{Parameters}
- hdf5_saver (Hdf5Saver) - Instance of the saving engine.
- h5gr (: class `Group`) - HDF5 group which is supposed to represent self.
- subpath (str) - The name of \(h 5 g r\) with a '/' in the end.
state_index (label)
Return index of a basis state from its label.
Parameters label (int | string)-eather the index directly or a label (string) set before.
Returns state_index - the index of the basis state associated with the label.
Return type int
state_indices (labels)
Same as state_index(), but for multiple labels.
test_sanity()
Sanity check, raises ValueErrors, if something is wrong.
valid_opname (name)
Check whether 'name' labels a valid onsite-operator.
Parameters name (str) - Label for the operator. Can be multiple operator(labels) separated by whitespace, indicating that they should be multiplied together.

Returns valid - True if name is a valid argument to get_op ().
Return type bool

\subsection*{20.1.2 FermionSite}
- full name: tenpy.networks.site.FermionSite
- parent module: tenpy.networks.site
- type: class

Inheritance Diagram


\section*{Methods}
\begin{tabular}{ll}
\hline FermionSite.__init__([conserve, filling]) & Initialize self. \\
\hline FermionSite.add_op(name, op[, need_JW, hc]) & Add one on-site operators. \\
\hline FermionSite.change_charge([new_leg_charge, & Change the charges of the site (in place). \\
\(\ldots\).\(] )\) & \\
\hline \begin{tabular}{l} 
FermionSite.from_hdf5(hdf5_loader, h5gr, sub- \\
path)
\end{tabular} & Load instance from a HDF5 file. \\
\hline FermionSite.get_hc_op_name(name) & Return the hermitian conjugate of a given operator. \\
\hline FermionSite.get_op(name) & Return operator of given name. \\
\hline FermionSite.multiply_op_names(names) & Multiply operator names together. \\
\hline FermionSite.multiply_operators(operators) & \begin{tabular}{l} 
Multiply local operators (possibly given by their names) \\
together.
\end{tabular} \\
\hline FermionSite.op_needs_Jw(name) & \begin{tabular}{l} 
Whether an (composite) onsite operator is fermionic and \\
needs a Jordan-Wigner string.
\end{tabular} \\
\hline FermionSite.remove_op(name) & Remove an added operator. \\
\hline FermionSite.rename_op(old_name, new_name) & Rename an added operator. \\
\hline \begin{tabular}{l} 
FermionSite.save_hdf5(hdf5_saver, h5gr, sub- \\
path)
\end{tabular} & Export self into a HDF5 file. \\
\hline FermionSite.state_index(label) & Return index of a basis state from its label. \\
\hline FermionSite.state_indices(labels) & Same as state_index (), but for multiple labels. \\
\hline FermionSite.test_sanity() & Sanity check, raises ValueErrors, if something is wrong. \\
\hline FermionSite.valid_opname(name) & Check whether 'name' labels a valid onsite-operator. \\
\hline
\end{tabular}

\section*{Class Attributes and Properties}
\begin{tabular}{ll}
\hline FermionSite.dim & Dimension of the local Hilbert space. \\
\hline FermionSite.onsite_ops & Dictionary of on-site operators for iteration. \\
\hline
\end{tabular}
class tenpy.networks.site.FermionSite (conserve \(=\) ' \(N\) ', filling=0.5)
Bases: tenpy.networks.site.Site
Create a Site for spin-less fermions.
Local states are empty and full.

Warning: Using the Jordan-Wigner string (JW) is crucial to get correct results, otherwise you just describe hardcore bosons! Further details in Fermions and the Jordan-Wigner transformation.
\begin{tabular}{|l|l|}
\hline operator & description \\
\hline Id & Identity \(\nVdash\) \\
\hline JW & Sign for the Jordan-Wigner string. \\
\hline C & Annihilation operator \(c\) (up to 'JW'-string left of it) \\
\hline Cd & Creation operator \(c^{\dagger}\) (up to 'JW'-string left of it) \\
\hline N & Number operator \(n=c^{\dagger} c\) \\
\hline dN & \(\delta n:=n-\) filling \\
\hline dNdN & \((\delta n)^{2}\) \\
\hline
\end{tabular}
\begin{tabular}{|l|l|l|}
\hline conserve & qmod & exluded onsite operators \\
\hline 'N' & {\([1]\)} & - \\
\hline 'parity' & {\([2]\)} & - \\
\hline None & {[]} & - \\
\hline
\end{tabular}

\section*{Parameters}
- conserve (str) - Defines what is conserved, see table above.
- filling (float) - Average filling. Used to define \(d N\).

\section*{conserve}

Defines what is conserved, see table above.
Type str
filling
Average filling. Used to define \(d N\).
Type float
add_op (name, op, need_JW=False, hc=None)
Add one on-site operators.

\section*{Parameters}
- name (str) - A valid python variable name, used to label the operator. The name under which \(o p\) is added as attribute to self.
- op (np.ndarray I Array) - A matrix acting on the local hilbert space representing the local operator. Dense numpy arrays are automatically converted to Array. LegCharges have to be [leg, leg.conj()]. We set labels 'p', 'p*'.
- need_JW (bool) - Whether the operator needs a Jordan-Wigner string. If True, add name to need_JW_string.
- hc (None / False / str) - The name for the hermitian conjugate operator, to be used for hc_ops. By default (None), try to auto-determine it. If False, disable adding antries to hc_ops.
```

change_charge (new_leg_charge=None,permute=None)

```

Change the charges of the site (in place).

\section*{Parameters}
- new_leg_charge (LegCharge I None) - The new charges to be used. If None, use trivial charges.
- permute (ndarray / None) - The permuation applied to the physical leg, which gets used to adjust state_labels and perm. If you sorted the previous leg with perm_qind, new_leg_charge \(=\) leg. sort(), use old_leg. perm_flat_from_perm_qind (perm_qind). Ignored if None.
```

property dim

```

Dimension of the local Hilbert space.
classmethod from_hdf5 (hdf5_loader, h5gr, subpath)
Load instance from a HDF5 file.
This method reconstructs a class instance from the data saved with save_hdf5().

\section*{Parameters}
- hdf5_loader (Hdf5Loader) - Instance of the loading engine.
- h5gr (Group) - HDF5 group which is represent the object to be constructed.
- subpath (str) - The name of \(h 5 g r\) with a ' /' in the end.

Returns obj - Newly generated class instance containing the required data.
Return type cls
get_hc_op_name (name)
Return the hermitian conjugate of a given operator.
Parameters name (str) - The name of the operator to be returned. Multiple operators separated by whitespace are interpreted as an operator product, exactly as get_op () does.
Returns hc_op_name - Operator name for the hermi such that get_op () of
Return type str
get_op (name)
Return operator of given name.
Parameters name (str) - The name of the operator to be returned. In case of multiple operator names separated by whitespace, we multiply them together to a single on-site operator (with the one on the right acting first).

Returns op - The operator given by name, with labels ' \(p\) ', ' \(p *\) '. If name already was an npc Array, it's directly returned.
Return type np_conserved
multiply_op_names (names)
Multiply operator names together.
Join the operator names in names such that get_op returns the product of the corresponding operators.
Parameters names (list of str) - List of valid operator labels.
Returns combined_opname - A valid operator name Operatorname representing the product of operators in names.

Return type str
multiply_operators (operators)
Multiply local operators (possibly given by their names) together.
Parameters operators (list of \(\{\operatorname{str} \mid\) Array \(\}\) ) - List of valid operator names (to be translated with get_op ()) or directly on-site operators in the form of npc arrays with 'p', 'p*' label. The operators are multiplied left-to-right.
Returns combined_operator - The product of the given operators in a left-to-right multiplication following the usual mathematical convention. For example, if operators=['Sz', 'Sp', 'Sx'], the final operator is equivalent to site.get_op('Sz Sp Sx'), with the ' \(S x\) ' operator acting first on any physical state.

\section*{Return type Array}
property onsite_ops
Dictionary of on-site operators for iteration.
Single operators are accessible as attributes.
op_needs_JW (name)
Whether an (composite) onsite operator is fermionic and needs a Jordan-Wigner string.

Parameters name (str) - The name of the operator, as in get_op ().
Returns needs_JW - Whether the operator needs a Jordan-Wigner string, judging from need_JW_string.
Return type bool
remove_op (name)
Remove an added operator.
Parameters name (str) - The name of the operator to be removed.
rename_op (old_name, new_name)
Rename an added operator.

\section*{Parameters}
- old_name (str) - The old name of the operator.
- new_name (str) - The new name of the operator.
save_hdf5 (hdf5_saver, h5gr, subpath)
Export self into a HDF5 file.
This method saves all the data it needs to reconstruct self with from_hdf5 ().
This implementation saves the content of __ dict__ with save_dict_content (), storing the format under the attribute 'format'.

\section*{Parameters}
- hdf5_saver (Hdf5Saver) - Instance of the saving engine.
- h5gr (: class `Group`) - HDF5 group which is supposed to represent self.
- subpath (str) - The name of \(h 5 g r\) with a '/' in the end.
state_index (label)
Return index of a basis state from its label.
Parameters label (int / string) - eather the index directly or a label (string) set before.
Returns state_index - the index of the basis state associated with the label.
Return type int
state_indices (labels)
Same as state_index(), but for multiple labels.
test_sanity()
Sanity check, raises ValueErrors, if something is wrong.
valid_opname (name)
Check whether 'name' labels a valid onsite-operator.
Parameters name (str) - Label for the operator. Can be multiple operator(labels) separated by whitespace, indicating that they should be multiplied together.
Returns valid - True if name is a valid argument to get_op ().
Return type bool

\subsection*{20.1.3 GroupedSite}
- full name: tenpy.networks.site.GroupedSite
- parent module: tenpy.networks.site
- type: class

\section*{Inheritance Diagram}


\section*{Methods}
\begin{tabular}{ll}
\hline GroupedSite.__init__(sites[, labels, charges]) & Initialize self. \\
\hline GroupedSite.add_op(name, op[, need_JW, hc]) & Add one on-site operators. \\
\hline \begin{tabular}{l} 
GroupedSite.change_charge([new_leg_charge,
\end{tabular} & Change the charges of the site (in place). \\
\(\ldots])\). & Load instance from a HDF5 file. \\
\hline \begin{tabular}{l} 
GroupedSite.from_hdf5(hdf5_loader, h5gr, sub- \\
path)
\end{tabular} & Return the hermitian conjugate of a given operator. \\
\hline GroupedSite.get_hc_op_name(name) & Return operator of given name. \\
\hline GroupedSite.get_op(name) & \begin{tabular}{l} 
Return the Kronecker product \(o p 0\) 0 \(\otimes\) op 1 of local oper- \\
ators.
\end{tabular} \\
\hline GroupedSite.kroneckerproduct(ops) & Multiply operator names together. \\
\hline GroupedSite.multiply_op_names(names) & \begin{tabular}{l} 
Multiply local operators (possibly given by their names) \\
together.
\end{tabular} \\
\hline GroupedSite.multiply_operators(operators) & \begin{tabular}{l} 
Whether an (composite) onsite operator is fermionic and \\
needs a Jordan-Wigner string.
\end{tabular} \\
\hline GroupedSite.op_needs_JW(name) & Remove an added operator. \\
\hline GroupedSite.remove_op(name) & Rename an added operator. \\
\hline GroupedSite.rename_op(old_name, new_name) & Export self into a HDF5 file. \\
\hline GroupedSite.save_hdf5(hdf5_saver, h5gr, sub- & Return index of a basis state from its label. \\
\hline GroupedSite.state_index(label) & continues on next page
\end{tabular}

Table 7 - continued from previous page
\begin{tabular}{ll}
\hline GroupedSite.state_indices(labels) & Same as state_index(), but for multiple labels. \\
\hline GroupedSite.test_sanity() & Sanity check, raises ValueErrors, if something is wrong. \\
\hline GroupedSite.valid_opname(name) & Check whether 'name' labels a valid onsite-operator. \\
\hline
\end{tabular}

\section*{Class Attributes and Properties}
\begin{tabular}{ll}
\hline GroupedSite.dim & Dimension of the local Hilbert space. \\
\hline GroupedSite.onsite_ops & Dictionary of on-site operators for iteration. \\
\hline class tenpy.networks.site. GroupedSite (sites, labels=None, charges='same') \\
Bases: tenpy.networks.site.Site
\end{tabular}

Group two or more Site into a larger one.
A typical use-case is that you want a NearestNeighborModel for TEBD although you have next-nearest neighbor interactions: you just double your local Hilbertspace to consist of two original sites. Note that this is a 'hack' at the cost of other things (e.g., measurements of 'local' operators) getting more complicated/computationally expensive.

If the individual sites indicate fermionic operators (with entries in need_JW_string), we construct the new onsite oerators of sitel to include the JW string of siteO, i.e., we use the Kronecker product of [ JW, op] instead of [Id, op] if necessary (but always [op, Id]). In that way the onsite operators of this DoubleSite automatically fulfill the expected commutation relations. See also Fermions and the Jordan-Wigner transformation.

\section*{Parameters}
- sites (list of Site) - The individual sites being grouped together. Copied before use if charges!='same'.
- labels - Include the Kronecker product of the each onsite operator op on sites [i] and identities on other sites with the name opname+labels [i]. Similarly, set state labels for ' '.join(state[i]+'_'+labels[i]). Defaults to [str(i) for i in range (n_sites)], which for example grouping two SpinSites gives operators name like "Sz0" and sites labels like 'up_0 down_1'.
- charges ('same' | 'drop' | 'independent') - How to handle charges, defaults to 'same'. 'same' means that all sites have the same ChargeInfo, and the total charge is the sum of the charges on the individual sites. 'independent' means that the sites have possibly different ChargeInfo, and the charges are conserved separately, i.e., we have n_sites conserved charges. For 'drop', we drop any charges, such that the remaining legcharges are trivial. For more complex situations, you can call multi_sites_combine_charges() beforehand.
n_sites
The number of sites grouped together, i.e. len(sites).
Type int
sites
The sites grouped together into self.
Type list of Site

\section*{labels}

The labels using which the single-site operators are added during construction.
Type list of str

\section*{kroneckerproduct (ops)}

Return the Kronecker product op0 \(\otimes o p 1\) of local operators.
Parameters ops (list of Array) - One operator (or operator name) on each of the ungrouped sites. Each operator should have labels ['p', 'p*'].

Returns prod - Kronecker product ops \([0] \otimes o p s[1] \otimes \cdots\), with labels ['p', 'p*'].
Return type Array
add_op (name, op, need_JW=False, \(h c=\) None)
Add one on-site operators.

\section*{Parameters}
- name (str) - A valid python variable name, used to label the operator. The name under which \(o p\) is added as attribute to self.
- op (np.ndarray I Array) - A matrix acting on the local hilbert space representing the local operator. Dense numpy arrays are automatically converted to Array. LegCharges have to be [leg, leg.conj()]. We set labels 'p', 'p*'.
- need_JW (bool) - Whether the operator needs a Jordan-Wigner string. If True, add name to need_JW_string.
- hc (None / False / str) - The name for the hermitian conjugate operator, to be used for hc_ops. By default (None), try to auto-determine it. If False, disable adding antries to hc_ops.
change_charge (new_leg_charge=None, permute=None)
Change the charges of the site (in place).

\section*{Parameters}
- new_leg_charge (LegCharge I None) - The new charges to be used. If None, use trivial charges.
- permute (ndarray / None) - The permuation applied to the physical leg, which gets used to adjust state_labels and perm. If you sorted the previous leg with perm_qind, new_leg_charge = leg.sort(), use old_leg. perm_flat_from_perm_qind(perm_qind). Ignored if None.
property dim
Dimension of the local Hilbert space.

\section*{classmethod from_hdf5 (hdf5_loader, h5gr, subpath)}

Load instance from a HDF5 file.
This method reconstructs a class instance from the data saved with save_hdf5 ().

\section*{Parameters}
- hdf5_loader (Hdf5Loader) - Instance of the loading engine.
- h5gr (Group) - HDF5 group which is represent the object to be constructed.
- subpath (str) - The name of \(h 5 g r\) with a '/' in the end.

Returns obj - Newly generated class instance containing the required data.
Return type cls
get_hc_op_name (name)
Return the hermitian conjugate of a given operator.

Parameters name (str) - The name of the operator to be returned. Multiple operators separated by whitespace are interpreted as an operator product, exactly as get_op () does.
Returns hc_op_name - Operator name for the hermi such that get_op () of
Return type str
get_op (name)
Return operator of given name.
Parameters name (str) - The name of the operator to be returned. In case of multiple operator names separated by whitespace, we multiply them together to a single on-site operator (with the one on the right acting first).

Returns op - The operator given by name, with labels ' \(\mathrm{p}^{\prime}\), ' \(\mathrm{p} *\) '. If name already was an npc Array, it's directly returned.
Return type np_conserved
multiply_op_names (names)
Multiply operator names together.
Join the operator names in names such that get_op returns the product of the corresponding operators.
Parameters names (list of str)- List of valid operator labels.
Returns combined_opname - A valid operator name Operatorname representing the product of operators in names.

\section*{Return type str}
multiply_operators (operators)
Multiply local operators (possibly given by their names) together.
Parameters operators (list of \(\{\operatorname{str} \mid \operatorname{Array}\})\) - List of valid operator names (to be translated with get_op ()) or directly on-site operators in the form of npc arrays with 'p', 'p*' label. The operators are multiplied left-to-right.

Returns combined_operator - The product of the given operators in a left-to-right multiplication following the usual mathematical convention. For example, if operators=['Sz', 'Sp', 'Sx'], the final operator is equivalent to site.get_op('Sz Sp Sx'), with the ' \(S x\) ' operator acting first on any physical state.

\section*{Return type Array}
property onsite_ops
Dictionary of on-site operators for iteration.
Single operators are accessible as attributes.
op_needs_JW (name)
Whether an (composite) onsite operator is fermionic and needs a Jordan-Wigner string.
Parameters name (str) - The name of the operator, as in get_op ().
Returns needs_JW - Whether the operator needs a Jordan-Wigner string, judging from need_JW_string.
Return type bool
remove_op (name)
Remove an added operator.
Parameters name (str) - The name of the operator to be removed.
rename_op (old_name, new_name)
Rename an added operator.

\section*{Parameters}
- old_name (str) - The old name of the operator.
- new_name (str) - The new name of the operator.
save_hdf5 (hdf5_saver, h5gr, subpath)
Export self into a HDF5 file.
This method saves all the data it needs to reconstruct self with \(f r o m \_h d f 5\) ().
This implementation saves the content of \(\qquad\) dict_
\(\qquad\) with save_dict_content (), storing the format under the attribute 'format'.

\section*{Parameters}
- hdf5_saver (Hdf5Saver) - Instance of the saving engine.
- h5gr (: class `Group \()\) - HDF5 group which is supposed to represent self.
- subpath (str) - The name of \(h 5 g r\) with a ' /' in the end.
state_index (label)
Return index of a basis state from its label.
Parameters label (int / string)-eather the index directly or a label (string) set before.
Returns state_index - the index of the basis state associated with the label.
Return type int
state_indices (labels)
Same as state_index (), but for multiple labels.
test_sanity()
Sanity check, raises ValueErrors, if something is wrong.
valid_opname (name)
Check whether 'name' labels a valid onsite-operator.
Parameters name (str) - Label for the operator. Can be multiple operator(labels) separated by whitespace, indicating that they should be multiplied together.
Returns valid - True if name is a valid argument to get_op ().
Return type bool

\subsection*{20.1.4 Site}
- full name: tenpy.networks.site.Site
- parent module: tenpy.networks.site
- type: class

\section*{Inheritance Diagram}


\section*{Methods}
\begin{tabular}{ll}
\hline Site.__init__(leg[, state_labels]) & Initialize self. \\
\hline Site.add_op(name, op[, need_JW, hc]) & Add one on-site operators. \\
\hline \begin{tabular}{l} 
Site. change_charge([new_leg_charge, \\
mute])
\end{tabular} & per--
\end{tabular} Change the charges of the site (in place). \(\quad\)\begin{tabular}{ll}
\hline Site.from_hdf5(hdf5_loader, h5gr, subpath) & Load instance from a HDF5 file. \\
\hline Site.get_hc_op_name(name) & Return the hermitian conjugate of a given operator. \\
\hline Site.get_op(name) & Return operator of given name. \\
\hline Site.multiply_op_names(names) & Multiply operator names together. \\
\hline Site.multiply_operators(operators) & \begin{tabular}{l} 
Multiply local operators (possibly given by their names) \\
together.
\end{tabular} \\
\hline Site.op_needs_JW(name) & \begin{tabular}{l} 
Whether an (composite) onsite operator is fermionic and \\
needs a Jordan-Wigner string.
\end{tabular} \\
\hline Site.remove_op(name) & Remove an added operator. \\
\hline Site.rename_op(old_name, new_name) & Rename an added operator. \\
\hline Site.save_hdf5(hdf5_saver, h5gr, subpath) & Export self into a HDF5 file. \\
\hline Site.state_index(label) & Return index of a basis state from its label. \\
\hline Site.state_indices(labels) & Same as state_index (), but for multiple labels. \\
\hline Site.test_sanity() & Sanity check, raises ValueErrors, if something is wrong. \\
\hline Site.valid_opname(name) & Check whether 'name' labels a valid onsite-operator. \\
\hline
\end{tabular}

\section*{Class Attributes and Properties}
\begin{tabular}{ll}
\hline Site.dim & Dimension of the local Hilbert space. \\
\hline Site.onsite_ops & Dictionary of on-site operators for iteration. \\
\hline
\end{tabular}
class tenpy.networks.site.Site (leg, state_labels=None, **site_ops)
Bases: tenpy.tools.hdf5_io.Hdf5Exportable
Collects necessary information about a single local site of a lattice.
This class defines what the local basis states are: it provides the leg defining the charges of the physical leg for this site. Moreover, it stores (local) on-site operators, which are directly available as attribute, e.g., self.Sz
is the Sz operator for the SpinSite. Alternatively, operators can be obained with get_op (). The operator names Id and JW are reserved for the identy and Jordan-Wigner strings.

Warning: The order of the local basis can change depending on the charge conservation! This is a necessary feature since we need to sort the basis by charges for efficiency. We use the state_labels and perm to keep track of these permutations.

\section*{Parameters}
- leg (LegCharge) - Charges of the physical states, to be used for the physical leg of MPS.
- state_labels (None / list of str) - Optionally a label for each local basis states. None entries are ignored / not set.
- **site_ops - Additional keyword arguments of the form name=op given to add_op(). The identity operator 'Id' is automatically included. If no 'JW' for the Jordan-Wigner string is given, 'JW' is set as an alias to 'Id'.

\section*{leg}

Charges of the local basis states.
Type LegCharge
state_labels
(Optional) labels for the local basis states.
Type \{str: int \}

\section*{opnames}

Labels of all onsite operators (i.e. self.op exists if 'op' in self.opnames). Note that get_op() allows arbitrary concatenations of them.

Type set

\section*{need_JW_string}

Labels of all onsite operators that need a Jordan-Wigner string. Used in op_needs_JW () to determine whether an operator anticommutes or commutes with operators on other sites.

Type set
ops
Onsite operators are added directly as attributes to self. For example after self.add_op ('Sz', Sz) you can use self.Sz for the \(S z\) operator. All onsite operators have labels 'p', 'p*'.

Type Array

\section*{perm}

Index permutation of the physical leg compared to conserve=None, i.e. \(\quad \mathrm{OP}^{\text {_conserved }=}\) OP_nonconserved[np.ix_(perm, perm)] and perm[state_labels_conserved[ "some_state"]] == state_labels_nonconserved["some_state"].

Type 1D array
JW_exponent
Exponents of the 'JW' operator, such that self.JW.to_ndarray() = np.diag(np.exp(1. j*np.pi* JW_exponent))

Type 1D array

\section*{hc_ops}

Mapping from operator names to their hermitian conjugates. Use get_hc_op_name () to obtain entries.

\section*{Type dict(str->str)}

\section*{Examples}

The following generates a site for spin- \(1 / 2\) with \(S z\) conservation. Note that \(S x=(S p+S m) / 2\) violates \(S z\) conservation and is thus not a valid on-site operator.
```

>>> chinfo = npc.ChargeInfo([1], ['Sz'])
>>> ch = npc.LegCharge.from_qflat(chinfo, [1, -1])
>>> Sp = [[0, 1.], [0, 0]]
>>> Sm=[[0, 0], [1., 0]]
>>> Sz = [[0.5, 0], [0, -0.5]]
>>> site = tenpy.networks.site.Site(ch, ['up', 'down'], Splus=Sp, Sminus=Sm,
Sz=Sz)
>>> print(site.Splus.to_ndarray())
[[0. 1.]
[0. 0.]]
>>> print(site.get_op('Sminus').to_ndarray())
[[0. 0.]
[1. 0.]]
>>> print(site.get_op('Splus Sminus').to_ndarray())
[[1. 0.]
[0. 0.]]

```
change_charge (new_leg_charge=None, permute=None)

Change the charges of the site (in place).

\section*{Parameters}
- new_leg_charge (LegCharge I None) - The new charges to be used. If None, use trivial charges.
- permute (ndarray / None) - The permuation applied to the physical leg, which gets used to adjust state_labels and perm. If you sorted the previous leg with perm_qind, new_leg_charge = leg. sort(), use old_leg. perm_flat_from_perm_qind (perm_qind). Ignored if None.

\section*{test_sanity()}

Sanity check, raises ValueErrors, if something is wrong.

\section*{property dim}

Dimension of the local Hilbert space.

\section*{property onsite_ops}

Dictionary of on-site operators for iteration.
Single operators are accessible as attributes.
add_op (name, op, need_JW=False, \(h c=\) None)
Add one on-site operators.

\section*{Parameters}
- name (str) - A valid python variable name, used to label the operator. The name under which \(o p\) is added as attribute to self.
- op (np.ndarray I Array) - A matrix acting on the local hilbert space representing the local operator. Dense numpy arrays are automatically converted to Array. LegCharges have to be [leg, leg.conj()]. We set labels 'p', 'p*'.
- need_JW (bool) - Whether the operator needs a Jordan-Wigner string. If True, add name to need_JW_string.
- hc (None / False / str) - The name for the hermitian conjugate operator, to be used for hc_ops. By default (None), try to auto-determine it. If False, disable adding antries to hc_ops.
rename_op (old_name, new_name)
Rename an added operator.

\section*{Parameters}
- old_name (str) - The old name of the operator.
- new_name (str) - The new name of the operator.
remove_op (name)
Remove an added operator.
Parameters name (str) - The name of the operator to be removed.

\section*{state_index (label)}

Return index of a basis state from its label.
Parameters label (int / string) - eather the index directly or a label (string) set before.
Returns state_index - the index of the basis state associated with the label.
Return type int
state_indices (labels)
Same as state_index(), but for multiple labels.
get_op (name)
Return operator of given name.
Parameters name (str) - The name of the operator to be returned. In case of multiple operator names separated by whitespace, we multiply them together to a single on-site operator (with the one on the right acting first).

Returns op - The operator given by name, with labels ' \(p\) ', ' \(p\) *'. If name already was an npc Array, it's directly returned.
Return type np_conserved
get_hc_op_name (name)
Return the hermitian conjugate of a given operator.
Parameters name (str) - The name of the operator to be returned. Multiple operators separated by whitespace are interpreted as an operator product, exactly as get_op () does.
Returns hc_op_name - Operator name for the hermi such that get_op () of
Return type str
op_needs_JW (name)
Whether an (composite) onsite operator is fermionic and needs a Jordan-Wigner string.
Parameters name (str) - The name of the operator, as in get_op ().
Returns needs_JW - Whether the operator needs a Jordan-Wigner string, judging from need_JW_string.
Return type bool

\section*{valid_opname (name)}

Check whether 'name' labels a valid onsite-operator.
Parameters name (str) - Label for the operator. Can be multiple operator(labels) separated by whitespace, indicating that they should be multiplied together.

Returns valid - True if name is a valid argument to get_op ().
Return type bool
multiply_op_names (names)
Multiply operator names together.
Join the operator names in names such that get_op returns the product of the corresponding operators.
Parameters names (list of str)-List of valid operator labels.
Returns combined_opname - A valid operator name Operatorname representing the product of operators in names.
Return type str
multiply_operators (operators)
Multiply local operators (possibly given by their names) together.
Parameters operators (list of \(\{\operatorname{str} \mid\) Array \(\}\) ) - List of valid operator names (to be translated with get_op ()) or directly on-site operators in the form of npc arrays with 'p', 'p*' label. The operators are multiplied left-to-right.

Returns combined_operator - The product of the given operators in a left-to-right multiplication following the usual mathematical convention. For example, if operators=['Sz', 'Sp', 'Sx'], the final operator is equivalent to site.get_op('Sz Sp Sx'), with the ' \(S x\) ' operator acting first on any physical state.
Return type Array
classmethod from_hdf5 (hdf5_loader, h5gr, subpath)
Load instance from a HDF5 file.
This method reconstructs a class instance from the data saved with save_hdf5 ().

\section*{Parameters}
- hdf5_loader (Hdf5Loader) - Instance of the loading engine.
- h5gr (Group) - HDF5 group which is represent the object to be constructed.
- subpath (str) - The name of \(h 5 g r\) with a ' / ' in the end.

Returns obj - Newly generated class instance containing the required data.
Return type cls
save_hdf5 (hdf5_saver, h5gr, subpath)
Export self into a HDF5 file.
This method saves all the data it needs to reconstruct self with from_hdf5 ().
This implementation saves the content of __ dict__ with save_dict_content (), storing the format under the attribute 'format'.

\section*{Parameters}
- hdf5_saver (Hdf5Saver) - Instance of the saving engine.
- h5gr (: class `Group ') - HDF5 group which is supposed to represent self.
- subpath (str) - The name of \(h 5 g r\) with a ' / ' in the end.

\subsection*{20.1.5 SpinHalfFermionSite}
- full name: tenpy.networks.site.SpinHalfFermionSite
- parent module: tenpy. networks.site
- type: class

\section*{Inheritance Diagram}


\section*{Methods}
\begin{tabular}{ll}
\hline \begin{tabular}{l} 
SpinHalfFermionSite.__init__([cons_N, \\
\(\ldots])\).
\end{tabular} & Initialize self. \\
\hline SpinHalfFermionSite.add_op(name, op[,...]) & Add one on-site operators. \\
\hline \begin{tabular}{l} 
SpinHalfFermionSite. \\
change_charge([...])
\end{tabular} & Change the charges of the site (in place). \\
\hline \begin{tabular}{l} 
SpinHalfFermionSite. \\
from_hdf5(hdf5_loader, .. \()\)
\end{tabular} & Load instance from a HDF5 file. \\
\hline \begin{tabular}{l} 
SpinHalfFermionSite. \\
get_hc_op_name(name \()\)
\end{tabular} & Return the hermitian conjugate of a given operator. \\
\hline SpinHalfFermionSite.get_op(name) & Return operator of given name. \\
\hline \begin{tabular}{l} 
SpinHalfFermionSite. \\
multiply_op_names(names)
\end{tabular} & Multiply operator names together. \\
\hline SpinHalfFermionSite. & \begin{tabular}{l} 
Multiply local operators (possibly given by their names) \\
together.
\end{tabular} \\
\hline multiply_operators(operators) & \begin{tabular}{l} 
Whether an (composite) onsite operator is fermionic and \\
SpinHalfFermionSite.op_needs_JW(name)
\end{tabular} \\
\hline SpinHalfFermionSite.remove_op(name) & Remove an added operator. \\
\hline
\end{tabular}

Table 11 - continued from previous page
SpinHalfFermionSite.rename_op(old_name, Rename an added operator.
new_name)
\begin{tabular}{ll}
\hline \begin{tabular}{l} 
SpinHalfFermionSite. \\
save_hdf5(hdf5_saver,...)
\end{tabular} & Export self into a HDF5 file. \\
\hline SpinHalfFermionSite.state_index(label) & Return index of a basis state from its label. \\
\hline \begin{tabular}{l} 
SpinHalfFermionSite. \\
state_indices(labels)
\end{tabular} & Same as state_index(), but for multiple labels. \\
\hline SpinHalffermionSite.test_sanity() & Sanity check, raises ValueErrors, if something is wrong. \\
\hline SpinHalfFermionSite.valid_opname(name) & Check whether 'name' labels a valid onsite-operator. \\
\hline
\end{tabular}

\section*{Class Attributes and Properties}
\begin{tabular}{ll}
\hline SpinHalfFermionSite.dim & Dimension of the local Hilbert space. \\
\hline SpinHalfFermionSite.onsite_ops & Dictionary of on-site operators for iteration. \\
\hline
\end{tabular}
class tenpy.networks.site.SpinHalfFermionSite (cons_ \(N={ }^{\prime} N^{\prime}\), cons_Sz='Sz', filling=1.0)
Bases: tenpy.networks.site.Site
Create a Site for spinful (spin-1/2) fermions.
Local states are: empty (vacuum), up (one spin-up electron), down (one spin-down electron), and full (both electrons)

Local operators can be built from creation operators.

