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# **pysisyphus Documentation**

*Release 0.7.6.post2*

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*pysisyphus* is a software-suite for the exploration of potential energy surfaces in ground- and **excited states**. It implements several methods to search for stationary points (minima and first order saddle points) and the calculation of minimum energy paths by means of IRC and Chain of States methods like Nudged Elastic Band and Growing String. Furthermore it provides tools to easily analyze & modify geometries (aligning, translating, **interpolating**, ...) and to visualize the calculation results/progress.

The required energies, gradients and hessians are calculated by calling external quantum chemistry codes. *pysisyphus* can also be used as a library to implement custom quantum chemistry workflows.

If any issues arise please open an [issue](#) and I'll see if it can be fixed and my time permits it. Contributions are welcome, so feel free to submit a PR.

**This software is still work in progress. Use at your own risk. Also take a look at the [license](#)**



## OVERVIEW OF PYSISYPHUS

*pysisyphus* is a software-suite for the exploration of potential energy surfaces in ground- and **excited states**. User input is read from YAML files, but it can also be used as a python library to set up custom workflows.

Below you can find a screencast of the transition state (TS) search for the famous alanine dipeptide isomerization at the *xtb* level of theory. It starts with pre-optimizations of the initial and final geometry, a subsequent growing string calculation in delocalized internal coordinates to generate a guess for the final TS optimization and with which it concludes.

```
14 0.000398 0.000109 0.008266 0.002214 0.1
Converged!

Final summary:
  max(forces,internal): 0.000398 hartree/(bohr,rad)
  rms(forces,internal): 0.000109 hartree/(bohr,rad)
  max(forces,cartesian): 0.000590 hartree/bohr
  rms(forces,cartesian): 0.000190 hartree/bohr
  energy: -32.97379138 hartree
Wrote final, hopefully optimized, geometry to 'last_prefinal_geometry.xyz'
Preoptimization of last geometry converged!
Saved final preoptimized structure to 'last_preopt.xyz'.

Read 2 geometries.
#####
# RUNNING GROWINGSTRING #
#####

Path with 2 moving images.
Spent 0.0 s preparing the first cycle.
cycle  max(force)  rms(force)  max(step)  rms(step)  s/cycle
0      0.020646    0.004532    0.016517   0.003626   0.3
      String=2+2 HEI=03/04 (E_max-E_0)=15.3 kJ/mol
1      0.054177    0.008406    0.043341   0.006724   0.5
      String=3+3 HEI=04/06 (E_max-E_0)=76.1 kJ/mol
2      0.091448    0.011890    0.073159   0.009512   0.6
      String=4+4 HEI=04/08 (E_max-E_0)=161.8 kJ/mol
```

Besides being able to track excited states over the course of an optimization the idea of *pysisyphus* is to provide tools for handling the whole process of calculating reaction paths, starting from pre-optimizing the given input geometries for chain of states methods and ending with the hessian calculation on the optimized IRC endpoints.

## 1.1 Entry points

pysisyphus provides several entry points that can be called from the shell (command line). The available commands of the entry points can be queried with the `-h` or `--help` arguments:

```
# Run calculations (Minima optimization, TS search, IRC, NEB, GS, ...)
pysis
# Plotting of path-energies, optimization progress, IRC progress, etc ...
pysisplot
# Manipulate a .trj file or multiple .xyz files
pysistrj
```

### 1.1.1 pysis

Called with a YAML-input file. Simple and more complex examples can be found in the *examples* directory of this repository. Some common arguments are given below:

```
usage: pysis [-h] [--clean] --cp CP
           [yaml]

positional arguments:
  yaml                Start pysisyphus with input from a YAML file.

optional arguments:
  --clean             Ask for confirmation before cleaning.
  --cp CP, --copy CP Copy .yaml file and corresponding geometries to a new
                    directory. Similar to TURBOMOLEs cpc command.
```

### 1.1.2 pysisplot

Visualization/plotting of the pysisyphus calculations. Some common arguments are given below:

```
usage: pysisplot (--cosforces | --cosens | --all_energies | --afir | --opt | --irc | --
↳overlaps)

optional arguments:
  --cosforces, --cf  Plot image forces along a COS.
  --cosens, --ce     Plot COS energies.
  --all_energies, -a Plot ground and excited state energies from 'overlap_data.h5'.
  --afir             Plot AFIR and true -energies and -forces from an AFIR calculation.
  --opt              Plot optimization progress.
  --irc              Plot IRC progress.
  --overlaps, -o
```

### 1.1.3 pysisstrj

Please see `pysisstrj --help` for a list of available arguments.

## 1.2 Available calculators

### 1.2.1 Excited state capabilities

Program	Gradient	Hessian	Exc. states	Version
ORCA	y	y	TD-DFT, TDA	4.2.x
Turbomole	y	y	TD-DFT, TDA, ricc2	7.2-7.4
Gaussian16	y	y	tested TD-DFT, TDA	
PySCF	y	y	tested TD-DFT	1.7.5
OpenMOLCAS	y	n	&rasscf	Not yet derived from OverlapCalculator

### 1.2.2 Ground states capabilities

Program	Gradient	Hessian	Version
MOPAC2016	y	y	
XTB	y	y	6.3.2
QCEngine	y	n	>= 0.16.0
Psi4	y	y	

### 1.2.3 Pure python calculators & Wrappers

Program	Gradient	Hessian	Comment
Sympy 2D	y	y	Many analytical potentials (LEPS, Rosenbrock, Cerjan-Miller, Muller-Brown, ...)
Lennard-Jones	y	n	<b>No</b> periodic boundary conditions
AFIR	y	n	
ONIOM	y	n	Arbitrary number of layers with multicenter-support in the highest layer.
FakeASE	y	n	Wraps <i>pysisyphus</i> calculators so they can be used with <i>ase</i> .

## 1.3 Available algorithms

### 1.3.1 Chain Of States Methods

Algorithm	Coordinates	Comment
Nudged Elastic Band (NEB)	Cartesian, DLC planned	Climbing Image variants, Doubly nudged variant
Adaptive NEB	Cartesian	Not well tested
Free-End NEB	Cartesian	Not well tested
Simple Zero-Temperature-String	Cartesian	Equal spacing, energy-dependent spacing
<b>Growing String Method</b>	Cartesian, <b>DLC</b>	

### 1.3.2 Chain Of States Optimizer

Algorithm	Comment	Links
Steepest Descent	Backtracking variant	<a href="#">NEB-Optimizers</a>
Conjugate Gradient	Backtracking variant	<a href="#">NEB-Optimizers</a>
QuickMin		<a href="#">NEB-Optimizers</a>
FIRE		<a href="#">NEB-Optimizers</a>
BFGS		<a href="#">NEB-Optimizers</a>

### 1.3.3 Transition state optimization

Algorithm	Comment	Links
RS-P-RFO	default	<a href="#">RFO-Paper</a> , <a href="#">RS-Paper</a>
RS-I-RFO		<a href="#">RFO-Paper</a> , <a href="#">RS-Paper</a>
TRIM		<a href="#">TRIM-Paper</a>
Dimer method		

### 1.3.4 Intrinsic Reaction Coordinate integrators

Algorithm	Comment	Links
Damped-Velocity-Verlet		<a href="#">DVV-Paper</a>
Euler	Not recommended	
EulerPC	default	<a href="#">Kaestner-PC</a> , <a href="#">Euler-PC</a>
Gonzales-Schlegel 2		<a href="#">GS2-Paper</a>
Local Quadratic Approximation		<a href="#">LQA-Paper</a>
Modified IMK		<a href="#">IMK-Paper</a>
Runge-Kutta-4	Not recommended	

## 1.4 Additional remarks

*pysisyphus* uses the *tempfile* module from the python stdlib. The location of the temporary directories can be controlled by setting the **\$TMPDIR** environment variable before executing *pysis*.

```
export TMPDIR=[tmpdir]
```



## INSTALLATION

### 2.1 Preparing an environment

It is good idea to install *pysisyphus* into a separate python environment, whether it is an [Anaconda](#) environment or a [virtualenv](#) environment, derived from the system python installation or a [pyenv](#) installation (preferred).

```
# Optional: Create separate Anaconda environment
conda create -n pysis-env python=3.9
# Activate conda environment
conda activate pysis-env

# Optional: Create separate virtual environment
python -m venv pysis-env
# On some linux distributions 'python3' is used instead of 'python', e.g. Ubuntu
# python3 -m venv pysis-env
# Activate virtual environment. Preferably, an absolute path pointing to the
# 'activate' script should be used.
source pysis-env/bin/activate
```

### 2.2 Installation from PyPI with pip

This installs the latest stable release as published on PyPI. If you don't want to do any development work this is the preferred way of installing *pysisyphus*.

```
python -m pip install pysisyphus
# Installation of additional optional dependencies is also possible
# python -m pip install pysisyphus[test]
```

If you run into any problems please make sure that your pip version is recent enough. A pip upgrade is achieved via:

```
python -m pip install --upgrade pip
```

## 2.3 Installation from source

Decide on an `$install_dir` and clone the repository from github. If you want to change the code after installation (develop) do an editable (-e) installation with pip.

```
git clone https://github.com/eljost/pysisyphus.git $install_dir
cd $install_dir
# Install with -e if you want an editable installation
python -m pip install [-e] .
# Installation of extras is also possible. 'sphinx' is only needed if you
# want to build the documentation.
# python -m pip install [-e] .[test]
```

With an editable installation it is also easy to use other branches, besides the *master* branch, e.g., the *dev* branch to test out a new feature.

```
git switch dev
```

## 2.4 Setting up .pysisyphusrc

*pysisyphus* interfaces several quantum chemistry codes and related software (Multiwfn, Jmol). Software available to *pysisyphus* is registered in `$HOME/.pysisyphusrc`, so *pysisyphus* knows which command to execute an/or where to find the binaries.

Depending on the software different choices were made how it is registered. An example `.pysisyphusrc` is given below, with a short comment for each software.

```
# Excited state calculators

[gaussian09]
# Cmd to execute. Please ensure that g09 is on your $PATH.
cmd=g09
formchk_cmd=formchk
unfchk_cmd=unfchk

[gaussian16]
# Cmds to execute. Please ensure that the binaries are found in your $PATH.
cmd=g16
formchk=formchk
unfchk=unfchk
rwfdump=rwfdump

[openmolcas]
# Cmd to execute. Please ensure that pymolcas is on your $PATH.
cmd=pymolcas

[orca]
# ORCA needs the full path to its binary, so please provide the full path.
cmd=/scratch/programme/orca_4_2_0_linux_x86-64_openmpi314/orca

#[pyscf]
```

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```
# pyscf must not have an explicit entry in the .pysisyphusrc. pysisyphus uses
# the python API of pyscf, so it is mandatory that is installed in the same environment
# as pysisyphus.

#[turbomole]
# Turbomole must not have an explicit entry in the .pysisyphusrc. The user has to take
# care that everything is set up correctly, e.g. TM-binaries are on the PATH etc...
# The respective commands are hardcoded into pysisyphus (dscf, ridft, ricc2, ...)

# Ground state calculators

[mopac]
# Similar to Psi4. An example is given below.
cmd=/user/johannes/bin/runmopac.sh

[psi4]
# As the Psi4 installation without conda is, to put it slightly, tricky it was
# decided to allow the installation of Psi4 into a separate conda environment.
# pysisyphus then creates a Psi4 input and sends it to the (bash)-script given below
# that accepts/expects one argument. It is the responsibility of the scrip to activate
# the appropriate conda environment and submit the Psi4 input. An example runpsi4.sh
# script is given below.
cmd=/user/johannes/bin/runpsi4.sh

#[qcengine]
# QCEngine must not have an entry explicit entry in the .pysisyphusrc. It is used
# via its python interface and can be installed as an extra with pip (see above).
# The user is referenced to the QCEngine-documentation for any further questions.

[xtb]
# Cmd to execute. Please ensure that xtb is on your $PATH.
cmd=xtb

[openmolcas]
# Make sure that the MOLCAS variable is set.
cmd=pymolcas

# Utilities

[wfoverlap]
# Cmd to execute. Please ensure that wfoverlap is on your $PATH. The binary/source
# can be obtained from https://github.com/sharc-md/sharc/tree/master/bin
cmd=/scratch/wfoverlap_1.0/bin/wfoverlap.x

[mwfn]
# Cmd to execute. Please ensure that Multiwfn is on your $PATH. Otherwise put an
# absolute path here. By default pysisyphus looks up "Multiwfn", so if you would
# put a relative path here you don't have to, as this is already covered by the
# defaults.
cmd=Multiwfn

[jmol]
```

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```
# Cmd to execute. The same arguments apply for jmol as for Multiwfn. "jmol" is
# already the default.
cmd=jmol
```

When the specified path/cmd is not absolute, but relative (e.g. for xtb, g16, ...) the corresponding binaries have to be available on the `$PATH` and all other environment variables have to be set up correctly by the user.

## 2.5 Example runpsi4.sh

```
#!/bin/bash

# Afaik this doesn't work in non-interactive shells ...
# See https://github.com/conda/conda/issues/8072
# conda activate psi4
source /scratch/programme/anaconda3/bin/activate psi4
#conda activate psi4
psi4 -o stdout $1
```

## 2.6 Example runmopac.sh

```
#!/bin/bash

module purge
module load mopac

MOPAC2016.exe $1
```

## 2.7 Verifying Installation

By executing `pytest -v --pyargs pysisyphus.tests` a series of quick tests can be executed, verifying successful calculator setup. Running these tests requires `pyscf` and `pytest` to be present (*pip install pyscf pytest*).

## INSTALLATION VIA NIX

An easy and reliable way to get a complete pysisyphus installation, including additional quantum chemistry (QC) codes, is possible through the [Nix package manager](#). Required Nix files for the QC codes and their dependencies (linear algebra & MPI libraries etc.) are provided by the [NixOS-QChem](#) overlay.

Nix installations are fully reproducible and extremely simple to accomplish for any potential pysisyphus user, as most of the necessary configuration is already described in the respective Nix files.

The [Nix Pills](#) series provides an informative introduction to the general concepts behind Nix.

Recent stable versions of pysisyphus are also directly included in [NixOS-QChem](#). If you do not require a development version of pysisyphus, using the [NixOS-QChem](#) overlay is the preferred option. Otherwise, continue with the instructions below.

### 3.1 Prerequisites

Please follow the installation instructions in the [nix manual](#) to set up a working Nix installation on your system. You may want to enable the `flakes` and `nix-command` features for Nix.

### 3.2 Caching

The build time can be reduced drastically, if the quantum chemistry codes are fetched from a binary cache.

Pysisyphus provides a binary cache on Cachix.

Nix  $\geq 2.6.0$  with flakes directly supports the binary cache without further action required.

Alternatively, if you are allowed to add binary caches in nix, you may simply execute:

```
nix run nixpkgs#cachix use pysisyphus
```

## 3.3 Usage

You may use pysisyphus directly via Flakes without cloning any repository or local installation. Some examples how to use pysisyphus with nix:

Inspect the structure of pysisyphus' flake:

```
nix flake show github:eljost/pysisyphus
```

Directly run the pysis command on an input :

```
nix run github:eljost/pysisyphus input.yml
```

Start an interactive shell with a pysisyphus installation

```
nix shell github:eljost/pysisyphus
```

For developing and hacking pysisyphus and for legacy Nix commands, first clone the pysisyphus repository

```
git clone https://github.com/eljost/pysisyphus.git && cd pysisyphus
```

You may obtain a development shell for pysisyphus via

```
nix develop
```

build pysisyphus in the legacy nix style

```
nix-build ./nix/default.nix
```

or obtain a development legacy-style dev-shell

```
nix-shell ./nix/shell.nix
```

The Flake also offers options to build pysisyphus with proprietary components such as ORCA or Turbomole, e.g.

```
nix run .#pysisyphusOrca
```

or build singularity or docker containers:

```
nix build .#pysisyphusSingularity
```

## WORKED EXAMPLE

This page provides a general overview of pysisyphus's capabilities for optimizing ground state reaction paths for the example of a tris-pericyclic reaction between a tropone derivative and dimethylfulvene.

You can find a short discussion of the reaction on Steven M. Bacharach's [blog](#). Here we focus on how to obtain TS1. Instead of using DFT as in the original [publication](#) we'll employ the semi-empirical tight binding code `xtb`. It allows us to run all necessary calculations on common desktop hardware in little time (2 min, using 6 physical CPU cores).

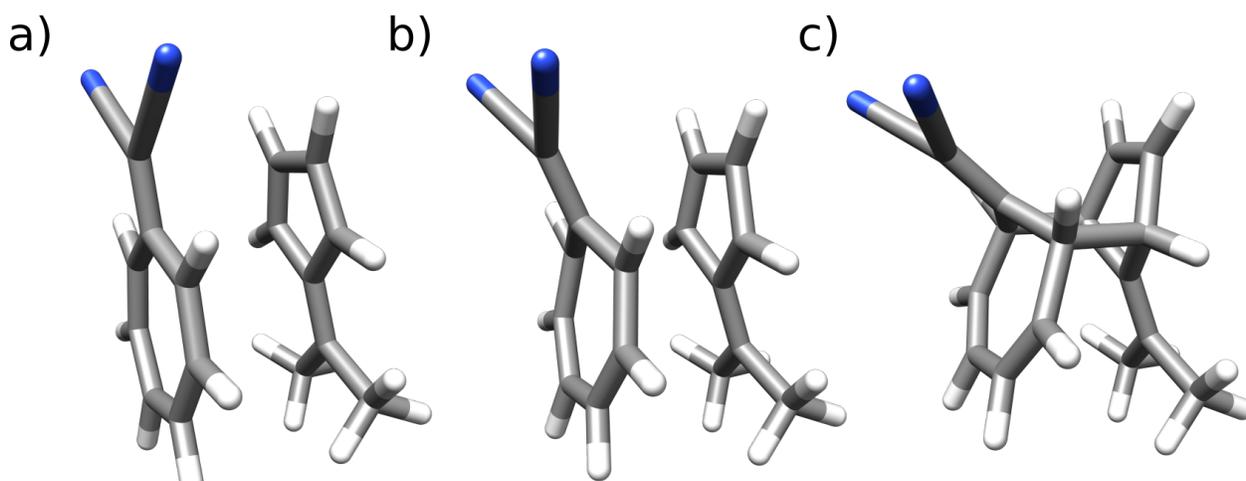


Fig. 4.1: Educts, TS and products of the tris-pericyclic reaction between a tropone derivative and dimethylfulvene.

At first, we have to create appropriate educt and product geometries using the molecular editor of our choice, e.g., [avogadro](#) or [TmoleX](#). Given a reasonable educt geometry a suitable product geometry can be constructed by decreasing the distance between the two ring systems and optimizing the result, e.g., by `xtb`. Sometimes it may be easier to start with the product geometry, and obtain the educt from it. **Consistent atom ordering in both geometries is mandatory!** Usually it's a bad idea to construct both geometries independently, as it is easy to mess up the atom ordering.

Given educts (`min_xtbopt.xyz`) and product (`prod_xtbopt.xyz`) we can create the YAML input. Our goal is to obtain the barrier heights for the reaction shown in [Fig. 4.1](#) by means of growing string (GS) calculation, transition state (TS) optimization, intrinsic reaction coordinate (IRC) calculation and subsequent optimization of the IRC endpoints. The full input is shown below. We'll go over it step by step.

```
geom:  
  type: dlc  
  fn: [min_xtbopt.xyz, prod_xtbopt.xyz]  
preopt:  
cos:  
  type: gs
```

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```

max_nodes: 18
climb: True
opt:
  type: string
  align: False
  max_cycles: 20
tsopt:
  type: rsirfo
  do_hess: True
  thresh: gau
  hessian_recalc: 3
irc:
  type: eulerpc
  rms_grad_thresh: 0.0005
endopt:
calc:
  type: xtb
  pal: 6

```

The desired coordinate system and the file names of the input geometries are given in the geom block.

```

geom:
  type: dlc
  fn: [min_xtbopt.xyz, prod_xtbopt.xyz]

```

Here we chose delocalized internal coordinates (DLC) for our GS, which is the preferred way. Alternatively Cartesian coordinates could be used by `type: cart`. If DLCs fail, Cartesian coordinates should be tried.

```
preopt:
```

The `preopt` block is given without any additional keywords, so sane defaults will be used for preoptimizing educt and product geometries (Rational function optimization (RFO) in redundant internal coordinates). Strictly, in our case pre-optimization is not necessary, as we already preoptimized the geometries using `xtb`. But in general it is advised to span chain of states (COS) like GS or nudged elastic band (NEB) between stationary points on the potential energy surface. If the educts and products are NOT covalently bound it may be a good idea to restrict the number of preoptimization cycles to a small number (`max_cycles: 10`), as these optimizations are sometimes hard to converge. Please see [Optimization of Minima](#) for a list of possible keywords in the `preopt` block.

```

cos:
  type: gs
  max_nodes: 18
  climb: True

```

The `cos` block configures COS calculations. Here we request a GS (`gs`) with 18 nodes (images) between educt and product, resulting in a total string length of 20 nodes. By default first and last node (educt and product) are kept fixed throughout the optimization. We enable a climbing image (CI) to obtain a better TS guess. Please see [Chain Of States Methods](#) for further information on COS methods.

```

opt:
  type: string
  align: False
  max_cycles: 20

```

COS/GS optimization is controlled via the `opt` block. For GS optimization one should always use `type: string`.

In internal coordinates we disable automated geometry alignment, as it is not needed. We also restrict the number of optimization cycles to 20 (default 50). The chosen optimizer will do steepest descent (SD) steps when the string grew in the previous cycle, otherwise conjugate gradient (CG) steps are used. When the GS is fully grown/connected the optimizer will use limited-memory Broyden-FletcherGoldfarb-Shanno (L-BFGS) to determine more sophisticated steps.

```
tsopt:
  type: rsirfo
  do_hess: True
  thresh: gau
  hessian_recalc: 3
```

After GS convergence the highest energy image (HEI) is determined by cubic splining and used as guess for a classical TS optimization using restricted step image RFO (RSIRFO). `do_hess: True` requests a frequency calculation after the TS optimization. The Hessian is recalculated every 3th step. When the Hessian for the chosen computational method is reasonably cheap it is a good idea to recalculate it periodically. Between recalculations it's updated using the Bofill-update. Convergence criteria are tightened from the default `thresh: gau_loose` to `thresh: gau`.

```
irc:
  type: eulerpc
  rms_grad_thresh: 0.0005
```

IRC integration is controlled in the `irc` block. By default the Euler-predictor-corrector (EulerPC) integrator is used. Integration is terminated when the root-mean-square (RMS) of the gradient is equal to or less than 0.0005 au. Possible inputs are given in *Intrinsic Reaction Coordinate (IRC)*.

```
endopt:
```

Similar to `preopt` the `endopt` will be executed with default arguments. It is used to optimize the IRC endpoints to stationary points and enables printing of additional information like RMS deviation of atomic positions (RMSD) between optimized endpoints and initial geometries. The RMSD values help in deciding if the obtained TS actually connects presumed educts and products.

```
calc:
  type: xtb
  pal: 6
```

The `calc` block configures the level of theory used in energy/gradient/Hessian calculations. Here we chose `xtb` and requested 6 CPU cores. Additional inputs for `xtb` can be found in the *xtb module documentation*

With everything set up we are ready to actually execute pysisyphus! Assuming the above YAML is saved to `01_pericyclic.yaml` just run

```
pysis 01_pericyclic.yaml | tee pysis.log
```

By default pysisyphus prints to STDOUT so you have to capture STDOUT explicitly. We use `tee` so everything is logged to a file and printed simulatenously. A lot of files and output will be produced so we will go over everything slowly.

```

                                d8b                                888
                                Y8P                                888
                                888                                888
888888b. 888 888 .d8888b 888 .d8888b 888 888 888888b. 888888b. 888 888 .d8888b
888 "88b 888 888 88K    888 88K    888 888 888 "88b 888 "88b 888 888 88K
888 888 888 888 "Y8888b. 888 "Y8888b. 888 888 888 888 888 888 888 "Y8888b.
```

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```

888 d88P Y88b 888      X88 888      X88 Y88b 888 888 d88P 888 888 Y88b 888      X88
888888P"  "Y88888 888888P' 888 888888P' "Y88888 888888P" 888 888 "Y88888 888888P'
888      888      888 888
888      Y8b d88P      Y8b d88P 888
888      "Y88P"      "Y88P" 888

Version 0.5.0.post1+450.g2c1654d3 (Python 3.8.5, NumPy 1.19.2, SciPy 1.5.2)
Git commit 2c1654d35d69e7b48ac4e9b00d38afd58a8bedd4
Executed at Tue Oct 6 10:08:30 2020 on 'your fancy hostname'

If pysisyphus benefitted your research please cite:

    https://doi.org/10.1002/qua.26390

Good luck!

```

You will be greeted by a banner and some information about your current installation, which hopefully aids in reproducing your results later on, if needed. Then your input is repeated, including default values that you did not explicitly set. There you can also see the default values chosen for `preopt` and `endopt`.

```

{'calc': {'pal': 6, 'type': 'xtb'},
'cos': {'climb': True, 'fix_ends': True, 'max_nodes': 18, 'type': 'gs'},
'endopt': {'dump': True,
           'fragments': False,
           'max_cycles': 100,
           'overachieve_factor': 3,
           'thresh': 'gau',
           'type': 'rfo'},
'geom': {'fn': ['min_xtbopt.xyz', 'prod_xtbopt.xyz'], 'type': 'dlc'},
'interpol': {'between': 0, 'type': None},
'irc': {'rms_grad_thresh': 0.0005, 'type': 'eulerpc'},
'opt': {'align': False, 'dump': True, 'max_cycles': 30, 'type': 'string'},
'preopt': {'dump': True,
           'max_cycles': 100,
           'overachieve_factor': 3,
           'preopt': 'both',
           'strict': False,
           'thresh': 'gau_loose',
           'trust_max': 0.3,
           'type': 'rfo'},
'tsopt': {'do_hess': True,
          'dump': True,
          'h5_group_name': 'tsopt',
          'hessian_recalc': 3,
          'overachieve_factor': 3,
          'thresh': 'gau',
          'type': 'rsirfo'}}

```

The whole run starts with preoptimizations of educt and product. Both optimizations converge quickly, as the geometries are already preoptimized.

```

#####

```

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```

# RUNNING FIRST PREOPTIMIZATION #
#####
Spent 0.0 s preparing the first cycle.
cycle  max(force)  rms(force)  max(step)  rms(step)  s/cycle
   0    0.000080   0.000010   0.027005   0.003934   0.1
Converged!

Final summary:
    max(forces, internal): 0.000080 hartree/(bohr,rad)
    rms(forces, internal): 0.000010 hartree/(bohr,rad)
    max(forces, cartesian): 0.000063 hartree/bohr
    rms(forces, cartesian): 0.000015 hartree/bohr
    energy: -52.35197394 hartree
Wrote final, hopefully optimized, geometry to 'first_pre_final_geometry.xyz'

Preoptimization of first geometry converged!
Saved final preoptimized structure to 'first_preopt.xyz'.
RMSD with initial geometry: 0.000000 au

#####
# RUNNING LAST PREOPTIMIZATION #
#####
Spent 0.0 s preparing the first cycle.
cycle  max(force)  rms(force)  max(step)  rms(step)  s/cycle
   0    0.000188   0.000025   0.002310   0.000592   0.1
Converged!

Final summary:
    max(forces, internal): 0.000188 hartree/(bohr,rad)
    rms(forces, internal): 0.000025 hartree/(bohr,rad)
    max(forces, cartesian): 0.000182 hartree/bohr
    rms(forces, cartesian): 0.000032 hartree/bohr
    energy: -52.39960456 hartree
Wrote final, hopefully optimized, geometry to 'last_pre_final_geometry.xyz'

Preoptimization of last geometry converged!
Saved final preoptimized structure to 'last_preopt.xyz'.
RMSD with initial geometry: 0.000000 au

```

After an optimization remaining RMS and max of the forces are reported for internal and Cartesian coordinates. If the internal force is zero, but a substantial Cartesian force remains something went wrong, e.g., the generated coordinate system is lacking important coordinates. In such cases the generated coordinates can be examined manually `pysistrj [geom file] --internals` to determine important missing coordinates.

Preoptimizations are followed by the GS optimization.

```

#####
# RUNNING GROWINGSTRING #
#####
Spent 0.0 s preparing the first cycle.
cycle  max(force)  rms(force)  max(step)  rms(step)  s/cycle
   0    0.006403   0.001545   0.006403   0.001545   0.2
String=2+2 HEI=02/04 (E_max-E_0)=1.4 kJ/mol

```

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```

1      0.009367      0.001734      0.009367      0.001734      0.3
String=3+3 HEI=03/06 (E_max-E_0)=4.1 kJ/mol
2      0.010394      0.001807      0.010394      0.001807      0.6
String=4+4 HEI=04/08 (E_max-E_0)=8.0 kJ/mol
3      0.011152      0.001790      0.011152      0.001790      1.1
String=5+5 HEI=05/10 (E_max-E_0)=12.5 kJ/mol
4      0.010687      0.001722      0.010687      0.001722      1.0
String=6+6 HEI=06/12 (E_max-E_0)=17.6 kJ/mol
5      0.008751      0.001620      0.008751      0.001620      1.1
String=7+7 HEI=07/14 (E_max-E_0)=22.8 kJ/mol
6      0.008007      0.001517      0.008007      0.001517      1.3
String=8+8 HEI=08/16 (E_max-E_0)=27.6 kJ/mol
7      0.006895      0.001423      0.006895      0.001423      1.4
String=9+9 HEI=09/18 (E_max-E_0)=31.3 kJ/mol
Starting to climb in next iteration.
8      0.006253      0.001313      0.006253      0.001313      1.6
String=Full HEI=10/20 (E_max-E_0)=32.8 kJ/mol
9      0.006134      0.001169      0.059438      0.011487      1.7
String=Full HEI=10/20 (E_max-E_0)=32.0 kJ/mol
10     0.009555      0.001504      0.022529      0.003827      1.6
String=Full HEI=10/20 (E_max-E_0)=31.6 kJ/mol
11     0.009792      0.001583      0.100000      0.011614      1.7
String=Full HEI=10/20 (E_max-E_0)=31.6 kJ/mol
12     0.008636      0.001534      0.100000      0.011518      1.7
String=Full HEI=10/20 (E_max-E_0)=30.7 kJ/mol
13     0.008708      0.001448      0.100000      0.011463      1.6
String=Full HEI=10/20 (E_max-E_0)=30.0 kJ/mol
14     0.008217      0.001363      0.100000      0.005660      1.7
String=Full HEI=10/20 (E_max-E_0)=29.4 kJ/mol
15     0.007743      0.001335      0.100000      0.005297      1.7
String=Full HEI=10/20 (E_max-E_0)=29.1 kJ/mol
16     0.007243      0.001307      0.100000      0.005143      1.6
String=Full HEI=10/20 (E_max-E_0)=28.8 kJ/mol
17     0.006897      0.001285      0.100000      0.004938      1.6
String=Full HEI=10/20 (E_max-E_0)=28.6 kJ/mol
18     0.006772      0.001277      0.100000      0.004992      1.6
String=Full HEI=10/20 (E_max-E_0)=28.5 kJ/mol
19     0.006714      0.001273      0.100000      0.005006      1.6
String=Full HEI=10/20 (E_max-E_0)=28.5 kJ/mol
Found sign 'converged'. Ending run.
Operator indicated convergence!
Wrote final, hopefully optimized, geometry to 'final_geometries.trj'
Splined HEI is at 8.07/19.00, between image 8 and 9 (0-based indexing).
Wrote splined HEI to 'splined_hei.xyz'

```

The string grows quickly and is fully grown in cycle 8. String size and barrier height between the first and HEI are reported in every cycle. From cycle 8 on, a CI is employed. The final HEI index is printed at the end. As we interpolate the HEI, the index may be a fractional number. The COS optimization is followed by a TS optimization.

```

#####
# RUNNING TS-OPTIMIZATION FROM COS #
#####

```

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```

Creating mixed HEI tangent, using tangents at images (8, 9).
Overlap of splined HEI tangent with these tangents:
  08: 0.982763
  09: 0.149474
Index of splined highest energy image (HEI) is 8.07.

Wrote animated HEI tangent to cart_hei_tangent.trj

Splined HEI (TS guess)
[xyz file printed here; removed for clarity]

Splined Cartesian HEI tangent
[xyz file printed here; removed for clarity]

Wrote splined HEI coordinates to 'splined_hei.xyz'
Calculating Hessian at splined TS guess.
Negative eigenvalues at splined HEI:
[-0.005069 -0.000317]
Overlaps between HEI tangent and imaginary modes:
  00: 0.856746
  01: 0.009826
Imaginary mode 0 has highest overlap with splined HEI tangent.
Spent 0.3 s preparing the first cycle.

```

cycle	max(force)	rms(force)	max(step)	rms(step)	s/cycle
0	0.009004	0.001343	0.056873	0.017930	0.2
1	0.000754	0.000227	0.158373	0.035866	0.2
2	0.001142	0.000156	0.316150	0.051562	3.6
3	0.000483	0.000082	0.339089	0.052646	0.2
4	0.000370	0.000067	0.321142	0.049527	0.2
5	0.000362	0.000050	0.243173	0.040395	3.6
6	0.000152	0.000023	0.174987	0.032910	0.2
7	0.000503	0.000063	0.092082	0.019454	0.2
8	0.000199	0.000031	0.121285	0.024502	3.5
9	0.000136	0.000019	0.061690	0.012096	0.2

```

Converged!

Final summary:
    max(forces, internal): 0.000136 hartree/(bohr,rad)
    rms(forces, internal): 0.000019 hartree/(bohr,rad)
    max(forces, cartesian): 0.000164 hartree/bohr
    rms(forces, cartesian): 0.000043 hartree/bohr
    energy: -52.34823514 hartree
Wrote final, hopefully optimized, geometry to 'ts_final_geometry.xyz'
Optimized TS coords:
[xyz file printed here; removed for clarity]

Wrote TS geometry to 'ts_opt.xyz'

-----
| HESSIAN AT FINAL GEOMETRY |
-----

```

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```

... mass-weighting cartesian hessian
... doing Eckart-projection

First 10 eigenvalues [-2.4303e-03 -3.9622e-17 -8.1858e-18 -7.8671e-19  2.1865e-18  5.
↪1589e-18
  2.6247e-17  1.3733e-05  7.3736e-05  1.2735e-04]
Imaginary frequencies: [-253.24] cm1

Wrote final (not mass-weighted) hessian to 'calculated_final_cart_hessian'.
Wrote HD5 Hessian to 'final_hessian.h5'.
Wrote imaginary mode with =-253.24 cm1 to 'imaginary_mode_000.trj'

Barrier between TS and first COS image: 9.8 kJ mol1
Barrier between TS and last COS image: 134.9 kJ mol1

```

The initial HEI TS guess features only two sizable imaginary frequencies, confirming that it is a suitable TS guess. Root 0 has the highest overlap (85%) with the HEI tangent and is chosen for maximization in the TS optimization, whereas the energy will be minimized along the remaining modes. The optimization converged quickly in 10 cycles. A final Hessian is computed at the optimized TS as we used `do_hess: True`. Only one imaginary frequency remains, which is the desired result for a first-order saddle point. All sizable imaginary modes are written to `.trj` files and can be viewed by tools like `jmol`.

Barrier heights between the TS and first and last COS image (no thermochemistry is included!) are reported. In this case the energy difference between the first COS image and the TS is very small, indicating an early TS, similar to the educts. This is also be confirmed by examining Fig. 4.1.

```

#####
# RUNNING IRC #
#####

Calculating energy and gradient at the TS
IRC length in mw. coords, max(|grad|) and rms(grad) in unweighted coordinates.
Norm of initial displacement step: 0.1974

```

```

#####
# IRC - FORWARD #
#####

Step  IRC length      dE / au  max(|grad|)  rms(grad)
-----
  0    0.298147      -0.000333  0.001413    0.000394
Integrator indicated convergence!

```

```

#####
# IRC - BACKWARD #
#####

Step  IRC length      dE / au  max(|grad|)  rms(grad)
-----
  0    0.332823      -0.000938  0.004607    0.001154
  1    0.662398      -0.001367  0.006116    0.001608
  2    0.991868      -0.001878  0.008670    0.002208
  3    1.320005      -0.002558  0.012581    0.002968
  4    1.644982      -0.003249  0.015719    0.003595

```

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5	1.966058	-0.003700	0.017023	0.003976
6	2.285006	-0.003972	0.016598	0.004230
7	2.596325	-0.004126	0.019112	0.004415
8	2.898296	-0.004206	0.021118	0.004439
9	3.186446	-0.004052	0.021182	0.004240
10	3.462246	-0.003697	0.019162	0.003793
11	3.723777	-0.003087	0.015042	0.003078
12	3.966373	-0.002263	0.009324	0.002161
13	4.173308	-0.001387	0.003822	0.001310
14	4.321559	-0.000781	0.002226	0.000838
15	4.420391	-0.000523	0.001616	0.000642
16	4.498118	-0.000418	0.001299	0.000548
17	4.571712	-0.000353	0.001967	0.000569
18	4.647481	-0.000298	0.004516	0.000754
19	4.722445	-0.000308	0.003469	0.000670
20	4.808189	-0.000290	0.003426	0.000682
21	4.881593	-0.000209	0.004155	0.000683

Integrator indicated convergence!

The imaginary mode is used to displace the TS towards educts and product. As the TS is very similar to the educt, forward IRC integration already terminates after one cycle. Maybe, further integration steps could be forced by tightening the threshold in the `irc:` block. Backward integration terminates after 22 cycles. At first, the gradient increases and after the inflection point is passed, falls off again as a stationary point is approached. In the end both IRC endpoints are fully optimized to stationary points.

```
#####
# RUNNING FORWARD_END OPTIMIZATION #
#####
Spent 0.0 s preparing the first cycle.
cycle  max(force)  rms(force)  max(step)  rms(step)  s/cycle
  0    0.000491    0.000126    0.099615    0.030780    0.1
  1    0.001769    0.000333    0.121277    0.033041    0.2
  2    0.002986    0.000525    0.228218    0.054622    0.2
  3    0.002455    0.000475    0.396872    0.096049    0.2
  4    0.003106    0.000502    0.061397    0.012734    0.2
  5    0.000676    0.000187    0.040565    0.007988    0.2
  6    0.000498    0.000144    0.048609    0.012294    0.2
  7    0.000908    0.000142    0.055895    0.016806    0.2
  8    0.000715    0.000102    0.037441    0.011352    0.2
  9    0.000226    0.000047    0.034467    0.006157    0.2
 10    0.000119    0.000032    0.034646    0.006157    0.1
Converged!

Final summary:
  max(forces, internal): 0.000119 hartree/(bohr,rad)
  rms(forces, internal): 0.000032 hartree/(bohr,rad)
  max(forces, cartesian): 0.000241 hartree/bohr
  rms(forces, cartesian): 0.000073 hartree/bohr
  energy: -52.35194671 hartree
Wrote final, hopefully optimized, geometry to 'forward_end_final_geometry.xyz'

Moved 'forward_end_final_geometry.xyz' to 'forward_end_opt.xyz'.
```

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```

#####
# RUNNING BACKWARD_END OPTIMIZATION #
#####
Spent 0.0 s preparing the first cycle.
cycle  max(force)  rms(force)  max(step)  rms(step)  s/cycle
  0     0.004485   0.001075   0.168005   0.031009   0.1
  1     0.003805   0.000731   0.158516   0.028598   0.2
  2     0.003345   0.000610   0.047413   0.008626   0.2
  3     0.002849   0.000531   0.093820   0.017252   0.2
  4     0.001571   0.000332   0.154150   0.028740   0.2
  5     0.000752   0.000148   0.034367   0.007031   0.2
  6     0.000630   0.000107   0.017362   0.004137   0.2
  7     0.000218   0.000047   0.019096   0.003624   0.1
  8     0.000165   0.000041   0.027156   0.004814   0.2
  9     0.000400   0.000065   0.044195   0.007618   0.2
 10     0.000460   0.000076   0.039724   0.006652   0.1
 11     0.000265   0.000059   0.021454   0.003444   0.2
 12     0.000194   0.000034   0.009649   0.001594   0.1
 13     0.000146   0.000025   0.009711   0.001835   0.1
Converged!

Final summary:
    max(forces, internal): 0.000146 hartree/(bohr,rad)
    rms(forces, internal): 0.000025 hartree/(bohr,rad)
    max(forces, cartesian): 0.000183 hartree/bohr
    rms(forces, cartesian): 0.000047 hartree/bohr
    energy: -52.39959356 hartree
Wrote final, hopefully optimized, geometry to 'backward_end_final_geometry.xyz'

Moved 'backward_end_final_geometry.xyz' to 'backward_end_opt.xyz'.

```

Both optimizations converge quickly without any problems. Finally, optimized endpoint geometries are compared to the initial geometries and final barrier heights are reported. Thermochemistry is not (yet) included. Even though both endpoints are reported as dissimilar to the initial geometries they are still very similar, confirming, that the obtained TS indeed connects presumed educts and product.

```

#####
# RMSDS AFTER END OPTIMIZATIONS #
#####
start geom 0 (first_preopt.xyz/min_xtbopt.xyz)
  end geom 0 ( forward_end_opt.xyz): RMSD=0.133727 au
  end geom 1 (backward_end_opt.xyz): RMSD=1.126059 au
  Optimized end geometries are dissimilar to 'first_preopt.xyz/min_xtbopt.xyz'!
start geom 1 (last_preopt.xyz/prod_xtbopt.xyz)
  end geom 0 ( forward_end_opt.xyz): RMSD=1.144102 au
  end geom 1 (backward_end_opt.xyz): RMSD=0.093044 au
  Optimized end geometries are dissimilar to 'last_preopt.xyz/prod_xtbopt.xyz'!

#####
# BARRIER HEIGHTS AFTER END OPTIMIZATIONS #
#####

```

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```
Thermochemical corrections are NOT included!
```

```
Minimum energy of 0.0 kJ mol1 at 'backward_end_opt.xyz'.
```

```
    forward_end_opt.xyz:  125.10 kJ mol1  
                        TS:  134.84 kJ mol1  
    backward_end_opt.xyz:   0.00 kJ mol1
```

```
Wrote optimized end-geometries and TS to 'end_geoms_and_ts.trj'
```

Visualization/plotting of running optimizations and IRC integrations is also possible. Please see *Plotting* for further information.

All inputs can be found in the *examples/complex/08\_trispericyclic* directory on github.



## COORDINATE SYSTEMS

Choice of coordinate system is often crucial, for the successful outcome of geometry optimizations. Pysisyphus supports different coordinate systems. By default, molecular optimizations are carried out in redundant internal coordinates (RIC).

### 5.1 Supported Coordinate Systems

#### 5.1.1 Cartesian Coordinates

- Strong coupling
- Unambiguously defined, if translation and rotation (TR) are removed
- Redundant set, if TR are not removed
- type: `cart`

#### 5.1.2 Redundant Internal Coordinates

- Comprising bond stretches, (linear) bends, (improper) torsions and other primitives
- Less coupling, compared to Cartesians
- Easy estimation of diagonal model Hessians
- **Support constraints**
- Require sophisticated setup algorithm
- Iterative internal-Cartesian backtransformation, which may fail
- Usually highly redundant set
- type: `redund`

### 5.1.3 Delocalized Internal Coordinates (DLC)

- Complicated linear combinations of primitive internals
- No coupling, coordinates are orthogonal to each other, at least at the geometry they were defined at
- Non redundant set
- More efficient compared to RIC for bigger systems (if initial DLC generation is feasible)
- Same comments apply, as for RICs
- type: `dlc`

### 5.1.4 Translation & Rotation Internal Coordinates (TRIC)

- Especially suited to optimize solvated/non-covalently bound systems
- Translation and rotation coordinates are assigned to every fragment
- Avoids error-prone assignment of interfragment coordinates
- See [10.1063/1.4952956](#) for a full discussion
- By default the B-Matrix is recalculated in every step of the internal-Cartesian backtransformation when TRIC is enabled
- type: `tric`

## 5.2 Supported File Formats

All formats can be read, at least to a certain extent. Coordinate files are expected to be in , if not otherwise dictated by the respective format.

Suffix	Write	Comment
<code>.xyz</code>	✓	Plain XYZ, single geometry.
<code>.trj</code>	✓	Plain XYZ, multiple geometries.
<code>.pdb</code>	✓	Protein Data Bank.
<code>.molden</code>		Restricted to [Geometries] block.
<code>.zmat</code>		Z-Matrix, see below for an example.
<code>.cjson</code>		As saved by Avogadro.
<code>.crd</code>		CHARMM card format
<code>.sdf</code>		Structure-data file

### 5.2.1 Z-Matrix example

```
C
N 1 1.35
H 1 1.0 2 105.0
H 2 1.4 1 105.0 3 150.0
H 2 1.4 1 110.0 3 -160.0
```

Indexing in Z-matrices is 1-based, so the first atom has index 1. Variable substitution is not supported in the current Z-matrix parser.

## 5.3 YAML Input

See below for an explanation of possible inputs in the *geom* section. Input related to internal coordinates is given mostly in the *coord\_kwargs* subgroup, whereas atom-related input (*isotops*, *freeze\_atoms*) is given one level above. See the example below:

```
geom:
  type: cart           # Coordinate system (cart/redund/dlc/tric)
  fn: [input]         # File name or inline input
  union: False        # Define same set of primitives at multiple geometries
  isotops: null       # Specify different isotopes
  freeze_atoms: null  # Freeze Cartesians of certain atoms
  coord_kwargs:      # Keywords that are passed to the internal coordinate_
  ↪class
    define_prims: null # Additionally define these primitives
    constrain_prims: null # Primitive internals to be constrained
    freeze_atoms_exclude: False # Whether to set up internal coordinates for frozen atoms
  preopt:
    geom:             # geom block in preopt takes same keywords as above
    ...              # no 'fn' key here!
  endopt:
    geom:             # geom block in endopt takes same keywords as above
    ...              # no 'fn' key here!
```

Employed coordinates and coordinate systems in *preopt* and *endopt* are similarly controlled by a *geom* block. Same keywords are supported, as for the *geom* block, at the top level, except the *fn* key.

### 5.3.1 Inline input

Inline coordinates are supported for XYZ format, and expected in `.`. Take care of proper indentation. The example below would yield RIC for the hydrogen molecule (1 bond).

```
geom:
  type: redund
  fn: |
    2

    H 0.0 0.0 0.0
    H 0.0 0.0 0.7
```

### 5.3.2 Types of Primitive Coordinates

Pysisyphus implements many different (primitive) internal coordinates. Every coordinate is defined by its type and a set of atom indices, e.g., 2 indices for a bond, 3 indices for a bend and 4 indices for a dihedral.

Specification of a type is necessary, as there are many different kinds of bonds, bends and dihedrals/out-of-plane. One can't just assume, that a coordinate comprised of 3 atom indices is always a regular bend, as it may also be a linear bend or a translational coordinate (TRANSLATION\_X, 14), describin the mean Cartesian X coordinate of 3 atoms.

Atom indices start at 0!

```

# Primitive types
BOND = 0
AUX_BOND = 1
HYDROGEN_BOND = 2
INTERFRAG_BOND = 3
AUX_INTERFRAG_BOND = 4
BEND = 5
LINEAR_BEND = 6
LINEAR_BEND_COMPLEMENT = 7
PROPER_DIHEDRAL = 8
IMPROPER_DIHEDRAL = 9
OUT_OF_PLANE = 10
LINEAR_DISPLACEMENT = 11
LINEAR_DISPLACEMENT_COMPLEMENT = 12
# TRANSLATION = 13 # Dummy coordinate
TRANSLATION_X = 14
TRANSLATION_Y = 15
TRANSLATION_Z = 16
# ROTATION = 17 # Dummy coordinate
ROTATION_A = 18
ROTATION_B = 19
ROTATION_C = 20
# CARTESIAN = 21 # Dummy coordinate
CARTESIAN_X = 22
CARTESIAN_Y = 23
CARTESIAN_Z = 24
BONDED_FRAGMENT = 25
DUMMY_TORSION = 26
DISTANCE_FUNCTION = 27
# atan2 based coordinates
BEND2 = 28
TORSION2 = 29

```

As some of these types are quite unwieldy, several shortcuts are supported, that can be used in place of the types above.

```

# Additional shortcuts
# Using Cartesians in the framework of internal coordinates is mainly
# useful if one wants to constrain certain atoms.
"X": [PT.CARTESIAN_X],
"Y": [PT.CARTESIAN_Y],
"Z": [PT.CARTESIAN_Z],
"XY": [PT.CARTESIAN_X, PT.CARTESIAN_Y],
"XZ": [PT.CARTESIAN_X, PT.CARTESIAN_Z],
"YZ": [PT.CARTESIAN_Y, PT.CARTESIAN_Z],
"XYZ": [PT.CARTESIAN_X, PT.CARTESIAN_Y, PT.CARTESIAN_Z],
"ATOM": [PT.CARTESIAN_X, PT.CARTESIAN_Y, PT.CARTESIAN_Z],
# Primitive aliases
"B": [PT.BOND],
"A": [PT.BEND],
"A2": [PT.BEND2],
"D": [PT.PROPER_DIHEDRAL],
"DIHEDRAL": [PT.PROPER_DIHEDRAL],
"TORSION": [PT.PROPER_DIHEDRAL],

```

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```
"D2": [PT.PROPER_DIHEDRAL2],
"DIHEDRAL2": [PT.PROPER_DIHEDRAL2],
"TORSION2": [PT.PROPER_DIHEDRAL2],
# Translation & Rotation coordinates
"TRANSLATION": [PT.TRANSLATION_X, PT.TRANSLATION_Y, PT.TRANSLATION_Z],
"ROTATION": [PT.ROTATION_A, PT.ROTATION_B, PT.ROTATION_C],
# Miscellaneous
"DIST_FUNC": [PT.DISTANCE_FUNCTION]
```

### 5.3.3 Define Additional Primitives

Pysisyphus tries its best, to automatically come up with a reasonable set of internal coordinates, but sometimes the algorithm misses an important one. Especially at transition state guesses, where increased atom distances are common, bonds may be missed.

In such cases, additional coordinates can be requested explicitly. If additional coordinates are requested, **a nested list is expected** `[[coord0], [coord1], ...]`.

```
# General structure (list of coordinate lists)
define_prims: [[PrimType or Shortcut], *[atom indices], ...]

# Examples

# Additional bond between atoms 4 and 7 (0-based indexing).
# All three lines below result in the same bond; the latter two use shortcuts.
define_prims: [[0, 4, 7]]
define_prims: [[B, 4, 7]]
define_prims: [[BOND, 4, 7]]

# Wrong specification (forgot outer list/brackets):
define_prims: [0, 4, 7]

# Also define an additional dihedral, beside the bond
define_prims: [[0, 4, 7], ["D", 0, 1, 2, 3]]
```

### 5.3.4 Freeze Atoms

All three Cartesian coordinates (X, Y, Z) of certain atoms can be frozen, so they always remain at their initial value. By setting `freeze_atoms_exclude` in `coord_kwargs`, frozen atoms can be excluded from the internal coordinate setup. By default frozen atoms are included.

```
geom:
  type: [type]
  fn: [fn]
  freeze_atoms: [*atom indices]
  coord_kwargs:
    freeze_atoms_exclude: False

# Example; fully freeze Cartesians of first and second atom.
geom:
```

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```

type: cart
fn: input.xyz
freeze_atoms: [0, 1]
coord_kwargs:
  freeze_atoms: True

```

### 5.3.5 Constraints

**Constraints beyond frozen atoms are currently only supported in conjunction with RIC (`type: redund`).** It is not (yet) possible to modify the value of the specified coordinate via YAML input; the internal coordinate is constrained at its initial value. The same syntax as for `define_prims` is used. If the coordinate of the requested constraint is not already defined, it will be defined subsequently. There is no need to also add the constrained coordinate to `define_prims`.

```

# General structure (nested list of coordinates)
constrain_prims: [[[PrimType or Shortcut], *[atom indices]], ...]

# Examples

# Constrain Cartesian coordinate of atom 0.
# Both lines result in the same constraint.
constrain_prims: [[XYZ, 0]]
constrain_prims: [[ATOM, 0]]

# Constrain only Cartesian X and Y component of atom 0.
constrain_prims: [[XY, 0]]

# Constraint bond between atoms 4 and 7 (0-based indexing).
# All three lines below result in the same constraint; the latter two use shortcuts.
constrain_prims: [[0, 4, 7]]
constrain_prims: [[B, 4, 7]]
constrain_prims: [[BOND, 4, 7]]

```

Constraining the Cartesian coordinates (X, Y and Z) of one atom does not affect the final energy of an optimization. **But constraining more than one atom does.**

Harmonic restraints to selected primitive internals can be specified in the `calc:` section (see the [Restraint](#) documentation).

### 5.3.6 Isotopes

Different isotope masses can be requested. The system works similar to Gaussians system. A list of pairs is expected, where the first number specifies the atom and the second number is either an integer or a float. If it is an integer, the isotope mass closest to this integer is looked up in an internal database. Floats are used as is.

```

# General structure (nested list of coordinates)
isotopes: [[[atom index], [new mass, integer/float], ...]]

# Modify the mass of atom with index 2 (hydrogen in this case)
# Both lines give identical results (deuterium).
# In the second line, the mass is given directly.

```

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```
isotopes: [[2, 2]]
isotopes: [[2, 2.014101778]]
```

Different isotope masses affect calculated frequencies and IRCs. Atoms can be fixed in IRC calculations by specifying a very high mass. **YAML does not recognize 1e9 as float**, take care to add a dot (**1.e9**).

```
# Fix atom 0 in IRC calculation.
isotopes: [[0, 1.e9]]
```

## 5.4 Geometry & RedundantCoords

The central class in pysisyphus, handling coordinates and delegating calculations to (external QC) codes, is the *Geometry* class, similar to ASE's *Atoms*.

```
class pysisyphus.Geometry.Geometry(atoms, coords, fragments=None, coord_type='cart',
                                   coord_kwargs=None, isotopes=None, freeze_atoms=None,
                                   comment="", name="")
```

```
__init__(atoms, coords, fragments=None, coord_type='cart', coord_kwargs=None, isotopes=None,
         freeze_atoms=None, comment="", name="")
```

Object representing atoms in a coordinate system.

The Geometry represents atoms and their positions in coordinate system. By default cartesian coordinates are used, but internal coordinates are also possible.

### Parameters

- **atoms** (*iterable*) -- Iterable of length N, containing element symbols.
- **coords** (*1d iterable*) -- 1d iterable of length 3N, containing the cartesian coordinates of N atoms.
- **fragments** (*dict, optional*) -- Dict with different keys denoting different fragments. The values contain lists of atom indices.
- **coord\_type** (*{"cart", "redund"}, optional*) -- Type of coordinate system to use. Right now cartesian (cart) and redundant (redund) are supported.
- **coord\_kwargs** (*dict, optional*) -- Dictionary containing additional arguments that get passed to the constructor of the internal coordinate class.
- **isotopes** (*iterable of pairs, optional*) -- Iterable of pairs consisting of 0-based atom index and either an integer or a float. If an integer is given the closest isotope mass will be selected. Given a float, this float will be directly used as mass.
- **freeze\_atoms** (*iterable of integers*) -- Specifies which atoms should remain fixed at their initial positions.
- **comment** (*str, optional*) -- Comment string.
- **name** (*str, optional*) -- Verbose name of the geometry, e.g. methanal or water. Used for printing

### align\_principal\_axes()

Align the principal axes to the cartesian axes.

<https://math.stackexchange.com/questions/145023>

**approximate\_radius()**

Approximate molecule radius from the biggest atom distance along an axis.

**as\_ase\_atoms()****as\_g98\_list()**

Returns data for fake Gaussian98 standard orientation output.

**Returns**

**g98\_list** -- List with one row per atom. Every row contains [center number, atomic number, atomic type (always 0 for now), X Y Z coordinates in Angstrom.

**Return type**

list

**as\_xyz(comment="", atoms=None, cart\_coords=None)**

Current geometry as a string in XYZ-format.

**Parameters**

- **comment** (*str*, *optional*) -- Will be written in the second line (comment line) of the XYZ-string.
- **cart\_coords** (*np.array*, *1d*, *shape* (3 \* *atoms.size*, )) -- Cartesians for dumping instead of *self.\_coords*.

**Returns**

**xyz\_str** -- Current geometry as string in XYZ-format.

**Return type**

str

**assert\_cart\_coords(coords)****assert\_compatibility(other)**

Assert that two Geometries can be subtracted from each other.

**Parameters**

**other** (*Geometry*) -- Geometry for comparison.

**atom\_indices()**

Dict with atom types as key and corresponding indices as values.

**Returns**

**inds\_dict** -- Unique atom types as keys, corresponding indices as values.

**Return type**

dict

**property atom\_types****atom\_xyz\_iter()****property atomic\_numbers****property bond\_sets****calc\_double\_ao\_overlap(geom2)****calc\_energy\_and\_forces()**

Force a calculation of the current energy and forces.

property `cart_coords`

property `cart_forces`

property `cart_gradient`

property `cart_hessian`

`center()`

property `center_of_mass`

Returns the center of mass.

**Returns**

**R** -- Center of mass.

**Return type**

np.array, shape(3, )

`center_of_mass_at(coords3d)`

Returns the center of mass at given coords3d.

**Parameters**

**coords3d** (np.array, shape(N, 3)) -- Cartesian coordinates.

**Returns**

**R** -- Center of mass.

**Return type**

np.array, shape(3, )

property `centroid`

Geometric center of the Geometry.

**Returns**

**R** -- Geometric center of the Geometry.

**Return type**

np.array, shape(3, )

`clear()`

Reset the object state.

property `comment`

```
coord_types = {'cart': None, 'cartesian': <class
'pysisyphus.intcoords.CartesianCoords.CartesianCoords'>, 'dlc': <class
'pysisyphus.intcoords.DLC.DLC'>, 'hdlc': <class 'pysisyphus.intcoords.DLC.HDLC'>,
'hredund': <class 'pysisyphus.intcoords.RedundantCoords.HybridRedundantCoords'>,
'mwcartesian': <class 'pysisyphus.intcoords.CartesianCoords.MWCartesianCoords'>,
'redund': <class 'pysisyphus.intcoords.RedundantCoords.RedundantCoords'>, 'tric':
<class 'pysisyphus.intcoords.RedundantCoords.TRIC'>}
```

property `coords`

1d vector of atomic coordinates.

**Returns**

**coords** -- 1d array holding the current coordinates.

**Return type**

np.array

**property coords3d**

Coordinates in 3d.

**Returns**

**coords3d** -- Coordinates of the Geometry as 2D array.

**Return type**

np.array

**property coords\_by\_type**

Coordinates in 3d by atom type and their corresponding indices.

**Returns**

- **cbt** (*dict*) -- Dictionary with the unique atom types of the Geometry as keys. It's values are the 3d coordinates of the corresponding atom type.
- **inds** (*dict*) -- Dictionary with the unique atom types of the Geometry as keys. It's values are the original indices of the 3d coordinates in the whole coords3d array.

**copy**(*coord\_type=None, coord\_kwargs=None*)

Returns a new Geometry object with same atoms and coordinates.

**Parameters**

- **coord\_type** (*str*) -- Desired coord\_type, defaults to current coord\_type.
- **coord\_kwargs** (*dict, optional*) -- Any desired coord\_kwargs that will be passed to the RedundantCoords object.

**Returns**

**geom** -- New Geometry object with the same atoms and coordinates.

**Return type**

*Geometry*

**copy\_all**(*coord\_type=None, coord\_kwargs=None*)**property covalent\_radii****del\_atoms**(*inds, \*\*kwargs*)**describe**()**dump\_xyz**(*fn*)**eckart\_projection**(*mw\_hessian, return\_P=False, full=False*)**property energy**

Energy of the current atomic configuration.

**Returns**

**energy** -- Energy of the current atomic configuration.

**Return type**

float

**fd\_coords3d\_gen**(*step\_size=0.001*)

Iterator returning 3d Cartesians for finite-differences.

**property forces**

Energy of the current atomic configuration.

**Returns**

**force** -- 1d array containing the forces acting on the atoms. Negative of the gradient.

**Return type**

np.array

**get\_energy\_and\_cart\_forces\_at**(*cart\_coords*)

**get\_energy\_and\_cart\_hessian\_at**(*cart\_coords*)

**get\_energy\_and\_forces\_at**(*coords*)

Calculate forces and energies at the given coordinates.

The results are not saved in the Geometry object.

**get\_energy\_at**(*coords*)

**get\_energy\_at\_cart\_coords**(*cart\_coords*)

**get\_fragments**(*regex*)

**get\_imag\_frequencies**(*hessian=None, thresh=1e-06*)

**get\_normal\_modes**(*cart\_hessian=None, full=False*)

Normal mode wavenumbers, eigenvalues and Cartesian displacements Hessian.

**get\_restart\_info**()

**get\_sphere\_radius**(*offset=4*)

**get\_subgeom**(*indices, coord\_type='cart', sort=False*)

Return a Geometry containing a subset of the current Geometry.

**Parameters**

- **indices** (*iterable of ints*) -- Atomic indices that the define the subset of the current Geometry.
- **coord\_type** (*str, ("cart", "redund"), optional*) -- Coordinate system of the new Geometry.

**Returns**

**sub\_geom** -- Subset of the current Geometry.

**Return type**

*Geometry*

**get\_subgeom\_without**(*indices, \*\*kwargs*)

**get\_temporary\_coords**(*coords*)

**get\_thermoanalysis**(*energy=None, cart\_hessian=None, T=298.15, p=101325, point\_group='c1'*)

**get\_trans\_rot\_projector**(*full=False*)

**property gradient**

Negative of the force.

**Returns**

**gradient** -- 1d array containing the negative of the current forces.

**Return type**

np.array

**property hessian**

Matrix of second derivatives of the energy in respect to atomic displacements.

**Returns**

**hessian** -- 2d array containing the second derivatives of the energy with respect to atomic/coordinate displacements depending on the type of coordinate system.

**Return type**

np.array

**property inertia\_tensor**

**property is\_analytical\_2d**

**jmol**(*atoms=None, cart\_coords=None*)

Show geometry in jmol.

**property layers**

**mass\_weigh\_hessian**(*hessian*)

**property masses**

**property masses\_rep**

**property mm\_inv**

Inverted mass matrix.

Returns a diagonal matrix containing the inverted atomic masses.

**property mm\_sqrt\_inv**

Inverted square root of the mass matrix.

**property moving\_atoms**

**moving\_atoms\_jmol**()

**property mw\_coords**

Mass-weighted coordinates.

**Returns**

**mw\_coords** -- 1d array containing the mass-weighted cartesian coordinates.

**Return type**

np.array

**property mw\_gradient**

Mass-weighted gradient.

**Returns**

**mw\_gradient** -- Returns the mass-weighted gradient.

**Return type**

np.array

**property mw\_hessian**

Mass-weighted hessian.

**Returns****mw\_hessian** -- 2d array containing the mass-weighted hessian  $M^{(-1/2)} H M^{(-1/2)}$ .**Return type**

np.array

**principal\_axes\_are\_aligned()**

Check if the principal axes are aligned with the cartesian axes.

**Returns****aligned** -- Wether the principal axes are aligned or not.**Return type**

bool

**reparametrize()****reset\_coords**(*new\_typed\_prims=None*)**rmsd**(*geom*)**rotate**(*copy=False*)**set\_calculator**(*calculator, clear=True*)

Reset the object and set a calculator.

**set\_coord**(*ind, coord*)

Set a coordinate by index.

**Parameters**

- **ind** (*int*) -- Index in of the coordinate to set in the self.coords array.
- **coord** (*float*) -- Coordinate value.

**set\_coords**(*coords, cartesian=False, update\_constraints=False*)**set\_h5\_hessian**(*fn*)**set\_restart\_info**(*restart\_info*)**set\_results**(*results*)

Save the results from a dictionary.

**Parameters****results** (*dict*) -- The keys in this dict will be set as attributes in the current object, with the corresponding item as value.**standard\_orientation()****property sum\_formula****property total\_mass**

**unweight\_mw\_hessian**(*mw\_hessian*)

Unweight a mass-weighted hessian.

**Parameters**

**mw\_hessian** (*np.array*) -- Mass-weighted hessian to be unweighted.

**Returns**

**hessian** -- 2d array containing the hessian.

**Return type**

*np.array*

**without\_hydrogens**()

**zero\_frozen\_forces**(*cart\_forces*)

`pysisyphus.Geometry.get_trans_rot_projector`(*cart\_coords*, *masses*, *full=False*)

`pysisyphus.Geometry.get_trans_rot_vectors`(*cart\_coords*, *masses*, *rot\_thresh=1e-06*)

Vectors describing translation and rotation.

These vectors are used for the Eckart projection by constructing a projector from them.

See Martin J. Field - A Practical Introduction to the simulation of Molecular Systems, 2007, Cambridge University Press, Eq. (8.23), (8.24) and (8.26) for the actual projection.

See also <https://chemistry.stackexchange.com/a/74923>.

**Parameters**

- **cart\_coords** (*np.array*, *1d*, *shape (3 \* atoms.size, )*) -- Atomic masses in amu.
- **masses** (*iterable*, *1d*, *shape (atoms.size, )*) -- Atomic masses in amu.

**Returns**

**ortho\_vecs** -- 2d array containing row vectors describing translations and rotations.

**Return type**

*np.array(6, 3\*atoms.size)*

`pysisyphus.Geometry.inertia_tensor`(*coords3d*, *masses*)

Inertia tensor.

$x^2 \quad xy \quad xz \quad |$

$(x \ y \ z)^T \cdot (x \ y \ z) = | \quad xy \quad y^2 \quad yz \quad |$

$xz \quad yz \quad z^2 \quad |$

Basic infrastructure to work with internal coordinates is provided by *RedundantCoords*.

`class pysisyphus.intcoords.RedundantCoords.HybridRedundantCoords`(\*args, \*\*kwargs)

```

class pysisyphus.intcoords.RedundantCoords.RedundantCoords(atoms, coords3d, masses=None,
bond_factor=1.3, typed_prims=None,
define_prims=None,
constrain_prims=None,
freeze_atoms=None,
freeze_atoms_exclude=False,
define_for=None, bonds_only=False,
check_bends=True, rebuild=True,
bend_min_deg=15,
dihed_max_deg=175,
lb_min_deg=175, weighted=False,
min_weight=0.3,
svd_inv_thresh=0.000316,
recalc_B=False, tric=False,
hybrid=False, hbond_angles=False)

```

**property B**

Wilson B-Matrix

**property B\_inv**

Generalized inverse of the Wilson B-Matrix.

**property B\_inv\_prim**

Generalized inverse of the primitive Wilson B-Matrix.

**property B\_prim**

Wilson B-Matrix

**property Bt\_inv**

Transposed generalized inverse of the Wilson B-Matrix.

**property Bt\_inv\_prim**

Transposed generalized inverse of the primitive Wilson B-Matrix.

**property C**

Diagonal matrix. Entries for constraints are set to one.

**property P**

Projection matrix onto B. See [1] Eq. (4).

**backtransform\_hessian(*redund\_hessian, int\_gradient=None*)**

Transform Hessian in internal coordinates to Cartesians.

**property bend\_atom\_indices****property bend\_indices****property bond\_atom\_indices****property bond\_indices****property bond\_typed\_prims****property cartesian\_indices****clear()**

**property constrained\_indices**

**property coords**

**property coords3d**

**property dihedral\_atom\_indices**

**property dihedral\_indices**

**eval**(*coords3d*, *attr=None*)

**get\_K\_matrix**(*int\_gradient=None*)

**get\_index\_of\_typed\_prim**(*typed\_prim*)  
Index in self.typed\_prims for the supplied typed\_prim.

**get\_prim\_internals\_by\_indices**(*indices*)

**inv\_B**(*B*)

**inv\_Bt**(*B*)

**property linear\_bend\_indices**

**log**(*message*)

**log\_int\_grad\_msg**(*int\_gradient*)

**property outofplane\_indices**

**property prim\_coords**

**property prim\_indices\_set**

**property prim\_internals**

**property primitives**

**print\_typed\_prims**()

**project\_hessian**(*H*, *shift=1000*)  
Expects a hessian in internal coordinates. See Eq. (11) in [1].

**project\_vector**(*vector*)  
Project supplied vector onto range of B.

**return\_inds**(*slice\_*)

**property rotation\_indices**

**set\_inds\_from\_typed\_prims**(*typed\_prims*)

**set\_primitive\_indices**(*atoms*, *coords3d*)

**transform\_forces**(*cart\_forces*)  
Combination of Eq. (9) and (11) in [1].

**transform\_hessian**(*cart\_hessian*, *int\_gradient=None*)  
Transform Cartesian Hessian to internal coordinates.

```
transform_int_step(int_step, update_constraints=False, pure=False)
```

```
property translation_indices
```

```
property typed_prims
```

```
class pysisyphus.intcoords.RedundantCoords.TRIC(*args, **kwargs)
```

## 5.5 Related Literature

1. The efficient optimization of molecular geometries using redundant internal coordinates
2. The generation and use of delocalized internal coordinates in geometry optimization
3. Geometry optimization in redundant internal coordinates
4. Geometry optimization made simple with translation and rotation coordinates



## CALCULATORS

Exploring potential energy surfaces (PESs) requires calculation of energies and its derivatives (gradients, Hessian matrices).

For testing purposes and method development, pysisyphus implements various analytical 2d potentials, as they allow fast evaluations of the aforementioned quantities. Actual (production) calculations are carried out by wrapping existing quantum chemistry codes (ORCA, TURBOMOLE, Gaussian etc.) and delegating the required calculations to them. Pysisyphus generates all necessary inputs, executes the QC code and parses their output for the requested quantities.

Furthermore pysisyphus provides several "meta"-calculators which wrap other (actual) calculators, to modify calculated energies and forces. Examples for this are the *Dimer* calculator, used for carrying out transition state searches or the *ONIOM* calculator, allowing multi-level calculations comprising different levels of theory.

External forces, e.g. a restraining spherical potential or harmonic restraints on primitive internal coordinates (stretches, bends, torsion) can be applied with *ExternalPotential*.

## 6.1 YAML input

Possible keywords for the YAML input can be derived from inspecting the possible arguments of the *Calculator* baseclass (see below) and the possible arguments of the respective calculator, e.g. ORCA or XTB.

The most commonly used keywords, derived from the *Calculator* baseclass are *mem*, handling the requested memory per core in MB, *pal*, handling the number of requested CPU cores, *charge*, the total charge of the system and *mult*, the systems multiplicity.

For (excited state, ES) calculations carried out with calculators derived from *OverlapCalculator* additional keywords are possible. The most common keywords controlling ES calculations are *track*, activating ES tracking, *ovlp\_type*, selecting the tracking algorithm and *ovlp\_with*, handling the selection of the reference cycle.

An example input highlighting the most important keywords is shown below.

```
geom:
  [... omitted ...]
calc:
  type: orca                # Calculator type, e.g. g09/g16/openmolcas/
                           # orca/pyscf/turbomole/dftb+/mopac/psi4/xtb

  pal: 1                   # Number of CPU cores
  mem: 1000                # Memory per core
  charge: 0                # Charge
  mult: 1                  # Multiplicity
  # Keywords for ES calculations
  track: False             # Activate ES tracking
```

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```

ovlp_type: tden                # Tracking algorithm
ovlp_with: previous            # Reference cycle selection
# Additional calculator specific keywords
[... omitted ...]
opt:
[... omitted ...]

```

## 6.2 Calculator base classes

```

class pysisyphus.calculators.Calculator.Calculator(
    calc_number=0, charge=0, mult=1,
    base_name='calculator', pal=1, mem=1000,
    check_mem=True, retry_calc=1,
    last_calc_cycle=None, clean_after=True,
    out_dir='qm_calcs')

```

```

__init__(calc_number=0, charge=0, mult=1, base_name='calculator', pal=1, mem=1000,
    check_mem=True, retry_calc=1, last_calc_cycle=None, clean_after=True, out_dir='qm_calcs')

```

Base-class of all calculators.

Meant to be extended.

### Parameters

- **calc\_number** (*int*, *default=0*) -- Identifier of the Calculator. Used in distinguishing it from other Calculators, e.g. in ChainOfStates calculations. Also used in the creation of filenames.
- **charge** (*int*, *default=0*) -- Molecular charge.
- **mult** (*int*, *default=1*) -- Molecular multiplicity (1 = singlet, 2 = doublet, ...)
- **base\_name** (*str*, *default=calculator*) -- Generated filenames will start with this string.
- **pal** (*int*, *default=1*) -- Positive integer that gives the number of physical cores to use on 1 node.
- **mem** (*int*, *default=1000*) -- Memory per core in MB. The total amount of memory is given as mem\*pal.
- **check\_mem** (*bool*, *default=True*) -- Whether to adjust the requested memory if too much is requested.
- **retry\_calc** (*int*, *default=0*) -- Number of additional retries when calculation failed.
- **last\_calc\_cycle** (*int*) -- Internal variable used in restarts.
- **clean\_after** (*bool*) -- Delete temporary directory were calculations were executed after a calculation.
- **out\_dir** (*str*) -- Path that is prepended to generated filenames.

