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# **PyCCE**

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**Mykyta Onizhuk**

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# GETTING STARTED

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## THEORETICAL BACKGROUND

This document contains a brief list of the coupling parameters between the central and the bath spins used in **PyCCE**, a description of the qubit dephasing, and a summary of the cluster correlation expansion (CCE) method. You can find more details in the following references<sup>1,2,3</sup>.

### 1.1 Hamiltonian

The **PyCCE** package allows one to simulate the dynamics of a central spin interacting with a spin bath through the following Hamiltonian:

$$\hat{H} = \hat{H}_S + \hat{H}_{SB} + \hat{H}_B$$

Where  $\hat{H}_S$  is the Hamiltonian of the free central spin,  $\hat{H}_{SB}$  denotes interactions between central spin and a spin belonging to the bath, and  $\hat{H}_B$  are intrinsic bath spin interactions:

$$\begin{aligned}\hat{H}_S &= \mathbf{S}\mathbf{D}\mathbf{S} + \mathbf{B}\gamma_S\mathbf{S} \\ \hat{H}_{SB} &= \sum_i \mathbf{S}\mathbf{A}_i\mathbf{I}_i \\ \hat{H}_B &= \sum_i \mathbf{I}_i\mathbf{P}_i\mathbf{I}_i + \mathbf{B}\gamma_i\mathbf{I}_i + \sum_{i>j} \mathbf{I}_i\mathbf{J}_{ij}\mathbf{I}_j\end{aligned}$$

Where  $\mathbf{S} = (\hat{S}_x, \hat{S}_y, \hat{S}_z)$  are the components of spin operators of the central spin,  $\mathbf{I} = (\hat{I}_x, \hat{I}_y, \hat{I}_z)$  are the components of the bath spin operators, and  $\mathbf{B} = (B_x, B_y, B_z)$  is an external applied magnetic field.

The interactions are described by the following tensors that are either required to be input by user or can be generated by the package itself (see *Hamiltonian Parameters Input* for details):

- **D** (**P**) is the self-interaction tensor of the central spin (bath spin). For the electron spin, the tensor corresponds to the zero-field splitting (ZFS) tensor. For nuclear spins corresponds to the quadrupole interactions tensor.
- $\gamma_i$ -spin describing the interaction of the spin and the external magnetic field  $B$ . We assume that for the bath spins, it is isotropic.
- **A** is the interaction tensor between central and bath spins. In the case of the nuclear spin bath, it corresponds to the hyperfine couplings.
- **J** is the interaction tensor between bath spins.

---

<sup>1</sup> Mykyta Onizhuk and Giulia Galli. "PyCCE: A Python Package for Cluster Correlation Expansion Simulations of Spin Qubit Dynamic". *Adv. Theory Simul.* 2021, 2100254, <https://onlinelibrary.wiley.com/doi/10.1002/adts.202100254>

<sup>2</sup> Wen Yang and Ren-Bao Liu. "Quantum many-body theory of qubit decoherence in a finite-size spin bath". *Phys. Rev. B* 78, p. 085315, <https://link.aps.org/doi/10.1103/PhysRevB.78.085315>

<sup>3</sup> Mykyta Onizhuk et al. "Probing the Coherence of Solid-State Qubits at Avoided Crossings". *PRX Quantum* 2, p. 010311. <https://link.aps.org/doi/10.1103/PRXQuantum.2.010311>.

## 1.2 Qubit dephasing

Usually, two coherence times are measured to characterize the loss of a qubit coherence -  $T_1$  and  $T_2$ .  $T_1$  defines the timescale over which the qubit population is thermalized;  $T_2$  describes a purely quantum phenomenon - the loss of the phase of the qubit's superposition state.

In the pure dephasing regime ( $T_1 \gg T_2$ ) the decoherence of the central spin is completely determined by the decay of the off diagonal element of the density matrix of the qubit.

Namely, if the qubit is initially prepared in the  $|\psi\rangle = \frac{1}{\sqrt{2}}(|0\rangle + e^{i\phi}|1\rangle)$  state, the loss of the relative phase of the  $|0\rangle$  and  $|1\rangle$  levels is characterized by the coherence function:

$$\mathcal{L}(t) = \frac{\langle 1 | \hat{\rho}_S(t) | 0 \rangle}{\langle 1 | \hat{\rho}_S(0) | 0 \rangle} = \frac{\langle \hat{\sigma}_-(t) \rangle}{\langle \hat{\sigma}_-(0) \rangle}$$

Where  $\hat{\rho}_S(t)$  is the density matrix of the central spin and  $|0\rangle$  and  $|1\rangle$  are qubit levels.

The cluster correlation expansion (CCE) method was first introduced in ref.<sup>2</sup>. The core idea of the CCE approach is that the spin bath-induced decoherence can be factorized into set of irreducible contributions from the bath spin clusters. Written in terms of the coherence function:

$$\mathcal{L}(t) = \prod_C \tilde{L}_C = \prod_i \tilde{L}_{\{i\}} \prod_{i,j} \tilde{L}_{\{ij\}} \dots$$

Where each cluster contribution is defined recursively as:

$$\tilde{L}_C = \frac{L_C}{\prod_{C'} \tilde{L}_{C' \subset C}}$$

Where  $L_C$  is a coherence function of the qubit, interacting only with the bath spins in a given cluster  $C$  (with the cluster Hamiltonian  $\hat{H}_C$ ), and  $\tilde{L}_{C'}$  are contributions of  $C'$  subcluster of  $C$ .

For example, the contribution of the single spin  $i$  is equal to the coherence function of the bath with one isolated spin  $i$ :

$$\tilde{L}_i = L_i$$

The contribution of pair of spins  $i$  and  $j$  is equal to:

$$\tilde{L}_{ij} = \frac{L_{ij}}{\tilde{L}_i \tilde{L}_j}$$

and so on.

Maximum size of the cluster included into the expansion determines the order of CCE approximation. For example, in the CCE2 approximation, only contributions up to spin pairs are included, and in CCE3 - up to triplets of bath spins are included, etc.

The way the coherence function for each cluster is computed slightly varies between depending on whether the conventional or generalized CCE method is used.

### 1.2.1 Conventional CCE

In the original formulation of the CCE method, the total Hamiltonian of the system is reduced to the sum of two effective Hamiltonians, conditioned on the qubit levels of the central spin:

$$\hat{H} = 00 \otimes \hat{H}^{(0)} + 11 \otimes \hat{H}^{(1)}$$

Where  $\hat{H}^{(\alpha)}$  is an effective Hamiltonian acting on the bath when the central qubit is in the  $\alpha$  state ( $\alpha = 0, 1$  is one of the two eigenstates of the  $\hat{H}_S$  chosen as qubit levels).

Given an initial qubit state  $\psi = \frac{1}{\sqrt{2}}(0 + e^{i\phi}1)$  and an initial state of the bath spin cluster  $C$  characterized by the density matrix  $\hat{\rho}_C$ , the coherence function of the qubit interacting with the cluster  $C$  is computed as:

$$L_C(t) = \text{Tr}[\hat{U}_C^{(0)}(t)\hat{\rho}_C\hat{U}_C^{(1)\dagger}(t)]$$

Where  $\hat{U}_C^{(\alpha)}(t)$  is time propagator defined in terms of the effective Hamiltonian  $\hat{H}_C^{(\alpha)}$  and the number of decoupling pulses. Note that  $\hat{H}_C^{(\alpha)}$  here includes only degrees of freedom of the given cluster.

For free induction decay (FID) the time propagators are trivial:

$$\hat{U}_C^{(0)} = e^{-\frac{i}{\hbar}\hat{H}_C^{(0)}t}; \hat{U}_C^{(1)} = e^{-\frac{i}{\hbar}\hat{H}_C^{(1)}t}$$

And for the generic decoupling sequence with  $N$  (even) decoupling pulses applied at  $t_1, t_2 \dots t_N$  we write:

$$\hat{U}^{(\alpha)}(t) = e^{-\frac{i}{\hbar}\hat{H}_C^{(\alpha)}(t_N - t_{N-1})} e^{-\frac{i}{\hbar}\hat{H}_C^{(\beta)}(t_{N-1} - t_{N-2})} \dots e^{-\frac{i}{\hbar}\hat{H}_C^{(\beta)}(t_2 - t_1)} e^{-\frac{i}{\hbar}\hat{H}_C^{(\alpha)}t_1}$$

Where  $\alpha = 0, 1$  and  $\beta = 1, 0$  accordingly (when  $\alpha = 0$  one should take  $\beta = 1$  and vice versa).  $t = \sum_i t_i$  is the total evolution time. In sequences with odd number of pulses  $N$ , the leftmost propagator is the exponent of  $\hat{H}_C^{(\beta)}$ .