Warning: Using the Jordan-Wigner string (JW) in the correct way is crucial to get correct results, otherwise you just describe hardcore bosons!
\begin{tabular}{|c|c|}
\hline operator & description \\
\hline Id & Identity \(\nVdash\) \\
\hline JW & Sign for the Jordan-Wigner string (-1) \({ }^{n_{\uparrow}+n_{\downarrow}}\) \\
\hline JWu & Partial sign for the Jordan-Wigner string \((-1)^{n \uparrow}\) \\
\hline JWd & Partial sign for the Jordan-Wigner string \((-1)^{n} \downarrow\) \\
\hline Cu & Annihilation operator spin-up \(c_{\uparrow}\) (up to 'JW'-string on sites left of it). \\
\hline Cdu & Creation operator spin-up \(c_{\uparrow}^{\dagger}\) (up to 'JW'-string on sites left of it). \\
\hline Cd & Annihilation operator spin-down \(c_{\downarrow}\) (up to 'JW'-string on sites left of it). Includes JWu such that it anti-commutes onsite with \(\mathrm{Cu}, \mathrm{Cdu}\). \\
\hline Cdd & Creation operator spin-down \(c_{\downarrow}^{\dagger}\) (up to 'JW'-string on sites left of it). Includes JWu such that it anti-commutes onsite with Cu , Cdu. \\
\hline Nu & Number operator \(n_{\uparrow}=c_{\uparrow}^{\dagger} c_{\uparrow}\) \\
\hline Nd & Number operator \(n_{\downarrow}=c_{\downarrow}^{\dagger} c_{\downarrow}\) \\
\hline NuNd & Dotted number operators \(n_{\uparrow} n_{\downarrow}\) \\
\hline Ntot & Total number operator \(n_{t}=n_{\uparrow}+n_{\downarrow}\) \\
\hline dN & Total number operator compared to the filling \(\Delta n=n_{t}-\) filling \\
\hline \[
\begin{aligned}
& \text { Sx, } \\
& \text { Sy, Sz }
\end{aligned}
\] & Spin operators \(S^{x, y, z}\), in particular \(S^{z}=\frac{1}{2}\left(n_{\uparrow}-n_{\downarrow}\right)\) \\
\hline Sp, Sm & Spin flips \(S^{ \pm}=S^{x} \pm i S^{y}\), e.g. \(S^{+}=c_{\uparrow}^{\dagger} c_{\downarrow}\) \\
\hline
\end{tabular}

The spin operators are defined as \(S^{\gamma}=\left(c_{\uparrow}^{\dagger}, c_{\downarrow}^{\dagger}\right) \sigma^{\gamma}\left(c_{\uparrow}, c_{\downarrow}\right)^{T}\), where \(\sigma^{\gamma}\) are spin- \(1 / 2\) matrices (i.e. half the pauli matrices).
\begin{tabular}{|l|l|l|l|}
\hline cons_N & Cons_Sz & qmod & excluded onsite operators \\
\hline 'N' & 'Sz' & {\([1,1]\)} & Sx, Sy \\
\hline 'N' & 'parity' & {\([1,2]\)} & - \\
\hline\(N^{\prime} N^{\prime}\) & None & {\([1]\)} & - \\
\hline 'parity' & 'Sz' & {\([2,1]\)} & Sx, Sy \\
\hline 'parity' & 'parity' & {\([2,2]\)} & - \\
\hline 'parity' & None & {\([2]\)} & - \\
\hline None & 'Sz' & {\([1]\)} & Sx, Sy \\
\hline None & 'parity' & {\([2]\)} & - \\
\hline None & None & {[]} & - \\
\hline
\end{tabular}

\section*{Parameters}
- cons_N('N' | 'parity' | None) - Whether particle number is conserved, c.f. table above.
- cons_Sz('Sz' | 'parity' | None)-Whether spin is conserved, c.f. table above.
- filling (float) - Average filling. Used to define dN .
cons_N
Whether particle number is conserved, c.f. table above.
Type 'N' | 'parity' | None
cons_Sz
Whether spin is conserved, c.f. table above.
Type 'Sz' | 'parity' | None
filling
Average filling. Used to define \(d N\).
Type float
add_op (name, op, need_JW=False, hc=None)
Add one on-site operators.

\section*{Parameters}
- name (str) - A valid python variable name, used to label the operator. The name under which \(o p\) is added as attribute to self.
- op (np.ndarray I Array) - A matrix acting on the local hilbert space representing the local operator. Dense numpy arrays are automatically converted to Array. LegCharges have to be [leg, leg.conj()]. We set labels 'p', 'p*'.
- need_JW (bool) - Whether the operator needs a Jordan-Wigner string. If True, add name to need_JW_string.
- hc (None / False / str) - The name for the hermitian conjugate operator, to be used for hc_ops. By default (None), try to auto-determine it. If False, disable adding antries to hc_ops.
change_charge (new_leg_charge=None, permute=None)
Change the charges of the site (in place).

\section*{Parameters}
- new_leg_charge (LegCharge I None) - The new charges to be used. If None, use trivial charges.
- permute (ndarray / None) - The permuation applied to the physical leg, which gets used to adjust state_labels and perm. If you sorted the previous leg with perm_qind, new_leg_charge = leg.sort(), use old_leg. perm_flat_from_perm_qind(perm_qind). Ignored if None.

\section*{property dim}

Dimension of the local Hilbert space.
classmethod from_hdf5 (hdf5_loader, h5gr, subpath)
Load instance from a HDF5 file.
This method reconstructs a class instance from the data saved with save_hdf5 ().

\section*{Parameters}
- hdf5_loader (Hdf5Loader) - Instance of the loading engine.
- h5gr (Group) - HDF5 group which is represent the object to be constructed.
- subpath (str) - The name of \(h 5 g r\) with a ' /' in the end.

Returns obj - Newly generated class instance containing the required data.
Return type cls
get_hc_op_name (name)
Return the hermitian conjugate of a given operator.
Parameters name (str) - The name of the operator to be returned. Multiple operators separated by whitespace are interpreted as an operator product, exactly as get_op () does.
Returns hc_op_name - Operator name for the hermi such that get_op () of
Return type str
get_op (name)
Return operator of given name.
Parameters name (str) - The name of the operator to be returned. In case of multiple operator names separated by whitespace, we multiply them together to a single on-site operator (with the one on the right acting first).

Returns op - The operator given by name, with labels ' p ', ' p ' . If name already was an npc Array, it's directly returned.
Return type np_conserved
multiply_op_names (names)
Multiply operator names together.
Join the operator names in names such that get_op returns the product of the corresponding operators.
Parameters names (list of str)- List of valid operator labels.
Returns combined_opname - A valid operator name Operatorname representing the product of operators in names.

\section*{Return type str}
multiply_operators (operators)
Multiply local operators (possibly given by their names) together.

Parameters operators (list of \(\{\operatorname{str} \mid\) Array \(\}\) ) - List of valid operator names (to be translated with get_op()) or directly on-site operators in the form of npe arrays with 'p', 'p*' label. The operators are multiplied left-to-right.

Returns combined_operator - The product of the given operators in a left-to-right multiplication following the usual mathematical convention. For example, if operators=['Sz', 'Sp', 'Sx'], the final operator is equivalent to site.get_op('Sz Sp Sx'), with the ' \(S x\) ' operator acting first on any physical state.

\section*{Return type Array}
property onsite_ops
Dictionary of on-site operators for iteration.
Single operators are accessible as attributes.
```

op_needs_JW(name)

```

Whether an (composite) onsite operator is fermionic and needs a Jordan-Wigner string.
Parameters name (str) - The name of the operator, as in get_op ().
Returns needs_JW - Whether the operator needs a Jordan-Wigner string, judging from need_JW_string.

\section*{Return type bool}
```

remove_op (name)

```

Remove an added operator.
Parameters name (str) - The name of the operator to be removed.
rename_op (old_name, new_name)
Rename an added operator.

\section*{Parameters}
- old_name (str) - The old name of the operator.
- new_name (str) - The new name of the operator.

\section*{save_hdf5 (hdf5_saver, h5gr, subpath)}

Export self into a HDF5 file.
This method saves all the data it needs to reconstruct self with from_hdf5 ().
This implementation saves the content of __dict__ with save_dict_content (), storing the format under the attribute 'format'.

\section*{Parameters}
- hdf5_saver (Hdf5Saver) - Instance of the saving engine.
- h5gr (: class`Group ') - HDF5 group which is supposed to represent self.
- subpath (str) - The name of \(h 5 g r\) with a '/' in the end.

\section*{state_index (label)}

Return index of a basis state from its label.
Parameters label (int / string)-eather the index directly or a label (string) set before.
Returns state_index - the index of the basis state associated with the label.
Return type int
state_indices (labels)
Same as state_index(), but for multiple labels.
```

test_sanity()

```

Sanity check, raises ValueErrors, if something is wrong.
```

valid_opname (name)

```

Check whether 'name' labels a valid onsite-operator.
Parameters name (str) - Label for the operator. Can be multiple operator(labels) separated by whitespace, indicating that they should be multiplied together.

Returns valid - True if name is a valid argument to get_op ().
Return type bool

\subsection*{20.1.6 SpinHalfSite}
- full name: tenpy.networks.site.SpinHalfSite
- parent module: tenpy.networks.site
- type: class

\section*{Inheritance Diagram}


\section*{Methods}
\begin{tabular}{ll}
\hline SpinHalfSite.__init__([conserve]) & Initialize self. \\
\hline SpinHalfSite.add_op(name, op[, need_JW, hc]) & Add one on-site operators. \\
\hline SpinHalfSite.change_charge([new_leg_charge, & Change the charges of the site (in place). \\
\(\ldots\).\(] )\) & \\
\hline SpinHalfSite.from_hdf5(hdf5_loader, & h5gr, \\
..) & Load instance from a HDF5 file. \\
\hline SpinHalfSite.get_hc_op_name(name) & Return the hermitian conjugate of a given operator. \\
\hline SpinHalfSite.get_op(name) & Return operator of given name. \\
\hline SpinHalfSite.multiply_op_names(names) & Multiply operator names together. \\
\hline
\end{tabular}

Table 13 - continued from previous page
SpinHalfSite.multiply_operators(operators) Multiply local operators (possibly given by their names) together.
SpinHalfSite.op_needs_JW(name) Whether an (composite) onsite operator is fermionic and needs a Jordan-Wigner string.
\begin{tabular}{ll}
\hline SpinHalfSite.remove_op(name) & Remove an added operator. \\
\hline \begin{tabular}{l} 
SpinHalfSite.rename_op(old_name, \\
new_name)
\end{tabular} & Rename an added operator. \\
\hline \begin{tabular}{l} 
SpinHalfSite.save_hdf5(hdf5_saver, \\
subpath)
\end{tabular} & h5gr,
\end{tabular} Export self into a HDF5 file. \(\quad\).

\section*{Class Attributes and Properties}
\begin{tabular}{ll}
\hline SpinHalfSite.dim & Dimension of the local Hilbert space. \\
\hline SpinHalfSite.onsite_ops & Dictionary of on-site operators for iteration. \\
\hline
\end{tabular}
class tenpy.networks.site.SpinHalfSite (conserve='Sz')
Bases: tenpy. networks.site. Site
Spin-1/2 site.
Local states are up (0) and down (1). Local operators are the usual spin- \(1 / 2\) operators, e.g. \(\mathrm{Sz}=[[0.5\), 0.], [0., -0.5]], Sx = 0.5*sigma_x for the Pauli matrix sigma_x.
\begin{tabular}{|l|l|}
\hline Operator & description \\
\hline Id, JW & Identity \(\nVdash\) \\
\hline Sx, Sy, Sz & Spin components \(S^{x, y, z}\), equal to half the Pauli matrices. \\
\hline Sigmax, Sigmay, Sigmaz & Pauli matrices \(\sigma^{x, y, z}\) \\
\hline Sp, Sm & Spin flips \(S^{ \pm}=S^{x} \pm i S^{y}\) \\
\hline
\end{tabular}
\begin{tabular}{|l|l|l|}
\hline Conserve & qmod & excluded onsite operators \\
\hline 'Sz' & {\([1]\)} & Sx, Sy, Sigmax, Sigmay \\
\hline 'parity' & {\([2]\)} & - \\
\hline None & {[]} & - \\
\hline
\end{tabular}

Parameters conserve (str) - Defines what is conserved, see table above.

\section*{conserve}

Defines what is conserved, see table above.
Type str
add_op (name, op, need_JW=False, hc=None)
Add one on-site operators.

\section*{Parameters}
- name (str) - A valid python variable name, used to label the operator. The name under which \(o p\) is added as attribute to self.
- op (np.ndarray I Array) - A matrix acting on the local hilbert space representing the local operator. Dense numpy arrays are automatically converted to Array. LegCharges have to be [leg, leg.conj()]. We set labels 'p', 'p*'.
- need_JW (bool) - Whether the operator needs a Jordan-Wigner string. If True, add name to need_JW_string.
- hc (None / False / str) - The name for the hermitian conjugate operator, to be used for hc_ops. By default (None), try to auto-determine it. If False, disable adding antries to hc_ops.
change_charge (new_leg_charge=None, permute=None)
Change the charges of the site (in place).

\section*{Parameters}
- new_leg_charge (LegCharge I None) - The new charges to be used. If None, use trivial charges.
- permute (ndarray / None) - The permuation applied to the physical leg, which gets used to adjust state_labels and perm. If you sorted the previous leg with perm_qind, new_leg_charge \(=\) leg. sort(), use old_leg. perm_flat_from_perm_qind(perm_qind). Ignored if None.

\section*{property dim}

Dimension of the local Hilbert space.
classmethod from_hdf5 (hdf5_loader, h5gr, subpath)
Load instance from a HDF5 file.
This method reconstructs a class instance from the data saved with save_hdf5().

\section*{Parameters}
- hdf5_loader (Hdf5Loader) - Instance of the loading engine.
- h5gr (Group) - HDF5 group which is represent the object to be constructed.
- subpath (str) - The name of \(h 5 g r\) with a '/' in the end.

Returns obj - Newly generated class instance containing the required data.
Return type cls
get_hc_op_name (name)
Return the hermitian conjugate of a given operator.
Parameters name (str) - The name of the operator to be returned. Multiple operators separated by whitespace are interpreted as an operator product, exactly as get_op () does.
Returns hc_op_name - Operator name for the hermi such that get_op () of
Return type str
get_op (name)
Return operator of given name.
Parameters name (str) - The name of the operator to be returned. In case of multiple operator names separated by whitespace, we multiply them together to a single on-site operator (with the one on the right acting first).
Returns op - The operator given by name, with labels ' p ', ' p *'. If name already was an npc Array, it's directly returned.

Return type np_conserved
multiply_op_names (names)
Multiply operator names together.
Join the operator names in names such that get_op returns the product of the corresponding operators.
Parameters names (list of str) - List of valid operator labels.
Returns combined_opname - A valid operator name Operatorname representing the product of operators in names.

Return type str
multiply_operators (operators)
Multiply local operators (possibly given by their names) together.
Parameters operators (list of \(\{\operatorname{str|Array}\}\) ) - List of valid operator names (to be translated with get_op ()) or directly on-site operators in the form of npc arrays with 'p', 'p*' label. The operators are multiplied left-to-right.

Returns combined_operator - The product of the given operators in a left-to-right multiplication following the usual mathematical convention. For example, if operators=['Sz', 'Sp', 'Sx'], the final operator is equivalent to site.get_op('Sz Sp Sx'), with the ' \(S x\) ' operator acting first on any physical state.

\section*{Return type Array}
property onsite_ops
Dictionary of on-site operators for iteration.
Single operators are accessible as attributes.
```

OP_needs_JW (name)

```

Whether an (composite) onsite operator is fermionic and needs a Jordan-Wigner string.
Parameters name (str) - The name of the operator, as in get_op ().
Returns needs_JW - Whether the operator needs a Jordan-Wigner string, judging from need_JW_string.

\section*{Return type bool}
```

remove_op (name)

```

Remove an added operator.
Parameters name (str) - The name of the operator to be removed.
rename_op (old_name, new_name)
Rename an added operator.

\section*{Parameters}
- old_name (str) - The old name of the operator.
- new_name (str) - The new name of the operator.
save_hdf5 (hdf5_saver, h5gr, subpath)
Export self into a HDF5 file.
This method saves all the data it needs to reconstruct self with from_hdf5 ().
This implementation saves the content of __dict__ with save_dict_content (), storing the format under the attribute 'format'.

\section*{Parameters}
- hdf5_saver (Hdf5Saver) - Instance of the saving engine.
- h5gr (: class `Group`) - HDF5 group which is supposed to represent self.
- subpath (str) - The name of \(h 5 g r\) with a '/' in the end.
state_index (label)
Return index of a basis state from its label.
Parameters label (int | string)-eather the index directly or a label (string) set before.
Returns state_index - the index of the basis state associated with the label.
Return type int
state_indices (labels)
Same as state_index(), but for multiple labels.
test_sanity()
Sanity check, raises ValueErrors, if something is wrong.
valid_opname (name)
Check whether 'name' labels a valid onsite-operator.
Parameters name (str) - Label for the operator. Can be multiple operator(labels) separated by whitespace, indicating that they should be multiplied together.

Returns valid - True if name is a valid argument to get_op ().
Return type bool

\subsection*{20.1.7 SpinSite}
- full name: tenpy.networks.site.SpinSite
- parent module: tenpy.networks.site
- type: class

Inheritance Diagram


\section*{Methods}
\begin{tabular}{ll}
\hline SpinSite.__init__([S, conserve]) & Initialize self. \\
\hline SpinSite.add_op(name, op[, need_JW, hc]) & Add one on-site operators. \\
\hline \begin{tabular}{l} 
SpinSite.change_charge([new_leg_charge, \\
permute])
\end{tabular} & Change the charges of the site (in place). \\
\hline SpinSite.from_hdf5(hdf5_loader, h5gr, subpath) & Load instance from a HDF5 file. \\
\hline SpinSite.get_hc_op_name(name) & Return the hermitian conjugate of a given operator. \\
\hline SpinSite.get_op(name) & Return operator of given name. \\
\hline SpinSite.multiply_op_names(names) & Multiply operator names together. \\
\hline SpinSite.multiply_operators(operators) & \begin{tabular}{l} 
Multiply local operators (possibly given by their names) \\
together.
\end{tabular} \\
\hline SpinSite.op_needs_JW(name) & \begin{tabular}{l} 
Whether an (composite) onsite operator is fermionic and \\
needs a Jordan-Wigner string.
\end{tabular} \\
\hline SpinSite.remove_op(name) & Remove an added operator. \\
\hline SpinSite.rename_op(old_name, new_name) & Rename an added operator. \\
\hline SpinSite.save_hdf5(hdf5_saver, h5gr, subpath) & Export self into a HDF5 file. \\
\hline SpinSite.state_index(label) & Return index of a basis state from its label. \\
\hline SpinSite.state_indices(labels) & Same as state_index(), but for multiple labels. \\
\hline SpinSite.test_sanity() & Sanity check, raises ValueErrors, if something is wrong. \\
\hline SpinSite.valid_opname(name) & Check whether 'name' labels a valid onsite-operator. \\
\hline
\end{tabular}

\section*{Class Attributes and Properties}
\begin{tabular}{ll}
\hline SpinSite.dim & Dimension of the local Hilbert space. \\
\hline SpinSite.onsite_ops & Dictionary of on-site operators for iteration. \\
\hline
\end{tabular}
class tenpy.networks.site.SpinSite ( \(S=0.5\), conserve='Sz')
Bases: tenpy.networks.site.Site
General Spin S site.
There are \(2 S+1\) local states range from down (0) to up \((2 S+1)\), corresponding to \(S z=-S,-S+1, \ldots\), \(S-1, S\) Local operators are the spin-S operators, e.g. \(S z=[[0.5,0],.[0 .,-0.5]], S x=0\). \(5 *\) sigma_x for the Pauli matrix sigma_x.
\begin{tabular}{|c|c|}
\hline operator & description \\
\hline Id, JW & Identity \(\nVdash\) \\
\hline Sx, Sy, Sz & Spin components \(S^{x, y, z}\), equal to half the Pauli matrices. \\
\hline Sp, Sm & Spin flips \(S^{ \pm}=S^{x} \pm i S^{y}\) \\
\hline
\end{tabular}
\begin{tabular}{|l|l|l|}
\hline conserve & qmod & excluded onsite operators \\
\hline 'Sz' & {\([1]\)} & Sx, Sy \\
\hline 'parity' & {\([2]\)} & - \\
\hline None & {[]} & - \\
\hline
\end{tabular}

Parameters conserve (str) - Defines what is conserved, see table above.

S
The \(2 \mathrm{~S}+1\) states range from \(\mathrm{m}=-\mathrm{S},-\mathrm{S}+1, \ldots+\mathrm{S}\).

Type \(\{0.5,1,1.5,2, .\).

\section*{conserve}

Defines what is conserved, see table above.
Type str
add_op (name, op, need_JW=False, \(h c=\) None)
Add one on-site operators.

\section*{Parameters}
- name (str) - A valid python variable name, used to label the operator. The name under which \(o p\) is added as attribute to self.
- op (np.ndarray I Array) - A matrix acting on the local hilbert space representing the local operator. Dense numpy arrays are automatically converted to Array. LegCharges have to be [leg, leg.conj()]. We set labels 'p', 'p*'.
- need_JW (bool) - Whether the operator needs a Jordan-Wigner string. If True, add name to need_JW_string.
- hc (None | False | str) - The name for the hermitian conjugate operator, to be used for hc_ops. By default (None), try to auto-determine it. If False, disable adding antries to hc_ops.
change_charge (new_leg_charge=None, permute=None)
Change the charges of the site (in place).

\section*{Parameters}
- new_leg_charge (LegCharge I None) - The new charges to be used. If None, use trivial charges.
- permute (ndarray | None) - The permuation applied to the physical leg, which gets used to adjust state_labels and perm. If you sorted the previous leg with perm_qind, new_leg_charge = leg.sort(), use old_leg. perm_flat_from_perm_qind (perm_qind). Ignored if None.

\section*{property dim}

Dimension of the local Hilbert space.
```

classmethod from_hdf5 (hdf5_loader, h5gr, subpath)

```

Load instance from a HDF5 file.
This method reconstructs a class instance from the data saved with save_hdf5 ().

\section*{Parameters}
- hdf5_loader (Hdf5Loader) - Instance of the loading engine.
- h5gr (Group) - HDF5 group which is represent the object to be constructed.
- subpath (str) - The name of \(h 5 g r\) with a ' / ' in the end.

Returns obj - Newly generated class instance containing the required data.
Return type cls
get_hc_op_name (name)
Return the hermitian conjugate of a given operator.
Parameters name (str) - The name of the operator to be returned. Multiple operators separated by whitespace are interpreted as an operator product, exactly as get_op () does.
Returns hc_op_name - Operator name for the hermi such that get_op () of

Return type str
get_op (name)
Return operator of given name.
Parameters name (str) - The name of the operator to be returned. In case of multiple operator names separated by whitespace, we multiply them together to a single on-site operator (with the one on the right acting first).

Returns op - The operator given by name, with labels 'p', 'p*'. If name already was an npc Array, it's directly returned.

Return type np_conserved
multiply_op_names (names)
Multiply operator names together.
Join the operator names in names such that get_op returns the product of the corresponding operators.
Parameters names (list of str) - List of valid operator labels.
Returns combined_opname - A valid operator name Operatorname representing the product of operators in names.

Return type str
multiply_operators (operators)
Multiply local operators (possibly given by their names) together.
Parameters operators (list of \(\{\operatorname{str|Array}\}\) ) - List of valid operator names (to be translated with \(\left.g e t \_o p()\right)\) or directly on-site operators in the form of npc arrays with 'p', 'p*' label. The operators are multiplied left-to-right.
Returns combined_operator - The product of the given operators in a left-to-right multiplication following the usual mathematical convention. For example, if operators=['Sz', 'Sp', 'Sx'], the final operator is equivalent to site.get_op('Sz Sp Sx'), with the ' \(S x\) ' operator acting first on any physical state.

Return type Array
property onsite_ops
Dictionary of on-site operators for iteration.
Single operators are accessible as attributes.
```

OP_needs_JW (name)

```

Whether an (composite) onsite operator is fermionic and needs a Jordan-Wigner string.
Parameters name (str) - The name of the operator, as in get_op ().
Returns needs_JW - Whether the operator needs a Jordan-Wigner string, judging from need_JW_string.

\section*{Return type bool}
```

remove_op (name)

```

Remove an added operator.
Parameters name (str) - The name of the operator to be removed.
rename_op (old_name, new_name)
Rename an added operator.

\section*{Parameters}
- old_name (str) - The old name of the operator.
- new_name (str) - The new name of the operator.
save_hdf5 (hdf5_saver, h5gr, subpath)
Export self into a HDF5 file.
This method saves all the data it needs to reconstruct self with from_hdf5 ().
This implementation saves the content of __ dict__ with save_dict_content (), storing the format under the attribute 'format'.

\section*{Parameters}
- hdf5_saver (Hdf5Saver) - Instance of the saving engine.
- h5gr (: class `Group \()\) - HDF5 group which is supposed to represent self.
- subpath (str) - The name of \(h 5 g r\) with a '/' in the end.
state_index (label)
Return index of a basis state from its label.
Parameters label (int | string)-eather the index directly or a label (string) set before.
Returns state_index - the index of the basis state associated with the label.
Return type int
state_indices (labels)
Same as state_index(), but for multiple labels.
test_sanity()
Sanity check, raises ValueErrors, if something is wrong.
valid_opname (name)
Check whether 'name' labels a valid onsite-operator.
Parameters name (str) - Label for the operator. Can be multiple operator(labels) separated by whitespace, indicating that they should be multiplied together.

Returns valid - True if name is a valid argument to get_op ().
Return type bool

\section*{Functions}
\begin{tabular}{ll}
\hline group_sites(sites[, n, labels, charges]) & Given a list of sites, group each \(n\) sites together. \\
\hline multi_sites_combine_charges(sites[,..]) & \begin{tabular}{l} 
Adjust the charges of the given sites (in place) such that \\
they can be used together.
\end{tabular} \\
\hline set_common_charges(sites[, new_charges,...]) & \begin{tabular}{l} 
Adjust the charges of the given sites in place such that \\
they can be used together.
\end{tabular} \\
\hline
\end{tabular}

\subsection*{20.1.8 group_sites}
- full name: tenpy.networks.site.group_sites
- parent module: tenpy.networks.site
- type: function
tenpy.networks.site.group_sites (sites, \(n=2\), labels=None, charges='same')
Given a list of sites, group each \(n\) sites together.

\section*{Parameters}
- sites (list of Site) - The sites to be grouped together.
- \(\mathbf{n}(i n t)\) - We group each \(n\) consecutive sites from sites together in a GroupedSite.
- labels - See GroupedSites.
- charges - See GroupedSites.

Returns grouped_sites - The grouped sites. Has length (len(sites)-1)//n + 1 .
Return type list of GroupedSite

\subsection*{20.1.9 multi_sites_combine_charges}
- full name: tenpy.networks.site.multi_sites_combine_charges
- parent module: tenpy. networks.site
- type: function
tenpy.networks.site.multi_sites_combine_charges (sites, same_charges=[])
Adjust the charges of the given sites (in place) such that they can be used together.
When we want to contract tensors corresponding to different Site instances, these sites need to share a single ChargeInfo. This function adjusts the charges of these sites such that they can be used together.

Deprecated since version 0.7.3: Deprecated in favore of the new, more powerful set_common_charges (). Be aware of the slightly different argument structure though, namely that this function keeps charges not included in same_charges, whereas you need to include them explicitly into the new_charges argument of set_common_charges.

\section*{Parameters}
- sites (list of Site) - The sites to be combined. Modified in place.
- same_charges ([[(int, int|str), (int, int|str), ...], ...]) Defines which charges actually are the same, i.e. their quantum numbers are added up. Each charge is specified by a tuple ( \(s, i\) ) = (int, int|str), where \(s\) gives the index of the site within sites and \(i\) the index or name of the charge in the ChargeInfo of this site.

Returns perms - For each site the permutation performed on the physical leg to sort by charges.
Return type list of ndarray

\section*{Examples}
```

>>> from tenpy.networks.site import *
>>> ferm = SpinHalfFermionSite(cons_N='N', cons_Sz='Sz')
>>> spin = SpinSite(1.0, 'Sz')
>>> ferm.leg.chinfo is spin.leg.chinfo
False
>>> print(spin.leg)
+1
0 [ [-2]
1 [ 0]
[ 2]]
3
>>> multi_sites_combine_charges([ferm, spin], same_charges=[[(0, 1), (1, 0)]])
[array([0, 1, 2, 3]), array([0, 1, 2])]
>>> \# no permutations where needed
>>> ferm.leg.chinfo is spin.leg.chinfo
True
>>> ferm.leg.chinfo.names
['N', '2*Sz']
>>> print(spin.leg)
+1
0 [[$$
\begin{array}{lll}{0}&{-2]}\end{array}
$$]
[ 0 0]
[ 0 2]]
3

```

\subsection*{20.1.10 set_common_charges}
- full name: tenpy.networks.site.set_common_charges
- parent module: tenpy. networks.site
- type: function
tenpy.networks.site.set_common_charges (sites, new_charges='same', new_names=None, new_mod=None)
Adjust the charges of the given sites in place such that they can be used together.
Before we can contract operators (and tensors) corresponding to different Site instances, we first need to define the overall conserved charges, i.e., we need to merge the ChargeInfo of them to a single, global chinfo and adjust the charges of the physical legs. That's what this function does.

A typical place to do this would be in tenpy.models.model. CouplingMPOModel.init_sites().
(This function replaces the now deprecated mutli_sites_combine_charges().)

\section*{Parameters}
- sites (list of Site) - The sites to be combined. The sites are modified in place.
- new_charges ('same'|'drop'|'independent' | list of list of tuple) - Defines the new, common charges in terms of the old ones.
list of lists of tuple If a list is given, each entry new_charge of the list defines one new charge, i.e. the new number of charges is qnumber=len (new_charges). Each entry new_charge of the outer list is itself a list of 3-tuples, new_charge = [ (factor, site_index, old_charge_index), ...]. where the value of the new charge is the sum of factor times the value of the old charge, (specified by the site_index and the old_charge_index within that site), and the sum runs over all entries in that list
new_charge. old_charge_index can be an integer (=the index) or a string (=the name) of the charge in the corresponding sites[site_index].leg.chinfo.
' same' defaults to charges with the same name to match, and charges with different names to be independently conserved (see example below); None-set names are considered different.
'drop' Drop/remove all charges, equivalent to new_charges=[].
'independent ' For the case that the charges of the different sites are independent and individually conserved, even if they have the same name.
- new_names (list of str) - Names for each of the new charges. Defaults to name of the first old charge specified.
- new_mod (list of int) - mod for the new charges, one entry for each list in new_charges. Defaults to the mod of the old charges, if not specified otherwise.