**clean**(*path*)

Delete the temporary directory.

### Parameters

**path** (*Path*) -- Directory to delete.

**conf\_key = None**

**get\_cmd**(*key='cmd'*)

**get\_energy**(*atoms, coords*)

Meant to be extended.

**get\_forces**(*atoms, coords*)

Meant to be extended.

**get\_hessian**(*atoms, coords*)

Meant to be extended.

**get\_restart\_info**()

Return a dict containing chkfiles.

**Returns**

**restart\_info** -- Dictionary holding the calculator state. Used for restoring calculators in restarted calculations.

**Return type**

dict

**keep**(*path*)

Backup calculation results.

**Parameters**

**path** (*Path*) -- Temporary directory of the calculation.

**Returns**

**kept\_fns** -- Dictionary holding the filenames that were backed up. The keys correspond to the type of file.

**Return type**

dict

**log**(*message=""*)

Write a log message.

Wraps the logger variable.

**Parameters**

**message** (*str*) -- Message to be logged.

**make\_fn**(*name, counter=None, return\_str=False*)

Make a full filename.

Return a full filename including the calculator name and the current counter given a suffix.

**Parameters**

- **name** (*str*) -- Suffix of the filename.
- **counter** (*int, optional*) -- If not given use the current calc\_counter.
- **return\_str** (*int, optional*) -- Return a string instead of a Path when True.

**Returns**

**fn** -- Filename.

**Return type**

str

**property name**

**prepare**(*inp*)

Prepare a temporary directory and write input.

Similar to `prepare_path`, but the input is also written into the prepared directory.

**inp**

[*str*] Input to be written into the file `self.inp_fn` in the prepared directory.

**Returns**

**path** -- Prepared directory.

**Return type**

Path

**prepare\_coords**(*atoms, coords*)

Get 3d coords in Angstrom.

Reshape internal 1d coords to 3d and convert to Angstrom.

**Parameters**

- **atoms** (*iterable*) -- Atom descriptors (element symbols).
- **coords** (*np.array, 1d*) -- 1D-array holding coordinates in Bohr.

**Returns**

**coords** -- 3D-array holding coordinates in Angstrom.

**Return type**

np.array, 3d

**prepare\_input**(*atoms, coords, calc\_type*)

Meant to be extended.

**prepare\_path**(*use\_in\_run=False*)

Get a temporary directory handle.

Create a temporary directory that can later be used in a calculation.

**Parameters**

**use\_in\_run** (*bool, option*) -- Sets the internal variable `self.path_already_prepared` that is later read by `self.run()`. No new temporary directory will be created in `self.run()`.

**Returns**

**path** -- Prepared directory.

**Return type**

Path

**prepare\_pattern**(*raw\_pat*)

Prepare globs.

Transforms an entry of `self.to_keep` into a glob and a key suitable for the use in `self.keep()`.

**Parameters**

**raw\_pat** (*str*) -- Entry of `self.to_keep`

**Returns**

- **pattern** (*str*) -- Glob that can be used in `Path.glob()`

- **multi** (*bool*) -- Flag if glob may match multiple files.
- **key** (*str*) -- A key to be used in the `kept_fns` dict.

**prepare\_turbo\_coords**(*atoms, coords*)

Get a Turbomole coords string.

**Parameters**

- **atoms** (*iterable*) -- Atom descriptors (element symbols).
- **coords** (*np.array, 1d*) -- 1D-array holding coordinates in Bohr.

**Returns**

**coords** -- String holding coordinates in Turbomole coords format.

**Return type**

str

**prepare\_xyz\_string**(*atoms, coords*)

Returns a xyz string in Angstrom.

**Parameters**

- **atoms** (*iterable*) -- Atom descriptors (element symbols).
- **coords** (*np.array, 1d*) -- 1D-array holding coordinates in Bohr.

**Returns**

**xyz\_str** -- Coordinates in .xyz format.

**Return type**

string

**print\_capabilities**()

**print\_out\_fn**(*path*)

Print calculation output.

Prints the output of a calculator after a calculation.

**Parameters**

**path** (*Path*) -- Temporary directory of the calculation.

**run**(*inp, calc, add\_args=None, env=None, shell=False, hold=False, keep=True, cmd=None, inc\_counter=True, run\_after=True, parser\_kwargs=None, symlink=True*)

Run a calculation.

The bread-and-butter method to actually run an external quantum chemistry code.

**Parameters**

- **inp** (*str*) -- Input for the external program that is written to the temp-dir.
- **calc** (*str, hashable*) -- Key (and more or less type of calculation) to select the right parsing function from `self.parser_funcs`.
- **add\_args** (*iterable, optional*) -- Additional arguments that will be appended to the program call.
- **env** (*Environment, optional*) -- A potentially modified environment for the subprocess call.
- **shell** (*bool, optional*) -- Use a shell to execute the program call. Need for Turbomole were we chain program calls like `dscf; escf`.

- **hold** (*bool, optional*) -- Wether to remove the temporary directory after the calculation.
- **keep** (*bool, optional*) -- Wether to backup files as specified in `self.to_keep()`. Usually you want this.
- **cmd** (*str or iterable, optional*) -- Overwrites `self.base_cmd`.
- **inc\_counter** (*bool, optional*) -- Wether to increment the counter after a calculation.

**Returns**

**results** -- Dictionary holding all applicable results of the calculations like the energy, a forces vector and/or excited state energies from TDDFT.

**Return type**

dict

**run\_after** (*path*)

Meant to be extended.

This method is called after a calculation was done, but before entering `self.keep()` and `self.clean()`. Can be used to call tools like `formchk` or `ricctools`.

**set\_restart\_info** (*restart\_info*)

Sets restart information (`chkfiles` etc.) on the calculator.

**Parameters**

**restart\_info** (*dict*) -- Dictionary holding the calculator state. Used for restoring calculators in restarted calculations.

**verify\_chkfiles** (*chkfiles*)

Checks if given `chkfiles` exist and return them as Paths

**Parameters**

**chkfiles** (*dict*) -- Dictionary holding the `chkfiles`. The keys correspond to the attribute names, the values are str's holding the (potentially full) filename (path).

**Returns**

**paths** -- Dictionary of Paths.

**Return type**

dict

## 6.3 OverlapCalculator base class

```
class pysisyphus.calculators.OverlapCalculator.NTos(ntos, lambdas)
```

Bases: tuple

**lambdas**

Alias for field number 1

**ntos**

Alias for field number 0

```
class pysisiphus.calculators.OverlapCalculator.OverlapCalculator(*args, track=False,
                                                                ovlp_type='tden',
                                                                double_mol=False,
                                                                ovlp_with='previous',
                                                                XY='X+Y', adapt_args=(0.5,
                                                                0.3, 0.6), use_ntos=4,
                                                                pr_nto=False, nto_thresh=0.3,
                                                                cdds=None, orient="",
                                                                dump_fn='overlap_data.h5',
                                                                h5_dump=False, ncore=0,
                                                                conf_thresh=0.001,
                                                                dyn_roots=0, mos_ref='cur',
                                                                mos_renorm=True, **kwargs)
```

Bases: *Calculator*

```
H5_MAP = {'all_energies': 'all_energies_list', 'ci_coeffs': 'ci_coeff_list',
          'coords': 'coords_list', 'mo_coeffs': 'mo_coeff_list', 'ref_roots':
          'reference_roots', 'roots': 'roots_list'}
```

```
OVL_P_TYPE_VERBOSE = {'nto': 'natural transition orbital overlap', 'nto_org':
                      'original natural transition orbital overlap', 'tden': 'transition density matrix
                      overlap', 'wf': 'wavefunction overlap'}
```

```
VALID_CDDS = (None, 'calc', 'render')
```

```
VALID_KEYS = ['wf', 'tden', 'nto', 'nto_org']
```

```
VALID_XY = ('X', 'X+Y', 'X-Y')
```

```
blowup_ci_coeffs(ci_coeffs)
```

```
calc_cdd_cube(root, cycle=-1)
```

```
calculate_state_ntos(state_ci_coeffs, mos)
```

```
property data_model
```

```
dump_overlap_data()
```

```
static from_overlap_data(h5_fn, set_wfow=False)
```

```
get_h5_group()
```

```
get_indices(indices=None)
```

A new root is determined by selecting the overlap matrix row corresponding to the reference root and checking for the root with the highest overlap (at the current geometry).

The overlap matrix is usually formed by a double loop like:

```
overlap_matrix = np.empty((ref_states, cur_states)) for i, ref_state in enumerate(ref_states):
```

```
    for j, cur_state in enumerate(cur_states):
        overlap_matrix[i, j] = make_overlap(ref_state, cur_state)
```

So the reference states run along the rows. That's why the ref\_state index comes first in the 'indices' tuple.

```
static get_mo_norms(mo_coeffs, ao_ovlp)
```

MOs are in rows.

**get\_nto\_overlaps**(*indices=None, ao\_ovlp=None, org=False*)

**get\_orbital\_matrices**(*indices=None, ao\_ovlp=None*)

Return MO coefficients and AO overlaps for the given indices.

If not provided, a AO overlap matrix is constructed from one of the MO coefficient matrices (controlled by `self.mos_ref`). Also, if requested one of the two MO coefficient matrices is re-normalized.

**get\_ref\_mos**(*ref\_mo\_coeffs, cur\_mo\_coeffs*)

**static get\_sao\_from\_mo\_coeffs**(*mo\_coeffs*)

Recover AO overlaps from given MO coefficients.

For MOs in the columns of `mo_coeffs`:

$$S_{AO} = C^T C^{-1} S_{AO} C = C^T (S_{AO} C)^T = C^T C^T S_{AO}^T = C^T C^T S_{AO} C = I$$

Here, MOs are expected to be in rows of `mo_coeffs`, yielding

$$C S_{AO} C^T = I$$

**get\_sao\_from\_mo\_coeffs\_and\_dump**(*mo\_coeffs*)

**get\_tden\_overlaps**(*indices=None, ao\_ovlp=None*)

**get\_wf\_overlaps**(*indices=None, ao\_ovlp=None*)

**nto\_org\_overlaps**(*ntos\_1, ntos\_2, ao\_ovlp, nto\_thresh=0.3*)

**nto\_overlaps**(*ntos\_1, ntos\_2, ao\_ovlp*)

**prepare\_overlap\_data**(*path*)

Implement calculator specific parsing of MO coefficients and CI coefficients here. Should return a filename pointing to TURBOMOLE like `mos`, a MO coefficient array and a CI coefficient array.

**render\_cdd\_cube**()

**static renorm\_mos**(*mo\_coeffs, ao\_ovlp*)

**property roots\_number**

**set\_ntos**(*mo\_coeffs, ci\_coeffs*)

**set\_wfow**(*ci\_coeffs*)

**store\_overlap\_data**(*atoms, coords, path=None, overlap\_data=None*)

**tden\_overlap\_with\_calculator**(*calc, ao\_ovlp=None*)

**tden\_overlaps**(*mo\_coeffs1, ci\_coeffs1, mo\_coeffs2, ci\_coeffs2, ao\_ovlp*)

#### Parameters

- **mo\_coeffs1** (*ndarray, shape (MOs, AOs)*) -- MO coefficient matrix. One row per MO, one column per basis function. Usually square.
- **mo\_coeffs2** (*ndarray*) -- See `mo_coeffs1`.
- **ci\_coeffs1** (*ndarray, shape (occ. MOs, MOs)*) -- CI-coefficient matrix.
- **ci\_coeffs2** (*ndarray, shape (occ. MOs, MOs)*) -- See `ci_coeffs1`.
- **ao\_ovlp** (*ndarray, shape (AOs1, AOs2)*) -- Double molecule AO overlaps.

**track\_root**(*ovlp\_type=None*)

Check if a root flip occurred compared to the previous cycle by calculating the overlap matrix wrt. a reference cycle.

**update\_array\_dims**()

Assure consistent array shapes when the number of calculated roots changed.

**wf\_overlap\_with\_calculator**(*calc, ao\_ovlp=None*)

**wf\_overlaps**(*mo\_coeffs1, ci\_coeffs1, mo\_coeffs2, ci\_coeffs2, ao\_ovlp=None*)

`pysisyphus.calculators.OverlapCalculator.get_data_model`(*exc\_state\_num, occ\_mo\_num, virt\_mo\_num, ovlp\_type, atoms, max\_cycles*)

## 6.4 Calculators with Excited state capabilities

### 6.4.1 Gaussian09

`class pysisyphus.calculators.Gaussian09.Gaussian09`(*\*args, \*\*kwargs*)

`conf_key = 'gaussian09'`

### 6.4.2 Gaussian16

`class pysisyphus.calculators.Gaussian16.Gaussian16`(*route, gbs="", gen="", keep\_chk=False, stable="", fchk=None, \*\*kwargs*)

Bases: `OverlapCalculator`

`conf_key = 'gaussian16'`

`get_chkfiles`()

`get_energy`(*atoms, coords, \*\*prepare\_kwargs*)

Meant to be extended.

`get_forces`(*atoms, coords, \*\*prepare\_kwargs*)

Meant to be extended.

`get_hessian`(*atoms, coords, \*\*prepare\_kwargs*)

Meant to be extended.

`keep`(*path*)

Backup calculation results.

#### Parameters

`path` (*Path*) -- Temporary directory of the calculation.

#### Returns

`kept_fns` -- Dictionary holding the filenames that were backed up. The keys correspond to the type of file.

#### Return type

dict

**make\_exc\_str()**

**make\_fchk**(*path*)

**make\_gbs\_str()**

**parse\_635r\_dump**(*dump\_path*, *roots*, *nmos*)

**parse\_charges**(*path=None*)

**parse\_double\_mol**(*path*, *out\_fn=None*)

**static parse\_fchk**(*fchk\_path*, *keys*)

**parse\_force**(*path*)

**parse\_hessian**(*path*)

**parse\_keyword**(*text*)

**parse\_log**(\**args*, \*\**kwargs*)

**parse\_stable**(*path*)

**parse\_tddft**(*path*)

**prepare\_input**(*atoms*, *coords*, *calc\_type*, *did\_stable=False*, *point\_charges=None*)

Meant to be extended.

**prepare\_overlap\_data**(*path*)

Implement calculator specific parsing of MO coefficients and CI coefficients here. Should return a filename pointing to TURBOMOLE like mos, a MO coefficient array and a CI coefficient array.

**reuse\_data**(*path*)

**run\_after**(*path*)

Meant to be extended.

This method is called after a calculation was done, but before entering `self.keep()` and `self.clean()`. Can be used to call tools like `formchk` or `ricctools`.

**run\_calculation**(*atoms*, *coords*, \*\**prepare\_kwargs*)

**run\_double\_mol\_calculation**(*atoms*, *coords1*, *coords2*)

**run\_rwf\_dump**(*path*, *rwf\_index*, *chk\_path=None*)

**run\_stable**(*atoms*, *coords*, \*\**prepare\_kwargs*)

**set\_chkfiles**(*chkfiles*)

### 6.4.3 OpenMolcas

**class** pysisiphus.calculators.OpenMolcas.**OpenMolcas**(*basis, inporb, roots, mdrlxroot, supsym=None, track=True, \*\*kwargs*)

Bases: *Calculator*

**build\_rassi\_str**()

**build\_supsym\_str**(*supsym*)

Can handle only one subgroup for now.

**conf\_key** = 'openmolcas'

**get\_energy**(*atoms, coords*)

Meant to be extended.

**get\_forces**(*atoms, coords*)

Meant to be extended.

**get\_pal\_env**()

**keep**(*path*)

Backup calculation results.

**Parameters**

**path** (*Path*) -- Temporary directory of the calculation.

**Returns**

**kept\_fns** -- Dictionary holding the filenames that were backed up. The keys correspond to the type of file.

**Return type**

dict

**parse\_energies**(*text*)

**parse\_gradient**(*path*)

**parse\_rassi\_track**(*path*)

**prepare\_coords**(*atoms, coords*)

Get 3d coords in Angstrom.

Reshape internal 1d coords to 3d and convert to Angstrom.

**Parameters**

- **atoms** (*iterable*) -- Atom descriptors (element symbols).
- **coords** (*np.array, 1d*) -- 1D-array holding coordinates in Bohr.

**Returns**

**coords** -- 3D-array holding coordinates in Angstrom.

**Return type**

np.array, 3d

**prepare\_input**(*atoms, coords*)

Meant to be extended.

**reattach**(*last\_calc\_cycle*)

## 6.4.4 ORCA 4.2.1 / 5.0.1

```
class pysisiphus.calculators.ORCA.ORCA(keywords, blocks="", gbw=None, do_stable=False,
                                       numfreq=False, **kwargs)
```

Bases: *OverlapCalculator*

```
__init__(keywords, blocks="", gbw=None, do_stable=False, numfreq=False, **kwargs)
```

ORCA calculator.

Wrapper for creating ORCA input files for energy, gradient and Hessian calculations. The PAL and memory inputs must not be given in the keywords and/or blocks, as they are handled by the 'pal' and 'memory' arguments.

### Parameters

- **keywords** (*str*) -- Keyword line, as normally given in ORCA, excluding the leading "!".
- **blocks** (*str*, *optional*) -- ORCA block input(s), e.g. for TD-DFT calculations (%tddft ... end). As the blocks start with a leading "%", wrapping the input in quotes (") is required, otherwise the parsing will fail.
- **gbw** (*str*, *optional*) -- Path to an input gbw file, which will be used as initial guess for the first calculation. Will be overridden later, with the path to the gbw file of a previous calculation.
- **do\_stable** (*bool*, *optional*) -- Run stability analysis until a stable wavefunction is obtained, before every calculation.
- **numfreq** (*boo*, *optional*) -- Use numerical frequencies instead of analytical ones.
- **mem** (*int*) -- Mememory per core in MB.

```
check_termination(*args, **kwargs)
```

```
clean_tmp(path)
```

```
conf_key = 'orca'
```

```
get_block_str()
```

```
get_chkfiles()
```

```
get_energy(atoms, coords, **prepare_kwargs)
```

Meant to be extended.

```
get_forces(atoms, coords, **prepare_kwargs)
```

Meant to be extended.

```
get_hessian(atoms, coords, **prepare_kwargs)
```

Meant to be extended.

```
get_moinp_str(gbw)
```

```
get_stable_wavefunction(atoms, coords)
```

```
keep(path)
```

Backup calculation results.

### Parameters

**path** (*Path*) -- Temporary directory of the calculation.

**Returns**

**kept\_fns** -- Dictionary holding the filenames that were backed up. The keys correspond to the type of file.

**Return type**

dict

**parse\_all\_energies**(*text=None, triplets=None*)

**static parse\_atoms\_coords**(*inp, \*args, \*\*kwargs*)

**parse\_cis**(*cis*)

**Read binary CI vector file from ORCA.**

Adapted from TheoDORE 1.7.1, Authors: S. Mai, F. Plasser <https://sourceforge.net/p/theodore-qc>

**parse\_energy**(*path*)

**parse\_engrad**(*path*)

**static parse\_engrad\_info**(*inp, \*args, \*\*kwargs*)

**parse\_gbw**(*gbw\_fn*)

Adapted from <https://orcaforum.kofo.mpg.de/viewtopic.php?f=8&t=3299>

The first 5 long int values represent pointers into the file:

Pointer @+0: Internal ORCA data structures Pointer @+8: Geometry Pointer @+16: BasisSet Pointer @+24: Orbitals Pointer @+32: ECP data

**static parse\_hess\_file**(*inp, \*args, \*\*kwargs*)

**parse\_hessian**(*path*)

**parse\_mo\_numbers**(*out\_fn*)

**parse\_stable**(*path*)

**prepare\_input**(*atoms, coords, calc\_type, point\_charges=None, do\_stable=False*)

Meant to be extended.

**prepare\_overlap\_data**(*path*)

Implement calculator specific parsing of MO coefficients and CI coefficients here. Should return a filename pointing to TURBOMOLE like mos, a MO coefficient array and a CI coefficient array.

**reattach**(*last\_calc\_cycle*)

**run\_after**(*path*)

Meant to be extended.

This method is called after a calculation was done, but before entering `self.keep()` and `self.clean()`. Can be used to call tools like `formchk` or `ricctools`.

**run\_calculation**(*atoms, coords, \*\*prepare\_kwargs*)

Basically some kind of dummy method that can be called to execute ORCA with the stored cmd of this calculator.

**set\_chkfiles**(*chkfiles*)

**set\_mo\_coeffs**(*mo\_coeffs=None, gbw=None*)

```
static set_mo_coeffs_in_gbw(in_gbw_fn, out_gbw_fn, mo_coeffs)
```

See `self.parse_gbw`.

```
store_and_track(results, func, atoms, coords, **prepare_kwargs)
```

```
pysisyphus.calculators.ORCA.make_sym_mat(table_block)
```

```
pysisyphus.calculators.ORCA.save_orca_pc_file(point_charges, pc_fn, hardness=None)
```

## 6.4.5 PySCF 1.7.6

## 6.4.6 Turbomole 7.x

Pysisyphus does not implement a wrapper for *define*, so the user has to manually prepare a directory containing a valid control file. An automated *define* wrapper, restricted to ground state functionality, is available via the [QCEngine project](#), to which I contributed the Turbomole harness.

Care should be taken to include only the minimum amount of necessary files in the *control\_path* directory, e.g., (*auxbasis*, *basis*, *control*, *coord*, *mos*) for a closed-shell calculation using RI. **A gradient file must not be present in control\_path, as well as other subdirectories and files with .out extension.** The *coord* file, while not strictly required, should be kept too, to facilitate testing of the setup with standalone Turbomole.

It may be a good idea to pre-converge the calculation in *control\_path*, to see if the setup is correct and actually works. Resulting files like *energy*, *statistics* can be deleted; *mos* should be kept, as the converged MOs are reused in pysisyphus.

If an excited-state optimization is desired, care has to be taken, to include `$exopt [n]` for TD-DFT/TDA or the **geoopt state=[n]** (ricc2)! Tracking of excited states is currently possible for closed shell *egrad* and *ricc2* calculations.

The current implementation was tested against Turbomole 7.4.1 and QCEngine 0.19.0. Please see [examples/complex/11\\_turbomole\\_gs\\_tsopt](#) for a full example where Turbomole is utilized in a growing string calculation. The same example, using QCEngine, is found in [examples/complex/12\\_qcengine\\_turbomole\\_gs\\_tsopt](#). MOs are not reused with the QCEngine calculator, so the native pysisyphus calculator is probably faster.

```
class pysisyphus.calculators.Turbomole.Turbomole(control_path, root=None, double_mol_path=None,
                                               **kwargs)
```

Bases: *OverlapCalculator*

```
append_control(to_append, log_msg="", **kwargs)
```

```
conf_key = 'turbomole'
```

```
get_chkfiles()
```

```
get_energy(atoms, coords, **prepare_kwargs)
```

Meant to be extended.

```
get_forces(atoms, coords, cmd=None, **prepare_kwargs)
```

Meant to be extended.

```
get_hessian(atoms, coords, **prepare_kwargs)
```

Meant to be extended.

```
get_pal_env()
```

```
get_ricc2_root(text)
```

**keep**(*path*)

Backup calculation results.

**Parameters**

**path** (*Path*) -- Temporary directory of the calculation.

**Returns**

**kept\_fns** -- Dictionary holding the filenames that were backed up. The keys correspond to the type of file.

**Return type**

dict

**make\_molden**(*path*)**parse\_cc2\_vectors**(*ccre*)**parse\_double\_mol**(*path*)

Parse a double molecule overlap matrix from Turbomole output to be used with WFOWrapper.

**parse\_energy**(*path*)**parse\_force**(*path*)**parse\_gs\_energy**()

Several places are possible: \$subenergy from control file total energy from turbomole.out Final MP2 energy from turbomole.out with ADC(2) Final CC2 energy from turbomole.out with CC(2)

**parse\_hessian**(*path, fn=None*)**parse\_mos**()**parse\_td\_vectors**(*text*)

For TDA calculations only the X vector is present in the *ciss\_a/etc.* file. In TDDFT calculations there are twice as much items compared with TDA. The first half corresponds to (X+Y) and the second half to (X-Y). X can be calculated as  $X = ((X+Y)+(X-Y))/2$ . Y is then given as  $Y = (X+Y)-X$ . The normalization can then be checked as

```
np.concatenate((X, Y)).dot(np.concatenate((X, -Y)))
```

and should be 1.

**prepare\_input**(*atoms, coords, calc\_type, point\_charges=None*)

To rectify this we have to construct the basemcd dynamically and construct it ad hoc. We could set a RI flag in the beginning and select the correct scf binary here from it. Then we select the following binary on demand, e.g. aoforce or rdgrad or egrad etc.

**prepare\_overlap\_data**(*path*)

Implement calculator specific parsing of MO coefficients and CI coefficients here. Should return a filename pointing to TURBOMOLE like mos, a MO coefficient array and a CI coefficient array.

**prepare\_point\_charges**(*point\_charges*)

```
$point_charges <x> <y> <z> <q>
```

**prepare\_td**(*text*)**run\_after**(*path*)

Meant to be extended.

This method is called after a calculation was done, but before entering `self.keep()` and `self.clean()`. Can be used to call tools like formchk or ricctools.

**run\_calculation**(*atoms, coords*)

Basically some kind of dummy method that can be called to execute Turbomole with the stored cmd of this calculator.

**run\_double\_mol\_calculation**(*atoms, coords1, coords2*)

**set\_chkfiles**(*chkfiles*)

**set\_occ\_and\_mo\_nums**(*text*)

**store\_and\_track**(*results, func, atoms, coords, \*\*prepare\_kwargs*)

**sub\_control**(*pattern, repl, log\_msg='', \*\*kwargs*)

### 6.4.7 DFTB+ 20.x

**class** pysisyphus.calculators.DFTBp.**DFTBp**(*parameter, \*args, slakos=None, root=None, \*\*kwargs*)

Bases: *OverlapCalculator*

**conf\_key** = 'dftbp'

**get\_energy**(*atoms, coords, \*\*prepare\_kwargs*)

Meant to be extended.

**static get\_excited\_state\_str**(*root, forces=False*)

**get\_forces**(*atoms, coords, \*\*prepare\_kwargs*)

Meant to be extended.

**static get\_gen\_str**(*atoms, coords*)

**max\_ang\_moms** = {'mio-ext': {'C': 'p', 'H': 's', 'N': 'p', 'O': 'p'}}

**parse\_energy**(*path*)

**static parse\_exc\_dat**(*text*)

**parse\_forces**(*path*)

**parse\_total\_energy**(*text*)

**prepare\_input**(*atoms, coords, calc\_type*)

Meant to be extended.

**prepare\_overlap\_data**(*path*)

Implement calculator specific parsing of MO coefficients and CI coefficients here. Should return a filename pointing to TURBOMOLE like mos, a MO coefficient array and a CI coefficient array.

**run\_calculation**(*atoms, coords, \*\*prepare\_kwargs*)

pysisyphus.calculators.DFTBp.**parse\_mo**(*eigvec*)

pysisyphus.calculators.DFTBp.**parse\_xplusy**(*text*)

## 6.5 Calculators with Ground state capabilities

### 6.5.1 MOPAC 2016

```
class pysisiphus.calculators.MOPAC.MOPAC(method='PM7', **kwargs)
```

Bases: *Calculator*

<http://openmopac.net/manual/>

```

CALC_TYPES = {'energy': '1SCF', 'gradient': '1SCF GRADIENTS', 'hessian': 'DFORCE
FORCE LET'}
```

```

METHODS = ['am1', 'pm3', 'pm6', 'pm6-dh2', 'pm6-d3', 'pm6-dh+', 'pm6-dh2',
'pm6-dh2x', 'pm6-d3h4', 'pm6-d3h4x', 'pm7', 'pm7-ts']
```

```

MULT_STRS = {1: 'SINGLET', 2: 'DOUBLET', 3: 'TRIPLET', 4: 'QUARTET', 5:
'QUINTET', 6: 'SEXTET', 7: 'SEPTET', 8: 'OCTET'}
```

**base\_cmd**

Do only SCF AUX: Creates a checkpoint file NOREO: Dont reorient geometry

**Type**

1SCF

**conf\_key = 'mopac'**

**get\_energy**(atoms, coords)

Meant to be extended.

**get\_forces**(atoms, coords)

Meant to be extended.

**get\_hessian**(atoms, coords)

Meant to be extended.

**parse\_energy**(path)

**static parse\_energy\_from\_aux**(inp, \*args, \*\*kwargs)

**parse\_grad**(path)

**parse\_hessian**(path)

**static parse\_hessian\_from\_aux**(inp, \*args, \*\*kwargs)

**prepare\_coords**(atoms, coords, opt=False)

Get 3d coords in Angstrom.

Reshape internal 1d coords to 3d and convert to Angstrom.

**Parameters**

- **atoms** (*iterable*) -- Atom descriptors (element symbols).
- **coords** (*np.array, 1d*) -- 1D-array holding coordinates in Bohr.

**Returns**

**coords** -- 3D-array holding coordinates in Angstrom.

**Return type**

np.array, 3d

**prepare\_input**(*atoms, coords, calc\_type, opt=False*)

Meant to be extended.

**read\_aux**(*path*)

## 6.5.2 Psi4

```
class pysisyphus.calculators.Psi4.Psi4(method, basis, to_set=None, pcm='iefpcm', solvent=None,
write_fchk=False, **kwargs)
```

Bases: *Calculator***conf\_key** = 'psi4'**get\_energy**(*atoms, coords, \*\*prepare\_kwargs*)

Meant to be extended.

**get\_fchk\_str**()**get\_forces**(*atoms, coords, \*\*prepare\_kwargs*)

Meant to be extended.

**get\_hessian**(*atoms, coords, \*\*prepare\_kwargs*)

Meant to be extended.

**parse\_energy**(*path*)**parse\_grad**(*path*)**parse\_hessian**(*path*)**prepare\_input**(*atoms, coords, calc\_type*)

Meant to be extended.

**run\_calculation**(*atoms, coords, \*\*prepare\_kwargs*)

## 6.5.3 QCEngine

## 6.5.4 XTB 6.x

```
class pysisyphus.calculators.XTB.OptResult(opt_geom, opt_log)
```

Bases: tuple

**opt\_geom**

Alias for field number 0

**opt\_log**

Alias for field number 1

```
class pysisyphus.calculators.XTB.XTB(gbsa="", alpb="", gfn=2, acc=1.0, etemp=None, retry_etemp=None,
topo=None, topo_update=None, quiet=False, **kwargs)
```

Bases: *Calculator*

```
__init__(gbsa="", alpb="", gfn=2, acc=1.0, etemp=None, retry_etemp=None, topo=None,
         topo_update=None, quiet=False, **kwargs)
```

XTB calculator.

Wrapper for running energy, gradient and Hessian calculations by XTB.

#### Parameters

- **gbsa** (*str*, *optional*) -- Solvent for GBSA calculation, by default no solvent model is used.
- **alpb** (*str*, *optional*) -- Solvent for ALPB calculation, by default no solvent model is used.
- **gfn** (*int or str*, *must be (0, 1, 2, or "ff")*) -- Hamiltonian for the XTB calculation (GFN0, GFN1, GFN2, or GFNFF).
- **acc** (*float*, *optional*) -- Accuracy control of the calculation, the lower the tighter several numerical thresholds are chosen.
- **topo** (*str*, *optional*) -- Path the a GFNFF-topolgy file. As setting up the topology may take some time for sizable systems, it may be desired to reuse the file.
- **topo\_update** (*int*) -- Integer controlling the update interval of the GFNFF topology update. If supplied, the topolgy will be recreated every N-th calculation.
- **mem** (*int*) -- Mememory per core in MB.
- **quiet** (*bool*, *optional*) -- Suppress creation of log files.

```
static check_termination(inp, *args, **kwargs)
```

```
conf_key = 'xtb'
```

```
get_energy(atoms, coords, **prepare_kwargs)
```

Meant to be extended.

```
get_forces(atoms, coords, **prepare_kwargs)
```

Meant to be extended.

```
get_hessian(atoms, coords, **prepare_kwargs)
```

Meant to be extended.

```
get_mdrestart_str(coords, velocities)
```

coords and velocities have to given in au!

```
get_pal_env()
```

```
get_retry_args()
```

```
keep(path)
```

Backup calculation results.

#### Parameters

**path** (*Path*) -- Temporary directory of the calculation.

#### Returns

**kept\_fns** -- Dictionary holding the filenames that were backed up. The keys correspond to the type of file.

#### Return type

dict

**parse\_charges**(*fn=None*)

**parse\_charges\_from\_json**(*fn=None*)

**parse\_energy**(*path*)

**parse\_gradient**(*path*)

**parse\_hessian**(*path*)

**parse\_md**(*path*)

**parse\_opt**(*path, keep\_log=False*)

**parse\_topo**(*path*)

**prepare\_add\_args**(*xcontrol=None*)

**prepare\_coords**(*atoms, coords*)

Get 3d coords in Angstrom.

Reshape internal 1d coords to 3d and convert to Angstrom.

**Parameters**

- **atoms** (*iterable*) -- Atom descriptors (element symbols).
- **coords** (*np.array, 1d*) -- 1D-array holding coordinates in Bohr.

**Returns**

**coords** -- 3D-array holding coordinates in Angstrom.

**Return type**

*np.array, 3d*

**prepare\_input**(*atoms, coords, calc\_type, point\_charges=None*)

Meant to be extended.

**reattach**(*last\_calc\_cycle*)

**run\_calculation**(*atoms, coords, \*\*prepare\_kwargs*)

**run\_md**(*atoms, coords, t, dt, velocities=None, dump=1*)

Expecting t and dt in fs, even though xtb wants t in ps!

**run\_opt**(*atoms, coords, keep=True, keep\_log=False*)

**run\_topo**(*atoms, coords*)

**write\_mdrestart**(*path, mdrestart\_str*)

## 6.5.5 Dalton

```
class pysisyphus.calculators.Dalton.Dalton(basis, method='hf', **kwargs)
```

```
    Bases: Calculator
```

```
    compute(mol, prop)
```

```
    conf_key = 'dalton'
```

**get\_energy**(*atoms, coords, \*\*prepare\_kwargs*)

Meant to be extended.

**get\_forces**(*atoms, coords, \*\*prepare\_kwargs*)

Meant to be extended.

**prepare\_input**(*atoms, coords*)

Meant to be extended.

## 6.5.6 OpenBabel

```
class pysisyphus.calculators.OBabel.OBabel(ff='gaff', mol=None, **kwargs)
```

Bases: *Calculator*

```
conv_dict = {'kcal/mol': 627.5094740630558, 'kJ/mol': 2625.4996394798254}
```

**get\_energy**(*atoms, coords, \*\*prepare\_kwargs*)

Meant to be extended.

**get\_forces**(*atoms, coords, \*\*prepare\_kwargs*)

Meant to be extended.

**setup**(*atoms, coords*)

## 6.6 Meta (wrapping) Calculators

### 6.6.1 ExternalPotential

### 6.6.2 Restraint

```
# General input structure for restraints
calc:
  type: ext
  # Multiple potentials could be specified here as a list
  potentials:
    # Primitive internal coordinate restraint
    - type: restraint
      # List of restraints; could also be multiple restraints. Each restraint is given as
      # list of 2 or 3 items.
      #
      # The first item always specifies an internal coordinate,
      # whereas the second argument is a force constant (in atomic units; actual units
      # depend on the coordinate). Optionally a reference value (third argument) can be
      # given. If omitted, the initial coordinate value is used as reference value.
      restraints: [[[BOND, 0, 1], 10, 3.0]]
      # The commented out input below would restrain the bond at its initial value.
      #restraints: [[[BOND, 0, 1], 10]]
      # Multiple restraints are specified as given below.
      #restraints: [[[BOND, 0, 1], 10], [[BEND, 0, 1, 2], 1.0]]
calc:
  type: [actual calculator that is wrapped by ExternalPotential]
```

### 6.6.3 HarmonicSphere

### 6.6.4 LogFermi

### 6.6.5 AFIR

```
class pysisyphus.calculators.AFIR.AFIR(calculator, fragment_indices, gamma, rho=1, p=6,
                                       ignore_hydrogen=True, complete_fragments=True, dump=True,
                                       h5_fn='afir.h5', h5_group_name='afir', **kwargs)
```

Bases: *Calculator*

```
__init__(calculator, fragment_indices, gamma, rho=1, p=6, ignore_hydrogen=True,
         complete_fragments=True, dump=True, h5_fn='afir.h5', h5_group_name='afir', **kwargs)
```

Artificial Force Induced Reaction calculator.

Currently, there are no automated drivers to run large-scale AFIR calculations with many different initial orientations and/or increasing collision energy parameter. Nonetheless, selected AFIR calculations can be carried out by hand. After convergence, artificial potential & forces, as well as real energies and forces can be plotted with 'pysisplot --afir'. The highest energy point along the AFIR path can then be selected for a subsequent TS-optimization, e.g. via 'pysistrj --get [index] optimization.trj'.

Future versions of pysisyphus may provide drivers for more automated AFIR calculations.

#### Parameters

- **calculator** (*Calculator*) -- Actual QC calculator that provides energies and its derivatives, that are modified by the AFIR calculator, e.g., ORCA or Psi4.
- **fragment\_indices** (List[List[int]]) -- List of lists of integers, specifying the separate fragments. If the indices in these lists don't comprise all atoms in the molecule, the remaining indices will be added as a separate fragment. If a AFIR calculation is carried out with 2 fragments and 'complete\_fragments' is True (see below) it is enough to specify only the indices of one fragment, e.g., for a system of 10 atoms 'fragment\_indices=[[0,1,2,3]]' is enough. The second system will be set up automatically with indices [4,5,6,7,8,9].
- **gamma** (Union[\_SupportsArray[dtype], \_NestedSequence[\_SupportsArray[dtype]], bool, int, float, complex, str, bytes, \_NestedSequence[Union[bool, int, float, complex, str, bytes]]]) -- Collision energy parameter in au. For 2 fragments it can be a single integer, while for > 2 fragments a list of gammas must be given, specifying the pair-wise collision energy parameters. For 3 fragments 3 gammas must be given [\_01, \_02, \_12], for 4 fragments 6 gammas would be required [\_01, \_02, \_03, \_12, \_13, \_23] and so on.
- **rho** (Union[\_SupportsArray[dtype], \_NestedSequence[\_SupportsArray[dtype]], bool, int, float, complex, str, bytes, \_NestedSequence[Union[bool, int, float, complex, str, bytes]]], default: 1) -- Direction of the artificial force, either 1 or -1. The same comments as for gamma apply. For 2 fragments a single integer is enough, for > 2 fragments a list of rhos must be given (see above). For rho=1 fragments are pushed together, for rho=-1 fragments are pulled apart.
- **p** (int, default: 6) -- Exponent p used in the calculation of the weight function. Defaults to 6 and probably does not have to be changed.
- **ignore\_hydrogen** (bool, default: True) -- Whether hydrogens are ignored in the calculation of the artificial force.
- **complete\_fragments** (bool, default: True) -- Whether an incomplete specification in 'fragment\_indices' is automatically completed.

- **dump** (bool, default: True) -- Whether an HDF5 file is created.
- **h5\_fn** (str, default: 'afir.h5') -- Filename of the HDF5 file used for dumping.
- **h5\_group\_name** (str, default: 'afir') -- HDF5 group name used for dumping.
- **\*\*kwargs** -- Keyword arguments passed to the Calculator baseclass.

**afir\_fd\_hessian\_wrapper**(*coords3d, afir\_grad\_func*)

**property charge**

**dump\_h5**(*atoms, coords, results*)

**get\_energy**(*atoms, coords, \*\*prepare\_kwargs*)

Meant to be extended.

**get\_forces**(*atoms, coords, \*\*prepare\_kwargs*)

Meant to be extended.

**get\_hessian**(*atoms, coords, \*\*prepare\_kwargs*)

Meant to be extended.

**init\_h5\_group**(*atoms, max\_cycles=None*)

**log\_fragments**()

**property mult**

**set\_atoms\_and\_funcs**(*atoms, coords*)

Initially atoms was also an argument to the constructor of AFIR. I removed it so creation becomes easier. The first time a calculation is requested with a proper atom set everything is set up (cov. radii, afir function and corresponding gradient). Afterwards there is only a check if atoms != None and it is expected that all functions are properly set.

fragment\_indices can also be incomplete w.r.t. to the number of atoms. If the sum of the specified fragment atoms is less than the number of atoms present then all remaining unspecified atoms will be gathered in one fragment.

**write\_fragment\_geoms**(*atoms, coords*)

**class** pysisyphus.calculators.AFIR.**AFIRPath**(*atoms, cart\_coords, energies, forces, opt\_is\_converged*)

Bases: object

**atoms:** tuple

**cart\_coords:** ndarray

**energies:** ndarray

**forces:** ndarray

**opt\_is\_converged:** bool

pysisyphus.calculators.AFIR.**afir\_closure**(*fragment\_indices, cov\_radii, gamma, rho=1, p=6, prefactor=1.0, logger=None*)

rho=1 pushes fragments together, rho=-1 pulls fragments apart.

pysisyphus.calculators.AFIR.**get\_data\_model**(*atoms, max\_cycles*)

### 6.6.6 ONIOM

**class** pysisyphus.calculators.ONIOMv2.**LayerCalc**(*models, total\_size, parent\_layer\_calc=None*)

Bases: object

**property** charge

**do\_parent**(*with\_parent*)

**static energy\_from\_results**(*model\_energies, parent\_energy=None*)

**get\_energy**(*atoms, coords, with\_parent=True*)

**get\_forces**(*atoms, coords, with\_parent=True*)

**get\_hessian**(*atoms, coords, with\_parent=True*)

**property** mult

**run\_calculations**(*atoms, coords, method*)

**class** pysisyphus.calculators.ONIOMv2.**Link**(*ind, parent\_ind, atom, g*)

Bases: tuple

**atom**

Alias for field number 2

**g**

Alias for field number 3

**ind**

Alias for field number 0

**parent\_ind**

Alias for field number 1

**class** pysisyphus.calculators.ONIOMv2.**Model**(*name, calc\_level, calc, parent\_name, parent\_calc\_level, parent\_calc, atom\_inds, parent\_atom\_inds, use\_link\_atoms=True*)

Bases: object

**as\_calculator**(*cap=False*)

**as\_geom**(*all\_atoms, all\_coords*)

**capped\_atoms\_coords**(*all\_atoms, all\_coords*)

**create\_bond\_vec\_getters**(*atoms*)

**create\_links**(*atoms, coords, debug=False*)

**get\_energy**(*atoms, coords, point\_charges=None, parent\_correction=True, cap=True*)

**get\_forces**(*atoms, coords, point\_charges=None, parent\_correction=True, cap=True*)

**get\_hessian**(*atoms, coords, point\_charges=None, parent\_correction=True, cap=True*)

**get\_jacobian**()

`get_sparse_jacobian()`

`log(message="")`

`parse_charges()`

`class pysisiphus.calculators.ONIOMv2.ModelDummyCalc(model, cap=False)`

Bases: object

`get_energy(atoms, coords)`

`get_forces(atoms, coords)`

`class pysisiphus.calculators.ONIOMv2.ONIOM(calcs, models, geom, layers=None, embedding="", real_key='real', use_link_atoms=True, *args, **kwargs)`

Bases: [Calculator](#)

`__init__(calcs, models, geom, layers=None, embedding="", real_key='real', use_link_atoms=True, *args, **kwargs)`

**layer: list of models**

len(layer) == 1: normal ONIOM, len(layer) >= 1: multicenter ONIOM.

**model:**

(sub)set of all atoms that resides in a certain layer and has a certain calculator.

`atom_inds_in_layer(index, exclude_inner=False)`

Returns list of atom indices in layer at index.

Atoms that also appear in inner layer can be excluded on request.

**Parameters**

- **index** (*int*) -- pasd
- **exclude\_inner** (*bool*, *default=False*, *optional*) -- Whether to exclude atom indices that also appear in inner layers.

**Returns**

**atom\_indices** -- List containing the atom indices in the selected layer.

**Return type**

list

`calc_layer(atoms, coords, index, parent_correction=True)`

**property charge**

```
embeddings = {'': '', 'electronic': 'Electronic embedding', 'electronic_rc':
'Electronic embedding with redistributed charges', 'electronic_rcd': 'Electronic
embedding with redistributed charges and dipoles'}
```

`get_energy(atoms, coords)`

Meant to be extended.

`get_forces(atoms, coords)`

Meant to be extended.

`get_hessian(atoms, coords)`

Meant to be extended.

`get_layer_calc(layer_ind)`

`property model_iter`

`property mult`

`run_calculation(atoms, coords)`

`run_calculations(atoms, coords, method)`

`pysisyphus.calculators.ONIOMv2.atom_inds_to_cart_inds(atom_inds)`

`pysisyphus.calculators.ONIOMv2.cap_fragment(atoms, coords, fragment, link_atom='H', g=0.709)`

`pysisyphus.calculators.ONIOMv2.get_embedding_charges(embedding, layer, parent_layer, coords3d)`

`pysisyphus.calculators.ONIOMv2.get_g_value(atom, parent_atom, link_atom)`

### 6.6.7 Dimer

```
class pysisyphus.calculators.Dimer.Dimer(calculator, *args, N_raw=None, length=0.0189,  
rotation_max_cycles=15, rotation_method='fourier',  
rotation_thresh=0.0001, rotation_tol=1,  
rotation_max_element=0.001, rotation_interpolate=True,  
rotation_disable=False, rotation_disable_pos_curv=True,  
rotation_remove_trans=True, trans_force_f_perp=True,  
bonds=None, N_hessian=None, bias_rotation=False,  
bias_translation=False, bias_gaussian_dot=0.1, seed=None,  
write_orientations=True, forward_hessian=True, **kwargs)
```

Bases: [Calculator](#)

**property C**

Shortcut for the curvature.

**property N**

**add\_gaussian**(*atoms, center, N, height=0.1, std=0.0529, max\_cycles=50, dot\_ref=None*)

**property can\_bias\_f0**

**property can\_bias\_f1**

**property coords0**

**property coords1**

**curvature**(*f1, f2, N*)

Curvature of the mode represented by the dimer.

**direct\_rotation**(*optimizer, prev\_step*)

**do\_dimer\_rotations**(*rotation\_thresh=None*)

**property energy0**

**property f0**

**property** `f1`

**property** `f1_bias`

**property** `f2`

Never calculated explicitly, but estimated from `f0` and `f1`.

**fourier\_rotation**(*optimizer, prev\_step*)

**get\_N\_raw\_from\_hessian**(*h5\_fn, root=0*)

**get\_forces**(*atoms, coords*)

Meant to be extended.

**get\_gaussian\_energies**(*coords, sum\_=True*)

**get\_gaussian\_forces**(*coords, sum\_=True*)

**get\_hessian**(*atoms, coords*)

Meant to be extended.

**remove\_translation**(*displacement*)

**property** `rot_force`

**rotate\_coords1**(*rad, theta*)

Rotate dimer and produce new `coords1`.

**set\_N\_raw**(*coords*)

**property** `should_bias_f0`

May lead to calculation of `f0` and/or `f1` if present!

**property** `should_bias_f1`

May lead to calculation of `f0` and/or `f1` if present!

**update\_orientation**(*coords*)

**class** `pysisyphus.calculators.Dimer.Gaussian`(*height, center, std, N*)

Bases: `object`

**energy**(*R, height=None*)

**forces**(*R, height=None*)

**exception** `pysisyphus.calculators.Dimer.RotationConverged`

Bases: `Exception`

## 6.7 Pure Python calculators

### 6.7.1 Sympy 2D Potentials

**class** `pysisyphus.calculators.AnaPotBase.AnaPotBase`(*V\_str, scale=1.0, xlim=(- 1, 1), ylim=(- 1, 1), levels=None, use\_sympify=True, minima=None, saddles=None*)

Bases: `Calculator`

**anim\_coords**(*coords*, *interval=50*, *show=False*, *title\_func=None*)

**anim\_opt**(*opt*, *energy\_profile=False*, *colorbar=False*, *figsize=(8, 6)*, *show=False*)

**get\_energy**(*atoms*, *coords*)  
 Meant to be extended.

**get\_forces**(*atoms*, *coords*)  
 Meant to be extended.

**classmethod get\_geom**(*coords*, *atoms='X'*), *V\_str=None*, *calc\_kwargs=None*, *geom\_kwargs=None*)

**get\_geoms\_from\_stored\_coords**(*coords\_list*, *i=None*, *calc\_kwargs=None*, *geom\_kwargs=None*)

**get\_hessian**(*atoms*, *coords*)  
 Meant to be extended.

**get\_minima**(*i=None*, *calc\_kwargs=None*, *geom\_kwargs=None*)

**get\_path**(*num*, *minima\_inds=None*)

**get\_saddles**(*i=None*, *calc\_kwargs=None*, *geom\_kwargs=None*)

**plot**(*levels=None*, *show=False*, *\*\*figkwargs*)

**plot3d**(*levels=None*, *show=False*, *zlim=None*, *vmin=None*, *vmax=None*, *resolution=100*, *rcount=50*, *ccount=50*, *nan\_above=None*, *init\_view=None*, *colorbar=False*, *\*\*figkwargs*)

**plot\_coords**(*xs*, *ys*, *enum=True*, *show=False*, *title=None*)

**plot\_eigenvalue\_structure**(*grid=50*, *levels=None*, *show=False*)

**plot\_geoms**(*geoms*, *\*\*kwargs*)

**plot\_irc**(*irc*, *\*args*, *\*\*kwargs*)

**plot\_opt**(*opt*, *\*args*, *\*\*kwargs*)

**statistics**()

## 6.7.2 Lennard-Jones

**class** pysisyphus.calculators.LennardJones.**LennardJones**(*sigma=1.8897261251*, *epsilon=1*, *rc=None*)  
 Bases: [Calculator](#)

**calculate**(*coords3d*)

**get\_energy**(*atoms*, *coords*)  
 Meant to be extended.

**get\_forces**(*atoms*, *coords*)  
 Meant to be extended.

### 6.7.3 FakeASE

```
class pysisyphus.calculators.FakeASE.FakeASE(calc)
    Bases: object
    get_atoms_coords(atoms)
    get_forces(atoms=None)
    get_potential_energy(atoms=None)
```

### 6.7.4 TIP3P

```
class pysisyphus.calculators.TIP3P.TIP3P(rc=9.44863062728914)
    Bases: Calculator
    Transferable Intermolecular Potential 3 Point
    aHOH = 104.52
    calculate(coords3d)
    charges
        coulomb_energy = (multiple of elem. charge * multiple of elem. charge)
            / (distance in bohr) * 1 / (4 * pi * vacuum permittivity)
        coulomb_prefactor converts everything to atomic units and it is ... drum roll ... 1. from scipy.constants
        import value as pcv self.coulomb_prefactor = (1 / (4 * np.pi) * pcv("elementary charge"))**2
            / pcv("Hartree energy") / pcv("Bohr radius") / pcv("vacuum electric permittivity")
    )
    coulomb(coords3d)
    epsilon = 0.0002423919586315716
    get_energy(atoms, coords)
        Meant to be extended.
    get_forces(atoms, coords)
        Meant to be extended.
    qH = 0.417
    qO = -0.834
    rOH = 1.8088458464917874
    sigma = 5.953790025507198
```



## MINIMIZATION

Searching for minimum energy geometries of molecules is preferably done using second order methods that employ hessian information. Using the plain hessian  $H$  for step prediction through  $p = -H^{-1}g$ , with  $g$  being the gradient vector, may yield erroneous uphill steps when  $H$  has negative eigenvalues.

This can be understood by calculating the step in the basis of the hessian eigenvectors  $V$ .

$$\begin{aligned} \tilde{H} &= V^T H V & (7.1) \\ \tilde{g} &= V^T g \\ \tilde{p}_i &= -\frac{\tilde{g}_i}{\tilde{H}_{ii}} \end{aligned} \tag{7.4}$$

$\tilde{H}$ ,  $\tilde{g}$  and  $\tilde{p}$  are transformed into the eigenvector basis and the subscript  $i$  indicates the component belongs to the  $i$ -th eigenvalue (-vector) of  $H$ . As the gradient always points into the direction of greater function values dividing it by a negative eigenvalues  $\tilde{H}_{ii}$  will lead to a step in uphill direction along the  $i$ -th eigenvector. The step in the original basis is obtained by a simple back-transformation:

$$p = V \tilde{p}$$

A step in downhill direction can be ensured by introducing an appropriate shift parameter  $\lambda$  that must be smaller than the smallest eigenvalue of  $H$ :

$$\tilde{p}_i = -\frac{\tilde{g}_i}{\tilde{H}_{ii} - \lambda}$$

In *pysisyphus* the shift parameter  $\lambda$  is obtained by the Rational Function Optimization approach.

When the root-mean-square of the predicted step  $p$  drops below a certain threshold (default is 0.0025 au or rad) controlled GDIIS is tried. If GDIIS fails or is not yet possible a polynomial line-search is conducted. Using the latest two energies and the projection of the latest two gradients onto the last step the coefficients of a cubic polynomial are defined unambiguously. By requiring  $d^2 f(x)/dx^2 \geq 0$  and the equality holding at exactly one point also a quartic polynomial can be determined.

For now line-searches using a cubic polynomial are disabled as they seem to degrade the optimizer performance, so only the constrained quartic polynomial is tried. By also using projected hessian information the coefficients of a quintic polynomial can be determined, but this also seems to be inferior compared to the constrained quartic polynomial.

Convergence is indicated when the root-mean-square and the maximum value of the gradient and step drop below a certain threshold. It is not uncommon that the gradient convergence is achieved before step convergence. By using *overachieve\_factor*:  $[n, \text{float} > 1]$  in the YAML input convergence will be signalled, when gradient convergence is overachieved by factor  $n$ .

## 7.1 YAML Example

Below you can find an example YAML-input including the most important options that the user may want to modify when using the RFOptimizer.

```

opt:
  type: rfo                                # Optimization algorithm
  max_cycles: 50                           # Maximum number of optimization cycles

  overachieve_factor: 2                    # Indicate convergence, regardless of the
                                           # proposed step when max(grad) and rms(grad)
                                           # are overachieved by factor [n]

  do_hess: True                             # Calculate the hessian at the final geometry
                                           # after the optimization.

  #hessian_recalc: None                     # Recalculate exact hessian every n-th cycle

  #hessian_recalc_adapt: None               # Expects a float. Recalculate exact hessian
                                           # whenever the gradient norms drops below
                                           # 1/[n] of the gradient norm at the last hessian
                                           # recalculation.

  #hessian_init: fischer                    # Type of model hessian. Other options are: 'calc,
                                           # simple, xtb, lindh, swart, unit'

  #hessian_update: bfgs                     # Hessian-update. Other options are: 'flowchart,
                                           # damped_bfgs, bofill'. bofill is not recommended
                                           # for minimum searches.

  #small_eigval_thresh: 1e-8               # Neglect eigenvalues and corresponding eigenvectors
                                           # below this threshold.

  #max_micro_cycles: 50                     # No. of micro cycles for the RS-variants. Does not apply
                                           # to TRIM.

  #trust_radius: 0.3                        # Initial trust radius.
  #trust_max: 1.0                           # Max. trust radius
  #trust_min: 0.1                           # Min. trust radius

  #line_search: True                       # Do line search

  #gdiis_thresh: 0.0025                     # May do GDIIS if rms(step) falls below this threshold
  #gediis_thresh: 0.01                      # May do GEDIIS if rms(grad) falls below this threshold
  #gdiis: True                               # Do controlled GDIIS after 'gdiis_thresh' is reached
  #gediis: False                             # Do GEDIIS after 'gediis_thresh' is reached

calc:
  type: turbomole
  control_path: control_path_pbe0_def2svp_s1 # Path to the prepared calculation
  track: True                                # Activate excited state tracking
  ovlp_type: tden                            # Track with transition density matrix.
  ↪ overlaps
  charge: 0

```

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```

mult: 1
pal: 4
mem: 2000

geom:
  type: redund
  fn: cytosin.xyz

```

Further examples for optimizations from `.yaml` input can be found [here](#).

## 7.2 Convergence Thresholds

Optimization convergence is indicated when four criteria are met. Each cycle the root-mean-square and the absolute maximum of force and proposed step vectors are checked. The default criteria in pysisyphus correspond to the `opt=loose` in Gaussian. Different thresholds can easily be requested by supplying the respective keyword in the YAML input.

Convergence is also indicated, when `max(force)` and `rms(force)` are overachieved, by a certain factor (`overachieve_factor`, default=3), similar to ORCA's optimizer. Further keywords, controlling the convergence check are found below.

```

opt:
  ...
  thresh: gau_loose           # See table below for different thresholds.
  overachieve_factor: 3
  rms_force: null           # Derive four criteria from this value
  rms_force_only: False     # Only check rms(force)
  max_force_only: False     # Only check max(force)
  # Chain-of-States specific
  coord_diff_thresh: 0.01   # Terminate on insignificant coordinate change
  reparam_thresh: 0.001     # Terminate on insignificant coordinate change upon
  ↔reparametrization
  ...

```

Thresh-old	max(force)	rms(forces)	max(step)	rms(step)	Comment
<code>gau_loose</code>	2.5e-3	1.7e-3	1.0e-2	6.7e-3	default
<code>gau</code>	4.5e-4	3.0e-4	1.8e-3	1.2e-3	
<code>gau_tight</code>	1.5e-5	1.0e-5	6.0e-5	4.0e-5	
<code>gau_vtight</code>	2.0e-6	1.0e-6	6.0e-6	4.0e-6	
<code>baker</code>	3.0e-4	2.0e-4	3.0e-4	2.0e-4	energy difference and step are also checked
<code>never</code>	2.0e-6	1.0e-6	6.0e-6	4.0e-6	dummy thresholds; never converges

## 7.3 Available Optimizers

Base class for optimizers that utilize Hessian information.

```
class pysisyphus.optimizers.Optimizer.Optimizer(geometry, thresh='gau_loose', max_step=0.04,
                                               max_cycles=100, min_step_norm=1e-08,
                                               assert_min_step=True, rms_force=None,
                                               rms_force_only=False, max_force_only=False,
                                               converge_to_geom_rms_thresh=0.05, align=False,
                                               dump=False, dump_restart=False, prefix='',
                                               reparam_thresh=0.001, reparam_check_rms=True,
                                               reparam_when='after', overachieve_factor=0.0,
                                               restart_info=None, check_coord_diffs=True,
                                               coord_diff_thresh=0.01, fragments=None,
                                               monitor_frag_dists=0, out_dir='',
                                               h5_fn='optimization.h5', h5_group_name='opt')
```

Bases: object

```
__init__(geometry, thresh='gau_loose', max_step=0.04, max_cycles=100, min_step_norm=1e-08,
         assert_min_step=True, rms_force=None, rms_force_only=False, max_force_only=False,
         converge_to_geom_rms_thresh=0.05, align=False, dump=False, dump_restart=False, prefix='',
         reparam_thresh=0.001, reparam_check_rms=True, reparam_when='after',
         overachieve_factor=0.0, restart_info=None, check_coord_diffs=True, coord_diff_thresh=0.01,
         fragments=None, monitor_frag_dists=0, out_dir='', h5_fn='optimization.h5',
         h5_group_name='opt')
```

Optimizer baseclass. Meant to be subclassed.

### Parameters

- **geometry** (*Geometry*) -- Geometry to be optimized.
- **thresh** (Literal['gau\_loose', 'gau', 'gau\_tight', 'gau\_vtight', 'baker', 'never'], default: 'gau\_loose') -- Convergence threshold.
- **max\_step** (float, default: 0.04) -- Maximum absolute component of the allowed step vector. Utilized in optimizers that don't support a trust region or line search.
- **max\_cycles** (int, default: 100) -- Maximum number of allowed optimization cycles.
- **min\_step\_norm** (float, default: 1e-08) -- Minimum norm of an allowed step. If the step norm drops below this value a ZeroStepLength-exception is raised. The unit depends on the coordinate system of the supplied geometry.
- **assert\_min\_step** (bool, default: True) -- Flag that controls whether the norm of the proposed step is check for being too small.
- **rms\_force** (Optional[float], default: None) -- Root-mean-square of the force from which user-defined thresholds are derived. When 'rms\_force' is given 'thresh' is ignored.
- **rms\_force\_only** (bool, default: False) -- When set, convergence is signalled only based on rms(forces).
- **max\_force\_only** (bool, default: False) -- When set, convergence is signalled only based on max(**forces**).
- **converge\_to\_geom\_rms\_thresh** (float, default: 0.05) -- Threshold for the RMSD with another geometry. When the RMSD drops below this threshold convergence is signalled. Only used with Growing Newton trajectories.

- **align** (bool, default: False) -- Flag that controls whether the geometry is aligned in every step onto the coordinates of the previous step. Must not be used with internal coordinates.
- **dump** (bool, default: False) -- Flag to control dumping/writing of optimization progress to the filesystem
- **dump\_restart** (bool, default: False) -- Flag to control whether restart information is dumped to the filesystem.
- **prefix** (str, default: ' ') -- Short string that is prepended to several files created by the optimizer. Allows distinguishing several optimizations carried out in the same directory.
- **reparam\_thresh** (float, default: 0.001) -- Controls the minimal allowed similarity between coordinates after two successive reparametrizations. Convergence is signalled if the coordinates did not change significantly.
- **reparam\_check\_rms** (bool, default: True) -- Whether to check for (too) similar coordinates after reparametrization.
- **reparam\_when** (Optional[Literal['before', 'after']], default: 'after') -- Reparametrize before or after calculating the step. Can also be turned off by setting it to None.
- **overachieve\_factor** (float, default: 0.0) -- Signal convergence when max(forces) and rms(forces) fall below the chosen threshold, divided by this factor. Convergence of max(step) and rms(step) is ignored.
- **restart\_info** -- Restart information. Undocumented.
- **check\_coord\_diffs** (bool, default: True) -- Whether coordinates of chain-of-sates images are checked for being too similar.
- **coord\_diff\_thresh** (float, default: 0.01) -- Unitless threshold for similiary checking of COS image coordinates. The first image is assigned 0, the last image is assigned to 1.
- **fragments** (Optional[Tuple], default: None) -- Tuple of lists containing atom indices, defining two fragments.
- **monitor\_frag\_dists** (int, default: 0) -- Monitor fragment distances for N cycles. The optimization is terminated when the interfragment distances falls below the initial value after N cycles.
- **out\_dir** (str, default: '. ') -- String poiting to a directory where optimization progress is dumped.
- **h5\_fn** (str, default: 'optimization.h5') -- Basename of the HDF5 file used for dumping.
- **h5\_group\_name** (str, default: 'opt') -- Groupname used for dumping of this optimization.

**check\_convergence**(*step=None, multiple=1.0, overachieve\_factor=None, energy\_thresh=1e-06*)

Check if the current convergence of the optimization is equal to or below the required thresholds, or a multiple thereof. The latter may be used in initiating the climbing image.

**dump\_restart\_info**()

**final\_summary**()

**get\_path\_for\_fn**(*fn, with\_prefix=True*)

**get\_restart\_info**()

```
log(message, level=50)
make_conv_dict(key, rms_force=None)
abstract optimize()
postprocess_opt()
prepare_opt()
print_opt_progress()
report_conv_thresholds()
run()
scale_by_max_step(steps)
set_restart_info(restart_info)
write_cycle_to_file()
write_image_trjs()
write_results()
write_to_out_dir(out_fn, content, mode='w')
```

```
pysisyphus.optimizers.Optimizer.get_data_model(geometry, is_cos, max_cycles)
```

```
class pysisyphus.optimizers.HessianOptimizer.HessianOptimizer(geometry, trust_radius=0.5,
                                                             trust_update=True, trust_min=0.1,
                                                             trust_max=1,
                                                             max_energy_incr=None,
                                                             hessian_update='bfgs',
                                                             hessian_init='fischer',
                                                             hessian_recalc=None,
                                                             hessian_recalc_adapt=None,
                                                             hessian_xtb=False,
                                                             hessian_recalc_reset=False,
                                                             small_eigval_thresh=1e-08,
                                                             line_search=False, alpha0=1.0,
                                                             max_micro_cycles=25,
                                                             rfo_overlaps=False, **kwargs)
```

Bases: [Optimizer](#)

```
__init__(geometry, trust_radius=0.5, trust_update=True, trust_min=0.1, trust_max=1,
         max_energy_incr=None, hessian_update='bfgs', hessian_init='fischer', hessian_recalc=None,
         hessian_recalc_adapt=None, hessian_xtb=False, hessian_recalc_reset=False,
         small_eigval_thresh=1e-08, line_search=False, alpha0=1.0, max_micro_cycles=25,
         rfo_overlaps=False, **kwargs)
```

Baseclass for optimizers utilizing Hessian information.

#### Parameters

- **geometry** ([Geometry](#)) -- Geometry to be optimized.
- **trust\_radius** (float, default: 0.5) -- Initial trust radius in whatever unit the optimization is carried out.

- **trust\_update** (bool, default: True) -- Whether to update the trust radius throughout the optimization.
- **trust\_min** (float, default: 0.1) -- Minimum trust radius.
- **trust\_max** (float, default: 1) -- Maximum trust radius.
- **max\_energy\_incr** (Optional[float], default: None) -- Maximum allowed energy increased after a faulty step. Optimization is aborted when the threshold is exceeded.
- **hessian\_update** (Literal['none', None, False, 'bfgs', 'damped\_bfgs', 'flowchart', 'bofill', 'ts\_bfgs', 'ts\_bfgs\_org', 'ts\_bfgs\_rev'], default: 'bfgs') -- Type of Hessian update. Defaults to BFGS for minimizations and Bofill for saddle point searches.
- **hessian\_init** (Literal['calc', 'unit', 'fischer', 'lindh', 'simple', 'swart', 'xtb', 'xtb1', 'xtbff'], default: 'fischer') -- Type of initial model Hessian.
- **hessian\_recalc** (Optional[int], default: None) -- Recalculate exact Hessian every n-th cycle instead of updating it.
- **hessian\_recalc\_adapt** (Optional[float], default: None) -- Use a more flexible scheme to determine Hessian recalculation. Undocumented.
- **hessian\_xtb** (bool, default: False) -- Recalculate the Hessian at the GFN2-XTB level of theory.
- **hessian\_recalc\_reset** (bool, default: False) -- Whether to skip Hessian recalculation after reset. Undocumented.
- **small\_eigval\_thresh** (float, default: 1e-08) -- Threshold for small eigenvalues. Eigenvectors belonging to eigenvalues below this threshold are discarded.
- **line\_search** (bool, default: False) -- Whether to carry out a line search. Not implemented by a subclassing optimizers.
- **alpha0** (float, default: 1.0) -- Initial alpha for restricted-step (RS) procedure.
- **max\_micro\_cycles** (int, default: 25) -- Maximum number of RS iterations.
- **rfo\_overlaps** (bool, default: False) -- Enable mode-following in RS procedure.
- **\*\*kwargs** -- Keyword arguments passed to the Optimizer baseclass.

**filter\_small\_eigvals**(*eigvals*, *eigvecs*, *mask=False*)

**get\_alpha\_step**(*cur\_alpha*, *rfo\_eigval*, *step\_norm*, *eigvals*, *gradient*)

**get\_augmented\_hessian**(*eigvals*, *gradient*, *alpha=1.0*)

**static get\_newton\_step**(*eigvals*, *eigvecs*, *gradient*)

**get\_newton\_step\_on\_trust**(*eigvals*, *eigvecs*, *gradient*, *transform=True*)

Step on trust-radius.

See Nocedal 4.3 Iterative solutions of the subproblem

**get\_rs\_step**(*eigvals*, *eigvecs*, *gradient*, *name='RS'*)

**static get\_shifted\_step\_trans**(*eigvals*, *gradient\_trans*, *shift*)

**get\_step\_func**(*eigvals*, *gradient*, *grad\_rms\_thresh=0.01*)

**housekeeping()**

Calculate gradient and energy. Update trust radius and hessian if needed. Return energy, gradient and hessian for the current cycle.

**log\_negative\_eigenvalues**(*eigvals*, *pre\_str=""*)

**prepare\_opt**(*hessian\_init=None*)

**property prev\_eigvec\_max**

**property prev\_eigvec\_min**

**static quadratic\_model**(*gradient*, *hessian*, *step*)

**reset()**

**rfo\_dict** = {'max': (-1, 'max'), 'min': (0, 'min')}

**static rfo\_model**(*gradient*, *hessian*, *step*)

**save\_hessian()**

**set\_new\_trust\_radius**(*coeff*, *last\_step\_norm*)

**solve\_rfo**(*rfo\_mat*, *kind='min'*, *prev\_eigvec=None*)

**update\_hessian()**

**update\_trust\_radius()**

`pysisyphus.optimizers.HessianOptimizer.dummy_hessian_update`(*H*, *dx*, *dg*)

**class** `pysisyphus.optimizers.RFOptimizer.RFOptimizer`(*geometry*, *line\_search=True*, *gediis=False*,  
*gdiis=True*, *gdiis\_thresh=0.0025*,  
*gediis\_thresh=0.01*, *gdiis\_test\_direction=True*,  
*max\_micro\_cycles=0*, *adapt\_step\_func=False*,  
*\*\*kwargs*)

Bases: `HessianOptimizer`

**\_\_init\_\_**(*geometry*, *line\_search=True*, *gediis=False*, *gdiis=True*, *gdiis\_thresh=0.0025*, *gediis\_thresh=0.01*,  
*gdiis\_test\_direction=True*, *max\_micro\_cycles=0*, *adapt\_step\_func=False*, *\*\*kwargs*)

Rational function Optimizer.

**Parameters**

- **geometry** (*Geometry*) -- Geometry to be optimized.
- **line\_search** (bool, default: True) -- Whether to carry out implicit line searches.
- **gediis** (bool, default: False) -- Whether to enable GEDIIS.
- **gdiis** (bool, default: True) -- Whether to enable GDIIS.
- **gdiis\_thresh** (float, default: 0.0025) -- Threshold for rms(forces) to enable GDIIS.
- **gediis\_thresh** (float, default: 0.01) -- Threshold for rms(step) to enable GEDIIS.
- **gdiis\_test\_direction** (bool, default: True) -- Whether to the overlap of the RFO step and the GDIIS step.
- **max\_micro\_cycles** (int, default: 0) -- Number of restricted-step microcycles. Disabled by default.

- **adapt\_step\_func** (bool, default: False) -- Whether to switch between shifted Newton and RFO-steps.
- **\*\*kwargs** -- Keyword arguments passed to the Optimizer/HessianOptimizer baseclass.

**optimize()**

**postprocess\_opt()**



## MINIMIZATION WITH MICROITERATIONS

Pysisyphus allows layered optimizations using microiterations via the *LayerOpt* class. Required energies and its derivatives (gradients, Hessians) can either be calculated with pysisyphus' own ONIOM implementation or supplied through Unix sockets via the i-PI protocol.

In ONIOM calculations a small (model) system is treated using a high level of theory, while its surroundings (real system) are described at a lower, more economic level of theory. There may also be several (optional) intermediate layers. Please see the figure below.

Fig. 8.1: General structure of a n-layer ONIOM calculation in pysisyphus. Every box corresponds to a layer. Indices of the respective layers are given as red numbers in the top right corner of each layer.

Searching stationary points of such systems using classical optimization approaches with simultaneous relaxation of model and real system can be computationally quite costly. In cases where the model system converges before the real system, many unnecessary model system energy and gradients evaluations may be required to fully relax the total system, as one step requires gradient evaluations in all layers. Also, it may be prohibitively expensive to optimize the whole system using internal coordinates, so it would be desirable to treat only the model system using internal coordinates.

Optimization with microiterations offers a way to potentially reduce the number of costly calculations in the innermost layer. By fully relaxing the outer layers before taking a step in the innermost layer, superfluous gradient evaluations can be avoided. As the overall calculation time of an ONIOM energy/gradient is usually dominated by the innermost layer, great computational saving can be realized by decreasing the required number of such calculations.

In the present approach, outer layers are optimized in Cartesian coordinates with an economic optimizer like (pre-conditioned) limited memory BFGS (L-BFGS) or conjugate gradient (CG). As L-BFGS avoids operations with steep scaling costs like matrix diagonalization, relaxation of large system comprising thousands of atoms becomes possible with negligible computational overhead. In contrast to the outer layers, the inner layer is usually optimized using internal coordinates and an optimizer that utilizes an explicit Hessian matrix, as this often allows for the most efficient optimizations.

As energy and gradient evaluations in the outer layers should be cheap the inferior performance of optimizations in Cartesian coordinates should not lead to an overall runtime increase.

## 8.1 YAML input

When pysisyphus' native ONIOM calculator is to be used, appropriate input has to be given in the `calc:` section, that is, layer composition and the respective calculators. If energies & gradients are sent via sockets and the i-PI-protocol `calc:` can be left empty but the appropriate socket addresses have to be specified later in the `opt:` section (see below). A separate socket/address is required for each layer.

Independent of the actual layer, pysisyphus always expects vectorial quantities with appropriate shapes of the total system. The expected size of a force vector for a system comprising  $N$  atoms is  $3N$ . Vector components not belonging to frozen atoms not belonging to a given layer are ignored and may be zero. **It is expected, that the full valid ONIOM forces for the total system are sent with the innermost layer.** These forces are then also used to check for optimization convergence.

A layered optimization is requested via `type: layer` in the `opt:` section. If no further input is given, appropriate defaults will be used. Microiterations for the outer layers  $0$  to  $n-2$  will be carried out using Cartesian coordinates and regularized L-BFGS, while the innermost layer is optimized in internal coordinates (`type: tric`) using rational function optimization (`type: rfo`).

If a more fine grained control is desired each layer can be controlled independently by further keywords in the `layers:` list. When `layers:` is present in the input all layers must be specified, but empty list entries are possible (see the lines only containing a comment in the example below). **One must start with the outmost layer;** the last list item corresponds to the innermost layer. Geometry and/or optimizer setup of each layer is controlled by the `geom:` and `opt:` sections in the list of layers with the usual keywords.

The usual keywords to control the optimization are supported, e.g., `max_cycles`, `thresh` etc.