## 1.2.2 Generalized CCE

Instead of projecting the total Hamiltonian on the qubit levels, one may directly include the central spin degrees of freedom to each clusters. We refer to such formulation as gCCE.

In this case we write the cluster Hamiltonian as:

$$\begin{aligned} \hat{H}_C = & \text{SDS} + \mathbf{B}\gamma_S\mathbf{S} + \sum_{i \in C} \mathbf{S}\mathbf{A}_i\mathbf{I}_i + \sum_{i \in C} \mathbf{I}_i\mathbf{P}_i\mathbf{I}_i + \mathbf{B}\gamma_i\mathbf{I}_i + \\ & \sum_{i < j \in C} \mathbf{I}_i\mathbf{J}_{ij}\mathbf{I}_j + \sum_{a \notin C} \mathbf{S}\mathbf{A}_a\langle \mathbf{I}_a \rangle + \sum_{i \in C, a \notin C} \mathbf{I}_i\mathbf{J}_{ia}\langle \mathbf{I}_a \rangle \end{aligned}$$

And the coherence function of the cluster  $L_C(t)$  is computed as:

$$L_C(t) = 0\hat{U}_C(t)\hat{\rho}_{C+S}\hat{U}_C^\dagger(t)1$$

Where  $\hat{\rho}_{C+S} = \hat{\rho}_C \otimes \hat{\rho}_S$  is the combined initial density matrix of the bath spins' cluster and central spin.

Further details on the theoretical background are available in the references below.





## QUICK START

The generic workflow of the simulation includes first the generation of the spin bath in the material, and second carrying the CCE dynamics calculations for the qubit interacting with this spin bath.

### 2.1 Base Units

- All coupling constants are given in kHz.
- Timesteps are in millisecond (ms).
- Distances are in angstrom (Å).
- Gyromagnetic ratios are given in  $\text{rad} \cdot \text{ms}^{-1} \cdot \text{G}^{-1}$ .
- Quadrupole constants are given in barn ( $10^{-28} \text{ m}^2$ ).
- Magnetic field is given in Gauss (G).

### 2.2 Simple Example

The simplest example includes the following steps:

1. Generate the BathCell object. Here we use the interface with ase which can effortlessly generate unit cells of many materials. As an example, we import the diamond structure.

```
import numpy as np
import pycce as pc
from ase.build import bulk

cell = pc.BathCell.from_ase(bulk('C', 'diamond', cubic=True))
```

2. Using the BathCell object, generate spin bath of the most common isotopes in the material. Here we generate the spin bath of size 200 Angstrom and remove one carbon, where the spin of interest is located, from the diamond crystal lattice.

```
atoms = cell.gen_supercell(200, remove=('C', [0, 0, 0]))
```

This function returns the BathArray instance, which contains names of the bath spins in 'N', their coordinates in angstrom in 'xyz', empty arrays of hyperfine couplings in kHz in 'A', and quadrupole couplings in kHz in 'Q' namefields. The hyperfine couplings will be generated by Simulator in the next step. For alternative ways to define hyperfine couplings see *Hamiltonian Parameters Input*.

3. Setup the `Simulator` using the generated spin bath. The first required argument is the total spin of the central spin, `r_bath`, `r_dipole` and `order` are convergence parameters (see the *Tutorials* for examples of convergence), `magnetic_field` is the external applied magnetic field along the z-axis, and `pulses` is the number of decoupling  $\pi$  pulses in Carr-Purcell-Meiboom-Gill (CPMG) sequence (0 - FID, 1 - Hahn-echo, 2 - Carr Purcell etc.).

```
calc = pc.Simulator(0.5, position=[0, 0, 0], bath=atoms, r_bath=40,  
                   r_dipole=6, order=2, magnetic_field=500, pulses=1)
```

The hyperfine couplings are automatically generated at this step assuming point dipole-dipole interactions between central spin and bath spins.

4. Compute the coherence function of the qubit using `.compute` method of the `Simulator` object with conventional CCE.

```
time_points = np.linspace(0, 2, 101)  
coherence = calc.compute(time_points)
```

This function outputs Numpy array with the same shape as the `time_points` and contains the coherence function computed at each time step. By default `compute` method uses the conventional CCE to compute the coherence function.

More detailed examples of **PyCCE** usage are available in the tutorials.

The examples below are available as Jupyter notebooks in the Github repository.

### 3.1 NV Center in Diamond

In this tutorial we will go over the main steps of running CCE calculations for the NV center in diamond with the **PyCCE** module. Those include:

- Generating the spin bath using the `pycce.BathCell` instance.
- Setting up properties of the `pycce.Simulator` instance.
- Running the calculations with the `Simulator.compute` function.

We will compute the Hahn-echo coherence function (with decoupling  $\pi$ -pulse applied) using the following available methods:

- Conventional CCE.
- Generalized CCE (gCCE).
- gCCE with Monte-Carlo bath sampling.

```
[1]: import numpy as np
import matplotlib.pyplot as plt
import pandas as pd

import sys
import pycce as pc
import ase

from mpl_toolkits import mplot3d

seed = 8805
np.random.seed(seed)
np.set_printoptions(suppress=True, precision=5)
```

### 3.1.1 Generate nuclear spin bath

Building a supercell of nuclear spins from the `ase.Atoms` object.

#### Build BathCell

To generate cell it from `ase.atoms` object, use classmethod `BathCell.from_ase`.

```
[2]: from ase.build import bulk

# Generate unitcell from ase
diamond = bulk('C', 'diamond', cubic=True)
diamond = pc.bath.BathCell.from_ase(diamond)
```

The following attributes are created with this initialization:

- `.cell` is ndarray containing information of lattice vectors. Each **column** is a lattice vector in cartesian coordinates.
- `.atoms` is a dictionary with keys corresponding to the atom name, and each item is a list of the coordinates in cell coordinates.

```
[3]: print('Cell\n', diamond.cell)
print('\nAtoms\n', diamond.atoms)

Cell
[[3.57 0.  0.  ]
 [0.  3.57 0.  ]
 [0.  0.  3.57]]

Atoms
defaultdict(<class 'list'>, {'C': [array([0., 0., 0.]), array([0.25, 0.25, 0.25]),
↪ array([0. , 0.5, 0.5]), array([0.25, 0.75, 0.75]), array([0.5, 0. , 0.5]), array([0.75,
↪ 0.25, 0.75]), array([0.5, 0.5, 0. ]), array([0.75, 0.75, 0.25])])})
```

#### Populate BathCell with isotopes

The **PyCCE** package uses EasySpin database of the concentrations of all common stable isotopes with non-zero spin, however the user can provide custom concentrations.

Use function `BathCell.add_isotopes` to add one (or several) isotopes of the element. Each isotope is initialized with tuple containing name of the isotope and its concentration.

Name of the isotope includes the number and element symbol, provided in the `atoms` object. As an output, the `BathCell.add_isotopes` method returns view on dictionary `BathCell.isotopes` which can be modified directly. Structure of the dictionary-like object:

```
{element_1: {isotope_1: concentration, isotope_2: concentration},
 element_2: {isotope_3: concentration ...}}
```

```
[4]: # Add types of isotopes
diamond.add_isotopes(('13C', 0.011))

[4]: defaultdict(dict, {'C': {'13C': 0.011}})
```

Isotopes may also be directly added to `BathCell.isotopes`. For example, below we are adding an isotope without the nuclear spin:

```
[5]: diamond.isotopes['C']['14C'] = 0.001
```

### Set z-direction of the bath (optional)

In the `Simulator` object everything is set in  $S_z$  basis. When the quantization axis of the defect does not align with the  $(0, 0, 1)$  direction of the crystal axis, the user needs to define the axis.