Returns perms - For each site the permutation performed on the physical leg to sort by charges.
Return type list of ndarray

\section*{Examples}

When we just initialize some sites, they will in general have different charges. For example, we could have a SpinHalfFermionSite a spin-1 SpinSite. For reference, let's also print the names and values of the charges.
```

>>> from tenpy.networks.site import *
>>> ferm = SpinHalfFermionSite(cons_N='N', cons_Sz='Sz')
>>> ferm.leg.chinfo.names
['N', '2*Sz']
>>> print(ferm.leg.to_qflat())
[[[$$
\begin{array}{lll}{1}&{1}&{-1}\end{array}
$$]
[ 0}00
[ 2 0]
[ [1 1]]
>>> spin = SpinSite(1.0, conserve='Sz')
>>> spin.leg.chinfo.names
['2*Sz']
>>> print(spin.leg.to_qflat())
[ [-2]
[ 0]
[ 2]]

```

With the default new_charges='same', this function will combine charges with the same name, and hence we will have two conserved quantities, namley the fermion particle number ' \(N\) ' \(=N\) _ \(\{\) up_fermions \} \(+N_{\_}\)\{down-fermions\}, and the total Sz spin ' \(2 *\) Sz' \(=N_{-}\)\{up-fermions\} + N_\{up-spins \} - N_\{down-fermions\} - N_\{down-spins\}. In this case, there will only appear an extra column of zeros for the charges of the spin leg.
```

>>> set_common_charges([ferm, spin], new_charges='same')
[array([0, 1, 2, 3]), array([0, 1, 2])]
>>> ferm.leg.chinfo.names
['N', '2*Sz']
>>> print(ferm.leg.to_qflat()) \# didn't change (except making a copy)
[[[$$
\begin{array}{lll}{1}&{1}&{-1}\end{array}
$$]
[ 0}00
[ 2 0]

```
```

[ 1 1]]
>>> spin.leg.chinfo.names \# additional 'N' chargename
['N', '2*Sz']
>>> print(spin.leg.to_qflat()) \# additional column of zeros for the 'N' charge
[[$$
\begin{array}{lll}{0}&{-2]}\end{array}
$$]
[ [ 0 0]
[ 0 2]]

```

With new_charges='independent', we preserve the charges of the old sites individually. In this example, we get 3 conserved quantities, namely the fermion particle number ' \(N\) _ferm' \(=\) N_\{up_fermions \} \(+N_{\_}\{d o w n-f e r m i o n s\}\), and the fermionic \(S z\) spin \(' 2 * S z \_f e r m '=N_{\text {l }}\{u p-f e r m i o n s\}-\) N_\{down-fermions\} and the Sz spin of the spin sites, ' \(2 *\) Sz_spin' = N_\{up-spins\} \(\mathrm{N}_{\text {_ }}\{\) down-spins \}. (We give the charges new names for clearer distinction.) Corresponding zero columns are added to the LegCharges.
```

>>> ferm = SpinHalfFermionSite(cons_N='N', cons_Sz='Sz')
>>> spin = SpinSite(1.0, conserve='Sz')
>>> set_common_charges([ferm, spin], new_charges='independent',
... new_names=['N_ferm', '2*Sz_ferm', '2*Sz_spin'])
[array([0, 1, 2, 3]), array([0, 1, 2])]
>>> print(ferm.leg.to_qflat()) \# didn't change (except making a copy)
[[[$$
\begin{array}{llll}{1}&{1}&{-1}&{0}\end{array}
$$]
[ 0}0000
[ 2 0 0]
[ 1 1 1 0]]
>>> print(spin.leg.to_qflat()) \# additional column of zeros for the 'N' charge
[[[ 0
[ 0
[ 0

```

With the full specification of the new_charges through a list of list of tuples, you can create new charges as linear combinations of the charges of the individual sites. For example, the SpinHalfFermionSite is essentially the product of two FermionSite, one for the up electrons, and one for the down electrons. The ' \(2 \star S z\) ' charge of the SpinHalfFermionSite is then equivalent to the difference of individual particle numbers, \({ }^{\prime} 2 * S z^{\prime}=\) N_\{up\} - N_\{down\}.
```

>>> f_up = FermionSite(conserve='N')
>>> f_down = FermionSite(conserve='N')
>>> print(f_up.leg.to_qflat())
[ [0]
[1]]
>>> print(f_down.leg.to_qflat())
[ [0]
[1]]
>>> f_down.state_labels
{'empty': 0, 'full': 1}
>>> set_common_charges([f_up, f_down],
... new_charges=[[(1, 0, 'N'), ( 1, 1, 'N')],
... [(1, 0, 'N'), (-1, 1, 'N')]],
... new_names=['N_tot', '2*Sz=(N__up-N_down)'])
[array([0, 1]), array([1, 0])]
>>> f_down.state_labels \# sorting charges caused permutation of local states
{'empty': 1, 'full': 0}
>>> print(f_up.leg.to_qflat())
[[[0 0]
[1 1]]

```
```

>>> print(f_down.leg.to_qflat()) \# top row = full, bottom row=empty
[[[$$
\begin{array}{lll}{1}&{1}&{-1}\end{array}
$$]
[ 0 0]]

```

Another example could be that you have both fermions and bosons, and that you have terms \(c_{i} c_{j} b_{k}^{\dagger}+c_{i}^{\dagger} c_{j}^{\dagger} b_{k}\), where two fermions can merge into a pair forming a boson. In this case, neither the fermion number nor the boson number is preserved individually, but the combination \(N_{\_}\{\)fermions \(\}+2 * N_{\_}\{\)bosons \(\}\)is preserved.
```

>>> ferm = FermionSite(conserve='N')
>>> bos = BosonSite(Nmax=3, conserve='N')
>>> set_common_charges([ferm, bos], [[(1, 0, 'N'), (2, 1, 'N')]], ['N_f + 2 N_b'])
[array([0, 1]), array([0, 1, 2, 3])]

```

Finally, it can sometimes be convenient to change the charges of the The new_charges='drop' or new_charges = [ ] option is a quick way to remove any charges.
```

>>> ferm = SpinHalfFermionSite(cons_N='N', cons_Sz='Sz')
>>> spin = SpinSite(1.0, conserve='Sz')
>>> set_common_charges([ferm, spin], new_charges='drop')
[array([0, 1, 2, 3]), array([0, 1, 2])]
>>> assert ferm.leg.chinfo.qnumber == spin.leg.chinfo.qnumber == 0 \# trivial: nos
->charges

```

\section*{Module description}

Defines a class describing the local physical Hilbert space.
The site is the prototype, read it's docstring.

\section*{20.2 mps}
- full name: tenpy.networks.mps
- parent module: tenpy. networks
- type: module

\section*{Classes}

\begin{tabular}{ll}
\hline InitialStateBuilder(lattice, options[,...]) & Class to simplify providing common sets of intial states. \\
\hline MPS(sites, Bs, SVs[, bc, form, norm]) & A Matrix Product State, finite (MPS) or infinite (iMPS). \\
\hline MPSEnvironment(bra, ket[, init_LP, init_RP, ...]) & \begin{tabular}{l} 
Stores partial contractions of \(<b r a|O p| k e t>\) for local \\
operators \(O p\).
\end{tabular} \\
\hline TransferMatrix(bra, ket[, shift_bra, ...]) & Transfer matrix of two MPS (bra \& ket). \\
\hline
\end{tabular}

\subsection*{20.2.1 MPSEnvironment}
- full name: tenpy.networks.mps.MPSEnvironment
- parent module: tenpy.networks.mps
- type: class

Inheritance Diagram

MPSEnvironment

\section*{Methods}
\begin{tabular}{ll}
\hline \begin{tabular}{l} 
MPSEnvironment.__init__(bra, ket[, init_LP, \\
\(\ldots])\).
\end{tabular} & Initialize self. \\
\hline MPSEnvironment.del_LP(i) & Delete stored part strictly to the left of site \(i\). \\
\hline MPSEnvironment.del_RP(i) & Delete storde part scrictly to the right of site \(i\). \\
\hline \begin{tabular}{l} 
MPSEnvironment.expectation_value(ops[,, \\
\(\ldots])\)
\end{tabular} & \begin{tabular}{l} 
Expectation value \(<\) bralops \(\mid\) ket \(>\) of \((n-\) site \()\) opera- \\
tor(s).
\end{tabular} \\
\hline
\end{tabular}

Table 19 - continued from previous page
\begin{tabular}{|c|c|}
\hline MPSEnvironment.full_contraction(i0) & Calculate the overlap by a full contraction of the network. \\
\hline MPSEnvironment.get_LP(i[, store]) & Calculate LP at given site from nearest available one. \\
\hline MPSEnvironment.get_LP_age(i) & Return number of physical sites in the contractions of get_LP(i). \\
\hline MPSEnvironment.get_RP(i[, store]) & Calculate RP at given site from nearest available one. \\
\hline MPSEnvironment.get_RP_age(i) & Return number of physical sites in the contractions of get_RP(i). \\
\hline \multicolumn{2}{|l|}{MPSEnvironment.get_initialization_data()Return data for (re-)initialization.} \\
\hline MPSEnvironment.init_LP(i) & Build initial left part LP. \\
\hline MPSEnvironment.init_RP(i) & Build initial right part
MPS \(/\) RPOEnvironment. for an \\
\hline MPSEnvironment.set_LP(i, LP, age) & Store part to the left of site \(i\). \\
\hline MPSEnvironment.set_RP(i, RP, age) & Store part to the right of site \(i\). \\
\hline MPSEnvironment.test_sanity() & Sanity check, raises ValueErrors, if something is wrong. \\
\hline
\end{tabular}
class tenpy.networks.mps.MPSEnvironment (bra, ket, init_LP=None, init_RP=None, age_LP \(=0\), age_ \(R P=0\) )
Bases: object
Stores partial contractions of \(<b r a|O p| k e t>\) for local operators \(O p\).
The network for a contraction \(<b r a|O p|\) ket \(>\) of a local operator \(O p\), say exemplary at sites \(i, i+l\) looks like:


Of course, we can also calculate the overlap <bralket> by using the special case \(\mathrm{Op}=\) Id.
We use the following label convention (where arrows indicate qconj):


To avoid recalculations of the whole network e.g. in the DMRG sweeps, we store the contractions up to some site index in this class. For bc='finite', ' segment ', the very left and right part LP [ 0 ] and RP [ -1 ] are trivial and don't change, but for \(\mathrm{bc}=\) ' infinite' they are might be updated (by inserting another unit cell to the left/right).

The MPS bra and ket have to be in canonical form. All the environments are constructed without the singular values on the open bond. In other words, we contract left-canonical \(A\) to the left parts \(L P\) and right-canonical \(B\) to the right parts \(R P\). Thus, the special case ket=bra should yield identity matrices for \(L P\) and \(R P\).

\section*{Parameters}
- bra (MPS) - The MPS to project on. Should be given in usual 'ket' form; we call conj() on the matrices directly. Stored in place, without making copies. If necessary to match charges, we call gauge_total_charge().
- ket (MPO I None) - The MPS on which the local operator acts. Stored in place, without making copies. If None, use bra.
- init_LP (None I Array) - Initial very left part LP. If None, build trivial one with init_LP().
- init_RP (None I Array) - Initial very right part RP. If None, build trivial one with init_RP().
- age_LP (int) - The number of physical sites involved into the contraction yielding firstLP.
- age_RP (int) - The number of physical sites involved into the contraction yielding lastRP.

L
Number of physical sites involved into the Environment, i.e. the least common multiple of bra.L and ket.L.

Type int

\section*{bra, ket}

The two MPS for the contraction.
Type MPS

\section*{dtype}

The data type.
Type type

\section*{_finite}

Whether the boundary conditions of the MPS are finite.
Type bool
_LP
Left parts of the environment, len \(L\). LP [i] contains the contraction strictly left of site \(i\) (or None, if we don't have it calculated).

Type list of \(\{\) None I Array \(\}\)
_RP
Right parts of the environment, len \(L\). RP [i] contains the contraction strictly right of site \(i\) (or None, if we don't have it calculated).

Type list of \(\{\) None I Array \(\}\)

\section*{_LP_age}

Used for book-keeping, how large the DMRG system grew: _LP_age [i] stores the number of physical sites invovled into the contraction network which yields self._LP [i].

Type list of int I None

\section*{_RP_age}

Used for book-keeping, how large the DMRG system grew: _RP_age [i] stores the number of physical sites invovled into the contraction network which yields self._RP [i].

Type list of int I None
test_sanity()
Sanity check, raises ValueErrors, if something is wrong.
init_LP (i)
Build initial left part LP.

Parameters i (int) - Build LP left of site \(i\).
Returns init_LP - Identity contractible with the \(v L\) leg of ket.get_B(i), labels 'vR*', 'vR'.
Return type Array
init_RP(i)
Build initial right part RP for an MPS/MPOEnvironment.
Parameters i (int) - Build RP right of site \(i\).
Returns init_RP - Identity contractible with the \(v R\) leg of ket.get_B(i), labels 'vL*', 'vL'.

\section*{Return type Array}
get_LP (i, store=True)
Calculate LP at given site from nearest available one.
The returned LP_i corresponds to the following contraction, where the M's and the N's are in the ' A ' form:


\section*{Parameters}
- i (int) - The returned \(L P\) will contain the contraction strictly left of site \(i\).
- store (bool) - Wheter to store the calculated LP in self (True) or discard them (False).

Returns LP_i - Contraction of everything left of site \(i\), with labels 'vR*', 'vR' for bra, ket.
Return type Array
get_RP (i, store=True)
Calculate RP at given site from nearest available one.
The returned RP_i corresponds to the following contraction, where the M's and the N's are in the 'B' form:


\section*{Parameters}
- i (int) - The returned \(R P\) will contain the contraction strictly right of site \(i\).
- store (b○○l) - Wheter to store the calculated RP in self (True) or discard them (False).
Returns RP_i - Contraction of everything left of site \(i\), with labels 'vL*', 'vL' for bra, ket.
Return type Array
get_LP_age (i)
Return number of physical sites in the contractions of get_LP(i).
Might be None.
get_RP_age (i)
Return number of physical sites in the contractions of get_RP(i).
Might be None.
set_LP (i,LP, age)
Store part to the left of site \(i\).
```

set_RP (i,RP,age)

```

Store part to the right of site \(i\).
del_LP (i)
Delete stored part strictly to the left of site \(i\).
del_RP (i)
Delete storde part scrictly to the right of site \(i\).
```

get_initialization_data()

```

Return data for (re-)initialization.
The returned parameters are collected in a dictionary with the following names.

\section*{Returns}
- init_LP, init_RP(Array) - \(L P\) on the left of site 0 and \(R P\) on the right of site \(L-1\), which can be used as init_LP and init_RP for the initialization of a new environment.
- age_LP, age_RP (int) - The number of physical sites involved into the contraction yielding init_LP and init_RP, respectively.
full_contraction (i0)
Calculate the overlap by a full contraction of the network.
The full contraction of the environments gives the overlap <bra|ket>, taking into account MPS.norm of both bra and ket. For this purpose, this function contracts get_LP (i0+1, store=False) and get_RP(i0, store=False) with appropriate singular values in between.

Parameters i0 (int) - Site index.
expectation_value (ops, sites \(=\) None, axes \(=\) None \()\)
Expectation value <bra|ops | ket> of (n-site) operator(s).
Calculates \(n\)-site expectation values of operators sandwiched between bra and ket. For examples the contraction for a two-site operator on site \(i\) would look like:


Here, the \(B\) are taken from ket, the \(B^{*}\) from bra. The call structure is the same as for MPS. expectation_value().

Warning: In contrast to MPS.expectation_value (), this funciton does not normalize, thus it also takes into account MPS . norm of both bra and ket.

\section*{Parameters}
- ops ((list of) \{ Array | str \}) - The operators, for wich the expectation value should be taken, All operators should all have the same number of legs (namely \(2 n\) ). If less than len (sites) operators are given, we repeat them periodically. Strings (like 'Id', ' Sz ') are translated into single-site operators defined by sites.
- sites (list) - List of site indices. Expectation values are evaluated there. If None (default), the entire chain is taken (clipping for finite b.c.)
- axes (None / (list of str, list of str))- Two lists of each \(n\) leg labels giving the physical legs of the operator used for contraction. The first \(n\) legs are contracted with conjugated \(B\), the second \(n\) legs with the non-conjugated \(B\). None defaults to (['p'], ['p*']) for single site (n=1), or (['p0', 'p1', ... 'p\{n-1\}'], ['p0*', 'p1*', .... 'p\{n-1\}*']) for \(n>1\).
Returns exp_vals - Expectation values, exp_vals[i] = <bra|ops[i]|ket>, where ops[i] acts on site(s) \(j, j+1, \ldots, j+\{n-1\}\) with \(j=s i t e s[i]\).
Return type 1D ndarray

\subsection*{20.2.2 TransferMatrix}
- full name: tenpy.networks.mps.TransferMatrix
- parent module: tenpy.networks.mps
- type: class

\section*{Inheritance Diagram}


\section*{Methods}
\begin{tabular}{ll}
\hline TransferMatrix.__init__(bra, ket[,...]) & Initialize self. \\
\hline TransferMatrix.adjoint() & Return the hermitian conjugate of self \\
\hline TransferMatrix.eigenvectors(*args, & Find (dominant) eigenvector(s) of self using scipy. \\
**kwargs) & sparse. \\
\hline TransferMatrix.initial_guess([diag]) & \begin{tabular}{l} 
Return a diagonal matrix as initial guess for the eigen- \\
\\
\\
vector.
\end{tabular} \\
\hline TransferMatrix.matvec(vec) & Given vec as an npc.Array, apply the transfer matrix. \\
\hline TransferMatrix.to_matrix() & Contract self to a matrix. \\
\hline
\end{tabular}

\section*{Class Attributes and Properties}

\section*{TransferMatrix.acts_on}
class tenpy.networks.mps.TransferMatrix (bra, ket, shift_bra=0, shift_ket=None, transpose \(=\) False, charge_sector \(=0\), form \(={ }^{\prime} B^{\prime}\) )
Bases: tenpy.linalg.sparse.NpcLinearOperator
Transfer matrix of two MPS (bra \& ket).
For an iMPS in the thermodynamic limit, we often need to find the 'dominant \(R P\) ' (and \(L P\) ). This mean nothing else than to take the transfer matrix of the unit cell and find the (right/left) eigenvector with the largest (magnitude) eigenvalue, since it will dominate \((T M)^{n} R P\) (or \(L P(T M)^{n}\) ) in the limit \(n \rightarrow \infty\) - whatever the initial \(R P\) is. This class provides exactly that functionality with eigenvectors ().

Given two MPS, we define the transfer matrix as:
```

| ---M[i]---M[i+1]- ... --M[i+L]---

```


Here the \(M\) denotes the matrices of the bra and \(N\) the ones of the ket, respectively. To view it as a matrix, we combine the left and right indices to pipes:


Note that we keep all M and N as copies.
Deprecated since version 0.6.0: The default for shift_ket was the value of shift_bra, this will be changed to 0 .

\section*{Parameters}
- bra (MPS) - The MPS which is to be (complex) conjugated.
- ket (MPS) - The MPS which is not (complex) conjugated.
- shift_bra (int) - We start the \(N\) of the bra at site shift_bra (i.e. the \(j\) in the above network).
- shift_ket (int / None) - We start the \(M\) of the ket at site shift_ket (i.e. the \(i\) in the above network). None is deprecated, default will be changed to 0 in the future.
- transpose (bool) - Wheter self.matvec acts on RP (False) or LP (True).
- charge_sector (None I charges I 0) - Selects the charge sector of the vector onto which the Linear operator acts. None stands for all sectors, 0 stands for the zero-charge sector. Defaults to 0 , i.e., assumes the dominant eigenvector is in charge sector 0 .
- form ('B' | 'A' | 'C' | 'G' | 'Th' | Noneltuple(float, float)) - In which canonical form we take the \(M\) and \(N\) matrices.

L
Number of physical sites involved in the transfer matrix, i.e. the least common multiple of bra.L and ket.L.
Type int
shift_bra
We start the \(N\) of the bra at site shift_bra.
Type int
shift_ket
We start the \(M\) of the ket at site shift_ket. None defaults to shift_bra.
Type int I None

\section*{transpose}

Wheter self.matvec acts on \(R P\) (True) or \(L P\) (False).
Type bool

\section*{qtotal}

Total charge of the transfer matrix (which is gauged away in matvec).
Type charges

\section*{form}

In which canonical form (all of) the \(M\) and \(N\) matrices are.
Type tuple(float, float) I None
flat_linop
Class lifting matvec () to ndarrays in order to use speigs ().
Type FlatLinearOperator
pipe
Pipe corresponding to '(vL.vL*)' for transpose=False or to '(vR.vR*)' for transpose=True.

Type LegPipe
label_split
['vL', 'vL*'] if tranpose=False or ['vR', 'vR*'] if transpose=True.
_bra_N
Complex conjugated matrices of the bra, transposed for fast matvec.
Type list of npc.Array
_ket_M
The matrices of the ket, transposed for fast matvec.
Type list of npc.Array
matvec (vec)
Given vec as an npc.Array, apply the transfer matrix.

Parameters vec (Array) - Vector to act on with the transfermatrix. If not transposed, vec is the right part \(R P\) of an environment, with legs ' \((v L \cdot v L \star)^{\prime}\) in a pipe or splitted. If transposed, the left part \(L P\) of an environment with legs ' ( \(\mathrm{vR} * . v \mathrm{v})^{\prime}\).
Returns mat_vec - The tranfer matrix acted on vec, in the same form as given.
Return type Array
initial_guess (diag=1.0)
Return a diagonal matrix as initial guess for the eigenvector.
Parameters diag (float / 1D ndarray) - Should be 1. for the identity or some singular values squared.
Returns mat - A 2D array with diag on the diagonal such that matvec () can act on it.
Return type Array
eigenvectors (*args, **kwargs)
Find (dominant) eigenvector(s) of self using scipy. sparse.
For arguments see eigenvectors().
If no charge_sector was selected, we look in all charge sectors. The returned eigenvectors have combined legs ' (vL.vL*)' or (vR*.vR).
```

adjoint()

```

Return the hermitian conjugate of self
If self is hermitian, subclasses can choose to implement this to define the adjoint operator of self.
to_matrix()
Contract self to a matrix.
If self represents an operator with very small shape, e.g. because the MPS bond dimension is very small, an algorithm might choose to contract self to a single tensor.

Returns matrix - Contraction of the represented operator.
Return type Array

\section*{Functions}
build_initial_state(size, states, filling[,...]) Build an "initial state" list.

\subsection*{20.2.3 build_initial_state}
- full name: tenpy.networks.mps.build_initial_state
- parent module: tenpy. networks.mps
- type: function
```

tenpy.networks.mps.build_initial_state (size, states, filling, mode='random', seed=None)

```

Build an "initial state" list.
Uses two iterables ('states' and 'filling') to determine how to fill the state. The two lists should have the same length as every element in 'filling' gives the filling fraction for the corresponding state in 'states'.

\section*{Example}
size \(=6\), states \(=[0,1,2]\), filling \(=[1 . / 3,2 . / 3,0\).\(] n_states =\) size \(*\) filling \(=[2,4,0]==>\) Two sites will get state 0,4 sites will get state 1,0 sites will get state 2 .

Todo: Make more general: it should be possible to specify states as strings.

\section*{Parameters}
- size (int) - length of state
- states (iterable) - Containing the possible local states
- filling (iterable) - Fraction of the total number of sites to get a certain state. If infinite fractions (e.g. 1/3) are needed, one should supply a fraction (1./3.)
- mode (str / None) - State filling pattern. Only 'random' is implemented
- seed (int / None) - Seed for random number generators

\section*{Returns initial_state (list)}

Return type the initial state

\section*{Raises}
- ValueError - If fractonal fillings are incommensurate with system size.
- AssertionError - If the total filling is not equal to 1 , or the length of filling does not equal the length of states.

\section*{Module description}

This module contains a base class for a Matrix Product State (MPS).
An MPS looks roughly like this:
```

| -- B[0] -- B[1] -- B B[2] -- ...

```

We use the following label convention for the \(B\) (where arrows indicate qconj):
```

: vL ->- B ->- vR

```

We store one 3-leg tensor_B[i] with labels 'vL', 'vR', 'p' for each of the \(L\) sites \(0<=i<L\). Additionally, we store \(L+1\) singular value arrays _S[ib] on each bond \(0<=i b<=L\), independent of the boundary conditions. _S [ib] gives the singlur values on the bond i-1, i. However, be aware that e.g. chi returns only the dimensions of the nontrivial_bonds depending on the boundary conditions.

The matrices and singular values always represent a normalized state (i.e. np.linalg.norm(psi._S[ib]) == 1 up to roundoff errors), but (for finite MPS) we keep track of the norm in norm (which is respected by overlap (), ...).

Valid MPS boundary conditions (not to confuse with bc_coupling of tenpy.models.model.CouplingModel) are the following:
\begin{tabular}{|c|c|}
\hline \(b c\) & description \\
\hline \[
\begin{aligned}
& \text { 'fi- } \\
& \text { nite' }
\end{aligned}
\] & Finite MPS, G0 s1 G1 ... \(s\{L-1\} G\{1-1\}\). This is acchieved by using a trivial left and right bond \(\mathrm{s}[0]=\mathrm{s}[-1]=\) np.array ([1.]). \\
\hline 'segment & Generalization of 'finite', describes an MPS embedded in left and right environments. The left environment is described by chi [0] orthonormal states which are weighted by the singular values s[0]. Similar, \(s[L]\) weight some right orthonormal states. You can think of the left and right states to be generated by additional MPS, such that the overall structure is something like ... s L s L [s0 G0 s1 G1 ... \(s\{L-1\} G\{L-1\} s\{L\}] R s R \quad s \quad\). . (where we save the part in the brackets [ . . . ] ). \\
\hline \[
\begin{aligned}
& \text { 'in- } \\
& \text { fi- } \\
& \text { nite' }
\end{aligned}
\] & infinite MPS (iMPS): we save a 'MPS unit cell' [s0 G0 s1 G1 ... s\{L-1\} G\{L-1\}] which is repeated periodically, identifying all indices modulo self.L. In particular, the last bond L is identified with 0 . (The MPS unit cell can differ from a lattice unit cell). bond is identified with the first one. \\
\hline
\end{tabular}

An MPS can be in different 'canonical forms' (see [[schollwoeck2011], [vidal2004]]). To take care of the different canonical forms, algorithms should use functions like get_theta (), get_B () and set_B () instead of accessing them directly, as they return the \(B\) in the desired form (which can be chosen as an argument). The values of the tuples for the form correspond to the exponent of the singular values on the left and right. To keep track of a "mixed" canonical form A A A S B B, we save the tuples for each site of the MPS in MPS. form.
\begin{tabular}{|c|c|c|}
\hline form & tuple & description \\
\hline 'B' & \((0,1)\) & right canonical: _B[i] = -- Gamma [i] -- s[i+1]-- The default form, which algorithms asssume. \\
\hline 'C' & \[
\begin{aligned}
& (0.5, \\
& 0.5)
\end{aligned}
\] & symmetric form:_B[i] = -- s[i]**0.5-- Gamma[i] -- s[i+1]**0.5-- \\
\hline 'A' & \((1,0)\) & left canonical: _B [i] = -- s[i] -- Gamma [i] --. \\
\hline 'G' & \((0,0)\) & Save only _B [i] = -- Gamma [i] --. \\
\hline 'Th & \((1,1)\) & Form of a local wave function theta with singular value on both sides. psi.get_B(i, 'Th') is equivalent to "`psi.get_theta(i, n=1). \\
\hline None & None & General non-canoncial form. Valid form for initialization, but you need to call canonical_form() (or similar) before using algorithms. \\
\hline
\end{tabular}

\section*{20.3 mpo}
- full name: tenpy.networks.mpo
- parent module: tenpy.networks
- type: module

\section*{Classes}

MPO

\begin{tabular}{ll}
\hline MPO(sites, Ws[, bc, IdL, IdR, max_range, \(\ldots\) ]) & \begin{tabular}{l} 
Matrix product operator, finite (MPO) or infinite \\
(iMPO).
\end{tabular} \\
\hline MPOEnvironment(bra, H, ket[, init_LP, \(\ldots\) ]) & \begin{tabular}{l} 
Stores partial contractions of \(<b r a|H| k e t ~>~ f o r ~ a n ~\) \\
MPO \(H\).
\end{tabular} \\
\hline MPOGraph(sites[, bc, max_range]) & \begin{tabular}{l} 
Representation of an MPO by a graph, based on a 'finite \\
state machine'.
\end{tabular} \\
\hline
\end{tabular}

\subsection*{20.3.1 MPOEnvironment}
- full name: tenpy.networks.mpo.MPOEnvironment
- parent module: tenpy. networks.mpo
- type: class

\section*{Inheritance Diagram}


\section*{Methods}
\begin{tabular}{|c|c|}
\hline MPOEnvironment.__init__(bra, H, ket[, ...]) & Initialize self. \\
\hline MPOEnvironment.del_IP(i) & Delete stored part strictly to the left of site \(i\). \\
\hline MPOEnvironment.del_RP(i) & Delete storde part scrictly to the right of site \(i\). \\
\hline MPOEnvironment.expectation_value(ops[, ...]) & (doesn't make sense) \\
\hline MPOEnvironment.full_contraction(i0) & Calculate the energy by a full contraction of the network. \\
\hline MPOEnvironment.get_LP(i[, store]) & Calculate LP at given site from nearest available one (including \(i\) ). \\
\hline MPOEnvironment.get_LP_age(i) & Return number of physical sites in the contractions of get_LP(i). \\
\hline MPOEnvironment.get_RP(i[, store]) & Calculate RP at given site from nearest available one (including \(i\) ). \\
\hline MPOEnvironment.get_RP_age(i) & Return number of physical sites in the contractions of get_RP(i). \\
\hline \multicolumn{2}{|l|}{MPOEnvironment.get_initialization_data()Return data for (re-)initialization.} \\
\hline MPOEnvironment.init_LP(i) & Build initial left part LP. \\
\hline MPOEnvironment.init_RP(i) & Build initial right part RP for an MPS/MPOEnvironment. \\
\hline MPOEnvironment.set_LP(i, LP, age) & Store part to the left of site \(i\). \\
\hline MPOEnvironment.set_RP(i, RP, age) & Store part to the right of site \(i\). \\
\hline MPOEnvironment.test_sanity() & Sanity check, raises ValueErrors, if something is wrong. \\
\hline
\end{tabular}
class tenpy.networks.mpo.MPOEnvironment (bra, \(H\), ket, init_LP=None, init_RP=None, age_LP=0, age_RP=0)
Bases: tenpy.networks.mps.MPSEnvironment
Stores partial contractions of \(<b r a|H|\) ket \(>\) for an MPO \(H\).
The network for a contraction \(<b r a|H| k e t>\) of an MPO \(H\) bewteen two MPS looks like:
\(\square\)
We use the following label convention (where arrows indicate qconj):
\begin{tabular}{|c|c|c|}
\hline & ->- vR & vL ->-. \\
\hline & | & | \\
\hline & LP->- wR & wL ->-RP \\
\hline & | & \\
\hline & <- vR* & vL* -<-. \\
\hline
\end{tabular}

To avoid recalculations of the whole network e.g. in the DMRG sweeps, we store the contractions up to some site index in this class. For \(b c=\) 'finite', 'segment ', the very left and right part LP [0] and RP [-1] are trivial and don't change in the DMRG algorithm, but for iDMRG ( \(\mathrm{bc}=\) 'infinite') they are also updated (by inserting another unit cell to the left/right).

The MPS bra and ket have to be in canonical form. All the environments are constructed without the singular values on the open bond. In other words, we contract left-canonical \(A\) to the left parts \(L P\) and right-canonical \(B\) to the right parts \(R P\).

\section*{Parameters}
- bra (MPS) - The MPS to project on. Should be given in usual 'ket' form; we call conj() on the matrices directly.
- H (MPO) - The MPO sandwiched between bra and ket. Should have 'IdL' and 'IdR' set on the first and last bond.
- ket (MPS) - The MPS on which \(H\) acts. May be identical with bra.
- init_LP (None I Array) - Initial very left part LP. If None, build trivial one with :meth init_LP`.
- init_RP (None I Array) - Initial very right part RP. If None, build trivial one with init_RP().
- age_LP (int) - The number of physical sites involved into the contraction yielding firstLP.
- age_RP (int) - The number of physical sites involved into the contraction yielding last \(R P\).