```
geom:
  type:
    cartesian
  fn: [filename of initial geometry]
calc:
  type:
    oniom:
      ...
opt:
  type:
    layer:
      [layers:
        - # dummy for layer 0 (real system)
        - # dummy for layer 1 layer
        - # ...
        # - # dummy for layer n-1 (model system)
        - geom:
            type:
              cartesian:
            opt:
              type:
                lbfgs:
          ]
      # thresh: gau_loose
      # max_cycles: 150
```

Below you can find a full example for the ONIOM2-optimization of Hexaphenylethane using pysisyphus' ONIOM implementation.

## 8.2 Example using pysisyphus' ONIOM

```
# 2-layer ONIOM optimization of Hexaphenylethan. The model
# system comprises the two carbons of the central ethan moiety.
# The model system is described GFN2-XTB while the real system is
# treated using GFN-FF.
```

```
geom:
```

```
  type: cart
```

```
  fn: |
```

```
    68
```

C	1.2639	0.6380	0.4785
C	2.1638	1.9009	0.1349
C	2.4167	2.3403	-1.2006
C	3.2096	3.4576	-1.4755
C	2.7467	2.7036	1.1595
C	3.5368	3.8191	0.8739
C	3.7741	4.1922	-0.4416
H	2.0100	1.8572	-2.0700
H	3.3815	3.7580	-2.5014
H	2.5944	2.5089	2.1979
H	3.9618	4.4009	1.6820
H	4.3848	5.0587	-0.6598
C	0.7891	0.0052	-0.8980
C	-1.0360	-0.5225	-2.5243
C	-0.0432	-1.1375	-3.4197
C	1.2448	-1.1452	-3.0690
C	-0.6577	0.0144	-1.3528
H	-1.4492	0.4671	-0.7765
H	-2.0793	-0.4955	-2.8238
H	-0.3570	-1.5703	-4.3642
C	1.6732	-0.5526	-1.7810
H	1.9838	-1.5872	-3.7297
H	2.7287	-0.5813	-1.6050
C	-0.0387	1.1726	1.2081
C	-0.2929	2.5599	1.4341
C	-1.0635	0.2863	1.6524
C	-2.2174	0.7426	2.2933
C	-2.4082	2.0998	2.5118
C	-1.4505	3.0060	2.0772
H	0.3724	3.3423	1.1178
H	-1.0153	-0.7691	1.5014
H	-2.9711	0.0356	2.6161
H	-1.6065	4.0659	2.2335
H	-3.3046	2.4506	3.0066
C	2.1016	-0.4883	1.4312
C	3.4003	-1.0269	0.6933
C	4.4239	-0.1425	0.2420
C	5.5730	-0.6012	-0.4056
C	3.6509	-2.4147	0.4669
C	4.8038	-2.8631	-0.1829
C	5.7602	-1.9589	-0.6243

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```

H      4.3778    0.9134    0.3928
H      6.3259    0.1044   -0.7337
H      2.9864   -3.1958    0.7882
H      4.9572   -3.9235   -0.3393
H      6.6530   -2.3115   -1.1243
C      2.5704    0.1468    2.8077
C      4.3159    0.7126    4.4630
C      3.3714    1.2844    5.3047
C      2.0334    1.2840    4.9354
C      3.9303    0.1495    3.2436
H      4.7333   -0.2864    2.6778
H      5.3575    0.6988    4.7582
H      3.6749    1.7164    6.2495
C      1.6376    0.7248    3.7185
H      1.2932    1.7141    5.5981
H      0.5868    0.7426    3.5290
C      1.2016   -1.7498    1.7785
C      0.9537   -2.1874    3.1154
C      0.6140   -2.5534    0.7571
C     -0.1758   -3.6680    1.0474
C     -0.4081   -4.0392    2.3643
C      0.1611   -3.3038    3.3950
H      1.3648   -1.7032    3.9822
H      0.7614   -2.3600   -0.2822
H     -0.6047   -4.2504    0.2418
H     -0.0069   -3.6027    4.4220
H     -1.0186   -4.9050    2.5861
calc:
type: oniom
calcs:
  real:
    type: xtb
    gfn: ff
    pal: 6
  high:
    type: xtb
    pal: 6
models:
  # The "real" does not have to be described, only the "real" system.
  high:
    # Use ethan-carbons for the model system
    inds: [0,34]
    # Use appropriate calculator for model system.
    calc: high
opt:
  # Use LayerOpt
  type: layers
  thresh: gau
  # When the 'layer' key is present, the user has to provide one
  # entry per layer in the optimization. When the layer is empty default
  # values are used. The layers are given from the outmost to the innermost
  # layer.

```

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```

# In the example above no other arguments for the outmost layer (# real dummy),
# while redundant internal coordinates are used for the innermost layer.
#
# When pysisyphus own ONIOM implementation is used no additional atom indices
# must be given here.
layers:
- # real dummy
- geom:
  type: redund
do_hess: True

```

### 8.3 Example with sockets & i-PI-protocol

Whereas pysisyphus can figure out the layer composition for the microcycles when its own ONIOM calculator is used, the user has to specify it when using sockets and the i-PI-protocol. When covalent bonds between layers exist link atoms (LAs) must be introduced. Given a layer with index  $n$ , atoms in layer  $n-1$  that are replaced by LAs in the given layer are called link atom hosts (LAH). A given layer and its LAHs in **higher** layers must be optimized together.

Fig. 8.2: Layer structure of a ONIOM2 optimization using microiterations with link atoms. A given layer as well as connected link atom hosts in **higher** layers must be optimized simultaneously.

A simple example for a ONIOM2 optimization with microiterations is found below. Here, ethanal is optimized, with the model system comprising the carbonyl group (atoms 4, 5 and 6). This introduces a link atom between the two carbons 4 and 0, with carbon 0 becoming a LAH.

Fig. 8.3: ONIOM2 optimization of ethanal. All atoms surrounded by the dashed blue line are optimized in the innermost layer. Hydrogens are shown white, carbons grey and oxygen red.

In contrast to the first example using the native ONIOM implementation the user can omit any input in the `calc:` section. Now the socket addresses have to be given for every layer, starting with the total system. For the total system the atom indices can be omitted, as it is assumed that it comprises all atoms. For the remaining layers, the indices of all moving atoms including LAHs in higher layers have to be given.

```

geom:
  type: cartesian
  fn: lib:acetaldehyd_oniom.xyz
calc:
opt:
  type: layers
  layers:
    # Atom indices of the total system / (first layer) don't have to be
    # specified.
    - address: ./layer0sock
    # Optimize atoms of the model system [4, 5, 6] and the link atom host 0.
    - indices: [0,4,5,6]
      address: ./layer1sock
  thresh: gau
# The assert: section can be left out in actual inputs,

```

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```
# it is here for testing purposes.
assert:
    opt_geom.energy: -115.53522653
```

Another sensible choice for optimizing outer layers besides (regularized) L-BFGS may be preconditioned L-BFGS (type: *plbfgs*).

```
class pysisyphus.optimizers.LayerOpt.LayerOpt(geometry, layers=None, **kwargs)
```

Bases: *Optimizer*

```
property layer_num: int
```

Return type  
int

```
optimize()
```

Return type  
None

```
postprocess_opt()
```

Return type  
None

```
class pysisyphus.optimizers.LayerOpt.Layers(geometry, opt_thresh, layers=None)
```

Bases: object

```
classmethod from_oniom_calculator(geometry, oniom_calc=None, layers=None, **kwargs)
```

```
pysisyphus.optimizers.LayerOpt.get_geom_kwargs(layer_ind, layer_mask)
```

```
pysisyphus.optimizers.LayerOpt.get_opt_kwargs(opt_key, layer_ind, thresh)
```

```
class pysisyphus.optimizers.LBFGS.LBFGS(geometry, keep_last=7, beta=1, max_step=0.2,
                                         double_damp=True, gamma_mult=False, line_search=False,
                                         mu_reg=None, max_mu_reg_adaptions=10, control_step=True,
                                         **kwargs)
```

Bases: *Optimizer*

```
__init__(geometry, keep_last=7, beta=1, max_step=0.2, double_damp=True, gamma_mult=False,
         line_search=False, mu_reg=None, max_mu_reg_adaptions=10, control_step=True, **kwargs)
```

Limited-memory BFGS optimizer.

See [1] Nocedal, Wright - Numerical Optimization, 2006 for a general discussion of LBFGS. See `pysisyphus.optimizers.hessian_updates` for the references related to double damping and `pysisyphus.optimizers.closures` for references related to regularized LBFGS.

#### Parameters

- **geometry** (*Geometry*) -- Geometry to be optimized.
- **keep\_last** (int, default: 7) -- History size. Keep last 'keep\_last' steps and gradient differences.
- **beta** (float, default: 1) -- Force constant in  $-(H + I)^{-1}g$ .
- **max\_step** (float, default: 0.2) -- Upper limit for the absolute component of the step vector in whatever unit the optimization is carried out.

- **double\_damp** (bool, default: True) -- Use double damping procedure to modify steps  $s$  and gradient differences  $y$  to ensure  $s_y > 0$ .
- **gamma\_mult** (bool, default: False) -- Estimate  $\gamma$  from previous cycle. Eq. (7.20) in [1]. See 'beta' argument.
- **line\_search** (bool, default: False) -- Enable implicit linesearches.
- **mu\_reg** (Optional[float], default: None) -- Initial guess for regularization constant in regularized LBFGS.
- **max\_mu\_reg\_adaptions** (int, default: 10) -- Maximum number of trial steps in regularized LBFGS.
- **control\_step** (bool, default: True) -- Wheter to scale down the proposed step its biggest absolute component is equal to or below 'max\_step'
- **\*\*kwargs** -- Keyword arguments passed to the Optimizer baseclass.

**get\_lbfgs\_step**(*forces*)

**optimize**()

**postprocess\_opt**()

**reset**()

```
class pysisiphus.optimizers.PreconLBFGS.PreconLBFGS(geometry, alpha_init=1.0, history=7,
                                                  precon=True, precon_update=1,
                                                  precon_getter_update=None, precon_kind='full',
                                                  max_step_element=None, line_search='armijo',
                                                  c_stab=None, **kwargs)
```

Bases: [Optimizer](#)

```
__init__(geometry, alpha_init=1.0, history=7, precon=True, precon_update=1,
         precon_getter_update=None, precon_kind='full', max_step_element=None, line_search='armijo',
         c_stab=None, **kwargs)
```

Preconditioned limited-memory BFGS optimizer.

See `pysisiphus.optimizers.precon` for related references.

#### Parameters

- **geometry** (*Geometry*) -- Geometry to be optimized.
- **alpha\_init** (float, default: 1.0) -- Initial scaling factor for the first trial step in the explicit line search.
- **history** (int, default: 7) -- History size. Keep last 'history' steps and gradient differences.
- **precon** (bool, default: True) -- Wheter to use preconditioning or not.
- **precon\_update** (int, default: 1) -- Recalculate preconditioner P in every n-th cycle with the same topology.
- **precon\_getter\_update** (Optional[int], default: None) -- Recalculate topology for preconditioner P in every n-th cycle. It is usually sufficient to only determine the topology once at the beginning.
- **precon\_kind** (Literal['full', 'full\_fast', 'bonds', 'bonds\_bends'], default: 'full') -- What types of primitive internal coordinates to consider in the preconditioner.

- **max\_step\_element** (Optional[float], default: None) -- Maximum component of the absolute step vector when no line search is carried out.
- **line\_search** (Literal['armijo', 'armijo\_fg', 'strong\_wolfe', 'hz', None, False], default: 'armijo') -- Whether to use explicit line searches and if so, which kind of line search.
- **c\_stab** (Optional[float], default: None) -- Regularization constant  $c$  in  $(H + cI)^1$  in atomic units.
- **\*\*kwargs** -- Keyword arguments passed to the Optimizer baseclass.

**get\_precon\_getter()**

**optimize()**

**prepare\_opt()**

**scale\_max\_element**(*step*, *max\_step\_element*)

## RELAXED SCANS

Pysisyphus currently supports 1d-relaxed scans, where one internal coordinate is kept frozen, while the remaining ones are relaxed. A sample input is given below.

```
geom:
  type: redund
  fn: lib:h2o2_hf_321g_opt.xyz
calc:
  type: pyscf
  basis: 321g
  pal: 4
scan:
  type: BOND
  indices: [2, 3]
  # start is optional; if not specified the initial value of the primitive
  # internal coordinate is used.
  start: 3.0
  end: 5.0
  # steps is optional and can be omitted, but then step_size must be given
  steps: 5
  # step_size: 0.4
  # Enable symmetric scan in both directions, starting from the initial coordinates
  # symmetric: False
  #
  # By specifying an 'opt' block, the default optimization values, e.g.
  # convergence thresholds can be altered.
  # opt:
  # thresh: gau
```

If no *start* value is specified in the *scan:* block, then the initial value is used. Specification of *steps* is mandatory, as it determines the number of optimizations to carry out. If not directly given (*step\_size*), a suitable step size is derived from an *end* value. Either *end* or *step\_size* must be given. The values *step\_size*, *start*, *end* have to be given in atomic units or radians.

Naive relaxed scan around the equilibrium value of a primitive internal may be affected by hysteresis, if the system under study is sufficiently complicated.

Relaxed scans are always carried out in redundant internal coordinates. Please see Optimization of Minima for the available optimization options. Supported primitive internals (*type*) are found in: *Types of Primitive Coordinates*.

All geometries, including the initial one are written to *relaxed\_scan.trj* in the current working directory.



## EXCITED STATE OPTIMIZATION

pysisyphus offers excited state (ES) tracking, to follow diadiabatic states along an optimization. ES tracking is enabled by putting `track: True` in the `calc` section of your YAML input. In ES optimizations pysisyphus may use additional programs (`wfoverlap`, `Multiwfn`, `jmol`) if requested. Please see the [installation instructions](#) for information on how to set them up.

**Please consider reading the relevant sections (2 and 3; 4 discusses examples) of the [pysisyphus paper](#).** Additionally the user should think about the relevance between equilibrium/non-equilibrium solvation when calculating ES-gradients with implicit solvation. Several programs, e.g., Gaussian use equilibrium solvation when doing ES-optimization. It is in the responsibility of the user to add the relevant keywords when using pysisyphus, e.g., `EqSolv` for Gaussian.

## 10.1 YAML Example

A bare-bone input for the  $S_1$  optimization of the 1H-amino-keto tautomer of cytosin at the TD-DFT/PBE0/def2-SVP level of theory using ORCA is shown below. The full example is found [here](#).

```
opt:
calc:
  type: orca
  keywords: pbe0 def2-svp rijcosx def2/J def2-svp/C
  # Calculate 2 ES by TD-DFT, follow the first one
  blocks: "%tddft nroots 2 iroot 1 tda false end"
  charge: 0
  mult: 1
  pal: 4
  mem: 2000
  # ES-tracking related keywords follow from here
  # Enable ES-tracking, this is important.
  track: True
  # Track ES by transition density overlaps
  ovlp_type: tden
geom:
  type: redund
  fn: cytosin.xyz
```

Additional keywords are possible in the `calc` section. The default values are shown below.

```
calc:
  # Controls calculation of charge-density-differences cubes and rendering
  # Cubes are calculated by Multiwfn, rendering is handled by jmol.
```

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```

#
# Possible values are: (None, calc, render).
# None: No cube calculation/rendering
# calc: Cube calculation by Multiwfn.
# render: Same as 'calc' and cubes are then rendered by jmol.
cdds: None
# Overlap type. Using 'wf' requires the external wfoverlap binary. The remaining
# options are implemented directly in pysisyphus.
#
# Possible values are (wf, tden, nto, nto_org).
# wf: Wavefunction overlaps using the external wfoverlap program.
# tden: Transition density matrix overlaps.
# nto: Natural transition orbital overlaps.
# nto_org: Natural transition orbital overlaps as described by Garcia.
ovlp_type: wf
# Controls the reference cycle that is used in the overlap calculation. The default
# 'adapt' is recommended.
#
# Possible values are (first, previous, adapt)
# first: Keep first calculation as reference cycle. Reliable when only minor
#         geometrical changes are expected.
# previous: Use previous cycle as reference.
# adapt: Use adaptive algorithm. Please see the pysisyphus paper for a discussion
#         at the end of section 3.
ovlp_with: adapt
# Thresholds controlling the update of the reference cycle.
# The first number specifies the minimum overlap that must be exceeded, for an update
# of the reference cycle. Assuming a value of 0.5 (50 %), the reference cycle update
# is skipped, if the overlaps between the current states and the reference state don't
# exceed 50 %.
# The last two numbers define an interval for the ratio between the second highest
# overlap, and the highest overlap. If the ratio is small, e.g., below 0.3, then both
# states are sufficiently different, and no reference cycle update is needed. If the
↳ratio
# is bigger (> 0.6), then the states are quite similar, and an update is currently not
# advised.
# Possible values [three positive floats between 0. and 1.]
adapt_args: [0.5, 0.3, 0.6]
# Explicitly calculate the AO-overlap matrix in a double molecule calculation. Only
# supported by Turbomole and Gaussian calculators. If False, the approximate AO
# overlap matrix is reconstructed from inverting the MO-coefficient matrix.
#
# Possible values: (True, False)
double_mol: False
# Absolute CI-coefficients below this threshold are ignored in the overlap calculation.
#
# Possible values: positive float
conf_thresh: 0.0001
#
# nto/natural transition orbital specific
#
# Number of NTOs to consider in the overlap calculation. Only relevant for 'nto'

```

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```

# and 'nto_org' ovlp_types.
#
# Possible values: positive integer
use_ntos: 4
# Dynamically decide on number of NTOs according to their participation ratio. Only
# relevant for 'nto_org'
#
# Possible values: boolean
pr_nto: False
#
# wfoverlap/wavefunction overlaps specific
#
# Number of core orbitals to neglect in a wfoverlap calculation. Only relevant
# for the 'wf' ovlp_type. Must be >= 0.
#
# Possible values: positive integer
ncore: 0
#
# tden/transition density matrix specific
#
# Controls which set of MO coefficients (at current cycle, or the reference cycle)
# is used to recover the AO overlap matrix.
#
# Possible values: (ref, cur)
mos_ref: cur
# Controls whether the set of MO coefficients that was NOT used for recovering the AO
# overlap matrix is re-normalized, using the recovered AO overlap matrix. If set to
# True and mos_ref = cur, then the MO coefficients at the reference cycle will be re-
# normalized, and vice versa.
#
# Possible values: (True, False)
mos_renorm: True

```

By brief reasoning it would seem that `mos_ref: ref` and `mos_renorm: True` are more sensible choices, which is possibly true. Right now the present defaults are kept for legacy reasons, and I'll update them after testing out the alternatives.

Please also see Example - Excited State Tracking for possible visualizations when optimizing ES.

## 10.2 Optimization of Conical Intersections

pysisyphus implements the projected gradient method using an updated branching plane, as developed by Maede, Ohno and Morokuma. Currently, CI-optimization is not enabled for YAML input. An illustrative example is found in `tests/test_conic_intersect`.



## CHAIN OF STATES METHODS

A chain of states (COS) comprises a set of distinct states (images) and is usually spanned between two minima on a potential energy surface (PES).

When properly relaxed, a COS coincides with a minimum energy path (MEP), or is a good approximation to it. Tangents can be defined for every COS image and together they make up a discretized path that describes the reaction/chemical transformation. The tangents are also used to divide the COS gradient into perpendicular and parallel components, w.r.t the tangents. Relaxation (optimization) of the COS is achieved by minimizing the perpendicular component of the gradient, so only the parallel component remains.

*pysisyphus* offers different COS implementations, namely Nudged Elastic Band (NEB), including its Adaptive and Free-End (and Free-End-Adaptive) modifications and different flavors of String methods (Growing String Method, GSM, and Simple Zero Temperature String, SZTS). The GSM implementation is also available for with internal coordinates.

### 11.1 String method

(When discussing String methods the COS 'images' are usually called 'nodes'.)

In the string method the whole COS is periodically (every  $n$ -th cycle) reparametrized by fitting a spline through all nodes and redistributing the nodes them along the spline, according to a predefined parametrization. By choosing between different parametrizations equal node spacing or higher resolution around the highest energy image (HEI) can be achieved. The tangents needed for the gradient projection are obtained as first derivatives of the spline.

Reparametrization every  $n$ -th cycle impedes efficient string optimization, and prevents the use of optimizers with some kind of history Conjugate Gradient (CG). The optimizer history is reset after each reparametrization and a simple Steepest Descent (SD) step is done after reparametrization.

### 11.2 Nudged Elastic Band

No reparametrization takes place in the NEB method. The parallel gradient component along the tangent is projected out and replaced by an artificial spring force. In principle, optimizing NEBs should be easier as there is no reparametrization and more sophisticated optimizers beyond SD can and should be employed.

## 11.3 General remarks

Converged COS produce good guesses for subsequent TS searches, when the (splined) HEI is determined. In *pysisyphus* subsequent TS searches are easily started by including *tsopt*: in the YAML input. Knowledge of the initial and final images of the COS is used to construct a more complete set of internal coordinates for the TS guess and it is less likely that important coordinates are missed. The initial imaginary mode to follow uphill is selected as the one featuring the highest overlap with HEI tangent.

## 11.4 YAML example(s)

Below you can find an example YAML-input including the most important options that the user may want to modify when running a GSM optimization.

```
precontr:           # Preconditioning of translation & rotation
preopt:            # Preoptimize initial and final geometry
cos:
  type: gs         # Do a growing string
  max_nodes: 9    # Total string will have 9 + 2 == 11 images
  climb: False    # Enable climbing image (CI), usually a good
↳idea.
  climb_rms: 0.005 # rms(forces) threshold for enabling CI
  climb_lanczos: False # Use tangent obtained from Lanczos algorithm
↳for CI.
  climb_lanczos_rms: 0.005 # rms(forces) threshold for enabling Lanczos
↳algorithm.
  reparam_check: rms # Criterion for growing new frontier nodes (rms/
↳norm).
  perp_thresh: 0.05 # Threshold for growing new frontier nodes.
  reparam_every: 2 # Reparametrize every n-th cycle when NOT fully
↳grown.
  reparam_every_full: 3 # Reparametrize every n-th cycle when fully
↳grown.
opt:
  type: string     # Optimizer for GrowingString
  stop_in_when_full: -1 # Stop string optimization N cycles after fully
↳grown
                    # Disabled by -1. Usually it's a better idea to
                    # further converge the string, after is fully
↳grown.
  align: False    # Disable Kabsch algorithm. Should be True with
                    # type: cart
scale_step: global # Scale down step as whole (global) or per image
                    # (per_image)
tsopt:
  type: rsprfo    # Continue with TS-optimization of highest
↳energy images
                    # (HEI) using the RS-P-RFO algorithm
  do_hess: True   # Calculate hessian at optimized TS geometry
  trust_max: 0.3
  thresh: gau_loose
calc:
```

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```

type: orca
keywords: "b3lyp 6-31G* rijcosx"
pal: 4
charge: 0
mult: 1
geom:
  type: dlc
  fn: [first_preopt.xyz, last_preopt.xyz] # Run GrowingString in delocalized internal_
↳coordinates
                                     # (preferred).

```

For NEB optimizations a different optimizer (not `type: string`) should be used, e.g., QuickMin `type: qm` in the beginning, and `type: lbfgs` later on. It is not yet possible to specify two different optimizers that are used in stages, so if it is desired it must be done manually.

```

# Taken from examples/complex/06_diels_alder...
geom:
  type: cart
  fn: diels_alder.trj
calc:
  type: xtb
  charge: 0
  mult: 1
  pal: 4
preopt:
  max_cycles: 5
interpol:
  type: redund
  between: 10
↳images,
                                     # In NEBs the whole path is interpolated beforehand.
                                     # Possible values: redund/idpp/lst/linear
                                     # Interpolate n-geometries between every pair of supplied
                                     # geometries. For two this yields 1 + 10 + 1 == 12_
↳== 23
                                     # for three geometries this yields 1 + 10 + 1 + 10 + 1_
                                     # geometries.
cos:
  type: neb
  climb: False
opt:
  type: lbfgs
  align: True
  rms_force: 0.01
  max_step: 0.04
tsopt:
  type: rsirfo
  do_hess: True
  max_cycles: 75
  thresh: gau_tight
  hessian_recalc: 7
irc:
  type: eulerpc
  rms_grad_thresh: 0.0005
endopt:

```

Further examples for COS optimizations from `.yaml` input can be found [here](#).

## 11.5 General advice for COS optimizations

- Start from optimized geometries or use the `preopt:` key in the YAML input.
- Consider fixing the initial and final images `fix_ends: True`
- Always use `align: True` when optimizing a COS in cartesian coordinates to remove translation and rotation. `align: True` must not be used when running a COS with DLC.
- Don't over-converge a COS. It is usually a better idea to converge a COS loosely and use the highest energy image (HEI) as a guess for a subsequent transition state (TS) search.
- If possible use a climbing image (CI) `climb: True`
- When running a growing string calculation (`type: gs`) use `stop_in_when_full: [n]` in the `opt:` section with a small integer `[n]` to stop the COS relaxation after the string is fully grown

## 11.6 Chain Of States base class

Base class for chain of state methods

```
class pysisyphus.cos.ChainOfStates.ChainOfStates(images, fix_ends=False, fix_first=True,
                                               fix_last=True, climb=False, climb_rms=0.005,
                                               climb_lanczos=False, climb_lanczos_rms=0.005,
                                               climb_fixed=True, energy_min_mix=False,
                                               scheduler=None, progress=False)

    as_xyz(comments=None)

    property atoms

    calculate_forces()

    property calculator

    property cart_coords
        Return a flat 1d array containing the cartesian coordinates of all images.

    check_for_climbing_start(ref_rms)

    clear()

    property coords
        Return a flat 1d array containing the coordinates of all images.

    property coords3d

    describe()

    property energy

    property forces

    get_climbing_forces(ind)
```

**get\_climbing\_indices()**

**get\_dask\_client()**

**get\_fixed\_indices()**

**get\_hei\_index(energies=None)**

Return index of highest energy image.

**get\_image\_calc\_counter\_sum()**

**get\_perpendicular\_forces(i)**

[1] Eq. 12

**get\_splined\_hei()**

**get\_tangent(i, kind='upwinding', lanczos\_guess=None, disable\_lanczos=False)**

[1] Equations (8) - (11)

**get\_tangents()**

**property gradient**

**property image\_inds**

**property last\_index**

**log(message)**

**logger = <Logger cos (DEBUG)>**

**property masses\_rep**

**property max\_image\_num**

**property moving\_images**

**property moving\_indices**

Returns the indices of the images that aren't fixed and can be optimized.

**par\_image\_calc(image)**

**property perpendicular\_forces**

**prepare\_opt\_cycle(last\_coords, last\_energies, last\_forces)**

Implements additional logic in preparation of the next optimization cycle.

Should be called by the optimizer at the beginning of a new optimization cycle. Can be used to implement additional logic as needed for AdaptiveNEB etc.

**property results**

**rms(arr)**

Root mean square

Returns the root mean square of the given array.

**Parameters**

**arr** (*iterable of numbers*) --

**Returns**

**rms** -- Root mean square of the given array.

**Return type**

float

**set\_climbing\_forces**(*forces*)**set\_coords\_at**(*i*, *coords*)

Called from helpers.procrustes with cartesian coordinates. Then tries to set cartesian coordinate as self.images[i].coords which will raise an error when coord\_type != "cart".

**set\_images**(*indices*, *images*)**set\_vector**(*name*, *vector*, *clear=False*)**set\_zero\_forces\_for\_fixed\_images**()

This is always done in cartesian coordinates, independent of the actual coord\_type of the images as setting forces only work with cartesian forces.

**valid\_coord\_types** = ('cart', 'cartesian', 'dlc')**zero\_fixed\_vector**(*vector*)

## 11.7 Chain Of State Methods

### 11.7.1 Nudged Elastic Band (NEB)

```
class pysisyphus.cos.NEB.NEB(images, variable_springs=False, k_max=0.3, k_min=0.1,  
                             perp_spring_forces=None, **kwargs)
```

Bases: *ChainOfStates***fmt\_k**()property **forces****get\_parallel\_forces**(*i*)**get\_quenched\_dneb\_forces**(*i*)

See [3], Sec. VI and [4] Sec. D.

**get\_spring\_forces**(*i*)property **parallel\_forces****set\_variable\_springs**()**update\_springs**()

## 11.7.2 Adaptive NEB

```
class pysisiphus.cos.AdaptiveNEB.AdaptiveNEB(images, adapt=True, adapt_fact=0.25, adapt_between=1,
                                             scale_fact=False, keep_hei=True, free_ends=True,
                                             **kwargs)
```

Bases: [NEB](#)

```
__init__(images, adapt=True, adapt_fact=0.25, adapt_between=1, scale_fact=False, keep_hei=True,
         free_ends=True, **kwargs)
```

(Free-End) Adaptive Nudged Elastic Band.

### Parameters

- **images** (*list of Geometry objects*) -- Images of the band.
- **adapt** (*bool, default True*) -- Whether to adapt the image number or not. This switch is included to support the FreeEndNEB class, that is just a thin wrapper around this class.
- **adapt\_fact** (*positive float*) -- Factor that is used to decide whether to adapt. The initial threshold is calculated by multiplying the RMS force of the band with this factor. When the RMS of the force falls below this threshold adaption takes place.
- **adapt\_between** (*positive integer*) -- Number of images to interpolate between the highest energy image and its neighbours. The number of starting images must be higher or equal then  $2 * \text{adapt\_between} + 3$ , as we reuse/transfer the calculators from the starting images onto the new ones.
- **scale\_fact** (*bool, default False*) -- Whether to increase `adapt_fact` in deeper levels. This may lead to earlier adaption.
- **keep\_hei** (*bool, optional*) -- Whether to keep the highest energy image (usually a very good idea) or to interpolate only between the neighbouring images.
- **free\_ends** (*bool, default True*) -- Whether to use modified forces on the end images.

```
adapt_this_cycle(forces)
```

Decide whether to adapt.

### Parameters

**forces** (*np.array*) -- Forces of the previous optimization cycle.

### Returns

**adapt** -- Flag that indicates if adaption should take place in this cycle.

### Return type

bool

```
property forces
```

See Eq. (7) in [2].

```
prepare_opt_cycle(*args, **kwargs)
```

Check for adaption and adapt if needed.

See `ChainOfStates.prepare_opt_cycle` for a complete docstring.

```
update_adapt_thresh(forces)
```

Update the adaption threshold.

### Parameters

**forces** (*np.array*) -- Forces of the previous optimization cycle.

### 11.7.3 Free-End NEB

```
class pysisyphus.cos.FreeEndNEB.FreeEndNEB(*args, fix_first=False, fix_last=False, **kwargs)
```

Bases: *AdaptiveNEB*

```
__init__(*args, fix_first=False, fix_last=False, **kwargs)
```

Simple Free-End-NEB method.

Derived from AdaptiveNEB with disabled adaptation. Only implements Eq. (7) from [2]. For other implementations please see the commit 01bc8812ca6f1cd3645d43e0337d9e3c5fb0ba55. There the other variants are present but I think Eq. (7) in [2] is the simplest & best bet.

### 11.7.4 Simple Zero-Temperature String

```
class pysisyphus.cos.SimpleZTS.SimpleZTS(images, param='equal', **kwargs)
```

Bases: *ChainOfStates*

```
reparametrize()
```

## 11.8 Growing Chain Of States base class

Base class for growing chain of state methods

```
class pysisyphus.cos.GrowingChainOfStates.GrowingChainOfStates(images, calc_getter,
                                                                max_nodes=10, **kwargs)
```

```
property arc_dims
```

```
get_new_image_from_coords(coords, index)
```

```
property max_image_num
```

```
new_node_coords(k)
```

```
prepare_opt_cycle(*args, **kwargs)
```

Implements additional logic in preparation of the next optimization cycle.

Should be called by the optimizer at the beginning of a new optimization cycle. Can be used to implement additional logic as needed for AdaptiveNEB etc.

```
set_new_node(k)
```

## 11.9 Growing Chain Of State Methods

### 11.9.1 Growing String Method

```
class pysisyphus.cos.GrowingString.GrowingString(images, calc_getter, perp_thresh=0.05,
                                                    param='equi', reparam_every=2,
                                                    reparam_every_full=3, reparam_tol=None,
                                                    reparam_check='rms', max_micro_cycles=5,
                                                    reset_dlc=True, climb=False, **kwargs)
```

Bases: *GrowingChainOfStates*

**property forces**

**property full\_string\_image\_inds**

**property fully\_grown**

Returns whether the string is fully grown. Don't count the first and last node.

**get\_additional\_print()**

**get\_cur\_param\_density(*kind=None*)**

**get\_new\_image(*ref\_index*)**

Get new image by taking a step from `self.images[ref_index]` towards the center of the string.

**get\_tangent(*i*)**

[1] Equations (8) - (11)

**property image\_inds**

**property left\_size**

**property lf\_ind**

Index of the left frontier node in `self.images`.

**property nodes\_missing**

Returns the number of nodes to be grown.

**reparam\_cart(*desired\_param\_density*)**

**reparam\_dlc(*desired\_param\_density, thresh=0.001*)**

**reparametrize()**

**reset\_geometries(*ref\_geometry*)**

**property rf\_ind**

Index of the right frontier node in `self.images`.

**property right\_size**

**set\_coords(*image, coords*)**

**spline(*tangents=False*)**

**property string\_size**



## TRANSITION STATE OPTIMIZATION

To cite Frank Jensen: "Locating transition states (TSs) is black magic, especially in internal coordinates" and I can say this is true. The most promising TS optimizers in *pysisyphus* employ second derivative information (hessian) but locating TS is also possible using only first derivatives by means of the *dimer method* (DM).

TS optimizations should preferably be carried out in internal coordinates (*type: redund|dlc*). Before starting the actual TS search, one should **always** check the internal coordinates that *pysisyphus* sets up, by executing *pysistrj [xyz] -internals*, with *[xyz]* corresponding to the geometry you plan to use for the optimization. **Check carefully** if all supposed reaction coordinates are present. **If they are missing** define them manually with the *add\_prims* key in the YAML input.

Calculation of the exact hessian can be avoided by using the DM by using *rx\_coords: [list of primitives]* in combination with *hessian\_init: fischer|lindh|swart|simple*. By using both options, a diagonal model hessian is calculated and modified for use in a TS optimization.

The DM can be used with a powerful preconditioned LBFGS optimizer (*type: lbfgs*).

### 12.1 Hessian Based TS Optimization

#### 12.1.1 YAML example

Below you can find an example YAML-input including the most important options that the user may want to modify for the Hessian-based optimizers (RS-I-RFO, RS-P-RFO, TRIM).

```

tsopt:
  type: rsirfo|rsprfo|trim      # Optimization algorithm

  do_hess: True                # Calculate the hessian at the final geometry
                              # after the optimization.

  hessian_recalc: 1            # Recalculate the exact hessian every n-th cycle
                              # Very costly but a small number like 5 may be a good
                              # idea.

  prim_coord: [BOND, 3, 18]    # Select the mode to follow uphill by overlap with
                              # this primitive internal coordinate. Expects exactly
                              # one typed primitive.

  rx_mode: [[[BOND, 3, 18], 1]] # Select initial mode based on overlaps with a
                              # constructed mode. Can be seen as generalization of
                              # prim_coord. Expects a list of pairs, with each pair

```

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```

# comprising a typed primitive and a phase factor.
#
# See examples/{05,06}... for examples.

#rx_coords: [[BOND, 3, 18],] # Obtain initial mode by modifying a model Hessian. Has
# to be used with 'hess_init: swart/fischer/lindh/xtb'
# to avoid calculation of exact hessian.
# Also requires 'redund' or 'dlc' to work.

#root: 0 # Follow the n-th imaginary mode uphill, minimize
# along the others.

#hessian_init: calc # Initial hessian is calculated exactly.

#hessian_update: bofill # Bofill hessian-update. Should not be modified for
# TS optimizations.

#hessian_ref: [path] # Expects a path to a precalculated hessian at any
# level of theory. The emode to maximize along is
↪selected # by highest overlap with imaginary mode from the
↪reference # hessian.

#max_micro_cycles: 50 # No. of micro cycles for the RS-variants. Does not apply
# to TRIM.

#trust_radius: 0.3 # Initial trust radius.
#trust_max: 1.0 # Max. trust radius
#trust_min: 0.1 # Min. trust radius
calc:
  type: xtb
  pal: 4
  charge: 0
  mult: 1
geom:
  type: redund
  fn: shaked.geom_000.xyz
  add_prims: [[24, 20], ] # If using internal coordinates ALWAYS check the
↪coordinates # that pysisyphus generates (pysistrj [xyz] --internals).
↪ If # some important (reaction) coordinates appears to be
↪missing # define them manually. In this example
# we add a bond. If additional internal coordinates can
# derived from the added primitives pysisyphus will do
# it.

```

Second-derivative-free TS optimization can be done using the *Dimer* method. Attached you can find a sample input.

Further examples for TS optimizations from *.yaml* input can be found [here](#).

## 12.2 Dimer Method

*pysisyphus* implements the dimer method (DM) as calculator, wrapping another calculator for the actual energy and gradient calculations. All DM configurations have to be done in the `calc:` section. The best results are obtained by optimizing the DM with LBFGS in connection with a preconditioner (PLBFGS), estimated from the Lindh model Hessian. In contrast to the TS optimizers mentioned above, PLBFGS is configured in the `opt` section of the YAML input. DM related information is logged to *dimer.log*. The current DM orientation is saved in files with extension *.N*. Animated *.trj* files of the DM orientation are saved to *.N.trj*.

### 12.2.1 YAML example

Below you can find an example for the DM, applied to the isomerization of HCN. Default values are commented out. See `examples/tsopt/02_hcn_tsopt_dimer` for the full example.

```
opt:
  type: plbfgs      # Preconditioned LBFGS
  thresh: baker    # Baker threshold, comparable to the Gaussian defaults
  do_hess: True    # Calculate Hessian at the end. May not be applicable to systems
                  # where Hessian calculation is infeasible.
calc:
  type: dimer      # Dimer calculator wrapping PySCF at HF/3-21G level of theory
  calc:
    type: pyscf
    basis: 321g
    pal: 2
    charge: 0
    mult: 1
    #N_raw: [file]      # Path to a file containing an initial dimer orientation
    #length: 0.0189    # Separation of dimer images from common midpoint
    #rotation_max_cycles: 15 # Maximum number of rotations
    #rotation_method: fourier
    #rotation_thresh: 1e-4 # rms(rotation force) threshold
    #rotation_tol: 1    # Rotation tolerance in degree. If a proposed trial
                      # rotation angle is below the tolerance the rotation
                      # will be skipped.
    #rotation_max_element: 0.001 # Max. step element for "rotation_method: direct"
    #rotation_interpolate: True # Interpolate force on image after rotation
    #rotation_remove_trans: True # Remove overall translation from N-vector
    #seed: 20182503     # Seed for the RNG for reproducibility reasons.
geom:
  type: cart
  fn: 01_hcn.xyz
```

## 12.3 General advice for TS optimizations

- Use as many exact Hessians as your computational budget allows (*hessian\_recalc*: [n])
- Use tight convergence settings for your SCF. In contrast to codes like ORCA *pysisyphus* does not enforce this automatically
- Grid-based methods (DFT, some RI-types) may need finer grids to reduce numerical noise
- Try to preoptimize the TS using a cheap method (xtb) that allows *hessian\_recalc*: 1
- Decrease current & allowed trust radius (*trust\_radius*: 0.1 and *trust\_max*: 0.3|0.5)
- Check for correct coordinate setup

## 12.4 TSHessianOptimizer base class

Base class for TS optimizers that employ a hessian.

```
class pysisyphus.tsoptimizers.TSHessianOptimizer.TSHessianOptimizer(geometry, root=0,
                                                                    hessian_ref=None,
                                                                    prim_coord=None,
                                                                    rx_coords=None,
                                                                    rx_mode=None,
                                                                    hessian_init='calc',
                                                                    hessian_update='bofill',
                                                                    hessian_recalc_reset=True,
                                                                    max_micro_cycles=50,
                                                                    trust_radius=0.3,
                                                                    trust_max=0.5,
                                                                    augment_bonds=False,
                                                                    min_line_search=False,
                                                                    max_line_search=False,
                                                                    assert_neg_eigval=False,
                                                                    **kwargs)
```

Bases: [HessianOptimizer](#)

Optimizer to find first-order saddle points.

```
__init__(geometry, root=0, hessian_ref=None, prim_coord=None, rx_coords=None, rx_mode=None,
          hessian_init='calc', hessian_update='bofill', hessian_recalc_reset=True, max_micro_cycles=50,
          trust_radius=0.3, trust_max=0.5, augment_bonds=False, min_line_search=False,
          max_line_search=False, assert_neg_eigval=False, **kwargs)
```

Baseclass for transition state optimizers utilizing Hessian information.

Several arguments expect a typed primitive or an iterable of typed primitives. A typed primitive is specified as (PrimType, int, int, ...), e.g., for a bond between atoms 0 and 1: (BOND, 0, 1) or for a bend between the atom triple 0, 1, 2 as (BEND, 0, 1, 2).

### Parameters

- **geometry** (*Geometry*) -- Geometry to be optimized.
- **root** (int, default: 0) -- Index of imaginary mode to maximize along.
- **hessian\_ref** (Optional[str], default: None) -- Filename pointing to a pysisyphus HDF5 Hessian.

- **prim\_coord** (*typed\_prim*) -- Select initial root/imaginary mode by overlap with this internal coordinate.
- **rx\_coords** (*iterable of (typed\_prim)*) -- Construct imaginary mode comprising the given typed primis by modifying a model Hessian.
- **rx\_mode** (*iterable of (typed\_prim, phase\_factor)*) -- Select initial root by overlap with a mode constructed from the given typed primitives.
- **hessian\_init** (Literal['calc', 'unit', 'fischer', 'lindh', 'simple', 'swart', 'xtb', 'xtb1', 'xtb1f'], default: 'calc') -- Type of initial model Hessian.
- **hessian\_update** (Literal['none', None, False, 'bfgs', 'damped\_bfgs', 'flowchart', 'bofill', 'ts\_bfgs', 'ts\_bfgs\_org', 'ts\_bfgs\_rev'], default: 'bofill') -- Type of Hessian update. Defaults to BFGS for minimizations and Bofill for saddle point searches.
- **hessian\_recalc\_reset** (bool, default: True) -- Whether to skip Hessian recalculation after reset. Undocumented.
- **max\_micro\_cycles** (int, default: 50) -- Maximum number of RS iterations.
- **trust\_radius** (float, default: 0.3) -- Initial trust radius in whatever unit the optimization is carried out.
- **trust\_max** (float, default: 0.5) -- Maximum trust radius.
- **augment\_bonds** (bool, default: False) -- Try to derive additional stretching coordinates from the imaginary mode.
- **min\_line\_search** (bool, default: False) -- Carry out line search along the imaginary mode.
- **max\_line\_search** (bool, default: False) -- Carry out line search in the subspace that is minimized.
- **assert\_neg\_eigval** (bool, default: False) -- Check for the existences for at least one significant negative eigenvalue. If enabled and no negative eigenvalue is present the optimization will be aborted.
- **\*\*kwargs** -- Keyword arguments passed to the HessianOptimizer/Optimizer baseclass.

```
static do_line_search(e0, e1, g0, g1, prev_step, maximize, logger=None)
```

```
prepare_opt(*args, **kwargs)
```

```
step_and_grad_from_line_search(energy, gradient_trans, eigvecs, min_indices)
```

```
update_ts_mode(eigvals, eigvecs)
```

```
valid_updates = ('bofill', 'ts_bfgs', 'ts_bfgs_org', 'ts_bfgs_rev')
```

## 12.5 TS-Optimizer using hessian information

### 12.5.1 Restricted-Step Partitioned-RFO

```
class pysisyphus.tsoptimizers.RSPRFOptimizer.RSPRFOptimizer(geometry, root=0, hessian_ref=None,
    prim_coord=None, rx_coords=None,
    rx_mode=None, hessian_init='calc',
    hessian_update='bofill',
    hessian_recalc_reset=True,
    max_micro_cycles=50,
    trust_radius=0.3, trust_max=0.5,
    augment_bonds=False,
    min_line_search=False,
    max_line_search=False,
    assert_neg_eigval=False, **kwargs)
```

Bases: *TSHessianOptimizer*

**optimize()**

### 12.5.2 Restricted-Step Image-Function-RFO

```
class pysisyphus.tsoptimizers.RSIRFOptimizer.RSIRFOptimizer(geometry, root=0, hessian_ref=None,
    prim_coord=None, rx_coords=None,
    rx_mode=None, hessian_init='calc',
    hessian_update='bofill',
    hessian_recalc_reset=True,
    max_micro_cycles=50,
    trust_radius=0.3, trust_max=0.5,
    augment_bonds=False,
    min_line_search=False,
    max_line_search=False,
    assert_neg_eigval=False, **kwargs)
```

Bases: *TSHessianOptimizer*

**optimize()**

### 12.5.3 Trust-Region Image-Function Method

```
class pysisyphus.tsoptimizers.TRIM.TRIM(geometry, root=0, hessian_ref=None, prim_coord=None,
    rx_coords=None, rx_mode=None, hessian_init='calc',
    hessian_update='bofill', hessian_recalc_reset=True,
    max_micro_cycles=50, trust_radius=0.3, trust_max=0.5,
    augment_bonds=False, min_line_search=False,
    max_line_search=False, assert_neg_eigval=False, **kwargs)
```

Bases: *TSHessianOptimizer*

**optimize()**

## 12.6 TS-Optimization without hessian information

### 12.6.1 Dimer method

```
class pysisyphus.calculators.Dimer.Dimer(calculator, *args, N_raw=None, length=0.0189,
                                         rotation_max_cycles=15, rotation_method='fourier',
                                         rotation_thresh=0.0001, rotation_tol=1,
                                         rotation_max_element=0.001, rotation_interpolate=True,
                                         rotation_disable=False, rotation_disable_pos_curv=True,
                                         rotation_remove_trans=True, trans_force_f_perp=True,
                                         bonds=None, N_hessian=None, bias_rotation=False,
                                         bias_translation=False, bias_gaussian_dot=0.1, seed=None,
                                         write_orientations=True, forward_hessian=True, **kwargs)

    property C
        Shortcut for the curvature.

    property N

    add_gaussian(atoms, center, N, height=0.1, std=0.0529, max_cycles=50, dot_ref=None)

    property can_bias_f0

    property can_bias_f1

    property coords0

    property coords1

    curvature(f1, f2, N)
        Curvature of the mode represented by the dimer.

    direct_rotation(optimizer, prev_step)

    do_dimer_rotations(rotation_thresh=None)

    property energy0

    property f0

    property f1

    property f1_bias

    property f2
        Never calculated explicitly, but estimated from f0 and f1.

    fourier_rotation(optimizer, prev_step)

    get_N_raw_from_hessian(h5_fn, root=0)

    get_forces(atoms, coords)
        Meant to be extended.

    get_gaussian_energies(coords, sum_=True)

    get_gaussian_forces(coords, sum_=True)
```

**get\_hessian**(*atoms, coords*)

Meant to be extended.

**remove\_translation**(*displacement*)

**property rot\_force**

**rotate\_coords1**(*rad, theta*)

Rotate dimer and produce new coords1.

**set\_N\_raw**(*coords*)

**property should\_bias\_f0**

May lead to calculation of f0 and/or f1 if present!

**property should\_bias\_f1**

May lead to calculation of f0 and/or f1 if present!

**update\_orientation**(*coords*)

**class** pysisyphus.calculators.Dimer.**Gaussian**(*height, center, std, N*)

**energy**(*R, height=None*)

**forces**(*R, height=None*)

**exception** pysisyphus.calculators.Dimer.**RotationConverged**

## INTRINSIC REACTION COORDINATE (IRC)

By default two paths are integrated in plus- and minus-direction of the imaginary mode (transition vector, TV) at the transition state (TS). If the IRC is started from a non-stationary point with non-vanishing gradient the *downhill: True* argument can be set to integrate only one path and skip the initial hessian calculation.

IRCs are integrated in mass-weighted coordinates and the default integrator is *EulerPC*.

The initial displacement from the TS is done by requiring a certain energy lowering  $dE$  from moving along the TV and calculating the corresponding step length from a quadratic potential:  $dE = \frac{1}{2}(k \cdot dq^2)$ , with  $k$  being the eigenvalue of the TV (imaginary mode) and  $dq$  the required step length. The default required energy lowering is 0.0005 au. Alternatively the initial can be done by a prescribed length: *displ: length* and *displ\_length: [desired step]* (default is  $0.1\sqrt{\text{amu}} \cdot \text{bohr}$ ).

The resulting endpoints of the IRC integration can be further optimized to stationary points by adding the *endopt:* section (vide infra). By setting *fragments: True* in *endopt* separate fragments will be detected and optimized individually. This may be useful if the molecules are only weakly bound. By setting *fragments: total* the total system, as well as the separate fragments will be optimized.

By default IRC path data is dumped to *dump\_fn: irc\_data.h5* every *dump\_every: 5* cycles. IRC paths can be plotted with *pysisplot --irc*.

### 13.1 YAML example(s)

Below you can find an example YAML-input including the most important options that the user may want to modify.

```

geom:
  fn: hfabstraction_ts_opt_xtb.xyz  # Input coordinates
calc:
  type: xtb                        # extended tight-binding calculator
  pal: 4
  charge: 0
  mult: 1
irc:
  type: eulerpc                    # Similar to EulerPC from Gaussian

#rms_grad_thresh: 0.001           # Convergence threshold
#displ: energy/length/energy_cubic # How to do the initial displacement
#displ_energy: 0.001              # Energy lowering in au (Hartree)
#displ_length: 0.1                # Step length along the TV
#forward: True
#backward: True
#downhill: False                  # Only integrate downhill, disables forward/backward

```

(continues on next page)

(continued from previous page)

```

#hessian_init: null           # Path to HDF5 Hessian file
#displ_third_h5: null        # Path to HDF5 file containing third derivative data
endopt:
#fragments: False|True|total # Detect & optimize fragments separately. Default is
                             # False. When set to 'total' the total system as well
                             # as the fragments are optimized.
do_hess: False               # Frequency calculation at the end

```

Further examples for IRC calculations from `.yaml` input can be found [here](#).

## 13.2 IRC base class

Base class for IRC integrators from which actual IRC integrators are derived.

```

class pysisyphus.irc.IRC.IRC(geometry, step_length=0.1, max_cycles=125, downhill=False, forward=True,
                              backward=True, root=0, hessian_init=None, displ='energy',
                              displ_energy=0.001, displ_length=0.1, displ_third_h5=None,
                              rms_grad_thresh=0.001, hard_rms_grad_thresh=None, energy_thresh=1e-06,
                              imag_below=0.0, force_inflection=True, check_bonds=True, out_dir='.',
                              prefix="", dump_fn='irc_data.h5', dump_every=5)

```

```

__init__(geometry, step_length=0.1, max_cycles=125, downhill=False, forward=True, backward=True,
         root=0, hessian_init=None, displ='energy', displ_energy=0.001, displ_length=0.1,
         displ_third_h5=None, rms_grad_thresh=0.001, hard_rms_grad_thresh=None,
         energy_thresh=1e-06, imag_below=0.0, force_inflection=True, check_bonds=True, out_dir='.',
         prefix="", dump_fn='irc_data.h5', dump_every=5)

```

Base class for IRC calculations.

### Parameters

- **geometry** (*Geometry*) -- Transition state geometry, or initial geometry for downhill run.
- **step\_length** (*float, optional*) -- Step length in unweighted coordinates.
- **max\_cycles** (*int, optional*) -- Positive integer, controlling the maximum number of IRC steps taken in a direction (forward/backward/downhill).
- **downhill** (*bool, default=False*) -- Downhill run from a non-stationary point with non-vanishing gradient. Disables forward and backward runs.
- **forward** (*bool, default=True*) -- Integrate IRC in positive *s* direction.
- **backward** (*bool, default=True*) -- Integrate IRC in negative *s* direction.
- **root** (*int, default=0*) -- Use *n*-th root for initial displacement from TS.
- **hessian\_init** (*str, default=None*) -- Path to Hessian HDF5 file, e.g., from a previous TS calculation.
- **displ** (*str, one of ("energy", "length", "energy\_cubic")*) -- Controls initial displacement from the TS. 'energy' assumes a quadratic model, from which a step length for a given energy lowering (see 'displ\_energy') is determined. 'length' corresponds to a displacement along the transition vector. 'energy\_cubic' considers 3rd derivatives of the energy along the transition vector.
- **displ\_energy** (*float, default=1e-3*) -- Required energy lowering from the TS in au (Hartree). Used with 'displ: energy|energy\_cubic'.

- **displ\_length** (*float*, *default=0.1*) -- Step length along the transition vector. Used only with 'displ: length'.
- **displ\_third\_h5** (*str*, *optional*) -- Path to HDF5 file containing 3rd derivative information. Used with 'displ: energy\_cubic'.
- **rms\_grad\_thresh** (*float*, *default=1e-3*,) -- Convergence is signalled when to root mean square of the unweighted gradient is less than or equal to this value
- **energy\_thresh** (*float*, *default=1e-6*,) -- Signal convergence when the energy difference between two points is equal to or less than 'energy\_thresh'.
- **imag\_below** (*float*, *default=0.0*) -- Require the wavenumber of the imaginary mode to be below the given threshold. If given, it should be a negative number.
- **force\_inflection** (*bool*, *optional*) -- Don't indicate convergence before passing an inflection point.
- **check\_bonds** (*bool*, *optional*, *default=True*) -- Report whether bonds are formed/broken along the IRC, w.r.t the TS.
- **out\_dir** (*str*, *optional*) -- Dump everything into 'out\_dir' directory instead of the CWD.
- **prefix** (*str*, *optional*) -- Short string that is prepended to all files that are created by this class, e.g., trajectories and HDF5 dumps.
- **dump\_fn** (*str*, *optional*) -- Base name for the HDF5 files.
- **dump\_every** (*int*, *optional*) -- Dump to HDF5 every n-th cycle.

#### property coords

**dump\_data**(*dump\_fn=None*, *full=False*)

**dump\_ends**(*path*, *prefix*, *coords=None*, *trj=False*)

#### property energy

**get\_conv\_fact**(*mw\_grad*, *min\_fact=2.0*)

**get\_endpoint\_and\_ts\_geoms**()

**get\_full\_irc\_data**()

**get\_irc\_data**()

**get\_path\_for\_fn**(*fn*)

#### property gradient

##### **initial\_displacement**()

Returns non-mass-weighted steps in +s and -s direction for initial displacement from the TS. Earlier version only returned one step, that was later multiplied by either 1 or -1, depending on the desired IRC direction (forward/backward). The current implementation directly returns two steps for forward and backward direction. Whereas for plus and minus steps for displ 'length' and displ 'energy'

$$\text{step\_plus} = -\text{step\_minus}$$

is valid, it is not valid for displ 'energy\_cubic' anymore. The latter step is formed as

$$x(ds) = ds * v0 + ds**2 * v1$$

```
so
    x(ds) != -x(ds)

as
    ds * v0 + ds**2 * v1 != -ds * v0 - ds**2 * v1 .
```

So, all required step are formed directly and later used as appropriate.

See

<https://aip.scitation.org/doi/pdf/10.1063/1.454172>    <https://pubs.acs.org/doi/10.1021/j100338a027>  
<https://aip.scitation.org/doi/pdf/10.1063/1.459634>

```
irc(direction)

log(msg)

property m_sqrt

mass_weigh_hessian(hessian)

property mw_coords

property mw_gradient

postprocess()

prepare(direction)

report_bonds(prefix, bonds)

report_conv_thresholds()

run()

set_data(prefix)

unweight_vec(vec)

valid_displs = ('energy', 'length', 'energy_cubic')
```

## 13.3 IRC Integrators

### 13.3.1 Damped-Velocity-Verlet integrator

```
class pysisyphus.irc.DampedVelocityVerlet.DampedVelocityVerlet(geometry, v0=0.04, dt0=0.5,  
                                                                error_tol=0.003,  
                                                                max_cycles=150, **kwargs)
```

Bases: *IRC*

```
damp_velocity(velocity)

estimate_error(new_mw_coords)

mw_grad_to_acc(mw_grad)
```

Takes care of the units for the mass-weighted gradient. Converts units of a mass-weighted gradient [Hartree/(Bohr\*amu)] to units of acceleration [ $\sqrt{\text{amu}} \cdot \text{Bohr}/\text{fs}^2$ ]. The 1e30 comes from converting second<sup>2</sup> to femto second<sup>2</sup>.

**prepare**(*direction*)

**step**()

### 13.3.2 Euler integrator

Not recommended as it only produces reasonable results with very small step sizes.

**class** pysisyphus.irc.Euler.**Euler**(*geometry, step\_length=0.01, \*\*kwargs*)

Bases: *IRC*

**step**()

### 13.3.3 Euler-Predictor-Corrector integrator

Recommended IRC integrator and default choice.

**class** pysisyphus.irc.EulerPC.**EulerPC**(*\*args, hessian\_recalc=None, hessian\_update='bofill',  
hessian\_init='calc', max\_pred\_steps=500, loose\_cycles=3,  
dump\_dwi=False, scipy\_method=None, corr\_func='mbs',  
\*\*kwargs*)

Bases: *IRC*

**corrector\_step**(*init\_mw\_coords, step\_length, dwi*)

**get\_integration\_length\_func**(*init\_mw\_coords*)

**prepare**(*\*args, \*\*kwargs*)

**scipy\_corrector\_step**(*init\_mw\_coords, step\_length, dwi*)

Solve IRC equation  $dx/ds = -g/|g|$  on DWI PES in mass-weighted coordinates. Integration done until `self.step_length` in unweighted coordinates is achieved.

**step**()

### 13.3.4 Gonzales-Schlegel-2 integrator

**class** pysisyphus.irc.GonzalezSchlegel.**GonzalezSchlegel**(*geometry, max\_micro\_cycles=20,  
micro\_step\_thresh=0.001,  
hessian\_recalc=None, line\_search=False,  
\*\*kwargs*)

Bases: *IRC*

**micro\_step**(*counter*)

Constrained optimization on a hypersphere.

**perp\_component**(*vec, perp\_to*)

**postprocess**()

**step**()

### 13.3.5 Local-Quadratic-Approximation integrator

```
class pysisyphus.irc.LQA.LQA(geometry, N_euler=5000, **kwargs)
```

Bases: *IRC*

```
step()
```

### 13.3.6 Modified-Ishida-Morokuma-Komornicki integrator

Similar to the algorithm implemented by ORCA 4.

```
class pysisyphus.irc.IMKMod.IMKMod(geometry, corr_first=True, corr_first_size=0.5,
                                   corr_first_energy=True, corr_bisec_size=0.0025,
                                   corr_bisec_energy=True, **kwargs)
```

Bases: *IRC*

```
fit_grad_and_energies(energy0, grad0, energy1, direction)
```

```
fit_parabola(x, y)
```

```
step()
```

### 13.3.7 Runge-Kutta-4 integrator

Not recommended, as it is very slow.

```
class pysisyphus.irc.RK4.RK4(geometry, step_length=0.1, max_cycles=125, downhill=False, forward=True,
                               backward=True, root=0, hessian_init=None, displ='energy',
                               displ_energy=0.001, displ_length=0.1, displ_third_h5=None,
                               rms_grad_thresh=0.001, hard_rms_grad_thresh=None, energy_thresh=1e-06,
                               imag_below=0.0, force_inflection=True, check_bonds=True, out_dir='.',
                               prefix='', dump_fn='irc_data.h5', dump_every=5)
```

Bases: *IRC*

```
get_k(mw_coords)
```

```
step()
```

## PLOTTING

`pysisyphus` offers extensive visualization capabilities via the `pysisplot` command. All relevant information produced by `pysisyphus` and the interfaced quantum chemistry programs is dumped to HDF5 files, namely:

Filename	Description
optimization.h5	Produced by all optimizers when <code>dump: True</code> (default).
afir.h5	Contains true and modified energies & forces in AFIR calculations.
forward_irc_data.h5	IRC data from forward run.
backward_irc_data.h5	IRC data from backward run.
finished_irc_data.h5	IRC data from forward & backward run.
overlap_data.h5	From excited state calculations by <code>OverlapCalculator</code> classes.

A help message that shows possible usages of `pysisplot` is displayed by

```
pysisplot --help
```

If needed, results from multiple optimizations are written to different HD5-groups of the same HDF5-file. Available groups can be listed either by the HDF5-tool `h5ls` or `pysisplot --h5_list [hd55 file]`. The desired group for visualization is then selected by the `--h5_group [group name]` argument.

## 14.1 Example - Diels-Alder reaction

The plotting capabilities of `pysisplot` are demonstrated for the example of a Diels-Alder reaction between ethene and 1-4-butadiene (see `examples/06_diels_alder_xtb_preopt_neb_tsopt_irc`) at `xtb-GFN2` level of theory. Educts and product of the reaction are preoptimized for 5 cycles, then interpolated in internal coordinates. A chain of states (COS) is optimized loosely via a LBFGS optimizer. The highest energy image (HEI) is employed for a subsequent transition state (TS) optimization. Finally the intrinsic reaction coordinate (IRC) is obtained by means of an Euler-Predictor-Corrector integrator.

### 14.1.1 Plotting optimization progress

The progress of an optimization can be plotted via

```
pysisplot --opt [--h5_fn optimization.h5] [--h5_group opt]
```

Arguments given in parantheses are the assumed defaults. The result is shown below for the preoptimizations of educts and product.

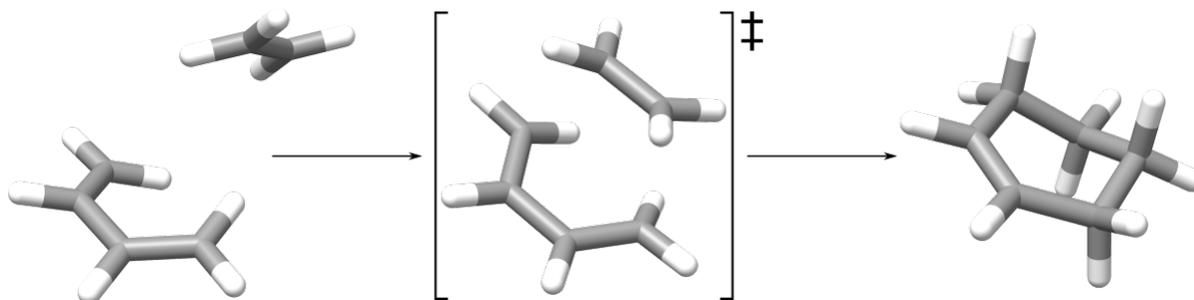


Fig. 14.1: Educts, TS and products of the Diels-Alder reaction between ethene and 1-4-butadiene.

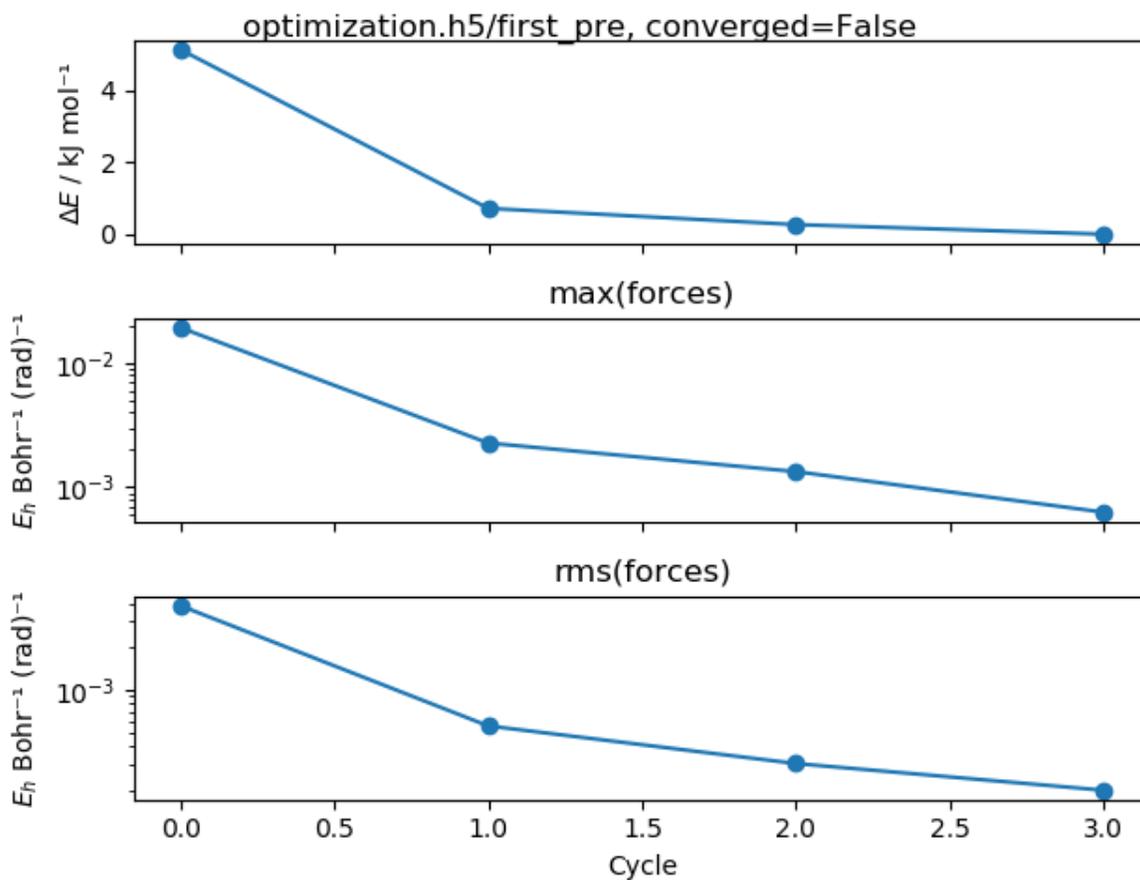


Fig. 14.2: Preoptimization of the Diels-Alder reaction educts ethene + 1,4-butadiene. The upper panel shows the energy change along the optimization. Middle and lower panel show the max and rms of the force. The HDF5 filename and group are noted in the image title.

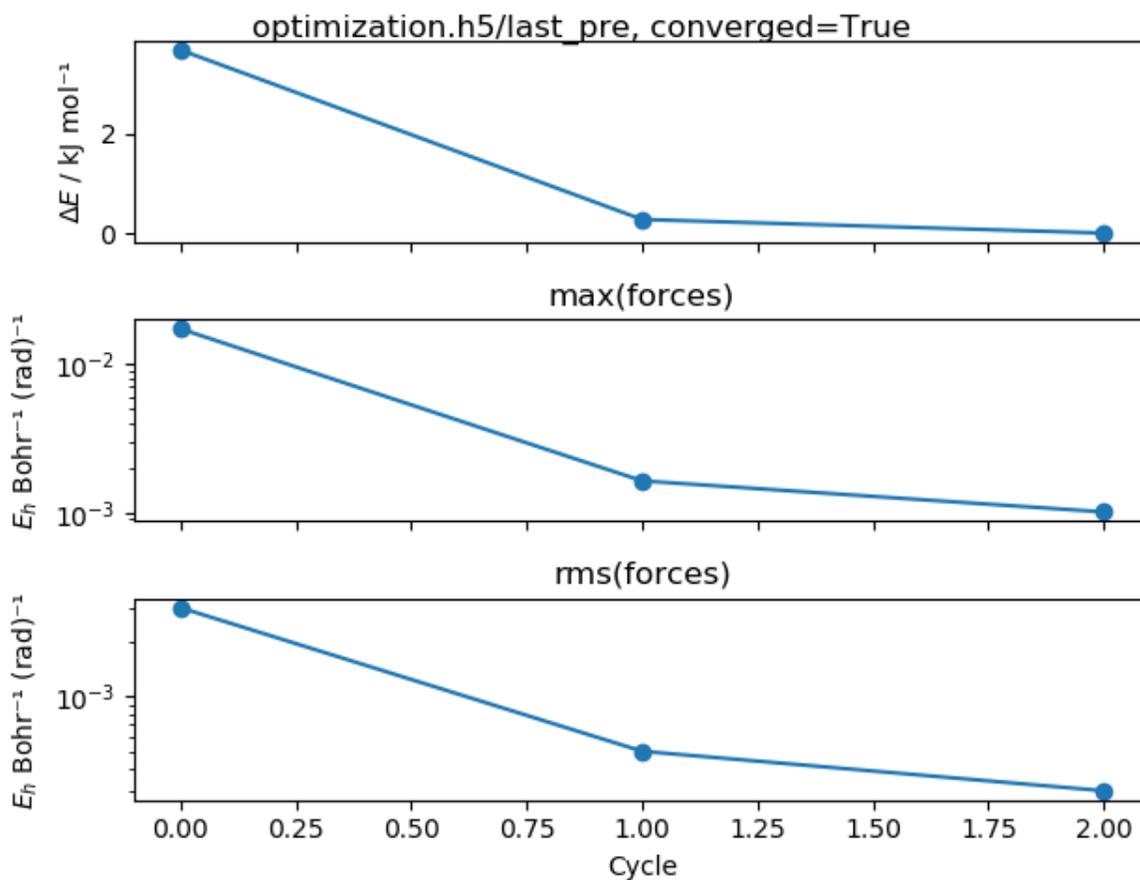


Fig. 14.3: Preoptimization of the Diels-Alder reaction product. Compared to the educts the optimization already converged after 4 cycles.

For COS optimizations all image energies of one optimization cycle are summed into a total energy, which is then plotted in the first panel. *pysisplot --opt* may not be the best choice in these cases. Use *pysisplot --cosens* and *pysisplot --cosforces* instead (see below).

### 14.1.2 Plotting COS optimization progress

Compared to simple surface-walking optimizations of single molecules, COSs consist of multiple images, each with its own energy and force vector. In this case a simple plot as shown above is not suitable. Instead of *pysisplot --opt* a better visualization is offered by *pysisplot --cosens* and *pysisplot --cosforces*. The latter two commands are compatible with all COS methods available in pysisyphus.

```
pysisplot --cosens [--h5_fn optimization.h5] [--h5_group opt]
```

This produces three plots.

1. Animated. COS image energies along the optimization.
2. Static. Energies of last optimization cycle.
3. Static. Energies of all cycles with earlier cycles given in a lighter shade and more recent cycles in a darker shade. The last cycle is splined and the position of the splined HEI is indicated.

Please note that equidistant image spacing is assumed for the latter two plots. Here only the two latter plots are shown.