If one wants to specify the complete rotation of cartesian axes, one can provide a rotation matrix to rotate the cartesian reference frame with respect to the cell coordinates by calling the `BathCell.rotate` method.

```
[6]: # set z direction of the defect
diamond.zdir = [1, 1, 1]
```

### Generate spin bath

To generate the spin bath, use the `BathCell.gen_supercell` method. First argument is the linear size of the supercell (minimum distance between any two faces of the supercell is equal to or larger than this parameter). Additional keyword arguments are `remove` and `add`.

`remove` takes a tuple or list of tuples as an argument. First element of each tuple is the name of the **atom** at that location, second element - coordinates in unit cell coordinates. If such atoms are found in the supercell, they are removed from it.

`add` takes a tuple or list of tuples as an argument. First element of each tuple is the name of the **isotope** at that location, second element - coordinates in unit cell coordinates. Each of the specified isotopes will be added in the final supercell at specified locations.

```
[7]: # Add the defect. remove and add atoms at the positions (in cell coordinates)
atoms = diamond.gen_supercell(200, remove=[('C', [0., 0, 0]),
                                           ('C', [0.5, 0.5, 0.5])],
                              add=('14N', [0.5, 0.5, 0.5]),
                              seed=seed)
```

```
/home/onizhuk/midway/codes_development/pyCCE/pycce/bath/array.py:168: UserWarning: Spin_
↪type for 14C was not provided and was not found in common isotopes.
obj[n] = array[n]
```

Note, that because the `14C` isotope doesn't have a spin, **PyCCE** does not find it in common isotopes, and raises a warning. We have to provide `SpinType` for it separately, or define the properties as follows:

```
[8]: atoms['14C'].gyro = 0
atoms['14C'].spin = 0
```

### 3.1.2 BathArray Structure

The bath spins are stored in the `BathArray` object - a subclass of `np.ndarray` with fixed datastructure:

- `N` field `dtype('<U16')` contains the names of bath spins.
- `xyz` field `dtype('<f8', (3,))` contains the positions of bath spins (in Å).
- `A` field `dtype('<f8', (3, 3))` contains the hyperfine coupling of bath spins (in kHz).
- `Q` field `dtype('<f8', (3, 3))` contains the quadrupole tensor of bath spins (in kHz) (Relevant for spin  $\geq 1$ ).

All of the fields are accessible as attributes of `BathArray`. Additionally, the subarrays of the specific spins are accessible with their name as indicated above.

Upon generation of the array from the cell, the `Q` and `A` fields are empty. The Hyperfine couplings will be automatically computed by the `Simulator` object, however the quadrupole couplings must be set by the user.

The additional attributes allow one to access `SpinType` properties:

- `name` returns the spin name or array of spin names;
- `spin` returns the value of the spin or array of ones;
- `gyro` returns gyromagnetic ratios of the spins;
- `q` returns quadrupole constants of the spins;
- `detuning` returns detunings of the spins (See definition below).

For example, below we print out the attributes of the first two spins in the `BathArray`.

```
[9]: print('Names\n', atoms[:2].N)
      print('\nCoordinates\n', atoms[:2].xyz)
      print('\nHyperfine tensors\n', atoms[:2].A)
      print('\nQuadrupole tensors\n', atoms[:2].Q)
```

Names

```
['13C' '13C']
```

Coordinates

```
[[-13.97678 -1.48178 -92.75132]
 [ 27.89939  42.17939 -45.86038]]
```

Hyperfine tensors

```
[[[0. 0. 0.]
 [0. 0. 0.]
 [0. 0. 0.]]
```

```
[[[0. 0. 0.]
 [0. 0. 0.]
 [0. 0. 0.]]]
```

Quadrupole tensors

```
[[[0. 0. 0.]
 [0. 0. 0.]
 [0. 0. 0.]]
```

```
[[[0. 0. 0.]
 [0. 0. 0.]
 [0. 0. 0.]]]
```

The properties of spin types (gyromagnetic ratio, quadrupole moment, etc) are stored in the `BathArray.types` attribute, which is an instance of `SpinDict` containing `SpinType` classes. For most known isotopes `SpinType` can be found in the `pycce.common_isotopes` dictionary, and is set by default (including electron spin-1/2, which is denoted by setting `N = e`). The user can add additional `SpinType` objects, by calling `BathArray.add_type` method or setting elements of `SpinDict` directly. For details of the first approach see documentation of `SpinDict.add_type` method.

The direct setting of types is rather simple. The user can set elements of `SpinDict` with tuple, containing:

- **(spin, gyromagnetic ratio, quadrupole moment \*(optional)\*, detuning \*(optional)\*, )**

OR

- **(isotope, spin, gyromagnetic ratio, quadrupole moment \*(optional)\*, detuning \*(optional)\*, )**

where:

- **isotope** (*str*) is the name of the given spin (same one as in `N` field of `BathArray`) to define new `SpinType` object. The key of `SpinDict` **has** to be the correct name of the spin (“isotope” field in the tuple).
- **spin** (*float*) is the total spin of the given bath spin.
- **gyromagnetic ratio** (*float*) is the gyromagnetic ratio of the given bath spin.
- **quadrupole moment** (*float*) is the quadrupole moment of the given bath spin. Relevant only when electric field gradient are used to generate quadrupole couplings for spins, stored in the `BathArray`, with `BathArray.from_efg` method.
- **detuning** (*float*) is an additional energy splitting for model spins, included as an extra  $+\omega\hat{S}_z$  term in the Hamiltonian, where  $\omega$  is the detuning.

Units of gyromagnetic ratio are rad / ms / G, quadrupole moments are given in barn, detunings are given in kHz.

```
[10]: # Several ways to set SpinDict elements
atoms.types['14C'] = 0, 0, 0
atoms.types['Y'] = ('Y', 0, 0, 0)
atoms.types['A'] = pc.SpinType('A', 0, 0, 0)

print(atoms.types)

SpinDict(13C: (0.5, 6.7283), 14N: (1.0, 1.9338, 0.0204), 14C: (0.0, 0.0000), ...)
```

### 3.1.3 Simulator class

The parameters of the CCE simulator engine.

Main parameters to consider:

- **alpha** — first qubit state in  $S_z$  basis.
- **beta** — second qubit state in  $S_z$  basis.
- **position** — coordinates of the central spin.
- **bath** — spin bath in any specified format. Can be either:
  - Instance of `BathArray` class;
  - `ndarray` with `dtype([(('N', np.unicode_, 16), ('xyz', np.float64, (3,)))])` containing names of bath spins (same ones as stored in `self.ntype`) and positions of the spins in angstroms;
  - The name of the `.xyz` text file containing 4 columns: name of the bath spin and xyz coordinates in A.
- **r\_bath** — cutoff radius around the central spin for the bath.

- `order` — maximum size of the cluster.
- `r_dipole` — cutoff radius for the pairwise distance to consider two nuclear spins to be connected.
- `magnetic_field` — applied magnetic field. Can also be provided during the simulation run.
- `pulses` — number of pulses in Carr-Purcell-Meiboom-Gill (CPMG) sequence or the pulse sequence itself.

Additional parameters, necessary for the generalized CCE:

- `D` — Zero field splitting (ZFS) tensor of the central spin, or axial ZFS.
- `E` — if `D` is provided as axial ZFS, then `E` sets transverse ZFS.

For the full description see the documentation of the `Simulator` object.