H
The MPO sandwiched between bra and ket.
Type MPO
test_sanity()
Sanity check, raises ValueErrors, if something is wrong.
init_LP (i)
Build initial left part LP.
Parameters i (int)-Build LP left of site \(i\).
Returns init_LP - Identity contractible with the \(v L\) leg of . ket.get_B (i), multiplied with a unit vector nonzero in \(\mathrm{H} . \mathrm{IdL}[\mathrm{i}]\), with labels 'vR*', 'wR', 'vR'.

\section*{Return type Array}
```

init_RP(i)

```

Build initial right part RP for an MPS/MPOEnvironment.
Parameters i (int) - Build RP right of site \(i\).
Returns init_RP - Identity contractible with the \(v R\) leg of self.get_B (i), multiplied with a unit vector nonzero in \(\mathrm{H} . \mathrm{IdR}[\mathrm{i}]\), with labels 'vL*', 'wL', 'vL'.

\section*{Return type Array}
get_LP (i, store=True)
Calculate LP at given site from nearest available one (including \(i\) ).
The returned LP_i corresponds to the following contraction, where the M's and the N's are in the 'A' form:
\(\square\)

\section*{Parameters}
- i (int) - The returned \(L P\) will contain the contraction strictly left of site \(i\).
- store (bool) - Wheter to store the calculated LP in self (True) or discard them (False).

Returns LP_i - Contraction of everything left of site \(i\), with labels 'vR*', 'wR', 'vR' for bra, \(H\), ket.

\section*{Return type Array}
get_RP (i, store=True)
Calculate RP at given site from nearest available one (including \(i\) ).
The returned RP_i corresponds to the following contraction, where the M's and the N's are in the 'B' form:
```

'vL' ->---M[i+1]-- ... --M[L-1]----.
'WL' ->---W[i+1]-- ... --W[L-1]-----RP[-1]
'vL*' -<---N[i+1]*- ... --N N[L-1]*----

```

\section*{Parameters}
- \(\mathbf{i}(i n t)\) - The returned \(R P\) will contain the contraction strictly rigth of site \(i\).
- store (bool) - Wheter to store the calculated \(R P\) in self (True) or discard them (False).

Returns RP_i - Contraction of everything right of site \(i\), with labels 'vL*', 'wL', 'vL' for \(b r a, H\), ket.

\section*{Return type Array}
full_contraction (i0)
Calculate the energy by a full contraction of the network.
The full contraction of the environments gives the value <bra|H|ket> / ( \(\operatorname{norm}(|\operatorname{bra}\rangle) * \operatorname{norm}(\mid k e t>))\), i.e. if bra is ket and normalized, the total energy. For this purpose, this function contracts get_LP (i0+1, store=False) and get_RP(i0, store=False).

Parameters i0 (int) - Site index.
expectation_value (ops, sites=None, axes=None)
(doesn't make sense)
del_LP (i)
Delete stored part strictly to the left of site \(i\).
del_RP (i)
Delete storde part scrictly to the right of site \(i\).
get_LP_age ( \(i\) )
Return number of physical sites in the contractions of get_LP(i).
Might be None.
get_RP_age (i)
Return number of physical sites in the contractions of get_RP(i).
Might be None.
get_initialization_data()
Return data for (re-)initialization.
The returned parameters are collected in a dictionary with the following names.

\section*{Returns}
- init_LP, init_RP (Array) - \(L P\) on the left of site 0 and \(R P\) on the right of site \(L-1\), which can be used as init_LP and init_ \(R P\) for the initialization of a new environment.
- age_LP, age_RP (int) - The number of physical sites involved into the contraction yielding init_LP and init_RP, respectively.
set_LP ( \(i, L P\), age \()\)
Store part to the left of site \(i\).
set_RP (i,RP, age)
Store part to the right of site \(i\).

\subsection*{20.3.2 MPOGraph}
- full name: tenpy.networks.mpo.MPOGraph
- parent module: tenpy.networks.mpo
- type: class

\section*{Inheritance Diagram}

\section*{MPOGraph}

\section*{Methods}
\begin{tabular}{ll}
\hline MPOGraph.__init__(sites[, bc, max_range]) & Initialize self. \\
\hline MPOGraph.add(i, keyL, keyR, opname, strength) & Insert an edge into the graph. \\
\hline MPOGraph.add_missing_IdL_IdR([insert_all_id]) Add missing identity ('Id') edges connecting \\
& 'IdL'->'IdL ' and ' 'IdR '->' IdR'.
\end{tabular}

\section*{Class Attributes and Properties}
\begin{tabular}{ll}
\hline MPOGraph.L & \begin{tabular}{l} 
Number of physical sites; for infinite boundaries the \\
length of the unit cell.
\end{tabular} \\
\hline
\end{tabular}
class tenpy.networks.mpo.MPOGraph (sites, bc='finite', max_range=None)
Bases: object
Representation of an MPO by a graph, based on a 'finite state machine'.
This representation is used for building H_MPO from the interactions. The idea is to view the MPO as a kind of 'finite state machine'. The states or keys of this finite state machine life on the MPO bonds between the Ws. They label the indices of the virtul bonds of the MPOs, i.e., the indices on legs wL and wR. They can be anything hash-able like a str, int or a tuple of them.

The edges of the graph are the entries \(W[k e y L, \quad k e y R]\), which itself are onsite operators on the local Hilbert space. The indices keyL and keyR correspond to the legs 'wL', 'wR' of the MPO. The entry \(w[k e y L\), keyR] connects the state keyL on bond (i-1, i) with the state keyR on bond (i, i+1).

The keys 'IdR' (for 'idenity left') and 'IdR' (for 'identity right') are reserved to represent only 'Id' (=identity) operators to the left and right of the bond, respectively.

Todo: might be useful to add a "cleanup" function which removes operators cancelling each other and/or unused states. Or better use a 'compress' of the MPO?

\section*{Parameters}
- sites (list of Site) - Local sites of the Hilbert space.
- bc(\{'finite', 'infinite'\})- MPO boundary conditions.
- max_range (int / np.inf / None) - Maximum range of hopping/interactions (in unit of sites) of the MPO. None for unknown.
sites
Defines the local Hilbert space for each site.
Type list of Site
chinfo
The nature of the charge.
Type ChargeInfo
bc
MPO boundary conditions.
Type \{'finite’, 'infinite'\}

\section*{max_range}

Maximum range of hopping/interactions (in unit of sites) of the MPO. None for unknown.
Type int I np.inf I None
states
states [i] gives the possible keys at the virtual bond (i-1, i) of the MPO. \(L+1\) enries.
Type list of set of keys
graph
For each site \(i\) a dictionary \{keyL: \{keyR: [(opname, strength)]\}\} with keyL in states[i] and keyR in states[i+1].

Type list of dict of dict of list of tuples
_grid_legs
The charges for the MPO
Type None I list of LegCharge
classmethod from_terms (terms, sites, bc, insert_all_id=True)
Initialize an MPOGraph from OnsiteTerms and CouplingTerms.

\section*{Parameters}
- terms (iterable of tenpy.networks.terms.*Terms classes) - Entries can be OnsiteTerms, CouplingTerms, MultiCouplingTerms or ExponentialCouplingTerms. All the entries get added to the new MPOGraph.
- sites (list of Site) - Local sites of the Hilbert space.
- bc('finite' | 'infinite')-MPO boundary conditions.
- insert_all_id (bool) - Whether to insert identities such that \(I d L\) and \(I d R\) are defined on each bond. See add_missing_IdL_IdR().

Returns graph - Initialized with the given terms.
Return type MPOGraph
See also:
from_term_list equivalent for representation by TermList.
classmethod from_term_list (term_list, sites, \(b c\), insert_all_id=True)
Initialize from a list of operator terms and prefactors.

\section*{Parameters}
- term_list (TermList) - Terms to be added to the MPOGraph.
- sites (list of Site) - Local sites of the Hilbert space.
- bc('finite' | 'infinite')-MPO boundary conditions.
- insert_all_id (bool) - Whether to insert identities such that \(I d L\) and \(I d R\) are defined on each bond. See add_missing_IdL_IdR().

Returns graph - Initialized with the given terms.
Return type MPOGraph
See also:
from_terms equivalent for other representation of terms.
test_sanity()
Sanity check, raises ValueErrors, if something is wrong.
property L
Number of physical sites; for infinite boundaries the length of the unit cell.
add (i,keyL, keyR, opname, strength, check_op=True, skip_existing=False)
Insert an edge into the graph.

\section*{Parameters}
- i (int) - Site index at which the edge of the graph is to be inserted.
- keyl (hashable) - The state at bond (i-1, i) to connect from.
- keyR (hashable) - The state at bond (i, \(\mathrm{i}+1\) ) to connect to.
- opname (str) - Name of the operator.
- strength (str) - Prefactor of the operator to be inserted.
- check_op (bool) - Whether to check that 'opname' exists on the given site.
- skip_existing (bool) - If True, skip adding the graph node if it exists (with same keys and opname).
add_string ( \(i, j\), key, opname \(=\) 'Id', check_op=True, skip_existing=True)
Insert a bunch of edges for an 'operator string' into the graph.
Terms like \(S_{i}^{z} S_{j}^{z}\) actually stand for \(S_{i}^{z} \otimes \prod_{i<k<j} \nVdash_{k} \otimes S_{j}^{z}\). This function adds the \(\nVdash\) terms to the graph.

\section*{Parameters}
- \(\mathbf{i}(i n t)\) - An edge is inserted on all bonds between \(i\) and \(j, i<j . j\) can be larger than \(L\), in which case the operators are supposed to act on different MPS unit cells.
- \(\mathbf{j}(i n t)\) - An edge is inserted on all bonds between \(i\) and \(j, i<j\). \(j\) can be larger than \(L\), in which case the operators are supposed to act on different MPS unit cells.
- key (hashable) - The state at bond (i-1, i) to connect from and on bond ( \(\mathrm{j}-1, \mathrm{j}\) ) to connect to. Also used for the intermediate states. No operator is inserted on a site \(i<k<\) \(j\) if has_edge (k, key, key).
- opname (str) - Name of the operator to be used for the string. Useful for the JordanWigner transformation to fermions.
- skip_existing (bool) - Whether existing graph nodes should be skipped.

Returns label_j - The key on the left of site \(\mathbf{j}\) to connect to. Usually the same as the parameter key, except if \(j-i>s e l f . L\), in which case we use the additional labels (key, 1), (key, 2),... to generate couplings over multiple unit cells.

Return type hashable
add_missing_IdL_IdR(insert_all_id=True)
Add missing identity ('Id') edges connecting 'IdL'->'IdL' and \({ }^{\prime}\) 'IdR'->'IdR'.
This function should be called after all other operators have been inserted.
Parameters insert_all_id (bool) - If True, insert 'Id' edges on all bonds. If False and boundary conditions are finite, only insert 'IdL'->'IdL' to the left of the rightmost existing 'IdL' and 'IdR'->'IdR' to the right of the leftmost existing 'IdR'. The latter avoid "dead ends" in the MPO, but some functions (like make_WI) expect 'IdL'/'IdR' to exist on all bonds.
has_edge ( \(i\), keyL, keyR)
True if there is an edge from keyL on bond (i-1, i\()\) to keyR on bond ( \(\mathrm{i}, \mathrm{i}+1\) ).
build_MPO (Ws_qtotal=None)
Build the MPO represented by the graph (self).
Parameters Ws_qtotal (None | (list of) charges)-The qtotal for each of the Ws to be generated, default (None) means 0 charge. A single qtotal holds for each site.

Returns mpo - the MPO which self represents.

Return type MPO

\section*{Functions}
\begin{tabular}{ll}
\hline grid_insert_ops(site, grid) & \begin{tabular}{l} 
Replaces entries representing operators in a grid of \\
W[i] with npc.Arrays.
\end{tabular} \\
\hline make_W_II(t, A, B, C, D) & W_II approx to \(\exp (\mathrm{t} H)\) from MPO parts (A, B, C, D). \\
\hline
\end{tabular}

\subsection*{20.3.3 grid_insert_ops}
- full name: tenpy.networks.mpo.grid_insert_ops
- parent module: tenpy. networks.mpo
- type: function
tenpy.networks.mpo.grid_insert_ops (site, grid)
Replaces entries representing operators in a grid of W [i] with npc.Arrays.

\section*{Parameters}
- site (site) - The site on which the grid acts.
- grid (list of list of entries) - Represents a single matrix \(W\) of an MPO, i.e. the lists correspond to the legs 'vL', 'vR', and entries to onsite operators acting on the given site. entries may be None, Array, a single string or of the form [('opname', strength), . . . ], where 'opname ' labels an operator in the site.

Returns grid - Copy of grid with entries [('opname', strength), ...] replaced by sum([strength*site.get_op('opname') for opname, strength in entry]) and entries 'opname' replaced by site.get_op('opname').
Return type list of list of \{None I Array\}

\subsection*{20.3.4 make_W_II}
- full name: tenpy.networks.mpo.make_W_II
- parent module: tenpy.networks.mpo
- type: function
tenpy.networks.mpo.make_W_II ( \(t, A, B, C, D\) )
W_II approx to \(\exp (\mathrm{t} \mathrm{H})\) from MPO parts (A, B, C, D).
Get the W_II approximation of [[zaletel2015]].
In the paper, we have two formal parameter "phi_\{r/c\}" which satisfies \(\phi_{r}^{2}=p h i_{c}^{2}=0\). To implement this, we temporarily extend the virtual Hilbert space with two hard-core bosons "br, bl". The components of Eqn (11) can be computed for each index of the virtual row/column independently The matrix exponential is done in the hard-core extended Hilbert space

\section*{Parameters}
- \(\boldsymbol{t}(f l o a t)\) - The time step per application of the propagator. Should be imaginary for real time evolution!
- A (numpy.ndarray) - Blocks of the MPO tensor to be exponentiated, as defined in [[zaletel2015]]. Legs 'wL', 'wR', 'p', 'p*'; legs projected to a single IdL/IdR can be dropped.
- B (numpy. ndarray) - Blocks of the MPO tensor to be exponentiated, as defined in [[zaletel2015]]. Legs 'wL', 'wR', 'p', 'p*'; legs projected to a single IdL/IdR can be dropped.
- C (numpy.ndarray) - Blocks of the MPO tensor to be exponentiated, as defined in [[zaletel2015]]. Legs 'wL', 'wR', 'p', 'p*'; legs projected to a single IdL/IdR can be dropped.
- D (numpy.ndarray) - Blocks of the MPO tensor to be exponentiated, as defined in [[zaletel2015]]. Legs 'wL', 'wR', 'p', ' p ''; legs projected to a single IdL/IdR can be dropped.

\section*{Module description}

Matrix product operator (MPO).
An MPO is the generalization of an MPS to operators. Graphically:


So each 'matrix' has two physical legs p, p* instead of just one, i.e. the entries of the 'matrices' are local operators. Valid boundary conditions of an MPO are the same as for an MPS (i.e. 'finite' | 'segment' | 'infinite'). (In general, you can view the MPO as an MPS with larger physical space and bring it into canoncial form. However, unlike for an MPS, this doesn't simplify calculations. Thus, an MPO has no form.)
We use the following label convention for the \(W\) (where arrows indicate qconj):


If an MPO describes a sum of local terms (e.g. most Hamiltonians), some bond indices correspond to 'only identities to the left/right'. We store these indices in \(I d L\) and \(I d R\) (if there are such indices).

Similar as for the MPS, a bond index \(i\) is left of site \(i\), i.e. between sites i-1 and \(i\).

\section*{20.4 terms}
- full name: tenpy.networks.terms
- parent module: tenpy.networks
- type: module

\section*{Classes}

\begin{tabular}{ll}
\hline CouplingTerms(L) & \begin{tabular}{l} 
Operator names, site indices and strengths representing \\
two-site coupling terms.
\end{tabular} \\
\hline ExponentiallyDecayingTerms(L) & \begin{tabular}{l} 
Represent a sum of exponentially decaying (long-range) \\
couplings.
\end{tabular} \\
\hline MultiCouplingTerms(L) & \begin{tabular}{l} 
Operator names, site indices and strengths representing \\
general \(M\)-site coupling terms.
\end{tabular} \\
\hline OnsiteTerms(L) & \begin{tabular}{l} 
Operator names, site indices and strengths representing \\
onsite terms.
\end{tabular} \\
\hline TermList(terms[, strength]) & \begin{tabular}{l} 
A list of terms (=operator names and sites they act on) \\
and associated strengths.
\end{tabular} \\
\hline
\end{tabular}

\subsection*{20.4.1 CouplingTerms}
- full name: tenpy.networks.terms.CouplingTerms
- parent module: tenpy.networks.terms
- type: class

\section*{Inheritance Diagram}


\section*{Methods}
\begin{tabular}{ll}
\hline CouplingTerms.__init__(L) & Initialize self. \\
\hline \begin{tabular}{ll} 
CouplingTerms.add_coupling_term(strength, & Add a two-site coupling term on given MPS sites. \\
i,...)
\end{tabular} & \\
\hline CouplingTerms.add_to_graph(graph) & Add terms from coupling_terms to an MPOGraph. \\
\hline \begin{tabular}{l} 
CouplingTerms.coupling_term_handle_JW(.. Helping function to \\
\(\ldots])\).
\end{tabular} & add_coupling_term ().
\end{tabular}
class tenpy.networks.terms.CouplingTerms \((L)\)
Bases: tenpy.tools.hdf5_io.Hdf5Exportable
Operator names, site indices and strengths representing two-site coupling terms.
Parameters L (int) - Number of sites.
L
Number of sites.
Type int
coupling_terms
Filled by add_coupling_term(). Nested dictionaries of the form \(\left\{i:\left\{\left(' o p n a m e \_i '\right.\right.\right.\), 'opname_string'): \{j: \{'opname_j': strength\}\}\}\}. Note that always i \(<j\), but entries with \(j>=L\) are allowed for bc_MPS == 'infinite', in which case they indicate couplings between different iMPS unit cells.

Type dict of dict
max_range ()
Determine the maximal range in coupling_terms.
Returns max_range - The maximum of \(j\) - i for the \(i, j\) occuring in a term of coupling_terms.
Return type int
add_coupling_term (strength, i, j, op_i, op_j, op_string='Id')
Add a two-site coupling term on given MPS sites.

\section*{Parameters}
- strength (float) - The strength of the coupling term.
- i (int) - The MPS indices of the two sites on which the operator acts. We require 0 <= i < N_sites and i < j, i.e., op_i acts "left" of op_j. If j >= N_sites, it indicates couplings between unit cells of an infinite MPS.
- \(\mathbf{j}\) (int) - The MPS indices of the two sites on which the operator acts. We require 0 <= i < N_sites and i < j, i.e., op_i acts "left" of \(o p_{-} j\). If \(\mathrm{j}>=\mathrm{N} \_\)sites, it indicates couplings between unit cells of an infinite MPS.
- op1 (str) - Names of the involved operators.
- op2 (str) - Names of the involved operators.
- op_string (str) - The operator to be inserted between \(i\) and \(j\).
coupling_term_handle_JW (strength, term, sites, op_string=None)
Helping function to call before add_coupling_term ().

\section*{Parameters}
- strength (float) - The strength of the coupling term.
- term ([(str, int), (str, int)]) - List of two tuples [(op_i, i), (op_j, j)] where \(i\) is the MPS index of the site the operator named op_i acts on; we require \(i<j\).
- sites (list of Site) - Defines the local Hilbert space for each site. Used to check whether the operators need Jordan-Wigner strings.
- op_string (None | str) - Operator name to be used as operator string between the operators, or None if the Jordan Wigner string should be figured out.

Warning: None figures out for each segment between the operators, whether a Jordan-Wigner string is needed. This is different from a plain ' JW ', which just applies a string on each segment!

Returns Arguments for MultiCouplingTerms.add_multi_coupling_term() such that the added term corresponds to the parameters of this function.
Return type strength, \(\mathrm{i}, \mathrm{j}\), op_i, op_j, op_string
plot_coupling_terms (ax, lat, style_map='default', common_style=\{'linestyle': '--'\}, text=None, text_pos=0.4)
"Plot coupling terms into a given lattice.
This function plots the coupling_terms

\section*{Parameters}
- ax (matplotlib.axes.Axes) - The axes on which we should plot.
- lat (Lattice) - The lattice for plotting the couplings, most probably the M. lat of the corresponding model M , see lat.
- style_map (function | None) - Function which get's called with arguments i, \(j\), op_i, op_string, op_j, strength for each two-site coupling and should return a keyword-dictionary with the desired plot-style for this coupling. By default (None), the linewidth is given by the absolute value of strength, and the linecolor depends on the phase of strength (using the \(h s v\) colormap).
- common_style (dict) - Common style, which overwrites values of the dictionary returned by style_map. A ' label' is only used for the first plotted line.
- text (format_string | None) - If not None, we add text labeling the couplings in the plot. Available keywords are i, j, op_i, op_string, op_j, strength as well as strength_abs, strength_angle, strength_real.
- text_pos (float) - Specify where to put the text on the line between \(i(0.0)\) and \(j\) (1.0), e.g. 0.5 is exactly in the middle between \(i\) and \(j\).

\section*{See also:}
tenpy.models.lattice.Lattice.plot_sites plot the sites of the lattice.
add_to_graph (graph)
Add terms from coupling_terms to an MPOGraph.
Parameters graph (MPOGraph) - The graph into which the terms from coupling_terms should be added.

\section*{to_nn_bond_Arrays (sites)}

Convert the coupling_terms into Arrays on nearest neighbor bonds.
Parameters sites (list of Site) - Defines the local Hilbert space for each site. Used to translate the operator names into Array.
Returns H_bond - The coupling_terms rewritten as sum_i H_bond[i] for MPS indices i. H_bond[i] acts on sites (i-1, i), None represents 0 . Legs of each H_bond[i] are ['p0', 'p0*', 'p1', 'p1*'].
Return type list of \{Array INone\}
remove_zeros (tol_zero=le-15)
Remove entries close to 0 from coupling_terms.
Parameters tol_zero (float)-Entries in coupling_terms with strength <tol_zero are considered to be zero and removed.
```

to_TermList()

```

Convert onsite_terms into a TermList.
Returns term_list - Representation of the terms as a list of terms.
Return type TermList
classmethod from_hdf5 (hdf5_loader, h5gr, subpath)
Load instance from a HDF5 file.
This method reconstructs a class instance from the data saved with save_hdf5().

\section*{Parameters}
- hdf5_loader (Hdf5Loader) - Instance of the loading engine.
- h5gr (Group) - HDF5 group which is represent the object to be constructed.
- subpath (str) - The name of \(h 5 g r\) with a '/' in the end.

Returns obj - Newly generated class instance containing the required data.
Return type cls
save_hdf5 (hdf5_saver, h5gr, subpath)
Export self into a HDF5 file.
This method saves all the data it needs to reconstruct self with from_hdf5 ().
This implementation saves the content of \(\qquad\) dict \(\qquad\) with save_dict_content (), storing the format under the attribute 'format'.

\section*{Parameters}
- hdf5_saver (Hdf5Saver) - Instance of the saving engine.
- h5gr (: class`Group`) - HDF5 group which is supposed to represent self.
- subpath (str) - The name of \(h 5 g r\) with a ' /' in the end.

\subsection*{20.4.2 ExponentiallyDecayingTerms}
- full name: tenpy.networks.terms.ExponentiallyDecayingTerms
- parent module: tenpy.networks.terms
- type: class

\section*{Inheritance Diagram}


\section*{Methods}
\begin{tabular}{|c|c|}
\hline ExponentiallyDecayingTerms.
\(\qquad\) init \(\qquad\) (L) & Initialize self. \\
\hline \begin{tabular}{l}
ExponentiallyDecayingTerms. \\
add_exponentially_decaying_coupling(...)
\end{tabular} & Add an exponentially decaying long-range coupling. \\
\hline \begin{tabular}{l}
ExponentiallyDecayingTerms. \\
add_to_graph(graph)
\end{tabular} & Add terms from onsite_terms to an MPOGraph. \\
\hline ExponentiallyDecayingTerms. from_hdf5(...) & Load instance from a HDF5 file. \\
\hline ExponentiallyDecayingTerms. max_range() & Maximum range of the couplings. \\
\hline ExponentiallyDecayingTerms. save_hdf5(...) & Export self into a HDF5 file. \\
\hline ```
ExponentiallyDecayingTerms.
to_TermList([...])
``` & Convert self into a TermList. \\
\hline
\end{tabular}
class tenpy.networks.terms.ExponentiallyDecayingTerms ( \(L\) )
Bases: tenpy.tools.hdf5_io.Hdf5Exportable
Represent a sum of exponentially decaying (long-range) couplings.
MPOs can represent translation invariant, exponentially decaying long-range terms of the following form with a single extra index of the virtual bonds:
\[
\operatorname{sum}_{i \neq j} l a m b d a a^{|i-j|} A_{i} B_{j}
\]

For 2D cylinders (or ladders), we need a slight generalization of this, where the operators act only on a subset of the sites in each unit cell, given by a 1D array subsites:
\[
\text { strengthsum }_{i<j} l a m b d a^{|i-j|} A_{\text {subsites }[i]} B_{\text {subsites }[j]}
\]

Note that we still have \(|i-j|\), such that this will give uniformly decaying interactions, independent of the way the MPS winds through the 2D lattice, as long as subsites is sorted. An easy example would be a ladder, where we want the long-range interactions on the first rung only, subsites = lat.mps_idx_fix_u(u=0), see mps_idx_fix_u().

Parameters L (int) - Number of sites.
L
Number of sites.
Type int
exp_decaying_terms
Each tuple (strength, opname_i, opname_j, lambda, subsites, opname_string) represents one of the terms as described above; see add_exponentially_decaying_coupling() for more details.

Type list of tuples
add_exponentially_decaying_coupling (strength, lambda_, op_i, op_j, subsites=None, \(o p \_\)string \(=\)' \(\mid d^{\prime}\) )
Add an exponentially decaying long-range coupling.
\[
\text { strengthsum }_{i<j} \text { lambda }^{|i-j|} A_{\text {subsites }[i]} B_{\text {subsites }[j]}
\]

Where the operator \(A\) is given by op_i, and \(B\) is given by op_j. Note that the sum over \(\mathrm{i}, \mathrm{j}\) is long-range, for infinite systems beyond the MPS unit cell.

\section*{Parameters}
- strength (float) - Overall prefactor.
- lambda (float) - Decay-rate
- op_i (string) - Names for the operators.
- op_j (string) - Names for the operators.
- subsites (None / 1D array) - Selects a subset of sites within the MPS unit cell on which the operators act. Needs to be sorted. None selects all sites.
- op_string (string) - The operator to be inserted between \(A\) and \(B\); for Fermions this should be "JW".
add_to_graph (graph, key='exp-decay')
Add terms from onsite_terms to an MPOGraph.

\section*{Parameters}
- graph (MPOGraph) - The graph into which the terms from exp_decaying_terms should be added.
- key (str) - Key to distinguish from other states in the MPOGraph. We find integers key_nr and use (key_nr, key) as state for the different entries in exp_decaying_terms.

\section*{to_TermList (cutoff \(=0.01, b c=\) 'finite')}

Convert self into a TermList.

\section*{Parameters}
- cutoff (float) - Drop terms where the overall prefactor is smaller then cutoff.
- bc ("finite" / "infinite") - Boundary conditions to be used.

Returns term_list - Representation of the terms as a list of terms. For "infinite" bc, only terms starting in the first MPS unit cell are included.

\section*{Return type TermList}
max_range ()
Maximum range of the couplings. In this case np.inf.
classmethod from_hdf5 (hdf5_loader, h5gr, subpath)
Load instance from a HDF5 file.
This method reconstructs a class instance from the data saved with save_hdf5 ().

\section*{Parameters}
- hdf5_loader (Hdf5Loader) - Instance of the loading engine.
- h5gr (Group) - HDF5 group which is represent the object to be constructed.
- subpath (str) - The name of \(h 5 g r\) with a '/' in the end.

Returns obj - Newly generated class instance containing the required data.
Return type cls
save_hdf5 (hdf5_saver, h5gr, subpath)
Export self into a HDF5 file.
This method saves all the data it needs to reconstruct self with from_hdf5 ().

This implementation saves the content of \(\qquad\) _dict \(\qquad\) with save_dict_content (), storing the format under the attribute 'format'.

\section*{Parameters}
- hdf5_saver (Hdf5Saver) - Instance of the saving engine.
- h5gr (: class `Group`) - HDF5 group which is supposed to represent self.
- subpath (str) - The name of \(h 5 g r\) with a ' /' in the end.

\subsection*{20.4.3 MultiCouplingTerms}
- full name: tenpy.networks.terms.MultiCouplingTerms
- parent module: tenpy.networks.terms
- type: class

\section*{Inheritance Diagram}


\section*{Methods}
\begin{tabular}{|c|c|}
\hline MultiCouplingTerms.__init__(L) & Initialize self. \\
\hline MultiCouplingTerms. add_coupling_term(...[,...]) & Add a two-site coupling term on given MPS sites. \\
\hline \begin{tabular}{l}
MultiCouplingTerms. \\
add_multi_coupling_term(...)
\end{tabular} & Add a multi-site coupling term. \\
\hline MultiCouplingTerms.add_to_graph(graph[,
\[
\ldots \text { i, ...]) }
\] & Add terms from coupling_terms to an MPOGraph. \\
\hline MultiCouplingTerms. coupling_term_handle_JW(...) & Helping function to call before
add_coupling_term(). \\
\hline MultiCouplingTerms.from_hdf5(hdf5_loader, ...) & Load instance from a HDF5 file. \\
\hline
\end{tabular}

Table 31 - continued from previous page
\begin{tabular}{ll}
\hline MultiCouplingTerms.max_range() & Determine the maximal range in coupling_terms. \\
\hline MultiCouplingTerms. & Helping function to call before \\
multi_coupling_term_handle_JW(...) & add_multi_coupling_term(). \\
\hline \begin{tabular}{ll} 
MultiCouplingTerms. \\
plot_coupling_terms(ax, lat)
\end{tabular} & "Plot coupling terms into a given lattice. \\
\hline MultiCouplingTerms. & \\
remove_zeros([tol_zero,_d0]) & Remove entries close to 0 from coupling_terms. \\
\hline MultiCouplingTerms.save_hdf5(hdf5_saver, & Export self into a HDF5 file. \\
\(\ldots\).. & \\
\hline MultiCouplingTerms.to_TermList() & Convert onsite_terms into a TermList. \\
\hline MultiCouplingTerms. & Convert the coupling_terms into Arrays on nearest \\
to_nn_bond_Arrays(sites) & neighbor bonds. \\
\hline
\end{tabular}
class tenpy.networks.terms.MultiCouplingTerms \((L)\)
Bases: tenpy.networks.terms.CouplingTerms
Operator names, site indices and strengths representing general \(M\)-site coupling terms.
Generalizes the coupling_terms of CouplingTerms to \(M\)-site couplings. The structure of the nested dictionary coupling_terms is similar, but we allow an arbitrary recursion depth of the dictionary.

Parameters L (int) - Number of sites.
L
Number of sites.
Type int

\section*{coupling_terms}

Nested dictionaries of the following form:
```