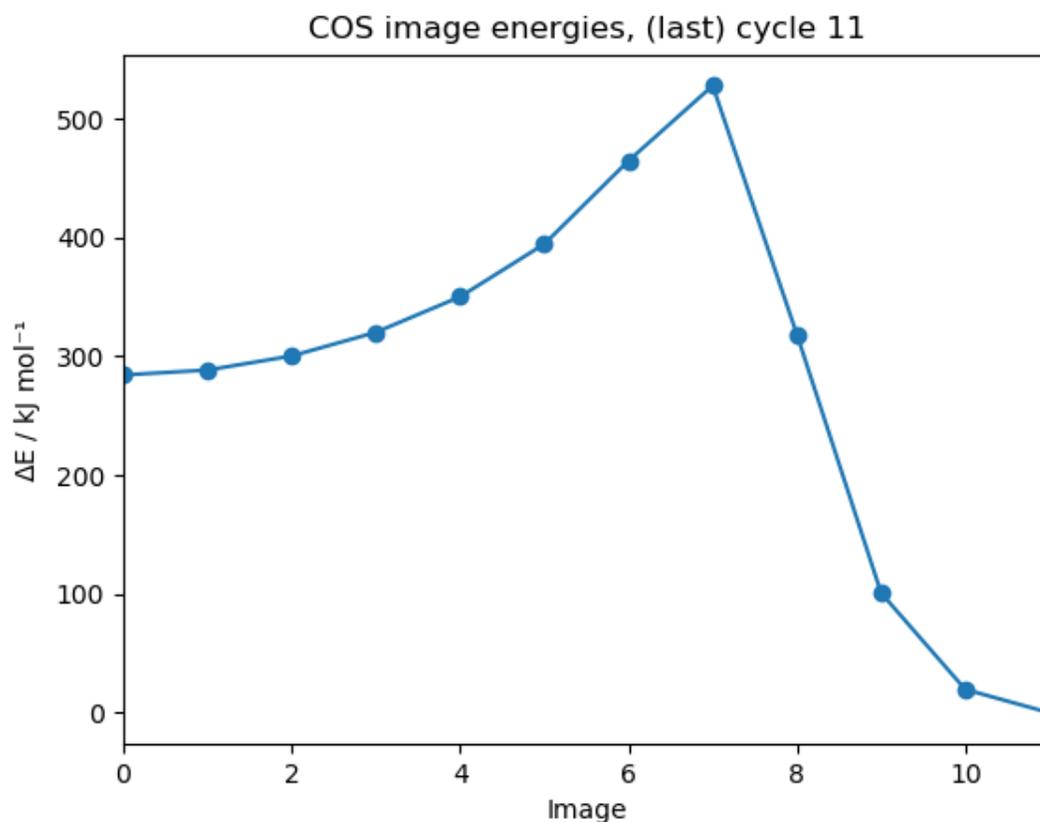


Fig. 14.4: COS image energies of the last (most recent) optimization cycle.

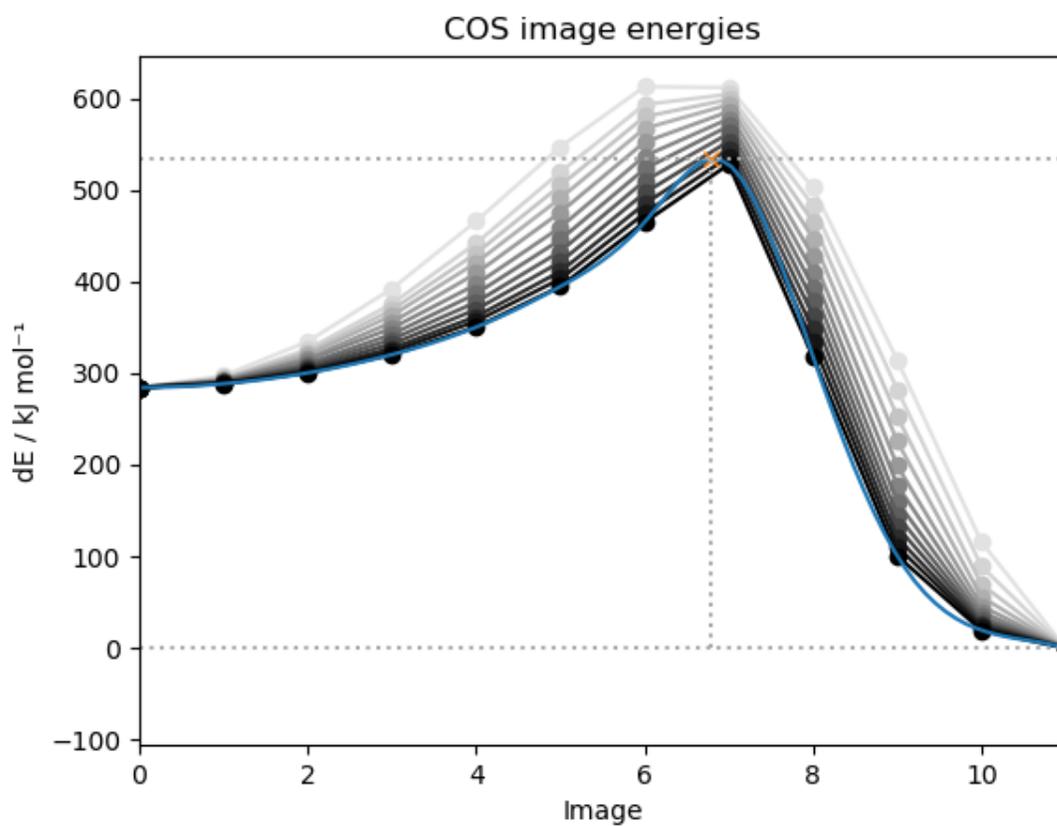


Fig. 14.5: COS image energies of all optimization cycles. Note that the actual difference between image geometries are not taken into account. Equidistance is assumed. Later (more recent) cycles are given in a darker shade.

The forces acting on the respective COS images can also be plotted.

```
pysisplot --coforces [--h5_fn optimization.h5] [--h5_group opt]
```

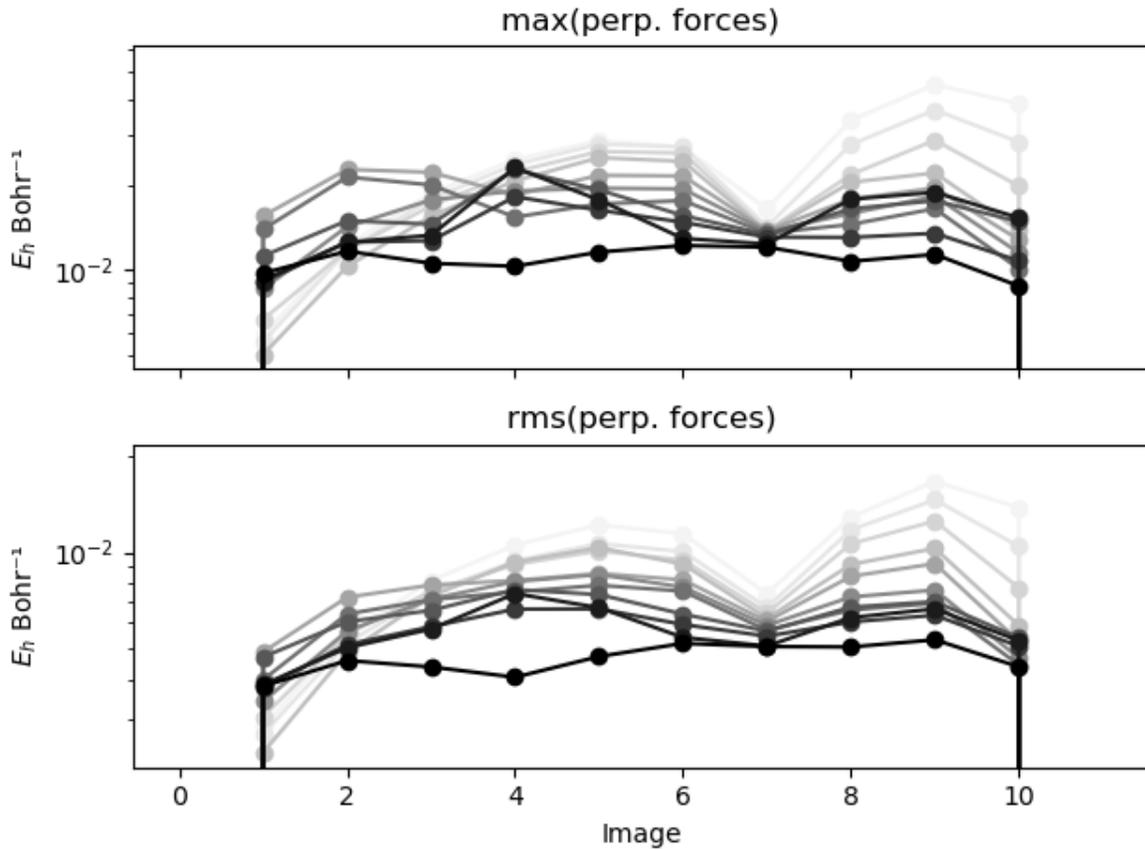


Fig. 14.6: Maximum component and root-mean-square (rms) of the perpendicular component of the force, acting on the COS images.

Please note that nothing is plotted for images 0 and 11, as they remained fixed in the optimization.

### 14.1.3 Plotting TS-optimization progress

The TS-optimization progress is plotted with `pysisplot --opt --h5_group tsopt`. Here we explicitly selected a different HDF5 group by `--h5_group`.

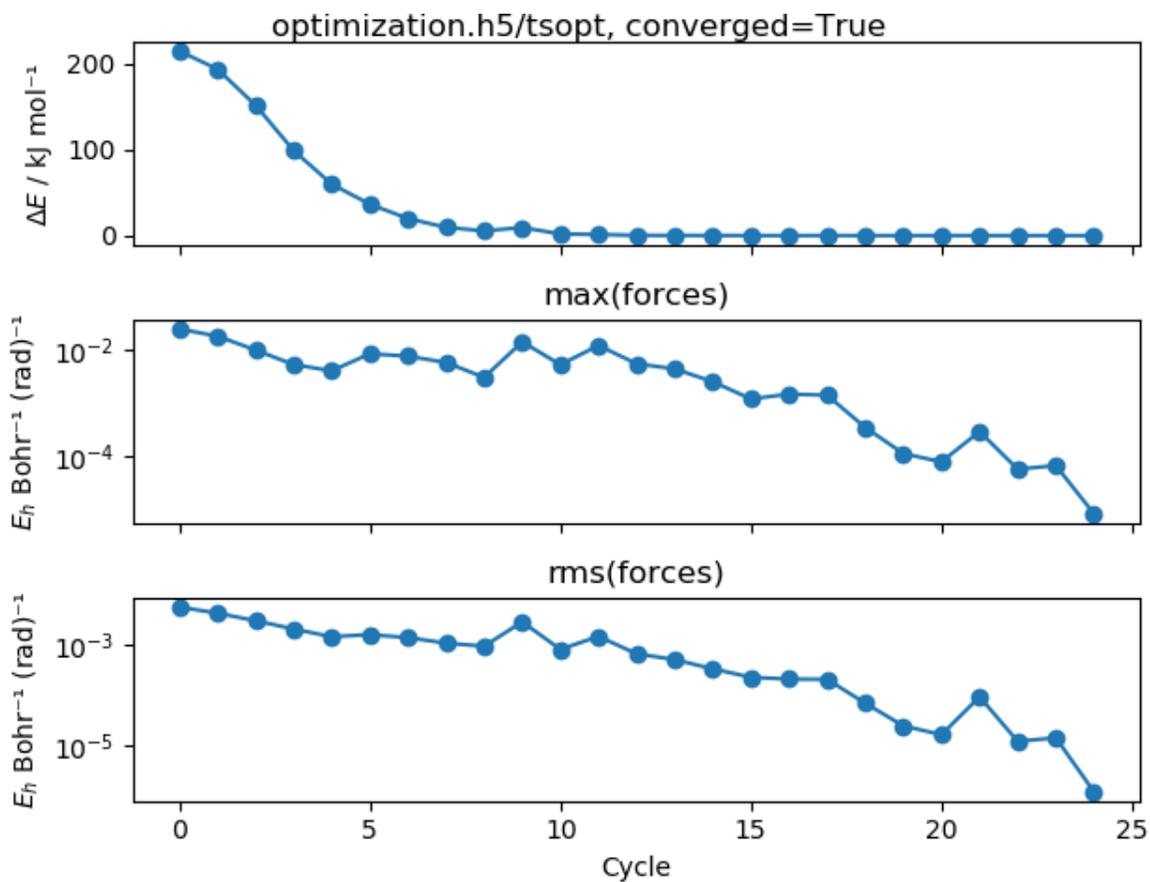


Fig. 14.7: Progress of the TS optimization, started from the HEI.

### 14.1.4 Plotting the Intrinsic Reaction Coordinate

IRC profiles are easily plotted by

```
pysisplot --irc
```

Multiple plots may appear, depending on the progress of the IRC. The IRC coordinate is given in mass-weighted cartesian coordinates, whereas gradients are given in non-mass-weighted units.

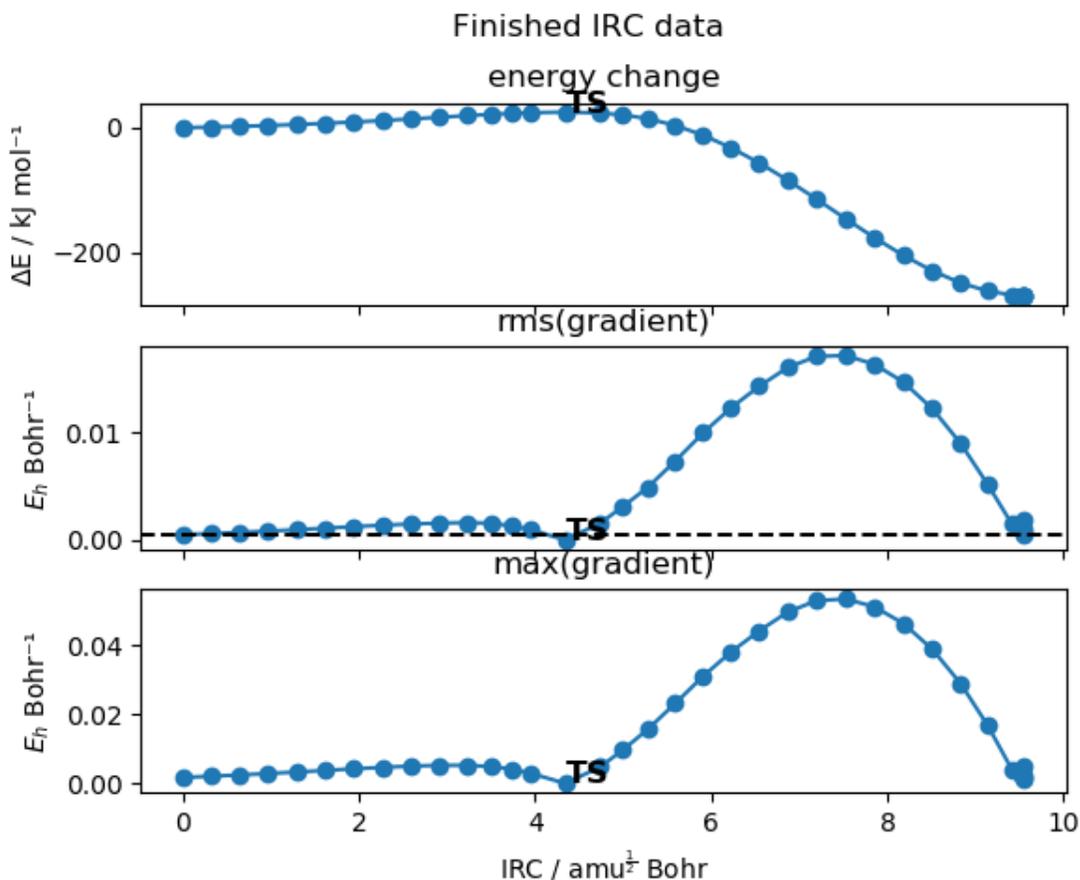
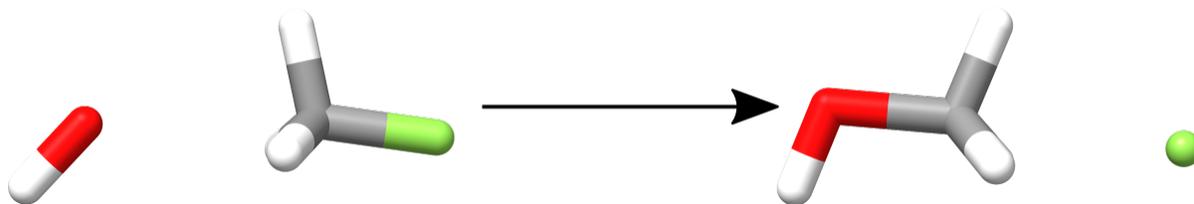
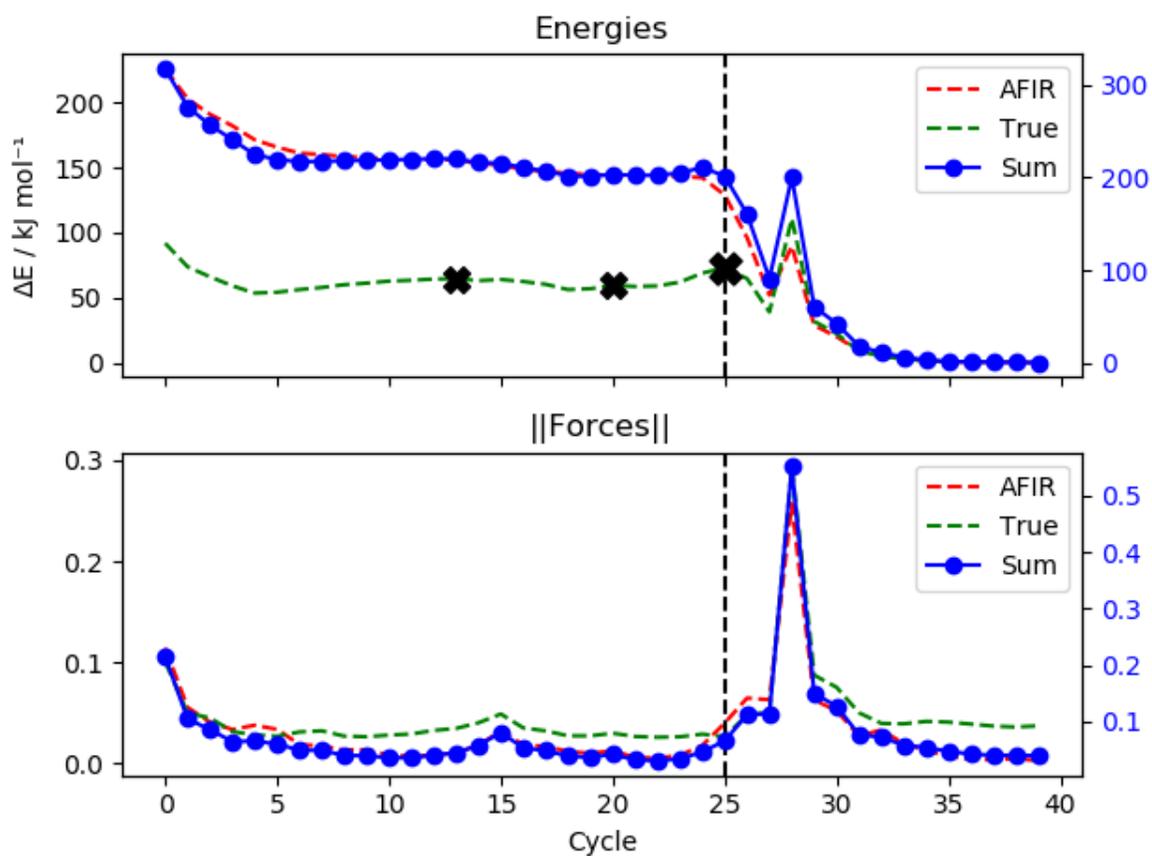


Fig. 14.8: IRC for the Diels-Alder reaction between ethene and 1,4-butadiene.

Evidently the IRC integration failed at the end, as can be seen from the the bunched up points, but unless you want to do some kind of transition-state-theory (TST; not supported by pysisyphus) calculations this should not be a problem.

## 14.2 Example - AFIR

pysisplot is able to visualize AFIR calculations and to highlight interesting geometries along the optimization. Shown below is an example taken from the [AFIR-Paper](#). By using AFIR the  $S_N2$  between  $\text{OH}^-$  and fluoromethylene can be forced, yielding methanol and the fluorine anion. The corresponding unit test can be found in the `tests/test_afir` directory of the repository.

Fig. 14.9: Formation of methanol by means of a  $S_N2$  reaction.Fig. 14.10: Energy profile and force norms along the  $S_N2$  reaction.

## 14.3 Example - Excited State Tracking

pysisyphus is aware of excited states (ES) and can track them using various approaches over the course of an optimization or an IRC. By calculating the overlap matrices between ESs at given geometry and a reference geometry pysisyphus can track the desired ES. All relevant data is stored in *overlap\_data.h5*.

Optimizing an ES is demonstrated for the  $S_1$  of the 1H-amino-keto tautomer of Cytosin at the PBE0/def-SVP level of theory. A corresponding test can be found under (*tests/test\_cytosin\_opt*). Right after the first optimization cycle a root flip occurs and the  $S_1$  and  $S_2$  switch. The potential energy curves along the optimization are plotted by:

```
pysisplot --all_energies
pysisplot -a
```

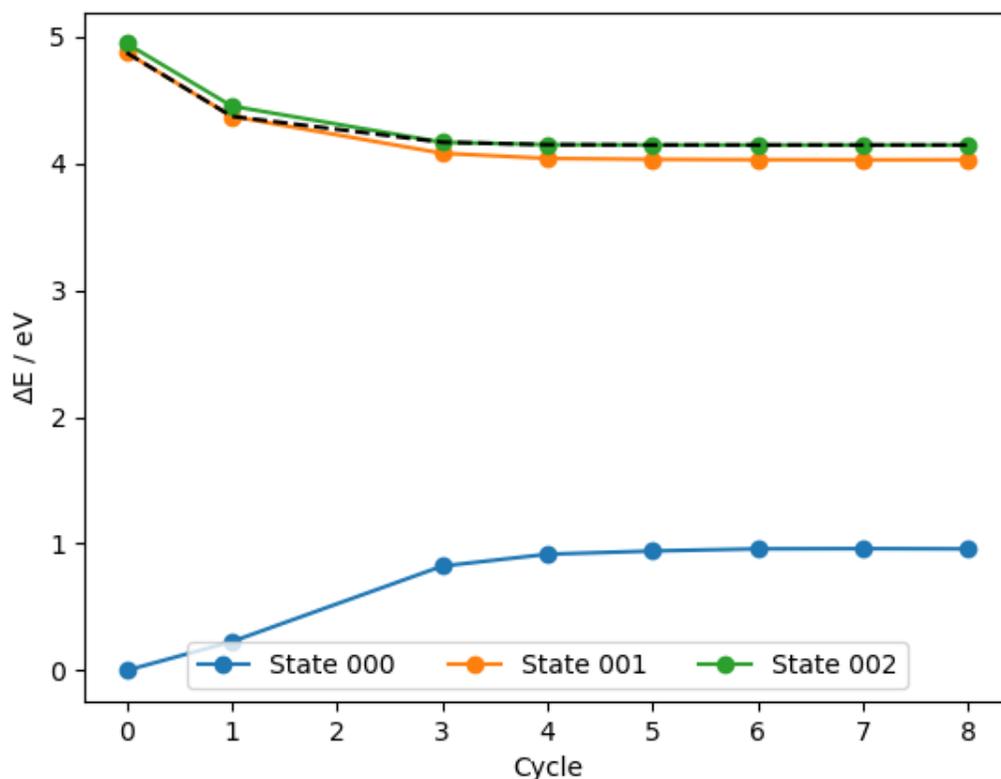


Fig. 14.11: Potential energy curves along the  $S_1$  optimization of the 1H-amino-keto tautomer of Cytosin at the PBE0/def2-SVP level of theory. The root actually followed is indicated by a dashed line.

The calculated overlap matrices can be plotted by:

```
pysisplot --overlaps
pysisplot -o
```

If the calculation was set up to calculate charge-density-differences (CDDs) via MultiWFN and to render them via Jmol then the CDD images displayed beside the overlap matrices.

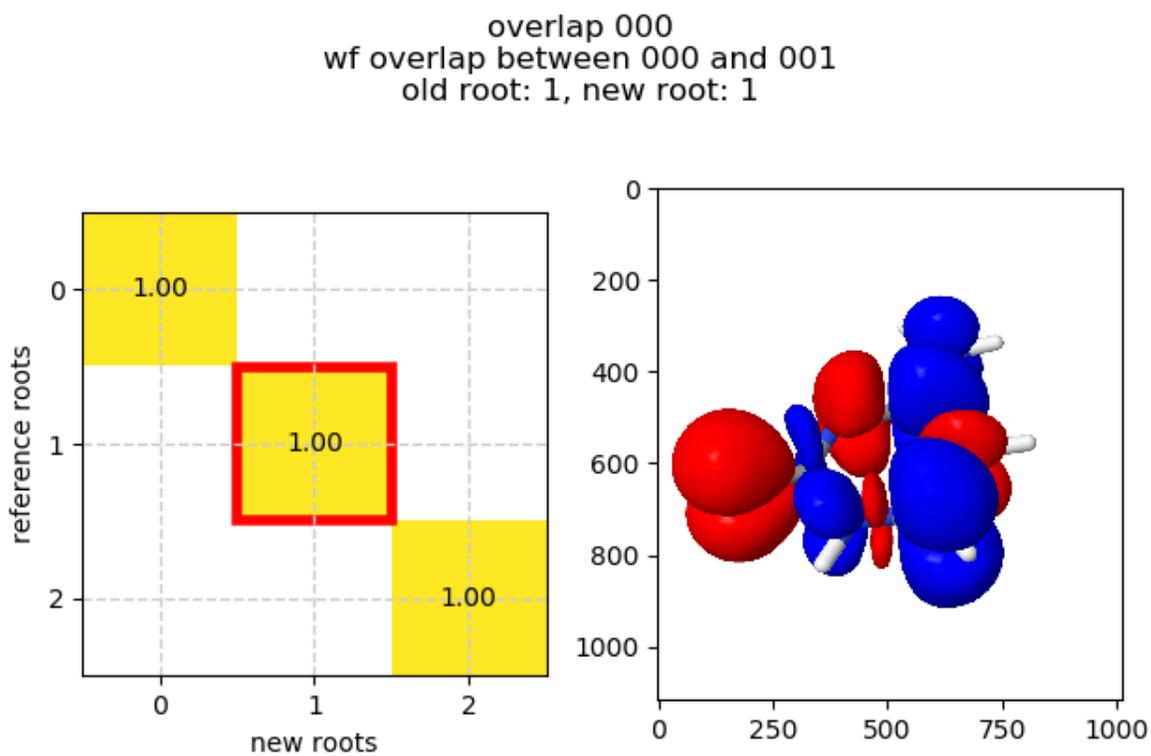


Fig. 14.12: Cytosin  $S_1$  optimization. Overlaps between first and second cycle. No root flips occurred. All ES are in the same order as in the reference geometry at cycle 0.

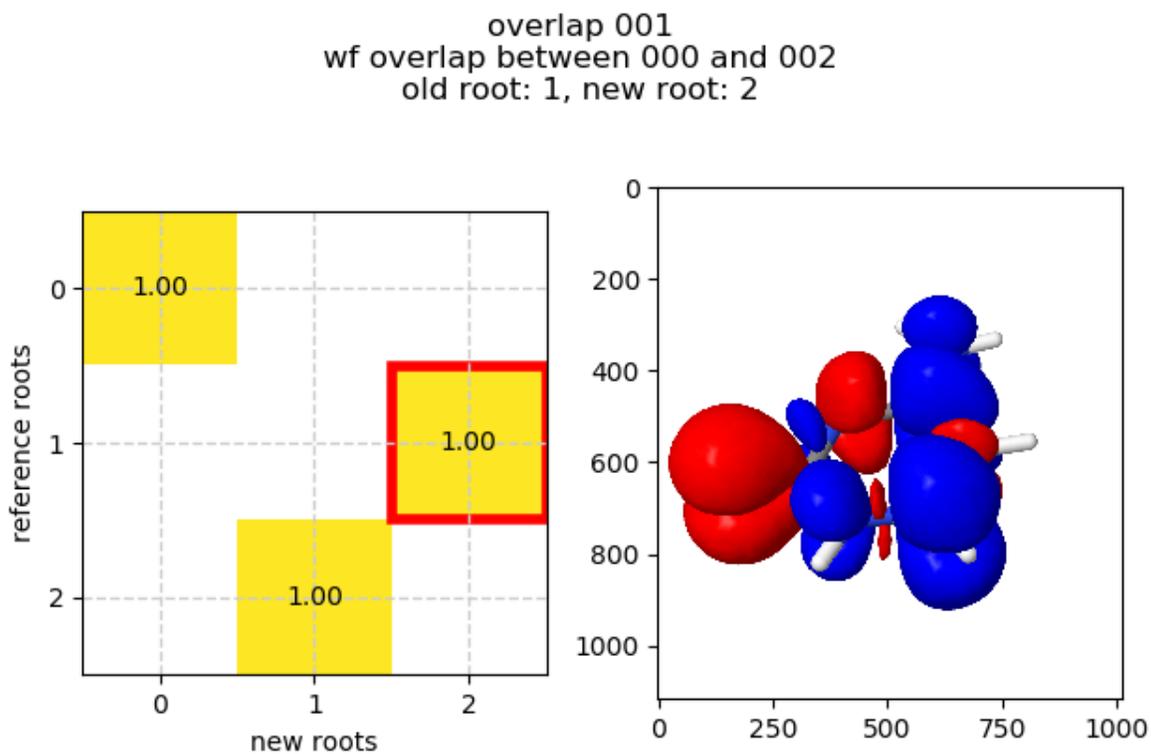


Fig. 14.13: Cytosin  $S_1$  optimization. Overlaps between second and third cycle with root flip. The  $S_1$  and  $S_2$  switch their order.

## WORKING WITH GEOMETRIES

### TBD

Please see *pysistrj --help* for a list of possible invocations.

```
usage: Utility to transform .xyz and .trj files. [-h]
        (--between BETWEEN | --align | --split
↪ | --reverse | --cleantrj | --spline | --first FIRST | --every EVERY | --center | --
↪ centerm | --translate TRANSLATE TRANSLATE TRANSLATE | --append | --join | --match | --
↪ std | --shake | --internals | --get GET | --origin | --fragsort)
        [--scale SCALE] [--seed SEED]
        [--idpp | --lst | --redund]
        [--noipalign] [--bohr]
        [--noxyz]
        [--atoms ATOMS [ATOMS ...]]
        [--add_prims ADD_PRIMS]
        fns [fns ...]
```

#### positional arguments:

fns                    Filenames of .xyz and/or .trj files (xyz and trj can be mixed).

#### optional arguments:

-h, --help            show this help message and exit  
--between BETWEEN    Interpolate additional images.  
--align                Align geometries onto the first geometry.  
--split                Split a supplied geometries in multiple .xyz files.  
--reverse              Reverse a .trj file.  
--cleantrj             Keep only the first four columns of xyz/trj files.  
--spline                Evenly redistribute geometries along a splined path.  
--first FIRST          Copy the first N geometries to a new .trj file.  
--every EVERY          Create new .trj with every N-th geometry. Always includes the first and last point.  
--center                Move the molecules centroid into the origin.  
--centerm              Move the molecules center of mass into the origin.  
--translate TRANSLATE TRANSLATE TRANSLATE    Translate the molecule by the given vector given in Ångström.  
--append                Combine the given .xyz files into one .xyz file.  
--join                  Combine the given .xyz/.trj files into one .trj file.  
--match                 Resort the second .xyz file so the atom order matches the first .xyz file. Uses the hungarian method.

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```
--std                Move supplied geometry to its standard orientation.
--shake             Shake (randomly displace) coordinates.
--internals        Print automatically generated internals.
--get GET          Get n-th geometry. Expects 0-based index input.
--origin           Translate geometry, so that min(X/Y/Z) == 0.
--fragsort         Resort atoms by fragments.
--idpp            Interpolate using Image Dependent Pair Potential.
--lst             Interpolate by linear synchronous transit.
--redund          Interpolate in internal coordinates.
--noipalign       Don't align geometries when interpolating.
--bohr            Input geometries are in Bohr instead of Angstrom.
--noxyz           Disable dumping of single .xyz files.
--atoms ATOMS [ATOMS ...]
                  Used with --internals. Only print primitives including
                  the given atoms.
--add_prims ADD_PRIMS
                  Used with --internals. Define additional primitives.
                  Expects a string representation of a nested list that
                  can be parsed as YAML e.g.
                  [[10,30],[1,2,3],[4,5,6,7]].

--scale SCALE      Scales the displacement in --shake.
--seed SEED        Initialize the RNG for reproducible results.
```

## PYSISYPHUS

### 16.1 pysisyphus package

#### 16.1.1 Subpackages

##### 16.1.1.1 pysisyphus.benchmarks package

###### Submodules

###### pysisyphus.benchmarks.Benchmark module

```
class pysisyphus.benchmarks.Benchmark.Benchmark(name, exclude=None, inv_exclude=False,  
                                                only=None, coord_type='cart', calc_getter=None)
```

Bases: object

```
data_funcs = {'baker': <function get_baker_data>, 'baker_ts': <function  
get_baker_ts_data>, 'birkholz_rx': <function get_birkholz_rx_data>,  
'precon_pos_rot': <function get_precon_pos_rot_data>, 's22': <function  
get_s22_data>, 'xtb_rx': <function get_xtb_rx_data>, 'zimmerman': <function  
get_zimmerman_data>, 'zimmerman_xtb': <function get_zimmerman_xtb_data>}
```

property geom\_iter

property geoms

```
get_geoms(id_, set_calculator=True)
```

###### pysisyphus.benchmarks.data module

```
pysisyphus.benchmarks.data.get_baker_data()
```

10.1002/jcc.540140910

HF/STO-3G

```
pysisyphus.benchmarks.data.get_baker_ts_data()
```

10.1002/(SICI)1096-987X(199605)17:7<888::AID-JCC12>3.0.CO;2-7

HF/3-21G

`pysisyphus.benchmarks.data.get_birkholz_rx_data()`

<https://doi.org/10.1002/jcc.23910>

`pysisyphus.benchmarks.data.get_precon_pos_rot_data()`

<https://doi.org/10.1002/jcc.23910>

`pysisyphus.benchmarks.data.get_s22_data()`

<https://doi.org/10.1039/B600027D>

`pysisyphus.benchmarks.data.get_xtb_rx_data()`

`pysisyphus.benchmarks.data.get_zimmerman_data()`

<https://dx.doi.org/10.1021/ct400319w>

`pysisyphus.benchmarks.data.get_zimmerman_xtb_data()`

#### Reoptimization of

<https://dx.doi.org/10.1021/ct400319w>

at the gfn2-xtb level of theory.

Includes set 1 (first 72 entries), minus (0-based) ids

7, 8, 13, 14, 15, 34, 39

## Module contents

### 16.1.1.2 pysisyphus.calculators package

#### Submodules

#### pysisyphus.calculators.AFIR module

```
class pysisyphus.calculators.AFIR.AFIR(calculator, fragment_indices, gamma, rho=1, p=6,
                                       ignore_hydrogen=True, complete_fragments=True, dump=True,
                                       h5_fn='afir.h5', h5_group_name='afir', **kwargs)
```

Bases: *Calculator*

```
__init__(calculator, fragment_indices, gamma, rho=1, p=6, ignore_hydrogen=True,
         complete_fragments=True, dump=True, h5_fn='afir.h5', h5_group_name='afir', **kwargs)
```

Artificial Force Induced Reaction calculator.

Currently, there are no automated drivers to run large-scale AFIR calculations with many different initial orientations and/or increasing collision energy parameter. Nonetheless, selected AFIR calculations can be carried out by hand. After convergence, artificial potential & forces, as well as real energies and forces can be plotted with 'pysisplot --afir'. The highest energy point along the AFIR path can then be selected for a subsequent TS-optimization, e.g. via 'pysistrj --get [index] optimization.trj'.

Future versions of pysisyphus may provide drivers for more automated AFIR calculations.

#### Parameters

- **calculator** (*Calculator*) -- Actual QC calculator that provides energies and its derivatives, that are modified by the AFIR calculator, e.g., ORCA or Psi4.
- **fragment\_indices** (List[List[int]]) -- List of lists of integers, specifying the separate fragments. If the indices in these lists don't comprise all atoms in the molecule, the remaining indices will be added as a separate fragment. If a AFIR calculation is carried out

with 2 fragments and 'complete\_fragments' is True (see below) it is enough to specify only the indices of one fragment, e.g., for a system of 10 atoms 'fragment\_indices=[[0,1,2,3]]' is enough. The second system will be set up automatically with indices [4,5,6,7,8,9].

- **gamma** (Union[\_SupportsArray[dtype], \_NestedSequence[\_SupportsArray[dtype]], bool, int, float, complex, str, bytes, \_NestedSequence[Union[bool, int, float, complex, str, bytes]]]) -- Collision energy parameter in au. For 2 fragments it can be a single integer, while for > 2 fragments a list of gammas must be given, specifying the pair-wise collision energy parameters. For 3 fragments 3 gammas must be given [\_01, \_02, \_12], for 4 fragments 6 gammas would be required [\_01, \_02, \_03, \_12, \_13, \_23] and so on.
- **rho** (Union[\_SupportsArray[dtype], \_NestedSequence[\_SupportsArray[dtype]], bool, int, float, complex, str, bytes, \_NestedSequence[Union[bool, int, float, complex, str, bytes]]], default: 1) -- Direction of the artificial force, either 1 or -1. The same comments as for gamma apply. For 2 fragments a single integer is enough, for > 2 fragments a list of rhos must be given (see above). For rho=1 fragments are pushed together, for rho=-1 fragments are pulled apart.
- **p** (int, default: 6) -- Exponent p used in the calculation of the weight function . Defaults to 6 and probably does not have to be changed.
- **ignore\_hydrogen** (bool, default: True) -- Whether hydrogens are ignored in the calculation of the artificial force.
- **complete\_fragments** (bool, default: True) -- Whether an incomplete specification in 'fragment\_indices' is automatically completed.
- **dump** (bool, default: True) -- Whether an HDF5 file is created.
- **h5\_fn** (str, default: 'afir.h5') -- Filename of the HDF5 file used for dumping.
- **h5\_group\_name** (str, default: 'afir') -- HDF5 group name used for dumping.
- **\*\*kwargs** -- Keyword arguments passed to the Calculator baseclass.

**afir\_fd\_hessian\_wrapper**(*coords3d, afir\_grad\_func*)

**property charge**

**dump\_h5**(*atoms, coords, results*)

**get\_energy**(*atoms, coords, \*\*prepare\_kwargs*)

Meant to be extended.

**get\_forces**(*atoms, coords, \*\*prepare\_kwargs*)

Meant to be extended.

**get\_hessian**(*atoms, coords, \*\*prepare\_kwargs*)

Meant to be extended.

**init\_h5\_group**(*atoms, max\_cycles=None*)

**log\_fragments**()

**property mult**

**set\_atoms\_and\_funcs**(*atoms, coords*)

Initially atoms was also an argument to the constructor of AFIR. I removed it so creation becomes easier. The first time a calculation is requested with a proper atom set everything is set up (cov. radii, afir function

and corresponding gradient). Afterwards there is only a check if atoms != None and it is expected that all functions are properly set.

fragment\_indices can also be incomplete w.r.t. to the number of atoms. If the sum of the specified fragment atoms is less than the number of atoms present then all remaining unspecified atoms will be gathered in one fragment.

**write\_fragment\_geoms**(atoms, coords)

**class** pysisyphus.calculators.AFIR.**AFIRPath**(atoms, cart\_coords, energies, forces, opt\_is\_converged)

Bases: object

**atoms:** tuple

**cart\_coords:** ndarray

**energies:** ndarray

**forces:** ndarray

**opt\_is\_converged:** bool

pysisyphus.calculators.AFIR.**afir\_closure**(fragment\_indices, cov\_radii, gamma, rho=1, p=6,  
prefactor=1.0, logger=None)

rho=1 pushes fragments together, rho=-1 pulls fragments apart.

pysisyphus.calculators.AFIR.**get\_data\_model**(atoms, max\_cycles)

### pysisyphus.calculators.AnaPot module

**class** pysisyphus.calculators.AnaPot.**AnaPot**(\*\*kwargs)

Bases: [AnaPotBase](#)

### pysisyphus.calculators.AnaPot2 module

**class** pysisyphus.calculators.AnaPot2.**AnaPot2**

Bases: [AnaPotBase](#)

We can't use sympify as it replaces 1/tan by cot and this isn't supported by numpy when we call lambdify.

**class** pysisyphus.calculators.AnaPot2.**AnaPot2\_**

Bases: [Calculator](#)

**get\_energy**(atoms, coords)

Meant to be extended.

**pysisyphus.calculators.AnaPot3 module****class** pysisyphus.calculators.AnaPot3.**AnaPot3**Bases: *AnaPotBase***pysisyphus.calculators.AnaPot4 module****class** pysisyphus.calculators.AnaPot4.**AnaPot4**Bases: *AnaPotBase***pysisyphus.calculators.AnaPotBase module****class** pysisyphus.calculators.AnaPotBase.**AnaPotBase**(*V\_str, scale=1.0, xlim=(- 1, 1), ylim=(- 1, 1), levels=None, use\_sympify=True, minima=None, saddles=None*)Bases: *Calculator***anim\_coords**(*coords, interval=50, show=False, title\_func=None*)**anim\_opt**(*opt, energy\_profile=False, colorbar=False, figsize=(8, 6), show=False*)**get\_energy**(*atoms, coords*)

Meant to be extended.

**get\_forces**(*atoms, coords*)

Meant to be extended.

**classmethod** **get\_geom**(*coords, atoms=('X'), V\_str=None, calc\_kwargs=None, geom\_kwargs=None*)**get\_geoms\_from\_stored\_coords**(*coords\_list, i=None, calc\_kwargs=None, geom\_kwargs=None*)**get\_hessian**(*atoms, coords*)

Meant to be extended.

**get\_minima**(*i=None, calc\_kwargs=None, geom\_kwargs=None*)**get\_path**(*num, minima\_inds=None*)**get\_saddles**(*i=None, calc\_kwargs=None, geom\_kwargs=None*)**plot**(*levels=None, show=False, \*\*figkwargs*)**plot3d**(*levels=None, show=False, zlim=None, vmin=None, vmax=None, resolution=100, rcount=50, ccount=50, nan\_above=None, init\_view=None, colorbar=False, \*\*figkwargs*)**plot\_coords**(*xs, ys, enum=True, show=False, title=None*)**plot\_eigenvalue\_structure**(*grid=50, levels=None, show=False*)**plot\_geoms**(*geoms, \*\*kwargs*)**plot\_irc**(*irc, \*args, \*\*kwargs*)**plot\_opt**(*opt, \*args, \*\*kwargs*)**statistics**()

### pysisyphus.calculators.AnaPotCBM module

**class** pysisyphus.calculators.AnaPotCBM.**AnaPotCBM**

Bases: *AnaPotBase*

### pysisyphus.calculators.AtomAtomTransTorque module

**class** pysisyphus.calculators.AtomAtomTransTorque.**AtomAtomTransTorque**(*geom, frags, A\_mats, kappa=2.0*)

Bases: object

**\_\_init\_\_**(*geom, frags, A\_mats, kappa=2.0*)

Atom-atom translational and torque forces.

See A.5. [1], Eq. (A6).

**get\_forces**(*atoms, coords*)

### pysisyphus.calculators.Calculator module

**class** pysisyphus.calculators.Calculator.**Calculator**(*calc\_number=0, charge=0, mult=1, base\_name='calculator', pal=1, mem=1000, check\_mem=True, retry\_calc=1, last\_calc\_cycle=None, clean\_after=True, out\_dir='qm\_calcs'*)

Bases: object

**\_\_init\_\_**(*calc\_number=0, charge=0, mult=1, base\_name='calculator', pal=1, mem=1000, check\_mem=True, retry\_calc=1, last\_calc\_cycle=None, clean\_after=True, out\_dir='qm\_calcs'*)

Base-class of all calculators.

Meant to be extended.

#### Parameters

- **calc\_number** (*int, default=0*) -- Identifier of the Calculator. Used in distinguishing it from other Calculators, e.g. in ChainOfStates calculations. Also used in the creation of filenames.
- **charge** (*int, default=0*) -- Molecular charge.
- **mult** (*int, default=1*) -- Molecular multiplicity (1 = singlet, 2 = doublet, ...)
- **base\_name** (*str, default=calculator*) -- Generated filenames will start with this string.
- **pal** (*int, default=1*) -- Positive integer that gives the number of physical cores to use on 1 node.
- **mem** (*int, default=1000*) -- Memory per core in MB. The total amount of memory is given as  $mem \cdot pal$ .
- **check\_mem** (*bool, default=True*) -- Whether to adjust the requested memory if too much is requested.
- **retry\_calc** (*int, default=0*) -- Number of additional retries when calculation failed.
- **last\_calc\_cycle** (*int*) -- Internal variable used in restarts.

- **clean\_after** (*bool*) -- Delete temporary directory were calculations were executed after a calculation.
- **out\_dir** (*str*) -- Path that is prepended to generated filenames.

**clean**(*path*)

Delete the temporary directory.

**Parameters**

**path** (*Path*) -- Directory to delete.

**conf\_key** = `None`

**get\_cmd**(*key='cmd'*)

**get\_energy**(*atoms, coords*)

Meant to be extended.

**get\_forces**(*atoms, coords*)

Meant to be extended.

**get\_hessian**(*atoms, coords*)

Meant to be extended.

**get\_restart\_info**()

Return a dict containing chkfiles.

**Returns**

**restart\_info** -- Dictionary holding the calculator state. Used for restoring calculators in restarted calculations.

**Return type**

dict

**keep**(*path*)

Backup calculation results.

**Parameters**

**path** (*Path*) -- Temporary directory of the calculation.

**Returns**

**kept\_fns** -- Dictionary holding the filenames that were backed up. The keys correspond to the type of file.

**Return type**

dict

**log**(*message=""*)

Write a log message.

Wraps the logger variable.

**Parameters**

**message** (*str*) -- Message to be logged.

**make\_fn**(*name, counter=None, return\_str=False*)

Make a full filename.

Return a full filename including the calculator name and the current counter given a suffix.

**Parameters**

- **name** (*str*) -- Suffix of the filename.

- **counter** (*int*, *optional*) -- If not given use the current `calc_counter`.
- **return\_str** (*int*, *optional*) -- Return a string instead of a `Path` when `True`.

**Returns**

**fn** -- Filename.

**Return type**

`str`

**property name****prepare**(*inp*)

Prepare a temporary directory and write input.

Similar to `prepare_path`, but the input is also written into the prepared directory.

**inp**

[`str`] Input to be written into the file `self.inp_fn` in the prepared directory.

**Returns**

**path** -- Prepared directory.

**Return type**

`Path`

**prepare\_coords**(*atoms*, *coords*)

Get 3d coords in Angstrom.

Reshape internal 1d coords to 3d and convert to Angstrom.

**Parameters**

- **atoms** (*iterable*) -- Atom descriptors (element symbols).
- **coords** (*np.array*, *1d*) -- 1D-array holding coordinates in Bohr.

**Returns**

**coords** -- 3D-array holding coordinates in Angstrom.

**Return type**

`np.array`, 3d

**prepare\_input**(*atoms*, *coords*, *calc\_type*)

Meant to be extended.

**prepare\_path**(*use\_in\_run=False*)

Get a temporary directory handle.

Create a temporary directory that can later be used in a calculation.

**Parameters**

**use\_in\_run** (*bool*, *option*) -- Sets the internal variable `self.path_already_prepared` that is later read by `self.run()`. No new temporary directory will be created in `self.run()`.

**Returns**

**path** -- Prepared directory.

**Return type**

`Path`

**prepare\_pattern**(*raw\_pat*)

Prepare globs.

Transforms an entry of `self.to_keep` into a glob and a key suitable for the use in `self.keep()`.

**Parameters**

**raw\_pat** (*str*) -- Entry of `self.to_keep`

**Returns**

- **pattern** (*str*) -- Glob that can be used in `Path.glob()`
- **multi** (*bool*) -- Flag if glob may match multiple files.
- **key** (*str*) -- A key to be used in the `kept_fns` dict.

**prepare\_turbo\_coords**(*atoms, coords*)

Get a Turbomole coords string.

**Parameters**

- **atoms** (*iterable*) -- Atom descriptors (element symbols).
- **coords** (*np.array, 1d*) -- 1D-array holding coordinates in Bohr.

**Returns**

**coords** -- String holding coordinates in Turbomole coords format.

**Return type**

str

**prepare\_xyz\_string**(*atoms, coords*)

Returns a xyz string in Angstrom.

**Parameters**

- **atoms** (*iterable*) -- Atom descriptors (element symbols).
- **coords** (*np.array, 1d*) -- 1D-array holding coordinates in Bohr.

**Returns**

**xyz\_str** -- Coordinates in .xyz format.

**Return type**

string

**print\_capabilities**()

**print\_out\_fn**(*path*)

Print calculation output.

Prints the output of a calculator after a calculation.

**Parameters**

**path** (*Path*) -- Temporary directory of the calculation.

**run**(*inp, calc, add\_args=None, env=None, shell=False, hold=False, keep=True, cmd=None, inc\_counter=True, run\_after=True, parser\_kwargs=None, symlink=True*)

Run a calculation.

The bread-and-butter method to actually run an external quantum chemistry code.

**Parameters**

- **inp** (*str*) -- Input for the external program that is written to the temp-dir.

- **calc** (*str*, *hashable*) -- Key (and more or less type of calculation) to select the right parsing function from `self.parser_funcs`.
- **add\_args** (*iterable*, *optional*) -- Additional arguments that will be appended to the program call.
- **env** (*Environment*, *optional*) -- A potentially modified environment for the subprocess call.
- **shell** (*bool*, *optional*) -- Use a shell to execute the program call. Need for Turbomole were we chain program calls like `dscf`; `escf`.
- **hold** (*bool*, *optional*) -- Wether to remove the temporary directory after the calculation.
- **keep** (*bool*, *optional*) -- Wether to backup files as specified in `self.to_keep()`. Usually you want this.
- **cmd** (*str or iterable*, *optional*) -- Overwrites `self.base_cmd`.
- **inc\_counter** (*bool*, *optional*) -- Wether to increment the counter after a calculation.

**Returns**

**results** -- Dictionary holding all applicable results of the calculations like the energy, a forces vector and/or excited state energies from TDDFT.

**Return type**

dict

**run\_after**(*path*)

Meant to be extended.

This method is called after a calculation was done, but before entering `self.keep()` and `self.clean()`. Can be used to call tools like `formchk` or `ricctools`.

**set\_restart\_info**(*restart\_info*)

Sets restart information (`chkfiles` etc.) on the calculator.

**Parameters**

**restart\_info** (*dict*) -- Dictionary holding the calculator state. Used for restoring calculators in restarted calculations.

**verify\_chkfiles**(*chkfiles*)

Checks if given `chkfiles` exist and return them as Paths

**Parameters**

**chkfiles** (*dict*) -- Dictionary holding the `chkfiles`. The keys correspond to the attribute names, the values are str's holding the (potentially full) filename (path).

**Returns**

**paths** -- Dictionary of Paths.

**Return type**

dict

**pysisyphus.calculators.CerjanMiller module**

```
class pysisyphus.calculators.CerjanMiller.CerjanMiller(a=1, b=1, c=1)
```

Bases: *AnaPotBase*

**pysisyphus.calculators.Composite module**

```
class pysisyphus.calculators.Composite.Composite(final, keys_calcs=None, calcs=None,
                                                remove_translation=False, **kwargs)
```

Bases: *Calculator*

```
get_energy(atoms, coords, **prepare_kwargs)
```

Meant to be extended.

```
get_forces(atoms, coords, **prepare_kwargs)
```

Meant to be extended.

```
run_calculation(atoms, coords, **prepare_kwargs)
```

**pysisyphus.calculators.ConicalIntersection module**

```
class pysisyphus.calculators.ConicalIntersection.CIQuantities(energy1, gradient1, energy2,
                                                              gradient2, energy_diff,
                                                              gradient_diff, gradient_mean, P, x,
                                                              y, energy, forces)
```

Bases: object

**P:** ndarray

**energy:** float

**energy1:** float

**energy2:** float

**energy\_diff:** float

**forces:** ndarray

**gradient1:** ndarray

**gradient2:** ndarray

**gradient\_diff:** ndarray

**gradient\_mean:** ndarray

**x:** ndarray

**y:** ndarray

```
class pysisyphus.calculators.ConicalIntersection.ConicalIntersection(calculator1, calculator2,  
                                                                **kwargs)
```

Bases: *Calculator*

Calculator for conical intersection optimization.

Based on [1].

```
get_ci_quantities(atoms, coords, **prepare_kwargs)
```

Relevant quantities including branching plane and projector P.

```
get_energy(atoms, coords, **prepare_kwargs)
```

Energy of calculator 1.

```
get_forces(atoms, coords, **prepare_kwargs)
```

Projected gradient for CI optimization.

```
get_hessian(atoms, coords, **prepare_kwargs)
```

Projected Hessian.

```
pysisyphus.calculators.ConicalIntersection.get_P(x, y)
```

Projector to project out components in branching plane.

```
pysisyphus.calculators.ConicalIntersection.update_y(x, x_prev, y_prev)
```

Update approximate coupling derivative vector y.

### pysisyphus.calculators.DFTBp module

```
class pysisyphus.calculators.DFTBp.DFTBp(parameter, *args, slakos=None, root=None, **kwargs)
```

Bases: *OverlapCalculator*

```
conf_key = 'dftbp'
```

```
get_energy(atoms, coords, **prepare_kwargs)
```

Meant to be extended.

```
static get_excited_state_str(root, forces=False)
```

```
get_forces(atoms, coords, **prepare_kwargs)
```

Meant to be extended.

```
static get_gen_str(atoms, coords)
```

```
max_ang_moms = {'mio-ext': {'C': 'p', 'H': 's', 'N': 'p', 'O': 'p'}}
```

```
parse_energy(path)
```

```
static parse_exc_dat(text)
```

```
parse_forces(path)
```

```
parse_total_energy(text)
```

```
prepare_input(atoms, coords, calc_type)
```

Meant to be extended.

**prepare\_overlap\_data**(*path*)

Implement calculator specific parsing of MO coefficients and CI coefficients here. Should return a filename pointing to TURBOMOLE like mos, a MO coefficient array and a CI coefficient array.

**run\_calculation**(*atoms, coords, \*\*prepare\_kwargs*)

`pysisyphus.calculators.DFTBp.parse_mo`(*eigvec*)

`pysisyphus.calculators.DFTBp.parse_xplusy`(*text*)

## pysisyphus.calculators.Dalton module

**class** `pysisyphus.calculators.Dalton.Dalton`(*basis, method='hf', \*\*kwargs*)

Bases: `Calculator`

**compute**(*mol, prop*)

**conf\_key** = 'dalton'

**get\_energy**(*atoms, coords, \*\*prepare\_kwargs*)

Meant to be extended.

**get\_forces**(*atoms, coords, \*\*prepare\_kwargs*)

Meant to be extended.

**prepare\_input**(*atoms, coords*)

Meant to be extended.

## pysisyphus.calculators.Dimer module

**class** `pysisyphus.calculators.Dimer.Dimer`(*calculator, \*args, N\_raw=None, length=0.0189, rotation\_max\_cycles=15, rotation\_method='fourier', rotation\_thresh=0.0001, rotation\_tol=1, rotation\_max\_element=0.001, rotation\_interpolate=True, rotation\_disable=False, rotation\_disable\_pos\_curv=True, rotation\_remove\_trans=True, trans\_force\_f\_perp=True, bonds=None, N\_hessian=None, bias\_rotation=False, bias\_translation=False, bias\_gaussian\_dot=0.1, seed=None, write\_orientations=True, forward\_hessian=True, \*\*kwargs*)

Bases: `Calculator`

**property** `C`

Shortcut for the curvature.

**property** `N`

**add\_gaussian**(*atoms, center, N, height=0.1, std=0.0529, max\_cycles=50, dot\_ref=None*)

**property** `can_bias_f0`

**property** `can_bias_f1`

**property** `coords0`

**property coords1**

**curvature**(*f1, f2, N*)

Curvature of the mode represented by the dimer.

**direct\_rotation**(*optimizer, prev\_step*)

**do\_dimer\_rotations**(*rotation\_thresh=None*)

**property energy0**

**property f0**

**property f1**

**property f1\_bias**

**property f2**

Never calculated explicitly, but estimated from f0 and f1.

**fourier\_rotation**(*optimizer, prev\_step*)

**get\_N\_raw\_from\_hessian**(*h5\_fn, root=0*)

**get\_forces**(*atoms, coords*)

Meant to be extended.

**get\_gaussian\_energies**(*coords, sum\_=True*)

**get\_gaussian\_forces**(*coords, sum\_=True*)

**get\_hessian**(*atoms, coords*)

Meant to be extended.

**remove\_translation**(*displacement*)

**property rot\_force**

**rotate\_coords1**(*rad, theta*)

Rotate dimer and produce new coords1.

**set\_N\_raw**(*coords*)

**property should\_bias\_f0**

May lead to calculation of f0 and/or f1 if present!

**property should\_bias\_f1**

May lead to calculation of f0 and/or f1 if present!

**update\_orientation**(*coords*)

**class** pysisyphus.calculators.Dimer.**Gaussian**(*height, center, std, N*)

Bases: object

**energy**(*R, height=None*)

**forces**(*R, height=None*)

**exception** pysisyphus.calculators.Dimer.**RotationConverged**

Bases: Exception

## pysisyphus.calculators.Dummy module

```
class pysisyphus.calculators.Dummy.Dummy(calc_number=0, charge=0, mult=1, base_name='calculator',
                                           pal=1, mem=1000, check_mem=True, retry_calc=1,
                                           last_calc_cycle=None, clean_after=True, out_dir='qm_calcs')
```

Bases: *Calculator*

**get\_energy**(\*args, \*\*kwargs)

Meant to be extended.

**get\_forces**(\*args, \*\*kwargs)

Meant to be extended.

**get\_hessian**(\*args, \*\*kwargs)

Meant to be extended.

**raise\_exception**()

**run\_calculation**(\*args, \*\*kwargs)

## pysisyphus.calculators.EGO module

```
class pysisyphus.calculators.EGO.EGO(calculator, ref_geom, max_force=0.175, **kwargs)
```

Bases: *Calculator*

**get\_energy**(*atoms, coords, \*\*prepare\_kwargs*)

Meant to be extended.

**get\_forces**(*atoms, coords, \*\*prepare\_kwargs*)

Meant to be extended.

**get\_mods**(*atoms, coords*)

**property ref\_hessian**

**property s**

## pysisyphus.calculators.EnergyMin module

```
class pysisyphus.calculators.EnergyMin.EnergyMin(calculator1, calculator2, mix=False, alpha=0.02,
                                                  sigma=3.5, min_energy_diff=0.0, check_after=0,
                                                  **kwargs)
```

Bases: *Calculator*

```
__init__(calculator1, calculator2, mix=False, alpha=0.02, sigma=3.5, min_energy_diff=0.0,
         check_after=0, **kwargs)
```

Use energy and derivatives of the calculator with lower energy.

This calculators carries out two calculations with different settings and returns the results of the lower energy one. This can be used to consider flips between a singlet and a triplet PES etc.

### Parameters

- **calculator1** (*Calculator*) -- Wrapped QC calculator that provides energies and its derivatives.

- **calculator2** (*Calculator*) -- Wrapped QC calculator that provides energies and its derivatives.
- **mix** (bool, default: False) -- Enable mixing of both forces, according to the approach outlined in [2]. Can be used to optimize guesses for MECPs. Pass
- **alpha** (float, default: 0.02) -- Smoothing parameter in Hartree. See [2] for a discussion.
- **sigma** (float, default: 3.5) -- Unitless gap size parameter. The final gap becomes smaller for bigger sigmas. Has to be adapted for each case. See [2] for a discussion (p. 407 right column and p. 408 left column.)
- **min\_energy\_diff** (float, default: 0.0) -- Energy difference in Hartree. When set to a value != 0 and the energy difference between both calculators drops below this value, execution of both calculations is disabled for 'check\_after' cycles. In these cycles the calculator choice remains fixed. After 'check\_after' cycles, both energies will be calculated and it is checked, if the previous calculator choice remains valid. In conjunction with 'check\_after' both arguments can be used to save computational resources.
- **check\_after** (int, default: 0) -- Amount of cycles in which the calculator choice remains fixed.
- **\*\*kwargs** -- Keyword arguments passed to the Calculator baseclass.

**do\_calculations**(*name, atoms, coords, \*\*prepare\_kwargs*)

**get\_chkfiles**()

**Return type**  
dict

**get\_energy**(*atoms, coords, \*\*prepare\_kwargs*)

Meant to be extended.

**get\_forces**(*atoms, coords, \*\*prepare\_kwargs*)

Meant to be extended.

**get\_hessian**(*atoms, coords, \*\*prepare\_kwargs*)

Meant to be extended.

**set\_chkfiles**(*chkfiles*)

## pysisyphus.calculators.ExternalPotential module

```
class pysisyphus.calculators.ExternalPotential.ExternalPotential(calculator=None,
                                                                potentials=None, geom=None,
                                                                **kwargs)
```

Bases: *Calculator*

```
available_potentials = {'harmonic_sphere': <class
'pysisyphus.calculators.ExternalPotential.HarmonicSphere'>, 'logfermi': <class
'pysisyphus.calculators.ExternalPotential.LogFermi'>, 'restraint': <class
'pysisyphus.calculators.ExternalPotential.Restraint'>}
```

**get\_energy**(*atoms, coords*)

Meant to be extended.

**get\_forces**(*atoms*, *coords*)

Meant to be extended.

**get\_hessian**(*atoms*, *coords*)

Meant to be extended.

**get\_potential\_energy**(*coords*)

**get\_potential\_forces**(*coords*)

**class** pysisyphus.calculators.ExternalPotential.**HarmonicSphere**(*k*, *radius*, *origin*=(0.0, 0.0, 0.0),  
*geom*=None)

Bases: object

**calc**(*coords3d*, *gradient*=False)

**instant\_pressure**(*coords3d*)

**property surface\_area**

In Bohr\*\*2

**class** pysisyphus.calculators.ExternalPotential.**LogFermi**(*beta*, *radius*, *T*=300, *origin*=(0.0, 0.0, 0.0),  
*geom*=None)

Bases: object

**\_\_init\_\_**(*beta*, *radius*, *T*=300, *origin*=(0.0, 0.0, 0.0), *geom*=None)

As described in the XTB docs.

<https://xtb-docs.readthedocs.io/en/latest/xcontrol.html#confining-in-a-cavity>

**calc**(*coords3d*, *gradient*=False)

**class** pysisyphus.calculators.ExternalPotential.**Restraint**(*restraints*, *geom*=None)

Bases: object

**calc**(*coords3d*, *gradient*=False)

**static calc\_prim\_restraint**(*prim*, *coords3d*, *force\_const*, *ref\_val*)

## pysisyphus.calculators.FakeASE module

**class** pysisyphus.calculators.FakeASE.**FakeASE**(*calc*)

Bases: object

**get\_atoms\_coords**(*atoms*)

**get\_forces**(*atoms*=None)

**get\_potential\_energy**(*atoms*=None)

### pysisyphus.calculators.FourWellAnaPot module

```
class pysisyphus.calculators.FourWellAnaPot.FourWellAnaPot
    Bases: AnaPotBase
```

### pysisyphus.calculators.FreeEndNEBPot module

```
class pysisyphus.calculators.FreeEndNEBPot.FreeEndNEBPot
    Bases: AnaPotBase
    __init__()
        Analytical potential as described in [1] Appendix A
```

### pysisyphus.calculators.Gaussian09 module

```
class pysisyphus.calculators.Gaussian09.Gaussian09(*args, **kwargs)
    Bases: Gaussian16
    conf_key = 'gaussian09'
```

### pysisyphus.calculators.Gaussian16 module

```
class pysisyphus.calculators.Gaussian16.Gaussian16(route, gbs="", gen="", keep_chk=False, stable="",
                                                    fchk=None, **kwargs)

    Bases: OverlapCalculator
    conf_key = 'gaussian16'
    get_chkfiles()
    get_energy(atoms, coords, **prepare_kwargs)
        Meant to be extended.
    get_forces(atoms, coords, **prepare_kwargs)
        Meant to be extended.
    get_hessian(atoms, coords, **prepare_kwargs)
        Meant to be extended.
    keep(path)
        Backup calculation results.

        Parameters
            path (Path) -- Temporary directory of the calculation.

        Returns
            kept_fns -- Dictionary holding the filenames that were backed up. The keys correspond to the
            type of file.

        Return type
            dict
    make_exc_str()
```

```

make_fchk(path)
make_gbs_str()
parse_635r_dump(dump_path, roots, nmos)
parse_charges(path=None)
parse_double_mol(path, out_fn=None)
static parse_fchk(fchk_path, keys)
parse_force(path)
parse_hessian(path)
parse_keyword(text)
parse_log(*args, **kwargs)
parse_stable(path)
parse_tddft(path)
prepare_input(atoms, coords, calc_type, did_stable=False, point_charges=None)
    Meant to be extended.
prepare_overlap_data(path)
    Implement calculator specific parsing of MO coefficients and CI coefficients here. Should return a filename
    pointing to TURBOMOLE like mos, a MO coefficient array and a CI coefficient array.
reuse_data(path)
run_after(path)
    Meant to be extended.

    This method is called after a calculation was done, but before entering self.keep() and self.clean().
    Can be used to call tools like formchk or ricctools.
run_calculation(atoms, coords, **prepare_kwargs)
run_double_mol_calculation(atoms, coords1, coords2)
run_rwf_dump(path, rwf_index, chk_path=None)
run_stable(atoms, coords, **prepare_kwargs)
set_chkfiles(chkfiles)

```

### pysisyphus.calculators.HardSphere module

```

class pysisyphus.calculators.HardSphere.HardSphere(geom, frags, kappa=1.0, permutations=False,
                                                frag_radii=None)

```

Bases: object

```

__init__(geom, frags, kappa=1.0, permutations=False, frag_radii=None)

```

Intra-Image Inter-Molecular Hard-Sphere force.

See A.2. in [1], Eq. (A1).

```

get_forces(atoms, coords, kappa=None)

```

### pysisyphus.calculators.IDPPCalculator module

**class** pysisyphus.calculators.IDPPCalculator.IDPPCalculator(*target*)

Bases: *Calculator*

**get\_forces**(*atoms, coords*)

Meant to be extended.

### pysisyphus.calculators.IPIClient module

pysisyphus.calculators.IPIClient.**calc\_ipi\_client**(*addr, atoms, calc, queue=None, \*\*kwargs*)

pysisyphus.calculators.IPIClient.**ipi\_client**(*addr, atoms, energy\_getter, forces\_getter, hessian\_getter=None, hdrlen=12*)

### pysisyphus.calculators.IPIServer module

**class** pysisyphus.calculators.IPIServer.IPIServer(*\*args, address=None, host=None, port=None, unlink=True, hdrlen=12, max\_retries=0, verbose=False, \*\*kwargs*)

Bases: *Calculator*

**get\_energy**(*atoms, coords*)

Meant to be extended.

**get\_forces**(*atoms, coords*)

Meant to be extended.

**get\_hessian**(*atoms, coords*)

Meant to be extended.

**listen\_for**(*atoms, coords, kind='forces'*)

**listen\_for\_client\_atom\_num**(*atom\_num*)

**listen\_for\_energy**()

**listen\_for\_forces**(*atom\_num*)

**listen\_for\_hessian**(*atom\_num*)

**listen\_kinds** = ('coords', 'energy', 'forces', 'hessian')

**reset\_client\_connection**()

**retried\_listen\_for**(*atoms, coords*)

**unlink**(*address*)

**pysisyphus.calculators.LEPSBase module**

**class** pysisyphus.calculators.LEPSBase.LEPSBase(*pot\_type='leps'*)

Bases: *AnaPotBase*

**pysisyphus.calculators.LEPSExpr module**

**class** pysisyphus.calculators.LEPSExpr.LEPSExpr

Bases: object

**G**(*a, b*)

Gaussian function.

**Gdimer**(*x, y, A, x0, y0, sx, sy*)

**J**(*d, alpha, r0, r*)

Quantum mechanical exchange interaction.

**Q**(*d, alpha, r0, r*)

Coulomb interactions.

**V\_LEPS**(*x=None, y=None, abc=None*)

Equation (A.1) in [1]. Mimics reaction involving three atoms confined to motion along a line.

**V\_dimer**()

III. Results Section A in [3]. Two additional saddle points from two added gaussians.

**V\_harmonic**()

Equation (A.2) in [1]. A and C are fixed, only B can move. A condensed phase environment is represented by adding a harmonic oscillator degree of freedom.

**V\_tot**()

Equation (A.3) in [1]. Additional saddle point.

**\_\_init\_\_**()

Generates sympy expression for various LEPS potentials.

**get\_dimer**()

**get\_expr**(*pot\_type='leps'*)

**get\_harmonic**()

**get\_leps**()

**get\_tot**()

### pysisyphus.calculators.LennardJones module

**class** pysisyphus.calculators.LennardJones.LennardJones(*sigma=1.8897261251, epsilon=1, rc=None*)

Bases: *Calculator*

**calculate**(*coords3d*)

**get\_energy**(*atoms, coords*)

Meant to be extended.

**get\_forces**(*atoms, coords*)

Meant to be extended.

### pysisyphus.calculators.MOPAC module

**class** pysisyphus.calculators.MOPAC.MOPAC(*method='PM7', \*\*kwargs*)

Bases: *Calculator*

<http://openmopac.net/manual/>

**CALC\_TYPES** = {'energy': '1SCF', 'gradient': '1SCF GRADIENTS', 'hessian': 'DFORCE FORCE LET'}

**METHODS** = ['am1', 'pm3', 'pm6', 'pm6-dh2', 'pm6-d3', 'pm6-dh+', 'pm6-dh2', 'pm6-dh2x', 'pm6-d3h4', 'pm6-d3h4x', 'pm7', 'pm7-ts']

**MULT\_STRS** = {1: 'SINGLET', 2: 'DOUBLET', 3: 'TRIPLET', 4: 'QUARTET', 5: 'QUINTET', 6: 'SEXTET', 7: 'SEPTET', 8: 'OCTET'}

**base\_cmd**

Do only SCF AUX: Creates a checkpoint file NOREO: Dont reorient geometry

**Type**

1SCF

**conf\_key** = 'mopac'

**get\_energy**(*atoms, coords*)

Meant to be extended.

**get\_forces**(*atoms, coords*)

Meant to be extended.

**get\_hessian**(*atoms, coords*)

Meant to be extended.

**parse\_energy**(*path*)

**static parse\_energy\_from\_aux**(*inp, \*args, \*\*kwargs*)

**parse\_grad**(*path*)

**parse\_hessian**(*path*)

**static parse\_hessian\_from\_aux**(*inp, \*args, \*\*kwargs*)

**prepare\_coords**(*atoms*, *coords*, *opt=False*)

Get 3d coords in Angstrom.

Reshape internal 1d coords to 3d and convert to Angstrom.

**Parameters**

- **atoms** (*iterable*) -- Atom descriptors (element symbols).
- **coords** (*np.array*, *1d*) -- 1D-array holding coordinates in Bohr.

**Returns**

**coords** -- 3D-array holding coordinates in Angstrom.

**Return type**

*np.array*, 3d

**prepare\_input**(*atoms*, *coords*, *calc\_type*, *opt=False*)

Meant to be extended.

**read\_aux**(*path*)

### pysisyphus.calculators.MullerBrownSympyPot module

**class** pysisyphus.calculators.MullerBrownSympyPot.**MullerBrownPot**

Bases: *AnaPotBase*

### pysisyphus.calculators.MultiCalc module

**class** pysisyphus.calculators.MultiCalc.**MultiCalc**(*calcs*, *\*\*kwargs*)

Bases: *Calculator*

**run\_calculation**(*atoms*, *coords*, *\*\*prepare\_kwargs*)

pysisyphus.calculators.MultiCalc.**calcs\_from\_dict**(*calc\_dict*, *base\_name*, *calc\_number*, *charge*, *mult*, *pal*, *mem*)

### pysisyphus.calculators.OBabel module

**class** pysisyphus.calculators.OBabel.**OBabel**(*ff='gaff'*, *mol=None*, *\*\*kwargs*)

Bases: *Calculator*

**conv\_dict** = {'kcal/mol': 627.5094740630558, 'kJ/mol': 2625.4996394798254}

**get\_energy**(*atoms*, *coords*, *\*\*prepare\_kwargs*)

Meant to be extended.

**get\_forces**(*atoms*, *coords*, *\*\*prepare\_kwargs*)

Meant to be extended.

**setup**(*atoms*, *coords*)

**pysisyphus.calculators.ONIOMv2 module****class** pysisyphus.calculators.ONIOMv2.**LayerCalc**(*models, total\_size, parent\_layer\_calc=None*)

Bases: object

**property** charge**do\_parent**(*with\_parent*)**static energy\_from\_results**(*model\_energies, parent\_energy=None*)**get\_energy**(*atoms, coords, with\_parent=True*)**get\_forces**(*atoms, coords, with\_parent=True*)**get\_hessian**(*atoms, coords, with\_parent=True*)**property** mult**run\_calculations**(*atoms, coords, method*)**class** pysisyphus.calculators.ONIOMv2.**Link**(*ind, parent\_ind, atom, g*)

Bases: tuple

**atom**

Alias for field number 2

**g**

Alias for field number 3

**ind**

Alias for field number 0

**parent\_ind**

Alias for field number 1

**class** pysisyphus.calculators.ONIOMv2.**Model**(*name, calc\_level, calc, parent\_name, parent\_calc\_level, parent\_calc, atom\_inds, parent\_atom\_inds, use\_link\_atoms=True*)

Bases: object

**as\_calculator**(*cap=False*)**as\_geom**(*all\_atoms, all\_coords*)**capped\_atoms\_coords**(*all\_atoms, all\_coords*)**create\_bond\_vec\_getters**(*atoms*)**create\_links**(*atoms, coords, debug=False*)**get\_energy**(*atoms, coords, point\_charges=None, parent\_correction=True, cap=True*)**get\_forces**(*atoms, coords, point\_charges=None, parent\_correction=True, cap=True*)**get\_hessian**(*atoms, coords, point\_charges=None, parent\_correction=True, cap=True*)**get\_jacobian**()

**get\_sparse\_jacobian()**

**log**(*message=""*)

**parse\_charges()**

**class** pysisyphus.calculators.ONIOMv2.**ModelDummyCalc**(*model, cap=False*)

Bases: object

**get\_energy**(*atoms, coords*)

**get\_forces**(*atoms, coords*)

**class** pysisyphus.calculators.ONIOMv2.**ONIOM**(*calcs, models, geom, layers=None, embedding="", real\_key='real', use\_link\_atoms=True, \*args, \*\*kwargs*)

Bases: [Calculator](#)

**\_\_init\_\_**(*calcs, models, geom, layers=None, embedding="", real\_key='real', use\_link\_atoms=True, \*args, \*\*kwargs*)

**layer: list of models**

len(layer) == 1: normal ONIOM, len(layer) >= 1: multicenter ONIOM.

**model:**

(sub)set of all atoms that resides in a certain layer and has a certain calculator.

**atom\_inds\_in\_layer**(*index, exclude\_inner=False*)

Returns list of atom indices in layer at index.

Atoms that also appear in inner layer can be excluded on request.

**Parameters**

- **index** (*int*) -- pasd
- **exclude\_inner** (*bool, default=False, optional*) -- Whether to exclude atom indices that also appear in inner layers.

**Returns**

**atom\_indices** -- List containing the atom indices in the selected layer.

**Return type**

list

**calc\_layer**(*atoms, coords, index, parent\_correction=True*)

**property charge**

```
embeddings = {'': '', 'electronic': 'Electronic embedding', 'electronic_rc':
'Electronic embedding with redistributed charges', 'electronic_rcd': 'Electronic
embedding with redistributed charges and dipoles'}
```

**get\_energy**(*atoms, coords*)

Meant to be extended.

**get\_forces**(*atoms, coords*)

Meant to be extended.

**get\_hessian**(*atoms, coords*)

Meant to be extended.