First we setup a “mock” instance of `Simulator` to visualize the smaller part of the bath around the central spin.

```
[11]: # Setting the runner engine
mock = pc.Simulator(spin=1, position=[0,0,0],
                   bath=atoms, r_bath=20,
                   r_dipole=6, order=3)
```

During the initialization, depending on the provided keyword arguments several methods may be called:

- `Simulator.read_bath` is called if keyword `bath` is provided. It may take several additional arguments:
  - `r_bath` - cutoff distance from the qubit for the bath.
  - `skiprows` - if `bath` is provided as `.xyz` file, this argument tells how many rows to skip when reading the file.
  - `external_bath` - `BathArray` instance, which contains bath spins with pre defined hyperfines to be used.
  - `hyperfine` - defines the way to compute hyperfine couplings. If it is not given and `bath` doesn't contain any predefined hyperfines (`bath['A'].any() == False`) the point dipole approximation is used. Otherwise it can be an instance of `pc.Cube` object, or callable with signature `func(coord, gyro, central_gyro)`, where `coord` is an array of the bath spin coordinate, `gyro` is the gyromagnetic ratio of bath spin, `central_gyro` is the gyromagnetic ratio of the central bath spin.
  - `types` - instance of `SpinDict` or input to create one.
  - `error_range` - maximum allowed distance between positions in `bath` and `external_bath` for two spins to be considered the same.
  - `ext_r_bath` - cutoff distance from the qubit for the `external_bath`. Useful if `external_bath` has very assymmetric shape and user wants to keep the precision level of the hyperfine at different distances consistent.
  - `imap` - instance of the `pc.InteractionMap` class, which contain tensor of bath spin interactions. If not provided, interactions between bath spins are assumed to be the same as one of point dipoles.

Generates `BathArray` object with hyperfine tensors to be used in the calculation.

- `Simulator.generate_clusters` is called if `order` and `r_dipole` are provided. It produces `dict` object, which contains the indexes of the bath spins in the clusters.

We implemented the following procedure to determine the clusters:

Each bath spin  $i$  forms a cluster of one. Bath spins  $i$  and  $j$  form cluster of two if there is an edge between them (distance  $d_{ij} \leq r\_dipole$ ). Bath spins  $i$ ,  $j$ , and  $k$  form a cluster of three if enough edges connect them (e.g., there are two edges  $ij$  and  $jk$ ) and so on. In general, we assume that spins  $\{i..n\}$  form clusters if they form a connected graph. Only clusters up to the size indicated by the `order` parameter (equal to CCE order) are included.



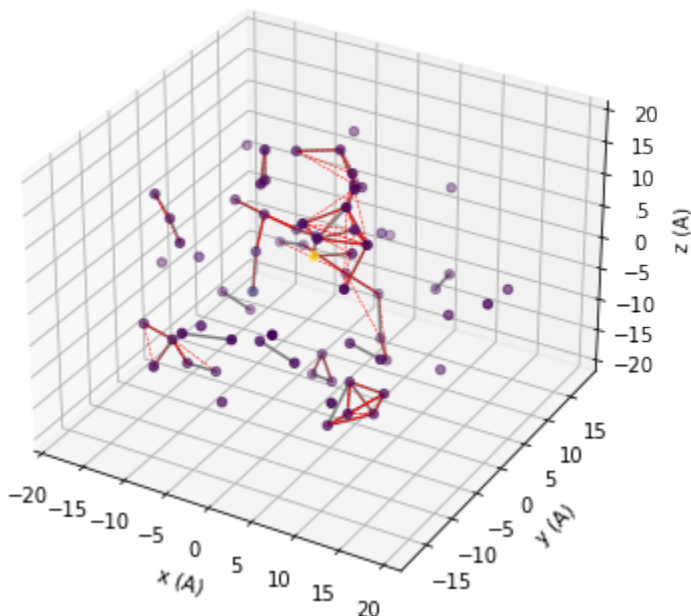
We use matplotlib to visualize the spatial distribution of the spin bath. The grey lines show connected pairs of nuclear spins, red dashed lines show clusters of three. You can try to increase `r_dipole`, `r_bath` parameters, or increase order and visualize.

```
[12]: # add 3D axis
fig = plt.figure(figsize=(6,6))
ax = fig.add_subplot(projection='3d')

# We want to visualize the smaller bath
data = mock.bath

# First plot the positions of the bath
colors = np.abs(data.A[:,2,2])/data.A[:,2,2].max()
ax.scatter3D(data.x, data.y, data.z, c=colors, cmap='viridis');
# Plot all pairs of nuclear spins, which are contained
# in the calc.clusters dictionary under the key 2
for c in mock.clusters[2]:
    ax.plot3D(data.x[c], data.y[c], data.z[c], color='grey')
# Plot all triplets of nuclear spins, which are contained
# in the calc.clusters dictionary under the key 3
for c in mock.clusters[3]:
    ax.plot3D(data.x[c], data.y[c], data.z[c], color='red', ls='--', lw=0.5)

ax.set(xlabel='x (A)', ylabel='y (A)', zlabel='z (A)');
```



Now we setup Simulator object for the actual simulation.

```
[13]: # Parameters of CCE calculations engine

# Order of CCE approximation
order = 2
# Bath cutoff radius
```

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```

r_bath = 40 # in A
# Cluster cutoff radius
r_dipole = 8 # in A
# position of central spin
position = [0, 0, 0]
# Qubit levels (in Sz basis)
alpha = [0, 0, 1]; beta = [0, 1, 0]
# ZFS Parameters of NV center in diamond
D = 2.88 * 1e6 # in kHz
E = 0 # in kHz

```

The code already knows the properties of the most common nuclear spins and of electron spin (accessible under the name 'e'), however the user can provide their own by providing types keyword argument of `Simulator.read_bath` method. The way to initialize `SpinType` objects is the same as in `SpinDict` above.

```

[14]: # The code already knows most existing isotopes.
#       Bath spin types
#       name      spin      gyro      quadrupole (for s>1/2)
spin_types = [('14N', 1, 1.9338, 20.44),
              ('13C', 1 / 2, 6.72828),
              ('29Si', 1 / 2, -5.3188),]

```

### Setting the Simulator object

All of the kwargs can be provided at the moment of creation. If all of the kwargs are provided, several methods of the `Simulator` class are called:

- `Simulator.read_bath`;
- `Simulator.generate_clusters`.

The details are available in the `Simulator` methods description.

```

[15]: # Setting the runner engine
calc = pc.Simulator(spin=1, position=position,
                   alpha=alpha, beta=beta,
                   types=spin_types, D=D, E=E,
                   bath=atoms, r_bath=r_bath,
                   r_dipole=r_dipole, order=order)

```

Taking advantage of subclassing `np.ndarray` we can change *in situ* the quadrupole tensor of the Nitrogen nuclear spin.

```

[16]: nspin = calc.bath
# Set model quadrupole tensor at N atom
quad = np.asarray([[ -2.5, 0, 0],
                  [ 0, -2.5, 0],
                  [ 0, 0, 5.0]]) * 1e3 * 2 * np.pi

nspin['Q'][nspin['N'] == '14N'] = quad

```

Note, that we need to apply the boolean mask **second** because of how structured arrays work.

## Compute coherence function with conventional CCE

The general interface to compute any property with PyCCE is implemented through the `Simulator.compute` method. It takes two keyword arguments to determine which quantity to compute and how:

- `method` can take 'cce' or 'gcce' values, and determines which method to use - conventional or generalized CCE.
- `quantity` can take 'coherence' or 'noise' values, and determines which quantity to compute - coherence function or autocorrelation function of the noise.

Each of the methods can be performed with Monte Carlo bath state sampling (if `nbstates` keyword is non zero) and with interlaced averaging (if `interlaced` keyword is set to `True`).

In the first example we use the conventional CCE method without Monte Carlo bath state sampling. In the conventional CCE method the Hamiltonian is projected on the qubit levels, and the coherence is computed from the overlap of the bath evolution, entangled with two different qubit states.

The conventional CCE requires one argument:

- `timespace` — time points at which the coherence function is computed.

Additionally, one can provide the following arguments now, instead of when initializing `Simulator` object:

- `pulses` — number of pulses in CPMG sequence (0 - FID, 1 - HE etc., default 0) or explicit sequence of pulses as `Sequence` class instance.
- `magnetic_field` — array of the magnetic field, default (0, 0, 0).