{i: {('opname_i', 'opname_string_ij'):
{j: {('opname_j', 'opname_string_jk'):
{k: {('opname_k', 'opname_string_kl'):
{l: {'opname_l':
strength
} }
} }
} }
} }

```

For a M-site coupling, this involves a nesting depth of \(2 \star\) M dictionaries. Note that always \(i<j<k\) \(<\ldots<l\), but entries with \(j, k, l>=L\) are allowed for the case of bc_MPS == 'infinite', when they indicate couplings between different iMPS unit cells.

Type dict of dict
add_multi_coupling_term (strength, ijkl,ops_ijkl,op_string='Id')
Add a multi-site coupling term.

\section*{Parameters}
- strength (float) - The strength of the coupling term.
- ijkl (list of int) - The MPS indices of the sites on which the operators acts. With \(i, j, k, \ldots=i j k l\), we require that they are ordered ascending, \(i<j<k<\ldots\) and
that \(0<=i<N \_s i t e s\). Inidces \(>=\) N_sites indicate couplings between different unit cells of an infinite MPS.
- ops_ijkl (list of str) - Names of the involved operators on sites \(i, j, k, \ldots\).
- op_string ((list of) str) - Names of the operator to be inserted between the operators, e.g., op_string[0] is inserted between \(i\) and \(j\). A single name holds for all inbetween segments.
multi_coupling_term_handle_JW (strength, term, sites, op_string=None)
Helping function to call before add_multi_coupling_term().
Handle/figure out Jordan-Wigner strings if needed.

\section*{Parameters}
- strength (float) - The strength of the term.
- term(list of (str, int))-List of tuples (op_i, i) where \(i\) is the MPS index of the site the operator named op_i acts on. We require the operators to be sorted (strictly ascending) by sites. If necessary, call order_combine_term() beforehand.
- sites (list of Site) - Defines the local Hilbert space for each site. Used to check whether the operators need Jordan-Wigner strings.
- op_string (None / str) - Operator name to be used as operator string between the operators, or None if the Jordan Wigner string should be figured out.

Warning: None figures out for each segment between the operators, whether a Jordan-Wigner string is needed. This is different from a plain ' JW ', which just applies a string on each segment!

Returns Arguments for MultiCouplingTerms.add_multi_coupling_term() such that the added term corresponds to the parameters of this function.

Return type strength, ijkl, ops_ijkl, op_string
max_range ()
Determine the maximal range in coupling_terms.
Returns max_range - The maximum of \(j-i\) for the \(i, j\) occuring in a term of coupling_terms.
Return type int
add_to_graph (graph,_i=None,_dl=None, _label_left=None)
Add terms from coupling_terms to an MPOGraph.

\section*{Parameters}
- graph (MPOGraph) - The graph into which the terms from coupling_terms should be added.
- _i (None) - Should not be given; only needed for recursion.
- _d1 (None) - Should not be given; only needed for recursion.
- _label_left (None) - Should not be given; only needed for recursion.
remove_zeros (tol_zero \(=1 e-15\),_d \(d 0=\) None \()\)
Remove entries close to 0 from coupling_terms.

\section*{Parameters}
- tol_zero (float) - Entries in coupling_terms with strength < tol_zero are considered to be zero and removed.
- _d0 (None) - Should not be given; only needed for recursion.
to_TermList ()
Convert onsite_terms into a TermList.
Returns term_list - Representation of the terms as a list of terms.
Return type TermList
add_coupling_term (strength, \(i, j, o p_{-} i, o p_{-} j, o p_{-}\)string \(\left.=' I d^{\prime}\right)\)
Add a two-site coupling term on given MPS sites.

\section*{Parameters}
- strength (float) - The strength of the coupling term.
- i (int) - The MPS indices of the two sites on which the operator acts. We require 0 <= i < N_sites and i < j, i.e., op_i acts "left" of op_j. If \(\mathrm{j}>=\mathrm{N}_{-}\)sites, it indicates couplings between unit cells of an infinite MPS.
- \(\mathbf{j}(i n t)\) - The MPS indices of the two sites on which the operator acts. We require 0 \(<=i<N \_s i t e s\) and \(i<j\), i.e., op_i acts "left" of \(o p_{-} j\). If \(\mathrm{j}>=\mathrm{N}_{-}\)sites, it indicates couplings between unit cells of an infinite MPS.
- op1 (str) - Names of the involved operators.
- op2 (str) - Names of the involved operators.
- op_string (str) - The operator to be inserted between \(i\) and \(j\).
coupling_term_handle_JW (strength, term, sites, op_string=None)
Helping function to call before add_coupling_term().

\section*{Parameters}
- strength (float) - The strength of the coupling term.
- term ([(str, int), (str, int)]) - List of two tuples [(op_i, i), (op_j, j)] where \(i\) is the MPS index of the site the operator named op_i acts on; we require \(i<j\).
- sites (list of Site) - Defines the local Hilbert space for each site. Used to check whether the operators need Jordan-Wigner strings.
- op_string (None / str) - Operator name to be used as operator string between the operators, or None if the Jordan Wigner string should be figured out.

Warning: None figures out for each segment between the operators, whether a Jordan-Wigner string is needed. This is different from a plain ' JW', which just applies a string on each segment!

Returns Arguments for MultiCouplingTerms.add_multi_coupling_term() such that the added term corresponds to the parameters of this function.
Return type strength, i, j, op_i, op_j, op_string
classmethod from_hdf5 (hdf5_loader, h5gr, subpath)
Load instance from a HDF5 file.
This method reconstructs a class instance from the data saved with save_hdf5().

\section*{Parameters}
- hdf5_loader (Hdf5Loader) - Instance of the loading engine.
- h5gr (Group) - HDF5 group which is represent the object to be constructed.
- subpath (str) - The name of \(h 5 g r\) with a ' / ' in the end.

Returns obj - Newly generated class instance containing the required data.
Return type cls
plot_coupling_terms (ax, lat, style_map='default', common_style=\{'linestyle': '--'\}, text=None, text_pos=0.4)
"Plot coupling terms into a given lattice.
This function plots the coupling_terms

\section*{Parameters}
- ax (matplotlib.axes.Axes) - The axes on which we should plot.
- lat (Lattice) - The lattice for plotting the couplings, most probably the M. lat of the corresponding model M , see lat.
- style_map (function | None) - Function which get's called with arguments i, \(j\), op_i, op_string, op_j, strength for each two-site coupling and should return a keyword-dictionary with the desired plot-style for this coupling. By default (None), the linewidth is given by the absolute value of strength, and the linecolor depends on the phase of strength (using the hsv colormap).
- common_style (dict) - Common style, which overwrites values of the dictionary returned by style_map. A ' label' is only used for the first plotted line.
- text (format_string | None) - If not None, we add text labeling the couplings in the plot. Available keywords are i, j, op_i, op_string, op_j, strength as well as strength_abs, strength_angle, strength_real.
- text_pos (float) - Specify where to put the text on the line between \(i(0.0)\) and \(j\) (1.0), e.g. 0.5 is exactly in the middle between \(i\) and \(j\).

\section*{See also:}
tenpy.models.lattice.Lattice.plot_sites plot the sites of the lattice.
save_hdf5 (hdf5_saver, h5gr, subpath)
Export self into a HDF5 file.
This method saves all the data it needs to reconstruct self with from_hdf5 ().
This implementation saves the content of __dict __ with save_dict_content (), storing the format under the attribute ' format'.

\section*{Parameters}
- hdf5_saver (Hdf5Saver) - Instance of the saving engine.
- h5gr (: class `Group \(\left.{ }^{\prime}\right)\) - HDF5 group which is supposed to represent self.
- subpath (str) - The name of \(h 5 g r\) with a ' \(/ '\) in the end.
to_nn_bond_Arrays (sites)
Convert the coupling_terms into Arrays on nearest neighbor bonds.
Parameters sites (list of Site) - Defines the local Hilbert space for each site. Used to translate the operator names into Array.

Returns H_bond - The coupling_terms rewritten as sum_i H_bond[i] for MPS indices i. H_bond[i] acts on sites (i-1, i), None represents 0 . Legs of each H_bond[i] are ['p0', 'p0*', 'p1', 'p1*'].
Return type list of \(\left\{\right.\) Array \({ }^{\prime}\) None \(\}\)

\subsection*{20.4.4 OnsiteTerms}
- full name: tenpy.networks.terms.OnsiteTerms
- parent module: tenpy.networks.terms
- type: class

\section*{Inheritance Diagram}


\section*{Methods}
\begin{tabular}{|c|c|}
\hline OnsiteTerms.__init__(L) & Initialize self. \\
\hline OnsiteTerms.add_onsite_term(strength, i, op) & Add a onsite term on a given MPS site. \\
\hline OnsiteTerms.add_to_graph(graph) & Add terms from onsite_terms to an MPOGraph. \\
\hline OnsiteTerms.add_to_nn_bond_Arrays(H_bon ...) & Add self.onsite_terms into nearest-neighbor bond arrays. \\
\hline OnsiteTerms.from_hdf5(hdf5_loader, h5gr, subpath) & Load instance from a HDF5 file. \\
\hline OnsiteTerms.max_range() & Maximum range of the terms. \\
\hline OnsiteTerms.remove_zeros([tol_zero]) & Remove entries close to 0 from onsite_terms. \\
\hline OnsiteTerms.save_hdf5(hdf5_saver, h5gr, subpath) & Export self into a HDF5 file. \\
\hline OnsiteTerms.to_Arrays(sites) & Convert the onsite_terms into a list of np_conserved Arrays. \\
\hline OnsiteTerms.to_TermList() & Convert onsite_terms into a TermList. \\
\hline \multicolumn{2}{|l|}{class tenpy.networks.terms.OnsiteTerms ( \(L\) )} \\
\hline \multicolumn{2}{|l|}{Bases: tenpy.tools.hdf5_io.Hdf5Exportable} \\
\hline \multicolumn{2}{|l|}{Operator names, site indices and strengths representing onsite terms.} \\
\hline
\end{tabular}

Represents a sum of onsite terms where the operators are only given by their name (in the form of a string). What the operator represents is later given by a list of Site with get_op ().

Parameters L (int) - Number of sites.
L
Number of sites.
Type int

\section*{onsite_terms}

Filled by meth:add_onsite_term. For each index \(i\) a dictionary \{'opname' : strength\} defining the onsite terms.

Type list of dict
max_range ()
Maximum range of the terms. In this case 0.
add_onsite_term (strength, \(i, o p\) )
Add a onsite term on a given MPS site.

\section*{Parameters}
- strength (float) - The strength of the term.
- \(\mathbf{i}(i n t)\) - The MPS index of the site on which the operator acts. We require \(0<=i<\) L.
- op (str) - Name of the involved operator.
add_to_graph (graph)
Add terms from onsite_terms to an MPOGraph.
Parameters graph (MPOGraph) - The graph into which the terms from onsite_terms should be added.
to_Arrays (sites)
Convert the onsite_terms into a list of np_conserved Arrays.
Parameters sites (list of Site) - Defines the local Hilbert space for each site. Used to translate the operator names into Array.
Returns onsite_arrays - Onsite terms represented by self. Entry \(i\) of the list lives on sites[i].
Return type list of Array
remove_zeros (tol_zero=le-15)
Remove entries close to 0 from onsite_terms.
Parameters tol_zero (float) - Entries in onsite_terms with strength <tol_zero are considered to be zero and removed.
add_to_nn_bond_Arrays (H_bond, sites, finite, distribute=(0.5, 0.5))
Add self.onsite_terms into nearest-neighbor bond arrays.

\section*{Parameters}
- H_bond (list of \{Array | None\}) - The coupling_terms rewritten as sum_i H_bond[i] for MPS indices i. H_bond[i] acts on sites (i-1, i), None represents 0. Legs of each H_bond[i] are ['p0', 'p0*', 'p1', 'p1*']. Modified in place.
- sites (list of Site) - Defines the local Hilbert space for each site. Used to translate the operator names into Array.
- distribute ((float, float)) - How to split the onsite terms (in the bulk) into the bond terms to the left (distribute [0]) and right (distribute [1]).
- finite (bool) - Boundary conditions of the MPS, MPS.finite. If finite, we distribute the onsite term of the
to_TermList()
Convert onsite_terms into a TermList.
Returns term_list - Representation of the terms as a list of terms.
Return type TermList
classmethod from_hdf5 (hdf5_loader, h5gr, subpath)
Load instance from a HDF5 file.
This method reconstructs a class instance from the data saved with save_hdf5 ().

\section*{Parameters}
- hdf5_loader (Hdf5Loader) - Instance of the loading engine.
- h5gr (Group) - HDF5 group which is represent the object to be constructed.
- subpath (str) - The name of \(h 5 g r\) with a '/' in the end.

Returns obj - Newly generated class instance containing the required data.
Return type cls
save_hdf5 (hdf5_saver, h5gr, subpath)
Export self into a HDF5 file.
This method saves all the data it needs to reconstruct self with from_hdf5 ().
This implementation saves the content of \(\qquad\) dict \(\qquad\) with save_dict_content (), storing the format under the attribute 'format'.

\section*{Parameters}
- hdf5_saver (Hdf5SAver) - Instance of the saving engine.
- h5gr (: class`Group`) - HDF5 group which is supposed to represent self.
- subpath (str) - The name of \(h 5 g r\) with a '/' in the end.

\subsection*{20.4.5 TermList}
- full name: tenpy.networks.terms.TermList
- parent module: tenpy.networks.terms
- type: class

\section*{Inheritance Diagram}


\section*{Methods}
\begin{tabular}{ll}
\hline TermList.__init__(terms[, strength]) & Initialize self. \\
\hline TermList.from_hdf5(hdf5_loader, h5gr, subpath) & Load instance from a HDF5 file. \\
\hline \begin{tabular}{l} 
TermList.from_lattice_locations(lattice, \\
terms)
\end{tabular} & \begin{tabular}{l} 
Initialize from a list of terms given in lattice indices in- \\
stead of MPS indices.
\end{tabular} \\
\hline TermList.order_combine(sites) & Order and combine operators in each term. \\
\hline TermList.save_hdf5(hdf5_saver, h5gr, subpath) & Export self into a HDF5 file. \\
\hline TermList.to_OnsiteTerms_CouplingTerms(sitesonvert to OnsiteTerms and CouplingTerms \\
\hline \\
class tenpy.networks.terms.TermList (terms, strength=1.0) \\
\(\quad\) Bases: tenpy.tools.hdf5_io.Hdf5Exportable
\end{tabular}

A list of terms (=operator names and sites they act on) and associated strengths.
A representation of terms, similar as OnsiteTerms, CouplingTerms and MultiCouplingTerms.
This class does not store operator strings between the sites. Jordan-Wigner strings of fermions are added during conversion to (Multi)CouplingTerms.

Warning: Since this class does not store the operator string between the sites, conversion from CouplingTerms or MultiCouplingTerms to TermList is lossy!

\section*{Parameters}
- terms (list of list of (str, int))-List of terms where each term is a list of tuples (opname, i) of an operator name and a site \(i\) it acts on. For Fermions, the order is the order in the mathematic sense, i.e., the right-most/last operator in the list acts last.
- strengths ((list of) float/complex) - For each term in terms an associated prefactor or strength. A single number holds for all terms equally.
terms
List of terms where each term is a tuple (opname, i) of an operator name and a site \(i\) it acts on.
Type list of list of (str, int)

\section*{strengths}

For each term in terms an associated prefactor or strength.

> Type 1D ndarray

\section*{Examples}

For fermions, the term \(0.5\left(c_{0}^{\dagger} c_{2}+\right.\) h.c. \()+1.3 * n_{1}\) can be represented by:
```

>>> t = TermList([[('Cd', 0), ('C', 2)], [('Cd', 2), ('C', 0)], [('N', 1)]],
.. [0.5, 0.5, 1.3])
>>> print(t)
0.50000 * Cd_0 C_2 +
0.50000 * Cd_2 C_0 +
1.30000 * N_1

```

If you have a Lattice, you might also want to specify the location of the operators by lattice indices insted of MPS indices. For example, you can obtain the nearest-neighbor density terms without double counting each pair) on a TriangularLattice:
```

>>> lat = tenpy.models.lattice.Triangular(6, 6, None, bc_MPS='infinite', bc=
\hookrightarrow'periodic')
>>> t2_terms = [[('N', [0, 0, u1]), ('N', [dx[0], dx[1], u2])]
... for (u1, u2, dx) in lat.pairs['nearest_neighbors']]
>>> t2 = TermList.from_lattice_locations(lat, t2_terms)
>>> print(t2)
1.00000 * N_O N_6 +
1.00000 * N_O N_-5 +
1.00000 * N_00 N__5

```

The negative index -5 here indicates a tensor left of the current MPS unit cell.
classmethod from_lattice_locations (lattice, terms, strength=1.0, shift=None)
Initialize from a list of terms given in lattice indices instead of MPS indices.

\section*{Parameters}
- lattice (Lattice) - The underlying lattice to be used for conversion, e.g. M.lat from a Model.
- terms (list of list of (str, tuple)) - List of terms, where each term is a tuple (opname, lat_idx) with lat_idx itself being a tuple ( \(x, y, u\) ) (for a 2D lattice) of the lattice corrdinates.
- strengths ((list of) float/complex) - For each term in terms an associated prefactor or strength. A single number holds for all terms equally.
- shift (None / tuple of int) - Overall shift added to all lattice coordinates lat_idx in terms before conversion. None defaults to no shift.
Returns term_list - Representation of the terms.
Return type TermList
to_OnsiteTerms_CouplingTerms (sites)
Convert to OnsiteTerms and CouplingTerms
Performs Jordan-Wigner transformation for fermionic operators.

Parameters sites (list of Site) - Defines the local Hilbert space for each site. Used to check whether the operators need Jordan-Wigner strings. The length is used as \(L\) for the onsite_terms and coupling_terms.

\section*{Returns}
- onsite_terms (OnsiteTerms) - Onsite terms.
- coupling_terms (CouplingTermsIMultiCouplingTerms) - Coupling terms. If self contains terms involving more than two operators, a MultiCouplingTerms instance, otherwise just CouplingTerms.
order_combine (sites)
Order and combine operators in each term.
Parameters sites (list of Site) - Defines the local Hilbert space for each site. Used to check whether the operators anticommute (= whether they need Jordan-Wigner strings) and for multiplication rules.

\section*{See also:}
order_and_combine_term does it for a single term.
classmethod from_hdf5 (hdf5_loader, h5gr, subpath)
Load instance from a HDF5 file.
This method reconstructs a class instance from the data saved with save_hdf5().

\section*{Parameters}
- hdf5_loader (Hdf5Loader) - Instance of the loading engine.
- h5gr (Group) - HDF5 group which is represent the object to be constructed.
- subpath (str) - The name of \(h 5 g r\) with a '/' in the end.

Returns obj - Newly generated class instance containing the required data.
Return type cls
save_hdf5 (hdf5_saver, h5gr, subpath)
Export self into a HDF5 file.
This method saves all the data it needs to reconstruct self with from_hdf5 ().
This implementation saves the content of __dict__ with save_dict_content (), storing the format under the attribute 'format'.

\section*{Parameters}
- hdf5_saver (Hdf5Saver) - Instance of the saving engine.
- h5gr (: class`Group \()\) - HDF5 group which is supposed to represent self.
- subpath (str) - The name of \(h 5 g r\) with a '/' in the end.

\section*{Functions}
order_combine_term(term, sites) Combine operators in a term to one terms per site.

\subsection*{20.4.6 order_combine_term}
- full name: tenpy.networks.terms.order_combine_term
- parent module: tenpy.networks.terms
- type: function
tenpy.networks.terms.order_combine_term (term, sites)
Combine operators in a term to one terms per site.
Takes in a term of operators and sites they acts on, commutes operators to order them by site and combines operators acting on the same site with multiply_op_names ().

\section*{Parameters}
- term (a list of (opname_i, i) tuples) - Represents a product of onsite operators with site indices \(i\) they act on. Needs not to be ordered and can have multiple entries acting on the same site.
- sites (list of Site) - Defines the local Hilbert space for each site. Used to check whether the operators anticommute (= whether they need Jordan-Wigner strings) and for multiplication rules.

\section*{Returns}
- combined_term - Equivalent to term but with at most one operator per site.
- overall_sign \((+1|-1| 0)\) - Comes from the (anti-)commutation relations. When the operators in term are multiplied from left to right, and then multiplied by overall_sign, the result is the same operator as the product of combined_term from left to right.

\section*{Module description}

Classes to store a collection of operator names and sites they act on, together with prefactors.
This modules collects classes which are not strictly speaking tensor networks but represent "terms" acting on them. Each term is given by a collection of (onsite) operator names and indices of the sites it acts on. Moreover, we associate a strength to each term, which corresponds to the prefactor when specifying e.g. a Hamiltonian.

\section*{20.5 purification_mps}
- full name: tenpy.networks.purification_mps
- parent module: tenpy. networks
- type: module

\section*{Classes}


\footnotetext{
PurificationMPS(sites, Bs, SVs[, bc, form, norm]) An MPS representing a finite-temperature ensemble using purification.
}

\section*{Module description}

This module contains an MPS class representing an density matrix by purification.
Usually, an MPS represents a pure state, i.e. the density matrix is \(\rho=|\psi><\psi|\), describing observables as \(<O>=\) \(\operatorname{Tr}(O|\psi><\psi|)=<\psi|O| \psi>\). Clearly, if \(\mid \psi>\) is the ground state of a Hamiltonian, this is the density matrix at \(T=0\).
At finite temperatures \(T>0\), we want to describe a non-pure density matrix \(\rho=\exp (-H / T)\). This can be accieved by the so-called purification: in addition to the physical space \(P\), we introduce a second 'auxiliar' space \(Q\) and define the density matrix of the physical system as \(\rho=\operatorname{Tr}_{Q}(|\phi><\phi|)\), where \(\mid \phi>\) is a pure state in the combined phyisical and auxiliar system.

For \(T=\infty\), the density matrix \(\rho_{\infty}\) is the identity matrix. In other words, expectation values are sums over all possible states \(<O>=\operatorname{Tr}_{P}\left(\rho_{\infty} O\right)=\operatorname{Tr}_{P}(O)\). Saying that each : on top is to be connected with the corresponding : on the bottom, the trace is simply a contraction:


Clearly, we get the same result, if we insert an identity operator, written as MPO, on the top and bottom:



We use the following label convention:
\begin{tabular}{|c|c|}
\hline & q \\
\hline & ^ \\
\hline & | \\
\hline \multicolumn{2}{|l|}{\multirow[t]{3}{*}{vL \(->-\mathrm{B}->-\mathrm{vR}\)}} \\
\hline & \\
\hline & \\
\hline & p \\
\hline
\end{tabular}

You can view the \(M P O\) as an MPS by combining the \(p\) and \(q\) leg and defining every physical operator to act trivial on the \(q\) leg. In expecation values, you would then sum over over the \(q\) legs, which is exactly what we need. In other words, the choice \(B=\delta_{p, q}\) with trivial (length-1) virtual bonds yields infinite temperature expectation values for operators action only on the \(p\) legs!

Now, you go a step further and also apply imaginary time evolution (acting only on \(p\) legs) to the initial infinite temperature state. For example, the normalized state \(|\psi>\alpha \exp (-\beta / 2 H)| \phi>\) yields expecation values
\[
<O>=\operatorname{Tr}(\exp (-\beta H) O) / \operatorname{Tr}(\exp (-\beta H)) \propto<\phi|\exp (-\beta / 2 H) O \exp (-\beta / 2 H)| \phi\rangle .
\]

An additional real-time evolution allows to calculate time correlation functions:
\[
<A(t) B(0)>\propto<\phi|\exp (-\beta H / 2) \exp (+i H t) A \exp (-i H t) B \exp (-\beta H / 2)| \phi>
\]

Time evolution algorithms (TEBD and MPO application) are adjusted in the module purification.
See also [[karrasch2013]] for additional tricks! On of their crucial observations is, that one can apply arbitrary unitaries on the auxiliar space (i.e. the \(q\) ) without changing the result. This can actually be used to reduce the necessary virtual bond dimensions: From the definition, it is easy to see that if we apply \(\exp (-i H t)\) to the \(p\) legs of \(|\phi\rangle\), and \(\exp (+i H t)\) to the \(q\) legs, they just cancel out! (They commute with \(\exp (-\beta H / 2) \ldots\) ) If the state is modified (e.g. by applying \(A\) or \(B\) to calculate correlation functions), this is not true any more. However, we still can find unitaries, which are 'optimal' in the sense of reducing the entanglement of the MPS/MPO to the minimal value. For a discussion of Disentanglers (implemented in disent anglers), see [[hauschild2018]].

Note: The classes MPSEnvironment and TransferMatrix should also work for the PurificationMPS defined here. For example, you can use expectation_value () for the expectation value of operators between different PurificationMPS. However, this makes only sense if the same disentangler was applied to the bra and ket PurificationMPS.

Note: The literature (e.g. section 7.2 of [[schollwoeck2011]] or [[karrasch2013]]) suggests to use a singlet as a maximally entangled state. Here, we use instead the identity \(\delta_{p, q}\), since it is easier to generalize for \(p\) running over more than two indices, and allows a simple use of charge conservation with the above qconj convention. Moreover, we don't split the physical and auxiliar space into separate sites, which makes TEBD as costly as \(O\left(d^{6} \chi^{3}\right)\).

\section*{SIMULATIONS}
- full name: tenpy.simulations
- parent module: tenpy
- type: module

\section*{Module description}

Simulation setup.
The classes provided here provide a structure for the whole setup of simulations.

\section*{Submodules}
\begin{tabular}{ll}
\hline simulation & This module contains base classes for simulations. \\
\hline measurement & Functions to perform measurments. \\
\hline ground_state_search & Simulations for ground state searches. \\
\hline time_evolution & Simulations for (real) time evolution. \\
\hline
\end{tabular}

\section*{21.1 simulation}
- full name: tenpy.simulations.simulation
- parent module: tenpy.simulations
- type: module

\section*{Classes}
```

Simulation

```
```

Skip

```
Simulation(options, *[, setup_logging]) Base class for simulations.

\section*{Exceptions}
Skip Error raised if simulation output already exists.

\subsection*{21.1.1 Skip}
- full name: tenpy.simulations.simulation.Skip
- parent module: tenpy.simulations.simulation
- type: exception
exception tenpy.simulations.simulation.Skip
Error raised if simulation output already exists.

\section*{Functions}
\begin{tabular}{ll}
\hline resume_from_checkpoint \((*[\), filename, \(\ldots])\) & Resume a simulation run from a given checkpoint. \\
\hline run_simulation \(([\) simulation_class_name,..\(])\) & Run the simulation with a simulation class. \\
\hline
\end{tabular}

\subsection*{21.1.2 resume_from_checkpoint}
- full name: tenpy.simulations.simulation.resume_from_checkpoint
- parent module: tenpy.simulations.simulation
- type: function
tenpy.simulations.simulation.resume_from_checkpoint(*, filename=None, checkpoint_results=None, update_sim_params \(=\) None, simulation_class_kwargs=None)
Resume a simulation run from a given checkpoint.
(All parameters have to be given as keyword arguments.)

\section*{Parameters}
- filename (None / str) - The filename of the checkpoint to be loaded. You can either specify the filename or the checkpoint_results.
- checkpoint_results (None / dict) - Alternatively to filename the results of the simulation so far, i.e. directly the data dicitonary saved at a simulation checkpoint.
- update_sim_params (None / dict) - Allows to update specific Simulation parameters, ignored if None. Uses update_recursive () to update values, such that the keys of update_sim_params can be recursive, e.g. algorithm_params/max_sweeps.
- simlation_class_kwargs (None / dict) - Further keyword arguemnts given to the simulation class, ignored if None.

Returns The results from running the simulation, i.e., what tenpy.simulations. Simulation.resume_run() returned.
Return type results

\section*{Notes}

The checkpoint_filename should be relative to the current working directory. If you use the Simulation. directory, the simulation class will attempt to change to that directory during initialization. Hence, either resume the simulation from the same directory where you originally started, or update the Simulation. directory (and :cfg:option`Simulation.output_filename`) parameter with update_sim_params.

\subsection*{21.1.3 run_simulation}
- full name: tenpy.simulations.simulation.run_simulation
- parent module: tenpy.simulations.simulation
- type: function
```

tenpy.simulations.simulation.run_simulation(simulation_class_name='GroundStateSearch',
simulation_class_kwargs=None, **simula- tion_params)

```

Run the simulation with a simulation class.

\section*{Parameters}
- simulation_class_name (str) - The name of a (sub)class of Simulation to be used for running the simulaiton.
- simulation_class_kwargs (dict / None) - A dictionary of keyword-arguments to be used for the initializing the simulation.
- **simulation_params - Further keyword arguments as documented in the corresponding simulation class, see :cfg:config`Simulation`.

Returns The results from running the simulation, i.e., what tenpy.simulations. Simulation.run() returned.
Return type results

\section*{Module description}

This module contains base classes for simulations.
The Simulation class tries to put everything need for a simulation in a structured form and collects task like initializing the tensor network state, model and algorithm classes, running the actual algorithm, possibly performing measurements and saving the results.

Todo: provide examples, give user guide

\section*{21.2 measurement}
- full name: tenpy.simulations.measurement
- parent module: tenpy.simulations
- type: module

\section*{Functions}
\begin{tabular}{ll}
\hline bond_dimension(results, psi, simulation[, key]) & 'Measure' the bond dimension of an MPS. \\
\hline bond_energies(results, psi, simulation[, key]) & Measure the energy of an MPS. \\
\hline correlation_length(results, psi, simulation) & Measure the correlaiton of an infinite MPS. \\
\hline energy_MPO(results, psi, simulation[, key]) & \begin{tabular}{l} 
Measure the energy of an MPS by evaluating the MPS \\
expectation value.
\end{tabular} \\
\hline entropy(results, psi, simulation[, key]) & Measure the entropy at all bonds of an MPS. \\
\hline evolved_time(results, psi, simulation[, key]) & \begin{tabular}{l} 
Measure the time evolved by the engine, engine. \\
evolved_time.
\end{tabular} \\
\hline measurement_index(results, psi, simulation) & \begin{tabular}{l} 
'Measure' the index of how many mearuements have \\
been performed so far.
\end{tabular} \\
\hline onsite_expectation_value(results, psi,...) & Measure expectation values of an onsite operator. \\
\hline
\end{tabular}

\subsection*{21.2.1 bond_dimension}
- full name: tenpy.simulations.measurement.bond_dimension
- parent module: tenpy.simulations.measurement
- type: function
tenpy.simulations.measurement.bond_dimension(results, psi, simulation, key='bond_dimension')
'Measure' the bond dimension of an MPS.

\section*{Parameters}
- results - See measurement_index().
- psi-See measurement_index().
- simulation - See measurement_index().
- key - See measurement_index ().

\subsection*{21.2.2 bond_energies}
- full name: tenpy.simulations.measurement.bond_energies
- parent module: tenpy.simulations.measurement
- type: function
```

tenpy.simulations.measurement.bond_energies(results, psi, simulation,
key='bond_energies')
Measure the energy of an MPS.

```

\section*{Parameters}
- results - See measurement_index ().
- psi-See measurement_index().
- simulation - See measurement_index().
- key - See measurement_index().

\subsection*{21.2.3 correlation_length}
- full name: tenpy.simulations.measurement.correlation_length
- parent module: tenpy.simulations.measurement
- type: function
tenpy.simulations.measurement.correlation_length(results, psi, simulation, key='correlation_length', **kwargs)
Measure the correlaiton of an infinite MPS.

\section*{Parameters}
- results - See measurement_index ().
- psi-See measurement_index().
- simulation - See measurement_index().
- key - See measurement_index ().
- **kwargs - Further keywoard arguments given to correlation_length().

\subsection*{21.2.4 energy_MPO}
- full name: tenpy.simulations.measurement.energy_MPO
- parent module: tenpy.simulations.measurement
- type: function
tenpy.simulations.measurement.energy_MPO (results, psi, simulation, key='energy_MPO') Measure the energy of an MPS by evaluating the MPS expectation value.

\section*{Parameters}
- results - See measurement_index().
- psi-See measurement_index().
- simulation - See measurement_index().
- key - See measurement_index ().

\subsection*{21.2.5 entropy}
- full name: tenpy.simulations.measurement.entropy
- parent module: tenpy.simulations.measurement
- type: function
tenpy.simulations.measurement.entropy (results, psi, simulation, key='entropy')
Measure the entropy at all bonds of an MPS.

\section*{Parameters}
- results - See measurement_index().
- psi-See measurement_index().
- simulation - See measurement_index().
- key - See measurement_index().

\subsection*{21.2.6 evolved_time}
- full name: tenpy.simulations.measurement.evolved_time
- parent module: tenpy.simulations.measurement
- type: function
tenpy.simulations.measurement.evolved_time (results, psi, simulation, key='evolved_time')
Measure the time evolved by the engine, engine.evolved_time.
See e.g. tenpy.algorithms.tebd.TEBDEngine.evolved_time.

\section*{Parameters}
- results - See measurement_index().
- psi-See measurement_index().
- simulation - See measurement_index().
- key - See measurement_index().

\subsection*{21.2.7 measurement_index}
- full name: tenpy.simulations.measurement.measurement_index
- parent module: tenpy.simulations.measurement
- type: function
```

tenpy.simulations.measurement.measurement_index(results, psi, simulation,
key='measurement_index')

```
'Measure' the index of how many mearuements have been performed so far.
The parameter description below also documents the common interface of all measurement functions, that can be registered to simulations.

\section*{Parameters}
- results (dict) - A dictionary with measurement results performed so far. Instead of returning the result, the output should be written into this dictionary under an appropriate key (or multiple keys, if applicable).
- psi - Tensor network state to be measured. Shorthand for simulation.psi.
- simulation (Simulation) - The simulation class. This gives also access to the model, algorithm engine, etc.
- key (str) - (Optional.) The key under which to save in results.
- **kwargs - Other optional keyword arguments for individual measurement functions. Those are documented inside each measurement function.

\subsection*{21.2.8 onsite_expectation_value}
- full name: tenpy.simulations.measurement.onsite_expectation_value
- parent module: tenpy.simulations.measurement
- type: function
```

tenpy.simulations.measurement.onsite_expectation_value (results, psi, simulation, op-
name, key=None)

```

Measure expectation values of an onsite operator.
The resulting array of measurements is indexed by lattice indices ( \(x, y, u\) ) (possibly dropping \(y\) and/or \(u\) if they are trivial), not by the MPS index. Note that this makes the result independent of the way the MPS winds through the lattice.
The key defaults to f " \(<\{\) opname \(\}>"\).

\section*{Parameters}
- results - See measurement_index ().
- psi-See measurement_index().
- simulation - See measurement_index().
- key - See measurement_index ().
- opname (str) - The operator to be measured. Passed on to expectation_value ().

\section*{Module description}

Functions to perform measurments.
All measurement functions provided in this module support the interface used by the simulation class, i.e. they take the parameters documented in measurement_index () and write the measurement results into the results dictionary taken as argument.

Todo: test, provide more.

\section*{21.3 ground_state_search}
- full name: tenpy.simulations.ground_state_search
- parent module: tenpy.simulations
- type: module

\section*{Classes}


GroundStateSearch(options, *[, setup_logging]) Simutions for variational ground state searches.

\section*{Module description}

Simulations for ground state searches.

\section*{21.4 time_evolution}
- full name: tenpy.simulations.time_evolution
- parent module: tenpy.simulations
- type: module

\section*{Classes}


\section*{Module description}

Simulations for (real) time evolution.
- full name: tenpy.tools
- parent module: tenpy
- type: module

\section*{Module description}

A collection of tools: mostly short yet quite useful functions.
Some functions are explicitly imported in other parts of the library, others might just be useful when using the libary. Common to all tools is that they are not just useful for a single algorithm but fairly general.

\section*{Submodules}
\begin{tabular}{ll}
\hline hdf5_io & Tools to save and load data (from TeNPy) to disk. \\
\hline params & Tools to handle config options/paramters for algorithms. \\
\hline events & Event handler. \\
\hline misc & \begin{tabular}{l} 
Miscellaneous tools, somewhat random mix yet often \\
helpful.
\end{tabular} \\
\hline math & \begin{tabular}{l} 
Different math functions needed at some point in the \\
library.
\end{tabular} \\
\hline fit & tools to fit to an algebraic decay. \\
\hline string & Tools for handling strings. \\
\hline process & \begin{tabular}{l} 
Tools to read out total memory usage and get/set the \\
number of threads.
\end{tabular} \\
\hline optimization & Optimization options for this library. \\
\hline
\end{tabular}

\section*{22.1 hdf5_io}
- full name: tenpy.tools.hdf5_io
- parent module: tenpy.tools
- type: module

\section*{Classes}

\begin{tabular}{ll}
\hline Hdf5Exportable() & \begin{tabular}{l} 
Interface specification for a class to be exportable to our \\
HDF5 format.
\end{tabular} \\
\hline Hdf5Ignored([name]) & \begin{tabular}{l} 
Placeholder for a dataset/group to be ignored during \\
both loading and saving.
\end{tabular} \\
\hline Hdf5Loader(h5group[, ignore_unknown, exclude]) & Class to load and import object from a HDF5 file. \\
\hline Hdf5Saver(h5group[, format_selection]) & Class to save simple enough objects into a HDF5 file. \\
\hline
\end{tabular}

\subsection*{22.1. Hdf5Exportable}
- full name: tenpy.tools.hdf5_io.Hdf5Exportable
- parent module: tenpy.tools.hdf5_io
- type: class

\section*{Inheritance Diagram}

\section*{Hdf5Exportable}

\section*{Methods}
```