```
get_layer_calc(layer_ind)
```

```
property model_iter
```

```
property mult
```

```
run_calculation(atoms, coords)
```

```
run_calculations(atoms, coords, method)
```

```
pysisyphus.calculators.ONIOMv2.atom_inds_to_cart_inds(atom_inds)
```

```
pysisyphus.calculators.ONIOMv2.cap_fragment(atoms, coords, fragment, link_atom='H', g=0.709)
```

```
pysisyphus.calculators.ONIOMv2.get_embedding_charges(embedding, layer, parent_layer, coords3d)
```

```
pysisyphus.calculators.ONIOMv2.get_g_value(atom, parent_atom, link_atom)
```

### pysisyphus.calculators.ORCA module

```
class pysisyphus.calculators.ORCA.ORCA(keywords, blocks="", gbw=None, do_stable=False,
                                       numfreq=False, **kwargs)
```

Bases: *OverlapCalculator*

```
__init__(keywords, blocks="", gbw=None, do_stable=False, numfreq=False, **kwargs)
```

ORCA calculator.

Wrapper for creating ORCA input files for energy, gradient and Hessian calculations. The PAL and memory inputs must not be given in the keywords and/or blocks, as they are handled by the 'pal' and 'memory' arguments.

#### Parameters

- **keywords** (*str*) -- Keyword line, as normally given in ORCA, excluding the leading "!".
- **blocks** (*str*, *optional*) -- ORCA block input(s), e.g. for TD-DFT calculations (%tddft ... end). As the blocks start with a leading "%", wrapping the input in quotes (") is required, otherwise the parsing will fail.
- **gbw** (*str*, *optional*) -- Path to an input gbw file, which will be used as initial guess for the first calculation. Will be overridden later, with the path to the gbw file of a previous calculation.
- **do\_stable** (*bool*, *optional*) -- Run stability analysis until a stable wavefunction is obtained, before every calculation.
- **numfreq** (*boo*, *optional*) -- Use numerical frequencies instead of analytical ones.
- **mem** (*int*) -- Mememory per core in MB.

```
check_termination(*args, **kwargs)
```

```
clean_tmp(path)
```

```
conf_key = 'orca'
```

```
get_block_str()
```

```
get_chkfiles()
```

**get\_energy**(*atoms*, *coords*, *\*\*prepare\_kwargs*)

Meant to be extended.

**get\_forces**(*atoms*, *coords*, *\*\*prepare\_kwargs*)

Meant to be extended.

**get\_hessian**(*atoms*, *coords*, *\*\*prepare\_kwargs*)

Meant to be extended.

**get\_moinp\_str**(*gbw*)

**get\_stable\_wavefunction**(*atoms*, *coords*)

**keep**(*path*)

Backup calculation results.

**Parameters**

**path** (*Path*) -- Temporary directory of the calculation.

**Returns**

**kept\_fns** -- Dictionary holding the filenames that were backed up. The keys correspond to the type of file.

**Return type**

dict

**parse\_all\_energies**(*text=None*, *triplets=None*)

**static parse\_atoms\_coords**(*inp*, *\*args*, *\*\*kwargs*)

**parse\_cis**(*cis*)

**Read binary CI vector file from ORCA.**

Adapted from TheoDORE 1.7.1, Authors: S. Mai, F. Plasser <https://sourceforge.net/p/theodore-qc>

**parse\_energy**(*path*)

**parse\_engrad**(*path*)

**static parse\_engrad\_info**(*inp*, *\*args*, *\*\*kwargs*)

**parse\_gbw**(*gbw\_fn*)

Adapted from <https://orcaforum.kofo.mpg.de/viewtopic.php?f=8&t=3299>

The first 5 long int values represent pointers into the file:

Pointer @+0: Internal ORCA data structures Pointer @+8: Geometry Pointer @+16: BasisSet Pointer @+24: Orbitals Pointer @+32: ECP data

**static parse\_hess\_file**(*inp*, *\*args*, *\*\*kwargs*)

**parse\_hessian**(*path*)

**parse\_mo\_numbers**(*out\_fn*)

**parse\_stable**(*path*)

**prepare\_input**(*atoms*, *coords*, *calc\_type*, *point\_charges=None*, *do\_stable=False*)

Meant to be extended.

**prepare\_overlap\_data**(*path*)

Implement calculator specific parsing of MO coefficients and CI coefficients here. Should return a filename pointing to TURBOMOLE like mos, a MO coefficient array and a CI coefficient array.

**reattach**(*last\_calc\_cycle*)**run\_after**(*path*)

Meant to be extended.

This method is called after a calculation was done, but before entering `self.keep()` and `self.clean()`. Can be used to call tools like `formchk` or `ricctools`.

**run\_calculation**(*atoms*, *coords*, **\*\*prepare\_kwargs**)

Basically some kind of dummy method that can be called to execute ORCA with the stored cmd of this calculator.

**set\_chkfiles**(*chkfiles*)**set\_mo\_coeffs**(*mo\_coeffs=None*, *gbw=None*)**static set\_mo\_coeffs\_in\_gbw**(*in\_gbw\_fn*, *out\_gbw\_fn*, *mo\_coeffs*)

See `self.parse_gbw`.

**store\_and\_track**(*results*, *func*, *atoms*, *coords*, **\*\*prepare\_kwargs**)

`pysisyphus.calculators.ORCA.make_sym_mat`(*table\_block*)

`pysisyphus.calculators.ORCA.save_orca_pc_file`(*point\_charges*, *pc\_fn*, *hardness=None*)

**pysisyphus.calculators.ORCA5 module**

```
class pysisyphus.calculators.ORCA5.ORCA5(keywords, blocks="", gbw=None, do_stable=False,  
                                         numfreq=False, **kwargs)
```

Bases: `ORCA`

`conf_key = 'orca5'`

**pysisyphus.calculators.OpenMM module**

```
class pysisyphus.calculators.OpenMM.OpenMM(topology, params, **kwargs)
```

Bases: `Calculator`

`get_charges`()

`get_dipole_moment`(*coords3d*, *reference=None*, *masses=None*)

`get_energy`(*atoms*, *coords*, **\*\*prepare\_kwargs**)

Meant to be extended.

`get_forces`(*atoms*, *coords*, **\*\*prepare\_kwargs**)

Meant to be extended.

`report_charges`()

Total charge of the system.

Adapted from `modeller.py _addIons()` in `OpenMM`.

**pysisyphus.calculators.OpenMolcas module**

**class** pysisyphus.calculators.OpenMolcas.**OpenMolcas**(*basis, inporb, roots, mdlxroot, supsym=None, track=True, \*\*kwargs*)

Bases: *Calculator*

**build\_rassi\_str**()

**build\_supsym\_str**(*supsym*)

Can handle only one subgroup for now.

**conf\_key** = 'openmolcas'

**get\_energy**(*atoms, coords*)

Meant to be extended.

**get\_forces**(*atoms, coords*)

Meant to be extended.

**get\_pal\_env**()

**keep**(*path*)

Backup calculation results.

**Parameters**

**path** (*Path*) -- Temporary directory of the calculation.

**Returns**

**kept\_fns** -- Dictionary holding the filenames that were backed up. The keys correspond to the type of file.

**Return type**

dict

**parse\_energies**(*text*)

**parse\_gradient**(*path*)

**parse\_rassi\_track**(*path*)

**prepare\_coords**(*atoms, coords*)

Get 3d coords in Angstrom.

Reshape internal 1d coords to 3d and convert to Angstrom.

**Parameters**

- **atoms** (*iterable*) -- Atom descriptors (element symbols).
- **coords** (*np.array, 1d*) -- 1D-array holding coordinates in Bohr.

**Returns**

**coords** -- 3D-array holding coordinates in Angstrom.

**Return type**

np.array, 3d

**prepare\_input**(*atoms, coords*)

Meant to be extended.

**reattach**(*last\_calc\_cycle*)

**pysisyphus.calculators.OverlapCalculator module****class** pysisyphus.calculators.OverlapCalculator.**NTOs**(*ntos, lambdas*)

Bases: tuple

**lambdas**

Alias for field number 1

**ntos**

Alias for field number 0

**class** pysisyphus.calculators.OverlapCalculator.**OverlapCalculator**(\*args, track=False, ovlp\_type='tden', double\_mol=False, ovlp\_with='previous', XY='X+Y', adapt\_args=(0.5, 0.3, 0.6), use\_ntos=4, pr\_nto=False, nto\_thresh=0.3, cdds=None, orient='', dump\_fn='overlap\_data.h5', h5\_dump=False, ncore=0, conf\_thresh=0.001, dyn\_roots=0, mos\_ref='cur', mos\_renorm=True, \*\*kwargs)Bases: *Calculator***H5\_MAP** = {'all\_energies': 'all\_energies\_list', 'ci\_coeffs': 'ci\_coeff\_list', 'coords': 'coords\_list', 'mo\_coeffs': 'mo\_coeff\_list', 'ref\_roots': 'reference\_roots', 'roots': 'roots\_list'}**OVLTYPE\_VERBOSE** = {'nto': 'natural transition orbital overlap', 'nto\_org': 'original natural transition orbital overlap', 'tden': 'transition density matrix overlap', 'wf': 'wavefunction overlap'}**VALID\_CDDS** = (None, 'calc', 'render')**VALID\_KEYS** = ['wf', 'tden', 'nto', 'nto\_org']**VALID\_XY** = ('X', 'X+Y', 'X-Y')**blowup\_ci\_coeffs**(*ci\_coeffs*)**calc\_cdd\_cube**(*root, cycle=-1*)**calculate\_state\_ntos**(*state\_ci\_coeffs, mos*)**property** data\_model**dump\_overlap\_data**()**static** **from\_overlap\_data**(*h5\_fn, set\_wfow=False*)**get\_h5\_group**()

**get\_indices**(*indices=None*)

A new root is determined by selecting the overlap matrix row corresponding to the reference root and checking for the root with the highest overlap (at the current geometry).

The overlap matrix is usually formed by a double loop like:

```
overlap_matrix = np.empty((ref_states, cur_states))
for i, ref_state in enumerate(ref_states):
```

```
    for j, cur_state in enumerate(cur_states):
        overlap_matrix[i, j] = make_overlap(ref_state, cur_state)
```

So the reference states run along the rows. That's why the ref\_state index comes first in the 'indices' tuple.

**static get\_mo\_norms**(*mo\_coeffs, ao\_ovlp*)

MOs are in rows.

**get\_nto\_overlaps**(*indices=None, ao\_ovlp=None, org=False*)

**get\_orbital\_matrices**(*indices=None, ao\_ovlp=None*)

Return MO coefficients and AO overlaps for the given indices.

If not provided, a AO overlap matrix is constructed from one of the MO coefficient matrices (controlled by self.mos\_ref). Also, if requested one of the two MO coefficient matrices is re-normalized.

**get\_ref\_mos**(*ref\_mo\_coeffs, cur\_mo\_coeffs*)

**static get\_sao\_from\_mo\_coeffs**(*mo\_coeffs*)

Recover AO overlaps from given MO coefficients.

For MOs in the columns of mo\_coeffs:

$$S_{AO} = C^{1^T} C^1 S_{AO} C = C^{1^T} (S_{AO} C)^T = C^1 C^{1^T} S_{AO}^T = C^1 C^{1^T} S_{AO} C = I$$

Here, MOs are expected to be in rows of mo\_coeffs, yielding

$$C S_{AO} C^{1^T} = I$$

**get\_sao\_from\_mo\_coeffs\_and\_dump**(*mo\_coeffs*)

**get\_tden\_overlaps**(*indices=None, ao\_ovlp=None*)

**get\_wf\_overlaps**(*indices=None, ao\_ovlp=None*)

**nto\_org\_overlaps**(*ntos\_1, ntos\_2, ao\_ovlp, nto\_thresh=0.3*)

**nto\_overlaps**(*ntos\_1, ntos\_2, ao\_ovlp*)

**prepare\_overlap\_data**(*path*)

Implement calculator specific parsing of MO coefficients and CI coefficients here. Should return a filename pointing to TURBOMOLE like mos, a MO coefficient array and a CI coefficient array.

**render\_cdd\_cube**()

**static renorm\_mos**(*mo\_coeffs, ao\_ovlp*)

**property roots\_number**

**set\_ntos**(*mo\_coeffs, ci\_coeffs*)

**set\_wfow**(*ci\_coeffs*)

**store\_overlap\_data**(*atoms, coords, path=None, overlap\_data=None*)

**tden\_overlap\_with\_calculator**(*calc, ao\_ovlp=None*)

**tden\_overlaps**(*mo\_coeffs1, ci\_coeffs1, mo\_coeffs2, ci\_coeffs2, ao\_ovlp*)

#### Parameters

- **mo\_coeffs1** (*ndarray, shape (MOs, AOs)*) -- MO coefficient matrix. One row per MO, one column per basis function. Usually square.
- **mo\_coeffs2** (*ndarray*) -- See `mo_coeffs1`.
- **ci\_coeffs1** (*ndarray, shape (occ. MOs, MOs)*) -- CI-coefficient matrix.
- **ci\_coeffs2** (*ndarray, shape (occ. MOs, MOs)*) -- See `ci_coeffs1`.
- **ao\_ovlp** (*ndarray, shape (AOs1, AOs2)*) -- Double molecule AO overlaps.

**track\_root**(*ovlp\_type=None*)

Check if a root flip occurred compared to the previous cycle by calculating the overlap matrix wrt. a reference cycle.

**update\_array\_dims**()

Assure consistent array shapes when the number of calculated roots changed.

**wf\_overlap\_with\_calculator**(*calc, ao\_ovlp=None*)

**wf\_overlaps**(*mo\_coeffs1, ci\_coeffs1, mo\_coeffs2, ci\_coeffs2, ao\_ovlp=None*)

`pysisyphus.calculators.OverlapCalculator.get_data_model`(*exc\_state\_num, occ\_mo\_num, virt\_mo\_num, ovlp\_type, atoms, max\_cycles*)

## pysisyphus.calculators.Psi4 module

**class** `pysisyphus.calculators.Psi4.Psi4`(*method, basis, to\_set=None, pcm='iefpcm', solvent=None, write\_fchk=False, \*\*kwargs*)

Bases: `Calculator`

**conf\_key** = 'psi4'

**get\_energy**(*atoms, coords, \*\*prepare\_kwargs*)

Meant to be extended.

**get\_fchk\_str**()

**get\_forces**(*atoms, coords, \*\*prepare\_kwargs*)

Meant to be extended.

**get\_hessian**(*atoms, coords, \*\*prepare\_kwargs*)

Meant to be extended.

**parse\_energy**(*path*)

**parse\_grad**(*path*)

**parse\_hessian**(*path*)

**prepare\_input**(*atoms, coords, calc\_type*)

Meant to be extended.

`run_calculation(atoms, coords, **prepare_kwargs)`

### pysisyphus.calculators.PyPsi4 module

`class pysisyphus.calculators.PyPsi4.PyPsi4(method, basis, **kwargs)`

Bases: *Calculator*

`get_forces(atoms, coords)`

Meant to be extended.

### pysisyphus.calculators.PySCF module

### pysisyphus.calculators.PyXTB module

`class pysisyphus.calculators.PyXTB.PyXTB(*args, gfn=2, acc=None, verbosity=0, keep_calculator=False, **kwargs)`

Bases: *Calculator*

`get_calculator(atoms, coords)`

`get_energy(atoms, coords, **prepare_kwargs)`

Meant to be extended.

`get_forces(atoms, coords, **prepare_kwargs)`

Meant to be extended.

### pysisyphus.calculators.QCEngine module

### pysisyphus.calculators.Rastrigin module

`class pysisyphus.calculators.Rastrigin.Rastrigin`

Bases: *AnaPotBase*

<http://www.sfu.ca/~ssurjano/rastr.html>

### pysisyphus.calculators.Remote module

`class pysisyphus.calculators.Remote.Remote(remote_calc, host, prefix="", **kwargs)`

Bases: *Calculator*

`get_energy(atoms, coords, **prepare_kwargs)`

Meant to be extended.

`get_forces(atoms, coords, **prepare_kwargs)`

Meant to be extended.

`get_hessian(atoms, coords, **prepare_kwargs)`

Meant to be extended.

`parse_results()`

`run_calculation(atoms, coords, run_func='get_energy')`

### pysisyphus.calculators.Rosenbrock module

**class** pysisyphus.calculators.Rosenbrock.**Rosenbrock**

Bases: *AnaPotBase*

### pysisyphus.calculators.SocketCalc module

**class** pysisyphus.calculators.SocketCalc.**SocketCalc**(\*args, host='localhost', port=8080, \*\*kwargs)

Bases: *Calculator*

**get\_energy**(atoms, coords)

Meant to be extended.

**get\_forces**(atoms, coords)

Meant to be extended.

**get\_hessian**(atoms, coords)

Meant to be extended.

**listen\_for**(atoms, coords, request)

**valid\_requests** = ('energy', 'forces', 'hessian')

### pysisyphus.calculators.TIP3P module

**class** pysisyphus.calculators.TIP3P.**TIP3P**(rc=9.44863062728914)

Bases: *Calculator*

Transferable Intermolecular Potential 3 Point

**aHOH** = 104.52

**calculate**(coords3d)

**charges**

**coulomb\_energy** = (multiple of elem. charge \* multiple of elem. charge)

/ (distance in bohr) \* 1 / (4 \* pi \* vacuum permittivity)

coulomb\_prefactor converts everything to atomic units and it is ... drum roll ... 1. from scipy.constants  
import value as pcv self.coulomb\_prefactor = (1 / (4 \* np.pi) \* pcv("elementary charge"))\*\*2

/ pcv("Hartree energy") / pcv("Bohr radius") / pcv("vacuum electric permittivity")

)

**coulomb**(coords3d)

**epsilon** = 0.0002423919586315716

**get\_energy**(atoms, coords)

Meant to be extended.

**get\_forces**(atoms, coords)

Meant to be extended.

```

qH = 0.417
q0 = -0.834
rOH = 1.8088458464917874
sigma = 5.953790025507198

```

### pysisyphus.calculators.TransTorque module

```

class pysisyphus.calculators.TransTorque.TransTorque(frags, iter_frags, a_mats, b_mats,
                                                    weight_func=None, skip=True, kappa=1.0,
                                                    b_coords3d=None, do_trans=True)

```

Bases: object

```

__init__(frags, iter_frags, a_mats, b_mats, weight_func=None, skip=True, kappa=1.0, b_coords3d=None,
        do_trans=True)

```

Translational and torque forces. See A.4. [1], Eqs. (A3) - (A5).

```

get_forces(atoms, coords, kappa=None)

```

```

get_forces_naive(atoms, coords, kappa=None)

```

```

set_N_invs()

```

```

pysisyphus.calculators.TransTorque.get_trans_torque_forces(mfrag, a_coords3d, b_coords3d,
                                                         a_mats, b_mats, m, frags, N_inv,
                                                         weight_func=None, skip=True,
                                                         kappa=1, do_trans=True)

```

### pysisyphus.calculators.Turbomole module

```

class pysisyphus.calculators.Turbomole.Turbomole(control_path, root=None, double_mol_path=None,
                                                **kwargs)

```

Bases: *OverlapCalculator*

```

append_control(to_append, log_msg="", **kwargs)

```

```

conf_key = 'turbomole'

```

```

get_chkfiles()

```

```

get_energy(atoms, coords, **prepare_kwargs)

```

Meant to be extended.

```

get_forces(atoms, coords, cmd=None, **prepare_kwargs)

```

Meant to be extended.

```

get_hessian(atoms, coords, **prepare_kwargs)

```

Meant to be extended.

```

get_pal_env()

```

```

get_ricc2_root(text)

```

**keep**(*path*)

Backup calculation results.

**Parameters**

**path** (*Path*) -- Temporary directory of the calculation.

**Returns**

**kept\_fns** -- Dictionary holding the filenames that were backed up. The keys correspond to the type of file.

**Return type**

dict

**make\_molden**(*path*)**parse\_cc2\_vectors**(*ccre*)**parse\_double\_mol**(*path*)

Parse a double molecule overlap matrix from Turbomole output to be used with WFOWrapper.

**parse\_energy**(*path*)**parse\_force**(*path*)**parse\_gs\_energy**()

Several places are possible: \$subenergy from control file total energy from turbomole.out Final MP2 energy from turbomole.out with ADC(2) Final CC2 energy from turbomole.out with CC(2)

**parse\_hessian**(*path, fn=None*)**parse\_mos**()**parse\_td\_vectors**(*text*)

For TDA calculations only the X vector is present in the *ciss\_a/etc.* file. In TDDFT calculations there are twice as much items compared with TDA. The first half corresponds to (X+Y) and the second half to (X-Y). X can be calculated as  $X = ((X+Y)+(X-Y))/2$ . Y is then given as  $Y = (X+Y)-X$ . The normalization can then be checked as

```
np.concatenate((X, Y)).dot(np.concatenate((X, -Y)))
```

and should be 1.

**prepare\_input**(*atoms, coords, calc\_type, point\_charges=None*)

To rectify this we have to construct the *basemcd* dynamically and construct it ad hoc. We could set a RI flag in the beginning and select the correct *scf* binary here from it. Then we select the following binary on demand, e.g. *aoforce* or *rdgrad* or *egrad* etc.

**prepare\_overlap\_data**(*path*)

Implement calculator specific parsing of MO coefficients and CI coefficients here. Should return a filename pointing to TURBOMOLE like *mos*, a MO coefficient array and a CI coefficient array.

**prepare\_point\_charges**(*point\_charges*)

```
$point_charges <x> <y> <z> <q>
```

**prepare\_td**(*text*)**run\_after**(*path*)

Meant to be extended.

This method is called after a calculation was done, but before entering `self.keep()` and `self.clean()`. Can be used to call tools like *formchk* or *ricctools*.

**run\_calculation**(*atoms, coords*)

Basically some kind of dummy method that can be called to execute Turbomole with the stored cmd of this calculator.

**run\_double\_mol\_calculation**(*atoms, coords1, coords2*)

**set\_chkfiles**(*chkfiles*)

**set\_occ\_and\_mo\_nums**(*text*)

**store\_and\_track**(*results, func, atoms, coords, \*\*prepare\_kwargs*)

**sub\_control**(*pattern, repl, log\_msg="", \*\*kwargs*)

## pysisyphus.calculators.WFOWrapper module

**class** pysisyphus.calculators.WFOWrapper.**WFOWrapper**(*occ\_mo\_num, virt\_mo\_num, conf\_thresh=0.001, calc\_number=0, out\_dir='./, wfow\_mem=8000, ncore=0, debug=False*)

Bases: object

**ci\_coeffs\_above\_thresh**(*ci\_coeffs, thresh=None*)

**property conf\_thresh**

**fake\_turbo\_mos**(*mo\_coeffs*)

Create a mos file suitable for TURBOMOLE input. All MO eigenvalues are set to 0.0. There is also a little deviation in the formatting (see `turbo_fmt()`) but it works ...

**generate\_all\_dets**(*occ\_set1, virt\_set1, occ\_set2, virt\_set2*)

Generate all possible single excitation determinant strings from union(occ\_mos) to union(virt\_mos).

**get\_from\_to\_sets**(*ci\_coeffs*)

**get\_gs\_line**(*ci\_coeffs\_with\_gs*)

**log**(*message*)

**logger** = <Logger wfoverlap (DEBUG)>

**make\_det\_string**(*inds*)

Return spin adapted strings.

**make\_dets\_header**(*cic, dets\_list*)

**make\_full\_dets\_list**(*all\_inds, det\_strings, ci\_coeffs*)

**matrix\_types** = {'ortho': 'Orthonormalized overlap matrix', 'ovlp': 'Overlap matrix', 'renorm': 'Renormalized overlap matrix'}

**parse\_wfoverlap\_out**(*text, type\_='ortho'*)

Returns overlap matrix.

**set\_from\_nested\_list**(*nested*)

**wf\_overlap**(*cycle1, cycle2, ao\_ovlp=None*)

## pysisyphus.calculators.WFOWrapper2 module

```
class pysisyphus.calculators.WFOWrapper2.WFOWrapper2(overlap_data, calc_number=0,  
                                                    conf_thresh=0.0001, out_dir='./')
```

Bases: object

**all\_overlaps()**

**ci\_coeffs\_above\_thresh**(*ci\_coeffs*, *thresh=1e-05*)

**static fake\_turbo\_mos**(*mo\_coeffs*)

Create a mos file suitable for TURBOMOLE input. All MO eigenvalues are set to 0.0. There is also a little deviation in the formatting (see `turbo_fmt()`) but it works ...

**generate\_all\_dets**(*occ\_set1*, *virt\_set1*, *occ\_set2*, *virt\_set2*)

Generate all possible single excitation determinant strings from `union(occ_mos)` to `union(virt_mos)`.

**get\_iteration**(*ind*)

**property last\_two\_coords**

**log**(*message*)

**logger** = <Logger wfoverlap (DEBUG)>

**make\_det\_string**(*inds*)

Return spin adapted strings.

**make\_dets\_header**(*cic*, *dets\_list*)

**make\_full\_dets\_list**(*all\_inds*, *det\_strings*, *ci\_coeffs*)

**matrix\_types** = {'ortho': 'Orthonormalized overlap matrix', 'ovlp': 'Overlap matrix', 'renorm': 'Renormalized overlap matrix'}

**parse\_wfoverlap\_out**(*text*, *type\_*='ortho')

Returns overlap matrix.

**set\_data**()

**set\_from\_nested\_list**(*nested*)

**wf\_overlap**(*ind1=- 2*, *ind2=- 1*, *ao\_ovlp=None*)

## pysisyphus.calculators.XTB module

```
class pysisyphus.calculators.XTB.OptResult(opt_geom, opt_log)
```

Bases: tuple

**opt\_geom**

Alias for field number 0

**opt\_log**

Alias for field number 1

```
class pysisyphus.calculators.XTB.XTB(gbsa="", alpb="", gfn=2, acc=1.0, etemp=None, retry_etemp=None,
                                     topo=None, topo_update=None, quiet=False, **kwargs)
```

Bases: *Calculator*

```
__init__(gbsa="", alpb="", gfn=2, acc=1.0, etemp=None, retry_etemp=None, topo=None,
         topo_update=None, quiet=False, **kwargs)
```

XTB calculator.

Wrapper for running energy, gradient and Hessian calculations by XTB.

#### Parameters

- **gbsa** (*str, optional*) -- Solvent for GBSA calculation, by default no solvent model is used.
- **alpb** (*str, optional*) -- Solvent for ALPB calculation, by default no solvent model is used.
- **gfn** (*int or str, must be (0, 1, 2, or "ff")*) -- Hamiltonian for the XTB calculation (GFN0, GFN1, GFN2, or GFNFF).
- **acc** (*float, optional*) -- Accuracy control of the calculation, the lower the tighter several numerical thresholds are chosen.
- **topo** (*str, optional*) -- Path the a GFNFF-topolgy file. As setting up the topology may take some time for sizable systems, it may be desired to reuse the file.
- **topo\_update** (*int*) -- Integer controlling the update interval of the GFNFF topology update. If supplied, the topolgy will be recreated every N-th calculation.
- **mem** (*int*) -- Mememory per core in MB.
- **quiet** (*bool, optional*) -- Suppress creation of log files.

```
static check_termination(inp, *args, **kwargs)
```

```
conf_key = 'xtb'
```

```
get_energy(atoms, coords, **prepare_kwargs)
```

Meant to be extended.

```
get_forces(atoms, coords, **prepare_kwargs)
```

Meant to be extended.

```
get_hessian(atoms, coords, **prepare_kwargs)
```

Meant to be extended.

```
get_mdrestart_str(coords, velocities)
```

coords and velocities have to given in au!

```
get_pal_env()
```

```
get_retry_args()
```

```
keep(path)
```

Backup calculation results.

#### Parameters

**path** (*Path*) -- Temporary directory of the calculation.

#### Returns

**kept\_fns** -- Dictionay holding the filenames that were backed up. The keys correspond to the type of file.

**Return type**

dict

**parse\_charges**(*fn=None*)

**parse\_charges\_from\_json**(*fn=None*)

**parse\_energy**(*path*)

**parse\_gradient**(*path*)

**parse\_hessian**(*path*)

**parse\_md**(*path*)

**parse\_opt**(*path, keep\_log=False*)

**parse\_topo**(*path*)

**prepare\_add\_args**(*xcontrol=None*)

**prepare\_coords**(*atoms, coords*)

Get 3d coords in Angstrom.

Reshape internal 1d coords to 3d and convert to Angstrom.

**Parameters**

- **atoms** (*iterable*) -- Atom descriptors (element symbols).
- **coords** (*np.array, 1d*) -- 1D-array holding coordinates in Bohr.

**Returns**

**coords** -- 3D-array holding coordinates in Angstrom.

**Return type**

np.array, 3d

**prepare\_input**(*atoms, coords, calc\_type, point\_charges=None*)

Meant to be extended.

**reattach**(*last\_calc\_cycle*)

**run\_calculation**(*atoms, coords, \*\*prepare\_kwargs*)

**run\_md**(*atoms, coords, t, dt, velocities=None, dump=1*)

Expecting t and dt in fs, even though xtb wants t in ps!

**run\_opt**(*atoms, coords, keep=True, keep\_log=False*)

**run\_topo**(*atoms, coords*)

**write\_mdrestart**(*path, mdrestart\_str*)

## pysisyphus.calculators.parser module

pysisyphus.calculators.parser.**make\_float\_class**(\*\*kwargs)

pysisyphus.calculators.parser.**parse\_turbo\_ccre0\_ascii**(text)

pysisyphus.calculators.parser.**parse\_turbo\_exstates**(text)

Parse excitation energies (first blocks) from an exstates file.

pysisyphus.calculators.parser.**parse\_turbo\_gradient**(path)

pysisyphus.calculators.parser.**parse\_turbo\_mos**(text)

pysisyphus.calculators.parser.**to\_float**(s, loc, toks)

## Module contents

**class** pysisyphus.calculators.**AFIR**(calculator, fragment\_indices, gamma, rho=1, p=6, ignore\_hydrogen=True, complete\_fragments=True, dump=True, h5\_fn='afir.h5', h5\_group\_name='afir', \*\*kwargs)

Bases: [Calculator](#)

**\_\_init\_\_**(calculator, fragment\_indices, gamma, rho=1, p=6, ignore\_hydrogen=True, complete\_fragments=True, dump=True, h5\_fn='afir.h5', h5\_group\_name='afir', \*\*kwargs)

Artificial Force Induced Reaction calculator.

Currently, there are no automated drivers to run large-scale AFIR calculations with many different initial orientations and/or increasing collision energy parameter. Nonetheless, selected AFIR calculations can be carried out by hand. After convergence, artificial potential & forces, as well as real energies and forces can be plotted with 'pysisplot --afir'. The highest energy point along the AFIR path can then be selected for a subsequent TS-optimization, e.g. via 'pysistrj --get [index] optimization.trj'.

Future versions of pysisyphus may provide drivers for more automated AFIR calculations.

### Parameters

- **calculator** ([Calculator](#)) -- Actual QC calculator that provides energies and its derivatives, that are modified by the AFIR calculator, e.g., ORCA or Psi4.
- **fragment\_indices** (List[List[int]]) -- List of lists of integers, specifying the separate fragments. If the indices in these lists don't comprise all atoms in the molecule, the remaining indices will be added as a separate fragment. If a AFIR calculation is carried out with 2 fragments and 'complete\_fragments' is True (see below) it is enough to specify only the indices of one fragment, e.g., for a system of 10 atoms 'fragment\_indices=[[0,1,2,3]]' is enough. The second system will be set up automatically with indices [4,5,6,7,8,9].
- **gamma** (Union[\_SupportsArray[dtype], \_NestedSequence[\_SupportsArray[dtype]], bool, int, float, complex, str, bytes, \_NestedSequence[Union[bool, int, float, complex, str, bytes]]]) -- Collision energy parameter in au. For 2 fragments it can be a single integer, while for > 2 fragments a list of gammas must be given, specifying the pair-wise collision energy parameters. For 3 fragments 3 gammas must be given [\_01, \_02, \_12], for 4 fragments 6 gammas would be required [\_01, \_02, \_03, \_12, \_13, \_23] and so on.
- **rho** (Union[\_SupportsArray[dtype], \_NestedSequence[\_SupportsArray[dtype]], bool, int, float, complex, str, bytes, \_NestedSequence[Union[bool, int, float, complex, str, bytes]]], default: 1) -- Direction of the artificial force, either 1 or -1. The

same comments as for gamma apply. For 2 fragments a single integer is enough, for > 2 fragments a list of rhos must be given (see above). For rho=1 fragments are pushed together, for rho=-1 fragments are pulled apart.

- **p** (int, default: 6) -- Exponent p used in the calculation of the weight function . Defaults to 6 and probably does not have to be changed.
- **ignore\_hydrogen** (bool, default: True) -- Whether hydrogens are ignored in the calculation of the artificial force.
- **complete\_fragments** (bool, default: True) -- Whether an incomplete specification in 'fragment\_indices' is automatically completed.
- **dump** (bool, default: True) -- Whether an HDF5 file is created.
- **h5\_fn** (str, default: 'afir.h5') -- Filename of the HDF5 file used for dumping.
- **h5\_group\_name** (str, default: 'afir') -- HDF5 group name used for dumping.
- **\*\*kwargs** -- Keyword arguments passed to the Calculator baseclass.

**afir\_fd\_hessian\_wrapper**(*coords3d, afir\_grad\_func*)

**property charge**

**dump\_h5**(*atoms, coords, results*)

**get\_energy**(*atoms, coords, \*\*prepare\_kwargs*)

Meant to be extended.

**get\_forces**(*atoms, coords, \*\*prepare\_kwargs*)

Meant to be extended.

**get\_hessian**(*atoms, coords, \*\*prepare\_kwargs*)

Meant to be extended.

**init\_h5\_group**(*atoms, max\_cycles=None*)

**log\_fragments**()

**property mult**

**set\_atoms\_and\_funcs**(*atoms, coords*)

Initially atoms was also an argument to the constructor of AFIR. I removed it so creation becomes easier. The first time a calculation is requested with a proper atom set everything is set up (cov. radii, afir function and corresponding gradient). Afterwards there is only a check if atoms != None and it is expected that all functions are properly set.

fragment\_indices can also be incomplete w.r.t. to the number of atoms. If the sum of the specified fragment atoms is less than the number of atoms present then all remaining unspecified atoms will be gathered in one fragment.

**write\_fragment\_geoms**(*atoms, coords*)

**class pysisyphus.calculators.AtomAtomTransTorque**(*geom, frags, A\_mats, kappa=2.0*)

Bases: object

**\_\_init\_\_**(*geom, frags, A\_mats, kappa=2.0*)

Atom-atom translational and torque forces.

See A.5. [1], Eq. (A6).

**get\_forces**(*atoms*, *coords*)

**class** pysisyphus.calculators.**Composite**(*final*, *keys\_calcs=None*, *calcs=None*, *remove\_translation=False*,  
\*\**kwargs*)

Bases: *Calculator*

**get\_energy**(*atoms*, *coords*, \*\**prepare\_kwargs*)

Meant to be extended.

**get\_forces**(*atoms*, *coords*, \*\**prepare\_kwargs*)

Meant to be extended.

**run\_calculation**(*atoms*, *coords*, \*\**prepare\_kwargs*)

**class** pysisyphus.calculators.**ConicalIntersection**(*calculator1*, *calculator2*, \*\**kwargs*)

Bases: *Calculator*

Calculator for conical intersection optimization.

Based on [1].

**get\_ci\_quantities**(*atoms*, *coords*, \*\**prepare\_kwargs*)

Relevant quantities including branching plane and projector P.

**get\_energy**(*atoms*, *coords*, \*\**prepare\_kwargs*)

Energy of calculator 1.

**get\_forces**(*atoms*, *coords*, \*\**prepare\_kwargs*)

Projected gradient for CI optimization.

**get\_hessian**(*atoms*, *coords*, \*\**prepare\_kwargs*)

Projected Hessian.

**class** pysisyphus.calculators.**DFTBp**(*parameter*, \**args*, *slakos=None*, *root=None*, \*\**kwargs*)

Bases: *OverlapCalculator*

**conf\_key** = 'dftbp'

**get\_energy**(*atoms*, *coords*, \*\**prepare\_kwargs*)

Meant to be extended.

**static get\_excited\_state\_str**(*root*, *forces=False*)

**get\_forces**(*atoms*, *coords*, \*\**prepare\_kwargs*)

Meant to be extended.

**static get\_gen\_str**(*atoms*, *coords*)

**max\_ang\_moms** = {'mio-ext': {'C': 'p', 'H': 's', 'N': 'p', 'O': 'p'}}

**parse\_energy**(*path*)

**static parse\_exc\_dat**(*text*)

**parse\_forces**(*path*)

**parse\_total\_energy**(*text*)

**prepare\_input**(*atoms*, *coords*, *calc\_type*)

Meant to be extended.

**prepare\_overlap\_data**(*path*)

Implement calculator specific parsing of MO coefficients and CI coefficients here. Should return a filename pointing to TURBOMOLE like mos, a MO coefficient array and a CI coefficient array.

**run\_calculation**(*atoms, coords, \*\*prepare\_kwargs*)

```
class pysisyphus.calculators.Dimer(calculator, *args, N_raw=None, length=0.0189,  
                                rotation_max_cycles=15, rotation_method='fourier',  
                                rotation_thresh=0.0001, rotation_tol=1, rotation_max_element=0.001,  
                                rotation_interpolate=True, rotation_disable=False,  
                                rotation_disable_pos_curv=True, rotation_remove_trans=True,  
                                trans_force_f_perp=True, bonds=None, N_hessian=None,  
                                bias_rotation=False, bias_translation=False, bias_gaussian_dot=0.1,  
                                seed=None, write_orientations=True, forward_hessian=True,  
                                **kwargs)
```

Bases: *Calculator*

**property C**

Shortcut for the curvature.

**property N**

**add\_gaussian**(*atoms, center, N, height=0.1, std=0.0529, max\_cycles=50, dot\_ref=None*)

**property can\_bias\_f0**

**property can\_bias\_f1**

**property coords0**

**property coords1**

**curvature**(*f1, f2, N*)

Curvature of the mode represented by the dimer.

**direct\_rotation**(*optimizer, prev\_step*)

**do\_dimer\_rotations**(*rotation\_thresh=None*)

**property energy0**

**property f0**

**property f1**

**property f1\_bias**

**property f2**

Never calculated explicitly, but estimated from f0 and f1.

**fourier\_rotation**(*optimizer, prev\_step*)

**get\_N\_raw\_from\_hessian**(*h5\_fn, root=0*)

**get\_forces**(*atoms, coords*)

Meant to be extended.

**get\_gaussian\_energies**(*coords, sum\_=True*)

**get\_gaussian\_forces**(*coords*, *sum\_=True*)

**get\_hessian**(*atoms*, *coords*)

Meant to be extended.

**remove\_translation**(*displacement*)

**property rot\_force**

**rotate\_coords1**(*rad*, *theta*)

Rotate dimer and produce new coords1.

**set\_N\_raw**(*coords*)

**property should\_bias\_f0**

May lead to calculation of f0 and/or f1 if present!

**property should\_bias\_f1**

May lead to calculation of f0 and/or f1 if present!

**update\_orientation**(*coords*)

**class** pysisyphus.calculators.**Dummy**(*calc\_number=0*, *charge=0*, *mult=1*, *base\_name='calculator'*, *pal=1*,  
*mem=1000*, *check\_mem=True*, *retry\_calc=1*, *last\_calc\_cycle=None*,  
*clean\_after=True*, *out\_dir='qm\_calcs'*)

Bases: [Calculator](#)

**get\_energy**(\**args*, \*\**kwargs*)

Meant to be extended.

**get\_forces**(\**args*, \*\**kwargs*)

Meant to be extended.

**get\_hessian**(\**args*, \*\**kwargs*)

Meant to be extended.

**raise\_exception**()

**run\_calculation**(\**args*, \*\**kwargs*)

**class** pysisyphus.calculators.**EGO**(*calculator*, *ref\_geom*, *max\_force=0.175*, \*\**kwargs*)

Bases: [Calculator](#)

**get\_energy**(*atoms*, *coords*, \*\**prepare\_kwargs*)

Meant to be extended.

**get\_forces**(*atoms*, *coords*, \*\**prepare\_kwargs*)

Meant to be extended.

**get\_mods**(*atoms*, *coords*)

**property ref\_hessian**

**property s**

**class** pysisyphus.calculators.**EnergyMin**(*calculator1*, *calculator2*, *mix=False*, *alpha=0.02*, *sigma=3.5*,  
*min\_energy\_diff=0.0*, *check\_after=0*, \*\**kwargs*)

Bases: [Calculator](#)

```
__init__(calculator1, calculator2, mix=False, alpha=0.02, sigma=3.5, min_energy_diff=0.0,
         check_after=0, **kwargs)
```

Use energy and derivatives of the calculator with lower energy.

This calculators carries out two calculations with different settings and returns the results of the lower energy one. This can be used to consider flips between a singlet and a triplet PES etc.

#### Parameters

- **calculator1** (*Calculator*) -- Wrapped QC calculator that provides energies and its derivatives.
- **calculator2** (*Calculator*) -- Wrapped QC calculator that provides energies and its derivatives.
- **mix** (bool, default: False) -- Enable mixing of both forces, according to the approach outlined in [2]. Can be used to optimize guesses for MECPs. Pass
- **alpha** (float, default: 0.02) -- Smoothing parameter in Hartree. See [2] for a discussion.
- **sigma** (float, default: 3.5) -- Unitless gap size parameter. The final gap becomes smaller for bigger sigmas. Has to be adapted for each case. See [2] for a discussion (p. 407 right column and p. 408 left column.)
- **min\_energy\_diff** (float, default: 0.0) -- Energy difference in Hartree. When set to a value != 0 and the energy difference between both calculators drops below this value, execution of both calculations is disabled for 'check\_after' cycles. In these cycles the calculator choice remains fixed. After 'check\_after' cycles, both energies will be calculated and it is checked, if the previous calculator choice remains valid. In conjunction with 'check\_after' both arguments can be used to save computational resources.
- **check\_after** (int, default: 0) -- Amount of cycles in which the calculator choice remains fixed.
- **\*\*kwargs** -- Keyword arguments passed to the Calculator baseclass.

```
do_calculations(name, atoms, coords, **prepare_kwargs)
```

```
get_chkfiles()
```

Return type  
dict

```
get_energy(atoms, coords, **prepare_kwargs)
```

Meant to be extended.

```
get_forces(atoms, coords, **prepare_kwargs)
```

Meant to be extended.

```
get_hessian(atoms, coords, **prepare_kwargs)
```

Meant to be extended.

```
set_chkfiles(chkfiles)
```

```
class pysisyphus.calculators.ExternalPotential(calculator=None, potentials=None, geom=None,
                                              **kwargs)
```

Bases: *Calculator*

```
available_potentials = {'harmonic_sphere': <class
'pysisyphus.calculators.ExternalPotential.HarmonicSphere'>, 'logfermi': <class
'pysisyphus.calculators.ExternalPotential.LogFermi'>, 'restraint': <class
'pysisyphus.calculators.ExternalPotential.Restraint'>}
```

```

get_energy(atoms, coords)
    Meant to be extended.
get_forces(atoms, coords)
    Meant to be extended.
get_hessian(atoms, coords)
    Meant to be extended.
get_potential_energy(coords)
get_potential_forces(coords)
class pysisyphus.calculators.FakeASE(calc)
    Bases: object
get_atoms_coords(atoms)
get_forces(atoms=None)
get_potential_energy(atoms=None)
class pysisyphus.calculators.Gaussian09(*args, **kwargs)
    Bases: Gaussian16
    conf_key = 'gaussian09'
class pysisyphus.calculators.Gaussian16(route, gbs="", gen="", keep_chk=False, stable="", fchk=None,
                                         **kwargs)
    Bases: OverlapCalculator
    conf_key = 'gaussian16'
get_chkfiles()
get_energy(atoms, coords, **prepare_kwargs)
    Meant to be extended.
get_forces(atoms, coords, **prepare_kwargs)
    Meant to be extended.
get_hessian(atoms, coords, **prepare_kwargs)
    Meant to be extended.
keep(path)
    Backup calculation results.
    Parameters
        path (Path) -- Temporary directory of the calculation.
    Returns
        kept_fns -- Dictionary holding the filenames that were backed up. The keys correspond to the
        type of file.
    Return type
        dict
make_exc_str()

```

**make\_fchk**(*path*)

**make\_gbs\_str**()

**parse\_635r\_dump**(*dump\_path*, *roots*, *nmos*)

**parse\_charges**(*path=None*)

**parse\_double\_mol**(*path*, *out\_fn=None*)

**static parse\_fchk**(*fchk\_path*, *keys*)

**parse\_force**(*path*)

**parse\_hessian**(*path*)

**parse\_keyword**(*text*)

**parse\_log**(\**args*, \*\**kwargs*)

**parse\_stable**(*path*)

**parse\_tddft**(*path*)

**prepare\_input**(*atoms*, *coords*, *calc\_type*, *did\_stable=False*, *point\_charges=None*)

Meant to be extended.

**prepare\_overlap\_data**(*path*)

Implement calculator specific parsing of MO coefficients and CI coefficients here. Should return a filename pointing to TURBOMOLE like mos, a MO coefficient array and a CI coefficient array.

**reuse\_data**(*path*)

**run\_after**(*path*)

Meant to be extended.

This method is called after a calculation was done, but before entering `self.keep()` and `self.clean()`. Can be used to call tools like `formchk` or `ricctools`.

**run\_calculation**(*atoms*, *coords*, \*\**prepare\_kwargs*)

**run\_double\_mol\_calculation**(*atoms*, *coords1*, *coords2*)

**run\_rwf\_dump**(*path*, *rwf\_index*, *chk\_path=None*)

**run\_stable**(*atoms*, *coords*, \*\**prepare\_kwargs*)

**set\_chkfiles**(*chkfiles*)

**class** pysisyphus.calculators.**HardSphere**(*geom*, *frags*, *kappa=1.0*, *permutations=False*, *frag\_radii=None*)

Bases: object

**\_\_init\_\_**(*geom*, *frags*, *kappa=1.0*, *permutations=False*, *frag\_radii=None*)

Intra-Image Inter-Molecular Hard-Sphere force.

See A.2. in [1], Eq. (A1).

**get\_forces**(*atoms*, *coords*, *kappa=None*)

```
class pysisyphus.calculators.IPIServer(*args, address=None, host=None, port=None, unlink=True,
                                       hdrlen=12, max_retries=0, verbose=False, **kwargs)
```

Bases: *Calculator*

```
get_energy(atoms, coords)
```

Meant to be extended.

```
get_forces(atoms, coords)
```

Meant to be extended.

```
get_hessian(atoms, coords)
```

Meant to be extended.

```
listen_for(atoms, coords, kind='forces')
```

```
listen_for_client_atom_num(atom_num)
```

```
listen_for_energy()
```

```
listen_for_forces(atom_num)
```

```
listen_for_hessian(atom_num)
```

```
listen_kinds = ('coords', 'energy', 'forces', 'hessian')
```

```
reset_client_connection()
```

```
retried_listen_for(atoms, coords)
```

```
unlink(address)
```

```
class pysisyphus.calculators.LennardJones(sigma=1.8897261251, epsilon=1, rc=None)
```

Bases: *Calculator*

```
calculate(coords3d)
```

```
get_energy(atoms, coords)
```

Meant to be extended.

```
get_forces(atoms, coords)
```

Meant to be extended.

```
class pysisyphus.calculators.MOPAC(method='PM7', **kwargs)
```

Bases: *Calculator*

<http://openmopac.net/manual/>

```
CALC_TYPES = {'energy': '1SCF', 'gradient': '1SCF GRADIENTS', 'hessian': 'DFORCE
FORCE LET'}
```

```
METHODS = ['am1', 'pm3', 'pm6', 'pm6-dh2', 'pm6-d3', 'pm6-dh+', 'pm6-dh2',
'pm6-dh2x', 'pm6-d3h4', 'pm6-d3h4x', 'pm7', 'pm7-ts']
```

```
MULT_STRS = {1: 'SINGLET', 2: 'DOUBLET', 3: 'TRIPLET', 4: 'QUARTET', 5:
'QUINTET', 6: 'SEXTET', 7: 'SEPTET', 8: 'OCTET'}
```

**base\_cmd**

Do only SCF AUX: Creates a checkpoint file NOREO: Dont reorient geometry

**Type**

1SCF

**conf\_key = 'mopac'**

**get\_energy**(*atoms, coords*)

Meant to be extended.

**get\_forces**(*atoms, coords*)

Meant to be extended.

**get\_hessian**(*atoms, coords*)

Meant to be extended.

**parse\_energy**(*path*)

**static parse\_energy\_from\_aux**(*inp, \*args, \*\*kwargs*)

**parse\_grad**(*path*)

**parse\_hessian**(*path*)

**static parse\_hessian\_from\_aux**(*inp, \*args, \*\*kwargs*)

**prepare\_coords**(*atoms, coords, opt=False*)

Get 3d coords in Angstrom.

Reshape internal 1d coords to 3d and convert to Angstrom.

**Parameters**

- **atoms** (*iterable*) -- Atom descriptors (element symbols).
- **coords** (*np.array, 1d*) -- 1D-array holding coordinates in Bohr.

**Returns**

**coords** -- 3D-array holding coordinates in Angstrom.

**Return type**

np.array, 3d

**prepare\_input**(*atoms, coords, calc\_type, opt=False*)

Meant to be extended.

**read\_aux**(*path*)

**class pysisyphus.calculators.MultiCalc**(*calcs, \*\*kwargs*)

Bases: [Calculator](#)

**run\_calculation**(*atoms, coords, \*\*prepare\_kwargs*)

**class pysisyphus.calculators.ONIOM**(*calcs, models, geom, layers=None, embedding="", real\_key='real', use\_link\_atoms=True, \*args, \*\*kwargs*)

Bases: [Calculator](#)

```
__init__(calcs, models, geom, layers=None, embedding='', real_key='real', use_link_atoms=True, *args,
          **kwargs)
```

**layer: list of models**

len(layer) == 1: normal ONIOM, len(layer) >= 1: multicenter ONIOM.

**model:**

(sub)set of all atoms that resides in a certain layer and has a certain calculator.

```
atom_inds_in_layer(index, exclude_inner=False)
```

Returns list of atom indices in layer at index.

Atoms that also appear in inner layer can be excluded on request.

**Parameters**

- **index** (*int*) -- pasd
- **exclude\_inner** (*bool, default=False, optional*) -- Whether to exclude atom indices that also appear in inner layers.

**Returns**

**atom\_indices** -- List containing the atom indices in the selected layer.

**Return type**

list

```
calc_layer(atoms, coords, index, parent_correction=True)
```

**property charge**

```
embeddings = {'': '', 'electronic': 'Electronic embedding', 'electronic_rc':
'Electronic embedding with redistributed charges', 'electronic_rcd': 'Electronic
embedding with redistributed charges and dipoles'}
```

```
get_energy(atoms, coords)
```

Meant to be extended.

```
get_forces(atoms, coords)
```

Meant to be extended.

```
get_hessian(atoms, coords)
```

Meant to be extended.

```
get_layer_calc(layer_ind)
```

**property model\_iter****property mult**

```
run_calculation(atoms, coords)
```

```
run_calculations(atoms, coords, method)
```

```
class pysisyphus.calculators.ORCA(keywords, blocks='', gbw=None, do_stable=False, numfreq=False,
                                   **kwargs)
```

Bases: *OverlapCalculator*

`__init__(keywords, blocks='', gbw=None, do_stable=False, numfreq=False, **kwargs)`

ORCA calculator.

Wrapper for creating ORCA input files for energy, gradient and Hessian calculations. The PAL and memory inputs must not be given in the keywords and/or blocks, as they are handled by the 'pal' and 'memory' arguments.

#### Parameters

- **keywords** (*str*) -- Keyword line, as normally given in ORCA, excluding the leading "!".
- **blocks** (*str*, *optional*) -- ORCA block input(s), e.g. for TD-DFT calculations (%tddft ... end). As the blocks start with a leading "%", wrapping the input in quotes (") is required, otherwise the parsing will fail.
- **gbw** (*str*, *optional*) -- Path to an input gbw file, which will be used as initial guess for the first calculation. Will be overridden later, with the path to the gbw file of a previous calculation.
- **do\_stable** (*bool*, *optional*) -- Run stability analysis until a stable wavefunction is obtained, before every calculation.
- **numfreq** (*boo*, *optional*) -- Use numerical frequencies instead of analytical ones.
- **mem** (*int*) -- Mememory per core in MB.

`check_termination(*args, **kwargs)`

`clean_tmp(path)`

`conf_key = 'orca'`

`get_block_str()`

`get_chkfiles()`

`get_energy(atoms, coords, **prepare_kwargs)`

Meant to be extended.

`get_forces(atoms, coords, **prepare_kwargs)`

Meant to be extended.

`get_hessian(atoms, coords, **prepare_kwargs)`

Meant to be extended.

`get_moinp_str(gbw)`

`get_stable_wavefunction(atoms, coords)`

`keep(path)`

Backup calculation results.

#### Parameters

**path** (*Path*) -- Temporary directory of the calculation.

#### Returns

**kept\_fns** -- Dictionary holding the filenames that were backed up. The keys correspond to the type of file.

#### Return type

dict

**parse\_all\_energies**(*text=None, triplets=None*)

**static parse\_atoms\_coords**(*inp, \*args, \*\*kwargs*)

**parse\_cis**(*cis*)

**Read binary CI vector file from ORCA.**

Adapted from TheoDORE 1.7.1, Authors: S. Mai, F. Plasser <https://sourceforge.net/p/theodore-qc>

**parse\_energy**(*path*)

**parse\_engrad**(*path*)

**static parse\_engrad\_info**(*inp, \*args, \*\*kwargs*)

**parse\_gbw**(*gbw\_fn*)

Adapted from <https://orcaforum.kofo.mpg.de/viewtopic.php?f=8&t=3299>

The first 5 long int values represent pointers into the file:

Pointer @+0: Internal ORCA data structures Pointer @+8: Geometry Pointer @+16: BasisSet Pointer @+24: Orbitals Pointer @+32: ECP data

**static parse\_hess\_file**(*inp, \*args, \*\*kwargs*)

**parse\_hessian**(*path*)

**parse\_mo\_numbers**(*out\_fn*)

**parse\_stable**(*path*)

**prepare\_input**(*atoms, coords, calc\_type, point\_charges=None, do\_stable=False*)

Meant to be extended.

**prepare\_overlap\_data**(*path*)

Implement calculator specific parsing of MO coefficients and CI coefficients here. Should return a filename pointing to TURBOMOLE like mos, a MO coefficient array and a CI coefficient array.

**reattach**(*last\_calc\_cycle*)

**run\_after**(*path*)

Meant to be extended.

This method is called after a calculation was done, but before entering `self.keep()` and `self.clean()`. Can be used to call tools like `formchk` or `ricctools`.

**run\_calculation**(*atoms, coords, \*\*prepare\_kwargs*)

Basically some kind of dummy method that can be called to execute ORCA with the stored cmd of this calculator.

**set\_chkfiles**(*chkfiles*)

**set\_mo\_coeffs**(*mo\_coeffs=None, gbw=None*)

**static set\_mo\_coeffs\_in\_gbw**(*in\_gbw\_fn, out\_gbw\_fn, mo\_coeffs*)

See `self.parse_gbw`.

**store\_and\_track**(*results, func, atoms, coords, \*\*prepare\_kwargs*)

```
class pysisyphus.calculators.ORCA5(keywords, blocks='', gbw=None, do_stable=False, numfreq=False,  
                                   **kwargs)
```

Bases: *ORCA*

```
conf_key = 'orca5'
```

```
class pysisyphus.calculators.OpenMolcas(basis, inporb, roots, mdrlxroot, supsym=None, track=True,  
                                         **kwargs)
```

Bases: *Calculator*

```
build_rassi_str()
```

```
build_supsym_str(supsym)
```

Can handle only one subgroup for now.

```
conf_key = 'openmolcas'
```

```
get_energy(atoms, coords)
```

Meant to be extended.

```
get_forces(atoms, coords)
```

Meant to be extended.

```
get_pal_env()
```

```
keep(path)
```

Backup calculation results.

**Parameters**

**path** (*Path*) -- Temporary directory of the calculation.

**Returns**

**kept\_fns** -- Dictionary holding the filenames that were backed up. The keys correspond to the type of file.

**Return type**

dict

```
parse_energies(text)
```

```
parse_gradient(path)
```

```
parse_rassi_track(path)
```

```
prepare_coords(atoms, coords)
```

Get 3d coords in Angstrom.

Reshape internal 1d coords to 3d and convert to Angstrom.

**Parameters**

- **atoms** (*iterable*) -- Atom descriptors (element symbols).
- **coords** (*np.array, 1d*) -- 1D-array holding coordinates in Bohr.

**Returns**

**coords** -- 3D-array holding coordinates in Angstrom.

**Return type**

np.array, 3d

**prepare\_input**(*atoms, coords*)

Meant to be extended.

**reattach**(*last\_calc\_cycle*)

**class** pysisyphus.calculators.**Psi4**(*method, basis, to\_set=None, pcm='iefpcm', solvent=None, write\_fchk=False, \*\*kwargs*)

Bases: *Calculator*

**conf\_key** = 'psi4'

**get\_energy**(*atoms, coords, \*\*prepare\_kwargs*)

Meant to be extended.

**get\_fchk\_str**()

**get\_forces**(*atoms, coords, \*\*prepare\_kwargs*)

Meant to be extended.

**get\_hessian**(*atoms, coords, \*\*prepare\_kwargs*)

Meant to be extended.

**parse\_energy**(*path*)

**parse\_grad**(*path*)

**parse\_hessian**(*path*)

**prepare\_input**(*atoms, coords, calc\_type*)

Meant to be extended.

**run\_calculation**(*atoms, coords, \*\*prepare\_kwargs*)

**class** pysisyphus.calculators.**PyPsi4**(*method, basis, \*\*kwargs*)

Bases: *Calculator*

**get\_forces**(*atoms, coords*)

Meant to be extended.

**class** pysisyphus.calculators.**PyXTB**(\**args, gfn=2, acc=None, verbosity=0, keep\_calculator=False, \*\*kwargs*)

Bases: *Calculator*

**get\_calculator**(*atoms, coords*)

**get\_energy**(*atoms, coords, \*\*prepare\_kwargs*)

Meant to be extended.

**get\_forces**(*atoms, coords, \*\*prepare\_kwargs*)

Meant to be extended.

**class** pysisyphus.calculators.**Remote**(*remote\_calc, host, prefix="", \*\*kwargs*)

Bases: *Calculator*

**get\_energy**(*atoms, coords, \*\*prepare\_kwargs*)

Meant to be extended.

**get\_forces**(*atoms, coords, \*\*prepare\_kwargs*)

Meant to be extended.

`get_hessian(atoms, coords, **prepare_kwargs)`

Meant to be extended.

`parse_results()`

`run_calculation(atoms, coords, run_func='get_energy')`

`class pysisyphus.calculators.TIP3P(rc=9.44863062728914)`

Bases: `Calculator`

Transferable Intermolecular Potential 3 Point

`aHOH = 104.52`

`calculate(coords3d)`

`charges`

`coulomb_energy = (multiple of elem. charge * multiple of elem. charge)`

`/ (distance in bohr) * 1 / (4 * pi * vacuum permittivity)`

`coulomb_prefactor` converts everything to atomic units and it is ... drum roll ... 1. from `scipy.constants`  
`import value as pcval self.coulomb_prefactor = (1 / (4 * np.pi) * pcval("elementary charge"))**2`

`/ pcval("Hartree energy") / pcval("Bohr radius") / pcval("vacuum electric permittivity")`

`)`

`coulomb(coords3d)`

`epsilon = 0.0002423919586315716`

`get_energy(atoms, coords)`

Meant to be extended.

`get_forces(atoms, coords)`

Meant to be extended.

`qH = 0.417`

`qO = -0.834`

`rOH = 1.8088458464917874`

`sigma = 5.953790025507198`

`class pysisyphus.calculators.TransTorque(frag, iter_frag, a_mats, b_mats, weight_func=None, skip=True, kappa=1.0, b_coords3d=None, do_trans=True)`

Bases: `object`

`__init__(frag, iter_frag, a_mats, b_mats, weight_func=None, skip=True, kappa=1.0, b_coords3d=None, do_trans=True)`

Translational and torque forces. See A.4. [1], Eqs. (A3) - (A5).

`get_forces(atoms, coords, kappa=None)`

`get_forces_naive(atoms, coords, kappa=None)`

`set_N_invs()`

```

class pysisyphus.calculators.Turbomole(control_path, root=None, double_mol_path=None, **kwargs)
    Bases: OverlapCalculator
    append_control(to_append, log_msg="", **kwargs)
    conf_key = 'turbomole'
    get_chkfiles()
    get_energy(atoms, coords, **prepare_kwargs)
        Meant to be extended.
    get_forces(atoms, coords, cmd=None, **prepare_kwargs)
        Meant to be extended.
    get_hessian(atoms, coords, **prepare_kwargs)
        Meant to be extended.
    get_pal_env()
    get_ricc2_root(text)
    keep(path)
        Backup calculation results.
        Parameters
            path (Path) -- Temporary directory of the calculation.
        Returns
            kept_fns -- Dictionary holding the filenames that were backed up. The keys correspond to the
            type of file.
        Return type
            dict
    make_molden(path)
    parse_cc2_vectors(ccre)
    parse_double_mol(path)
        Parse a double molecule overlap matrix from Turbomole output to be used with WFOWrapper.
    parse_energy(path)
    parse_force(path)
    parse_gs_energy()
        Several places are possible: $subenergy from control file total energy from turbomole.out Final MP2 energy
        from turbomole.out with ADC(2) Final CC2 energy from turbomole.out with CC(2)
    parse_hessian(path, fn=None)
    parse_mos()
    parse_td_vectors(text)
        For TDA calculations only the X vector is present in the ciss_a/etc. file. In TDDFT calculations there are
        twice as much items compared with TDA. The first half corresponds to (X+Y) and the second half to (X-Y).
        X can be calculated as  $X = ((X+Y)+(X-Y))/2$ . Y is then given as  $Y = (X+Y)-X$ . The normalization can then
        be checked as
        
$$\text{np.concatenate}((X, Y)).\text{dot}(\text{np.concatenate}((X, -Y)))$$


```

and should be 1.

**prepare\_input**(*atoms, coords, calc\_type, point\_charges=None*)

To rectify this we have to construct the basemcd dynamically and construct it ad hoc. We could set a RI flag in the beginning and select the correct scf binary here from it. Then we select the following binary on demand, e.g. aoforce or rdgrad or egrad etc.

**prepare\_overlap\_data**(*path*)

Implement calculator specific parsing of MO coefficients and CI coefficients here. Should return a filename pointing to TURBOMOLE like mos, a MO coefficient array and a CI coefficient array.

**prepare\_point\_charges**(*point\_charges*)

\$point\_charges <x> <y> <z> <q>

**prepare\_td**(*text*)

**run\_after**(*path*)

Meant to be extended.

This method is called after a calculation was done, but before entering `self.keep()` and `self.clean()`. Can be used to call tools like formchk or ricctools.

**run\_calculation**(*atoms, coords*)

Basically some kind of dummy method that can be called to execute Turbomole with the stored cmd of this calculator.

**run\_double\_mol\_calculation**(*atoms, coords1, coords2*)

**set\_chkfiles**(*chkfiles*)

**set\_occ\_and\_mo\_nums**(*text*)

**store\_and\_track**(*results, func, atoms, coords, \*\*prepare\_kwargs*)

**sub\_control**(*pattern, repl, log\_msg="", \*\*kwargs*)

```
class pysisyphus.calculators.XTB(gbsa="", alpb="", gfn=2, acc=1.0, etemp=None, retry_etemp=None,
                                topo=None, topo_update=None, quiet=False, **kwargs)
```

Bases: [Calculator](#)

```
__init__(gbsa="", alpb="", gfn=2, acc=1.0, etemp=None, retry_etemp=None, topo=None,
         topo_update=None, quiet=False, **kwargs)
```

XTB calculator.

Wrapper for running energy, gradient and Hessian calculations by XTB.

#### Parameters

- **gbsa** (*str, optional*) -- Solvent for GBSA calculation, by default no solvent model is used.
- **alpb** (*str, optional*) -- Solvent for ALPB calculation, by default no solvent model is used.
- **gfn** (*int or str, must be (0, 1, 2, or "ff")*) -- Hamiltonian for the XTB calculation (GFN0, GFN1, GFN2, or GFNFF).
- **acc** (*float, optional*) -- Accuracy control of the calculation, the lower the tighter several numerical thresholds are chosen.

- **topo** (*str*, *optional*) -- Path the a GFNFF-topolgy file. As setting up the topology may take some time for sizable systems, it may be desired to reuse the file.
- **topo\_update** (*int*) -- Integer controlling the update interval of the GFNFF topology update. If supplied, the topolgy will be recreated every N-th calculation.
- **mem** (*int*) -- Mememory per core in MB.
- **quiet** (*bool*, *optional*) -- Suppress creation of log files.

**static check\_termination**(*inp*, *\*args*, *\*\*kwargs*)

**conf\_key** = 'xtb'

**get\_energy**(*atoms*, *coords*, *\*\*prepare\_kwargs*)

Meant to be extended.

**get\_forces**(*atoms*, *coords*, *\*\*prepare\_kwargs*)

Meant to be extended.

**get\_hessian**(*atoms*, *coords*, *\*\*prepare\_kwargs*)

Meant to be extended.

**get\_mdrestart\_str**(*coords*, *velocities*)

*coords* and *velocities* have to given in au!

**get\_pal\_env**()

**get\_retry\_args**()

**keep**(*path*)

Backup calculation results.

#### Parameters

**path** (*Path*) -- Temporary directory of the calculation.

#### Returns

**kept\_fns** -- Dictionary holding the filenames that were backed up. The keys correspond to the type of file.

#### Return type

dict

**parse\_charges**(*fn=None*)

**parse\_charges\_from\_json**(*fn=None*)

**parse\_energy**(*path*)

**parse\_gradient**(*path*)

**parse\_hessian**(*path*)

**parse\_md**(*path*)

**parse\_opt**(*path*, *keep\_log=False*)

**parse\_topo**(*path*)

**prepare\_add\_args**(*xcontrol=None*)

**prepare\_coords**(*atoms, coords*)

Get 3d coords in Angstrom.

Reshape internal 1d coords to 3d and convert to Angstrom.

**Parameters**

- **atoms** (*iterable*) -- Atom descriptors (element symbols).
- **coords** (*np.array, 1d*) -- 1D-array holding coordinates in Bohr.

**Returns**

**coords** -- 3D-array holding coordinates in Angstrom.

**Return type**

np.array, 3d

**prepare\_input**(*atoms, coords, calc\_type, point\_charges=None*)

Meant to be extended.

**reattach**(*last\_calc\_cycle*)

**run\_calculation**(*atoms, coords, \*\*prepare\_kwargs*)

**run\_md**(*atoms, coords, t, dt, velocities=None, dump=1*)

Expecting t and dt in fs, even though xtb wants t in ps!

**run\_opt**(*atoms, coords, keep=True, keep\_log=False*)

**run\_topo**(*atoms, coords*)

**write\_mdrestart**(*path, mdrestart\_str*)

### 16.1.1.3 pysisyphus.cos package

#### Submodules

#### pysisyphus.cos.AdaptiveNEB module

**class** pysisyphus.cos.AdaptiveNEB.**AdaptiveNEB**(*images, adapt=True, adapt\_fact=0.25, adapt\_between=1, scale\_fact=False, keep\_hei=True, free\_ends=True, \*\*kwargs*)

Bases: [NEB](#)

**\_\_init\_\_**(*images, adapt=True, adapt\_fact=0.25, adapt\_between=1, scale\_fact=False, keep\_hei=True, free\_ends=True, \*\*kwargs*)

(Free-End) Adaptive Nudged Elastic Band.

**Parameters**

- **images** (*list of Geometry objects*) -- Images of the band.
- **adapt** (*bool, default True*) -- Whether to adapt the image number or not. This switch is included to support the FreeEndNEB class, that is just a thin wrapper around this class.
- **adapt\_fact** (*positive float*) -- Factor that is used to decide whether to adapt. The initial threshold is calculated by multiplying the RMS force of the band with this factor. When the RMS of the force falls below this threshold adaption takes place.

- **adapt\_between** (*positive integer*) -- Number of images to interpolate between the highest energy image and its neighbours. The number of starting images must be higher or equal then  $2 \cdot \text{adapt\_between} + 3$ , as we reuse/transfer the calculators from the starting images onto the new ones.
- **scale\_fact** (*bool, default False*) -- Whether to increase `adapt_fact` in deeper levels. This may lead to earlier adaption.
- **keep\_hei** (*bool, optional*) -- Whether to keep the highest energy image (usually a very good idea) or to interpolate only between the neighbouring images.
- **free\_ends** (*bool, default True*) -- Whether to use modified forces on the end images.

**adapt\_this\_cycle**(*forces*)

Decide wether to adapt.

**Parameters**

**forces** (*np.array*) -- Forces of the previous optimization cycle.

**Returns**

**adapt** -- Flag that indicates if adaption should take place in this cycle.

**Return type**

bool

**property forces**

See Eq. (7) in [2].

**prepare\_opt\_cycle**(\*args, \*\*kwargs)

Check for adaption and adapt if needed.

See `ChainOfStates.prepare_opt_cycle` for a complete docstring.

**update\_adapt\_thresh**(*forces*)

Update the adaption threshold.

**Parameters**

**forces** (*np.array*) -- Forces of the previous optimization cycle.

## pysisyphus.cos.ChainOfStates module