```
[17]: # Time points
time_space = np.linspace(0, 2, 201) # in ms
# Number of pulses in CPMG seq (0 = FID, 1 = HE)
N = 1
# Mag. Field (Bx By Bz)
B = np.array([0, 0, 500]) # in G

l_conv = calc.compute(time_space, pulses=N, magnetic_field=B,
                      method='cce', quantity='coherence', as_delay=False)
```

```
[18]: %%timeit
calc.compute(time_space, pulses=N, magnetic_field=B,
             method='cce', quantity='coherence', as_delay=False)

1.09 s ± 1.69 ms per loop (mean ± std. dev. of 7 runs, 1 loop each)
```

## Generalized CCE (gCCE)

In contrast to the conventional CCE method, in generalized CCE approach each cluster includes the central spin explicitly.

`Simulator` can take `pulses` argument as an actual pulse sequence with a list of tuples or `Sequence` class instance. In each tuple the first entry is the axis, the second entry - rotational angle.

For example:

```
[('x', np.pi), ('y', np.pi), ('x', np.pi), ('y', np.pi)]
```

will define XY-4 pulse sequence.

An integer number to define the number of pulses is also accepted as in the case of conventional CCE. If the integer is provided, the code assumes the CPMG sequence.

```
[19]: # Hahn-echo pulse sequence
pulse_sequence = [('x', np.pi)]

# Calculate coherence function
l_generatitze = calc.compute(time_space, magnetic_field=B,
                             pulses=pulse_sequence,
                             method='gcce', quantity='coherence')
```

```
[20]: %%timeit
calc.compute(time_space, magnetic_field=B,
             pulses=pulse_sequence, D=D, E=E,
             method='gcce',
             quantity='coherence')
```

3.31 s ± 2.32 ms per loop (mean ± std. dev. of 7 runs, 1 loop each)

### gCCE with random sampling of bath states

Using this approach, one may carry out generalized CCE calculations for the set of random bath states. This functionality can be turned on by by setting the keyword argument `nbstates` to a number of bath states to sample over. Recommended number of bath states is above 100, but the convergence should be checked for each system. Note, that this computation is roughly `nbstates` times longer than an equivalent generalized CCE calculation, as it computes everything `nbstates` times.

For details see `help(calc.compute)`.

```
[21]: # Number of random bath states to sample over
n_bath_states = 20

# Calculate coherence function
l_gcce = calc.compute(time_space, magnetic_field=B,
                     pulses=pulse_sequence,
                     nbstates=n_bath_states,
                     method='gcce', quantity='coherence', seed=seed)
```

```
[22]: %%timeit
n_bath_states = 5
calc.compute(time_space, magnetic_field=B,
             pulses=pulse_sequence,
             nbstates=n_bath_states,
             method='gcce', quantity='coherence', seed=seed)
```

19.2 s ± 12.9 ms per loop (mean ± std. dev. of 7 runs, 1 loop each)

Take a look at the results of three different methods, and check that they produce similar coherence decay. Note that the results obtained using gCCE with bath states sampling deviates from other ones (generalized and conventional CCE), as the chosen number of states (20) is not enough to converge.

```
[23]: plt.plot(time_space, l_conv.real,
              label='conventional CCE')
plt.plot(time_space, l_generatitze.real,
         label='generalized CCE', ls='--')
plt.plot(time_space, l_gcce.real,
```

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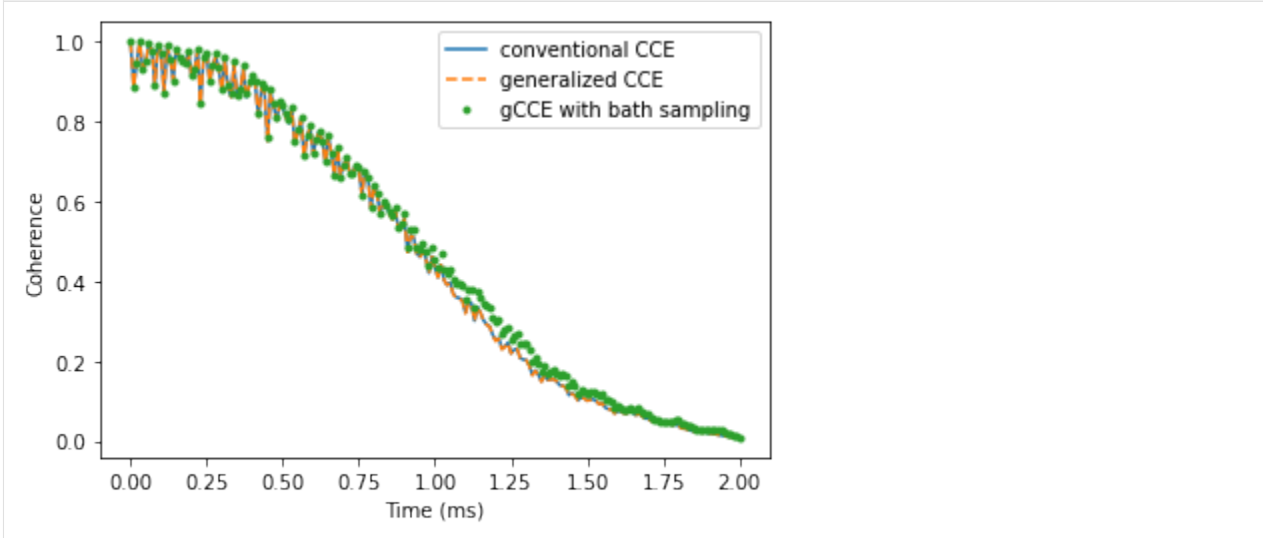
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```

        label='gCCE with bath sampling', ls='', marker='.')
plt.legend()
plt.xlabel('Time (ms)')
plt.ylabel('Coherence')

```

```
[23]: Text(0, 0.5, 'Coherence')
```



### Convergence parameters

Having confirmed that all methods produce the same results, we check the convergence of the conventional CCE with respect to order, `r_bath`, `r_dipole` parameters of the `Simulator` object.

First, define all of the parameters.

```
[24]: parameters = dict(
    order=2, # CCE order
    r_bath=40, # Size of the bath in A
    r_dipole=8, # Cutoff of pairwise clusters in A
    position=[0, 0, 0], # Position of central Spin
    alpha=[0, 0, 1],
    beta=[0, 1, 0],
    pulses = 1, # N pulses in CPMG sequence
    magnetic_field=[0,0,500]
) # Qubit levels)

time_space = np.linspace(0, 2, 201) # Time points in ms

```

We can define a little helper function to streamline the process. Note that resetting the parameters automatically recomputes the properties of the bath.

```
[25]: def runner(variable, values):
    inval = parameters[variable]
    calc = pc.Simulator(spin=1, bath=atoms, **parameters)
    ls = []

```

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```

for v in values:
    setattr(calc, variable, v)
    l = calc.compute(time_space, method='cce',
                    quantity='coherence')

    ls.append(l.real)

parameters[variable] = invalue
ls = pd.DataFrame(ls, columns=time_space, index=values).T
return ls

```

Now we can compute the coherence function at different values of the parameters:

```

[26]: orders = runner('order', [1, 2, 3, 4])
      rbs = runner('r_bath', [20, 30, 40, 50, 60])
      rds = runner('r_dipole', [4, 6, 8, 10])

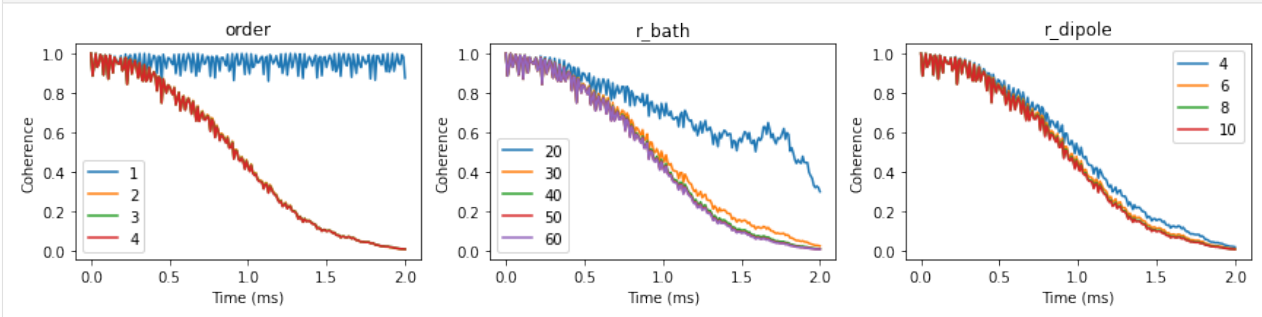
```

We can visualize the convergence of the coherence function with respect to different parameters:

```

[27]: fig, axes = plt.subplots(1, 3, figsize=(12, 3))
      orders.plot(ax=axes[0], title='order')
      rbs.plot(ax=axes[1], title='r_bath')
      rds.plot(ax=axes[2], title='r_dipole')
      for ax in axes:
          ax.set(xlabel='Time (ms)', ylabel='Coherence')
      fig.tight_layout()

```



## 3.2 VV in SiC

An example of computing Free Induction Decay (FID) and Hahn-echo (HE) with hyperfine couplings from GIPAW for axial and basal divacancies.