Hdf5Exportable.__init__() Initialize self.
Hdf5Exportable.from_hdf5(hdf5_loader, h5gr, Load instance from a HDF5 file.
...)
Hdf5Exportable.save_hdf5(hdf5_saver, h5gr, Export self into a HDF5 file.
...)
class tenpy.tools.hdf5_io.Hdf5Exportable
Bases: object

```

Interface specification for a class to be exportable to our HDF5 format.
To allow a class to be exported to HDF5 with save_to_hdf5 (), it only needs to implement the save_hdf5 () method as documented below. To allow import, a class should implement the classmethod from_hdf5 (). During the import, the class already needs to be defined; loading can only initialize instances, not define classes.

The implementation given works for sufficiently simple (sub-)classes, for which all data is stored in __ dict __ In particular, this works for python-defined classes which simply store data using self.data \(=\) data in their methods.
save_hdf5 (hdf5_saver, h5gr, subpath)
Export self into a HDF5 file.
This method saves all the data it needs to reconstruct self with from_hdf5 ().
This implementation saves the content of __dict__ with save_dict_content (), storing the format under the attribute 'format'.

\section*{Parameters}
- hdf5_saver (Hdf5Saver) - Instance of the saving engine.
- h5gr (: class`Group \()\) - HDF5 group which is supposed to represent self.
- subpath (str) - The name of \(h 5 g r\) with a ' /' in the end.
classmethod from_hdf5 (hdf5_loader, h5gr, subpath)
Load instance from a HDF5 file.
This method reconstructs a class instance from the data saved with save_hdf5().

\section*{Parameters}
- hdf5_loader (Hdf5Loader) - Instance of the loading engine.
- h5gr (Group) - HDF5 group which is represent the object to be constructed.
- subpath (str) - The name of \(h 5 g r\) with a '/' in the end.

Returns obj - Newly generated class instance containing the required data.
Return type cls

\subsection*{22.1.2 Hdf5Ignored}
- full name: tenpy.tools.hdf5_io.Hdf5Ignored
- parent module: tenpy.tools.hdf5_io
- type: class

\section*{Inheritance Diagram}

\section*{Hdf5Ignored}

\section*{Methods}
Hdf5Ignored.__init__([name]) Initialize self.
class tenpy.tools.hdf5_io.Hdf5Ignored (name='unknown')
Bases: object
Placeholder for a dataset/group to be ignored during both loading and saving.
Objects of this type are not saved. Moreover, if a saved dataset/group has the type attribute matching REPR_IGNORED, instance of this class are returned instead of loading the data.

Parameters name (str) - The name of the dataset during loading; just for reference.

\section*{name}

See above.
Type str

\subsection*{22.1.3 Hdf5Loader}
- full name: tenpy.tools.hdf5_io.Hdf5Loader
- parent module: tenpy.tools.hdf5_io
- type: class

\section*{Inheritance Diagram}

\author{
Hdf5Loader
}

\section*{Methods}
\begin{tabular}{|c|c|}
\hline Hdf5Loader.__init__(h5group[,...]) & Initialize self. \\
\hline Hdf5Loader.get_attr(h5gr, attr_name) & Return attribute h5gr.attrs[attr_name], if existent. \\
\hline Hdf5Loader.load([path]) & Load a Python ob ject from the dataset. \\
\hline Hdf5Loader.load_dataset(h5gr, type_info,
subpath) & Load a h5py Dataset and convert it into the desired type. \\
\hline Hdf5Loader.load_dict(h5gr, type_info, subpath) & Load a dictionary in the format according to type_info. \\
\hline Hdf5Loader.load_dtype(h5gr, type_info, subpath) & Load a numpy.dtype. \\
\hline Hdf5Loader.load_general_dict(h5gr, ...) & Load a dictionary with general keys. \\
\hline Hdf5Loader.load_global(h5gr, type_info, subpath) & Load a global object like a class or function from its qualified name and module. \\
\hline Hdf5Loader.load_hdf5exportable(h5gr, ...) & Load an instance of a userdefined class. \\
\hline Hdf5Loader.load_ignored(h5gr, type_info, subpath) & Ignore the group to be loaded. \\
\hline Hdf5Loader.load_list(h5gr, type_info, subpath) & Load a list. \\
\hline Hdf5Loader. load_none(h5gr, type_info, subpath) & Load the None object from a dataset. \\
\hline Hdf5Loader. load_range(h5gr, type_info, subpath) & Load a range. \\
\hline Hdf5Loader. load_reduce(h5gr, type_info, subpath) & Load an object where the return values of obj.
\(\qquad\) reduce \(\qquad\) has been exported. \\
\hline Hdf5Loader.load_set(h5gr, type_info, subpath) & Load a set. \\
\hline Hdf5Loader.load_simple_dict(h5gr, type_info, ...) & Load a dictionary with simple keys. \\
\hline Hdf5Loader.load_str(h5gr, type_info, subpath) & Load a string from a h5py Dat aset. \\
\hline Hdf5Loader. load_tuple(h5gr, type_info, subpath) & Load a tuple. \\
\hline Hdf5Loader.memorize_load(h5gr, obj) & Store objects already loaded in the memo_load. \\
\hline
\end{tabular}

\section*{Class Attributes and Properties}
```

Hdf5Loader.dispatch_load

```
class tenpy.tools.hdf5_io.Hdf5Loader (h5group, ignore_unknown=True, exclude=None)
Bases: ob ject
Class to load and import object from a HDF5 file.
The intended use of this class is through load_from_hdf5(), which is simply an alias for Hdf5Loader (h5group). load (path).

It can load data exported with save_to_hdf5 () or the Hdf5Saver, respectively.
The basic structure of this class is similar as the Unpickler from pickle.
See Saving to disk: input/output for a specification of what can be saved and what the resulting datastructure is.

\section*{Parameters}
- h5group (Group) - The HDF5 group (or file) where to save the data.
- ignore_unknown (bool) - Whether to just warn (True) or raise an Error (False) if a class to be loaded is not found.
- exclude (list of str) - List of paths (possibly relative to h5group) for objects to be excluded from loading. References to the corresponding object are replaced by an instance of Hdf5Ignored. Of course, this might break other functions expecting correctly loaded data.

\section*{h5group}

The HDF5 group (or HDF5 File) where to save the data.
Type Group

\section*{ignore_unknown}

Whether to just warn (True) or raise an Error (False) if a class to be loaded is not found.
Type bool

\section*{dispatch_load}

Mapping from one of the global REPR_* variables to (unbound) methods \(f\) of this class. The method is called as f (self, h5gr, type_info, subpath). The call to \(f\) should load and return an object obj from the h5py Group or Dataset \(h 5 g r\); and memorize the loaded obj with memorize_load(). subpath is just the name of \(h 5 g r\) with a guaranteed ' / ' in the end. type_info is often the REPR_* variable of the type or some other information about the type, which allows to use a single dispatch_load function for different datatypes.

Type dict

\section*{memo_load}

A dictionary to remember all the objects which we already loaded from h5group. The dictionary key is a h5py group- or dataset id; the value is the loaded object. See memorize_load().

Type dict
load (path=None)
Load a Python ob ject from the dataset.
See load_from_hdf5 () for more details.
Parameters path (None I str I Reference) - Path within h5group to be used for loading.
Defaults to the name of \(h 5\) group itself.
Returns obj - The Python object loaded from h5group (specified by path).
Return type object
memorize_load (h5gr,obj)
Store objects already loaded in the memo_load.
This allows to avoid copies, if the same dataset appears multiple times in the hdf5 group of obj. Examples can be shared LegCharge objects or even shared Array.

To handle cyclic references correctly, this function should be called before loading data from subgroups with new calls of load ().
static get_attr (h5gr, attr_name)
Return attribute h5gr.attrs [attr_name], if existent.
Raises Hdf5ImportError - If the attribute does not exist.
load_none ( \(h 5 g r\), type_info, subpath)
Load the None object from a dataset.
load_dataset (h5gr, type_info, subpath)
Load a h5py Dataset and convert it into the desired type.
load_str (h5gr, type_info, subpath)
Load a string from a h5py Dat aset.
load_list (h5gr, type_info, subpath) Load a list.
load_set (h5gr, type_info, subpath)
Load a set.
load_tuple (h5gr, type_info, subpath)
Load a tuple.
load_dict (h5gr, type_info, subpath)
Load a dictionary in the format according to type_info.
load_general_dict ( h 5 gr , type_info, subpath) Load a dictionary with general keys.
load_simple_dict (h5gr, type_info, subpath) Load a dictionary with simple keys.
load_range (h5gr, type_info, subpath) Load a range.
load_dtype (h5gr, type_info, subpath)
Load a numpy.dtype.
load_hdf5exportable (h5gr, type_info, subpath)
Load an instance of a userdefined class.
load_ignored (h5gr, type_info, subpath) Ignore the group to be loaded.
load_global (h5gr, type_info, subpath) Load a global object like a class or function from its qualified name and module.
load_reduce (h5gr, type_info, subpath)
Load an object where the return values of obj.__reduce__ has been exported.

\subsection*{22.1.4 Hdf5Saver}
- full name: tenpy.tools.hdf5_io.Hdf5Saver
- parent module: tenpy.tools.hdf5_io
- type: class

\section*{Inheritance Diagram}

\section*{Hdf5Saver}

\section*{Methods}
\begin{tabular}{ll}
\hline \begin{tabular}{l} 
Hdf5Saver.__init__(h5group[, \\
mat_selection])
\end{tabular} & for- \\
\hline Hdf5Saver.create_group_for_obj(path, obj) & Create an HDF5 group self.h5group [path] to \\
store obj.
\end{tabular}

\section*{Class Attributes and Properties}
```

Hdf5Saver.dispatch_save

```
class tenpy.tools.hdf5_io.Hdf5Saver (h5group, format_selection=None)
Bases: ob ject
Class to save simple enough objects into a HDF5 file.
The intended use of this class is through save_to_hdf5(), which is simply an alias for Hdf5Saver(h5group).save (obj, path).

It exports python objects to a HDF5 file such that they can be loaded with the Hdf5Loader, or a call to load_from_hdf5 (), respectively.

The basic structure of this class is similar as the Pickler from pickle.
See Saving to disk: input/output for a specification of what can be saved and what the resulting datastructure is.

\section*{Parameters}
- h5group (Group) - The HDF5 group (or HDF5 File) where to save the data.
- format_selection (dict) - This dictionary allows to set a output format selection for user-defined Hdf5Exportable.save_hdf5() implementations. For example, LegCharge checks it for the key "LegCharge".

\section*{h5group}

The HDF5 group (or HDF5 File) where to save the data.
Type Group

\section*{dispatch_save}

Mapping from a type keytype to methods \(f\) of this class. The method is called as f (self, obj, path, type_repr). The call to \(f\) should save the object obj in self.h5group [path], call memorize_save (), and set h5gr.attr[ATTR_TYPE] = type_repr to a string type_repr in order to allow loading with the dispatcher in Hdf5Loader. dispatch_save [type_repr].

\section*{Type dict}

\section*{memo_save}

A dictionary to remember all the objects which we already stored to h5group. The dictionary key is the object id; the value is a two-tuple of the hdf5 group or dataset where an object was stored, and the object itself. See memorize_save ().

Type dict
format_selection
This dictionary allows to set a output format selection for user-defined Hdf5Exportable. save_hdf5 () implementations. For example, LegCharge checks it for the key "LegCharge".

Type dict
save (obj, path='/')
Save obj in self.h5group [path].

\section*{Parameters}
- obj (object) - The object (=data) to be saved.
- path (str) - Path within h5group under which the obj should be saved. To avoid unwanted overwriting of important data, the group/object should not yet exist, except if path is the default '/'.
Returns \(\mathbf{h 5 g r}\) - The h5py group or dataset in which obj was saved.
Return type Group I Dataset
create_group_for_obj (path,obj)
Create an HDF5 group self.h5group [path] to store obj.
Also handle ending of path with '/', and memorize obj in memo_save.

\section*{Parameters}
- path (str) - Path within h5group under which the obj should be saved. To avoid unwanted overwriting of important data, the group/object should not yet exist, except if path is the default '/'.
- obj (object) - The object (=data) to be saved.

\section*{Returns}
- h5group (Group) - Newly created h5py (sub)group self.h5group [path], unless path is '/', in which case it is simply the existing self.h5group ['/'].
- subpath (str) - The group.name ending with '/', such that other names can be appended to get the path for subgroups or datasets in the group.
:raises ValueError : if self.h5group[path]` already existed and path is not ' / ' .:
```

memorize_save (h5gr,obj)

```

Store objects already saved in the memo_save.
This allows to avoid copies, if the same python object appears multiple times in the data of obj. Examples can be shared LegCharge objects or even shared Array. Using the memo also avoids crashes from cyclic references, e.g., when a list contains a reference to itself.

\section*{Parameters}
- h5gr (Group I Dataset) - The h5py group or dataset in which obj was saved.
- obj (ob ject) - The object saved.
save_reduce (func, args, state=None, listitems=None, dictitems=None, state_setter=None, obj=None, path=None)
Save the return values of obj.__reduce \(\qquad\) following the pickle protocol.
save_none (obj, path, type_repr)
Save the None object as a string (dataset); in dispatch table.
save_dataset (obj, path, type_repr)
Save obj as a hdf5 dataset; in dispatch table.
save_iterable (obj, path, type_repr)
Save an iterable obj like a list, tuple or set; in dispatch table.
save_iterable_content (obj, h5gr, subpath)
Save contents of an iterable obj in the existing \(h 5 g r\).

\section*{Parameters}
- obj (dict) - The data to be saved
- h5gr (Group) - h5py Group under which the keys and values of obj should be saved.
- subpath (str) - Name of h5gr with ' / ' in the end.
save_dict (obj, path, type_repr)
Save the dictionary obj; in dispatch table.
save_dict_content (obj, h5gr, subpath)
Save contents of a dictionary obj in the existing \(h 5 g r\).
The format depends on whether the dictionary obj has simple keys valid for hdf5 path components (see valid_hdf5_path_component ()) or not. For simple keys: directly use the keys as path. For nonsimple keys: save list of keys und "keys" and list of values und "values".

\section*{Parameters}
- obj (dict) - The data to be saved
- h5gr (Group) - h5py Group under which the keys and values of obj should be saved.
- subpath (str) - Name of h5gr with ' / ' in the end.

Returns type_repr - Indicates whether the data was saved in the format for a dictionary with simple keys or general keys, see comment above.

\section*{Return type REPR_DICT_SIMPLE I REPR_DICT_GENERAL}
save_range (obj, path, type_repr)
Save a range object; in dispatch table.
save_dtype (obj, path, type_repr)
Save a dtype object; in dispatch table.
save_ignored (obj, path, type_repr)
Don't save the Hdf5Ignored object; just return None.
save_global (obj, path, type_repr)
Save a global object like a function or class.

\section*{Exceptions}
\begin{tabular}{ll}
\hline Hdf5ExportError & \begin{tabular}{l} 
This exception is raised when something went wrong \\
during export to hdf5.
\end{tabular} \\
\hline Hdf5FormatError & \begin{tabular}{l} 
Common base class for errors regarding our HDF5 for- \\
mat.
\end{tabular} \\
\hline Hdf5 ImportError & \begin{tabular}{l} 
This exception is raised when something went wrong \\
during import from hdf5.
\end{tabular} \\
\hline
\end{tabular}

\subsection*{22.1.5 Hdf5ExportError}
- full name: tenpy.tools.hdf5_io.Hdf5ExportError
- parent module: tenpy.tools.hdf5_io
- type: exception
exception tenpy.tools.hdf5_io.Hdf5ExportError
This exception is raised when something went wrong during export to hdf5.

\subsection*{22.1.6 Hdf5FormatError}
- full name: tenpy.tools.hdf5_io.Hdf5FormatError
- parent module: tenpy.tools.hdf5_io
- type: exception
exception tenpy.tools.hdf5_io. Hdf5FormatError Common base class for errors regarding our HDF5 format.

\subsection*{22.1.7 Hdf5ImportError}
- full name: tenpy.tools.hdf5_io.Hdf5ImportError
- parent module: tenpy.tools.hdf5_io
- type: exception
exception tenpy.tools.hdf5_io. Hdf5ImportError
This exception is raised when something went wrong during import from hdf5.

\section*{Functions}
\begin{tabular}{ll}
\hline find_global(module, qualified_name) & \begin{tabular}{l} 
Get the object of the qualified_name in a given python \\
module.
\end{tabular} \\
\hline load(filename) & Load data from file with given filename. \\
\hline load_from_hdf5(h5group[, path, ...]) & Load an object from hdf5 file or group. \\
\hline save(data, filename[, mode]) & Save data to file with given filename. \\
\hline save_to_hdf5(h5group, obj[, path]) & Save an object obj into a hdf5 file or group. \\
\hline valid_hdf5_path_component(name) & Determine if name is a valid HDF5 path component. \\
\hline
\end{tabular}

\subsection*{22.1.8 find_global}
- full name: tenpy.tools.hdf5_io.find_global
- parent module: tenpy.tools.hdf5_io
- type: function
tenpy.tools.hdf5_io.find_global (module, qualified_name)
Get the object of the qualified_name in a given python module.

\section*{Parameters}
- module (str) - Name of the module containing the object. The module gets imported.
- qualified_name (str) - Name of the object to be retrieved. May contain dots if the object is part of a class etc.

\subsection*{22.1.9 load}
- full name: tenpy.tools.hdf5_io.load
- parent module: tenpy.tools.hdf5_io
- type: function
tenpy.tools.hdf5_io.load (filename)
Load data from file with given filename.
Guess the type of the file from the filename ending, see save () for possible endings.
Parameters filename (str) - The name of the file to load.
Returns data - The object loaded from the file.
Return type obj

\subsection*{22.1.10 load_from_hdf5}
- full name: tenpy.tools.hdf5_io.load_from_hdf5
- parent module: tenpy.tools.hdf5_io
- type: function
```

tenpy.tools.hdf5_io.load_from_hdf5(h5group, path=None, ignore_unknown=True, ex-

```

Load an object from hdf5 file or group.
Roughly equivalent to obj \(=\) h5group [path] [. . .], but handle more complicated objects saved as hdf5 groups and/or datasets with save_to_hdf5 (). For example, dictionaries are handled recursively. See Saving to disk: input/output for a specification of what can be saved/loaded and what the corresponding datastructure is.

\section*{Parameters}
- h5group (Group) - The HDF5 group (or h5py File) to be loaded.
- path (None I str I Reference) - Path within h5group to be used for loading. Defaults to the \(h 5\) group itself specified.
- ignore_unknown (b○○1) - Whether to just warn (True) or raise an Error (False) if a class to be loaded is not found.
- exclude (list of str) - List of paths (possibly relative to h5group) for objects to be excluded from loading. References to the corresponding object are replaced by an instance of Hdf5Ignored. For example, you could load a saved dictionary \{'big_data': [...], 'small_data': small_data\} with exclude=['/big_data'] to get \{'big_data': Hdf5Ignored('/big_data'), 'small_data': small_data\}. Of course, this might break other functions expecting correctly loaded data.

Returns obj - The Python object loaded from h5group (specified by path).
Return type object

\subsection*{22.1.11 save}
- full name: tenpy.tools.hdf5_io.save
- parent module: tenpy.tools.hdf5_io
- type: function
tenpy.tools.hdf5_io.save (data, filename, mode \(=\) ' \(w^{\prime}\) )
Save data to file with given filename.
This function guesses the type of the file from the filename ending. Supported endings:
\begin{tabular}{|l|l|}
\hline ending & description \\
\hline. pkl & Pickle without compression \\
\hline .pklz & Pickle with gzip compression. \\
\hline .hdf5, .h5 & HDF5 file (using h5py). \\
\hline
\end{tabular}

\section*{Parameters}
- filename (str) - The name of the file where to save the data.
- mode (str) - File mode for opening the file. ' w ' for write (discard existing file), ' a ' for append (add data to exisiting file). See open () for more details.

\subsection*{22.1.12 save_to_hdf5}
- full name: tenpy.tools.hdf5_io.save_to_hdf5
- parent module: tenpy.tools.hdf5_io
- type: function
tenpy.tools.hdf5_io.save_to_hdf5 (h5group, obj, path='/')
Save an object obj into a hdf5 file or group.
Roughly equivalent to h5group [path] = obj, but handle different types of obj. For example, dictionaries are handled recursively. See Saving to disk: input/output for a specification of what can be saved and what the resulting datastructure is.

\section*{Parameters}
- h5group (Group) - The HDF5 group (or h5py File) to which obj should be saved.
- obj (object) - The object (=data) to be saved.
- path (str) - Path within h5group under which the obj should be saved. To avoid unwanted overwriting of important data, the group/object should not yet exist, except if path is the default '/'.

Returns h5obj - The h5py group or dataset under which obj was saved.
Return type Group I Dataset

\subsection*{22.1.13 valid_hdf5_path_component}
- full name: tenpy.tools.hdf5_io.valid_hdf5_path_component
- parent module: tenpy.tools.hdf5_io
- type: function
```

tenpy.tools.hdf5_io.valid_hdf5_path_component (name)

```

Determine if name is a valid HDF5 path component.
Conditions: String, no '/', and overall name != '.'.

\section*{Module description}

Tools to save and load data (from TeNPy) to disk.

Note: This module is maintained in the repository https://github.com/tenpy/hdf5_io.git

See Saving to disk: input/output for a motivation and specification of the HDF5 format implemented below. .. online at https://tenpy.readthedocs.io/en/latest/intro/input_output.html

The functions save () and load () are convenience functions for saving and loading quite general python objects (like dictionaries) to/from files, guessing the file type (and hence protocol for reading/writing) from the file ending.

On top of that, this function provides support for saving python objects to [HDF5] files with the Hdf5Saver and Hdf5Loader classes and the wrapper functions save_to_hdf5(), load_from_hdf5().

Note: To use the export/import features to HDF5, you need to install the h5py python package (and hence some version of the HDF5 library).

Warning: Like loading a pickle file, loading data from a manipulated HDF5 file with the functions provided below has the potential to cause arbitrary code execution. Only load data from trusted sources!

\section*{Global module constants used for our HDF5 format}

Names of HDF5 attributes:
```

tenpy.tools.hdf5_io.ATTR_TYPE = 'type'

```

Attribute name for type of the saved object, should be one of the REPR_*
tenpy.tools.hdf5_io.ATTR_CLASS = 'class'
Attribute name for the class name of an HDF5Exportable
tenpy.tools.hdf5_io.ATTR_MODULE = 'module'
Attribute name for the module where ATTR_CLASS can be retrieved
tenpy.tools.hdf5_io.ATTR_LEN = 'len'
Attribute name for the length of iterables, e.g, list, tuple
tenpy.tools.hdf5_io.ATTR_FORMAT = 'format' indicates the \(A T T R \_T Y P E\) format used by Hdf5Exportable

Names for the ATTR_TYPE attribute:
```

tenpy.tools.hdf5_io.REPR_HDF5EXPORTABLE = 'instance'

```
saved object is instance of a user-defined class following the Hdf5Exportable style.
tenpy.tools.hdf5_io.REPR_ARRAY = 'array'
saved object represents a numpy array
tenpy.tools.hdf5_io.REPR_INT = 'int'
saved object represents a (python) int
tenpy.tools.hdf5_io.REPR_FLOAT = 'float'
saved object represents a (python) float
tenpy.tools.hdf5_io.REPR_STR = 'str'
saved object represents a (python unicode) string
tenpy.tools.hdf5_io.REPR_COMPLEX = 'complex'
saved object represents a complex number
tenpy.tools.hdf5_io.REPR_INT64 = 'np.int64'
saved object represents a np.int64
tenpy.tools.hdf5_io.REPR_FLOAT64 = 'np.float64'
saved object represents a np.float64
tenpy.tools.hdf5_io.REPR_INT32 = 'np.int32'
saved object represents a np.int32
tenpy.tools.hdf5_io.REPR_FLOAT32 = 'np.float32'
saved object represents a np.float 32
```

tenpy.tools.hdf5_io.REPR_BOOI = 'bool'
saved object represents a boolean
tenpy.tools.hdf5_io.REPR_NONE = 'None'
saved object is None
tenpy.tools.hdf5_io.REPR_RANGE = 'range'
saved object is a range
tenpy.tools.hdf5_io.REPR_LIST = 'list'
saved object represents a list
tenpy.tools.hdf5_io.REPR_TUPLE = 'tuple'
saved object represents a tuple
tenpy.tools.hdf5_io.REPR_SET = 'set'
saved object represents a set
tenpy.tools.hdf5_io.REPR_DICT_GENERAL = 'dict'
saved object represents a dict with complicated keys
tenpy.tools.hdf5_io.REPR_DICT_SIMPLE = 'simple_dict'
saved object represents a dict with simple keys
tenpy.tools.hdf5_io.REPR_DTYPE = 'dtype'
saved object represents a np.dtype
tenpy.tools.hdf5_io.REPR_IGNORED = 'ignore'
ignore the object/dataset during loading and saving
tenpy.tools.hdf5_io.TYPES_FOR_HDF5_DATASETS = ((<class 'numpy.ndarray'>, 'array'), (<class
tuple of (type, type_repr) which h5py can save as datasets; one entry for each type.

```

Todo: For memory caching with big MPO environments, we need a Hdf5Cacher clearing the memo's every now and then (triggered by what?).

\section*{22.2 params}
- full name: tenpy.tools.params
- parent module: tenpy.tools
- type: module

\section*{Classes}

\begin{tabular}{ll}
\hline Config(config, name) & \begin{tabular}{l} 
Dict-like wrapper class for parameter/configuration dic- \\
tionaries.
\end{tabular} \\
\hline
\end{tabular}

\section*{Functions}
\begin{tabular}{ll}
\hline asConfig(config, name) & Convert a dict-like config to a Config. \\
\hline get_parameter(params, key, default, descr[, ,.]) & \begin{tabular}{l} 
Read out a parameter from the dictionary and/or provide \\
default values.
\end{tabular} \\
\hline unused_parameters(params[, warn]) & \begin{tabular}{l} 
Returns a set of the parameters which have not been read \\
out with get_parameters.
\end{tabular} \\
\hline
\end{tabular}

\subsection*{22.2.1 asConfig}
- full name: tenpy.tools.params.asConfig
- parent module: tenpy.tools.params
- type: function
tenpy.tools.params.asConfig (config, name)
Convert a dict-like config to a config.

\section*{Parameters}
- config (dictl Config) - If this is a Config, just return it. Otherwise, create a Config from it and return that.
- name (str) - Name to be used for the Config.

Returns config - Either directly config or Config(config, name).
Return type Config

\subsection*{22.2.2 get_parameter}
- full name: tenpy.tools.params.get_parameter
- parent module: tenpy.tools.params
- type: function
tenpy.tools.params.get_parameter (params, key, default, descr, asarray=False)
Read out a parameter from the dictionary and/or provide default values.
Deprecated since version 0.6.0: Use the Config instead.
This function provides a similar functionality as params.get (key, default). Unlike dict.get this function writes the default value into the dictionary (i.e. in other words it's more similar to params. setdefault (key, default)).

This allows the user to save the modified dictionary as meta-data, which gives a concrete record of the actually used parameters and simplifies reproducing the results and restarting simulations.

Moreover, a special entry with the key 'verbose' in the params can trigger this function to also print the used value. A higer verbose level implies more output. If verbose \(>=100\), it is printed every time it's used. If verbose \(>=2\)., its printed for the first time time its used. and for verbose \(>=1\), non-default values are printed the first time they are used. otherwise only for the first use.

Internally, whether a parameter was used is saved in the set params ['_used_param']. This is used in unused_parameters () to print a warning if the key wasn't used at the end of the algorithm, to detect mis-spelled parameters.

\section*{Parameters}
- params (dict) - A dicionary of the parameters as provided by the user. If key is not a valid key, params [key] is set to default.
- key (string) - The key for the parameter which should be read out from the dictionary.
- default - The default value for the parameter.
- descr (str) - A short description for verbose output, like 'TEBD', 'XXZ_model', 'truncation'.
- asarray (bool) - If True, convert the result to a numpy array with np. asarray (. . .) before returning.
Returns params [key] if the key is in params, otherwise default. Converted to a numpy array, if asarray.
Return type value

\section*{Examples}

In the algorithm TEBDEngine gets a dictionary of parameters. Beside doing other stuff, it calls tenpy. models.model.NearestNeighborModel.calc_U_bond() with the dictionary as argument, which looks similar like:
```