```
class pysisyphus.cos.ChainOfStates.ChainOfStates(images, fix_ends=False, fix_first=True,
                                                fix_last=True, climb=False, climb_rms=0.005,
                                                climb_lanczos=False, climb_lanczos_rms=0.005,
                                                climb_fixed=True, energy_min_mix=False,
                                                scheduler=None, progress=False)
```

Bases: object

**as\_xyz**(*comments=None*)

**property atoms**

**calculate\_forces**()

**property calculator**

**property cart\_coords**

Return a flat 1d array containing the cartesian coordinates of all images.

`check_for_climbing_start(ref_rms)`

`clear()`

**property coords**

Return a flat 1d array containing the coordinates of all images.

**property coords3d**

`describe()`

**property energy**

**property forces**

`get_climbing_forces(ind)`

`get_climbing_indices()`

`get_dask_client()`

`get_fixed_indices()`

`get_hei_index(energies=None)`

Return index of highest energy image.

`get_image_calc_counter_sum()`

`get_perpendicular_forces(i)`

[1] Eq. 12

`get_splined_hei()`

`get_tangent(i, kind='upwinding', lanczos_guess=None, disable_lanczos=False)`

[1] Equations (8) - (11)

`get_tangents()`

**property gradient**

**property image\_inds**

**property last\_index**

`log(message)`

`logger = <Logger cos (DEBUG)>`

**property masses\_rep**

**property max\_image\_num**

**property moving\_images**

**property moving\_indices**

Returns the indices of the images that aren't fixed and can be optimized.

`par_image_calc(image)`

**property perpendicular\_forces**

**prepare\_opt\_cycle**(*last\_coords*, *last\_energies*, *last\_forces*)

Implements additional logic in preparation of the next optimization cycle.

Should be called by the optimizer at the beginning of a new optimization cycle. Can be used to implement additional logic as needed for AdaptiveNEB etc.

### property results

**rms**(*arr*)

Root mean square

Returns the root mean square of the given array.

#### Parameters

**arr** (*iterable of numbers*) --

#### Returns

**rms** -- Root mean square of the given array.

#### Return type

float

**set\_climbing\_forces**(*forces*)

**set\_coords\_at**(*i*, *coords*)

Called from helpers.procrustes with cartesian coordinates. Then tries to set cartesian coordinate as self.images[i].coords which will raise an error when coord\_type != "cart".

**set\_images**(*indices*, *images*)

**set\_vector**(*name*, *vector*, *clear=False*)

**set\_zero\_forces\_for\_fixed\_images**()

This is always done in cartesian coordinates, independent of the actual coord\_type of the images as setting forces only work with cartesian forces.

**valid\_coord\_types** = ('cart', 'cartesian', 'dlc')

**zero\_fixed\_vector**(*vector*)

## pysisyphus.cos.FreeEndNEB module

**class** pysisyphus.cos.FreeEndNEB.**FreeEndNEB**(\*args, fix\_first=False, fix\_last=False, \*\*kwargs)

Bases: *AdaptiveNEB*

**\_\_init\_\_**(\*args, fix\_first=False, fix\_last=False, \*\*kwargs)

Simple Free-End-NEB method.

Derived from AdaptiveNEB with disabled adaptation. Only implements Eq. (7) from [2]. For other implementations please see the commit 01bc8812ca6f1cd3645d43e0337d9e3c5fb0ba55. There the other variants are present but I think Eq. (7) in [2] is the simplest & best bet.

### pysisyphus.cos.FreezingString module

```
class pysisyphus.cos.FreezingString.FreezingString(images, calc_getter, max_nodes=10,  
                                                    opt_steps=3)
```

Bases: object

property allcoords

as\_xyz()

property coords

property energy

property forces

property fully\_grown

get\_new\_image(coords, index)

get\_tangent()

property left\_frontier

reparametrize()

property right\_frontier

set\_new\_frontier\_nodes()

### pysisyphus.cos.GrowingChainOfStates module

```
class pysisyphus.cos.GrowingChainOfStates.GrowingChainOfStates(images, calc_getter,  
                                                                max_nodes=10, **kwargs)
```

Bases: *ChainOfStates*

property arc\_dims

get\_new\_image\_from\_coords(coords, index)

property max\_image\_num

new\_node\_coords(k)

prepare\_opt\_cycle(\*args, \*\*kwargs)

Implements additional logic in preparation of the next optimization cycle.

Should be called by the optimizer at the beginning of a new optimization cycle. Can be used to implement additional logic as needed for AdaptiveNEB etc.

set\_new\_node(k)

**pysisyphus.cos.GrowingNT module**

```
class pysisyphus.cos.GrowingNT.GrowingNT(geom, step_len=0.5, rms_thresh=0.0017, r=None,
final_geom=None, between=None, bonds=None,
r_update=True, r_update_thresh=1.0, stop_after_ts=False,
require_imag_freq=0.0, hessian_at_ts=False, out_dir='.',
dump=True)
```

Bases: object

**property** P

Projector that keeps perpendicular component.

**as\_xyz**()

**property** atoms

**calc\_hessian\_for**(*other\_geom*)

**property** cart\_coords

**property** cart\_forces

**check\_convergence**(\*args, \*\*kwargs)

**clear\_passed**()

**property** coords

**property** energy

**property** forces

**get\_additional\_print**()

**get\_energy\_and\_forces\_at**(*coords*)

**get\_energy\_at**(*coords*)

**get\_path**(*fn*)

**static get\_r**(*geom, final\_geom, bonds, r*)

**grow\_image**()

**initialize**()

**log**(*message*)

**logger** = <Logger cos (DEBUG)>

**property** r

Parallel/search direction.

**reparametrize**()

Check if GNT can be grown.

**pysisyphus.cos.GrowingString module**

```
class pysisyphus.cos.GrowingString.GrowingString(images, calc_getter, perp_thresh=0.05,  
param='equi', reparam_every=2,  
reparam_every_full=3, reparam_tol=None,  
reparam_check='rms', max_micro_cycles=5,  
reset_dlc=True, climb=False, **kwargs)
```

Bases: *GrowingChainOfStates*

**property forces**

**property full\_string\_image\_inds**

**property fully\_grown**

Returns whether the string is fully grown. Don't count the first and last node.

**get\_additional\_print()**

**get\_cur\_param\_density**(kind=None)

**get\_new\_image**(ref\_index)

Get new image by taking a step from self.images[ref\_index] towards the center of the string.

**get\_tangent**(i)

[1] Equations (8) - (11)

**property image\_inds**

**property left\_size**

**property lf\_ind**

Index of the left frontier node in self.images.

**property nodes\_missing**

Returns the number of nodes to be grown.

**reparam\_cart**(desired\_param\_density)

**reparam\_dlc**(desired\_param\_density, thresh=0.001)

**reparametrize**()

**reset\_geometries**(ref\_geometry)

**property rf\_ind**

Index of the right frontier node in self.images.

**property right\_size**

**set\_coords**(image, coords)

**spline**(tangents=False)

**property string\_size**

## pysisyphus.cos.NEB module

```
class pysisyphus.cos.NEB.NEB(images, variable_springs=False, k_max=0.3, k_min=0.1,
                             perp_spring_forces=None, **kwargs)
```

Bases: *ChainOfStates*

**fmt\_k()**

**property forces**

**get\_parallel\_forces(*i*)**

**get\_quenched\_dneb\_forces(*i*)**

See [3], Sec. VI and [4] Sec. D.

**get\_spring\_forces(*i*)**

**property parallel\_forces**

**set\_variable\_springs()**

**update\_springs()**

## pysisyphus.cos.SimpleZTS module

```
class pysisyphus.cos.SimpleZTS.SimpleZTS(images, param='equal', **kwargs)
```

Bases: *ChainOfStates*

**reparametrize()**

## Module contents

### 16.1.1.4 pysisyphus.db package

#### Submodules

#### pysisyphus.db.generate\_db module

pysisyphus.db.generate\_db.**clean()**

pysisyphus.db.generate\_db.**generate\_db()**

pysisyphus.db.generate\_db.**parse\_args(*args*)**

pysisyphus.db.generate\_db.**run()**

### pysisyphus.db.helpers module

pysisyphus.db.helpers.**full\_name**(*molecule\_name*, *level\_name*)

pysisyphus.db.helpers.**get\_molecules\_levels**()

pysisyphus.db.helpers.**get\_path**(*molecule\_name*, *level\_name*)

### pysisyphus.db.level module

### pysisyphus.db.molecules module

**class** pysisyphus.db.molecules.**Molecule**(*name*, *fn*, *charge*, *mult*, *density*, *resname*)

Bases: tuple

**charge**

Alias for field number 2

**density**

Alias for field number 4

**fn**

Alias for field number 1

**mult**

Alias for field number 3

**name**

Alias for field number 0

**resname**

Alias for field number 5

### Module contents

#### 16.1.1.5 pysisyphus.drivers package

##### Submodules

### pysisyphus.drivers.afir module

pysisyphus.drivers.afir.**analyze\_afir\_path**(*energies*)

pysisyphus.drivers.afir.**automatic\_fragmentation**(*atoms*, *coords3d*, *frag1*, *frag2*, *cycles=2*, *p=6*,  
*bond\_factor=1.25*)

Automatic fragmentation scheme as described in SC-AFIR paper [1].

pysisyphus.drivers.afir.**coordinates\_similar**(*test\_coords3d*, *ref\_coords3d*, *rmsd\_thresh=0.01*)

**Return type**

Tuple[bool, int]

pysisyphus.drivers.afir.**decrease\_distance**(*coords3d*, *m*, *n*, *frac=0.9*)

`pysisyphus.drivers.afir.determine_target_pairs(atoms, coords3d, min_=1.25, max_=5.0)`

Determine possible target m, n atom pairs for SC-AFIR calculations.

**Return type**

List[Tuple[int]]

`pysisyphus.drivers.afir.determine_target_pairs_for_geom(geom, **kwargs)`

Determine possible target m, n atom pairs for SC-AFIR calculations from geom.

**Return type**

List[Tuple[int]]

`pysisyphus.drivers.afir.find_candidates(center, bond_sets)`

`pysisyphus.drivers.afir.generate_random_union(geoms, offset=1.0, copy=True)`

`pysisyphus.drivers.afir.geom_similar(test_geom, ref_geoms, **kwargs)`

**Return type**

bool

`pysisyphus.drivers.afir.lstsqsWithReference(coords3d, ref_coords3d, freeze_atoms=None)`

Least-squares w.r.t. reference coordinates while keeping some atoms frozen.

`pysisyphus.drivers.afir.multicomponent_afir(geom, calc_getter, fragments, gamma, out_dir=None)`

`pysisyphus.drivers.afir.prepare_single_component_afir(geom, m, n, calc_getter, afir_kwargs)`

Create perturbed geometry, determine fragments and set AFIR calculator.

`pysisyphus.drivers.afir.relax_afir_path(atoms, cart_coords, calc_getter, images=15, out_dir=None)`

`pysisyphus.drivers.afir.run_mc_afir_paths(geoms, calc_getter, num=5, gamma=None, t=None, T=298.15)`

`pysisyphus.drivers.afir.weight_function(atoms, coords3d, i, j, p=6)`

### pysisyphus.drivers.barriers module

`pysisyphus.drivers.barriers.do_endopt_ts_barriers(ts_geom, left_geoms, right_geoms=None, left_fns=None, right_fns=None, do_thermo=False, T=298.15, p=101325, calc_getter=None, solv_calc_getter=None, do_standard_state_corr=False)`

### pysisyphus.drivers.birkholz module

`pysisyphus.drivers.birkholz.birkholz_interpolation(geoms, calc_getter, recreate=True)`

`pysisyphus.drivers.birkholz.bond_order(r, r0, b=2)`

Bond order for given bond length and reference length.

Eq. (3) in [1].

`pysisyphus.drivers.birkholz.bond_orders(coords3d, bond_indices, r0s)`

List of bond orders.

`pysisyphus.drivers.birkholz.bond_orders_for_geom(geom, bond_indices)`

Wrapper for `bond_orders` for simple use with Geometry.

`pysisyphus.drivers.birkholz.get_r0s(geom, bond_indices)`

Reference bond lengths as sum of covalent radii.

`pysisyphus.drivers.birkholz.length_for_bond_order(bo, r0, b=2)`

Return bond length for given bond order and reference length.

Eq. (3) in [1].

## pysisyphus.drivers.merge module

`pysisyphus.drivers.merge.align_on_subset(geom1, union, del1=None)`

Align 'union' onto subset of 'geom1'

`pysisyphus.drivers.merge.hardsphere_merge(geom1, geom2)`

`pysisyphus.drivers.merge.merge_geoms(geom1, geom2, geom1_del=None, geom2_del=None,  
make_bonds=None)`

Merge `geom1` and `geom2` while keeping the original coordinates.

Supports deleting certain atoms.

`pysisyphus.drivers.merge.merge_opt(union, bond_diff, ff='uff')`

Fragment merging along given bond by forcefield optimization.

`pysisyphus.drivers.merge.merge_with_frozen_geom(frozen_geom, lig_geom, make_bonds, frozen_del,  
lig_del)`

`pysisyphus.drivers.merge.parse_args(args)`

`pysisyphus.drivers.merge.prepare_merge(geom1, bond_diff, geom2=None, del1=None, del2=None,  
dump=False)`

`pysisyphus.drivers.merge.run_merge()`

## pysisyphus.drivers.opt module

`class pysisyphus.drivers.opt.OptResult(opt, geom, fn)`

Bases: object

**fn:** Path

**geom:** Geometry

**opt:** Optimizer

`pysisyphus.drivers.opt.opt_davidson(opt, tsopt=True, res_rms_thresh=0.0001)`

`pysisyphus.drivers.opt.run_opt(geom, calc_getter, opt_key, opt_kwargs=None, iterative=False,  
iterative_max_cycles=5, iterative_thresh=-15, iterative_scale=2.0,  
cart_hessian=None, print_thermo=False, title='Optimization',  
copy_final_geom=None, level=0)`

## pysisyphus.drivers.perf module

pysisyphus.drivers.perf.**print\_perf\_results**(*results*)

pysisyphus.drivers.perf.**run\_perf**(*geom*, *calc\_getter*, *pals=None*, *mems=None*, *pal\_range=None*,  
*mem\_range=None*, *repeat=1*, *kind='forces'*)

## pysisyphus.drivers.pka module

pysisyphus.drivers.pka.**G\_aq\_from\_h5\_hessian**(*h5\_hessian*, *solvent*, *T=298.15*, *p=101325*)

pysisyphus.drivers.pka.**direct\_cycle**(*acid\_h5*, *base\_h5*, *acid\_solvent*, *base\_solvent*, *G\_aq\_H=None*,  
*G\_gas\_H=- 6.28*, *dG\_solvent\_H=- 265.9*, *T=298.15*, *p=101325*)

## pysisyphus.drivers.precon\_pos\_rot module

prp a901cdfacc579eb63b193cbc9043212e8b57746f pysis 340ab6105ac4156f0613b4d0e8f080d9f195530c do\_trans  
accidentally disabled in transtorque

**class** pysisyphus.drivers.precon\_pos\_rot.**SteepestDescent**(*geom*, *max\_cycles=1000*, *max\_step=0.05*,  
*rms\_force=0.05*, *rms\_force\_only=True*,  
*prefix=None*, *dump=False*, *print\_mod=25*)

Bases: object

**run**()

pysisyphus.drivers.precon\_pos\_rot.**center\_fragments**(*frag\_list*, *geom*)

pysisyphus.drivers.precon\_pos\_rot.**form\_A**(*frags*, *which\_frag*, *formed\_bonds*)

Construct the A-matrices.

AR[(m, n)] (AP[(m, n)]) contains the subset of atoms in Rm (Pm) that forms bonds with Rn (Pn).

pysisyphus.drivers.precon\_pos\_rot.**get\_fragments\_and\_bonds**(*geoms*)

pysisyphus.drivers.precon\_pos\_rot.**get\_rot\_mat**(*coords3d\_1*, *coords3d\_2*, *center=False*)

pysisyphus.drivers.precon\_pos\_rot.**get\_steps\_to\_active\_atom\_mean**(*frag\_lists*, *iter\_frag\_lists*,  
*ind\_dict*, *coords3d*, *skip=True*)

pysisyphus.drivers.precon\_pos\_rot.**get\_which\_frag**(*frags*)

pysisyphus.drivers.precon\_pos\_rot.**precon\_pos\_rot**(*reactants*, *products*, *prefix=None*,  
*config={'s2\_hs\_kappa': 1.0, 's4\_hs\_kappa': 50.0,*  
*'s4\_v\_kappa': 1.0, 's4\_w\_kappa': 1.0,*  
*'s5\_hs\_kappa': 10.0, 's5\_rms\_force': 0.01,*  
*'s5\_trans': True, 's5\_v\_kappa': 1.0, 's5\_w\_kappa':*  
*3.0, 's5\_z\_kappa': 2.0}*)

pysisyphus.drivers.precon\_pos\_rot.**report\_frags**(*rgeom*, *pgeom*, *rfrags*, *pfrags*, *rbond\_diff*, *pbond\_diff*)

pysisyphus.drivers.precon\_pos\_rot.**report\_mats**(*name*, *mats*)

pysisyphus.drivers.precon\_pos\_rot.**run\_precontr**(*reactant\_geom*, *product\_geom*, *\*\*kwargs*)

**pysisyphus.drivers.rates module**

```
class pysisyphus.drivers.rates.ReactionRates(from_, barrier, barrier_si, temperature,  
                                             imag_wavenumber, imag_frequency, rate_eyring,  
                                             kappa_eyring, rate_wigner=None, kappa_wigner=None,  
                                             rate_bell=None, kappa_bell=None, rate_eckart=None,  
                                             kappa_eckart=None)
```

Bases: object

**barrier:** float

**barrier\_si:** float

**from\_:** str

**imag\_frequency:** float

**imag\_wavenumber:** float

**kappa\_bell:** Optional[float] = None

**kappa\_eckart:** Optional[float] = None

**kappa\_eyring:** float

**kappa\_wigner:** Optional[float] = None

**rate\_bell:** Optional[float] = None

**rate\_eckart:** Optional[float] = None

**rate\_eyring:** float

**rate\_wigner:** Optional[float] = None

**temperature:** float

**pysisyphus.drivers.rates.bell\_corr**(*temperature*, *imag\_frequency*)

Tunneling correction according to Bell.

See <https://onlinelibrary.wiley.com/doi/10.1002/anie.201708489> eq. (1) and eq. (2).

**Parameters**

- **temperature** (float) -- Temperature in Kelvin.
- **imag\_frequency** (float) -- Imaginary frequency in 1/s.

**Returns**

Unitless tunneling correction according to Bell. Negative kappas are meaningless.

**Return type**

kappa

**pysisyphus.drivers.rates.eckart\_corr**(*fw\_barrier\_height*, *bw\_barrier\_height*, *temperature*,  
 *imag\_frequency*)

Tunneling correction according to Eckart.

See [3], [4] and [5]. The correction should be independent of the order *fw\_barrier\_height/bw\_barrier\_height*.

**Parameters**

- **fw\_barrier\_height** (float) -- Barrier height in forward direction in Hartree.
- **bw\_barrier\_height** (float) -- Barrier height in backward direction in Hartree.
- **temperature** (float) -- Temperature in Kelvin.
- **imag\_frequency** (float) -- Frequency in 1/s of the imaginary mode at the TS.

**Returns**

Unitless tunneling correction according to Eckart.

**Return type**

kappa

`pysisyphus.drivers.rates.eckart_corr_brown`(*fw\_barrier\_height*, *bw\_barrier\_height*, *temperature*, *imag\_frequency*)

Tunneling correction according to Eckart.

Wrapper for the TUNL subroutine as given in the appendix of [5].

**Parameters**

- **fw\_barrier\_height** (float) -- Barrier height in forward direction in Hartree.
- **bw\_barrier\_height** (float) -- Barrier height in backward direction in Hartree.
- **temperature** (float) -- Temperature in Kelvin.
- **imag\_frequency** (float) -- Frequency in 1/s of the imaginary mode at the TS.

**Returns**

Unitless tunneling correction according to Eckart.

**Return type**

kappa

`pysisyphus.drivers.rates.eyring_rate`(*barrier\_height*, *temperature*, *trans\_coeff*=1.0)

Rate constant in 1/s from the Eyring equation.

See <https://pubs.acs.org/doi/10.1021/acs.organomet.8b00456> "Reaction Rates and Barriers" on p. 3234 and eq. (8).

**Parameters**

- **barrier\_height** (float) -- Barrier height (energy, enthalpy, gibbs energy, ...) in Hartree.
- **temperature** (Union[\_SupportsArray[dtype], \_NestedSequence[\_SupportsArray[dtype]], bool, int, float, complex, str, bytes, \_NestedSequence[Union[bool, int, float, complex, str, bytes]]]) -- Temperature in Kelvin.
- **trans\_coeff** (float, default: 1.0) -- Unitless transmission coefficient, e.g., obtained from Wigner or Eckart correction.

**Returns**

Eyring reaction rate in 1/s.

**Return type**

rate

`pysisyphus.drivers.rates.get_rates`(*temperature*, *reactant\_thermos*, *ts\_thermo*, *product\_thermos*=None)

`pysisyphus.drivers.rates.get_rates_for_geoms`(*temperature*, *reactant\_geoms*, *ts\_geom*, *product\_geoms*)

`pysisyphus.drivers.rates.get_rates_for_hessians`(*temperature*, *reactant\_h5s*, *ts\_h5*, *product\_h5s*)

`pysisyphus.drivers.rates.harmonic_tst_rate(barrier_height, temperature, rs_part_func, ts_part_func, trans_coeff=1.0)`

Rate constant in 1/s from harmonic TST.

See <http://dx.doi.org/10.18419/opus-9841>, chapter 5. Contrary to the Eyring rate this function does only takes a scalar temperature as the partition functions are also functions of the temperature and would have to be recalculated for different temperatures.

A possible extension would be to also support multiple rs/ts partition functions, one for each temperature.

#### Parameters

- **barrier\_height** (float) -- Barrier height (energy, enthalpy, gibbs energy, ...) in Hartree.
- **rs\_part\_func** (float) -- Partition function of the reactant state.
- **ts\_part\_func** (float) -- Partition function of the transition state.
- **temperature** (float) -- Temperature in Kelvin.
- **trans\_coeff** (float, default: 1.0) -- Unitless transmission coefficient, e.g., obtained from Wigner or Eckart correction.

#### Returns

HTST reaction rate in 1/s.

#### Return type

rate

`pysisyphus.drivers.rates.render_rx_rates(rx_rates)`

#### Return type

str

`pysisyphus.drivers.rates.tunl(alpha1, alpha2, U, strict=False)`

Eckart correction factor for rate constants according to Brown.

Python adaption of the TUNL subroutine in 4. Appendix of [5].

#### Parameters

- **alph1** (float) -- Unitless barrier height descriptor.  $2 V_1 / (h \nu^*)$ ; see (2) in [5].
- **alph2** (float) -- Unitless barrier height descriptor.  $2 V_2 / (h \nu^*)$ ; see (2) in [5].
- **u** -- Unitless curvature descriptor.  $h \nu^* / kT$ ; see (2) in [5].
- **strict** (bool, default: False) -- If enabled, arguments are bound checked. Will raise `AssertionError` if they are out of bounds. TUNL was found to yield accurate results when the arguments are within bounds.

#### Returns

Unitless tunneling correction according to Eckart.

#### Return type

G

`pysisyphus.drivers.rates.wigner_corr(temperature, imag_frequency)`

Tunneling correction according to Wigner.

See <https://doi.org/10.1002/qua.25686> eq. (12) and [https://doi.org/10.1007/978-3-642-59033-7\\_9](https://doi.org/10.1007/978-3-642-59033-7_9) for the original publication.

#### Parameters

- **temperature** (float) -- Temperature in Kelvin.
- **imag\_frequency** (float) -- Imaginary frequency in 1/s.

**Returns**

Unitless tunneling correction according to Wigner.

**Return type**

kappa

**pysisyphus.drivers.replace module**

`pysisyphus.drivers.replace.get_bond_subgeom(geom, ind, invert=False)`

`pysisyphus.drivers.replace.normalize_replacements(replacements)`

`pysisyphus.drivers.replace.parse_args(args)`

`pysisyphus.drivers.replace.replace_atom(geom, ind, repl_geom, repl_ind, return_full=True, opt=False, use_xtb=False, charge=0, mult=1)`

Replace atom with fragment.

`pysisyphus.drivers.replace.replace_atoms(geom, replacements, opt=False, use_xtb=False, charge=0, mult=1)`

`pysisyphus.drivers.replace.run()`

`pysisyphus.drivers.replace.run_opt(geom, freeze_inds, ff='uff', use_xtb=False, charge=0, mult=1)`

**pysisyphus.drivers.scan module**

`pysisyphus.drivers.scan.relaxed_1d_scan(geom, calc_getter, constrain_prims, start, step_size, steps, opt_key, opt_kwargs, pref=None, callback=None)`

`pysisyphus.drivers.scan.relaxed_scan(geom, calc_getter, constrain_prims, target_values, title, max_cycles=25, trust_radius=0.5, thresh=0.01, dump=True)`

Relaxed scan, allowing fixing of multiple primitive internals.

**pysisyphus.drivers.thermo module**

`pysisyphus.drivers.thermo.parse_args(args)`

`pysisyphus.drivers.thermo.run_thermo()`

## Module contents

### 16.1.1.6 pysisyphus.dynamics package

#### Submodules

#### pysisyphus.dynamics.Gaussian module

**class** pysisyphus.dynamics.Gaussian.DummyColvar

Bases: object

**eval**(*x*)

**class** pysisyphus.dynamics.Gaussian.Gaussian(*w=1, s=1, x0=None, colvar=None, dump\_name=None*)

Bases: object

**\_\_init\_\_**(*w=1, s=1, x0=None, colvar=None, dump\_name=None*)

**See:**

<https://doi.org/10.1016/j.cpc.2018.02.017>

$V(f(x)) = w * \exp(-(f(x) - f_0)**2 / (2*s**2))$

$F(x) = -dV/dx = -dV/df * df/dx$

**See also:**

<https://doi.org/10.1103/PhysRevLett.90.238302>

**calculate**(*coords, x0=None, gradient=False*)

Return potential and gradient for Gaussian(s) centered at *x0*.

**dump**(*step, s, w, center*)

**eval**(*coords, x0=None*)

**gradient**(*coords, x0=None*)

**property** *s*

**value**(*coords, x0=None*)

**property** *w*

#### pysisyphus.dynamics.colvars module

**class** pysisyphus.dynamics.colvars.CVBend(*indices, \*\*kwargs*)

Bases: *Colvar*

**value**(*c3d*)

**class** pysisyphus.dynamics.colvars.CVDistance(*indices, \*\*kwargs*)

Bases: *Colvar*

**value**(*c3d*)

**class** pysisyphus.dynamics.colvars.CVTorsion(*indices, \*\*kwargs*)

Bases: *Colvar*

**value**(*c3d*)

**class** pysisyphus.dynamics.colvars.**Colvar**(*force\_agrad=False*)

Bases: object

**eval**(*coords*)

**gradient**(*c3d*)

**abstract value**(*c3d*)

pysisyphus.dynamics.colvars.**get\_colvar**(*key, kwargs*)

## pysisyphus.dynamics.driver module

**class** pysisyphus.dynamics.driver.**MDResult**(*coords, t\_ps, step, terminated, T, E\_tot*)

Bases: tuple

**E\_tot**

Alias for field number 5

**T**

Alias for field number 4

**coords**

Alias for field number 0

**step**

Alias for field number 2

**t\_ps**

Alias for field number 1

**terminated**

Alias for field number 3

pysisyphus.dynamics.driver.**get\_data\_model**(*atoms, dump\_steps*)

pysisyphus.dynamics.driver.**md**(*geom, v0, steps, dt, remove\_com\_v=True, thermostat=None, T=298.15, timecon=100, term\_funcs=None, constraints=None, constraint\_kwargs=None, gaussians=None, verbose=True, print\_stride=50, dump\_stride=None, h5\_group\_name='run'*)

Velocity verlet integrator.

### Parameters

- **geom** (*Geometry*) -- The system for which the dynamics are to be run.
- **v0** (*np.array, floats*) -- Initial velocities in Bohr/fs.
- **steps** (*float*) -- Number of simulation steps.
- **dt** (*float*) -- Timestep in fs.
- **remove\_com\_v** (*bool, default=True*) -- Remove center-of-mass velocity.
- **thermostat** (*str, optional, default None*) -- Which and whether to use a thermostat.
- **T** (*float, optional, default=None*) -- Desired temperature in thermostated runs.

- **timecon** (*float*) -- Timeconstant of the thermostat in fs.
- **term\_funcs** (*dict, optional*) -- Iterable of functions that are called with the atomic coordinates in every MD cycle and result in termination
- **constraints** (*2d iterable, optional*) -- 2D iterable containing atom indices describing constrained bond lengths. of the MD integration when they evaluate to true.
- **constraint\_kwargs** (*dict, optional*) -- Keyword arguments for the constraint algorithm.
- **gaussians** (*list, optional, default=None*) -- List of Gaussians to be used in a metadynamics run.
- **verbose** (*bool, default=True*) -- Do additional printing when True.
- **print\_stride** (*int, default=50*) -- Report every n-th step.
- **dump\_stride** (*int, default=None*) -- If given, MD data will be dumped to a HDF5 file every n-th step.
- **str** (*h5\_group\_name =*) -- Name of the HDF5 group, used for dumping.
- **optional** -- Name of the HDF5 group, used for dumping.

### pysisyphus.dynamics.helpers module

`pysisyphus.dynamics.helpers.dump_coords(atoms, coords, trj_fn)`

`pysisyphus.dynamics.helpers.energy_forces_getter_closure(geom)`

`pysisyphus.dynamics.helpers.get_mb_velocities(masses, cart_coords, T, remove_com_v=True, remove_rot_v=True, seed=None)`

Initial velocities from Maxwell-Boltzmann distribution.

#### Parameters

- **masses** (*np.array, 1d, shape (number of atoms, )*) -- Atomic masses in amu.
- **cart\_coords** (*iterable, 1d, shape (3 \* number of atoms, )*) -- Atomic cartesian coordinates. Needed for removal of rotation.
- **T** (*float*) -- Temperature in Kelvin.
- **remove\_com\_v** (*bool, default=True, optional*) -- Whether to remove center-of-mass velocity.
- **remove\_rot\_v** (*bool, default=True, optional*) -- Whether to remove rotational velocity.
- **seed** (*int, default=None, optional*) -- Seed for the random-number-generator.

#### Returns

**v** -- Initial velocities in Bohr/fs.

#### Return type

`np.array, 2d, shape (number of atoms, 3)`

`pysisyphus.dynamics.helpers.get_mb_velocities_for_geom(geom, T, remove_com_v=True, remove_rot_v=True, seed=None)`

Initial velocities from Maxwell-Boltzmann distribution.

See 'get\_mb\_velocities' for explanation.

`pysisyphus.dynamics.helpers.kinetic_energy_for_temperature(atom_number, T, fixed_dof=0)`

Kinetic energy for given temperature and number of atoms.

Each atom has three degrees of freedom ( $1/2 * 3 == 3/2$ ).

#### Parameters

- **atom\_number** (*int*) -- Number of atoms. Each atom has three degrees of freedom.
- **T** (*float*) -- Temperature in Kelvin.
- **fixed\_dof** (*int, optional, default=0*) -- Number of fixed degrees of freedom, e.g. 3 when the center-of-mass velocity is removed.

#### Returns

**E\_kin** -- Kinetic energy in Hartree.

#### Return type

float

`pysisyphus.dynamics.helpers.kinetic_energy_from_velocities(masses, velocities)`

Kinetic energy for given velocities and masses.

#### Parameters

- **masses** (*1d array, shape (number of atoms, )*) -- Atomic masses in amu.
- **velocities** (*2d array, (number of atoms, 3)*) -- Atomic velocities in Bohr/fs.

#### Returns

**E\_kin** -- Kinetic energy in Hartree.

#### Return type

float

`pysisyphus.dynamics.helpers.remove_com_velocity(v, masses, keep_norm=True)`

Remove center-of-mass velocity.

Returned units vary with the given input units.

#### Parameters

- **v** (*np.array, 2d, shape (number of atoms, 3)*) -- Velocities.
- **masses** (*np.array, 1d, shape (number of atoms, )*) -- Atomic masses.
- **keep\_norm** (*bool, default=True*) -- Whether to rescale v to its original norm, after removal of v\_com.

#### Returns

**v** -- Velocities without center-of-mass velocity.

#### Return type

np.array

`pysisyphus.dynamics.helpers.scale_velocities_to_energy(masses, v, E_desired)`

Scale velocities to a given temperature.

#### Parameters

- **masses** (*np.array, 1d, shape (number of atoms, )*) -- Atomic masses in amu.
- **v** (*np.array, 2d, shape (number of atoms, 3)*) -- (Unscaled) velocities in Bohr/fs.
- **E\_desired** (*float*) -- Desired kinetic energy in Hartree.

**Returns**

**v** -- Scaled velocities in Bohr/fs.

**Return type**

np.array, 2d, shape (number of atoms, 3)

pysisyphus.dynamics.helpers.**scale\_velocities\_to\_temperatue**(masses, v, T\_desired, fixed\_dof=0)

Scale velocities to a given temperature.

**Parameters**

- **masses** (np.array, 1d, shape (number of atoms, )) -- Atomic masses in amu.
- **v** (np.array, 2d, shape (number of atoms, 3)) -- (Unscaled) velocities in Bohr/fs.
- **T\_desired** (float) -- Desired temperature in Kelvin.
- **fixed\_dof** (int, optional, default=0) -- Number of fixed degrees of freedom, e.g. 3 when the center-of-mass velocity is removed.

**Returns**

**v** -- Scaled velocities in Bohr/fs.

**Return type**

np.array, 2d, shape (number of atoms, 3)

pysisyphus.dynamics.helpers.**temperature\_for\_kinetic\_energy**(atom\_number, E\_kin, fixed\_dof=0)

Temperature for given kinetic energy and atom number.

Each atom has three degrees of freedom ( $1/2 * 3 == 3/2$ ).

**Parameters**

- **atom\_number** (int) -- Number of atoms. Each atom has three degrees of freedom.
- **E\_kin** (float) -- Kinetic energy in Hartree.
- **fixed\_dof** (int, optional, default=0) -- Number of fixed degrees of freedom, e.g. 3 when the center-of-mass velocity is removed.

**Returns**

**Temperature** -- Temperature in Kelvin.

**Return type**

float

pysisyphus.dynamics.helpers.**unscaled\_velocity\_distribution**(masses, T, seed=None)

$\exp(-v^2 * m / (2 * kT)) \ln() - 1/2 * v^2 * m / kT v^2 - 2 * \ln() * kT / m v \pm (-2 * \ln() * kT / m)**0.5 v \pm (-2 * k)**0.5 * (\ln() * T / m)**0.5$

The first term of the RHS is constant. As we later scale the velocities we neglect it. Don't use these velocities unscaled!

$v \pm (\ln() * T / m)**0.5$

## pysisyphus.dynamics.lincs module

`pysisyphus.dynamics.lincs.lincs_closure`(*geom*, *constraints*, *order=4*)

Drop `conn_nums` and keep `conns_` in a list of list, instead of an array. We could do everything with

```
for i, conn_constr in enumerate(conns)
    pass
```

instead of the pseudo-code way as given in the paper.

## pysisyphus.dynamics.mdp module

`pysisyphus.dynamics.mdp.MDPResult`

alias of `MDResult`

`pysisyphus.dynamics.mdp.mdp`(*geom*, *steps*, *dt*, *term\_funcs=None*, *steps\_init=None*, *E\_excess=0.0*,  
*displ\_length=0.1*, *epsilon=0.0005*, *ascent\_alpha=0.05*, *max\_ascent\_steps=25*,  
*max\_init\_trajs=10*, *dump=True*, *seed=None*, *external\_md=False*)

`pysisyphus.dynamics.mdp.parse_raw_term_func`(*raw\_term\_func*)

`pysisyphus.dynamics.mdp.parse_raw_term_funcs`(*raw\_term\_funcs*)

`pysisyphus.dynamics.mdp.run_md`(*geom*, *dt*, *steps*, *v0=None*, *term\_funcs=None*, *external=False*)

## pysisyphus.dynamics.rattle module

`pysisyphus.dynamics.rattle.rattle_closure`(*geom*, *constraints*, *dt*, *tol=0.001*, *max\_cycles=25*,  
*energy\_forces\_getter=None*, *remove\_com\_v=True*)

## pysisyphus.dynamics.thermostats module

`pysisyphus.dynamics.thermostats.berendsen_closure`(*sigma*, *dof*, *dt*, *tau=100*, *rng=None*)

<https://doi.org/10.1063/1.448118>

`pysisyphus.dynamics.thermostats.csvr_closure`(*sigma*, *dof*, *dt*, *tau=100*, *rng=None*)

### Parameters

- **sigma** (*float*) -- Target average value of the kinetic energy ( $1/2 \text{ dof } k_b T$ ) in the same units as `cur_kinetic_energy`.
- **dof** (*int*) -- Degrees of freedom.
- **tau** (*float*) -- Timeconstant of the thermostat. `tau` : float Timeconstant of the thermostat.
- **rng** (*numpy.random.Generator, optional*) -- Instances of a random number generator (RNG). If it is not provided the module-level RNG will be used.

`pysisyphus.dynamics.thermostats.csvr_closure_2`(*sigma*, *dof*, *dt*, *tau=100*, *rng=None*)

pysisyphus.dynamics.thermostats.**sum\_noises**(*num*, *rng=None*)

#### Parameters

- **num** (*int*) -- Number of independent Gaussian noises to be squared.
- **rng** (*numpy.random.Generator*, *optional*) -- Instances of a random number generator (RNG). If it is not provided the module-level RNG will be used.

## Module contents

### 16.1.1.7 pysisyphus.intcoords package

#### Submodules

##### pysisyphus.intcoords.Bend module

**class** pysisyphus.intcoords.Bend.**Bend**(*indices*, *periodic=False*, *calc\_kwargs=None*, *cache=False*)  
Bases: *Primitive*

##### pysisyphus.intcoords.Bend2 module

**class** pysisyphus.intcoords.Bend2.**Bend2**(*indices*, *periodic=False*, *calc\_kwargs=None*, *cache=False*)  
Bases: *Bend*

##### pysisyphus.intcoords.BondedFragment module

**class** pysisyphus.intcoords.BondedFragment.**BondedFragment**(*indices*, *bond\_indices*, *\*\*kwargs*)  
Bases: *Primitive*

##### pysisyphus.intcoords.Cartesian module

**class** pysisyphus.intcoords.Cartesian.**Cartesian**(\**args*, *\*\*kwargs*)  
Bases: *Primitive*

**class** pysisyphus.intcoords.Cartesian.**CartesianX**(\**args*, *\*\*kwargs*)  
Bases: *Cartesian*  
**cart\_axis** = 0

**class** pysisyphus.intcoords.Cartesian.**CartesianY**(\**args*, *\*\*kwargs*)  
Bases: *Cartesian*  
**cart\_axis** = 1

**class** pysisyphus.intcoords.Cartesian.**CartesianZ**(\**args*, *\*\*kwargs*)  
Bases: *Cartesian*  
**cart\_axis** = 2

## pysisyphus.intcoords.CartesianCoords module

```
class pysisyphus.intcoords.CartesianCoords.CartesianCoords(atoms, coords3d, masses,
                                                         freeze_atoms=None, *,
                                                         mass_weighted=False, **kwargs)
```

Bases: *CoordSys*

**property coords:** `ndarray[Any, dtype[ScalarType]]`

Getter for coordinates in this coordinate system.

**Return type**

`ndarray[Any, dtype[TypeVar(ScalarType, bound= generic, covariant=True)]]`

**property coords3d:** `ndarray[Any, dtype[ScalarType]]`

Getter for 3d Cartesian coordinates.

**Return type**

`ndarray[Any, dtype[TypeVar(ScalarType, bound= generic, covariant=True)]]`

**property inv\_masses\_rep\_sqrt:** `ndarray[Any, dtype[ScalarType]]`

**Return type**

`ndarray[Any, dtype[TypeVar(ScalarType, bound= generic, covariant=True)]]`

**property masses:** `ndarray[Any, dtype[ScalarType]]`

**Return type**

`ndarray[Any, dtype[TypeVar(ScalarType, bound= generic, covariant=True)]]`

**property masses\_sqrt:** `ndarray[Any, dtype[ScalarType]]`

**Return type**

`ndarray[Any, dtype[TypeVar(ScalarType, bound= generic, covariant=True)]]`

**project\_hessian**(*hessian*)

Project Hessian in the current coordinate system.

**transform\_forces**(*cart\_forces*)

Transform Cartesian forces to this coordinate system.

**Return type**

`ndarray[Any, dtype[TypeVar(ScalarType, bound= generic, covariant=True)]]`

**transform\_hessian**(*cart\_hessian, int\_gradient=None*)

Transform Cartesian Hessian to this coordinate system.

**transform\_int\_step**(*step, update\_constraints=False, pure=False*)

Transform step in this coordinate system to Cartesians.

**Return type**

`ndarray[Any, dtype[TypeVar(ScalarType, bound= generic, covariant=True)]]`

**property typed\_prims:** `List`

List of (primitive) internal coordinates.

May be empty, e.g., when the coordinate system is Cartesian.

**Return type**

`List`

```
class pysisyphus.intcoords.CartesianCoords.MWCartesianCoords(*args, **kwargs)
    Bases: CartesianCoords
```

### pysisyphus.intcoords.Coords module

```
class pysisyphus.intcoords.Coords.CoordSys(*args, **kwargs)
    Bases: Protocol

    abstract property coords: ndarray[Any, dtype[ScalarType]]
        Getter for coordinates in this coordinate system.

        Return type
            ndarray[Any, dtype[TypeVar(ScalarType, bound= generic, covariant=True)]]

    abstract property coords3d: ndarray[Any, dtype[ScalarType]]
        Getter for 3d Cartesian coordinates.

        Return type
            ndarray[Any, dtype[TypeVar(ScalarType, bound= generic, covariant=True)]]

    abstract project_hessian(hessian)
        Project Hessian in the current coordinate system.

        Return type
            ndarray[Any, dtype[TypeVar(ScalarType, bound= generic, covariant=True)]]

    abstract transform_forces(cart_forces)
        Transform Cartesian forces to this coordinate system.

        Return type
            ndarray[Any, dtype[TypeVar(ScalarType, bound= generic, covariant=True)]]

    abstract transform_hessian(cart_hessian, int_gradient)
        Transform Cartesian Hessian to this coordinate system.

        Return type
            ndarray[Any, dtype[TypeVar(ScalarType, bound= generic, covariant=True)]]

    abstract transform_int_step(step, update_constraints=False, pure=False)
        Transform step in this coordinate system to Cartesians.

        Return type
            ndarray[Any, dtype[TypeVar(ScalarType, bound= generic, covariant=True)]]

    abstract property typed_prims: List
        List of (primitive) internal coordinates.

        May be empty, e.g., when the coordinate system is Cartesian.

        Return type
            List
```

**pysisyphus.intcoords.DLC module**

**class** pysisyphus.intcoords.DLC.DLC(\*args, full\_set=True, \*\*kwargs)

Bases: *RedundantCoords*

**property B**

Wilson B-Matrix in the non-redundant subspace.

**property U**

**backtransform\_hessian**(\*args, \*\*kwargs)

Transform Hessian in internal coordinates to Cartesians.

**property constraints****property coords**

**freeze\_primitives**(typed\_prims)

Freeze primitive internal coordinates.

**Parameters**

**typed\_prims** (iterable of typed primitives) -- Iterable containing typed\_primitives, starting with a PrimType and followed by atom indices.

**get\_active\_set**(B, inv\_thresh=None)

See [5] between Eq. (7) and Eq. (8) for advice regarding the threshold.

**get\_constrained\_U**(constraint\_vecs)

**project\_hessian**(H)

As we already work in the non-redundant subspace we don't have to project/shift the hessian as we do it for simple redundant internal coordinates.

**project\_primitive\_on\_active\_set**(prim\_ind)

**reset\_constraints**()

**set\_active\_set**()

**transform\_hessian**(cart\_hessian, int\_gradient=None)

Transform Cartesian Hessian to DLC.

**transform\_int\_step**(step, \*args, \*\*kwargs)

As the transformation is done in primitive internal coordinates we convert the DLC back to primitive coordinates.

**class** pysisyphus.intcoords.DLC.HDLC(\*args, \*\*kwargs)

Bases: *DLC*

### pysisyphus.intcoords.DistanceFunction module

**class** pysisyphus.intcoords.DistanceFunction.**DistanceFunction**(*indices, coeff=-1, \*\*kwargs*)

Bases: *Primitive*

### pysisyphus.intcoords.DummyTorsion module

**class** pysisyphus.intcoords.DummyTorsion.**DummyTorsion**(*indices, \*args, fix\_inner=True, \*\*kwargs*)

Bases: *Torsion*

**static** **get\_coords3d\_and\_indices\_ext**(*coords3d, indices*)

**static** **get\_fourth\_coords**(*coords3d, indices, r=1.889, theta=90*)

**M N** <- **add**

**/**

**u v**

**/**

**P**

m, o, p, n = indices

### pysisyphus.intcoords.LinearBend module

**class** pysisyphus.intcoords.LinearBend.**LinearBend**(*\*args, complement=False, \*\*kwargs*)

Bases: *Primitive*

**calculate**(*coords3d, indices=None, gradient=False*)

**jacobian**(*coords3d, indices=None*)

### pysisyphus.intcoords.LinearDisplacement module

**class** pysisyphus.intcoords.LinearDisplacement.**LinearDisplacement**(*\*args, complement=False, \*\*kwargs*)

Bases: *Primitive*

**calculate**(*coords3d, indices=None, gradient=False*)

**jacobian**(*coords3d, indices=None*)

**pysisyphus.intcoords.OutOfPlane module**

**class** pysisyphus.intcoords.OutOfPlane.**OutOfPlane**(*indices, periodic=False, calc\_kwargs=None, cache=False*)

Bases: *Primitive*

[1] [https://doi.org/10.1002/\(SICI\)1096-987X\(19990730\)20:10<1067::AID-JCC9>3.0.CO;2-V](https://doi.org/10.1002/(SICI)1096-987X(19990730)20:10<1067::AID-JCC9>3.0.CO;2-V)  
Lee, 1999

**pysisyphus.intcoords.PrimTypes module**

pysisyphus.intcoords.PrimTypes.**PT**

alias of *PrimTypes*

**class** pysisyphus.intcoords.PrimTypes.**PrimTypes**(*value*)

Bases: *OrderedEnum*

An enumeration.

**AUX\_BOND** = 1

**AUX\_INTERFRAG\_BOND** = 4

**BEND** = 5

**BEND2** = 28

**BOND** = 0

**BONDED\_FRAGMENT** = 25

**CARTESIAN** = 21

**CARTESIAN\_X** = 22

**CARTESIAN\_Y** = 23

**CARTESIAN\_Z** = 24

**DISTANCE\_FUNCTION** = 27

**DUMMY\_TORSION** = 26

**HYDROGEN\_BOND** = 2

**IMPROPER\_DIHEDRAL** = 9

**INTERFRAG\_BOND** = 3

**LINEAR\_BEND** = 6

**LINEAR\_BEND\_COMPLEMENT** = 7

**LINEAR\_DISPLACEMENT** = 11

**LINEAR\_DISPLACEMENT\_COMPLEMENT** = 12

**OUT\_OF\_PLANE** = 10

PROPER\_DIHEDRAL = 8

PROPER\_DIHEDRAL2 = 29

ROTATION = 17

ROTATION\_A = 18

ROTATION\_B = 19

ROTATION\_C = 20

TRANSLATION = 13

TRANSLATION\_X = 14

TRANSLATION\_Y = 15

TRANSLATION\_Z = 16

`pysisyphus.intcoords.PrimTypes.get_bonded_frag_coord()`

`pysisyphus.intcoords.PrimTypes.get_dist_func()`

`pysisyphus.intcoords.PrimTypes.get_rot_coord(cls)`

`pysisyphus.intcoords.PrimTypes.normalize_prim_input(prim_inp)`

Normalize input for `define_prims` and `constrain_prims`

The `intcoords.RedundantCoords` constructor expects lists of integer lists (tuples) for arguments like `'define_prims'` and `'constrain_prims'`. The first item of every list determines the type of primitive coordinate. Currently there are about 20 different types and it is hard to remember all of them.

So we also allow a more human friendly input, that is normalized here. The most common primitives are:

0: BOND 5: BEND 8: PROPER\_DIHEDRAL

This function maps inputs like `["BOND", 1, 2]` to `[PrimTypes.BOND, 1, 2]` etc.

Always returns a list of tuples, as some `prim_inps` expand to multiple coordinates, e.g., XYZ or ATOM.

`pysisyphus.intcoords.PrimTypes.normalize_prim_inputs(prim_inps)`

`pysisyphus.intcoords.PrimTypes.prims_from_prim_inputs(prim_inps)`

## **pysisyphus.intcoords.Primitive module**

`class pysisyphus.intcoords.Primitive.Primitive(indices, periodic=False, calc_kwargs=None, cache=False)`

Bases: object

`calculate(coords3d, indices=None, gradient=False)`

`jacobian(coords3d, indices=None)`

`log(msg, lvl=10)`

`log_dbg(msg)`

```

static parallel(u, v, thresh=1e-06)
static rho(atoms, coords3d, indices)
set_cross_vec(coords3d, indices)
weight(atoms, coords3d, f_damping=0.12)

```

### pysisyphus.intcoords.RedundantCoords module

```

class pysisyphus.intcoords.RedundantCoords.HybridRedundantCoords(*args, **kwargs)

```

Bases: *RedundantCoords*

```

class pysisyphus.intcoords.RedundantCoords.RedundantCoords(atoms, coords3d, masses=None,
                                                             bond_factor=1.3, typed_prims=None,
                                                             define_prims=None,
                                                             constrain_prims=None,
                                                             freeze_atoms=None,
                                                             freeze_atoms_exclude=False,
                                                             define_for=None, bonds_only=False,
                                                             check_bends=True, rebuild=True,
                                                             bend_min_deg=15,
                                                             dihed_max_deg=175,
                                                             lb_min_deg=175, weighted=False,
                                                             min_weight=0.3,
                                                             svd_inv_thresh=0.000316,
                                                             recalc_B=False, tric=False,
                                                             hybrid=False, hbond_angles=False)

```

Bases: object

#### **property B**

Wilson B-Matrix

#### **property B\_inv**

Generalized inverse of the Wilson B-Matrix.

#### **property B\_inv\_prim**

Generalized inverse of the primitive Wilson B-Matrix.

#### **property B\_prim**

Wilson B-Matrix

#### **property Bt\_inv**

Transposed generalized inverse of the Wilson B-Matrix.

#### **property Bt\_inv\_prim**

Transposed generalized inverse of the primitive Wilson B-Matrix.

#### **property C**

Diagonal matrix. Entries for constraints are set to one.

#### **property P**

Projection matrix onto B. See [1] Eq. (4).

```

backtransform_hessian(redund_hessian, int_gradient=None)

```

Transform Hessian in internal coordinates to Cartesians.

`property bend_atom_indices`  
`property bend_indices`  
`property bond_atom_indices`  
`property bond_indices`  
`property bond_typed_prims`  
`property cartesian_indices`  
`clear()`  
`property constrained_indices`  
`property coords`  
`property coords3d`  
`property dihedral_atom_indices`  
`property dihedral_indices`  
`eval(coords3d, attr=None)`  
`get_K_matrix(int_gradient=None)`  
`get_index_of_typed_prim(typed_prim)`  
    Index in self.typed\_prims for the supplied typed\_prim.  
`get_prim_internals_by_indices(indices)`  
`inv_B(B)`  
`inv_Bt(B)`  
`property linear_bend_indices`  
`log(message)`  
`log_int_grad_msg(int_gradient)`  
`property outofplane_indices`  
`property prim_coords`  
`property prim_indices_set`  
`property prim_internals`  
`property primitives`  
`print_typed_prims()`  
`project_hessian(H, shift=1000)`  
    Expects a hessian in internal coordinates. See Eq. (11) in [1].  
`project_vector(vector)`  
    Project supplied vector onto range of B.

`return_inds(slice_)`

property `rotation_indices`

`set_inds_from_typed_prims(typed_prims)`

`set_primitive_indices(atoms, coords3d)`

`transform_forces(cart_forces)`

Combination of Eq. (9) and (11) in [1].

`transform_hessian(cart_hessian, int_gradient=None)`

Transform Cartesian Hessian to internal coordinates.

`transform_int_step(int_step, update_constraints=False, pure=False)`

property `translation_indices`

property `typed_prims`

`class pysisyphus.intcoords.RedundantCoords.TRIC(*args, **kwargs)`

Bases: `RedundantCoords`

## pysisyphus.intcoords.Rotation module

`class pysisyphus.intcoords.Rotation.Rotation(indices, *args, ref_coords3d, **kwargs)`

Bases: `Primitive`

See (II. Theory) in [1], Eq. (3) - (14)

`index = None`

`static to_origin(coords3d, indices)`

`class pysisyphus.intcoords.Rotation.RotationA(indices, *args, ref_coords3d, **kwargs)`

Bases: `Rotation`

`index = 0`

`class pysisyphus.intcoords.Rotation.RotationB(indices, *args, ref_coords3d, **kwargs)`

Bases: `Rotation`

`index = 1`

`class pysisyphus.intcoords.Rotation.RotationC(indices, *args, ref_coords3d, **kwargs)`

Bases: `Rotation`

`index = 2`

`pysisyphus.intcoords.Rotation.compare_to_geometric(c3d, ref_c3d, dR, dF, dqdx, dvdx, atol=1e-14)`

### pysisyphus.intcoords.Stretch module

```
class pysisyphus.intcoords.Stretch.Stretch(indices, periodic=False, calc_kwargs=None, cache=False)
    Bases: Primitive
```

### pysisyphus.intcoords.Torsion module

```
class pysisyphus.intcoords.Torsion.Torsion(indices, periodic=False, calc_kwargs=None, cache=False)
    Bases: Primitive
```

### pysisyphus.intcoords.Torsion2 module

```
class pysisyphus.intcoords.Torsion2.Torsion2(indices, periodic=False, calc_kwargs=None,
                                               cache=False)
    Bases: Torsion
```

### pysisyphus.intcoords.Translation module

```
class pysisyphus.intcoords.Translation.Translation(*args, **kwargs)
    Bases: Primitive
```

See (II. Theory) in [1], Eq. (2)

```
class pysisyphus.intcoords.Translation.TranslationX(*args, **kwargs)
    Bases: Translation
```

```
    cart_axis = 0
```

```
class pysisyphus.intcoords.Translation.TranslationY(*args, **kwargs)
    Bases: Translation
```

```
    cart_axis = 1
```

```
class pysisyphus.intcoords.Translation.TranslationZ(*args, **kwargs)
    Bases: Translation
```

```
    cart_axis = 2
```

### pysisyphus.intcoords.augment\_bonds module

```
pysisyphus.intcoords.augment_bonds.augment_bonds(geom, root=0, proj=False)
```

```
pysisyphus.intcoords.augment_bonds.find_missing_bonds_by_projection(geom, hessian,
                                                                    bond_factor=2.0,
                                                                    bond_thresh=0.35,
                                                                    concerted_thresh=0.35,
                                                                    root=0)
```

```
pysisyphus.intcoords.augment_bonds.find_missing_strong_bonds(geom, hessian, bond_factor=1.7,
                                                                thresh=0.3, root=0)
```

**pysisyphus.intcoords.derivatives module**

`pysisyphus.intcoords.derivatives.d2q_a(m0, m1, m2, o0, o1, o2, n0, n1, n2)`

Bend, 2nd derivative wrt. Cartesians

`pysisyphus.intcoords.derivatives.d2q_a2(m0, m1, m2, o0, o1, o2, n0, n1, n2)`

Bend2, 2nd derivative wrt. Cartesians

`pysisyphus.intcoords.derivatives.d2q_b(m0, m1, m2, n0, n1, n2)`

Stretch, 2nd derivative wrt. Cartesians

`pysisyphus.intcoords.derivatives.d2q_d(m0, m1, m2, o0, o1, o2, p0, p1, p2, n0, n1, n2)`

Torsion, 2nd derivative wrt. Cartesians

`pysisyphus.intcoords.derivatives.d2q_d2(m0, m1, m2, o0, o1, o2, p0, p1, p2, n0, n1, n2)`

Torsion2, 2nd derivative wrt. Cartesians

`pysisyphus.intcoords.derivatives.d2q_lb(m0, m1, m2, o0, o1, o2, n0, n1, n2, p0, p1, p2)`

Linear Bend, 2nd derivative wrt. Cartesians

`pysisyphus.intcoords.derivatives.d2q_ld(m0, m1, m2, o0, o1, o2, n0, n1, n2, p0, p1, p2)`

Linear Displacement, 2nd derivative wrt. Cartesians

`pysisyphus.intcoords.derivatives.d2q_oop(m0, m1, m2, n0, n1, n2, o0, o1, o2, p0, p1, p2)`

OutOfPlane, 2nd derivative wrt. Cartesians

`pysisyphus.intcoords.derivatives.dq_a(m0, m1, m2, o0, o1, o2, n0, n1, n2)`

Bend, first derivative wrt. Cartesians

`pysisyphus.intcoords.derivatives.dq_a2(m0, m1, m2, o0, o1, o2, n0, n1, n2)`

Bend2, first derivative wrt. Cartesians

`pysisyphus.intcoords.derivatives.dq_b(m0, m1, m2, n0, n1, n2)`

Stretch, first derivative wrt. Cartesians

`pysisyphus.intcoords.derivatives.dq_d(m0, m1, m2, o0, o1, o2, p0, p1, p2, n0, n1, n2)`

Torsion, first derivative wrt. Cartesians

`pysisyphus.intcoords.derivatives.dq_d2(m0, m1, m2, o0, o1, o2, p0, p1, p2, n0, n1, n2)`

Torsion2, first derivative wrt. Cartesians

`pysisyphus.intcoords.derivatives.dq_lb(m0, m1, m2, o0, o1, o2, n0, n1, n2, p0, p1, p2)`

Linear Bend, first derivative wrt. Cartesians

`pysisyphus.intcoords.derivatives.dq_ld(m0, m1, m2, o0, o1, o2, n0, n1, n2, p0, p1, p2)`

Linear Displacement, first derivative wrt. Cartesians

`pysisyphus.intcoords.derivatives.dq_oop(m0, m1, m2, n0, n1, n2, o0, o1, o2, p0, p1, p2)`

OutOfPlane, first derivative wrt. Cartesians

`pysisyphus.intcoords.derivatives.q_a(m0, m1, m2, o0, o1, o2, n0, n1, n2)`

Bend

`pysisyphus.intcoords.derivatives.q_a2(m0, m1, m2, o0, o1, o2, n0, n1, n2)`

Bend2

`pysisyphus.intcoords.derivatives.q_b(m0, m1, m2, n0, n1, n2)`

Stretch

`pysisyphus.intcoords.derivatives.q_d(m0, m1, m2, o0, o1, o2, p0, p1, p2, n0, n1, n2)`

Torsion

`pysisyphus.intcoords.derivatives.q_d2(m0, m1, m2, o0, o1, o2, p0, p1, p2, n0, n1, n2)`

Torsion2

`pysisyphus.intcoords.derivatives.q_lb(m0, m1, m2, o0, o1, o2, n0, n1, n2, p0, p1, p2)`

Linear Bend

`pysisyphus.intcoords.derivatives.q_ld(m0, m1, m2, o0, o1, o2, n0, n1, n2, p0, p1, p2)`

Linear Displacement

`pysisyphus.intcoords.derivatives.q_oop(m0, m1, m2, n0, n1, n2, o0, o1, o2, p0, p1, p2)`

OutOfPlane

### **pysisyphus.intcoords.eval module**

**class** `pysisyphus.intcoords.eval.PrimInternal(inds, val, grad=None)`

Bases: object

`pysisyphus.intcoords.eval.augment_primitives(missing_prims, coords3d, prim_indices, fragments)`

`pysisyphus.intcoords.eval.check_primitives(coords3d, primitives, B=None, thresh=1e-06, logger=None)`

`pysisyphus.intcoords.eval.eval_B(coords3d, primitives)`

`pysisyphus.intcoords.eval.eval_primitives(coords3d, primitives)`

### **pysisyphus.intcoords.exceptions module**

**exception** `pysisyphus.intcoords.exceptions.DifferentCoordLengthsException`

Bases: Exception

**exception** `pysisyphus.intcoords.exceptions.DifferentPrimitivesException`

Bases: Exception

**exception** `pysisyphus.intcoords.exceptions.NeedNewInternalsException(coords3d, *args, invalid_inds=None, invalid_prims=None, **kwargs)`

Bases: Exception

**exception** `pysisyphus.intcoords.exceptions.PrimitiveNotDefinedException(typed_prim, *args, **kwargs)`

Bases: Exception

**exception** `pysisyphus.intcoords.exceptions.RebuiltInternalsException(*args, typed_prims=None, **kwargs)`

Bases: Exception

## pysisyphus.intcoords.findiffs module

`pysisyphus.intcoords.findiffs.fin_diff_B`(*primitive*, *coords3d*, *delta=1e-06*)

Derivatives of a primitive internal gradient wrt its defining cartesian coordinates.

`pysisyphus.intcoords.findiffs.fin_diff_prim`(*primitive*, *coords3d*, *delta=1e-06*)

Derivatives of a primitive internal gradient wrt its defining cartesian coordinates.

## pysisyphus.intcoords.generate\_derivatives module

**class** `pysisyphus.intcoords.generate_derivatives.FuncResult`(*d0*, *d1*, *d2*, *f0*, *f1*, *f2*)

Bases: tuple

**d0**

Alias for field number 0

**d1**

Alias for field number 1

**d2**

Alias for field number 2

**f0**

Alias for field number 3

**f1**

Alias for field number 4

**f2**

Alias for field number 5

`pysisyphus.intcoords.generate_derivatives.generate_wilson`(*generate=None*, *out\_fn='derivatives.py'*,  
*use\_mpmath=False*)

`pysisyphus.intcoords.generate_derivatives.make_deriv_funcs`(*base\_expr*, *dx*, *args*, *names*, *comment*,  
*use\_mpmath=True*)

`pysisyphus.intcoords.generate_derivatives.make_py_func`(*exprs*, *args=None*, *name=None*, *comment=""*,  
*use\_mpmath=False*)

## pysisyphus.intcoords.helpers module

`pysisyphus.intcoords.helpers.form_coordinate_union`(*geom1*, *geom2*)

`pysisyphus.intcoords.helpers.get_bond_difference`(*geom1*, *geom2*, *bond\_factor=1.3*)

Return formed and broken bonds when going from *geom1* to *geom2*.

`pysisyphus.intcoords.helpers.get_bond_differences_verbose`(*geom1*, *geom2*, *bond\_factor=1.3*,  
*key1='geom1'*, *key2='geom2'*)

`pysisyphus.intcoords.helpers.get_step`(*geom*, *coords*)

`pysisyphus.intcoords.helpers.get_tangent(prim1, prim2, dihedral_inds, normalize=False)`

Normalized tangent between primitive internal coordinates.

Tangent pointing from prim1 to prim2 in primitive internal coordinates, taking into account the periodicity of dihedral angles.

**Parameters**

- **prim1** (*np.array*) -- 1d-array of primitive internal coordinates in the order (stretches, bends, dihedrals).
- **prim2** (*np.array*) -- See prim1.
- **dihedral\_inds** (*list of int*) -- Dihedral indices in prim1 and prim2.

**Returns**

**tangent** -- 1d array containing the normalized tangent pointing from prim1 to prim2.

**Return type**

`np.array`

`pysisyphus.intcoords.helpers.get_weighted_bond_mode(weighted_bonds, coords3d, remove_translation=True)`

`pysisyphus.intcoords.helpers.get_weighted_bond_mode_getter(target_weighted_bonds, bond_factor=1.2, fractional=False)`

Create input for `intcoords.helpers.get_weighted_bond_mode`.

Compared to the rest of `pysisyphus` this method uses a slightly lowered bond factor, so it is more strict regarding what is considered a bond and what not.

`pysisyphus.intcoords.helpers.interfragment_distance(frag1, frag2, coords3d)`

**Return type**

`float`

`pysisyphus.intcoords.helpers.merge_coordinate_definitions(geom1, geom2)`

`pysisyphus.intcoords.helpers.verbose_bond_difference(formed, broken, key1, key2, atoms=None)`

**pysisyphus.intcoords.mp\_derivatives module**

`pysisyphus.intcoords.mp_derivatives.d2q_a(m0, m1, m2, o0, o1, o2, n0, n1, n2)`

Bend, 2nd derivative wrt. Cartesians

`pysisyphus.intcoords.mp_derivatives.d2q_a2(m0, m1, m2, o0, o1, o2, n0, n1, n2)`

Bend2, 2nd derivative wrt. Cartesians

`pysisyphus.intcoords.mp_derivatives.d2q_b(m0, m1, m2, n0, n1, n2)`

Stretch, 2nd derivative wrt. Cartesians

`pysisyphus.intcoords.mp_derivatives.d2q_d(m0, m1, m2, o0, o1, o2, p0, p1, p2, n0, n1, n2)`

Torsion, 2nd derivative wrt. Cartesians

`pysisyphus.intcoords.mp_derivatives.d2q_d2(m0, m1, m2, o0, o1, o2, p0, p1, p2, n0, n1, n2)`

Torsion2, 2nd derivative wrt. Cartesians

`pysisyphus.intcoords.mp_derivatives.d2q_lb(m0, m1, m2, o0, o1, o2, n0, n1, n2, p0, p1, p2)`

Linear Bend, 2nd derivative wrt. Cartesians

`pysisyphus.intcoords.mp_derivatives.d2q_ld(m0, m1, m2, o0, o1, o2, n0, n1, n2, p0, p1, p2)`  
 Linear Displacement, 2nd derivative wrt. Cartesians

`pysisyphus.intcoords.mp_derivatives.d2q_oop(m0, m1, m2, n0, n1, n2, o0, o1, o2, p0, p1, p2)`  
 OutOfPlane, 2nd derivative wrt. Cartesians

`pysisyphus.intcoords.mp_derivatives.dq_a(m0, m1, m2, o0, o1, o2, n0, n1, n2)`  
 Bend, first derivative wrt. Cartesians

`pysisyphus.intcoords.mp_derivatives.dq_a2(m0, m1, m2, o0, o1, o2, n0, n1, n2)`  
 Bend2, first derivative wrt. Cartesians

`pysisyphus.intcoords.mp_derivatives.dq_b(m0, m1, m2, n0, n1, n2)`  
 Stretch, first derivative wrt. Cartesians

`pysisyphus.intcoords.mp_derivatives.dq_d(m0, m1, m2, o0, o1, o2, p0, p1, p2, n0, n1, n2)`  
 Torsion, first derivative wrt. Cartesians

`pysisyphus.intcoords.mp_derivatives.dq_d2(m0, m1, m2, o0, o1, o2, p0, p1, p2, n0, n1, n2)`  
 Torsion2, first derivative wrt. Cartesians

`pysisyphus.intcoords.mp_derivatives.dq_lb(m0, m1, m2, o0, o1, o2, n0, n1, n2, p0, p1, p2)`  
 Linear Bend, first derivative wrt. Cartesians

`pysisyphus.intcoords.mp_derivatives.dq_ld(m0, m1, m2, o0, o1, o2, n0, n1, n2, p0, p1, p2)`  
 Linear Displacement, first derivative wrt. Cartesians

`pysisyphus.intcoords.mp_derivatives.dq_oop(m0, m1, m2, n0, n1, n2, o0, o1, o2, p0, p1, p2)`  
 OutOfPlane, first derivative wrt. Cartesians

`pysisyphus.intcoords.mp_derivatives.q_a(m0, m1, m2, o0, o1, o2, n0, n1, n2)`  
 Bend

`pysisyphus.intcoords.mp_derivatives.q_a2(m0, m1, m2, o0, o1, o2, n0, n1, n2)`  
 Bend2

`pysisyphus.intcoords.mp_derivatives.q_b(m0, m1, m2, n0, n1, n2)`  
 Stretch

`pysisyphus.intcoords.mp_derivatives.q_d(m0, m1, m2, o0, o1, o2, p0, p1, p2, n0, n1, n2)`  
 Torsion

`pysisyphus.intcoords.mp_derivatives.q_d2(m0, m1, m2, o0, o1, o2, p0, p1, p2, n0, n1, n2)`  
 Torsion2

`pysisyphus.intcoords.mp_derivatives.q_lb(m0, m1, m2, o0, o1, o2, n0, n1, n2, p0, p1, p2)`  
 Linear Bend

`pysisyphus.intcoords.mp_derivatives.q_ld(m0, m1, m2, o0, o1, o2, n0, n1, n2, p0, p1, p2)`  
 Linear Displacement

`pysisyphus.intcoords.mp_derivatives.q_oop(m0, m1, m2, n0, n1, n2, o0, o1, o2, p0, p1, p2)`  
 OutOfPlane

**pysisyphus.intcoords.setup module**

```
class pysisyphus.intcoords.setup.CoordInfo(bonds, hydrogen_bonds, interfrag_bonds,  
aux_interfrag_bonds, bends, linear_bends,  
linear_bend_complements, proper_dihedrals,  
improper_dihedrals, translation_inds, rotation_inds,  
cartesian_inds, typed_prims, fragments)
```

Bases: tuple

**aux\_interfrag\_bonds**

Alias for field number 3

**bends**

Alias for field number 4

**bonds**

Alias for field number 0

**cartesian\_inds**

Alias for field number 11

**fragments**

Alias for field number 13

**hydrogen\_bonds**

Alias for field number 1

**improper\_dihedrals**

Alias for field number 8

**interfrag\_bonds**

Alias for field number 2

**linear\_bend\_complements**

Alias for field number 6

**linear\_bends**

Alias for field number 5

**proper\_dihedrals**

Alias for field number 7

**rotation\_inds**

Alias for field number 10

**translation\_inds**

Alias for field number 9

**typed\_prims**

Alias for field number 12

```
pysisyphus.intcoords.setup.connect_fragments(cdm, fragments, max_aux=3.78, aux_factor=1.3,  
logger=None)
```

Determine the smallest interfragment bond for a list of fragments and a condensed distance matrix.

```
pysisyphus.intcoords.setup.get_bend_inds(coords3d, bond_inds, min_deg, max_deg, logger=None)
```

```
pysisyphus.intcoords.setup.get_bond_mat(geom, bond_factor=1.3)
```

```
pysisyphus.intcoords.setup.get_bond_sets(atoms, coords3d, bond_factor=1.3, return_cdm=False,
                                          return_cbm=False)
```

I'm sorry, but this function does not return sets, but an int ndarray.

```
pysisyphus.intcoords.setup.get_dihedral_inds(coords3d, bond_inds, bend_inds, max_deg, logger=None)
```

```
pysisyphus.intcoords.setup.get_fragments(atoms, coords, bond_inds=None, bond_factor=1.3)
```

This misses unconnected single atoms!