```

[1]: import numpy as np
      import matplotlib.pyplot as plt
      import sys
      import ase
      import pandas as pd
      import warnings
      sys.path.append('/home/onizhuk/midway/codes_development/pyCCE')

```

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```
import pycce as pc

np.set_printoptions(suppress=True, precision=5)
warnings.simplefilter("ignore")

seed = 8805
```

### 3.2.1 Axial kk-VV

First we compute FID and HE for axial divacancy.

#### Build BathCell from the ground

One can set up an BathCell instance by providing the parameters of the unit cell, or cell argument as 3x3 tensor, where each column defines a, b, c unit cell vectors in cartesian coordinates.

In this tutorial we use the first approach.

```
[2]: # Set up unit cell with (a, b, c, alpha, beta, gamma)
sic = pc.BathCell(3.073, 3.073, 10.053, 90, 90, 120, 'deg')
# z axis in cell coordinates
sic.zdir = [0, 0, 1]
```

Next, user has to define positions of atoms in the unit cell. It is done with BathCell.add\_atoms function. It takes an unlimited number of arguments, each argument is a tuple. First element of the tuple is the name of the atom, second - list of xyz coordinates either in cell units (if keyword type='cell', default value) or in Angstrom (if keyword type='angstrom'). Returns BathCell.atoms dictionary, which contains list of coordinates for each type of elements.

```
[3]: # position of atoms
sic.add_atoms(('Si', [0.00000000, 0.00000000, 0.1880]),
              ('Si', [0.00000000, 0.00000000, 0.6880]),
              ('Si', [0.33333333, 0.66666667, 0.4380]),
              ('Si', [0.66666667, 0.33333333, 0.9380]),
              ('C', [0.00000000, 0.00000000, 0.0000]),
              ('C', [0.00000000, 0.00000000, 0.5000]),
              ('C', [0.33333333, 0.66666667, 0.2500]),
              ('C', [0.66666667, 0.33333333, 0.7500]));
```

Two types of isotopes present in SiC:  $^{29}\text{Si}$  and  $^{13}\text{C}$ . We add this information with the BathCell.add\_isotopes function. The code knows most of the concentrations, so this step is actually unnecessary. If no isotopes is provided, the natural concentration of common magnetic isotopes is assumed.

```
[4]: # isotopes
sic.add_isotopes(('29Si', 0.047), ('13C', 0.011))

# defect position in cell units
vsi_cell = [0, 0, 0.1880]
vc_cell = [0, 0, 0]

# Generate bath spin positions
```

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```
atoms = sic.gen_supercell(200, remove=[('Si', vsi_cell),
                                       ('C', vc_cell)],
                        seed=seed)
```

## Read Quantum Espresso output

PyCCE provides a helper function `read_qe` in `pycce.io` module to read hyperfine couplings from quantum espresso output. `read_qe` takes from 1 to 3 positional arguments:

- `pwfile` name of the pw input/output file;
- `hyperfine` name of the gipaw output file containing hyperfine couplings;
- `efg` name of the gipaw output file containing electric field tensor calculations.

During its call, `read_qe` will read the cell matrix in pw file and apply it to the coordinates is necessary. However, usually we still need to rotate and translate the Quantum Espresso supercell to align it with our `BathArray`. To do so we can provide additional keywords arguments `center` and `rotation_matrix`. `center` is the position of (0, 0, 0) point in coordinates of pw file, and `rotation_matrix` is rotation matrix which aligns z-direction of the GIPAW output. This matrix, acting on the (0, 0, 1) in Cartesian coordinates of GIPAW output should produce (a, b, c) vector, alligned with zdirection of the BathCell. Keyword argument `rm_style` shows whether `rotation_matrix` contains coordinates of new basis set as rows ('row', common in physics) or columns ('col', common in math).

```
[5]: # Prepare rotation matrix to alling with z axis of generated atoms
# This matrix, acting on the [0, 0, 1] in Cartesian coordinates of GIPAW output
# Should produce [a, b, c] vector, alligned with zdirection of the BathCell
M = np.array([[0, 0, -1],
              [0, -1, 0],
              [-1, 0, 0]])

# Position of (0,0,0) point in cell coordinates
center = [0.6, 0.5, 0.5]
# Read GIPAW results
exatoms = pc.read_qe('axial/pw.in',
                    hyperfine='axial/gipaw.out',
                    center=center, rotation_matrix=M,
                    rm_style='col',
                    isotopes={'C':'13C', 'Si':'29Si'})
```

`pc.read_qe` produces instance of `BathArray`, with names of bath spins as the most common isotopes of the following elements (if keyword `isotopes` set to `None`) or from the mapping provided by the `isotopes` argument.

## Set up CCE Simulator

In this example we set up a bare Simulator and add properties of the spin bath later.

```
[6]: # Setting up CCE calculations
pos = sic.to_cartesian(vsi_cell)
CCE_order = 2
r_bath = 40
r_dipole = 8
B = np.array([0, 0, 500])
```

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```
calc = pc.Simulator(1, pos, alpha=[0, 0, 1], beta=[0, 1, 0], magnetic_field=B)
```

Function `Simulator.read_bath` can be called explicitly to initialize spin bath. Additional keyword argument `external_bath` takes instance of `BathArray` with hyperfine couplings read from Quantum Espresso. The program then finds the spins with the same name at the same positions (within the range defined by `error_range` keyword argument) in the total bath and sets their hyperfine couplings.

Finally, we call `Simulator.generate_clusters` to find the bath spin clusters in the provided bath.

```
[7]: calc.read_bath(atoms, r_bath, external_bath=exatoms);
      calc.generate_clusters(CCE_order, r_dipole=r_dipole);
```

### FID with DFT hyperfine couplings

We provide `pulses` argument directly to the compute function instead of during initialization of the `Simulator` object.

```
[8]: time_space = np.linspace(0, 0.01, 501)
      N = 0

      ldft = calc.compute(time_space, pulses=N, as_delay=False)
```

```
[9]: len(calc.pulses)
```

```
[9]: 0
```

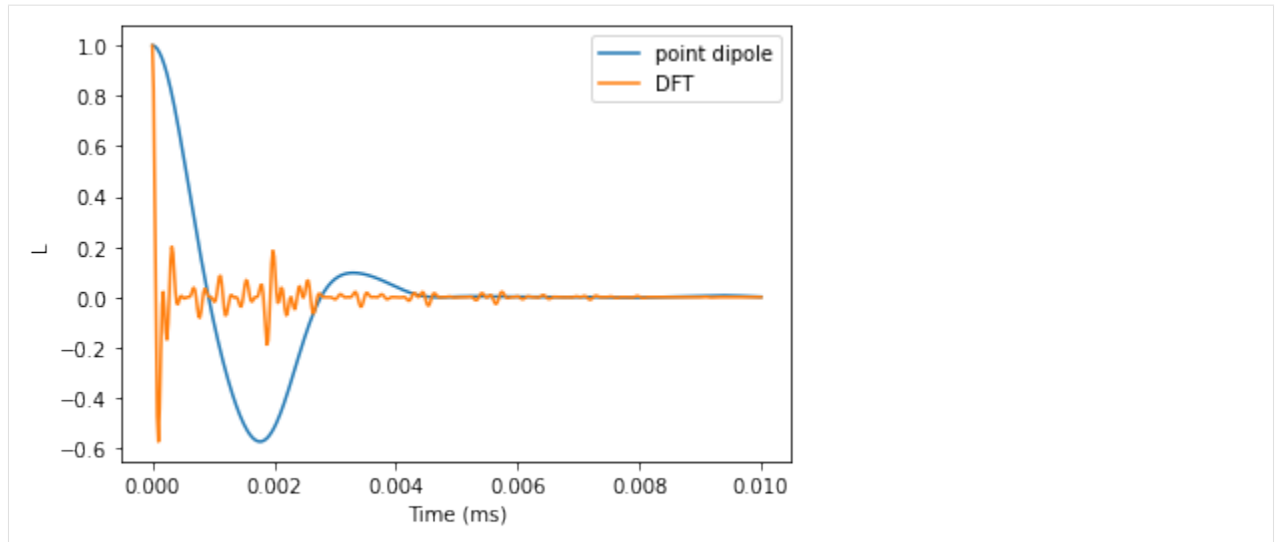
### FID with hyperfine couplings from point dipole approximation

```
[10]: pdcalc = pc.Simulator(1, pos, alpha=[0, 0, 1], beta=[0, 1, 0], magnetic_field=B,
                           bath=atoms, r_bath=r_bath, order=CCE_order, r_dipole=r_dipole)
      lpd = pdcalc.compute(time_space, pulses=N, as_delay=False)
```