>>> from tenpy.tools.params import get_parameter
>>> def model_calc_U(params):
... dt = get_parameter(params, 'dt', 0.01, 'TEBD')
... order = get_parameter(params, 'order', 1, 'TEBD')
... print("calc U with dt =", dt, "and order =", order )
... \# ... calculate exp(-i * dt* H) ....

```

Then, when you call it without any parameters, it just uses the default value:
```

>>> model_calc_U(dict())
calc U with dt = 0.01 and order = 1

```

Of course you can also provide the parameter to use a non-default value:
```

>>> model_calc_U(dict(dt=0.02))
calc U with dt = 0.02 and order = 1

```

Increasing the special keyword 'verbose' generally prints more:
```

>>> model_calc_U(dict(dt=0.02, verbose=1))
parameter 'dt'=0.02 for TEBD
calc U with dt = 0.02 and order = 1
>>> model_calc_U(dict(dt=0.02, verbose=2))
parameter 'dt'=0.02 for TEBD
parameter 'order'=1 (default) for TEBD
calc U with dt = 0.02 and order = 1

```

\subsection*{22.2.3 unused_parameters}
- full name: tenpy.tools.params.unused_parameters
- parent module: tenpy.tools.params
- type: function
tenpy.tools.params.unused_parameters (params, warn=None)
Returns a set of the parameters which have not been read out with get_parameters.
This function might be useful to check for typos in the parameter keys.
Deprecated since version 0.6.0: Use the Config instead.

\section*{Parameters}
- params (dict) - A dictionary of parameters which was given to (functions using) get_parameter()
- warn (None | str) - If given, print a warning "unused parameter for \{warn!s\}: \{unused_keys!s\}".
Returns unused_keys - The set of keys of the params which was not used
Return type set

\section*{Module description}

Tools to handle config options/paramters for algorithms.
See the doc-string of Config for details.

\section*{22.3 events}
- full name: tenpy.tools.events
- parent module: tenpy.tools
- type: module

\section*{Classes}
EventHandler
Listener
EventHandler([arg_descr]) \begin{tabular}{l} 
Handler for an event represented by an instance of this \\
class.
\end{tabular}

Listener(listener_id, callback, priority)

\subsection*{22.3.1 EventHandler}
- full name: tenpy.tools.events.EventHandler
- parent module: tenpy.tools.events
- type: class

\section*{Inheritance Diagram}

\section*{EventHandler}

\section*{Methods}
\begin{tabular}{ll}
\hline EventHandler.__init__([arg_descr]) & Initialize self. \\
\hline EventHandler.connect([callback, priority]) & Register a callback function as a listener to the event. \\
\hline EventHandler.connect_by_name(module_name, & Connect to a function given by the name in a module, \\
\(\ldots)\). & optionally inserting arguments. \\
\hline EventHandler.copy() & Make a (shallow) copy. \\
\hline EventHandler.disconnect(listener_id) & De-register a listener. \\
\hline EventHandler.emit(*args, **kwargs) & Call the callback functions of all listeners. \\
\hline \begin{tabular}{l} 
EventHandler.emit_until_result(*args, \\
**kwargs)
\end{tabular} & Call the listeners callback until one returns not None. \\
\hline
\end{tabular}

\section*{Class Attributes and Properties}
```

EventHandler.id_of_last_connected

```
```

class tenpy.tools.events.EventHandler(arg_descr=None)

```

Bases: ob ject
Handler for an event represented by an instance of this class.
All in all, events provide a flexible extension mechanism for classes to define "checkpoints" in the code where the user of a class might want to run something else, for example doing some measurements or saving intermediate results.

Parameters arg_descr (str) - An informative description how the callback function is called.
An empty string indicates no arguments.

\section*{arg_descr}

An informative description how the callback function is called.
Type str

\section*{listeners}

Entries are tuples (listener_id, callback, priority).
Type list of (int, function, int)

\section*{Examples}

Instances of this class typically get defined during class initialization and define an event. The event "happens" each time emit () or emit_until_result () is called, typically inside a method of the class defining the event. Example:
```

>>> class MyAlgorithm:
... def ___init__(self):
... self.checkpoint = EventHandler("algorithm, iteration")
... self.data = 0
... def run(self):
... for i in range(4):
... self.data += i \# do some complicated stuff
... self.checkpoint.emit(self, i)

```

Other code with access to the event can then connect a listener to the event, i.e., give a function to the event that should be called each time the event is emit ()-ed.
```

>>> my_alg = MyAlgorithm()
>>> def my_listener(algorithm, iteration):
... print("my_listener called: iteration", iteration, "with data", algorithm.
@data)
>>> my_alg.checkpoint.connect(my_listener)
<function my_listener at 0x...>
>>> my_alg.run()
my_listener called: iteration 0 with data 0
my_listener called: iteration 1 with data 1
my_listener called: iteration 2 with data 3
my_listener called: iteration 3 with data 6

```

As you can see, the function my_listener has been called during the MyAlgorithm.run () and had full access to the current status of the algorithm class. This is convenient to e.g. perform measurements of the state so far, print a status message of the progress or save intermediate results.
If the EventHandler is already initialized when you define the function, you can also use connect () as a function property like this:
```

>>> @my_alg.checkpoint.connect
... def another_one(algorithm, iteration):
... print("another_one called: iteration", iteration)
>>> @my_alg.checkpoint.connect(priority=5)
... def high_priority(algorithm, iteration):
... print("high_priority call: iteration", iteration)
>>> my_alg.run()
high_priority call: iteration 0
my_listener called: iteration 0 with data 6
another_one called: iteration 0
high_priority call: iteration 1
my_listener called: iteration 1 with data 7
another_one called: iteration 1
high_priority call: iteration 2
my_listener called: iteration 2 with data 9
another_one called: iteration 2
high_priority call: iteration 3
my_listener called: iteration 3 with data 12
another_one called: iteration 3

```
copy ()

Make a (shallow) copy.
```

connect $($ callback $=$ None, priority $=0$ )

```

Register a callback function as a listener to the event.
You can either call this function directly or use it as a function decorator, see the example in EventHandler.
If you ever plan to disconnect () again, you can read it out with id_of_last_connected right after connecting, i.e., right after calling this method.

\section*{Parameters}
- callback (callable) - A function to be called during each emit () of the event.
- priority (int) - Higher priority indicates that the callback function should be called before other possibly registered callback functions.
Returns callback - The callback function exactly as given.
Return type callable
connect_by_name (module_name, func_name, \(k w a r g s=N o n e, ~ p r i o r i t y=0) ~\)
Connect to a function given by the name in a module, optionally inserting arguments.

\section*{Parameters}
- module_name (str) - The name of the module containing the function to be used. Gets imported.
- func_name (str) - The (qualified) name of the function inside the module.
- kwargs (dict) - Optional extra keyword-arguments to be given to the function.
- priority (int) - Higher priority indicates that the callback function should be called before other possibly registered callback functions.
disconnect (listener_id)
De-register a listener.
Parameters listener_id (int) - The id of the listener, as given by id_of_last_connected right after calling connect ().
emit (*args, **kwargs)
Call the callback functions of all listeners.
Returns results - List of results returned by the individual callback functions.
Return type list
emit_until_result ( \(\left.{ }^{\arg s,}{ }^{* *} k w a r g s\right)\)
Call the listeners callback until one returns not None.

\subsection*{22.3.2 Listener}
- full name: tenpy.tools.events.Listener
- parent module: tenpy.tools.events
- type: class

\section*{Inheritance Diagram}
```

Listener

```

\section*{Methods}
\begin{tabular}{ll}
\hline Listener.__init__() & Initialize self. \\
\hline Listener.count(value, /) & Return number of occurrences of value. \\
\hline Listener. index(value[, start, stop] \()\) & Return first index of value. \\
\hline
\end{tabular}

\section*{Class Attributes and Properties}
\begin{tabular}{ll}
\hline Listener.callback & Alias for field number 1 \\
\hline Listener.listener_id & Alias for field number 0 \\
\hline Listener.priority & Alias for field number 2 \\
\hline \\
class tenpy.tools.events. Listener (listener_id, callback, priority) \\
Bases: tuple \\
property callback \\
Alias for field number 1 \\
count (value,/) \\
Return number of occurrences of value. \\
index (value, start=0, stop=9223372036854775807, /) \\
Return first index of value. \\
Raises ValueError if the value is not present. \\
property listener_id \\
Alias for field number 0 \\
property priority \\
Alias for field number 2
\end{tabular}

\section*{Module description}

Event handler.
The EventHandler is basically just holds a list of functions which can get called once a certain "event" happens. Examples are given in the class doc-string.

\section*{22.4 misc}
- full name: tenpy.tools.misc
- parent module: tenpy.tools
- type: module

\section*{Functions}
\begin{tabular}{|c|c|}
\hline add_with_None_O(a, b) & Return \(\mathrm{a}+\mathrm{b}\), treating None as zero. \\
\hline any_nonzero(params, keys[, verbose_msg]) & Check for any non-zero or non-equal entries in some parameters. \\
\hline anynan(a) & check whether any entry of a ndarray \(a\) is ' NaN '. \\
\hline argsort(a[, sort]) & wrapper around np.argsort to allow sorting ascending/descending and by magnitude. \\
\hline atleast_2d_pad(a[, pad_item]) & Transform \(a\) into a 2D array, filling missing places with pad_item. \\
\hline \multicolumn{2}{|l|}{build_initial_state(size, states, filling[,...])} \\
\hline \multicolumn{2}{|l|}{chi_list(chi_max[, dchi, nsweeps, verbose])} \\
\hline find_subclass(base_class, subclass_name) & For a given base class, recursively find the subclass with the given name. \\
\hline flatten(mapping[, separator]) & Obtain a flat dictionary with all key/value pairs of a nested data structure. \\
\hline get_close(values, target[, default, eps]) & Iterate through values and return first entry closer than eps. \\
\hline get_recursive(nested_data, recursive_key[,...]) & Extract specific value from a nested data structure. \\
\hline group_by_degeneracy(E, *args[, subset, cutoff]) & Find groups of indices for which (energy) values are degenerate. \\
\hline inverse_permutation(perm) & reverse sorting indices. \\
\hline lexsort(a[, axis]) & wrapper around \(n p\). lexsort: allow for trivial case a.shape[0] = 0 without sorting \\
\hline list_to_dict_list(l) & Given a list \(l\) of objects, construct a lookup table. \\
\hline pad(a[, w_l, v_l, w_r, v_r, axis]) & Pad an array along a given axis. \\
\hline set_recursive(nested_data, recursive_key, value) & Same as get_recursive (), but set the data entry to value. \\
\hline setup_executable(mod, run_defaults[, ...]) & Read command line arguments and turn into useable dicts. \\
\hline setup_logging([options, output_filename]) & Configure the logging module. \\
\hline to_array(a[, shape, dtype]) & Convert \(a\) to an numpy array and tile to matching dimension/shape. \\
\hline
\end{tabular}

Table 18 - continued from previous page
\begin{tabular}{|c|c|}
\hline to_iterable(a) & If \(a\) is a not iterable or a string, return [a], else return a. \\
\hline to_iterable_of_len( \(\mathrm{a}, \mathrm{L}\) ) & If a is a non-string iterable of length \(L\), return \(a\), otherwise return [a]*L. \\
\hline transpose_list_list(D[, pad]) & Returns a list of lists T , such that \(\mathrm{T}[\mathrm{i}][\mathrm{j}]=\) D[j][i]. \\
\hline ```
update_recursive(nested_data, update_data[,
...])
``` & Wrapper around set_recursive () to allow updating multiple values at once. \\
\hline zero_if_close(a[, tol]) & set real and/or imaginary part to 0 if their absolute value is smaller than tol. \\
\hline
\end{tabular}

\subsection*{22.4.1 add_with_None_0}
- full name: tenpy.tools.misc.add_with_None_0
- parent module: tenpy.tools.misc
- type: function
tenpy.tools.misc.add_with_None_0 \((a, b)\)
Return \(\mathrm{a}+\mathrm{b}\), treating None as zero.

\section*{Parameters}
- a - The two things to be added, or None.
- \(\mathbf{b}\) - The two things to be added, or None.

Returns \(a+b\), except if \(a\) or \(b\) is None, in which case the other variable is returned.

\section*{Return type sum}

\subsection*{22.4.2 any_nonzero}
- full name: tenpy.tools.misc.any_nonzero
- parent module: tenpy.tools.misc
- type: function
tenpy.tools.misc.any_nonzero (params, keys, verbose_msg=None)
Check for any non-zero or non-equal entries in some parameters.
Deprecated since version 0.8.0: This method will be removed in version 1.0.0. Use tenpy.toosl. params. Config.any_nonzero() instead.

\section*{Parameters}
- params (dict / Config)-A dictionary of parameters, or a Config instance.
- keys (list of \(\{\) key / tuple of keys \}) - For a single key, check params[key] for non-zero entries. For a tuple of keys, all the params[key] have to be equal (as numpy arrays).
- verbose_msg (None / str) - If params['verbose'] >= 1, we print verbose_msg before checking, and a short notice with the key, if a non-zero entry is found.
Returns match - False, if all params[key] are zero or None and True, if any of the params[key] for single key in keys, or if any of the entries for a tuple of keys

Return type bool

\subsection*{22.4.3 anynan}
- full name: tenpy.tools.misc.anynan
- parent module: tenpy.tools.misc
- type: function
tenpy.tools.misc.anynan (a)
check whether any entry of a ndarray \(a\) is ' NaN '.

\subsection*{22.4.4 argsort}
- full name: tenpy.tools.misc.argsort
- parent module: tenpy.tools.misc
- type: function
tenpy.tools.misc.argsort (a, sort=None, **kwargs)
wrapper around np.argsort to allow sorting ascending/descending and by magnitude.

\section*{Parameters}
- a (array_like) - The array to sort.
- sort ('m>', 'm<', '>', '<', None) - Specify how the arguments should be sorted.
\begin{tabular}{|c|c|}
\hline Sort & order \\
\hline 'm>', 'LM' & Largest magnitude first \\
\hline 'm<', 'SM' & Smallest magnitude first \\
\hline '>', 'LR', 'LA' & Largest real part first \\
\hline '<', 'SR', 'SA' & Smallest real part first \\
\hline 'LI' & Largest imaginary part first \\
\hline 'SI' & Smallest imaginary part first \\
\hline None & numpy default: same as '<' \\
\hline
\end{tabular}
- **kwargs - Further keyword arguments given directly to numpy . argsort ().

Returns index_array - Same shape as \(a\), such that a [index_array] is sorted in the specified way.
Return type ndarray, int

\subsection*{22.4.5 atleast_2d_pad}
- full name: tenpy.tools.misc.atleast_2d_pad
- parent module: tenpy.tools.misc
- type: function
tenpy.tools.misc.atleast_2d_pad (a, pad_item=0)
Transform \(a\) into a 2D array, filling missing places with pad_item.
Given a list of lists, turn it to a 2D array (pad with 0), or turn a 1D list to 2D.
Parameters a (list of lists) - to be converted into ad 2D array.
Returns a_2D - a converted into a numpy array.
Return type 2D ndarray

\section*{Examples}
```

>>> atleast_2d_pad([3, 4, 0])
array([[3, 4, 0]])

```
```

>>> atleast_2d_pad([[3, 4], [1, 6, 7]])
array([[3., 4., 0.],
[1., 6., 7.]])

```

\subsection*{22.4.6 build_initial_state}
- full name: tenpy.tools.misc.build_initial_state
- parent module: tenpy.tools.misc
- type: function
```

tenpy.tools.misc.build_initial_state(size, states, filling,mode='random', seed=None)

```

\subsection*{22.4.7 chi_list}
- full name: tenpy.tools.misc.chi_list
- parent module: tenpy.tools.misc
- type: function
```

tenpy.tools.misc.chi_list (chi_max,dchi=20,nsweeps=20,verbose=0)

```

\subsection*{22.4.8 find_subclass}
- full name: tenpy.tools.misc.find_subclass
- parent module: tenpy.tools.misc
- type: function
tenpy.tools.misc.find_subclass (base_class, subclass_name)
For a given base class, recursively find the subclass with the given name.

\section*{Parameters}
- base_class (class) - The base class of which subclass_name is supposed to be a subclass.
- subclass_name (str) - Name of the class to be found.

Returns subclass - Class with name subclass_name which is a subclass of the base_class. None, if no subclass of the given name is found.

Return type None I class

\subsection*{22.4.9 flatten}
- full name: tenpy.tools.misc.flatten
- parent module: tenpy.tools.misc
- type: function
tenpy.tools.misc.flatten (mapping, separator='/')
Obtain a flat dictionary with all key/value pairs of a nested data structure.
Parameters separator (str) - Separator for merging keys to a single string.
Returns flat_config - A single dictionary with all key-value pairs.
Return type dict

\section*{Examples}
```

>>> sample_data = {'some': {'nested': {'entry': 100, 'structure': 200},
... 'subkey': 10},
... 'topentry': 1}
>>> flat = flatten(sample_data)
>>> for k in sorted(flat):
... print(repr(k), ':', flat[k])
'some/nested/entry' : 100
'some/nested/structure' : 200
'some/subkey' : 10
'topentry' : 1

```

\section*{See also:}
get_recursive Useful to obtain a single entry from a nested data structure.

\subsection*{22.4.10 get_close}
- full name: tenpy.tools.misc.get_close
- parent module: tenpy.tools.misc
- type: function
tenpy.tools.misc.get_close (values, target, default=None, eps=le-13) Iterate through values and return first entry closer than eps.

\section*{Parameters}
- values (interable of float) - Values to compare to.
- target (float) - Value to find.
- default - Returned if no value close to target is found.
- eps (float) - Tolerance what counts as "close", namely everything with abs(val-target) < eps.
Returns value - An entry of values, if one close to target is found, otherwise default.
Return type float

\subsection*{22.4.11 get_recursive}
- full name: tenpy.tools.misc.get_recursive
- parent module: tenpy.tools.misc
- type: function
tenpy.tools.misc.get_recursive (nested_data, recursive_key, separator='/')
Extract specific value from a nested data structure.

\section*{Parameters}
- nested_data (dict of dict (-like)) - Some nested data structure supporting a dict-like interface.
- recursive_key (str) - The key(-parts) to be extracted, separated by separator. A leading separator is ignored.
- separator (str) - Separator for splitting recursive_key into subkeys.

Returns For example, recursive_key="/some/sub/key" will result in extracing nested_data["some"]["sub"]["key"].

Return type entry
See also:
set_recursive same for changing/setting a value.
flatten Get a completely flat structure.

\subsection*{22.4.12 group_by_degeneracy}
- full name: tenpy.tools.misc.group_by_degeneracy
- parent module: tenpy.tools.misc
- type: function
tenpy.tools.misc.group_by_degeneracy (E, *args, subset=None, cutoff=1e-12)
Find groups of indices for which (energy) values are degenerate.

\section*{Parameters}
- values (1D array) - Values (e.g. energies) which need to be close to count as degenerate.
- *args (1D array) - Additional vectors (with same length as values), which also need to be close (up to cutoff) to count as degenerate.
- subset (1D array) - Optionally selects a subset of the indices
- cutoff (float) - Precision up to which values still count as degenerate.

\section*{Returns}
- idx_groups (list of tuple of int) - Each tuple group contains indices i, j, k, ... for which the values are closer than cutoff, i.e., |E[j, k, ...] - E[i]| <= cutoff. Each index appears exactly once (if it is containted in subset).
- .. testsetup \(: \because\) - from tenpy.tools.misc import *
- >>> E = [2., 2.4, 1.9999, 1.8, 2.3999, 5, 1.8]
-... \#-> 0123456
- \(\ggg k=[0,1,2,2,1,2,1]\)
- >>> group_by_degeneracy \((E\), cutoff \(=0.001)\)
- \([(0,2),(1,4),(3,6),(5)\),
- >>> group_by_degeneracy(E, \(k\), cutoff=0.001) \# \(k\) and \(E\) need to be close
- \([(0),,(1,4),(2),,(3),,(5),,(6)\),

\subsection*{22.4.13 inverse_permutation}
- full name: tenpy.tools.misc.inverse_permutation
- parent module: tenpy.tools.misc
- type: function
tenpy.tools.misc.inverse_permutation (perm)
reverse sorting indices.
Sort functions (as LegCharge. sort ()) return a (1D) permutation perm array, such that sorted_array \(=\) old_array[perm]. This function inverts the permutation perm, such that old_array \(=\) sorted_array[inverse_permutation(perm)].

Parameters perm (1D array_like) - The permutation to be reversed. Assumes that it is a permutation with unique indices. If it is, inverse_permutation(inverse_permutation(perm)) == perm.

Returns inv_perm - The inverse permutation of perm such that inv_perm[perm[j]]=j= perm[inv_perm[j]].

Return type 1D array (int)

\subsection*{22.4.14 lexsort}
- full name: tenpy.tools.misc.lexsort
- parent module: tenpy.tools.misc
- type: function
tenpy.tools.misc.lexsort (a, axis=-1)
wrapper around np. lexsort: allow for trivial case a.shape [0] = 0 without sorting

\subsection*{22.4.15 list_to_dict_list}
- full name: tenpy.tools.misc.list_to_dict_list
- parent module: tenpy.tools.misc
- type: function
tenpy.tools.misc.list_to_dict_list (l)
Given a list \(l\) of objects, construct a lookup table.
This function will handle duplicate entries in \(l\).
Parameters l (iterable of iterabele of immutable) - A list of objects that can be converted to tuples to be used as keys for a dictionary.

Returns lookup - A dictionary with (key, value) pairs (key): [i1,i2,...] where i1, i2, ... are the indices where key is found in \(l\) : i.e. key \(==\) tuple(l[i1]) == tuple(l[i2]) == ...
Return type dict

\subsection*{22.4.16 pad}
- full name: tenpy.tools.misc.pad
- parent module: tenpy.tools.misc
- type: function
tenpy.tools.misc.pad ( \(a, w_{-} l=0, v_{-} l=0, w_{-} r=0, v_{-} r=0, a x i s=0\) )
Pad an array along a given axis.

\section*{Parameters}
- a (ndarray) - the array to be padded
- w_l (int) - the width to be padded in the front
- \(\mathbf{v} \_\mathbf{l}\) ( \(d t y p e\) ) - the value to be inserted before \(a\)
- w_r (int) - the width to be padded after the last index
- \(\mathbf{v} \_\mathbf{r}(d t y p e)\) - the value to be inserted after \(a\)
- axis (int) - the axis along which to pad

Returns padded - a copy of \(a\) with enlarged axis, padded with the given values.
Return type ndarray

\subsection*{22.4.17 set_recursive}
- full name: tenpy.tools.misc.set_recursive
- parent module: tenpy.tools.misc
- type: function
tenpy.tools.misc.set_recursive(nested_data, recursive_key, value, separator='/', insert_dicts=False)
Same as get_recursive (), but set the data entry to value.

\subsection*{22.4.18 setup_executable}
- full name: tenpy.tools.misc.setup_executable
- parent module: tenpy.tools.misc
- type: function
tenpy.tools.misc.setup_executable (mod, run_defaults, identifier_list=None)
Read command line arguments and turn into useable dicts.

Warning: this is a deprecated interface. Use the Simulation interface in combination with console_main() instead. You can invoce that from the command line as python \(-m\) tenpy ..

Uses default values defined at: - model class for model_par - here for sim_par - executable file for run_par Alternatively, a model_defaults dictionary and identifier_list can be supplied without the model

\section*{NB: for setup_executable to work with a model class, the model class needs to define two things:}
- defaults, a static (class level) dictionary with (key, value) pairs that have the name of the parameter (as string) as key, and the default value as value.
- identifier, a static (class level) list or other iterable with the names of the parameters to be used in filename identifiers.

\section*{Parameters}
- mod (model / dict) - Model class (or instance) OR a dictionary containing model defaults
- run_defaults (dict) - default values for executable file parameters
- identifier_list (ieterable) - Used only if mod is a dict. Contains the identifier variables

\section*{Returns}
- model_par, sim_par, run_par (dict) - containing all parameters.
- args - namespace with raw arguments for some backwards compatibility with executables.

\subsection*{22.4.19 to_array}
- full name: tenpy.tools.misc.to_array
- parent module: tenpy.tools.misc
- type: function
tenpy.tools.misc.to_array (a, shape=(None), dtype=None)
Convert \(a\) to an numpy array and tile to matching dimension/shape.
This function provides similar functionality as numpys broadcast, but not quite the same: Only scalars are broadcasted to higher dimensions, for a non-scalar, we require the number of dimension to match. If the shape does not match, we repeat periodically, e.g. we tile \((3,4) \rightarrow>(6,16)\), but \((4,4) \rightarrow(6,16)\) will raise an error.

\section*{Parameters}
- a (scalar / array_like) - The input to be converted to an array. A scalar is reshaped to the desired dimension.
- shape (tuple of \{None / int \}) - The desired shape of the array. An entry None indicates arbitrary len \(>=1\). For int entries, tile the array periodically to fit the len.
- dtype - Optionally specifies the data type.

Returns a_array - A copy of \(a\) converted to a numpy ndarray of desired dimension and shape.
Return type ndarray

\subsection*{22.4.20 to_iterable}
- full name: tenpy.tools.misc.to_iterable
- parent module: tenpy.tools.misc
- type: function
tenpy.tools.misc.to_iterable (a)
If \(a\) is a not iterable or a string, return [a], else return a.

\subsection*{22.4.21 to_iterable_of_len}
- full name: tenpy.tools.misc.to_iterable_of_len
- parent module: tenpy.tools.misc
- type: function
tenpy.tools.misc.to_iterable_of_len \((a, L)\)
If a is a non-string iterable of length \(L\), return \(a\), otherwise return \([\mathrm{a}]^{*} \mathrm{~L}\).
Raises ValueError if \(a\) is already an iterable of different length.

\subsection*{22.4.22 transpose_list_list}
- full name: tenpy.tools.misc.transpose_list_list
- parent module: tenpy.tools.misc
- type: function
tenpy.tools.misc.transpose_list_list ( \(D\), pad=None)
Returns a list of lists T, such that \(\mathrm{T}[\mathrm{i}][\mathrm{j}]=\mathrm{D}[\mathrm{j}][\mathrm{i}]\).

\section*{Parameters}
- D (list of list) - to be transposed
- pad - Used to fill missing places, if D is not rectangular.

Returns \(\mathbf{T}\) - transposed, rectangular version of \(D\). constructed such that \(T[i][j]=D[j][i]\) if i < len (D[j]) else pad
Return type list of lists

\subsection*{22.4.23 update_recursive}
- full name: tenpy.tools.misc.update_recursive
- parent module: tenpy.tools.misc
- type: function
tenpy.tools.misc.update_recursive (nested_data, update_data, separator='/', insert_dicts=True)
Wrapper around set_recursive () to allow updating multiple values at once.
It simply calls set_recursive() for each recursive_key, value in update_data.items().

\subsection*{22.4.24 zero_if_close}
- full name: tenpy.tools.misc.zero_if_close
- parent module: tenpy.tools.misc
- type: function
tenpy.tools.misc.zero_if_close ( \(a\), tol=le-15)
set real and/or imaginary part to 0 if their absolute value is smaller than tol.

\section*{Parameters}
- a (ndarray) - numpy array to be rounded
- tol (float) - the threashold which values to consider as ' 0 '.

\section*{Module description}

Miscellaneous tools, somewhat random mix yet often helpful.

\section*{22.5 math}
- full name: tenpy.tools.math
- parent module: tenpy.tools
- type: module

\section*{Functions}
\begin{tabular}{|c|c|}
\hline entropy (p[, n]) & Calculate the entropy of a distribution. \\
\hline \(\operatorname{gcd}(\mathrm{a}, \mathrm{b})\) & Computes the greatest common divisor (GCD) of two numbers. \\
\hline gcd_array(a) & Return the greatest common divisor of all of entries in \(a\) \\
\hline \(\operatorname{lcm}(\mathrm{a}, \mathrm{b})\) & Returns the least common multiple (LCM) of two positive numbers. \\
\hline matvec_to_array (H) & transform an linear operator with a matvec method into a dense numpy array. \\
\hline perm_sign(p) & Given a permutation \(p\) of numbers, returns its sign. \\
\hline qr_li(A[, cutoff]) & QR decomposition with cutoff to discard nearly linear dependent columns in \(Q\). \\
\hline rq_li(A[, cutoff]) & RQ decomposition with cutoff to discard nearly linear dependent columns in \(Q\). \\
\hline speigs(A, k, *args, **kwargs) & Wrapper around scipy.sparse.linalg. eigs(), lifting the restriction \(k<\operatorname{rank}(A)-1\). \\
\hline speigsh(A, k, *args, **kwargs) & Wrapper around scipy.sparse.linalg. eigsh(), lifting the restriction \(k<r a n k(A)-1\). \\
\hline
\end{tabular}

\subsection*{22.5.1 entropy}
- full name: tenpy.tools.math.entropy
- parent module: tenpy.tools.math
- type: function
```

tenpy.tools.math.entropy ( }p,n=1\mathrm{ )

```

Calculate the entropy of a distribution.
Assumes that \(p\) is a normalized distribution ( \(n \mathrm{p} . \operatorname{sum}(\mathrm{p})==1\).).

\section*{Parameters}
- p (1D array) - A normalized distribution.
- n(1 | float | np.inf)-Selects the entropy, see below.

Returns entropy - Shannon-entropy \(-\sum_{i} p_{i} \log \left(p_{i}\right)(\mathrm{n}=1)\) or Renyi-entropy \(\frac{1}{1-n} \log \left(\sum_{i} p_{i}^{n}\right)(\mathrm{n}!=\) 1) of the distribution \(p\).

Return type float

\section*{22.5 .2 gcd}
- full name: tenpy.tools.math.gcd
- parent module: tenpy.tools.math
- type: function
tenpy.tools.math.gcd \((a, b)\)
Computes the greatest common divisor (GCD) of two numbers.
Return 0 if both \(\mathrm{a}, \mathrm{b}\) are zero, otherwise always return a non-negative number.

\subsection*{22.5.3 gcd_array}
- full name: tenpy.tools.math.gcd_array
- parent module: tenpy.tools.math
- type: function
tenpy.tools.math.gcd_array (a)
Return the greatest common divisor of all of entries in \(a\)

\subsection*{22.5.4 Icm}
- full name: tenpy.tools.math.lcm
- parent module: tenpy.tools.math
- type: function
tenpy.tools.math.lcm \((a, b)\)
Returns the least common multiple (LCM) of two positive numbers.

\subsection*{22.5.5 matvec_to_array}
- full name: tenpy.tools.math.matvec_to_array
- parent module: tenpy.tools.math
- type: function
tenpy.tools.math.matvec_to_array \((H)\)
transform an linear operator with a matvec method into a dense numpy array.
Parameters H (linear operator) - should have shape, dtype attributes and a matvec method.
Returns H_dense - a dense array version of \(H\).
Return type ndarray, shape (H.dim, H.dim)

\subsection*{22.5.6 perm_sign}
- full name: tenpy.tools.math.perm_sign
- parent module: tenpy.tools.math
- type: function
tenpy.tools.math.perm_sign \((p)\)
Given a permutation \(p\) of numbers, returns its sign. ( +1 or -1 )
Assumes that all the elements are distinct, if not, you get crap.

\section*{Examples}
```

>>> import itertools
>>> for p in itertools.permutations(range(3)):
... print('{p!s}: {sign!s}'.format(p=p, sign=tenpy.tools.math.perm_sign(p)))
(0, 1, 2): 1
(0, 2, 1): -1
(1, 0, 2): -1
(1, 2, 0): 1
(2, 0, 1): 1
(2, 1, 0): -1

```

\subsection*{22.5.7 qr_li}
- full name: tenpy.tools.math.qr_li
- parent module: tenpy.tools.math
- type: function
```

tenpy.tools.math.qr_li (A, cutoff=1e-15)

```

QR decomposition with cutoff to discard nearly linear dependent columns in \(Q\).
Perform a QR decomposition with pivoting, discard columns where R[i,i] < cuttoff, reverse the permututation from pivoting and perform another QR decomposition to ensure that \(R\) is upper right.