```
pysisyphus.intcoords.setup.get_hydrogen_bond_inds(atoms, coords3d, bond_inds, logger=None)
```

```
pysisyphus.intcoords.setup.get_linear_bend_inds(coords3d, cbm, bends, min_deg=175, max_bonds=4,
                                                logger=None)
```

```
pysisyphus.intcoords.setup.get_pair_covalent_radii(atoms)
```

```
pysisyphus.intcoords.setup.get_primitives(coords3d, typed_prims, logger=None)
```

```
pysisyphus.intcoords.setup.setup_redundant(atoms, coords3d, factor=1.3, define_prims=None,
                                           min_deg=15, dihed_max_deg=175.0, lb_min_deg=None,
                                           lb_max_bonds=4, min_weight=None, tric=False,
                                           hybrid=False, interfrag_hbonds=True,
                                           hbond_angles=False, freeze_atoms=None, define_for=None,
                                           logger=None)
```

```
pysisyphus.intcoords.setup.setup_redundant_from_geom(geom, *args, **kwargs)
```

```
pysisyphus.intcoords.setup.sort_by_prim_type(to_sort=None)
```

### pysisyphus.intcoords.setup\_fast module

```
pysisyphus.intcoords.setup_fast.find_bends(coords3d, bonds, min_deg, max_deg, logger=None)
```

```
pysisyphus.intcoords.setup_fast.find_bonds(atoms, coords3d, covalent_radii=None, bond_factor=1.3)
```

```
pysisyphus.intcoords.setup_fast.find_bonds_bends(geom, bond_factor=1.3, min_deg=15,
                                                  max_deg=175)
```

```
pysisyphus.intcoords.setup_fast.find_bonds_for_geom(geom, bond_factor=1.3)
```

```
pysisyphus.intcoords.setup_fast.find_dihedrals(coords3d, bonds, bends, max_deg, logger=None)
```

```
pysisyphus.intcoords.setup_fast.get_bond_vec_getter(atoms, covalent_radii, bonds_for_inds,
                                                    no_bonds_with=None, bond_factor=1.3)
```

```
pysisyphus.intcoords.setup_fast.get_max_bond_dists(atoms, bond_factor, covalent_radii=None)
```

## pysisyphus.intcoords.update module

`pysisyphus.intcoords.update.correct_dihedrals`(*new\_dihedrals*, *old\_dihedrals*)

Dihedrals are periodic. Going from  $-179^\circ$  to  $179^\circ$  is not a step of  $358^\circ$ , but a step of  $2^\circ$ . By considering the actual distance of the dihedrals from the correct step can be calculated.

$$\text{dihedral step length} = \text{abs}(\text{abs}(\text{new\_dihedral}) - ) + \text{abs}(\text{abs}(\text{old\_dihedral}) - )$$

or put differently

$$\text{dihedral step length} = \text{abs}(\text{abs}(\text{new\_dihedral} - \text{old\_dihedral}) - 2^*)$$

The sign is left to be determined. Going from  $-179^\circ$  to  $179^\circ$  (roughly  $-- = 2$ ) is a counter clockwise rotation and the dihedral has to decrease below  $-$ . Going from  $179^\circ$  to  $-179^\circ$  (roughly  $-- = -2$ ) is a clockwise rotation and the dihedral increases above  $.$  So the correct sign corresponds to the negative sign of the original difference.

original difference  $2 \rightarrow$  dihedral must decrease  $\rightarrow$  sign =  $-1$  original difference  $-2 \rightarrow$  dihedral must increase  $\rightarrow$  sign =  $+1$

Overall, the old dihedral is modified by the actual step length with the correct sign.

`pysisyphus.intcoords.update.inds_from_prim_types`(*typed\_prims*, *prim\_types*)

`pysisyphus.intcoords.update.transform_int_step`(*int\_step*, *old\_cart\_coords*, *cur\_internals*, *Bt\_inv\_prim*, *primitives*, *typed\_prims=None*, *dihedral\_inds=None*, *rotation\_inds=None*, *bend\_inds=None*, *check\_dihedrals=False*, *check\_bends=False*, *bend\_min\_deg=15*, *bend\_max\_deg=175*, *freeze\_atoms=None*, *constrained\_inds=None*, *update\_constraints=False*, *cart\_rms\_thresh=1e-06*, *Bt\_inv\_prim\_getter=None*, *max\_cycles=25*, *logger=None*)

Transformation is done in primitive internals, so `int_step` must be given in primitive internals and not in DLC!

`pysisyphus.intcoords.update.update_internals`(*new\_coords3d*, *old\_internals*, *primitives*, *dihedral\_inds*, *rotation\_inds*, *bend\_inds*, *check\_dihedrals=False*, *check\_bends=False*, *bend\_min\_deg=15*, *bend\_max\_deg=175*, *rotation\_thresh=0.9*, *logger=None*)

## pysisyphus.intcoords.valid module

`pysisyphus.intcoords.valid.are_collinear`(*vec1*, *vec2*, *deg\_thresh=179.5*)

`pysisyphus.intcoords.valid.bend_valid`(*coords3d*, *indices*, *min\_deg*, *max\_deg*)

`pysisyphus.intcoords.valid.check_typed_prims`(*coords3d*, *typed\_prims*, *bend\_min\_deg*, *dihed\_max\_deg*, *lb\_min\_deg*, *logger=None*, *check\_bends=True*)

`pysisyphus.intcoords.valid.dihedral_valid`(*coords3d*, *inds*, *deg\_thresh=177.5*)

`pysisyphus.intcoords.valid.dihedrals_are_valid`(*coords3d*, *dihedral\_inds*, *logger=None*)

## Module contents

**class** pysisyphus.intcoords.**Bend**(*indices, periodic=False, calc\_kwargs=None, cache=False*)

Bases: *Primitive*

**class** pysisyphus.intcoords.**Bend2**(*indices, periodic=False, calc\_kwargs=None, cache=False*)

Bases: *Bend*

**class** pysisyphus.intcoords.**CartesianCoords**(*atoms, coords3d, masses, freeze\_atoms=None, \*, mass\_weighted=False, \*\*kwargs*)

Bases: *CoordSys*

**property** **coords**: **ndarray**[**Any**, **dtype**[**ScalarType**]]

Getter for coordinates in this coordinate system.

**Return type**

**ndarray**[**Any**, **dtype**[**TypeVar**(**ScalarType**, bound= generic, covariant=True)]]

**property** **coords3d**: **ndarray**[**Any**, **dtype**[**ScalarType**]]

Getter for 3d Cartesian coordinates.

**Return type**

**ndarray**[**Any**, **dtype**[**TypeVar**(**ScalarType**, bound= generic, covariant=True)]]

**property** **inv\_masses\_rep\_sqrt**: **ndarray**[**Any**, **dtype**[**ScalarType**]]

**Return type**

**ndarray**[**Any**, **dtype**[**TypeVar**(**ScalarType**, bound= generic, covariant=True)]]

**property** **masses**: **ndarray**[**Any**, **dtype**[**ScalarType**]]

**Return type**

**ndarray**[**Any**, **dtype**[**TypeVar**(**ScalarType**, bound= generic, covariant=True)]]

**property** **masses\_sqrt**: **ndarray**[**Any**, **dtype**[**ScalarType**]]

**Return type**

**ndarray**[**Any**, **dtype**[**TypeVar**(**ScalarType**, bound= generic, covariant=True)]]

**project\_hessian**(*hessian*)

Project Hessian in the current coordinate system.

**transform\_forces**(*cart\_forces*)

Transform Cartesian forces to this coordinate system.

**Return type**

**ndarray**[**Any**, **dtype**[**TypeVar**(**ScalarType**, bound= generic, covariant=True)]]

**transform\_hessian**(*cart\_hessian, int\_gradient=None*)

Transform Cartesian Hessian to this coordinate system.

**transform\_int\_step**(*step, update\_constraints=False, pure=False*)

Transform step in this coordinate system to Cartesians.

**Return type**

**ndarray**[**Any**, **dtype**[**TypeVar**(**ScalarType**, bound= generic, covariant=True)]]

**property typed\_prims:** List

List of (primitive) internal coordinates.

May be empty, e.g., when the coordinate system is Cartesian.

**Return type**

List

**class** pysisyphus.intcoords.CartesianX(\*args, \*\*kwargs)

Bases: *Cartesian*

**cart\_axis** = 0

**class** pysisyphus.intcoords.CartesianY(\*args, \*\*kwargs)

Bases: *Cartesian*

**cart\_axis** = 1

**class** pysisyphus.intcoords.CartesianZ(\*args, \*\*kwargs)

Bases: *Cartesian*

**cart\_axis** = 2

**class** pysisyphus.intcoords.DLC(\*args, full\_set=True, \*\*kwargs)

Bases: *RedundantCoords*

**property** B

Wilson B-Matrix in the non-redundant subspace.

**property** U

**backtransform\_hessian**(\*args, \*\*kwargs)

Transform Hessian in internal coordinates to Cartesians.

**property** constraints

**property** coords

**freeze\_primitives**(typed\_prims)

Freeze primitive internal coordinates.

**Parameters**

**typed\_prims** (iterable of typed primitives) -- Iterable containing typed\_primitives, starting with a PrimType and followed by atom indices.

**get\_active\_set**(B, inv\_thresh=None)

See [5] between Eq. (7) and Eq. (8) for advice regarding the threshold.

**get\_constrained\_U**(constraint\_vecs)

**project\_hessian**(H)

As we already work in the non-redundant subspace we don't have to project/shift the hessian as we do it for simple redundant internal coordinates.

**project\_primitive\_on\_active\_set**(prim\_ind)

**reset\_constraints**()

**set\_active\_set**()

**transform\_hessian**(*cart\_hessian*, *int\_gradient=None*)

Transform Cartesian Hessian to DLC.

**transform\_int\_step**(*step*, *\*args*, *\*\*kwargs*)

As the transformation is done in primitive internal coordinates we convert the DLC back to primitive coordinates.

**class** pysisyphus.intcoords.**DistanceFunction**(*indices*, *coeff=-1*, *\*\*kwargs*)

Bases: *Primitive*

**class** pysisyphus.intcoords.**DummyTorsion**(*indices*, *\*args*, *fix\_inner=True*, *\*\*kwargs*)

Bases: *Torsion*

**static get\_coords3d\_and\_indices\_ext**(*coords3d*, *indices*)

**static get\_fourth\_coords**(*coords3d*, *indices*, *r=1.889*, *theta=90*)

**M N <- add**

**/**

**u v**

**/**

**P**

m, o, p, n = indices

**class** pysisyphus.intcoords.**HDLC**(*\*args*, *\*\*kwargs*)

Bases: *DLC*

**class** pysisyphus.intcoords.**HybridRedundantCoords**(*\*args*, *\*\*kwargs*)

Bases: *RedundantCoords*

**class** pysisyphus.intcoords.**LinearBend**(*\*args*, *complement=False*, *\*\*kwargs*)

Bases: *Primitive*

**calculate**(*coords3d*, *indices=None*, *gradient=False*)

**jacobian**(*coords3d*, *indices=None*)

**class** pysisyphus.intcoords.**LinearDisplacement**(*\*args*, *complement=False*, *\*\*kwargs*)

Bases: *Primitive*

**calculate**(*coords3d*, *indices=None*, *gradient=False*)

**jacobian**(*coords3d*, *indices=None*)

**class** pysisyphus.intcoords.**MWCartesianCoords**(*\*args*, *\*\*kwargs*)

Bases: *CartesianCoords*

**class** pysisyphus.intcoords.**OutOfPlane**(*indices*, *periodic=False*, *calc\_kwargs=None*, *cache=False*)

Bases: *Primitive*

[1] [https://doi.org/10.1002/\(SICI\)1096-987X\(19990730\)20:10<1067::AID-JCC9>3.0.CO;2-V](https://doi.org/10.1002/(SICI)1096-987X(19990730)20:10<1067::AID-JCC9>3.0.CO;2-V)

Lee, 1999

**exception** pysisyphus.intcoords.**PrimitiveNotDefinedException**(*typed\_prim*, *\*args*, *\*\*kwargs*)

Bases: *Exception*

```
class pysisyphus.intcoords.RedundantCoords(atoms, coords3d, masses=None, bond_factor=1.3,  
typed_prims=None, define_prims=None,  
constrain_prims=None, freeze_atoms=None,  
freeze_atoms_exclude=False, define_for=None,  
bonds_only=False, check_bends=True, rebuild=True,  
bend_min_deg=15, dihed_max_deg=175, lb_min_deg=175,  
weighted=False, min_weight=0.3,  
svd_inv_thresh=0.000316, recalc_B=False, tric=False,  
hybrid=False, hbond_angles=False)
```

Bases: object

**property B**

Wilson B-Matrix

**property B\_inv**

Generalized inverse of the Wilson B-Matrix.

**property B\_inv\_prim**

Generalized inverse of the primitive Wilson B-Matrix.

**property B\_prim**

Wilson B-Matrix

**property Bt\_inv**

Transposed generalized inverse of the Wilson B-Matrix.

**property Bt\_inv\_prim**

Transposed generalized inverse of the primitive Wilson B-Matrix.

**property C**

Diagonal matrix. Entries for constraints are set to one.

**property P**

Projection matrix onto B. See [1] Eq. (4).

**backtransform\_hessian**(*redund\_hessian, int\_gradient=None*)

Transform Hessian in internal coordinates to Cartesians.

**property bend\_atom\_indices**

**property bend\_indices**

**property bond\_atom\_indices**

**property bond\_indices**

**property bond\_typed\_prims**

**property cartesian\_indices**

**clear()**

**property constrained\_indices**

**property coords**

**property coords3d**

**property** `dihedral_atom_indices`  
**property** `dihedral_indices`  
`eval(coords3d, attr=None)`  
`get_K_matrix(int_gradient=None)`  
`get_index_of_typed_prim(typed_prim)`  
 Index in `self.typed_prims` for the supplied `typed_prim`.  
`get_prim_internals_by_indices(indices)`  
`inv_B(B)`  
`inv_Bt(B)`  
**property** `linear_bend_indices`  
`log(message)`  
`log_int_grad_msg(int_gradient)`  
**property** `outofplane_indices`  
**property** `prim_coords`  
**property** `prim_indices_set`  
**property** `prim_internals`  
**property** `primitives`  
`print_typed_prims()`  
`project_hessian(H, shift=1000)`  
 Expects a hessian in internal coordinates. See Eq. (11) in [1].  
`project_vector(vector)`  
 Project supplied vector onto range of B.  
`return_inds(slice_)`  
**property** `rotation_indices`  
`set_inds_from_typed_prims(typed_prims)`  
`set_primitive_indices(atoms, coords3d)`  
`transform_forces(cart_forces)`  
 Combination of Eq. (9) and (11) in [1].  
`transform_hessian(cart_hessian, int_gradient=None)`  
 Transform Cartesian Hessian to internal coordinates.  
`transform_int_step(int_step, update_constraints=False, pure=False)`  
**property** `translation_indices`  
**property** `typed_prims`

```
class pysisyphus.intcoords.RotationA(indices, *args, ref_coords3d, **kwargs)
    Bases: Rotation
    index = 0

class pysisyphus.intcoords.RotationB(indices, *args, ref_coords3d, **kwargs)
    Bases: Rotation
    index = 1

class pysisyphus.intcoords.RotationC(indices, *args, ref_coords3d, **kwargs)
    Bases: Rotation
    index = 2

class pysisyphus.intcoords.Stretch(indices, periodic=False, calc_kwargs=None, cache=False)
    Bases: Primitive

class pysisyphus.intcoords.TRIC(*args, **kwargs)
    Bases: RedundantCoords

class pysisyphus.intcoords.Torsion(indices, periodic=False, calc_kwargs=None, cache=False)
    Bases: Primitive

class pysisyphus.intcoords.Torsion2(indices, periodic=False, calc_kwargs=None, cache=False)
    Bases: Torsion

class pysisyphus.intcoords.TranslationX(*args, **kwargs)
    Bases: Translation
    cart_axis = 0

class pysisyphus.intcoords.TranslationY(*args, **kwargs)
    Bases: Translation
    cart_axis = 1

class pysisyphus.intcoords.TranslationZ(*args, **kwargs)
    Bases: Translation
    cart_axis = 2
```

#### 16.1.1.8 pysisyphus.interpolate package

##### Submodules

##### pysisyphus.interpolate.IDPP module

```
class pysisyphus.interpolate.IDPP.IDPP(geoms, between, extrapolate=0, extrapolate_before=0,
                                       extrapolate_after=0, extrapolate_damp=1.0, align=False)
    Bases: Interpolator
    interpolate(initial_geom, final_geom, **kwargs)
```

**pysisyphus.interpolate.Interpolator module**

```
class pysisyphus.interpolate.Interpolator.Interpolator(geoms, between, extrapolate=0,
                                                    extrapolate_before=0, extrapolate_after=0,
                                                    extrapolate_damp=1.0, align=False)
```

Bases: object

```
all_geoms_to_trj(trj_fn)
```

```
interpolate(initial_geom, final_geom, interpolate_only=0, extrapolate=False)
```

```
interpolate_all()
```

**pysisyphus.interpolate.LST module**

```
class pysisyphus.interpolate.LST.LST(*args, align=True, gtol=0.0001, silent=False, **kwargs)
```

Bases: *Interpolator*

```
cost_function(wa_c, f, rab, wab)
```

```
interpolate(initial_geom, final_geom, **kwargs)
```

**pysisyphus.interpolate.Redund module**

```
class pysisyphus.interpolate.Redund.Redund(*args, align=True, **kwargs)
```

Bases: *Interpolator*

```
dump_progress(geoms, out_fn='redund_interpol_fail.trj')
```

```
interpolate(initial_geom, final_geom, interpolate_only=0, extrapolate=False, typed_prims=None)
```

```
restart_interpolate(initial_geom, final_geom, geoms, new_geom)
```

```
step_along_tangent(geom, prim_tangent, step_size)
```

**pysisyphus.interpolate.helpers module**

```
pysisyphus.interpolate.helpers.interpolate(initial_geom, final_geom, between, kind='linear',
                                             only_between=False, align=False, interpol_kwargs=None)
```

```
pysisyphus.interpolate.helpers.interpolate_all(geoms, between, kind='linear', align=False,
                                                  **interpol_kwargs)
```

## Module contents

### 16.1.1.9 pysisyphus.io package

#### Submodules

##### pysisyphus.io.cjson module

`pysisyphus.io.cjson.geom_from_cjson(fn, **kwargs)`

`pysisyphus.io.cjson.parse_cjson(fn)`

##### pysisyphus.io.crd module

`pysisyphus.io.crd.a8(str_)`

`pysisyphus.io.crd.atoms_coords_to_crd_str(atoms, coords, resno=1, res='UNL1', segid=None, resid=1, ref_atoms=None, del_atoms=None)`

`pysisyphus.io.crd.f20(float_)`

`pysisyphus.io.crd.geom_from_crd(fn, **kwargs)`

`pysisyphus.io.crd.geom_to_crd_str(geom, **kwargs)`

`pysisyphus.io.crd.i10(int_)`

##### pysisyphus.io.hdf5 module

`pysisyphus.io.hdf5.get_h5_group(fn, group_name, data_model=None, reset=False)`

Return (and create if necessary) group with given name and data model.

`pysisyphus.io.hdf5.init_h5_group(f, group_name, data_model)`

Create group with given name and data model.

`pysisyphus.io.hdf5.resize_h5_group(group, max_cycles)`

Increase size of first dimension of datasets in the given group.

##### pysisyphus.io.hessian module

`pysisyphus.io.hessian.geom_from_hessian(h5_fn, **geom_kwargs)`

`pysisyphus.io.hessian.save_hessian(h5_fn, geom, cart_hessian=None, energy=None, mult=None)`

`pysisyphus.io.hessian.save_third_deriv(h5_fn, geom, third_deriv_result, H_mw)`

### pysisyphus.io.mol2 module

pysisyphus.io.mol2.geom\_from\_mol2(*text*, *\*\*kwargs*)

### pysisyphus.io.molden module

pysisyphus.io.molden.geoms\_from\_molden(*fn*, *\*\*kwargs*)

### pysisyphus.io.pdb module

pysisyphus.io.pdb.atoms\_coords\_to\_pdb\_str(*atoms*, *coords*, *fragments=None*, *resname=""*, *conect=True*)

pysisyphus.io.pdb.geom\_from\_pdb(*fn*, *\*\*kwargs*)

pysisyphus.io.pdb.geom\_to\_pdb\_str(*geom*, *detect\_fragments=False*, *\*\*kwargs*)

pysisyphus.io.pdb.get\_conect\_lines(*atoms*, *coords*)

pysisyphus.io.pdb.get\_parser(*widths*)

pysisyphus.io.pdb.parse\_atom\_name(*name*)

pysisyphus.io.pdb.parse\_pdb(*fn*)

### pysisyphus.io.pubchem module

pysisyphus.io.pubchem.cid\_from\_name(*name*)

pysisyphus.io.pubchem.geom\_from\_pubchem\_name(*name*, *\*\*kwargs*)

pysisyphus.io.pubchem.sdf\_from\_cid(*cid*)

### pysisyphus.io.sdf module

pysisyphus.io.sdf.geom\_from\_sdf(*text*, *\*\*kwargs*)

pysisyphus.io.sdf.parse\_coord\_line(*line*)

pysisyphus.io.sdf.parse\_sdf(*text*)

### pysisyphus.io.xyz module

pysisyphus.io.xyz.geom\_from\_xyz(*fn*, *\*\*kwargs*)

pysisyphus.io.xyz.geoms\_from\_inline\_xyz(*inline\_xyz*, *\*\*kwargs*)

pysisyphus.io.xyz.geoms\_from\_xyz(*fn*, *\*\*kwargs*)

## pysisyphus.io.zmat module

**class** pysisyphus.io.zmat.ZLine(*atom, rind, r, aind, a, dind, d*)

Bases: tuple

**a**

Alias for field number 4

**aind**

Alias for field number 3

**atom**

Alias for field number 0

**d**

Alias for field number 6

**dind**

Alias for field number 5

**r**

Alias for field number 2

**rind**

Alias for field number 1

pysisyphus.io.zmat.geom\_from\_zmat(*zmat, atoms=None, coords3d=None, geom=None, start\_at=None, drop\_dummy=True, \*\*geom\_kwargs*)

Adapted from <https://github.com/robashaw/geomConvert> by Robert Shaw.

pysisyphus.io.zmat.geom\_from\_zmat\_fn(*fn, \*\*geom\_kwargs*)

pysisyphus.io.zmat.geom\_from\_zmat\_str(*text, coord\_type='cart', coord\_kwargs=None*)

pysisyphus.io.zmat.zmat\_from\_fn(*fn*)

pysisyphus.io.zmat.zmat\_from\_str(*text*)

## Module contents

pysisyphus.io.geom\_from\_cjson(*fn, \*\*kwargs*)

pysisyphus.io.geom\_from\_crd(*fn, \*\*kwargs*)

pysisyphus.io.geom\_from\_hessian(*h5\_fn, \*\*geom\_kwargs*)

pysisyphus.io.geom\_from\_mol2(*text, \*\*kwargs*)

pysisyphus.io.geom\_from\_pdb(*fn, \*\*kwargs*)

pysisyphus.io.geom\_from\_pubchem\_name(*name, \*\*kwargs*)

pysisyphus.io.geom\_from\_zmat(*zmat, atoms=None, coords3d=None, geom=None, start\_at=None, drop\_dummy=True, \*\*geom\_kwargs*)

Adapted from <https://github.com/robashaw/geomConvert> by Robert Shaw.

`pysisyphus.io.geoms_from_molden(fn, **kwargs)`

`pysisyphus.io.geoms_from_xyz(fn, **kwargs)`

`pysisyphus.io.save_hessian(h5_fn, geom, cart_hessian=None, energy=None, mult=None)`

`pysisyphus.io.save_third_deriv(h5_fn, geom, third_deriv_result, H_mw)`

### 16.1.1.10 pysisyphus.irc package

#### Submodules

#### pysisyphus.irc.DWI module

**class** `pysisyphus.irc.DWI.DWI(n=4, maxlen=2)`

Bases: `object`

**\_\_init\_\_**(n=4, maxlen=2)

Distance weighted interpolation.

**dump**(fn)

**static from\_h5**(fn)

**interpolate**(at\_coords, gradient=False)

See [1] Eq. (25) - (29)

**update**(coords, energy, gradient, hessian)

`pysisyphus.irc.DWI.taylor(energy, gradient, hessian, step)`

Taylor series expansion of the energy to second order.

`pysisyphus.irc.DWI.taylor_grad(gradient, hessian, step)`

Gradient of a Taylor series expansion of the energy to second order.

#### pysisyphus.irc.DampedVelocityVerlet module

**class** `pysisyphus.irc.DampedVelocityVerlet.DampedVelocityVerlet(geometry, v0=0.04, dt0=0.5, error_tol=0.003, max_cycles=150, **kwargs)`

Bases: `IRC`

**damp\_velocity**(velocity)

**estimate\_error**(new\_mw\_coords)

**mw\_grad\_to\_acc**(mw\_grad)

Takes care of the units for the mass-weighted gradient. Converts units of a mass-weighted gradient [Hartree/(Bohr\*amu)] to units of acceleration [ $\sqrt{\text{amu}} \cdot \text{Bohr}/\text{fs}^2$ ]. The 1e30 comes from converting second<sup>2</sup> to femto second<sup>2</sup>.

**prepare**(direction)

**step**()

**pysisyphus.irc.Euler module**

```
class pysisyphus.irc.Euler.Euler(geometry, step_length=0.01, **kwargs)
```

Bases: *IRC*

```
step()
```

**pysisyphus.irc.EulerPC module**

```
class pysisyphus.irc.EulerPC.EulerPC(*args, hessian_recalc=None, hessian_update='bofill',
                                       hessian_init='calc', max_pred_steps=500, loose_cycles=3,
                                       dump_dwi=False, scipy_method=None, corr_func='mbs',
                                       **kwargs)
```

Bases: *IRC*

```
corrector_step(init_mw_coords, step_length, dwi)
```

```
get_integration_length_func(init_mw_coords)
```

```
prepare(*args, **kwargs)
```

```
scipy_corrector_step(init_mw_coords, step_length, dwi)
```

Solve IRC equation  $dx/ds = -g/|g|$  on DWI PES in mass-weighted coordinates. Integration done until `self.step_length` in unweighted coordinates is achieved.

```
step()
```

**pysisyphus.irc.GonzalezSchlegel module**

```
class pysisyphus.irc.GonzalezSchlegel.GonzalezSchlegel(geometry, max_micro_cycles=20,
                                                         micro_step_thresh=0.001,
                                                         hessian_recalc=None, line_search=False,
                                                         **kwargs)
```

Bases: *IRC*

```
micro_step(counter)
```

Constrained optimization on a hypersphere.

```
perp_component(vec, perp_to)
```

```
postprocess()
```

```
step()
```

## pysisyphus.irc.IMKMod module

```
class pysisyphus.irc.IMKMod.IMKMod(geometry, corr_first=True, corr_first_size=0.5,
                                   corr_first_energy=True, corr_bisec_size=0.0025,
                                   corr_bisec_energy=True, **kwargs)
```

Bases: *IRC*

```
fit_grad_and_energies(energy0, grad0, energy1, direction)
```

```
fit_parabola(x, y)
```

```
step()
```

## pysisyphus.irc.IRC module

```
class pysisyphus.irc.IRC.IRC(geometry, step_length=0.1, max_cycles=125, downhill=False, forward=True,
                              backward=True, root=0, hessian_init=None, displ='energy',
                              displ_energy=0.001, displ_length=0.1, displ_third_h5=None,
                              rms_grad_thresh=0.001, hard_rms_grad_thresh=None, energy_thresh=1e-06,
                              imag_below=0.0, force_inflection=True, check_bonds=True, out_dir='.',
                              prefix='', dump_fn='irc_data.h5', dump_every=5)
```

Bases: *object*

```
__init__(geometry, step_length=0.1, max_cycles=125, downhill=False, forward=True, backward=True,
         root=0, hessian_init=None, displ='energy', displ_energy=0.001, displ_length=0.1,
         displ_third_h5=None, rms_grad_thresh=0.001, hard_rms_grad_thresh=None,
         energy_thresh=1e-06, imag_below=0.0, force_inflection=True, check_bonds=True, out_dir='.',
         prefix='', dump_fn='irc_data.h5', dump_every=5)
```

Base class for IRC calculations.

### Parameters

- **geometry** (*Geometry*) -- Transition state geometry, or initial geometry for downhill run.
- **step\_length** (*float, optional*) -- Step length in unweighted coordinates.
- **max\_cycles** (*int, optional*) -- Positive integer, controlling the maximum number of IRC steps taken in a direction (forward/backward/downhill).
- **downhill** (*bool, default=False*) -- Downhill run from a non-stationary point with non-vanishing gradient. Disables forward and backward runs.
- **forward** (*bool, default=True*) -- Integrate IRC in positive s direction.
- **backward** (*bool, default=True*) -- Integrate IRC in negative s direction.
- **root** (*int, default=0*) -- Use n-th root for initial displacement from TS.
- **hessian\_init** (*str, default=None*) -- Path to Hessian HDF5 file, e.g., from a previous TS calculation.
- **displ** (*str, one of ("energy", "length", "energy\_cubic")*) -- Controls initial displacement from the TS. 'energy' assumes a quadratic model, from which a step length for a given energy lowering (see 'displ\_energy') is determined. 'length' corresponds to a displacement along the transition vector. 'energy\_cubic' considers 3rd derivatives of the energy along the transition vector.
- **displ\_energy** (*float, default=1e-3*) -- Required energy lowering from the TS in au (Hartree). Used with 'displ: energy|energy\_cubic'.

- **displ\_length** (*float*, *default=0.1*) -- Step length along the transition vector. Used only with 'displ: length'.
- **displ\_third\_h5** (*str*, *optional*) -- Path to HDF5 file containing 3rd derivative information. Used with 'displ: energy\_cubic'.
- **rms\_grad\_thresh** (*float*, *default=1e-3*,) -- Convergence is signalled when to root mean square of the unweighted gradient is less than or equal to this value
- **energy\_thresh** (*float*, *default=1e-6*,) -- Signal convergence when the energy difference between two points is equal to or less than 'energy\_thresh'.
- **imag\_below** (*float*, *default=0.0*) -- Require the wavenumber of the imaginary mode to be below the given threshold. If given, it should be a negative number.
- **force\_inflection** (*bool*, *optional*) -- Don't indicate convergence before passing an inflection point.
- **check\_bonds** (*bool*, *optional*, *default=True*) -- Report whether bonds are formed/broken along the IRC, w.r.t the TS.
- **out\_dir** (*str*, *optional*) -- Dump everything into 'out\_dir' directory instead of the CWD.
- **prefix** (*str*, *optional*) -- Short string that is prepended to all files that are created by this class, e.g., trajectories and HDF5 dumps.
- **dump\_fn** (*str*, *optional*) -- Base name for the HDF5 files.
- **dump\_every** (*int*, *optional*) -- Dump to HDF5 every n-th cycle.

#### property coords

**dump\_data**(*dump\_fn=None*, *full=False*)

**dump\_ends**(*path*, *prefix*, *coords=None*, *trj=False*)

#### property energy

**get\_conv\_fact**(*mw\_grad*, *min\_fact=2.0*)

**get\_endpoint\_and\_ts\_geoms**()

**get\_full\_irc\_data**()

**get\_irc\_data**()

**get\_path\_for\_fn**(*fn*)

#### property gradient

##### **initial\_displacement**()

Returns non-mass-weighted steps in +s and -s direction for initial displacement from the TS. Earlier version only returned one step, that was later multiplied by either 1 or -1, depending on the desired IRC direction (forward/backward). The current implementation directly returns two steps for forward and backward direction. Whereas for plus and minus steps for displ 'length' and displ 'energy'

$\text{step\_plus} = -\text{step\_minus}$

is valid, it is not valid for displ 'energy\_cubic' anymore. The latter step is formed as

$x(ds) = ds * v0 + ds**2 * v1$

```

so
    x(ds) != -x(ds)

as
    ds * v0 + ds**2 * v1 != -ds * v0 - ds**2 * v1 .

```

So, all required step are formed directly and later used as appropriate.

See

<https://aip.scitation.org/doi/pdf/10.1063/1.454172>    <https://pubs.acs.org/doi/10.1021/j100338a027>  
<https://aip.scitation.org/doi/pdf/10.1063/1.459634>

```

irc(direction)

log(msg)

property m_sqrt

mass_weigh_hessian(hessian)

property mw_coords

property mw_gradient

postprocess()

prepare(direction)

report_bonds(prefix, bonds)

report_conv_thresholds()

run()

set_data(prefix)

unweight_vec(vec)

valid_displs = ('energy', 'length', 'energy_cubic')

```

### pysisyphus.irc.IRCDummy module

```

class pysisyphus.irc.IRCDummy.IRCDummy(all_coords, atoms, forward=True, backward=True,
                                         downhill=False)

```

```

Bases: object

all_coords: list

atoms: tuple

backward: bool = True

downhill: bool = False

forward: bool = True

```

## pysisyphus.irc.Instanton module

**class** pysisyphus.irc.Instanton.**Instanton**(*images*, *calc\_getter*, *T*)

Bases: object

**property** P

**P\_bh**

the first image is connected to the last image. At k=0 the index k-1 will be -1, which points to the last image.

**Below we pre-calculate some indices (assuming N images).**

unshifted indices ks: k = {0, 1, .. , N-1} shifted indices ksm1: k-1 = {-1, 0, 1, .. , N-2} shifted indices ksp1: k+1 = {1, 2, .. , N-1, 0}

**Type**

The Instanton is periodic

**action()**

Action in au / fs, Hartree per femtosecond.

**action\_gradient()**

kin\_grad corresponds to the gradient of S\_0 in (Eq. 1 in [1], or first term in Eq. 6 in [2].) It boils down to the derivative of a sum of vector norms

$$d \text{sum}_k (\|y_k - y_{(k-1)}\|_2)^2 \text{ --- } d y_k$$

The derivative of a norm of a vector difference is quite simple, but care has to be taken to recognize, that y\_k appears two times in the sum. It appears in the first summand for k and in the second summand for k+1.

$$\text{sum}_k (\|y_k - y_{(k-1)}\|_2)^2$$

1. term 2. term

$$= (\|y_k - y_{(k-1)}\|_2)^2 + (\|y_{(k+1)} - y_k\|_2)^2 + \dots \text{ and so on}$$

The derivative of the first term is

$$2 * (y_k - y_{(k-1)})$$

and the derivative of the second term is

$$-2 * (y_{(k+1)} - y_k)$$

which is equal to

$$2 * (y_k - y_{(k+1)}) .$$

To summarize:

$$\begin{aligned} d \text{sum}_k (\|y_k - y_{(k-1)}\|_2)^2 \text{ --- } d y_k \\ = 2 * (2 * y_k - y_{(k-1)} - y_{(k+1)}) . \end{aligned}$$

**action\_hessian()**

**as\_xyz()**

**property** cart\_coords

**property** cart\_forces

property `cart_hessian`

property `coords`

property `energy`

property `forces`

classmethod `from_instanton(other, **kwargs)`

classmethod `from_ts(ts_geom, P, dr=0.4, delta_T=25, cart_hessian=None, **kwargs)`

`get_additional_print()`

property `gradient`

property `hessian`

`is_analytical_2d()`

property `path_length`

`pysisyphus.irc.Instanton.T_crossover_from_eigval(eigval)`

`pysisyphus.irc.Instanton.T_crossover_from_ts(ts_geom)`

`pysisyphus.irc.Instanton.log_progress(val, key, i)`

### pysisyphus.irc.LQA module

`class pysisyphus.irc.LQA.LQA(geometry, N_euler=5000, **kwargs)`

Bases: `IRC`

`step()`

### pysisyphus.irc.ModeKill module

`class pysisyphus.irc.ModeKill.ModeKill(geometry, kill_inds, nu_thresh=- 5.0, do_hess=True, hessian_update=True, **kwargs)`

Bases: `IRC`

`get_additional_print()`

`postprocess()`

`prepare(*args, **kwargs)`

`step()`

`update_mw_down_step()`

### pysisyphus.irc.ParamPlot module

**class** pysisyphus.irc.ParamPlot.**ParamPlot**(*coords\_list, param1\_inds, param2\_inds*)

Bases: object

**calc\_angle**(*coords, ind1, ind2, ind3*)

**calc\_bond**(*coords, ind1, ind2*)

**get\_param**(*coords, param\_inds*)

**plot**()

**save**(*path, prefix*)

**save\_coords**(*path, prefix*)

**show**()

### pysisyphus.irc.RK4 module

**class** pysisyphus.irc.RK4.**RK4**(*geometry, step\_length=0.1, max\_cycles=125, downhill=False, forward=True, backward=True, root=0, hessian\_init=None, displ='energy', displ\_energy=0.001, displ\_length=0.1, displ\_third\_h5=None, rms\_grad\_thresh=0.001, hard\_rms\_grad\_thresh=None, energy\_thresh=1e-06, imag\_below=0.0, force\_inflection=True, check\_bonds=True, out\_dir='.', prefix='', dump\_fn='irc\_data.h5', dump\_every=5*)

Bases: *IRC*

**get\_k**(*mw\_coords*)

**step**()

### pysisyphus.irc.initial\_displ module

**class** pysisyphus.irc.initial\_displ.**ThirdDerivResult**(*coords\_plus, energy\_plus, H\_plus, coords\_minus, energy\_minus, H\_minus, G\_vec, vec, ds*)

Bases: tuple

**G\_vec**

Alias for field number 6

**H\_minus**

Alias for field number 5

**H\_plus**

Alias for field number 2

**coords\_minus**

Alias for field number 3

**coords\_plus**

Alias for field number 0

**ds**

Alias for field number 8

**energy\_minus**

Alias for field number 4

**energy\_plus**

Alias for field number 1

**vec**

Alias for field number 7

pysisyphus.irc.initial\_displ.**cubic\_displ**(*H, v0, w0, Gv, dE*)

According to Eq. (26) in [2] **v1 does not depend on the sign of v0.**

$$v1 = (F0 - 2v0^T F0 v0 I)^1 \times ([G0v0] - v0^T [G0v0] v0 I) v0$$

The first term is obviously independent of v0's sign. Using v0' = -v0 the second term becomes

$$([G0v0'] - v0'^T [G0v0'] v0' I) v0' (-[G0v0] - v0^T [G0v0'] v0 I) v0' (-[G0v0] + v0^T [G0v0] v0 I) v0' - (-[G0v0] + v0^T [G0v0] v0 I) v0 ([G0v0] - v0^T [G0v0] v0 I) v0$$

Strictly speaking the contraction [G0v0] should depend on the sign of v0. In the current implementation, the direction of v0 is not taken into account, but

$$\text{get\_curv\_vec}(H, Gv, v0, w0) == \text{get\_curv\_vec}(H, -Gv, -v0, w0) .$$

But somehow the Taylor expansion gives bogus results when called with -Gv and -v0...

pysisyphus.irc.initial\_displ.**cubic\_displ\_for\_geom**(*geom, dE=- 0.0005*)

pysisyphus.irc.initial\_displ.**cubic\_displ\_for\_h5**(*h5\_fn='third\_deriv.h5', dE=- 0.0005*)

pysisyphus.irc.initial\_displ.**get\_curv\_vec**(*H, Gv, v0, w0*)

pysisyphus.irc.initial\_displ.**taylor\_closure**(*H, Gv, v0, v1, w0*)

Taylor expansion of energy to 3rd order.

$$dx(ds) = ds*v0 + ds**2*v1 / 2 \quad dE(ds) = dx^T H dx / 2 + dx^T [Gv] dx / 6$$

H = Hessian Gv = 3rd derivative of energy along v0 v0 = Transition vector v1 = Curvature vector w0 = Eigenvalue belonging to v0

pysisyphus.irc.initial\_displ.**third\_deriv\_fd**(*geom, vec, ds=0.001*)

Third derivative of the energy in direction 'vec'.

**Module contents**

**class** pysisyphus.irc.**DampedVelocityVerlet**(*geometry, v0=0.04, dt0=0.5, error\_tol=0.003, max\_cycles=150, \*\*kwargs*)

Bases: *IRC*

**damp\_velocity**(*velocity*)

**estimate\_error**(*new\_mw\_coords*)

**mw\_grad\_to\_acc**(*mw\_grad*)

Takes care of the units for the mass-weighted gradient. Converts units of a mass-weighted gradient [Hartree/(Bohr\*amu)] to units of acceleration [sqrt(amu)\*Bohr/fs<sup>2</sup>]. The 1e30 comes from converting second<sup>2</sup> to femto second<sup>2</sup>.

**prepare**(*direction*)

**step**()

**class** pysisyphus.irc.**Euler**(*geometry, step\_length=0.01, \*\*kwargs*)

Bases: [IRC](#)

**step**()

**class** pysisyphus.irc.**EulerPC**(\**args, hessian\_recalc=None, hessian\_update='bofill', hessian\_init='calc', max\_pred\_steps=500, loose\_cycles=3, dump\_dwi=False, scipy\_method=None, corr\_func='mbs', \*\*kwargs*)

Bases: [IRC](#)

**corrector\_step**(*init\_mw\_coords, step\_length, dwi*)

**get\_integration\_length\_func**(*init\_mw\_coords*)

**prepare**(\**args, \*\*kwargs*)

**scipy\_corrector\_step**(*init\_mw\_coords, step\_length, dwi*)

Solve IRC equation  $dx/ds = -g/|g|$  on DWI PES in mass-weighted coordinates. Integration done until `self.step_length` in unweighted coordinates is achieved.

**step**()

**class** pysisyphus.irc.**GonzalezSchlegel**(*geometry, max\_micro\_cycles=20, micro\_step\_thresh=0.001, hessian\_recalc=None, line\_search=False, \*\*kwargs*)

Bases: [IRC](#)

**micro\_step**(*counter*)

Constrained optimization on a hypersphere.

**perp\_component**(*vec, perp\_to*)

**postprocess**()

**step**()

**class** pysisyphus.irc.**IMKMod**(*geometry, corr\_first=True, corr\_first\_size=0.5, corr\_first\_energy=True, corr\_bisec\_size=0.0025, corr\_bisec\_energy=True, \*\*kwargs*)

Bases: [IRC](#)

**fit\_grad\_and\_energies**(*energy0, grad0, energy1, direction*)

**fit\_parabola**(*x, y*)

**step**()

**class** pysisyphus.irc.**LQA**(*geometry, N\_euler=5000, \*\*kwargs*)

Bases: [IRC](#)

**step**()

**class** pysisyphus.irc.**ModeKill**(*geometry, kill\_inds, nu\_thresh=- 5.0, do\_hess=True, hessian\_update=True, \*\*kwargs*)

Bases: [IRC](#)

**get\_additional\_print**()

`postprocess()`

`prepare(*args, **kwargs)`

`step()`

`update_mw_down_step()`

```
class pysisyphus.irc.RK4(geometry, step_length=0.1, max_cycles=125, downhill=False, forward=True,
                        backward=True, root=0, hessian_init=None, displ='energy', displ_energy=0.001,
                        displ_length=0.1, displ_third_h5=None, rms_grad_thresh=0.001,
                        hard_rms_grad_thresh=None, energy_thresh=1e-06, imag_below=0.0,
                        force_inflection=True, check_bonds=True, out_dir='.', prefix='',
                        dump_fn='irc_data.h5', dump_every=5)
```

Bases: *IRC*

`get_k(mw_coords)`

`step()`

### 16.1.1.11 pysisyphus.line\_searches package

#### Submodules

#### pysisyphus.line\_searches.Backtracking module

```
class pysisyphus.line_searches.Backtracking.Backtracking(*args, rho_lo=0.05, rho_hi=0.9,
                                                         use_grad=False, **kwargs)
```

Bases: *LineSearch*

```
__init__(*args, rho_lo=0.05, rho_hi=0.9, use_grad=False, **kwargs)
```

Backtracking line search enforcing Armijo conditions.

Uses only energy evaluations.

See [1], Chapter 3, Line Search methods, Section 3.1 p. 31 and Section 3.5 p. 56.

```
alpha_new_from_phi(cycle, phi0, dphi0, alpha, alpha_prev)
```

```
alpha_new_from_phi_dphi(cycle, phi0, dphi0, alpha)
```

```
run_line_search()
```

#### pysisyphus.line\_searches.HagerZhang module

```
class pysisyphus.line_searches.HagerZhang.HagerZhang(*args, alpha_prev=None, f_prev=None,
                                                       dphi0_prev=None, quad_step=False,
                                                       eps=1e-06, theta=0.5, gamma=0.5, rho=5,
                                                       psi_0=0.01, psi_1=0.1, psi_2=2.0,
                                                       psi_low=0.1, psi_hi=10, Delta=0.7,
                                                       omega=0.001, max_bisects=10, **kwargs)
```

Bases: *LineSearch*

**bisect**(*a*, *b*)

Bisect interval [*a*, *b*].

**bracket**(*c*)

Generate initial interval [*a*, *b*] that satisfies the opposite slope condition ( $\text{dphi}(a) < 0$ ,  $\text{dphi}(b) > 0$ ).

**double\_secant**(*a*, *b*)

Take secant<sup>2</sup> step.

**initial**()

Get an initial guess for alpha.

**interval\_update**(*a*, *b*, *c*)

Narrows down the bracketing interval.

**norm\_inf**(*arr*)

Returns infinity norm of given array.

**prepare\_line\_search**()

**run\_line\_search**()

**secant**(*a*, *b*)

Take secant step.

**take\_quad\_step**(*alpha*, *g0*\_)

Try to get alpha for minimum step from quadratic interpolation.

### pysisyphus.line\_searches.LineSearch module

```
class pysisyphus.line_searches.LineSearch.LineSearch(p, cond='armijo', x0=None, geometry=None,
f=None, df=None, alpha_init=None, f0=None,
g0=None, c1=0.1, c2=0.9, max_cycles=10,
alpha_min=1e-06)
```

Bases: object

**check\_alpha**(*alpha*)

**curvature\_condition**(*alpha*)

**get\_fg**(*what*, *alpha*)

Lookup raw function/gradient values for a given alpha.

**get\_phi\_dphi**(*what*, *alpha*, *check*=True)

Wrapper that handles function/gradient evaluations.

**got\_alpha\_phi\_dphi**(*alpha*)

**log**(*message*)

**prepare\_line\_search**()

**run**()

**abstract run\_line\_search**()

**strong\_curvature\_condition**(*alpha*)

**strong\_wolfe\_condition**(*alpha*)

Strong wolfe condition.

**sufficiently\_decreased**(*alpha*)

Sufficient decrease/Armijo condition.

**wolfe\_condition**(*alpha*)

Normal, not strong, Wolfe condition.

**exception** pysisyphus.line\_searches.LineSearch.LineSearchConverged(*alpha*)

Bases: Exception

**exception** pysisyphus.line\_searches.LineSearch.LineSearchNotConverged

Bases: Exception

**class** pysisyphus.line\_searches.LineSearch.LineSearchResult(*converged, alpha, f\_new, g\_new, f\_evals, df\_evals, dphi0*)

Bases: tuple

**alpha**

Alias for field number 1

**converged**

Alias for field number 0

**df\_evals**

Alias for field number 5

**dphi0**

Alias for field number 6

**f\_evals**

Alias for field number 4

**f\_new**

Alias for field number 2

**g\_new**

Alias for field number 3

### pysisyphus.line\_searches.StrongWolfe module

**class** pysisyphus.line\_searches.StrongWolfe.StrongWolfe(\**args, alpha\_max=10.0, fac=2, \*\*kwargs*)

Bases: [LineSearch](#)

**\_\_init\_\_**(\**args, alpha\_max=10.0, fac=2, \*\*kwargs*)

Wolfe line search.

Uses only energy & gradient evaluations.

See [1], Chapter 3, Line Search methods, Section 3.5 p. 60.

**run\_line\_search**()

**zoom**(*alpha\_lo, alpha\_hi, phi\_lo, phi\_alpha\_=None, alpha\_0\_=None, max\_cycles=10*)

## pysisyphus.line\_searches.interpol module

pysisyphus.line\_searches.interpol.**interpol\_alpha\_cubic**(*f\_0, df\_0, f\_alpha\_0, f\_alpha\_1, alpha\_0, alpha\_1*)

pysisyphus.line\_searches.interpol.**interpol\_alpha\_quad**(*f\_0, df\_0, f\_alpha\_0, alpha\_0*)

## Module contents

**class** pysisyphus.line\_searches.**Backtracking**(\*args, rho\_lo=0.05, rho\_hi=0.9, use\_grad=False, \*\*kwargs)

Bases: *LineSearch*

**\_\_init\_\_**(\*args, rho\_lo=0.05, rho\_hi=0.9, use\_grad=False, \*\*kwargs)

Backtracking line search enforcing Armijo conditions.

Uses only energy evaluations.

See [1], Chapter 3, Line Search methods, Section 3.1 p. 31 and Section 3.5 p. 56.

**alpha\_new\_from\_phi**(*cycle, phi0, dphi0, alpha, alpha\_prev*)

**alpha\_new\_from\_phi\_dphi**(*cycle, phi0, dphi0, alpha*)

**run\_line\_search**()

**class** pysisyphus.line\_searches.**HagerZhang**(\*args, alpha\_prev=None, f\_prev=None, dphi0\_prev=None, quad\_step=False, eps=1e-06, theta=0.5, gamma=0.5, rho=5, psi\_0=0.01, psi\_1=0.1, psi\_2=2.0, psi\_low=0.1, psi\_hi=10, Delta=0.7, omega=0.001, max\_bisects=10, \*\*kwargs)

Bases: *LineSearch*

**bisect**(*a, b*)

Bisect interval [a, b].

**bracket**(*c*)

Generate initial interval [a, b] that satisfies the opposite slope condition ( $dphi(a) < 0$ ,  $dphi(b) > 0$ ).

**double\_secant**(*a, b*)

Take secant<sup>2</sup> step.

**initial**()

Get an initial guess for alpha.

**interval\_update**(*a, b, c*)

Narrows down the bracketing interval.

**norm\_inf**(*arr*)

Returns infinity norm of given array.

**prepare\_line\_search**()

**run\_line\_search**()

**secant**(*a, b*)

Take secant step.

**take\_quad\_step**(*alpha*, *g0\_*)

Try to get alpha for minimum step from quadratic interpolation.

**class** pysisyphus.line\_searches.**StrongWolfe**(\*args, alpha\_max=10.0, fac=2, \*\*kwargs)

Bases: *LineSearch*

**\_\_init\_\_**(\*args, alpha\_max=10.0, fac=2, \*\*kwargs)

Wolfe line search.

Uses only energy & gradient evaluations.

See [1], Chapter 3, Line Search methods, Section 3.5 p. 60.

**run\_line\_search**()

**zoom**(alpha\_lo, alpha\_hi, phi\_lo, phi\_alpha\_=None, alpha\_0\_=None, max\_cycles=10)

### 16.1.1.12 pysisyphus.modelfollow package

#### Submodules

#### pysisyphus.modelfollow.NormalMode module

**class** pysisyphus.modelfollow.NormalMode.**NormalMode**(*l*, *masses*)

Bases: object

See <http://gaussian.com/vib/>

**\_\_init\_\_**(*l*, *masses*)

NormalMode class.

Cartesian displacements are normalized to 1.

#### Parameters

- **l** (*np.array*) -- Cartesian, non-mass-weighted displacements.
- **masses** (*np.array*) -- Atomic masses.

**property** **l\_mw**

**mw\_norm\_for\_norm**(*norm=0.01*)

**property** **red\_mass**

#### pysisyphus.modelfollow.davidson module

**class** pysisyphus.modelfollow.davidson.**DavidsonResult**(*cur\_cycle*, *converged*, *final\_modes*, *qs*, *nus*, *mode\_inds*, *res\_rms*)

Bases: tuple

**converged**

Alias for field number 1

**cur\_cycle**

Alias for field number 0

**final\_modes**

Alias for field number 2

**mode\_inds**

Alias for field number 5

**nus**

Alias for field number 4

**qs**

Alias for field number 3

**res\_rms**

Alias for field number 6

```
pysisyphus.modefollow.davidson.block_davidson(cart_coords, masses, forces_getter, guess_modes,  
lowest=None, trial_step_size=0.01,  
hessian_precon=None, max_cycles=25,  
res_rms_thresh=0.0001, start_precon=5,  
remove_trans_rot=True, print_level=1)
```

```
pysisyphus.modefollow.davidson.forces_fin_diff(forces_getter, coords, b, step_size)
```

```
pysisyphus.modefollow.davidson.geom_davidson(geom, *args, **kwargs)
```

### pysisyphus.modefollow.lanczos module

```
pysisyphus.modefollow.lanczos.geom_lanczos(geom, *args, **kwargs)
```

Wraps Lanczos algorithm for use with Geometry objects.

```
pysisyphus.modefollow.lanczos.lanczos(coords, grad_getter, dx=0.005, dl=0.01, guess=None,  
max_cycles=25, reortho=True, logger=None)
```

Lanczos method to determine smallest eigenvalue & -vector.

See [1] for description of algorithm.

### Module contents

```
class pysisyphus.modefollow.NormalMode(l, masses)
```

Bases: object

See <http://gaussian.com/vib/>

```
__init__(l, masses)
```

NormalMode class.

Cartesian displacements are normalized to 1.

**Parameters**

- **l** (*np.array*) -- Cartesian, non-mass-weighted displacements.
- **masses** (*np.array*) -- Atomic masses.

```
property l_mw
```

**mw\_norm\_for\_norm**(*norm=0.01*)

**property red\_mass**

`pysisyphus.modelfollow.geom_davidson`(*geom, \*args, \*\*kwargs*)

`pysisyphus.modelfollow.geom_lanczos`(*geom, \*args, \*\*kwargs*)

Wraps Lanczos algorithm for use with Geometry objects.

### 16.1.1.13 pysisyphus.optimizers package

#### Submodules

#### pysisyphus.optimizers.BFGS module

**class** `pysisyphus.optimizers.BFGS.BFGS`(*geometry, \*args, update='bfgs', \*\*kwargs*)

Bases: *Optimizer*

**bfgs\_update**(*s, y*)

**damped\_bfgs\_update**(*s, y, mu\_1=0.2*)

Damped BFGS update of inverse Hessian.

Potentially updates *s*. See Section 3.2 of [2], Eq. (30) - (33). There is a typo ;) It should be

$$H_{\{k+1\}} = V_k H_k V_k^T + \dots$$

instead of

$$H_{\{k+1\}} = V_k^T H_k V_k + \dots$$

**double\_damped\_bfgs\_update**(*s, y, mu\_1=0.2, mu\_2=0.2*)

Double damped BFGS update of inverse Hessian.

See [3]. Potentially updates *s* and *y*.

**property eye**

**optimize**()

**prepare\_opt**()

#### pysisyphus.optimizers.BacktrackingOptimizer module

**class** `pysisyphus.optimizers.BacktrackingOptimizer.BacktrackingOptimizer`(*geometry, alpha, bt\_force=5, dont\_skip\_after=2, bt\_max\_scale=4, bt\_disable=False, \*\*kwargs*)

Bases: *Optimizer*

**backtrack**(*cur\_forces, prev\_forces, reset\_hessian=None*)

Accelerated backtracking line search.

**reset**()

### pysisyphus.optimizers.ConjugateGradient module

```
class pysisyphus.optimizers.ConjugateGradient.ConjugateGradient(geometry, alpha=0.1,  
                                                                formula='FR', dont_skip=True,  
                                                                **kwargs)
```

Bases: *BacktrackingOptimizer*

**get\_beta**(cur\_forces, prev\_forces)

**optimize**()

**prepare\_opt**()

**reset**()

### pysisyphus.optimizers.CubicNewton module

```
class pysisyphus.optimizers.CubicNewton.CubicNewton(geometry, **kwargs)
```

Bases: *HessianOptimizer*

**optimize**()

**postprocess\_opt**()

### pysisyphus.optimizers.FIRE module

```
class pysisyphus.optimizers.FIRE.FIRE(geometry, dt=0.1, dt_max=1, N_acc=2, f_inc=1.1, f_acc=0.99,  
                                         f_dec=0.5, n_reset=0, a_start=0.1, **kwargs)
```

Bases: *Optimizer*

**optimize**()

**reset**()

### pysisyphus.optimizers.HessianOptimizer module

```
class pysisyphus.optimizers.HessianOptimizer.HessianOptimizer(geometry, trust_radius=0.5,  
                                                                trust_update=True, trust_min=0.1,  
                                                                trust_max=1,  
                                                                max_energy_incr=None,  
                                                                hessian_update='bfgs',  
                                                                hessian_init='fischer',  
                                                                hessian_recalc=None,  
                                                                hessian_recalc_adapt=None,  
                                                                hessian_xtb=False,  
                                                                hessian_recalc_reset=False,  
                                                                small_eigval_thresh=1e-08,  
                                                                line_search=False, alpha0=1.0,  
                                                                max_micro_cycles=25,  
                                                                rfo_overlaps=False, **kwargs)
```

Bases: *Optimizer*

```
__init__(geometry, trust_radius=0.5, trust_update=True, trust_min=0.1, trust_max=1,
max_energy_incr=None, hessian_update='bfgs', hessian_init='fischer', hessian_recalc=None,
hessian_recalc_adapt=None, hessian_xtb=False, hessian_recalc_reset=False,
small_eigval_thresh=1e-08, line_search=False, alpha0=1.0, max_micro_cycles=25,
rfo_overlaps=False, **kwargs)
```

Baseclass for optimizers utilizing Hessian information.

### Parameters

- **geometry** (*Geometry*) -- Geometry to be optimized.
- **trust\_radius** (float, default: 0.5) -- Initial trust radius in whatever unit the optimization is carried out.
- **trust\_update** (bool, default: True) -- Whether to update the trust radius throughout the optimization.
- **trust\_min** (float, default: 0.1) -- Minimum trust radius.
- **trust\_max** (float, default: 1) -- Maximum trust radius.
- **max\_energy\_incr** (Optional[float], default: None) -- Maximum allowed energy increased after a faulty step. Optimization is aborted when the threshold is exceeded.
- **hessian\_update** (Literal['none', None, False, 'bfgs', 'damped\_bfgs', 'flowchart', 'bofill', 'ts\_bfgs', 'ts\_bfgs\_org', 'ts\_bfgs\_rev'], default: 'bfgs') -- Type of Hessian update. Defaults to BFGS for minimizations and Bofill for saddle point searches.
- **hessian\_init** (Literal['calc', 'unit', 'fischer', 'lindh', 'simple', 'swart', 'xtb', 'xtb1', 'xtbff'], default: 'fischer') -- Type of initial model Hessian.
- **hessian\_recalc** (Optional[int], default: None) -- Recalculate exact Hessian every n-th cycle instead of updating it.
- **hessian\_recalc\_adapt** (Optional[float], default: None) -- Use a more flexible scheme to determine Hessian recalculation. Undocumented.
- **hessian\_xtb** (bool, default: False) -- Recalculate the Hessian at the GFN2-XTB level of theory.
- **hessian\_recalc\_reset** (bool, default: False) -- Whether to skip Hessian recalculation after reset. Undocumented.
- **small\_eigval\_thresh** (float, default: 1e-08) -- Threshold for small eigenvalues. Eigenvectors belonging to eigenvalues below this threshold are discarded.
- **line\_search** (bool, default: False) -- Whether to carry out a line search. Not implemented by a subclassing optimizers.
- **alpha0** (float, default: 1.0) -- Initial alpha for restricted-step (RS) procedure.
- **max\_micro\_cycles** (int, default: 25) -- Maximum number of RS iterations.
- **rfo\_overlaps** (bool, default: False) -- Enable mode-following in RS procedure.
- **\*\*kwargs** -- Keyword arguments passed to the Optimizer baseclass.

```
filter_small_eigvals(eigvals, eigvecs, mask=False)
```

```
get_alpha_step(cur_alpha, rfo_eigval, step_norm, eigvals, gradient)
```

```
get_augmented_hessian(eigvals, gradient, alpha=1.0)
```

```

static get_newton_step(eigvals, eigvecs, gradient)
get_newton_step_on_trust(eigvals, eigvecs, gradient, transform=True)
    Step on trust-radius.
    See Nocedal 4.3 Iterative solutions of the subproblem
get_rs_step(eigvals, eigvecs, gradient, name='RS')
static get_shifted_step_trans(eigvals, gradient_trans, shift)
get_step_func(eigvals, gradient, grad_rms_thresh=0.01)
housekeeping()
    Calculate gradient and energy. Update trust radius and hessian if needed. Return energy, gradient and
    hessian for the current cycle.
log_negative_eigenvalues(eigvals, pre_str="")
prepare_opt(hessian_init=None)
property prev_eigvec_max
property prev_eigvec_min
static quadratic_model(gradient, hessian, step)
reset()
rfo_dict = {'max': (-1, 'max'), 'min': (0, 'min')}
static rfo_model(gradient, hessian, step)
save_hessian()
set_new_trust_radius(coeff, last_step_norm)
solve_rfo(rfo_mat, kind='min', prev_eigvec=None)
update_hessian()
update_trust_radius()

```

`pysisyphus.optimizers.HessianOptimizer.dummy_hessian_update(H, dx, dg)`

### pysisyphus.optimizers.LBFGS module

```

class pysisyphus.optimizers.LBFGS.LBFGS(geometry, keep_last=7, beta=1, max_step=0.2,
    double_damp=True, gamma_mult=False, line_search=False,
    mu_reg=None, max_mu_reg_adaptions=10, control_step=True,
    **kwargs)

```

Bases: *Optimizer*

```
__init__(geometry, keep_last=7, beta=1, max_step=0.2, double_damp=True, gamma_mult=False,
         line_search=False, mu_reg=None, max_mu_reg_adaptions=10, control_step=True, **kwargs)
```

Limited-memory BFGS optimizer.

See [1] Nocedal, Wright - Numerical Optimization, 2006 for a general discussion of LBFGS. See `pysisyphus.optimizers.hessian_updates` for the references related to double damping and `pysisyphus.optimizers.closures` for references related to regularized LBFGS.

#### Parameters

- **geometry** (*Geometry*) -- Geometry to be optimized.
- **keep\_last** (int, default: 7) -- History size. Keep last 'keep\_last' steps and gradient differences.
- **beta** (float, default: 1) -- Force constant in  $-(H + I)^{-1}g$ .
- **max\_step** (float, default: 0.2) -- Upper limit for the absolute component of the step vector in whatever unit the optimization is carried out.
- **double\_damp** (bool, default: True) -- Use double damping procedure to modify steps and gradient differences  $y$  to ensure  $sy > 0$ .
- **gamma\_mult** (bool, default: False) -- Estimate  $\gamma$  from previous cycle. Eq. (7.20) in [1]. See 'beta' argument.
- **line\_search** (bool, default: False) -- Enable implicit line searches.
- **mu\_reg** (Optional[float], default: None) -- Initial guess for regularization constant in regularized LBFGS.
- **max\_mu\_reg\_adaptions** (int, default: 10) -- Maximum number of trial steps in regularized LBFGS.
- **control\_step** (bool, default: True) -- Whether to scale down the proposed step its biggest absolute component is equal to or below 'max\_step'
- **\*\*kwargs** -- Keyword arguments passed to the Optimizer baseclass.

```
get_lbfgs_step(forces)
```

```
optimize()
```

```
postprocess_opt()
```

```
reset()
```

### pysisyphus.optimizers.LayerOpt module

```
class pysisyphus.optimizers.LayerOpt.LayerOpt(geometry, layers=None, **kwargs)
```

Bases: *Optimizer*

```
property layer_num: int
```

Return type  
int

```
optimize()
```

Return type  
None

**postprocess\_opt()**

**Return type**

None

**class** pysisyphus.optimizers.LayerOpt.**Layers**(*geometry, opt\_thresh, layers=None*)

Bases: object

**classmethod** **from\_oniom\_calculator**(*geometry, oniom\_calc=None, layers=None, \*\*kwargs*)

pysisyphus.optimizers.LayerOpt.**get\_geom\_kwargs**(*layer\_ind, layer\_mask*)

pysisyphus.optimizers.LayerOpt.**get\_opt\_kwargs**(*opt\_key, layer\_ind, thresh*)

### pysisyphus.optimizers.MicroOptimizer module

**class** pysisyphus.optimizers.MicroOptimizer.**MicroOptimizer**(*geom, step='lbfgs', line\_search=True, max\_cycles=10000000, max\_step=0.2, keep\_last=10, rms\_force=None, double\_damp=True, dump=False, \*\*kwargs*)

Bases: object

**cg\_step**(*forces*)

**lbfgs\_step**(*forces*)

**log**(*msg*)

**optimize**()

**run**()

**sd\_step**(*forces*)

**take\_step**(*energy, forces, return\_step=False*)

### pysisyphus.optimizers.NCOptimizer module

**class** pysisyphus.optimizers.NCOptimizer.**NCOptimizer**(*geometry, \*args, freeze\_modes=None, \*\*kwargs*)

Bases: *HessianOptimizer*

**optimize**()

### pysisyphus.optimizers.Optimizer module

```
class pysisyphus.optimizers.Optimizer.Optimizer(geometry, thresh='gau_loose', max_step=0.04,
max_cycles=100, min_step_norm=1e-08,
assert_min_step=True, rms_force=None,
rms_force_only=False, max_force_only=False,
converge_to_geom_rms_thresh=0.05, align=False,
dump=False, dump_restart=False, prefix="",
reparam_thresh=0.001, reparam_check_rms=True,
reparam_when='after', overachieve_factor=0.0,
restart_info=None, check_coord_diffs=True,
coord_diff_thresh=0.01, fragments=None,
monitor_frag_dists=0, out_dir='.',
h5_fn='optimization.h5', h5_group_name='opt')
```

Bases: object

```
__init__(geometry, thresh='gau_loose', max_step=0.04, max_cycles=100, min_step_norm=1e-08,
assert_min_step=True, rms_force=None, rms_force_only=False, max_force_only=False,
converge_to_geom_rms_thresh=0.05, align=False, dump=False, dump_restart=False, prefix="",
reparam_thresh=0.001, reparam_check_rms=True, reparam_when='after',
overachieve_factor=0.0, restart_info=None, check_coord_diffs=True, coord_diff_thresh=0.01,
fragments=None, monitor_frag_dists=0, out_dir='.', h5_fn='optimization.h5',
h5_group_name='opt')
```

Optimizer baseclass. Meant to be subclassed.

#### Parameters

- **geometry** (*Geometry*) -- Geometry to be optimized.
- **thresh** (Literal['gau\_loose', 'gau', 'gau\_tight', 'gau\_vtight', 'baker', 'never'], default: 'gau\_loose') -- Convergence threshold.
- **max\_step** (float, default: 0.04) -- Maximum absolute component of the allowed step vector. Utilized in optimizers that don't support a trust region or line search.
- **max\_cycles** (int, default: 100) -- Maximum number of allowed optimization cycles.
- **min\_step\_norm** (float, default: 1e-08) -- Minimum norm of an allowed step. If the step norm drops below this value a ZeroStepLength-exception is raised. The unit depends on the coordinate system of the supplied geometry.
- **assert\_min\_step** (bool, default: True) -- Flag that controls whether the norm of the proposed step is check for being too small.
- **rms\_force** (Optional[float], default: None) -- Root-mean-square of the force from which user-defined thresholds are derived. When 'rms\_force' is given 'thresh' is ignored.
- **rms\_force\_only** (bool, default: False) -- When set, convergence is signalled only based on rms(forces).
- **max\_force\_only** (bool, default: False) -- When set, convergence is signalled only based on max(|forces|).
- **converge\_to\_geom\_rms\_thresh** (float, default: 0.05) -- Threshold for the RMSD with another geometry. When the RMSD drops below this threshold convergence is signalled. Only used with Growing Newton trajectories.
- **align** (bool, default: False) -- Flag that controls whether the geometry is aligned in every step onto the coordinates of the previous step. Must not be used with internal coordinates.
- **dump** (bool, default: False) -- Flag to control dumping/writing of optimization progress to the filesystem

- **dump\_restart** (bool, default: False) -- Flag to control whether restart information is dumped to the filesystem.
- **prefix** (str, default: ' ') -- Short string that is prepended to several files created by the optimizer. Allows distinguishing several optimizations carried out in the same directory.
- **reparam\_thresh** (float, default: 0.001) -- Controls the minimal allowed similarity between coordinates after two successive reparametrizations. Convergence is signalled if the coordinates did not change significantly.
- **reparam\_check\_rms** (bool, default: True) -- Whether to check for (too) similar coordinates after reparametrization.
- **reparam\_when** (Optional[Literal['before', 'after']], default: 'after') -- Reparametrize before or after calculating the step. Can also be turned off by setting it to None.
- **overachieve\_factor** (float, default: 0.0) -- Signal convergence when max(forces) and rms(forces) fall below the chosen threshold, divided by this factor. Convergence of max(step) and rms(step) is ignored.
- **restart\_info** -- Restart information. Undocumented.
- **check\_coord\_diffs** (bool, default: True) -- Whether coordinates of chain-of-sates images are checked for being too similar.
- **coord\_diff\_thresh** (float, default: 0.01) -- Unitless threshold for similiary checking of COS image coordinates. The first image is assigned 0, the last image is assigned to 1.
- **fragments** (Optional[Tuple], default: None) -- Tuple of lists containing atom indices, defining two fragments.
- **monitor\_frag\_dists** (int, default: 0) -- Monitor fragment distances for N cycles. The optimization is terminated when the interfragment distances falls below the initial value after N cycles.
- **out\_dir** (str, default: '. ') -- String poiting to a directory where optimization progress is dumped.
- **h5\_fn** (str, default: 'optimization.h5') -- Basename of the HDF5 file used for dumping.
- **h5\_group\_name** (str, default: 'opt') -- Groupname used for dumping of this optimization.

**check\_convergence**(*step=None, multiple=1.0, overachieve\_factor=None, energy\_thresh=1e-06*)

Check if the current convergence of the optimization is equal to or below the required thresholds, or a multiple thereof. The latter may be used in initiating the climbing image.

**dump\_restart\_info**()

**final\_summary**()

**get\_path\_for\_fn**(*fn, with\_prefix=True*)

**get\_restart\_info**()

**log**(*message, level=50*)

**make\_conv\_dict**(*key, rms\_force=None*)

**abstract optimize**()

```

postprocess_opt()
prepare_opt()
print_opt_progress()
report_conv_thresholds()
run()
scale_by_max_step(steps)
set_restart_info(restart_info)
write_cycle_to_file()
write_image_trjs()
write_results()
write_to_out_dir(out_fn, content, mode='w')

```

```
pysisyphus.optimizers.Optimizer.get_data_model(geometry, is_cos, max_cycles)
```

### pysisyphus.optimizers.PreconLBFGS module

```

class pysisyphus.optimizers.PreconLBFGS.PreconLBFGS(geometry, alpha_init=1.0, history=7,
                                                    precon=True, precon_update=1,
                                                    precon_getter_update=None, precon_kind='full',
                                                    max_step_element=None, line_search='armijo',
                                                    c_stab=None, **kwargs)

```

Bases: [Optimizer](#)

```

__init__(geometry, alpha_init=1.0, history=7, precon=True, precon_update=1,
         precon_getter_update=None, precon_kind='full', max_step_element=None, line_search='armijo',
         c_stab=None, **kwargs)

```

Preconditioned limited-memory BFGS optimizer.

See `pysisyphus.optimizers.precon` for related references.

#### Parameters

- **geometry** (*Geometry*) -- Geometry to be optimized.
- **alpha\_init** (float, default: 1.0) -- Initial scaling factor for the first trial step in the explicit line search.
- **history** (int, default: 7) -- History size. Keep last 'history' steps and gradient differences.
- **precon** (bool, default: True) -- Whether to use preconditioning or not.
- **precon\_update** (int, default: 1) -- Recalculate preconditioner P in every n-th cycle with the same topology.
- **precon\_getter\_update** (Optional[int], default: None) -- Recalculate topology for preconditioner P in every n-th cycle. It is usually sufficient to only determine the topology once at the beginning.
- **precon\_kind** (Literal['full', 'full\_fast', 'bonds', 'bonds\_bends'], default: 'full') -- What types of primitive internal coordinates to consider in the preconditioner.

- **max\_step\_element** (Optional[float], default: None) -- Maximum component of the absolute step vector when no line search is carried out.
- **line\_search** (Literal['armijo', 'armijo\_fg', 'strong\_wolfe', 'hz', None, False], default: 'armijo') -- Whether to use explicit line searches and if so, which kind of line search.
- **c\_stab** (Optional[float], default: None) -- Regularization constant  $c$  in  $(H + cI)^1$  in atomic units.
- **\*\*kwargs** -- Keyword arguments passed to the Optimizer baseclass.