Plot the results and verify that the predictions are significantly different.

```
[11]: plt.plot(time_space, lpd.real, label='point dipole')
      plt.plot(time_space, ldft.real, label='DFT')
      plt.legend()
      plt.xlabel('Time (ms)')
      plt.ylabel('L')
```

```
[11]: Text(0, 0.5, 'L')
```



### Hahn-echo comparison

Now we compare the predictions for Hahn-echo signal with different hyperfine couplings.

```
[12]: he_time = np.linspace(0, 2.5, 501)
      B = np.array([0, 0, 500])
      N = 1

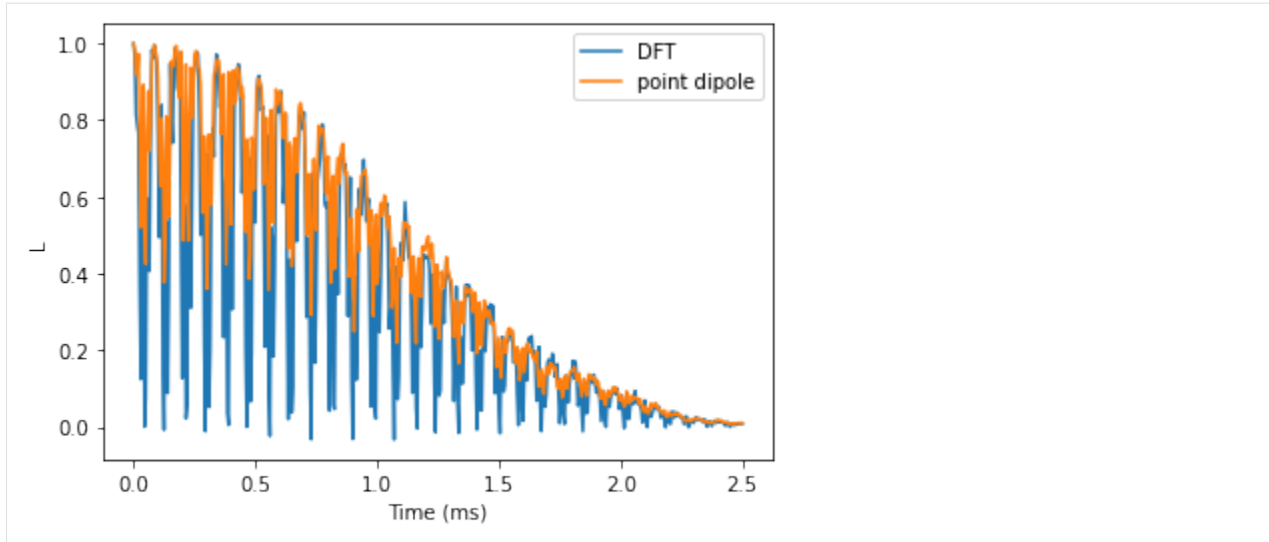
      he_ldft = calc.compute(he_time, magnetic_field=B, pulses=N, as_delay=False)
      he_lpd = pdcalc.compute(he_time, magnetic_field=B, pulses=N, as_delay=False)
```

Plot the results and compare. We observe that electron spin echo modulations differ significantly, while the observed decay is about the same.

```
[13]: plt.plot(he_time, he_ldft.real, label='DFT')
      plt.plot(he_time, he_lpd.real, label='point dipole')

      plt.legend()
      plt.xlabel('Time (ms)')
      plt.ylabel('L')
```

```
[13]: Text(0, 0.5, 'L')
```



### 3.2.2 Basal kh-VV in SiC

The basal divacancy's Hamiltonian includes both D and E terms, which allows for mixing between +1 and -1 spin levels at zero field.

Thus, either the generalized CCE should be used, or additional perturbational Hamiltonian terms are to be added. Here we consider the generalized CCE framework.

First, prepare rotation matrix for DFT results. The same supercell was used to compute hyperfine couplings, however z-axis of the electron spin qubit is aligned with Si-C bond, therefore we will need to rotate the DFT supercell accordingly.

```
[14]: # Coordinates of vacancies in cell coordinates (note that Vsi is not located in the
      ↪ first unitcell)
      vsi_cell = -np.array([1 / 3, 2 / 3, 0.0620])
      vc_cell = np.array([0, 0, 0])

      sic.zdir = [0, 0, 1]

      # Rotation matrix for DFT supercell
      R = pc.rotmatrix([0, 0, 1], sic.to_cartesian(vsi_cell - vc_cell))
```

Total spin bath can be initialized by simply setting z direction of the BathCell object.

```
[15]: sic.zdir = vsi_cell - vc_cell

      # Generate bath spin positions
      sic.add_isotopes(('29Si', 0.047), ('13C', 0.011))
      atoms = sic.gen_supercell(200, remove=[('Si', vsi_cell),
                                             ('C', vc_cell)],
                               seed=seed)
```

Read DFT results with `read_qe` function. To rotate in the correct frame we need to apply both changes of basis consequently

```
[16]: M = np.array([[0, 0, -1],
                 [0, -1, 0],
                 [-1, 0, 0]])

# Position of (0,0,0) point in cell coordinates
center = np.array([0.59401, 0.50000, 0.50000])

# Read GIPAW results
exatoms = pc.read_qe('basal/pw.in',
                    hyperfine='basal/gipaw.out',
                    center=center, rotation_matrix=(M.T @ R),
                    rm_style='col',
                    isotopes={'C':'13C', 'Si':'29Si'})
```

To check whether our rotations produced correct results we can find the indexes of the `BathArray` and DFT output with `pc.same_bath_indexes` function. It returns a tuple, containing the indexes of elements in the two `BathArray` instances with the same position and name. First element of the tuple - indexes of first argument, second - of the second. For that we generate supercell with `BathCell` class, containing 100% isotopes, and count the number of found indexes - iut should be equal to the size of DFT supercell.

```
[17]: # isotopes
sic.add_isotopes(('29Si', 1), ('13C', 1))
allcell = sic.gen_supercell(50, remove=[('Si', vsi_cell),
                                       ('C', vc_cell)],
                           seed=seed)

indexes, ext_indexes = pc.same_bath_indexes(allcell, exatoms, 0.2, True)
print(f"There are {indexes.size} same elements."
      f" Size of the DFT supercell is {exatoms.size}")
```

There are 1438 same elements. Size of the DFT supercell is 1438

## Setting up calculations

Now we can safely setup calculations of coherence function with DFT couplings. We will compare results with or without bath state sampling.

```
[18]: D = 1.334 * 1e6
E = 0.0184 * 1e6
magnetic_field = 0

calc = pc.Simulator(1, pos, bath=atoms, external_bath=exatoms, D=D, E=E,
                   magnetic_field=magnetic_field,
                   r_bath=r_bath, order=CCE_order, r_dipole=r_dipole)
```

The code automatically picks up the two lowest eigenstates of the central spin hamiltonian as qubit states.

```
[19]: print(calc)

Simulator for spin-1.
alpha: 0
beta: 1
gyromagnetic ratio: -17608.59705 kHz * rad / G
```

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```
zero field splitting:
array([[ -426266.66667,    0.    ,    0.    ],
       [    0.    , -463066.66667,    0.    ],
       [    0.    ,    0.    , 889333.33333]])
```

```
magnetic field:
array([0., 0., 0.])
```

```
Parameters of cluster expansion:
r_bath: 40
r_dipole: 8
order: 2
```

Bath consists of 761 spins.