Parameters A (numpy.ndarray) - Matrix to be decomposed as \(A=Q . R\)
Returns \(\mathbf{Q}, \mathbf{R}\) - Decomposition of \(A\) into isometry \(Q^{\wedge} d Q=1\) and upper right \(R\) with diagonal entries larger than cutoff.

Return type numpy.ndarray

\subsection*{22.5.8 rq_li}
- full name: tenpy.tools.math.rq_li
- parent module: tenpy.tools.math
- type: function
tenpy.tools.math.rq_li (A, cutoff=le-15)
RQ decomposition with cutoff to discard nearly linear dependent columns in \(Q\).
Uses qr_li () on tranpose of \(A\). Note that \(R\) is nonzero in the lowest left corner; \(R\) has entries below the diagonal for non-square \(R\).

Parameters A (numpy. ndarray) - Matrix to be decomposed as \(A=Q \cdot R\)
Returns \(\mathbf{R}, \mathbf{Q}\) - Decomposition of \(A\) into isometry \(Q Q^{\wedge} d=l\) and upper right \(R\) with diagonal entries larger than cutoff. If \(M, N=A\). shape, then \(R\). shape \(=M, K\) and Q. shape \(=K, N\) with \(K<=\min (M, N)\).

Return type numpy. ndarray

\subsection*{22.5.9 speigs}
- full name: tenpy.tools.math.speigs
- parent module: tenpy.tools.math
- type: function
tenpy.tools.math.speigs ( \(A, k\), *args, **kwargs)
Wrapper around scipy.sparse.linalg.eigs(), lifting the restriction \(k<r a n k(A)-1\).

\section*{Parameters}
- A (MxM ndarray or like scipy.sparse.linalg.Linearoperator) - the (square) linear operator for which the eigenvalues should be computed.
- \(\mathbf{k}(i n t)\) - the number of eigenvalues to be computed.
- *args - Further arguments directly given to scipy.sparse.linalg.eigs()
- **kwargs - Further keyword arguments directly given to scipy.sparse.linalg. eigs()

\section*{Returns}
- \(\mathbf{w}\) (ndarray) - array of \(\min (k\), A.shape[0]) eigenvalues
- \(\mathbf{v}\) (ndarray) - array of \(\min (k\), A.shape[0]) eigenvectors, \(\mathrm{v}[:, ~ i]\) is the \(i\)-th eigenvector. Only returned if kwargs ['return_eigenvectors'] == True.

\subsection*{22.5.10 speigsh}
- full name: tenpy.tools.math.speigsh
- parent module: tenpy.tools.math
- type: function
tenpy.tools.math.speigsh ( \(A, k\), *args, **kwargs)
Wrapper around scipy.sparse.linalg.eigsh(), lifting the restriction \(k<r a n k(A)-1\).

\section*{Parameters}
- A (MxM ndarray or like scipy.sparse.linalg. LinearOperator) - The (square) hermitian linear operator for which the eigenvalues should be computed.
- \(\mathbf{k}\) (int) - The number of eigenvalues to be computed.
- *args - Further arguments directly given to scipy. sparse.linalg.eigsh().
- **kwargs - Further keyword arguments directly given to scipy.sparse.linalg. eigsh().

\section*{Returns}
- w (ndarray) - Array of \(\min (k\), A.shape[0]) eigenvalues.
- \(\mathbf{v}\) (ndarray) - Array of \(\min (k\), A.shape[0]) eigenvectors, \(\mathrm{v}[:, ~ i]\) is the \(i\)-th eigenvector. Only returned if kwargs['return_eigenvectors'] == True.

\section*{Module description}

Different math functions needed at some point in the library.
```

tenpy.tools.math.LeviCivita3 = array([[[ 0, 0, 0], [ 0, 0, 1], [ 0, -1, 0]], [[ 0, 0, -1],

```
    Levi-Civita Symbol of int type

\section*{22.6 fit}
- full name: tenpy.tools.fit
- parent module: tenpy.tools
- type: module

\section*{Functions}
\begin{tabular}{|c|c|}
\hline alg_decay (x, a, b, c) & define the algebraic decay. \\
\hline alg_decay_fit(x, y[, npts, power_range, ...]) & Fit y to the form \(\mathrm{a} * \mathrm{x} * *(-\mathrm{b})+\mathrm{c}\). \\
\hline alg_decay_fit_res(log_b, x, y) & Returns the residue of an algebraic decay fit of the form x** (-np.exp (log_b)). \\
\hline alg_decay_fits(x, ys[, npts, power_range, ...]) & Fit arrays of y's to the form a * \(\mathrm{x}^{* *}(-\mathrm{b})+\mathrm{c}\). \\
\hline ```
central_charge_from_S_profile(psi[,
exclude])
``` & Fit the entanglement entropy of a finite MPS to the expected profile for critical models. \\
\hline entropy_profile_from_CFT(size_A, L, ...) & Expected profile for the entanglement entropy at a critical point. \\
\hline fit_with_sum_of_exp(f, \(\mathrm{n}[\mathrm{N}, \mathrm{N}])\) & Approximate a decaying function \(f\) with a sum of exponentials. \\
\hline lin_fit_res(x, y) & Returns the least-square residue of a linear fit y vs x . \\
\hline linear_fit(x, y) & Perform a linear fit of y to \(\mathrm{ax}+\mathrm{b}\). \\
\hline plot_alg_decay_fit(plot_module, \(\mathbf{x}\), \(\mathbf{y}\), fit_par) & Given \(x, y\), and fit_par (output from alg_decay_fit), produces a plot of the algebraic decay fit. \\
\hline sum_of_exp(lambdas, prefactors, \(\mathbf{x}\) ) & Evaluate sum_i prefactor[i] *
lambda[i]**x for different \(x\). \\
\hline
\end{tabular}

\subsection*{22.6.1 alg_decay}
- full name: tenpy.tools.fit.alg_decay
- parent module: tenpy.tools.fit
- type: function
tenpy.tools.fit.alg_decay \((x, a, b, c)\) define the algebraic decay.

\subsection*{22.6.2 alg_decay_fit}
- full name: tenpy.tools.fit.alg_decay_fit
- parent module: tenpy.tools.fit
- type: function
tenpy.tools.fit.alg_decay_fit ( \(x, y, n p t s=5\), power_range=(0.01, 4.0), power_mesh=[60, 10])
Fit \(y\) to the form \(a * x * *(-b)+c\).
Returns a triplet \([\mathrm{a}, \mathrm{b}, \mathrm{c}]\).
npts specifies the maximum number of points to fit. If npts < len(x), then alg_decay_fit() will only fit to the last npts points. power_range is a tuple that gives that restricts the possible ranges for b. power_mesh is a list of numbers, which specifies how fine to search for the optimal b. E.g., if power_mesh \(=[60,10]\), then it'll first divide the power_range into 60 intervals, and then divide those intervals by 10 .

\subsection*{22.6.3 alg_decay_fit_res}
- full name: tenpy.tools.fit.alg_decay_fit_res
- parent module: tenpy.tools.fit
- type: function
tenpy.tools.fit.alg_decay_fit_res \(\left(\log _{2} b, x, y\right)\)
Returns the residue of an algebraic decay fit of the form \(\mathrm{x} * *\left(-\mathrm{np} . \exp \left(\log \_\mathrm{b}\right)\right)\).

\subsection*{22.6.4 alg_decay_fits}
- full name: tenpy.tools.fit.alg_decay_fits
- parent module: tenpy.tools.fit
- type: function
tenpy.tools.fit.alg_decay_fits \((x, y s, n p t s=5\), power_range=(0.01, 4.0), power_mesh=[60, 10])
Fit arrays of \(y\) 's to the form \(\mathrm{a} * \mathrm{x}^{* *}(\mathrm{~b})+\mathrm{c}\).
Returns arrays of \([a, b, c]\).

\subsection*{22.6.5 central_charge_from_S_profile}
- full name: tenpy.tools.fit.central_charge_from_S_profile
- parent module: tenpy.tools.fit
- type: function
tenpy.tools.fit.central_charge_from_S_profile (psi, exclude=None)
Fit the entanglement entropy of a finite MPS to the expected profile for critical models.
See entropy_profile_from_CFT () for the function we fit to.

\section*{Parameters}
- psi (MPS) - Ground state of a finite system at a critical point (i.e. gapless!). The bond dimension should be large enough to be converged!
- exclude (int) - How many sites at the left (and at the right) boundary to exclude from the fit (to avoid boundary effects). Defaults to psi.L // 4

\section*{Returns}
- central_charge, const (float) - Central charge and constant offset as in entropy_profile_from_CFT().
- res \((\) float \()\) - Residuum of the error.

\subsection*{22.6.6 entropy_profile_from_CFT}
- full name: tenpy.tools.fit.entropy_profile_from_CFT
- parent module: tenpy.tools.fit
- type: function
```

tenpy.tools.fit.entropy_profile_from_CFT(size_A,L,central_charge,const)

```

Expected profile for the entanglement entropy at a critical point.
Conformal field theory predicts the entanglement entropy for cutting a ground state of a finite, critical (i.e. gapless) system of length \(L\) into the left \(l\) and right \(L-l\) sites to be (eq. 2 of [[calabrese2004]]):
\[
S(l, L)=\frac{c}{6} \log \left(\frac{2 L}{\pi a} \sin \left(\frac{\pi l}{L}\right)\right)+\text { const }
\]

Here, \(c\) is the central charge of the system, and \(a\) is the lattice spacing, which we set to 1 , and const is a non-universal constant.

Returns exactly that formula.

\subsection*{22.6.7 fit_with_sum_of_exp}
- full name: tenpy.tools.fit.fit_with_sum_of_exp
- parent module: tenpy.tools.fit
- type: function
```

tenpy.tools.fit.fit_with_sum_of_exp (f,n,N=50)

```

Approximate a decaying function \(f\) with a sum of exponentials.
MPOs can naturally represent long-range interactions with an exponential decay. A common technique for other (e.g. powerlaw) long-range interactions is to approximate them by sums of exponentials and to include them into the MPOs. This funciton allows to do that.

The algorithm/implementation follows the appendix of [[murg2010]].

\section*{Parameters}
- \(\mathbf{f}(\) function \()-\) Decaying function to be approximated. Needs to accept a 1D numpy array \(x\)
- \(\mathbf{n}(i n t)\) - Number of exponentials to be used.
- \(\mathbf{N}(i n t)\) - Number of points at which to evaluate/fit \(f\); we evaluate and fit \(f\) at the points x = np.arange (1, N+1).

Returns lambdas, prefactors - Such that \(f(k) \approx \sum_{i} x_{i} \lambda_{i}^{k}\) for (integer) \(1<=k<=N\). The function sum_of_exp () evaluates this for given \(x\).
Return type 1D arrays

\subsection*{22.6.8 lin_fit_res}
- full name: tenpy.tools.fit.lin_fit_res
- parent module: tenpy.tools.fit
- type: function
tenpy.tools.fit.lin_fit_res \((x, y)\)
Returns the least-square residue of a linear fit y vs x .

\subsection*{22.6.9 linear_fit}
- full name: tenpy.tools.fit.linear_fit
- parent module: tenpy.tools.fit
- type: function
tenpy.tools.fit.linear_fit ( \(x, y\) )
Perform a linear fit of \(y\) to \(a x+b\).
Returns \(a, b\), res.

\subsection*{22.6.10 plot_alg_decay_fit}
- full name: tenpy.tools.fit.plot_alg_decay_fit
- parent module: tenpy.tools.fit
- type: function
tenpy.tools.fit.plot_alg_decay_fit(plot_module, x, y, fit_par, xfunc=None, kwargs=\{\}, plot_fit_args=\{\})
Given \(x, y\), and fit_par (output from alg_decay_fit), produces a plot of the algebraic decay fit.
plot_module is matplotlib.pyplot, or a subplot. x , y are the data (real, 1-dimensional np.ndarray) fit_par is a triplet of numbers [a, b, c] that describes and algebraic decay (see alg_decay()). xfunc is an optional parameter that scales the x -axis in the resulting plot. kwargs is a dictionary, whoses key/items are passed to the plot function. plot_fit_args is a dictionary that controls how the fit is shown.

\subsection*{22.6.11 sum_of_exp}
- full name: tenpy.tools.fit.sum_of_exp
- parent module: tenpy.tools.fit
- type: function
tenpy.tools.fit.sum_of_exp (lambdas, prefactors, \(x\) )
Evaluate sum_i prefactor[i] * lambda[i]**x for different \(x\).
See fit_sum_of_exp () for more details.

\section*{Module description}
tools to fit to an algebraic decay.

\section*{22.7 string}
- full name: tenpy.tools.string
- parent module: tenpy.tools
- type: module

\section*{Functions}
\begin{tabular}{ll}
\hline is_non_string_iterable(x) & \begin{tabular}{l} 
Check if x is a non-string iterable, (e.g., list, tuple, dic- \\
tionary, np.ndarray)
\end{tabular} \\
\hline to_mathematica_lists(a) & \begin{tabular}{l} 
convert nested \(a\) to string readable by mathematica us- \\
ing curly brackets ' \(\{\ldots\}\) '.
\end{tabular} \\
\hline vert_join(strlist[, valign, halign, delim]) & \begin{tabular}{l} 
Join strings with multilines vertically such that they ap- \\
pear next to each other.
\end{tabular} \\
\hline
\end{tabular}

\subsection*{22.7.1 is_non_string_iterable}
- full name: tenpy.tools.string.is_non_string_iterable
- parent module: tenpy.tools.string
- type: function
```

tenpy.tools.string.is_non_string_iterable(x)

```

Check if x is a non-string iterable, (e.g., list, tuple, dictionary, np.ndarray)

\subsection*{22.7.2 to_mathematica_lists}
- full name: tenpy.tools.string.to_mathematica_lists
- parent module: tenpy.tools.string
- type: function
tenpy.tools.string.to_mathematica_lists \((a)\)
convert nested \(a\) to string readable by mathematica using curly brackets ' \(\{\ldots\}\) '.

\subsection*{22.7.3 vert_join}
- full name: tenpy.tools.string.vert_join
- parent module: tenpy.tools.string
- type: function
tenpy.tools.string.vert_join (strlist, valign='t', halign='l', delim=' ')
Join strings with multilines vertically such that they appear next to each other.

\section*{Parameters}
- strlist (list of str) - the strings to be joined vertically
- valing ('t', 'c', 'b') - vertical alignment of the strings: top, center, or bottom
- halign ('l', 'c', 'r')-horizontal alignment of the strings: left, center, or right
- delim (str) - field separator between the strings

Returns joined - a string where the strings of strlist are aligned vertically
Return type str

\section*{Examples}
```

>>> from tenpy.tools.string import vert_join
>>> print(vert_join(['a\nsample\nmultiline\nstring', str(np.arange(9).reshape(3, ь
->3))],
\cdots. delim=' | '))
a | [l[llll
sample | [llll}
multiline | [ll 7 8]]
string |

```

\section*{Module description}

Tools for handling strings.

\section*{22.8 process}
- full name: tenpy.tools.process
- parent module: tenpy.tools
- type: module

\section*{Functions}
\begin{tabular}{ll}
\hline load_omp_library([libs, verbose]) & Tries to load openMP library. \\
\hline memory_usage () & Return memory usage of the running python process. \\
\hline mkl_get_nthreads () & wrapper around MKL get_max_threads. \\
\hline mkI_set_nthreads \((\mathrm{n})\) & wrapper around MKL set_num_threads. \\
\hline omp_get_nthreads () & wrapper around OpenMP get_max_threads. \\
\hline omp_set_nthreads \((\mathrm{n})\) & wrapper around OpenMP set_nthreads. \\
\hline
\end{tabular}

\subsection*{22.8.1 load_omp_library}
- full name: tenpy.tools.process.load_omp_library
- parent module: tenpy.tools.process
- type: function
tenpy.tools.process.load_omp_library (libs=['libiomp5.so', None, 'libgomp.so.l'], verbose \(=\) True )
Tries to load openMP library.

\section*{Parameters}
- libs - list of possible library names we should try to load (with ctypes.CDLL).
- verbose ( \(b \circ \circ 1\) ) - wheter to print the name of the loaded library.

Returns omp - OpenMP shared libary if found, otherwise None. Once it was sucessfully imported, no re-imports are tried.

Return type CDLL I None

\subsection*{22.8.2 memory_usage}
- full name: tenpy.tools.process.memory_usage
- parent module: tenpy.tools.process
- type: function

\section*{tenpy.tools.process.memory_usage()}

Return memory usage of the running python process.
You can pip install psutil if you get only \(-1 .\).
Returns mem - Currently used memory in megabytes. - 1 . if no way to read out.
Return type float

\subsection*{22.8.3 mkl_get_nthreads}
- full name: tenpy.tools.process.mkl_get_nthreads
- parent module: tenpy.tools.process
- type: function
tenpy.tools.process.mkl_get_nthreads ()
wrapper around MKL get_max_threads.
Returns max_threads - The maximum number of threads used by MKL. - 1 if unable to read out.
Return type int

\subsection*{22.8.4 mkl_set_nthreads}
- full name: tenpy.tools.process.mkl_set_nthreads
- parent module: tenpy.tools.process
- type: function
tenpy.tools.process.mkl_set_nthreads ( \(n\) )
wrapper around MKL set_num_threads.
Parameters \(\mathbf{n}(i n t)\) - the number of threads to use
Returns success - whether the shared library was found and set.
Return type bool

\subsection*{22.8.5 omp_get_nthreads}
- full name: tenpy.tools.process.omp_get_nthreads
- parent module: tenpy.tools.process
- type: function
tenpy.tools.process.omp_get_nthreads () wrapper around OpenMP get_max_threads.

Returns max_threads - The maximum number of threads used by OpenMP (and thus MKL). -1 if unable to read out.

Return type int

\subsection*{22.8.6 omp_set_nthreads}
- full name: tenpy.tools.process.omp_set_nthreads
- parent module: tenpy.tools.process
- type: function
tenpy.tools.process.omp_set_nthreads ( \(n\) )
wrapper around OpenMP set_nthreads.
Parameters \(\mathbf{n}\) (int) - the number of threads to use
Returns success - whether the shared library was found and set.

Return type bool

\section*{Module description}

Tools to read out total memory usage and get/set the number of threads.
If your python is compiled against MKL (e.g. if you use anaconda as recommended in INSTALL), it will by default use as many threads as CPU cores are available. If you run a job on a cluster, you should limit this to the number of cores you reserved - otherwise your colleagues might get angry... A simple way to achieve this is to set a suitable enviornment variable before calling your python program, e.g. on the linux bash export OMP_NUM_THREADS=4 for 4 threads. (MKL used OpenMP and thus respects its settings.)

Alternatively, this module provides omp_get_nthreads () and omp_set_nthreads (), which give their best to get and set the number of threads at runtime, while still being failsave if the shared OpenMP library is not found. In the latter case, you might also try the equivalent \(m k l_{\text {_ }}\) get_nthreads () and mkI_set_nthreads ().

\section*{22.9 optimization}
- full name: tenpy.tools.optimization
- parent module: tenpy.tools
- type: module

\section*{Classes}

\begin{tabular}{ll}
\hline OptimizationFlag(value) & \begin{tabular}{l} 
Options for the global 'optimization level' used for dy- \\
namical optimizations.
\end{tabular} \\
\hline temporary_level(temporary_level) & \begin{tabular}{l} 
Context manager to temporarily set the optimization \\
level to a different value.
\end{tabular} \\
\hline
\end{tabular}

\subsection*{22.9.1 OptimizationFlag}
- full name: tenpy.tools.optimization.OptimizationFlag
- parent module: tenpy.tools.optimization
- type: class

\section*{Inheritance Diagram}


\section*{Class Attributes and Properties}
```

    OptimizationFlag.default
    OptimizationFlag.none
    OptimizationFlag.safe
    OptimizationFlag.skip_arg_checks
    class tenpy.tools.optimization.OptimizationFlag(value)
Bases: enum.IntEnum
Options for the global 'optimization level' used for dynamical optimizations.
Whether we optimize dynamically is decided by comparison of the global "optimization level" with one of the following flags. A higher level includes all the previous optimizations.

```
\begin{tabular}{|l|l|l|}
\hline Levell Flag & Description \\
\hline 0 & none & Don't do any optimizations, i.e., run many sanity checks. Used for testing. \\
\hline 1 & default & \begin{tabular}{l} 
Skip really unnecessary sanity checks, but also don't try any optional optimizations if \\
they might give an overhead.
\end{tabular} \\
\hline 2 & safe & \begin{tabular}{l} 
Activate safe optimizations in algorithms, even if they might give a small overhead. Ex- \\
ample: Try to compress the MPO representing the hamiltonian.
\end{tabular} \\
\hline 3 & skip_arg_chellnsafe! Skip (some) class sanity tests and (function) argument checks. \\
\hline
\end{tabular}

Warning: When unsafe optimizations are enabled, errors will not be detected that easily, debugging is much harder, and you might even get segmentation faults in the compiled parts. Use this kind of optimization only for code which you succesfully ran before with (very) similar parmeters and disabled optimiztions! Enable this optimization only during the parts of the code where it is really necessary. Check whether it actually helps - if it doesn't, keep the optimization disabled!

\subsection*{22.9.2 temporary_level}
- full name: tenpy.tools.optimization.temporary_level
- parent module: tenpy.tools.optimization
- type: class

\section*{Inheritance Diagram}
temporary_level

\section*{Methods}
temporary_level.__init__(temporary_level) Initialize self.
class tenpy.tools.optimization.temporary_level (temporary_level)
Bases: object
Context manager to temporarily set the optimization level to a different value.
Parameters temporary_level (int | OptimizationFlag | str | None) - The optimization level to be set during the context. None defaults to the current value of the optimization level.

\section*{temporary_level}

The optimization level to be set during the context.
Type None I OptimizationFlag

\section*{_old_level}

Optimization level to be restored at the end of the context manager.

\section*{Type OptimizationFlag}

\section*{Examples}

It is recommended to use this context manager in a with statement:
```


# optimization level default

with temporary_level(OptimizationFlag.safe):
do_some_stuff() \# temporarily have Optimization level `safe`
\# you can even change the optimization level to something else:
set_level(OptimizationFlag.skip_args_check)
do_some_really_heavy_stuff()

# here we are back to the optimization level as before the ``with ... `` statement

```

\section*{Functions}
\begin{tabular}{ll}
\hline get_level() & Return the global optimization level. \\
\hline optimize([level_compare]) & \begin{tabular}{l} 
Called by algorithms to check whether it should (try to) \\
do some optimizations.
\end{tabular} \\
\hline set_level([level]) & Set the global optimization level. \\
\hline to_OptimizationFlag(level) & Convert strings and int to a valid OptimizationFlag. \\
\hline use_cython([func, replacement, check_doc]) & \begin{tabular}{l} 
Decorator to replace a function with a Cython- \\
equivalent from_npc_helper.pyx.
\end{tabular} \\
\hline
\end{tabular}

\subsection*{22.9.3 get_level}
- full name: tenpy.tools.optimization.get_level
- parent module: tenpy.tools.optimization
- type: function
tenpy.tools.optimization.get_level()
Return the global optimization level.

\subsection*{22.9.4 optimize}
- full name: tenpy.tools.optimization.optimize
- parent module: tenpy.tools.optimization
- type: function
tenpy.tools.optimization.optimize (level_compare \(=<\) OptimizationFlag.default: \(1>\) )
Called by algorithms to check whether it should (try to) do some optimizations.
Parameters level_compare (OptimizationFlag) - At which level to start optimization, i.e., how safe the suggested optimization is.

Returns optimize - True if the algorithms should try to optimize, i.e., whether the global "optimization level" is equal or higher than the level to compare to.

Return type bool

\subsection*{22.9.5 set_level}
- full name: tenpy.tools.optimization.set_level
- parent module: tenpy.tools.optimization
- type: function
tenpy.tools.optimization.set_level (level=1)
Set the global optimization level.
Parameters level (int | OptimizationFlag | str | None) - The new global optimization level to be set. None defaults to keeping the current level.

\subsection*{22.9.6 to_OptimizationFlag}
- full name: tenpy.tools.optimization.to_OptimizationFlag
- parent module: tenpy.tools.optimization
- type: function
tenpy.tools.optimization.to_OptimizationFlag(level)
Convert strings and int to a valid OptimizationFlag.
None defaults to the current level.

\subsection*{22.9.7 use_cython}
- full name: tenpy.tools.optimization.use_cython
- parent module: tenpy.tools.optimization
- type: function
```

tenpy.tools.optimization.use_cython(func=None, replacement=None, check_doc=True)

```

Decorator to replace a function with a Cython-equivalent from _npc_helper.pyx.
This is a decorator, which is supposed to be used in front of function definitions with an @ sign, for example:
```

@use_cython
def my_slow_function(a):
"some example function with slow python loops"
result = 0.
for i in range(a.shape[0]):
for j in range(a.shape[1]):
\#... heavy calculations ...
result += np.cos(a[i, j]**2) * (i + j)
return result

```

This decorator indicates that there is a Cython implementation in the file tenpy/linalg/_npc_helper. pyx, which should have the same signature (i.e. same arguments and return values) as the decorated function, and can be used as a replacement for the decorated function. However, if the cython code could not be compiled on your system (or if the environment variable TENPY_OPTIMIZE is set to negative values, or the environment variable TENPY_NO_CYTHON is "true"), we just pass the previous function.

Note: In case that the decorator is used for a class method, the corresponding Cython version needs to have an @cython.binding(True).

\section*{Parameters}
- func (function) - The defined function
- replacement (string / None) - The name of the function defined in tenpy/ linalg/_npc_helper.pyx which should replace the decorated function. None defaults to the name of the decorated function, e.g., in the above example my_slow_function.
- check_doc (bool) - If True, we check that the cython version of the function has the exact same doc string (up to a possible first line containing the function signature) to exclude typos and inconsistent versions.
Returns replacement_func - The function replacing the decorated function func. If the cython code can not be loaded, this is just func, otherwise it's the cython version specified by replacement.

\section*{Return type function}

\section*{Module description}

Optimization options for this library.
Let me start with a quote of "Micheal Jackson" (a programmer, not the musician):
```

First rule of optimization: "Don't do it."
Second rule of optimization (for experts only): "Don't do it yet."
Third rule of optimization: "Profile before optimizing."

```

Luckily, following the third optimization rule, namely profiling code, is fairly simple in python, see the documentation. If you have a python skript running your code, you can simply call it with python -m "cProfile" -s "tottime" your_skript.py. Alternatively, save the profiling statistics with python -m "cProfile" -o "profile_data.stat" your_skript.py and run these few lines of python code:
```

import pstats
p = pstats.Pstats("profile_data.stat")
p.sort_stats('cumtime') \# sort by 'cumtime' column
p.print_stats(30) \# prints first 30 entries

```

That being said, I actually did profile and optimize (parts of) the library; and there are a few knobs you can turn to tweak the most out of this library, explained in the following.
1) Simply install the 'bottleneck' python package, which allows to optimize slow parts of numpy, most notably ' NaN ' checking.
2) Figure out which numpy/scipy/python you are using. As explained in Installation instructions, we recommend to use the Python distributed provided by Intel or Anaconda. They ship with numpy and scipy which use Intels MKL library, such that e.g. np.tensordot is parallelized to use multiple cores.
3) In case you didn't do that yet: some parts of the library are written in both python and Cython with the same interface, so you can simply compile the Cython code, as explained in Installation instructions. Then everything should work the same way from a user perspective, while internally the faster, pre-compiled cython code from tenpy/linalg/_npc_helper.pyx is used. This should also be a safe thing to do. The replacement of the optimized functions is done by the decorator use_cython ().

Note: By default, the compilation will link against the BLAS functions provided by scipy.linalg. cython_blas. Whether they use MKL depends on the scipy version you installed. However, you can explicitly link against a given MKL by providing the path during compilation, as explained in Compile linking agains MKL.
4) One of the great things about python is its dynamical nature - anything can be done at runtime. In that spirit, this module allows to set a global "optimization level" which can be changed dynamically (i.e., during runtime) with set_level (). The library will then try some extra optimiztion, most notably skip sanity checks of arguments. The possible choices for this global level are given by the Optimizationflag. The default initial value for the global optimization level can be adjusted by the environment variable TENPY_OPTIMIZE.

Warning: When this optimizing is enabled, we skip (some) sanity checks. Thus, errors will not be detected that easily, and debugging is much harder! We recommend to use this kind of optimization only for code which you succesfully have run before with (very) similar parmeters! Enable this optimization only during the parts of the code where it is really necessary. The context manager temporary_level can help with that. Check whether it actually helps - if it doesn't, keep the optimization disabled! Some parts of the library already do that as well (e.g. DMRG after the first sweep).
5) You might want to try some different compile time options for the cython code, set in the setup.py in the top directory of the repository. Since the setup.py reads out the TENPY_OPTIMIZE environment variable, you can simple use an export TENPY_OPTIMIZE=3 (in your bash/terminal) right before compilation. An export TENPY_OPTIMIZE=0 activates profiling hooks instead.

Warning: This increases the probability of getting segmentation faults and anyway might not help that much; in the crucial parts of the cython code, these optimizations are already applied. We do not recommend using this!
tenpy.tools.optimization.bottleneck \(=\) None
tenpy.tools.optimization.have_cython_functions = False
bool whether the import of the cython file tenpy/linalg/_npc_helper.pyx succeeded.
The value is set in the first call of use_cython().
tenpy.tools.optimization.compiled_with_MKL = False
bool whether the cython file was compiled with \(H A V E \_M K L\).
The value is set in the first call of use_cython ().

\section*{VERSION}
- full name: tenpy.version
- parent module: tenpy
- type: module

\section*{Module description}

Access to version of this library.
The version is provided in the standard python format major.minor.revision as string. Use pkg_resources.parse_version before comparing versions.
tenpy.version.version \(=\) '0.8.0'
current release version as a string
tenpy.version.released \(=\) True
whether this is a released version or modified
tenpy.version.short_version = 'v0.8.0'
same as version, but with ' \(v\) ' in front
tenpy.version.git_revision = 'e1a0b3df552a8f9599cb66ef8cf356e49b0dcc49'
the hash of the last git commit (if available)
tenpy.version.full_version = '0.8.0'
if not released additional info with part of git revision
tenpy.version.version_summary = 'tenpy 0.8.0 (not compiled), \ngit revision e1a0b3df552a8f9! summary of the tenpy, python, numpy and scipy versions used

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[TeNPySource] https://github.com/tenpy/tenpy
[TeNPyNotebooks] Collection of example [jupyter] notebooks using TeNPy: https://github.com/tenpy/tenpy_ notebooks
[TeNPyDoc] Online documentation, https://tenpy.readthedocs.io/
[TeNPyForum] Community forum for discussions, FAQ and announcements, https://tenpy.johannes-hauschild.de
[git] "git version control system", https://git-scm.com A software which we use to keep track of changes in the source code.
[conda] "conda package manger", https://docs.conda.io/en/latest/ A package and environment management system that allows to easily install (multiple version of) various software, and in particular python packages like TeNPy.
[pip] "pip - the Python Package installer", https://pip.pypa.io/en/stable/ Traditional way to handle installed python packages with pip install ... and pip uninstall ... on the command line.
[matplotlib] "Matplotlib", https://matplotlib.org/ A Python 2D plotting library. Some TeNPy functions expect matplotlib.axes.Axes as arguments to plot into.
[HDF5] "Hierarchical Data Format 5 (R)", https://portal.hdfgroup.org/display/HDF5/HDF5 A file format and library for saving data (including metadata). We use it through the python interface of the h5py library, see Saving to disk: input/output.
[yaml] "YAML Ain’t Markup Language", https://yaml.org A human-readable file format for configuration files. TeNpy (optionally) uses it through pyyaml for reading in simulation parameters, and in some places in the documentation to keep things more readable.
[jupyter] Jupyter notebooks, https://jupyter.org/ An amazing interface for (python) notebooks which can contain both source code, text and outputs in a single file. They provide a good way to get started with python, we use them for examples.
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