```

get_precon_getter()

optimize()

prepare_opt()

scale_max_element(step, max_step_element)

```

### pysisyphus.optimizers.PreconSteepestDescent module

```

class pysisyphus.optimizers.PreconSteepestDescent.PreconSteepestDescent(geometry,
                                                                    alpha_init=0.5,
                                                                    **kwargs)

```

Bases: *PreconLBFGS*

### pysisyphus.optimizers.QuickMin module

```

class pysisyphus.optimizers.QuickMin.QuickMin(geometry, dt=0.35, **kwargs)

```

Bases: *Optimizer*

```

optimize()

prepare_opt()

reset()

```

### pysisyphus.optimizers.RFOptimizer module

```

class pysisyphus.optimizers.RFOptimizer.RFOptimizer(geometry, line_search=True, gdiis=False,
                                                       gdiis=True, gdiis_thresh=0.0025,
                                                       gdiis_thresh=0.01, gdiis_test_direction=True,
                                                       max_micro_cycles=0, adapt_step_func=False,
                                                       **kwargs)

```

Bases: *HessianOptimizer*

```

__init__(geometry, line_search=True, gdiis=False, gdiis=True, gdiis_thresh=0.0025, gdiis_thresh=0.01,
         gdiis_test_direction=True, max_micro_cycles=0, adapt_step_func=False, **kwargs)

```

Rational function Optimizer.

#### Parameters

- **geometry** (*Geometry*) -- Geometry to be optimized.
- **line\_search** (bool, default: True) -- Whether to carry out implicit line searches.

- **gediis** (bool, default: False) -- Whether to enable GEDIIS.
- **gdiis** (bool, default: True) -- Whether to enable GDIIS.
- **gdiis\_thresh** (float, default: 0.0025) -- Threshold for rms(forces) to enable GDIIS.
- **gediis\_thresh** (float, default: 0.01) -- Threshold for rms(step) to enable GEDIIS.
- **gdiis\_test\_direction** (bool, default: True) -- Whether to the overlap of the RFO step and the GDIIS step.
- **max\_micro\_cycles** (int, default: 0) -- Number of restricted-step microcycles. Disabled by default.
- **adapt\_step\_func** (bool, default: False) -- Whether to switch between shifted Newton and RFO-steps.
- **\*\*kwargs** -- Keyword arguments passed to the Optimizer/HessianOptimizer baseclass.

**optimize()**

**postprocess\_opt()**

### pysisyphus.optimizers.RSA module

```
class pysisyphus.optimizers.RSA.RSA(geometry, trust_radius=0.5, trust_update=True, trust_min=0.1,
    trust_max=1, max_energy_incr=None, hessian_update='bfgs',
    hessian_init='fischer', hessian_recalc=None,
    hessian_recalc_adapt=None, hessian_xtb=False,
    hessian_recalc_reset=False, small_eigval_thresh=1e-08,
    line_search=False, alpha0=1.0, max_micro_cycles=25,
    rfo_overlaps=False, **kwargs)
```

Bases: *HessianOptimizer*

The Importance of Step Control in Optimization Methods, del Campo, 2009.

**optimize()**

### pysisyphus.optimizers.StabilizedQNMethod module

```
class pysisyphus.optimizers.StabilizedQNMethod.StabilizedQNMethod(geometry, alpha=0.5,
    alpha_max=1,
    alpha_stretch=0.5,
    alpha_stretch_max=1,
    eps=0.0001, hist_max=10,
    E_thresh=1e-06, bio=True,
    trust_radius=0.1,
    linesearch=True, **kwargs)
```

Bases: *Optimizer*

**adjust\_alpha**(gradient, precon\_gradient)

**adjust\_alpha\_stretch**()

**bio\_mode**(gradient)

```
property n_hist
optimize()
precondition_gradient(gradient, steps, grad_diffs, eps)
prepare_opt()
```

### pysisyphus.optimizers.SteepestDescent module

```
class pysisyphus.optimizers.SteepestDescent.SteepestDescent(geometry, alpha=0.1, **kwargs)
    Bases: BacktrackingOptimizer
    optimize()
    prepare_opt()
```

### pysisyphus.optimizers.StringOptimizer module

```
class pysisyphus.optimizers.StringOptimizer.StringOptimizer(geometry, max_step=0.1,
    stop_in_when_full=-1, keep_last=10,
    lbfgs_when_full=True,
    gamma_mult=False,
    double_damp=True,
    scale_step='global', **kwargs)
    Bases: Optimizer
    check_convergence(*args, **kwargs)
        Check if the current convergence of the optimization is equal to or below the required thresholds, or a
        multiple thereof. The latter may be used in initiating the climbing image.
    optimize()
    prepare_opt()
    reset()
    restrict_step_components(steps)
```

### pysisyphus.optimizers.closures module

```
pysisyphus.optimizers.closures.bfgs_multiply(s_list, y_list, vector, beta=1, P=None, logger=None,
    gamma_mult=True, mu_reg=None, inds=None,
    cur_size=None)
```

Matrix-vector product H·v.

Multiplies given vector with inverse Hessian, obtained from repeated BFGS updates calculated from steps in 's\_list' and gradient differences in 'y\_list'.

Based on algorithm 7.4 Nocedal, Num. Opt., p. 178.

```
pysisyphus.optimizers.closures.get_update_mu_reg(mu_min=0.001, gamma_1=0.1, gamma_2=5.0,
    eta_1=0.01, eta_2=0.9, logger=None)
```

See 5.1 in [1]

`pysisyphus.optimizers.closures.lbfgs_closure(force_getter, M=10, beta=1, restrict_step=None)`

`pysisyphus.optimizers.closures.modified_broyden_closure(force_getter, M=5, beta=1, restrict_step=None)`

<https://doi.org/10.1006/jcph.1996.0059> F corresponds to the residual gradient, so we after calling `force_getter` we multiply the force by -1 to get the gradient.

`pysisyphus.optimizers.closures.small_lbfgs_closure(history=5, gamma_mult=True)`

Compact LBFGS closure.

The returned function takes two arguments: `forces` and `prev_step`. `forces` are the forces at the current iterate and `prev_step` is the previous step that lead us to the current iterate. In this way step restriction/line search can be done outside of the `lbfgs` function.

### pysisyphus.optimizers.cls\_map module

`pysisyphus.optimizers.cls_map.get_opt_cls(opt_key)`

`pysisyphus.optimizers.cls_map.key_is_tsopt(opt_key)`

### pysisyphus.optimizers.exceptions module

**exception** `pysisyphus.optimizers.exceptions.OptimizationError`

Bases: `Exception`

**exception** `pysisyphus.optimizers.exceptions.ZeroStepLength`

Bases: `Exception`

### pysisyphus.optimizers.gdiis module

**class** `pysisyphus.optimizers.gdiis.DIISResult(coeffs, coords, forces, energy, N, type)`

Bases: `tuple`

**N**

Alias for field number 4

**coeffs**

Alias for field number 0

**coords**

Alias for field number 1

**energy**

Alias for field number 3

**forces**

Alias for field number 2

**type**

Alias for field number 5

`pysisyphus.optimizers.gdiis.diis_result(coeffs, coords, forces, energy=None, prefix='')`

```
pysisyphus.optimizers.gdiis.from_coeffs(vec, coeffs)
pysisyphus.optimizers.gdiis.gdiis(err_vecs, coords, forces, ref_step, max_vecs=5, test_direction=True)
pysisyphus.optimizers.gdiis.gediis(coords, energies, forces, hessian=None, max_vecs=3)
pysisyphus.optimizers.gdiis.log(msg)
pysisyphus.optimizers.gdiis.valid_diis_direction(diis_step, ref_step, use)
```

### pysisyphus.optimizers.guess\_hessians module

```
pysisyphus.optimizers.guess_hessians.fischer_guess(geom)
pysisyphus.optimizers.guess_hessians.get_guess_hessian(geometry, hessian_init, int_gradient=None,
cart_gradient=None, h5_fn=None)
```

Obtain/calculate (model) Hessian.

For `hessian_init="calc"` the Hessian will be in the `coord_type` of the geometry, otherwise a Hessian in primitive internals will be returned.

```
pysisyphus.optimizers.guess_hessians.get_lindh_alpha(atom1, atom2)
pysisyphus.optimizers.guess_hessians.improved_guess(geom, bond_func, bend_func, dihedral_func)
pysisyphus.optimizers.guess_hessians.lindh_guess(geom)
```

Slightly modified Lindh model hessian as described in [1].

Instead of using the tabulated `r_ref,ij` values from [1] we will use the 'true' covalent radii as pyberny. The tabulated `r_ref,ij` value for two carbons (2nd period) is 2.87 Bohr. Carbons covalent radius is ~ 1.44 Bohr, so  $2 * 1.44 \text{ Bohr} = 2.88 \text{ Bohr}$  which fits nicely with the tabulate value. If values for elements > 3rd are requested the alpha values for the 3rd period will be (re)used.

```
pysisyphus.optimizers.guess_hessians.lindh_style_guess(geom, ks, rhos)
```

Approximate force constants according to Lindh.[1]

Bonds:  $k_{ij} = k_r * \rho_{ij}$  Bends:  $k_{ijk} = k_b * \rho_{ij} * \rho_{jk}$  Dihedrals:  $k_{ijkl} = k_d * \rho_{ij} * \rho_{jk} * \rho_{kl}$

```
pysisyphus.optimizers.guess_hessians.simple_guess(geom)
```

Default force constants.

```
pysisyphus.optimizers.guess_hessians.swart_guess(geom)
```

```
pysisyphus.optimizers.guess_hessians.ts_hessian(hessian, coord_inds, damp=0.25)
```

According to [3]

```
pysisyphus.optimizers.guess_hessians.xtb_hessian(geom, gfn=None)
```

**pysisyphus.optimizers.hessian\_updates module**

`pysisyphus.optimizers.hessian_updates.bfgs_update(H, dx, dg)`

`pysisyphus.optimizers.hessian_updates.bofill_update(H, dx, dg)`

`pysisyphus.optimizers.hessian_updates.damped_bfgs_update(H, dx, dg)`

See [5]

`pysisyphus.optimizers.hessian_updates.double_damp(s, y, H=None, s_list=None, y_list=None, mu_1=0.2, mu_2=0.2, logger=None)`

Double damped step 's' and gradient differences 'y'.

H is the inverse Hessian! See [6]. Potentially updates s and y. y is only updated if mu\_2 is not None.

**Parameters**

- **s** (*np.array, shape (N, ), floats*) -- Coordinate differences/step.
- **y** (*np.array, shape (N, ), floats*) -- Gradient differences
- **H** (*np.array, shape (N, N), floats, optional*) -- Inverse Hessian.
- **s\_list** (*list of nd.array, shape (K, N), optional*) -- List of K previous steps. If no H is supplied and prev\_ys is given the matrix-vector product Hy will be calculated through the two-loop LBFGS-recursion.
- **y\_list** (*list of nd.array, shape (K, N), optional*) -- List of K previous gradient differences. See s\_list.
- **mu\_1** (*float, optional*) -- Parameter for 's' damping.
- **mu\_2** (*float, optional*) -- Parameter for 'y' damping.
- **logger** (*logging.Logger, optional*) -- Logger to be used.

**Returns**

- **s** (*np.array, shape (N, ), floats*) -- Damped coordinate differences/step.
- **y** (*np.array, shape (N, ), floats*) -- Damped gradient differences

`pysisyphus.optimizers.hessian_updates.flowchart_update(H, dx, dg)`

`pysisyphus.optimizers.hessian_updates.mod_flowchart_update(H, dx, dg)`

`pysisyphus.optimizers.hessian_updates.psb_update(z, dx)`

`pysisyphus.optimizers.hessian_updates.sr1_update(z, dx)`

`pysisyphus.optimizers.hessian_updates.ts_bfgs_update(H, dx, dg)`

As described in [7]

`pysisyphus.optimizers.hessian_updates.ts_bfgs_update_org(H, dx, dg)`

Do not use! Implemented as described in the 1998 bofill paper [8].

This does not seem to work too well.

`pysisyphus.optimizers.hessian_updates.ts_bfgs_update_revised(H, dx, dg)`

TS-BFGS update as described in [9].

Better than the original formula of Bofill, worse than the implementation in [7]. a is calculated as described in the footnote 1 on page 38. Eq. (8) looks suspicious as it contains the inverse of a vector?! As also outlined in the paper abs(a) is used (**|a|** in the paper).

## pysisyphus.optimizers.poly\_fit module

**class** pysisyphus.optimizers.poly\_fit.**FitResult**(*x, y, polys*)

Bases: tuple

**polys**

Alias for field number 2

**x**

Alias for field number 0

**y**

Alias for field number 1

pysisyphus.optimizers.poly\_fit.**cubic\_fit**(*e0, e1, g0, g1*)

pysisyphus.optimizers.poly\_fit.**gen\_solutions**()

Given two energies (*e0, e1*) and corresponding gradients (*g0, g1*) we can (try to) fit a quartic polynomial

$$f(x) = a_0 + a_1*x + a_2*x**2 + a_3*x**3 + a_4*x**4$$

s.t. the constraint  $f'(x) \geq 0$ , with the equality being fulfilled at only one point. There are five unknowns ( $a_0 - a_4$ ) to be determined. Four equations can be derived from  $f(x)$  and its first derivative

$$f'(x) = a_1 + 2*a_2*x + 3*a_3*x**2 + 4*a_4*x**3 .$$

With (*e0, g0*) being given at  $x=0$  and (*e1, g1*) being given at  $x=1$  we can setup the following equations:

$$f(0) = a_0 \quad (1) \quad f'(0) = a_1 \quad (2)$$

**using  $e_0$  and  $g_0$  at  $x=0$ , and**

$$f(1) = a_0 + a_1 + a_2 + a_3 + a_4 \quad (3) \quad f'(1) = a_1 + 2*a_2 + 3*a_3 + 4*a_4 . \quad (4)$$

The missing last equation can be derived from the constraint. The second derivative of  $f(x)$  is

$$f''(x) = 2*a_2 + 6*a_3*x + 12*a_4*x**2$$

and shall be positive except at one point where it is allowed to be 0, that its two roots ( $f''(x) = 0$ ) must be degenerate. This is fulfilled when the discriminant  $D$  of the quadratic polynomial  $a*x**2 + b*x + c$  is zero.

$$D = b**2 - 4*a*c = 0$$

**With**

$$a = 12*a_4 \quad b = 6*a_3 \quad c = 2*a_2$$

**we get**

$$0 = (6*a_3)**2 - 4*12*a_4*2*a_2 \quad 0 = 36*a_3**2 - 96*a_4*a_2 \quad 0 = 3*a_3**2 - 8*a_4*a_2 \quad (5) \quad \text{or} \quad a_4 = 3/8 * a_3**2 / a_2$$

Using (1) - (5) we can solve the set of equations for  $a_0 - a_4$ .

pysisyphus.optimizers.poly\_fit.**get\_maximum**(*poly*)

pysisyphus.optimizers.poly\_fit.**get\_minimum**(*poly*)

pysisyphus.optimizers.poly\_fit.**poly\_line\_search**(*cur\_energy, prev\_energy, cur\_grad, prev\_grad, prev\_step, cubic\_max\_x=2.0, quartic\_max\_x=4.0, logger=None*)

Generate directional gradients by projecting them on the previous step.

`pysisyphus.optimizers.poly_fit.quartic_fit(e0, e1, g0, g1, maximize=False)`

See `gen_solutions()` for derivation.

`pysisyphus.optimizers.poly_fit.quintic_fit(e0, e1, g0, g1, H0, H1)`

### **pysisyphus.optimizers.precon module**

`pysisyphus.optimizers.precon.get_lindh_k(atoms, coords3d, bonds=None, angles=None, torsions=None)`

`pysisyphus.optimizers.precon.get_lindh_precon(atoms, coords, bonds=None, bends=None, dihedrals=None, c_stab=0.0103, logger=None)`

`c_stab = 0.00103 hartree/bohr2` corresponds to `0.1 eV/Å2` as given in the paper.

`pysisyphus.optimizers.precon.precon_getter(geom, c_stab=0.0103, kind='full', logger=None)`

### **pysisyphus.optimizers.restrict\_step module**

`pysisyphus.optimizers.restrict_step.get_scale_max(max_element)`

`pysisyphus.optimizers.restrict_step.restrict_step(steps, max_step)`

`pysisyphus.optimizers.restrict_step.scale_by_max_step(steps, max_step)`

### **Module contents**

**class** `pysisyphus.optimizers.CubicNewton(geometry, **kwargs)`

Bases: `HessianOptimizer`

**optimize()**

**postprocess\_opt()**

**class** `pysisyphus.optimizers.MicroOptimizer(geom, step='lbfgs', line_search=True, max_cycles=100000000, max_step=0.2, keep_last=10, rms_force=None, double_damp=True, dump=False, **kwargs)`

Bases: `object`

**cg\_step(*forces*)**

**lbfgs\_step(*forces*)**

**log(*msg*)**

**optimize()**

**run()**

**sd\_step(*forces*)**

**take\_step(*energy, forces, return\_step=False*)**

#### 16.1.1.14 pysisyphus.plotters package

##### Submodules

##### pysisyphus.plotters.AnimPlot module

```
class pysisyphus.plotters.AnimPlot.AnimPlot(calculator, optimizer, xlim=None, ylim=None,
                                             levels=None, num=100, figsize=(8, 8), interval=250,
                                             energy_profile=True, colorbar=True, save=None,
                                             title=True, tight_layout=False)
```

Bases: object

**animate**()

**as\_html5**(out\_fn)

**func**(frame)

**on\_keypress**(event)

Pause on SPACE press.

##### Module contents

#### 16.1.1.15 pysisyphus.stocastic package

##### Submodules

##### pysisyphus.stocastic.FragmentKick module

```
class pysisyphus.stocastic.FragmentKick.FragmentKick(geom, fragments, fix_fragments=None,
                                                       displace_from=None,
                                                       random_displacement=False, **kwargs)
```

Bases: *Kick*

**get\_fragments**(fragments)

**get\_input\_geom**(geom)

**get\_origin**()

**kick\_fragment**(frag\_coords)

**print\_fragments**()

**pysisyphus.stochastic.Kick module**

```
class pysisyphus.stochastic.Kick.Kick(geom, radius=0.5, **kwargs)
```

Bases: *Pipeline*

```
get_input_geom(geom)
```

```
get_kick()
```

```
run_cycle(geom)
```

**pysisyphus.stochastic.Pipeline module**

```
class pysisyphus.stochastic.Pipeline.Pipeline(geom, calc_getter=None, seed=None, max_cycles=5,
                                             cycle_size=15, rmsd_thresh=0.1, energy_thresh=0.001,
                                             energy_range=0.125, compare_num=25, break_after=2,
                                             calc_kwargs=None)
```

Bases: object

```
atoms_are_too_close(geom, factor=0.7)
```

Determine if atoms are too close.

```
geom_is_close_in_energy(geom)
```

```
geom_is_new(geom)
```

Determine if geometry is not already known.

```
geom_is_valid(geom)
```

Filter out geometries that are None, or were the atoms are too close or when they are already known.

```
get_input_geom(geom)
```

```
get_unique_geometries(geoms)
```

```
get_valid_index_set(to_intersect)
```

```
log(message)
```

Write a log message.

Wraps the logger variable.

**Parameters**

**message** (*str*) -- Message to be logged.

```
run()
```

```
run_geom_opt(geom)
```

```
write_geoms_to_trj(geoms, fn, comments=None)
```

## pysisyphus.stochastic.align module

`pysisyphus.stochastic.align.apply_transform`(*coords3d*, *swap*, *reflection*)

Apply axis swaps and reflections to a coordinate array.

`pysisyphus.stochastic.align.match_geom_atoms`(*ref\_geom*, *geom\_to\_match*, *hydrogen=True*)

Apply the hungarian method to *geom\_to\_match*.

Uses the scipy implementation of the Hungarian method/ Kuhn-Munkres algorithm in `scipy.optimize.linear_sum_assignment`.

### See

[1] 10.1021/ci400534h [2] 10.1021/acs.jcim.6b00516

`pysisyphus.stochastic.align.matched_rmsd`(*geom1*, *geom2*, *thresh=0.05*)

RMSD for optimally aligned and matched geometries.

### Returns

- **matched\_rmsd** (*float*) -- RMSD of optimally aligned and matched geometries.
- **matched\_geoms** (*tuple*(*Geometry*, *Geometry*)) -- Tuple of the optimally aligned and matched geometries.

## Module contents

`class pysisyphus.stochastic.FragmentKick`(*geom*, *fragments*, *fix\_fragments=None*, *displace\_from=None*, *random\_displacement=False*, *\*\*kwargs*)

Bases: *Kick*

`get_fragments`(*fragments*)

`get_input_geom`(*geom*)

`get_origin`()

`kick_fragment`(*frag\_coords*)

`print_fragments`()

`class pysisyphus.stochastic.Kick`(*geom*, *radius=0.5*, *\*\*kwargs*)

Bases: *Pipeline*

`get_input_geom`(*geom*)

`get_kick`()

`run_cycle`(*geom*)

### 16.1.1.16 pysisyphus.tests package

#### Submodules

pysisyphus.tests.test\_calculators module

#### Module contents

### 16.1.1.17 pysisyphus.tsoptimizers package

#### Submodules

pysisyphus.tsoptimizers.RSIRFOptimizer module

```
class pysisyphus.tsoptimizers.RSIRFOptimizer.RSIRFOptimizer(geometry, root=0, hessian_ref=None,
    prim_coord=None, rx_coords=None,
    rx_mode=None, hessian_init='calc',
    hessian_update='bofill',
    hessian_recalc_reset=True,
    max_micro_cycles=50,
    trust_radius=0.3, trust_max=0.5,
    augment_bonds=False,
    min_line_search=False,
    max_line_search=False,
    assert_neg_eigval=False, **kwargs)
```

Bases: *TSHessianOptimizer*

**optimize()**

pysisyphus.tsoptimizers.RSPRFOptimizer module

```
class pysisyphus.tsoptimizers.RSPRFOptimizer.RSPRFOptimizer(geometry, root=0, hessian_ref=None,
    prim_coord=None, rx_coords=None,
    rx_mode=None, hessian_init='calc',
    hessian_update='bofill',
    hessian_recalc_reset=True,
    max_micro_cycles=50,
    trust_radius=0.3, trust_max=0.5,
    augment_bonds=False,
    min_line_search=False,
    max_line_search=False,
    assert_neg_eigval=False, **kwargs)
```

Bases: *TSHessianOptimizer*

**optimize()**

### pysisyphus.tsoptimizers.TRIM module

```
class pysisyphus.tsoptimizers.TRIM.TRIM(geometry, root=0, hessian_ref=None, prim_coord=None, rx_coords=None, rx_mode=None, hessian_init='calc', hessian_update='bofill', hessian_recalc_reset=True, max_micro_cycles=50, trust_radius=0.3, trust_max=0.5, augment_bonds=False, min_line_search=False, max_line_search=False, assert_neg_eigval=False, **kwargs)
```

Bases: *TSHessianOptimizer*

**optimize()**

### pysisyphus.tsoptimizers.TSHessianOptimizer module

```
class pysisyphus.tsoptimizers.TSHessianOptimizer.TSHessianOptimizer(geometry, root=0, hessian_ref=None, prim_coord=None, rx_coords=None, rx_mode=None, hessian_init='calc', hessian_update='bofill', hessian_recalc_reset=True, max_micro_cycles=50, trust_radius=0.3, trust_max=0.5, augment_bonds=False, min_line_search=False, max_line_search=False, assert_neg_eigval=False, **kwargs)
```

Bases: *HessianOptimizer*

Optimizer to find first-order saddle points.

```
__init__(geometry, root=0, hessian_ref=None, prim_coord=None, rx_coords=None, rx_mode=None, hessian_init='calc', hessian_update='bofill', hessian_recalc_reset=True, max_micro_cycles=50, trust_radius=0.3, trust_max=0.5, augment_bonds=False, min_line_search=False, max_line_search=False, assert_neg_eigval=False, **kwargs)
```

Baseclass for transition state optimizers utilizing Hessian information.

Several arguments expect a typed primitive or an iterable of typed primitives. A typed primitive is specified as (PrimType, int, int, ...), e.g., for a bond between atoms 0 and 1: (BOND, 0, 1) or for a bend between the atom triple 0, 1, 2 as (BEND, 0, 1, 2).

#### Parameters

- **geometry** (*Geometry*) -- Geometry to be optimized.
- **root** (int, default: 0) -- Index of imaginary mode to maximize along.
- **hessian\_ref** (Optional[str], default: None) -- Filename pointing to a pysisyphus HDF5 Hessian.
- **prim\_coord** (*typed\_prim*) -- Select initial root/imaginary mode by overlap with this internal coordinate.

- **rx\_coords** (*iterable of (typed\_prim)*) -- Construct imaginary mode comprising the given typed prims by modifying a model Hessian.
- **rx\_mode** (*iterable of (typed\_prim, phase\_factor)*) -- Select initial root by overlap with a mode constructed from the given typed primitives.
- **hessian\_init** (`Literal['calc', 'unit', 'fischer', 'lindh', 'simple', 'swart', 'xtb', 'xtb1', 'xtbff']`, default: `'calc'`) -- Type of initial model Hessian.
- **hessian\_update** (`Literal['none', None, False, 'bfgs', 'damped_bfgs', 'flowchart', 'bofill', 'ts_bfgs', 'ts_bfgs_org', 'ts_bfgs_rev']`, default: `'bofill'`) -- Type of Hessian update. Defaults to BFGS for minimizations and Bofill for saddle point searches.
- **hessian\_recalc\_reset** (`bool`, default: `True`) -- Whether to skip Hessian recalculation after reset. Undocumented.
- **max\_micro\_cycles** (`int`, default: `50`) -- Maximum number of RS iterations.
- **trust\_radius** (`float`, default: `0.3`) -- Initial trust radius in whatever unit the optimization is carried out.
- **trust\_max** (`float`, default: `0.5`) -- Maximum trust radius.
- **augment\_bonds** (`bool`, default: `False`) -- Try to derive additional stretching coordinates from the imaginary mode.
- **min\_line\_search** (`bool`, default: `False`) -- Carry out line search along the imaginary mode.
- **max\_line\_search** (`bool`, default: `False`) -- Carry out line search in the subspace that is minimized.
- **assert\_neg\_eigval** (`bool`, default: `False`) -- Check for the existences for at least one significant negative eigenvalue. If enabled and no negative eigenvalue is present the optimization will be aborted.
- **\*\*kwargs** -- Keyword arguments passed to the HessianOptimizer/Optimizer baseclass.

```
static do_line_search(e0, e1, g0, g1, prev_step, maximize, logger=None)
```

```
prepare_opt(*args, **kwargs)
```

```
step_and_grad_from_line_search(energy, gradient_trans, eigvecs, min_indices)
```

```
update_ts_mode(eigvals, eigvecs)
```

```
valid_updates = ('bofill', 'ts_bfgs', 'ts_bfgs_org', 'ts_bfgs_rev')
```

## Module contents

```
class pysisyphus.tsoptimizers.RSIRFOptimizer(geometry, root=0, hessian_ref=None, prim_coord=None,
rx_coords=None, rx_mode=None, hessian_init='calc',
hessian_update='bofill', hessian_recalc_reset=True,
max_micro_cycles=50, trust_radius=0.3, trust_max=0.5,
augment_bonds=False, min_line_search=False,
max_line_search=False, assert_neg_eigval=False,
**kwargs)
```

Bases: *TSHessianOptimizer*

```
optimize()
```

```
class pysisyphus.tsoptimizers.RSPRFOptimizer(geometry, root=0, hessian_ref=None, prim_coord=None,
                                             rx_coords=None, rx_mode=None, hessian_init='calc',
                                             hessian_update='bofill', hessian_recalc_reset=True,
                                             max_micro_cycles=50, trust_radius=0.3, trust_max=0.5,
                                             augment_bonds=False, min_line_search=False,
                                             max_line_search=False, assert_neg_eigval=False,
                                             **kwargs)
```

Bases: *TSHessianOptimizer*

**optimize()**

```
class pysisyphus.tsoptimizers.TRIM(geometry, root=0, hessian_ref=None, prim_coord=None,
                                     rx_coords=None, rx_mode=None, hessian_init='calc',
                                     hessian_update='bofill', hessian_recalc_reset=True,
                                     max_micro_cycles=50, trust_radius=0.3, trust_max=0.5,
                                     augment_bonds=False, min_line_search=False,
                                     max_line_search=False, assert_neg_eigval=False, **kwargs)
```

Bases: *TSHessianOptimizer*

**optimize()**

### 16.1.1.18 pysisyphus.wrapper package

#### Submodules

##### pysisyphus.wrapper.jmol module

`pysisyphus.wrapper.jmol.call_jmol(spt_str, show=False)`

`pysisyphus.wrapper.jmol.render_cdd_cube(cdd_cube, isoval=0.002, orient='')`

`pysisyphus.wrapper.jmol.render_geom_and_charges(geom, point_charges)`

##### pysisyphus.wrapper.mwfn module

`pysisyphus.wrapper.mwfn.call_mwfn(inp_fn, stdin, cwd=None)`

`pysisyphus.wrapper.mwfn.get_mwfn_exc_str(energies, ci_coeffs, dexc_ci_coeffs=None, thresh=0.001)`

`pysisyphus.wrapper.mwfn.log(msg)`

`pysisyphus.wrapper.mwfn.make_cdd(inp_fn, state, log_fn, cwd=None, keep=False, quality=2, prefix='S')`

Create CDD cube in cwd.

#### Parameters

- **inp\_fn** (*str*) -- Filename of a .molden/.fchk file.
- **state** (*int*) -- CDD cubes will be generated up to this state.
- **log\_fn** (*str*) -- Filename of the .log file.
- **cwd** (*str* or *Path*, *optional*) -- If a different cwd should be used.
- **keep** (*bool*) -- Whether to keep electron.cub and hole.cub, default is False.

- **quality** (*int*) -- Quality of the cube. (1=low, 2=medium, 3=high).

`pysisyphus.wrapper.mwfn.wrap_stdin(stdin)`

## pysisyphus.wrapper.packmol module

`pysisyphus.wrapper.packmol.call_packmol(inp)`

`pysisyphus.wrapper.packmol.make_input(output_fn, solvent_fn, solvent_num, sphere_radius, solute_fn=None, solute_num=None, tolerance=2.0)`

## Module contents

### 16.1.2 Submodules

### 16.1.3 pysisyphus.Geometry module

**class** `pysisyphus.Geometry.Geometry`(*atoms, coords, fragments=None, coord\_type='cart', coord\_kwargs=None, isotopes=None, freeze\_atoms=None, comment="", name=""*)

Bases: `object`

**\_\_init\_\_**(*atoms, coords, fragments=None, coord\_type='cart', coord\_kwargs=None, isotopes=None, freeze\_atoms=None, comment="", name=""*)

Object representing atoms in a coordinate system.

The Geometry represents atoms and their positions in coordinate system. By default cartesian coordinates are used, but internal coordinates are also possible.

#### Parameters

- **atoms** (*iterable*) -- Iterable of length N, containing element symbols.
- **coords** (*1d iterable*) -- 1d iterable of length 3N, containing the cartesian coordinates of N atoms.
- **fragments** (*dict, optional*) -- Dict with different keys denoting different fragments. The values contain lists of atom indices.
- **coord\_type** (*{"cart", "redund"}, optional*) -- Type of coordinate system to use. Right now cartesian (cart) and redundand (redund) are supported.
- **coord\_kwargs** (*dict, optional*) -- Dictionary containing additional arguments that get passed to the constructor of the internal coordinate class.
- **isotopes** (*iterable of pairs, optional*) -- Iterable of pairs consisting of 0-based atom index and either an integer or a float. If an integer is given the closest isotope mass will be selected. Given a float, this float will be directly used as mass.
- **freeze\_atoms** (*iterable of integers*) -- Specifies which atoms should remain fixed at their initial positions.
- **comment** (*str, optional*) -- Comment string.
- **name** (*str, optional*) -- Verbose name of the geometry, e.g. methanal or water. Used for printing

**align\_principal\_axes()**

Align the principal axes to the cartesian axes.

<https://math.stackexchange.com/questions/145023>

**approximate\_radius()**

Approximate molecule radius from the biggest atom distance along an axis.

**as\_ase\_atoms()****as\_g98\_list()**

Returns data for fake Gaussian98 standard orientation output.

**Returns**

**g98\_list** -- List with one row per atom. Every row contains [center number, atomic number, atomic type (always 0 for now), X Y Z coordinates in Angstrom.

**Return type**

list

**as\_xyz(comment="", atoms=None, cart\_coords=None)**

Current geometry as a string in XYZ-format.

**Parameters**

- **comment** (*str*, *optional*) -- Will be written in the second line (comment line) of the XYZ-string.
- **cart\_coords** (*np.array*, *1d*, *shape* (3 \* *atoms.size*, )) -- Cartesians for dumping instead of *self.\_coords*.

**Returns**

**xyz\_str** -- Current geometry as string in XYZ-format.

**Return type**

str

**assert\_cart\_coords(coords)****assert\_compatibility(other)**

Assert that two Geometries can be subtracted from each other.

**Parameters**

**other** (*Geometry*) -- Geometry for comparison.

**atom\_indices()**

Dict with atom types as key and corresponding indices as values.

**Returns**

**inds\_dict** -- Unique atom types as keys, corresponding indices as values.

**Return type**

dict

**property atom\_types****atom\_xyz\_iter()****property atomic\_numbers****property bond\_sets**

`calc_double_ao_overlap(geom2)`

`calc_energy_and_forces()`

Force a calculation of the current energy and forces.

**property** `cart_coords`

**property** `cart_forces`

**property** `cart_gradient`

**property** `cart_hessian`

`center()`

**property** `center_of_mass`

Returns the center of mass.

**Returns**

**R** -- Center of mass.

**Return type**

np.array, shape(3, )

`center_of_mass_at(coords3d)`

Returns the center of mass at given coords3d.

**Parameters**

**coords3d** (np.array, shape(*N*, 3)) -- Cartesian coordiantes.

**Returns**

**R** -- Center of mass.

**Return type**

np.array, shape(3, )

**property** `centroid`

Geometric center of the Geometry.

**Returns**

**R** -- Geometric center of the Geometry.

**Return type**

np.array, shape(3, )

`clear()`

Reset the object state.

**property** `comment`

```
coord_types = {'cart': None, 'cartesian': <class
'pysisyphus.intcoords.CartesianCoords.CartesianCoords'>, 'dlc': <class
'pysisyphus.intcoords.DLC.DLC'>, 'hdlc': <class 'pysisyphus.intcoords.DLC.HDLC'>,
'hredund': <class 'pysisyphus.intcoords.RedundantCoords.HybridRedundantCoords'>,
'mwcartesian': <class 'pysisyphus.intcoords.CartesianCoords.MWCartesianCoords'>,
'redund': <class 'pysisyphus.intcoords.RedundantCoords.RedundantCoords'>, 'tric':
<class 'pysisyphus.intcoords.RedundantCoords.TRIC'>}
```

**property coords**

1d vector of atomic coordinates.

**Returns**

**coords** -- 1d array holding the current coordinates.

**Return type**

np.array

**property coords3d**

Coordinates in 3d.

**Returns**

**coords3d** -- Coordinates of the Geometry as 2D array.

**Return type**

np.array

**property coords\_by\_type**

Coordinates in 3d by atom type and their corresponding indices.

**Returns**

- **cbt** (*dict*) -- Dictionary with the unique atom types of the Geometry as keys. It's values are the 3d coordinates of the corresponding atom type.
- **inds** (*dict*) -- Dictionary with the unique atom types of the Geometry as keys. It's values are the original indices of the 3d coordinates in the whole coords3d array.

**copy**(*coord\_type=None, coord\_kwargs=None*)

Returns a new Geometry object with same atoms and coordinates.

**Parameters**

- **coord\_type** (*str*) -- Desired coord\_type, defaults to current coord\_type.
- **coord\_kwargs** (*dict, optional*) -- Any desired coord\_kwargs that will be passed to the RedundantCoords object.

**Returns**

**geom** -- New Geometry object with the same atoms and coordinates.

**Return type**

*Geometry*

**copy\_all**(*coord\_type=None, coord\_kwargs=None*)**property covalent\_radii****del\_atoms**(*inds, \*\*kwargs*)**describe**()**dump\_xyz**(*fn*)**eckart\_projection**(*mw\_hessian, return\_P=False, full=False*)**property energy**

Energy of the current atomic configuration.

**Returns**

**energy** -- Energy of the current atomic configuration.

**Return type**

float

**fd\_coords3d\_gen**(*step\_size=0.001*)

Iterator returning 3d Cartesians for finite-differences.

**property forces**

Energy of the current atomic configuration.

**Returns****force** -- 1d array containing the forces acting on the atoms. Negative of the gradient.**Return type**

np.array

**get\_energy\_and\_cart\_forces\_at**(*cart\_coords*)**get\_energy\_and\_cart\_hessian\_at**(*cart\_coords*)**get\_energy\_and\_forces\_at**(*coords*)

Calculate forces and energies at the given coordinates.

The results are not saved in the Geometry object.

**get\_energy\_at**(*coords*)**get\_energy\_at\_cart\_coords**(*cart\_coords*)**get\_fragments**(*regex*)**get\_imag\_frequencies**(*hessian=None, thresh=1e-06*)**get\_normal\_modes**(*cart\_hessian=None, full=False*)

Normal mode wavenumbers, eigenvalues and Cartesian displacements Hessian.

**get\_restart\_info**()**get\_sphere\_radius**(*offset=4*)**get\_subgeom**(*indices, coord\_type='cart', sort=False*)

Return a Geometry containing a subset of the current Geometry.

**Parameters**

- **indices** (*iterable of ints*) -- Atomic indices that the define the subset of the current Geometry.
- **coord\_type** (*str, ("cart", "redund"), optional*) -- Coordinate system of the new Geometry.

**Returns****sub\_geom** -- Subset of the current Geometry.**Return type***Geometry***get\_subgeom\_without**(*indices, \*\*kwargs*)**get\_temporary\_coords**(*coords*)**get\_thermoanalysis**(*energy=None, cart\_hessian=None, T=298.15, p=101325, point\_group='c1'*)

**get\_trans\_rot\_projector**(*full=False*)

**property gradient**

Negative of the force.

**Returns**

**gradient** -- 1d array containing the negative of the current forces.

**Return type**

np.array

**property hessian**

Matrix of second derivatives of the energy in respect to atomic displacements.

**Returns**

**hessian** -- 2d array containing the second derivatives of the energy with respect to atomic/coordinate displacements depending on the type of coordiante system.

**Return type**

np.array

**property inertia\_tensor**

**property is\_analytical\_2d**

**jmol**(*atoms=None, cart\_coords=None*)

Show geometry in jmol.

**property layers**

**mass\_weigh\_hessian**(*hessian*)

**property masses**

**property masses\_rep**

**property mm\_inv**

Inverted mass matrix.

Returns a diagonal matrix containing the inverted atomic masses.

**property mm\_sqrt\_inv**

Inverted square root of the mass matrix.

**property moving\_atoms**

**moving\_atoms\_jmol**()

**property mw\_coords**

Mass-weighted coordinates.

**Returns**

**mw\_coords** -- 1d array containing the mass-weighted cartesian coordiantes.

**Return type**

np.array

**property mw\_gradient**

Mass-weighted gradient.

**Returns**

**mw\_gradient** -- Returns the mass-weighted gradient.

**Return type**

np.array

**property mw\_hessian**

Mass-weighted hessian.

**Returns****mw\_hessian** -- 2d array containing the mass-weighted hessian  $M^{(-1/2)} H M^{(-1/2)}$ .**Return type**

np.array

**principal\_axes\_are\_aligned()**

Check if the principal axes are aligned with the cartesian axes.

**Returns****aligned** -- Wether the principal axes are aligned or not.**Return type**

bool

**reparametrize()****reset\_coords**(*new\_typed\_prims=None*)**rmsd**(*geom*)**rotate**(*copy=False*)**set\_calculator**(*calculator, clear=True*)

Reset the object and set a calculator.

**set\_coord**(*ind, coord*)

Set a coordinate by index.

**Parameters**

- **ind** (*int*) -- Index in of the coordinate to set in the self.coords array.
- **coord** (*float*) -- Coordinate value.

**set\_coords**(*coords, cartesian=False, update\_constraints=False*)**set\_h5\_hessian**(*fn*)**set\_restart\_info**(*restart\_info*)**set\_results**(*results*)

Save the results from a dictionary.

**Parameters****results** (*dict*) -- The keys in this dict will be set as attributes in the current object, with the corresponding item as value.**standard\_orientation()****property sum\_formula****property total\_mass**

**unweight\_mw\_hessian**(*mw\_hessian*)

Unweight a mass-weighted hessian.

**Parameters**

**mw\_hessian** (*np.array*) -- Mass-weighted hessian to be unweighted.

**Returns**

**hessian** -- 2d array containing the hessian.

**Return type**

*np.array*

**without\_hydrogens**()

**zero\_frozen\_forces**(*cart\_forces*)

`pysisyphus.Geometry.get_trans_rot_projector`(*cart\_coords*, *masses*, *full=False*)

`pysisyphus.Geometry.get_trans_rot_vectors`(*cart\_coords*, *masses*, *rot\_thresh=1e-06*)

Vectors describing translation and rotation.

These vectors are used for the Eckart projection by constructing a projector from them.

See Martin J. Field - A Pratical Introduction to the simulation of Molecular Systems, 2007, Cambridge University Press, Eq. (8.23), (8.24) and (8.26) for the actual projection.

See also <https://chemistry.stackexchange.com/a/74923>.

**Parameters**

- **cart\_coords** (*np.array*, *1d*, *shape* (*3 \* atoms.size*, )) -- Atomic masses in amu.
- **masses** (*iterable*, *1d*, *shape* (*atoms.size*, )) -- Atomic masses in amu.

**Returns**

**ortho\_vecs** -- 2d array containing row vectors describing translations and rotations.

**Return type**

*np.array*(6, 3\**atoms.size*)

`pysisyphus.Geometry.inertia_tensor`(*coords3d*, *masses*)

Inertita tensor.

$x^2 \quad xy \quad xz \quad |$

$(x \ y \ z)^T \cdot (x \ y \ z) = | \quad xy \quad y^2 \quad yz \quad |$

$xz \quad yz \quad z^2 \quad |$

## 16.1.4 pysisyphus.TableFormatter module

`class pysisyphus.TableFormatter.TableFormatter`(*header*, *fmts*, *min\_width=7*, *space=3*)

Bases: object

**property header**

**join\_format**(*fmts*, *lst*)

Format a given iterable *lst* with formats given in the iterable *lst* and return the joined items of the formatted list.

**line**(\*args)

**min\_width\_fmts**(raw\_fmts=None)

pysisyphus.TableFormatter.run()

### 16.1.5 pysisyphus.TablePrinter module

**class** pysisyphus.TablePrinter.**TablePrinter**(header, col\_fmts, width=12, sub\_underline=True, shift\_left=0, fmts\_update=None)

Bases: object

**print**(\*args, \*\*kwargs)

**print\_header**(with\_sep=True)

**print\_row**(args)

**print\_sep**()

### 16.1.6 pysisyphus.color module

pysisyphus.color.**bool\_color**(bool\_, str\_=None)

pysisyphus.color.**green**(str\_)

pysisyphus.color.**red**(str\_)

### 16.1.7 pysisyphus.config module

pysisyphus.config.**detect\_paths**()

pysisyphus.config.**get\_cmd**(section, key='cmd', use\_defaults=True)

pysisyphus.config.**parse\_args**(args)

pysisyphus.config.**run\_detect\_paths**()

### 16.1.8 pysisyphus.constants module

### 16.1.9 pysisyphus.elem\_data module

### 16.1.10 pysisyphus.exceptions module

**exception** pysisyphus.exceptions.**HEIIsFirstOrLastException**

Bases: Exception

### 16.1.11 pysisyphus.filtertrj module

`pysisyphus.filtertrj.get_unique_internals(geom)`

`pysisyphus.filtertrj.parse_args(args)`

`pysisyphus.filtertrj.parse_filter(raw_filter)`

`pysisyphus.filtertrj.run()`

### 16.1.12 pysisyphus.helpers module

`class pysisyphus.helpers.FinalHessianResult(neg_eigvals, eigvals, nus, imag_fns, thermo)`

Bases: tuple

**eigvals**

Alias for field number 1

**imag\_fns**

Alias for field number 3

**neg\_eigvals**

Alias for field number 0

**nus**

Alias for field number 2

**thermo**

Alias for field number 4

`pysisyphus.helpers.align_coords(coords_list)`

`pysisyphus.helpers.align_geoms(geoms)`

`pysisyphus.helpers.check_for_end_sign(check_user=True, cwd='.)`

`pysisyphus.helpers.complete_fragments(atoms, fragments)`

`pysisyphus.helpers.confirm_input(message)`

`pysisyphus.helpers.do_final_hessian(geom, save_hessian=True, write_imag_modes=False, is_ts=False, prefix='', T=298.15, p=101325, ev_thresh=- 1e-06, print_thermo=False, out_dir=None)`

`pysisyphus.helpers.fit_rigid(geometry, vectors=(), vector_lists=(), hessian=None)`

`pysisyphus.helpers.geom_from_xyz_file(xyz_fn, coord_type='cart', **coord_kwargs)`

`pysisyphus.helpers.geom_loader(fn, coord_type='cart', iterable=False, **coord_kwargs)`

After introducing the pubchem functionality I don't like this function anymore :) Too complicated.

`pysisyphus.helpers.geoms_from_trj(trj_fn, first=None, coord_type='cart', **coord_kwargs)`

`pysisyphus.helpers.get_coords_diffs(coords, align=False, normalize=True)`

`pysisyphus.helpers.get_geom_getter(ref_geom, calc_setter)`

`pysisyphus.helpers.get_tangent_trj_str(atoms, coords, tangent, comment=None, points=10, displ=None)`

`pysisyphus.helpers.imag_modes_from_geom(geom, freq_thresh=-10, points=10, displ=None)`

`pysisyphus.helpers.index_array_from_overlaps(overlaps, axis=1)`

It is assumed that the overlaps between two points with indices  $i$  and  $j$  with ( $j > i$ ) are computed and that  $i$  changes along the first axis ( $axis=0$ ) and  $j$  changes along the second axis ( $axis=1$ ).

So the first row of the overlap matrix (`overlaps[0]`) should contain the overlaps between state 0 at index  $i$  and all states at index  $j$ .

`argmax` along axis 1 returns the indices of the most overlapping states at index  $j$  with the states at index  $i$ , given by the item index in the indices array. E.g.:

[0 1 3 2] indicates a root flip in a system with four states when going from index  $i$  to index  $j$ . Root 2 at  $i$  became root 3 at  $j$  and vice versa.

`pysisyphus.helpers.match_geoms(ref_geom, geom_to_match, hydrogen=False)`

See

[1] 10.1021/ci400534h [2] 10.1021/acs.jcim.6b00516

`pysisyphus.helpers.norm_max_rms(arr)`

`pysisyphus.helpers.np_print(func, precision=2, suppress=True, linewidth=120)`

`pysisyphus.helpers.pick_image_inds(cart_coords, images)`

Pick approx. evenly distributed images from given Cartesian coordinates.

`pysisyphus.helpers.print_barrier(ref_energy, comp_energy, ref_str, comp_str)`

`pysisyphus.helpers.procrustes(geometry)`

`pysisyphus.helpers.rms(arr)`

`pysisyphus.helpers.shake_coords(coords, scale=0.1, seed=None)`

`pysisyphus.helpers.simple_peaks(data)`

`pysisyphus.helpers.slugify_worker(dask_worker)`

### 16.1.13 pysisyphus.helpers\_pure module

`class pysisyphus.helpers_pure.OrderedEnum(value)`

Bases: Enum

An enumeration.

`pysisyphus.helpers_pure.approx_float(num, expected, abs_tol=1e-06, rel_tol=1e-12)`

**Return type**

bool

`pysisyphus.helpers_pure.check_mem(mem, pal, avail_frac=0.85, logger=None)`

`pysisyphus.helpers_pure.chunks(l, n)`

Yield successive  $n$ -sized chunks from  $l$ . <https://stackoverflow.com/a/312464>

`pysisyphus.helpers_pure.describe(arr)`

`pysisyphus.helpers_pure.eigval_to_wavenumber(ev)`

`pysisyphus.helpers_pure.expand(to_expand)`

`pysisyphus.helpers_pure.file_or_str(*args, method=False)`

`pysisyphus.helpers_pure.filter_fixture_store(test_name)`

`pysisyphus.helpers_pure.find_closest_sequence(str_, comp_strs)`

**Return type**

Tuple[str, float]

`pysisyphus.helpers_pure.full_expand(to_expand)`

`pysisyphus.helpers_pure.get_clock()`

`pysisyphus.helpers_pure.get_input(data, prompt, lbl_func=None)`

`pysisyphus.helpers_pure.get_molecular_radius(coords3d, min_offset=0.9452)`

`pysisyphus.helpers_pure.get_ratio(str_, comp_str)`

See <https://stackoverflow.com/a/17388505>

**Return type**

str

`pysisyphus.helpers_pure.hash_args(*args, precision=4)`

`pysisyphus.helpers_pure.hash_arr(arr, precision=4)`

`pysisyphus.helpers_pure.highlight_text(text, width=80, level=0)`

`pysisyphus.helpers_pure.increment_fn(org_fn, suffix=None)`

Append, or increase a suffixed counter on a given filename. If no counter is present it will be set to zero. Otherwise it is incremented by one.

```
>>> increment_fn("opt", "rebuilt")
'opt_rebuilt_000'
>>> increment_fn("opt")
'opt_000'
>>> increment_fn("opt_rebuilt_000", "rebuilt")
'opt_rebuilt_001'
```

**Parameters**

- **org\_fn** (str) -- The original, unaltered filename.
- **suffix** (Optional[str], default: None) -- Optional suffix to be append.

**Returns**

Modified filename with optional suffix and incremented counter.

**Return type**

incr\_fn

`pysisyphus.helpers_pure.interpolate_colors(values, c1, c2, num=32)`

Expects two RGB colors c1 and c2.

`pysisyphus.helpers_pure.json_to_results(as_json)`

`pysisyphus.helpers_pure.log(logger, msg, level=10)`

`pysisyphus.helpers_pure.merge_sets(fragments)`

Merge a list of iterables.

`pysisyphus.helpers_pure.recursive_update(d, u)`

Recursive update of *d* with keys/values from *u*.

From <https://stackoverflow.com/questions/3232943>

`pysisyphus.helpers_pure.remove_duplicates(seq)`

`pysisyphus.helpers_pure.report_frozen_atoms(geom)`

`pysisyphus.helpers_pure.report_isotopes(geom, affect_str)`

`pysisyphus.helpers_pure.results_to_json(results)`

`pysisyphus.helpers_pure.sort_by_central(set1, set2)`

Determines a common index in two sets and returns a length 3 tuple with the central index at the middle position and the two terminal indices as first and last indices.

`pysisyphus.helpers_pure.standard_state_corr(T=298.15, p=101325, n=1)`

dG for change of standard state from gas to solution of 1 mol/l

`pysisyphus.helpers_pure.timed(logger)`

`pysisyphus.helpers_pure.touch(fn)`

### 16.1.14 pysisyphus.init\_logging module

`pysisyphus.init_logging.get_fh_logger(name, log_fn)`

Initialize a logger with 'name', level DEBUG and a FileHandler.

`pysisyphus.init_logging.init_logging(log_dir='.', scheduler=None)`

Prepare the logger in *log\_path*. When called with scheduler loggers for every worker are prepared.

`pysisyphus.init_logging.init_logging_base(dask_worker, log_path)`

Prepare individual loggers for one *dask\_worker*.

### 16.1.15 pysisyphus.linalg module

`pysisyphus.linalg.cross3(a, b)`

10x as fast as `np.cross` for two 1d arrays of size 3.

`pysisyphus.linalg.eigvec_grad(w, v, ind, mat_grad)`

Gradient of 'ind'-th eigenvector.

$dv_i / dx_i = (w_i I - mat)^{-1} dmat/dx_i v_i$

`pysisyphus.linalg.finite_difference_hessian`(*coords*, *grad\_func*, *step\_size=0.01*, *acc=2*)

Numerical Hessian from central finite gradient differences.

**See central differences in**

[https://en.wikipedia.org/wiki/Finite\\_difference\\_coefficient](https://en.wikipedia.org/wiki/Finite_difference_coefficient)

for the different accuracies.

**Return type**

`ndarray[Any, dtype[float]]`

`pysisyphus.linalg.get_rot_mat`(*abc=None*)

`pysisyphus.linalg.get_rot_mat_for_coords`(*coords3d\_1*, *coords3d\_2*)

`pysisyphus.linalg.gram_schmidt`(*vecs*, *thresh=1e-08*)

`pysisyphus.linalg.make_unit_vec`(*vec1*, *vec2*)

Return unit vector pointing from *vec2* to *vec1*.

`pysisyphus.linalg.norm3`(*a*)

5x as fast as `np.linalg.norm` for a 1d array of size 3.

`pysisyphus.linalg.orthogonalize_against`(*mat*, *vecs*, *max\_cycles=5*, *thresh=1e-10*)

Orthogonalize rows of 'mat' against rows in 'vecs'

Returns a (modified) copy of *mat*.

`pysisyphus.linalg.perp_comp`(*vec*, *along*)

Return the perpendicular component of *vec* along *along*.

`pysisyphus.linalg.svd_inv`(*array*, *thresh*, *hermitian=False*)

## 16.1.16 pysisyphus.pack module

`pysisyphus.pack.as_pdb`(*fn*)

`pysisyphus.pack.parse_args`(*args*)

`pysisyphus.pack.print_info`(*title*, *geom*)

`pysisyphus.pack.run`()

`pysisyphus.pack.sphere_radius_from_volume`(*volume*)

`pysisyphus.pack.volume_for_density`(*molecule\_num*, *mol\_mass*, *density*)

## 16.1.17 pysisyphus.peakdetect module

`pysisyphus.peakdetect.peakdetect`(*y\_axis*, *x\_axis=None*, *lookahead=200*, *delta=0*)

Converted from/based on a MATLAB script at: <http://billauer.co.il/peakdet.html>

function for detecting local maxima and minima in a signal. Discovers peaks by searching for values which are surrounded by lower or larger values for maxima and minima respectively

keyword arguments: *y\_axis* -- A list containing the signal over which to find peaks

**x\_axis** -- A x-axis whose values correspond to the y\_axis list and is used

in the return to specify the position of the peaks. If omitted an index of the y\_axis is used. (default: None)

**lookahead** -- distance to look ahead from a peak candidate to determine if

it is the actual peak (default: 200) '(samples / period) / f' where '4 >= f >= 1.25' might be a good value

**delta** -- this specifies a minimum difference between a peak and

the following points, before a peak may be considered a peak. Useful to hinder the function from picking up false peaks towards to end of the signal. To work well delta should be set to  $\text{delta} \geq \text{RMSnoise} * 5$ . (default: 0)

When omitted delta function causes a 20% decrease in speed. When used Correctly it can double the speed of the function

**return:** two lists [max\_peaks, min\_peaks] containing the positive and

negative peaks respectively. Each cell of the lists contains a tuple of: (position, peak\_value) to get the average peak value do: `np.mean(max_peaks, 0)[1]` on the results to unpack one of the lists into x, y coordinates do: `x, y = zip(*max_peaks)`

`pysisyphus.peakdetect.peakdetect_fft(y_axis, x_axis, pad_len=20)`

Performs a FFT calculation on the data and zero-pads the results to increase the time domain resolution after performing the inverse fft and send the data to the 'peakdetect' function for peak detection.

Omitting the x\_axis is forbidden as it would make the resulting x\_axis value silly if it was returned as the index 50.234 or similar.

Will find at least 1 less peak than the 'peakdetect\_zero\_crossing' function, but should result in a more precise value of the peak as resolution has been increased. Some peaks are lost in an attempt to minimize spectral leakage by calculating the fft between two zero crossings for n amount of signal periods.

The biggest time eater in this function is the ifft and thereafter it's the 'peakdetect' function which takes only half the time of the ifft. Speed improvements could include to check if  $2^{**n}$  points could be used for fft and ifft or change the 'peakdetect' to the 'peakdetect\_zero\_crossing', which is maybe 10 times faster than 'peakdetct'. The pro of 'peakdetect' is that it results in one less lost peak. It should also be noted that the time used by the ifft function can change greatly depending on the input.

keyword arguments: y\_axis -- A list containing the signal over which to find peaks

**x\_axis** -- A x-axis whose values correspond to the y\_axis list and is used

in the return to specify the position of the peaks.

**pad\_len** -- By how many times the time resolution should be

increased by, e.g. 1 doubles the resolution. The amount is rounded up to the nearest  $2^{**n}$  amount (default: 20)

**return:** two lists [max\_peaks, min\_peaks] containing the positive and

negative peaks respectively. Each cell of the lists contains a tuple of: (position, peak\_value) to get the average peak value do: `np.mean(max_peaks, 0)[1]` on the results to unpack one of the lists into x, y coordinates do: `x, y = zip(*max_peaks)`

`pysisyphus.peakdetect.peakdetect_parabola(y_axis, x_axis, points=31)`

Function for detecting local maxima and minima in a signal. Discovers peaks by fitting the model function:  $y = k(x - \tau)^{**2} + m$  to the peaks. The amount of points used in the fitting is set by the points argument.

Omitting the x\_axis is forbidden as it would make the resulting x\_axis value silly, if it was returned as index 50.234 or similar.

will find the same amount of peaks as the 'peakdetect\_zero\_crossing' function, but might result in a more precise value of the peak.

keyword arguments: y\_axis -- A list containing the signal over which to find peaks

**x\_axis** -- A x-axis whose values correspond to the y\_axis list and is used in the return to specify the position of the peaks.

**points** -- How many points around the peak should be used during curve fitting (default: 31)

**return:** two lists [max\_peaks, min\_peaks] containing the positive and negative peaks respectively. Each cell of the lists contains a tuple of: (position, peak\_value) to get the average peak value do: `np.mean(max_peaks, 0)[1]` on the results to unpack one of the lists into x, y coordinates do: `x, y = zip(*max_peaks)`

`pysisyphus.peakdetect.peakdetect_sine(y_axis, x_axis, points=31, lock_frequency=False)`

Function for detecting local maxima and minima in a signal. Discovers peaks by fitting the model function:  $y = A * \sin(2 * \pi * f * (x - \tau))$  to the peaks. The amount of points used in the fitting is set by the points argument.

Omitting the x\_axis is forbidden as it would make the resulting x\_axis value silly if it was returned as index 50.234 or similar.

will find the same amount of peaks as the 'peakdetect\_zero\_crossing' function, but might result in a more precise value of the peak.

The function might have some problems if the sine wave has a non-negligible total angle i.e. a  $k*x$  component, as this messes with the internal offset calculation of the peaks, might be fixed by fitting a  $y = k * x + m$  function to the peaks for offset calculation.

keyword arguments: y\_axis -- A list containing the signal over which to find peaks

**x\_axis** -- A x-axis whose values correspond to the y\_axis list and is used in the return to specify the position of the peaks.

**points** -- How many points around the peak should be used during curve fitting (default: 31)

**lock\_frequency** -- Specifies if the frequency argument of the model function should be locked to the value calculated from the raw peaks or if optimization process may tinker with it. (default: False)

**return:** two lists [max\_peaks, min\_peaks] containing the positive and negative peaks respectively. Each cell of the lists contains a tuple of: (position, peak\_value) to get the average peak value do: `np.mean(max_peaks, 0)[1]` on the results to unpack one of the lists into x, y coordinates do: `x, y = zip(*max_peaks)`

`pysisyphus.peakdetect.peakdetect_sine_locked(y_axis, x_axis, points=31)`

Convenience function for calling the 'peakdetect\_sine' function with the lock\_frequency argument as True.

keyword arguments: y\_axis -- A list containing the signal over which to find peaks x\_axis -- A x-axis whose values correspond to the y\_axis list and is used

in the return to specify the position of the peaks.

**points** -- How many points around the peak should be used during curve fitting (default: 31)

return: see the function 'peakdetect\_sine'

`pysisyphus.peakdetect.peakdetect_spline(y_axis, x_axis, pad_len=20)`

Performs a b-spline interpolation on the data to increase resolution and send the data to the 'peakdetect\_zero\_crossing' function for peak detection.

Omitting the x\_axis is forbidden as it would make the resulting x\_axis value silly if it was returned as the index 50.234 or similar.

will find the same amount of peaks as the 'peakdetect\_zero\_crossing' function, but might result in a more precise value of the peak.

keyword arguments: `y_axis` -- A list containing the signal over which to find peaks

**x\_axis** -- A x-axis whose values correspond to the y\_axis list and is used in the return to specify the position of the peaks. x-axis must be equally spaced.

**pad\_len** -- By how many times the time resolution should be increased by, e.g. 1 doubles the resolution. (default: 20)

**return:** two lists [`max_peaks`, `min_peaks`] containing the positive and negative peaks respectively. Each cell of the lists contains a tuple of: (position, peak\_value) to get the average peak value do: `np.mean(max_peaks, 0)[1]` on the results to unpack one of the lists into x, y coordinates do: `x, y = zip(*max_peaks)`

`pysisiphus.peakdetect.peakdetect_zero_crossing(y_axis, x_axis=None, window=11)`

Function for detecting local maxima and minima in a signal. Discovers peaks by dividing the signal into bins and retrieving the maximum and minimum value of each the even and odd bins respectively. Division into bins is performed by smoothing the curve and finding the zero crossings.

Suitable for repeatable signals, where some noise is tolerated. Executes faster than 'peakdetect', although this function will break if the offset of the signal is too large. It should also be noted that the first and last peak will probably not be found, as this function only can find peaks between the first and last zero crossing.

keyword arguments: `y_axis` -- A list containing the signal over which to find peaks

**x\_axis** -- A x-axis whose values correspond to the y\_axis list and is used in the return to specify the position of the peaks. If omitted an index of the y\_axis is used. (default: None)

**window** -- the dimension of the smoothing window; should be an odd integer (default: 11)

**return:** two lists [`max_peaks`, `min_peaks`] containing the positive and negative peaks respectively. Each cell of the lists contains a tuple of: (position, peak\_value) to get the average peak value do: `np.mean(max_peaks, 0)[1]` on the results to unpack one of the lists into x, y coordinates do: `x, y = zip(*max_peaks)`

`pysisiphus.peakdetect.zero_crossings(y_axis, window_len=11, window_f='hanning', offset_corrected=False)`

Algorithm to find zero crossings. Smooths the curve and finds the zero-crossings by looking for a sign change.

keyword arguments: `y_axis` -- A list containing the signal over which to find zero-crossings

**window\_len** -- the dimension of the smoothing window; should be an odd integer (default: 11)

**window\_f** -- the type of window from 'flat', 'hanning', 'hamming', 'bartlett', 'blackman' (default: 'hanning')

`offset_corrected` -- Used for recursive calling to remove offset when needed

return: the index for each zero-crossing

`pysisiphus.peakdetect.zero_crossings_sine_fit(y_axis, x_axis, fit_window=None, smooth_window=11)`

Detects the zero crossings of a signal by fitting a sine model function around the zero crossings:  $y = A * \sin(2 * \pi * Hz * (x - \tau)) + k * x + m$  Only tau (the zero crossing) is varied during fitting.

Offset and a linear drift of offset is accounted for by fitting a linear function the negative respective positive raw peaks of the wave-shape and the amplitude is calculated using data from the offset calculation i.e. the 'm' constant from the negative peaks is subtracted from the positive one to obtain amplitude.

Frequency is calculated using the mean time between raw peaks.

Algorithm seems to be sensitive to first guess e.g. a large `smooth_window` will give an error in the results.

keyword arguments: `y_axis` -- A list containing the signal over which to find peaks

**`x_axis`** -- A x-axis whose values correspond to the `y_axis` list

and is used in the return to specify the position of the peaks. If omitted an index of the `y_axis` is used.  
(default: None)

**`fit_window`** -- Number of points around the approximate zero crossing that

should be used when fitting the sine wave. Must be small enough that no other zero crossing will be seen.  
If set to none then the mean distance between zero crossings will be used (default: None)

**`smooth_window`** -- the dimension of the smoothing window; should be an odd

integer (default: 11)

return: A list containing the positions of all the zero crossings.

### 16.1.18 pysisyphus.plot module

`pysisyphus.plot.CDD_PNG_FNS = 'cdd_png_fns'`

Default to  $\text{kJ mol}^{-1}$ . `DE_LABEL` and `get_en_conv` will be overwritten when `run()` is executed. Both are still defined here, to make the functions from `plot.py` importable.

`pysisyphus.plot.anim_cos(cart_coords, energies)`

`pysisyphus.plot.get_en_conv()`

`pysisyphus.plot.get_force_unit(coord_type)`

`pysisyphus.plot.list_h5_groups(h5_fn)`

`pysisyphus.plot.load_h5(h5_fn, h5_group, datasets=None, attrs=None)`

`pysisyphus.plot.parse_args(args)`

`pysisyphus.plot.plot_afir(h5_fn='afir.h5', h5_group='afir')`

`pysisyphus.plot.plot_all_energies(h5)`

`pysisyphus.plot.plot_cos_energies(h5_fn='optimization.h5', h5_group='opt')`

`pysisyphus.plot.plot_cos_forces(h5_fn='optimization.h5', h5_group='opt', last=15)`

`pysisyphus.plot.plot_cycle(cart_coords, energies)`

`pysisyphus.plot.plot_gau(gau_fns, num=50)`

`pysisyphus.plot.plot_irc()`

`pysisyphus.plot.plot_irc_h5(h5, title=None)`

`pysisyphus.plot.plot_md(h5_group='run')`

`pysisyphus.plot.plot_opt(h5_fn='optimization.h5', h5_group='opt')`

`pysisyphus.plot.plot_overlaps(h5, thresh=0.1)`

`pysisyphus.plot.plot_scan(h5_fn='scan.h5')`

`pysisyphus.plot.plot_trj_energies(trj)`

Parse comments of .xyz/trj as energies and plot.

`pysisyphus.plot.render_cdds(h5)`

`pysisyphus.plot.run()`

`pysisyphus.plot.spline_plot_cycles(cart_coords, energies)`

### 16.1.19 pysisyphus.run module

**class** `pysisyphus.run.RunResult` (*preopt\_first\_geom, preopt\_last\_geom, cos, cos\_opt, ts\_geom, ts\_opt, end\_geoms, irc, irc\_geom, mdp\_result, opt\_geom, opt, calced\_geoms, calced\_results, stochastic, calc\_getter, scan\_geoms, scan\_vals, scan\_energies, perf\_results*)

Bases: tuple

**calc\_getter**

Alias for field number 15

**calced\_geoms**

Alias for field number 12

**calced\_results**

Alias for field number 13

**cos**

Alias for field number 2

**cos\_opt**

Alias for field number 3

**end\_geoms**

Alias for field number 6

**irc**

Alias for field number 7

**irc\_geom**

Alias for field number 8

**mdp\_result**

Alias for field number 9

**opt**

Alias for field number 11

**opt\_geom**

Alias for field number 10

**perf\_results**

Alias for field number 19

**preopt\_first\_geom**

Alias for field number 0

**preopt\_last\_geom**

Alias for field number 1

**scan\_energies**

Alias for field number 18

**scan\_geoms**

Alias for field number 16

**scan\_vals**

Alias for field number 17

**stochastic**

Alias for field number 14

**ts\_geom**

Alias for field number 4

**ts\_opt**

Alias for field number 5

`pysisyphus.run.check_asserts(results, run_dict)`

`pysisyphus.run.copy_yaml_and_geometries(run_dict, yaml_fn, dest_and_add_cp, new_yaml_fn=None)`

`pysisyphus.run.do_clean(force=False)`

Deletes files from previous runs in the cwd. A similar function could be used to store everything ...

`pysisyphus.run.do_rmsds(xyz, geoms, end_fns, end_geoms, preopt_map=None, similar_thresh=0.25)`

`pysisyphus.run.get_calc_closure(base_name, calc_key, calc_kwargs, iter_dict=None, index=None)`

`pysisyphus.run.get_defaults(conf_dict, T_default=298.15, p_default=101325)`

`pysisyphus.run.get_last_calc_cycle()`

`pysisyphus.run.main(run_dict, restart=False, yaml_dir='.', scheduler=None)`

`pysisyphus.run.parse_args(args)`

`pysisyphus.run.print_bibtex()`

`pysisyphus.run.print_header()`

Generated from <https://asciartgen.now.sh/?s=pysisyphus&style=colossal>

`pysisyphus.run.run()`

`pysisyphus.run.run_calculations(geoms, calc_getter, scheduler=None, assert_track=False, run_func=None)`

`pysisyphus.run.run_endopt(irc, endopt_key, endopt_kwargs, calc_getter)`

`pysisyphus.run.run_from_dict(run_dict, cwd=None, set_defaults=True, yaml_fn=None, cp=None, scheduler=None, clean=False, fclean=False, version=False, restart=False)`

`pysisyphus.run.run_irc(geom, irc_key, irc_kwargs, calc_getter)`

`pysisyphus.run.run_md(geom, calc_getter, md_kwargs)`

`pysisyphus.run.run_mdp(geom, calc_getter, mdp_kwargs)`

```
pysisyphus.run.run_preopt(first_geom, last_geom, calc_getter, preopt_key, preopt_kwargs)
pysisyphus.run.run_scan(geom, calc_getter, scan_kwargs, callback=None)
pysisyphus.run.run_stochastic(stoc)
pysisyphus.run.run_tsopt_from_cos(cos, tsopt_key, tsopt_kwargs, calc_getter=None, ovlp_thresh=0.4,
coordinate_union='bonds')
pysisyphus.run.setup_run_dict(run_dict)
```

### 16.1.20 pysisyphus.socket\_helper module

```
pysisyphus.socket_helper.get_fmts(cartesians)
pysisyphus.socket_helper.recv_closure(sock, hdrlen, fmts, verbose=False)
pysisyphus.socket_helper.send_closure(sock, hdrlen, fmts, verbose=False)
```

### 16.1.21 pysisyphus.testing module

```
class pysisyphus.testing.DummyMark(available)
    Bases: object
pysisyphus.testing.available(calculator, **kwargs)
pysisyphus.testing.module_available(calculator)
pysisyphus.testing.using(calculator, set_pytest_mark=True)
    Calling disabling set_pytest_mark avoids a runtime dependency on pytest.
```

### 16.1.22 pysisyphus.thermo module

```
pysisyphus.thermo.get_thermoanalysis_from_hess_h5(h5_fn, T=298.15, p=101325, point_group='c1',
return_geom=False)
pysisyphus.thermo.print_thermoanalysis(thermo, geom=None, is_ts=False, unit='joule', level=0,
title=None)
    Print thermochemical analysis.
```

### 16.1.23 pysisyphus.trj module

```
exception pysisyphus.trj.GotNoGeometryException
    Bases: Exception
pysisyphus.trj.align(geoms)
    Align all geometries onto the first using partial procrustes.
pysisyphus.trj.append(geoms)
pysisyphus.trj.center(geoms)
```

`pysisyphus.trj.centerm(geoms)`

`pysisyphus.trj.dump_geoms(geoms, fn_base, trj_infix=", dump_trj=True, dump_xyz=True, dump_pdb=False, ang=False)`

`pysisyphus.trj.every(geoms, every_nth)`

`pysisyphus.trj.frag_sort(geoms)`

`pysisyphus.trj.get(geoms, index)`

`pysisyphus.trj.get_geoms(xyz_fns, coord_type='cart', geom_kwargs=None, union=False, same_prims=True, quiet=False, interpol_kwargs=None)`

Returns a list of Geometry objects in the given coordinate system and interpolates if necessary.

`pysisyphus.trj.get_interactively(geoms)`

`pysisyphus.trj.match(ref_geom, geom_to_match)`

`pysisyphus.trj.origin(geoms)`

`pysisyphus.trj.parse_args(args)`

`pysisyphus.trj.print_internals(geoms, filter_atoms=None, add_prims="")`

`pysisyphus.trj.read_geoms(xyz_fns, coord_type='cart', geom_kwargs=None)`

`pysisyphus.trj.run()`

`pysisyphus.trj.shake(geoms, scale=0.1, seed=None)`

`pysisyphus.trj.spline_redistribute(geoms)`

`pysisyphus.trj.standard_orientation(geoms)`

`pysisyphus.trj.standardize_geoms(geoms, coord_type, geom_kwargs, same_prims=True, union=False)`

`pysisyphus.trj.translate(geoms, trans)`

## 16.1.24 pysisyphus.version module

## 16.1.25 pysisyphus.xyzloader module

`pysisyphus.xyzloader.coords_to_trj(trj_fn, atoms, coords_list, comments=None)`

`pysisyphus.xyzloader.make_trj_str(atoms, coords_list, comments=None)`

`pysisyphus.xyzloader.make_trj_str_from_geoms(geoms, comments=None, energy_comments=False)`

`pysisyphus.xyzloader.make_xyz_str(atoms, coords, comment="")`

`pysisyphus.xyzloader.parse_trj_file(trj_fn, with_comments=False)`

`pysisyphus.xyzloader.parse_trj_str(trj_str, with_comments=False)`

`pysisyphus.xyzloader.parse_xyz_file(xyz_fn, with_comment=False)`

`pysisyphus.xyzloader.parse_xyz_str(xyz_str, with_comment)`

Parse a xyz string.

**xyz\_str**

[str] The contents of a .xyz file.

**with\_comment**

[bool] Return comment line if True.

**Returns**

- **atoms** (*list*) -- List of length N (N = number of atoms) holding the element symbols.
- **coords** (*np.array*) -- An array of shape (N, 3) holding the xyz coordinates.
- **comment\_line** (*str*, *optional*) -- Comment line if `with_comment` argument was True.

`pysisyphus.xyzloader.split_xyz_str(xyz_str)`

Example:

**xyz:**

1

X -1.2 1.4 0.0 1

X 2.0 4.0 0.0

`pysisyphus.xyzloader.write_geoms_to_trj(geoms, fn, comments=None)`

### 16.1.26 pysisyphus.yaml\_mods module

`pysisyphus.yaml_mods.get_constructor(unit)`

`pysisyphus.yaml_mods.get_loader(units=['Eh', 'kJpermol', 'kcalpermol', 'eV', 'fs', 'ps', 'ns', 'a0', 'angstrom', 'nm'])`

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