```
Clusters include:
761 clusters of order 1.
1870 clusters of order 2.
```

We can use the `Simulator.eigenstates` function to generate qubit states.

```
[20]: calc.eigenstates(D=D, E=E, alpha=-1, beta=0)
print(f'0 state: {calc.alpha}; 1 state: {calc.beta}')

0 state: [-0.70711+0.j  0.    +0.j -0.70711+0.j]; 1 state: [ 0.+0.j -1.+0.j  0.+0.j]
```

## FID Decay

Now, use the generalized CCE to compute FID decay of the coherence function at different CCE orders.

```
[21]: N = 0 # Number of pulses
time_space = np.linspace(0, 1, 101) # Time points at which to compute

orders = [1, 2, 3]
lgen = []

r_bath = 30
r_dipole = 8

calc = pc.Simulator(1, pos, bath=atoms, external_bath=exatoms,
                   D=D, E=E, pulses=N, alpha=-1, beta=0,
                   r_bath=r_bath, r_dipole=r_dipole)
```

```
[22]: for o in orders:
    calc.generate_clusters(o)
    l = calc.compute(time_space, method='gcce',
                    quantity='coherence', as_delay=False)

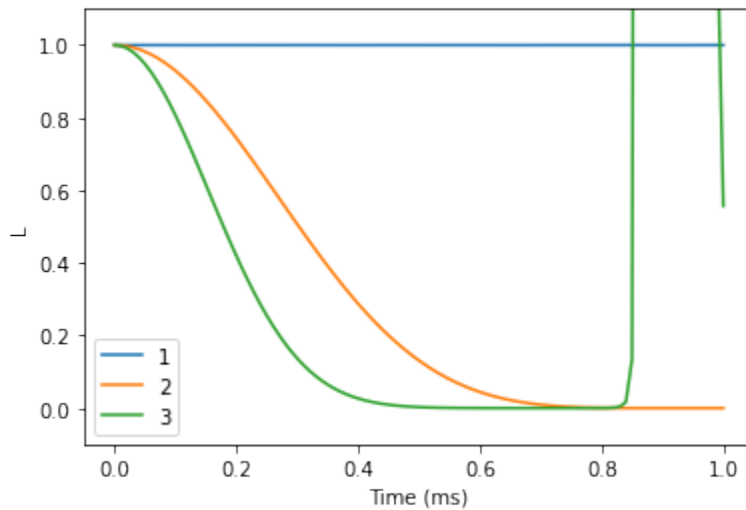
    lgen.append(np.abs(l))

lgen = pd.DataFrame(lgen, columns=time_space, index=orders).T
```

We see that the results do not converge, but rather start to diverge. Bath sampling (setting nbstates to some value) will help with that.

```
[23]: lgen.plot()
plt.xlabel('Time (ms)')
plt.ylabel('L')
plt.ylim(-0.1,1.1)
```

```
[23]: (-0.1, 1.1)
```



Note, that this approach is nbstates times more expensive than the gCCE, therefore the following calculation will take a couple of minutes.

```
[24]: orders = [1, 2, 3]
lgcce = []

r_bath = 30
r_dipole = 6

for o in orders:
    calc.generate_clusters(o)

    l = calc.compute(time_space, nbstates=30, seed=seed,
                    method='gcce',
                    quantity='coherence', as_delay=False)

    lgcce.append(np.abs(l))

lgcce = pd.DataFrame(lgcce, columns=time_space, index=orders).T
```

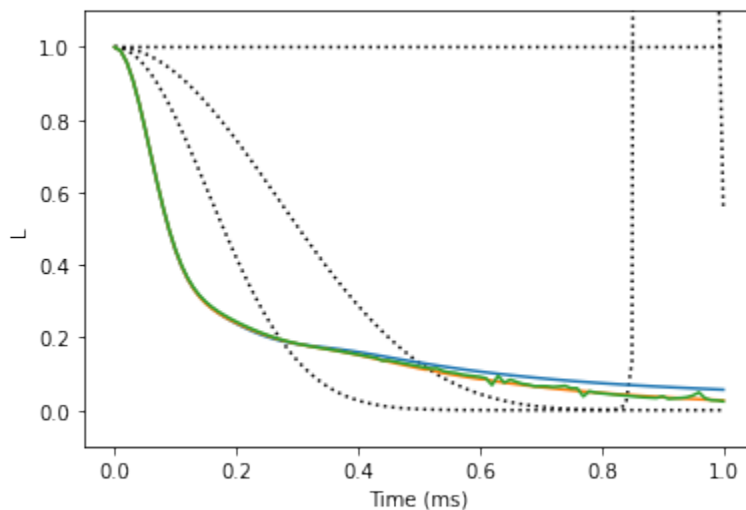
### Compare the two results

The gCCE results are converged at 1st order. Note that we used only a small number of bath states (30), thus the calculations are not converged with respect to the number of bath states. Calculations with higher number of bath states (~100) will produce correct results.

```
[25]: plt.plot(lgen, color='black', ls=':')
plt.plot(lgcce)

plt.xlabel('Time (ms)')
plt.ylabel('L')
plt.ylim(-0.1,1.1)
```

```
[25]: (-0.1, 1.1)
```



### Hahn-echo decay

Using the similar procedure to the one used for FID, we can compute the Hahn-echo decay.

```
[26]: r_bath = 40
r_dipole = 8
order = 2
N = 1 # Number of pulses

calc = pc.Simulator(1, pos, bath=atoms, external_bath=exatoms,
                    pulses=N, D=D, E=E, alpha=-1, beta=0,
                    r_bath=r_bath, order=order, r_dipole=r_dipole)
```

```
[27]: ts = np.linspace(0, 4, 101) # time points (in ms)
```

```
[28]: helgen = calc.compute(ts,
                             method='gcce', quantity='coherence')
```

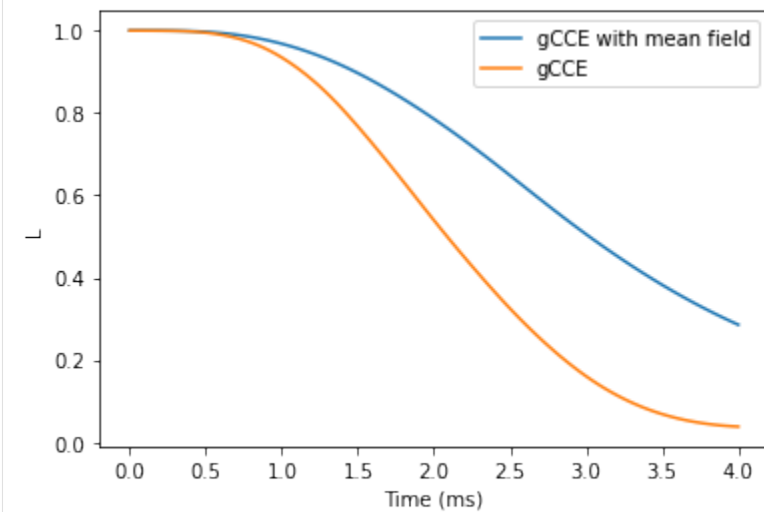
Note the number of nbstates leads to significantly increased time of the calculation. The interface to mpi implementation is provided with keyword `parallel_states`. However it requires `mpi4py` installed and a run on several cores.

```
[29]: helgcce = calc.compute(time_space, nbstates=30, seed=seed,
                             method='gcce', quantity='coherence')
```

```
[30]: plt.plot(ts, helgcce, label='gCCE with mean field')
plt.plot(ts, helgen, label='gCCE')

plt.legend()
plt.xlabel('Time (ms)')
plt.ylabel('L')
```

```
[30]: Text(0, 0.5, 'L')
```



### 3.3 Shallow donor in Si

Example of more complicated simulations, in which we compare the coherence predicted with point-dipole hyperfine couplings and one obtained using the hyperfines from model wavefunction of the shallow donor in Si (P:Si).

```
[1]: import numpy as np
import matplotlib.pyplot as plt
import sys
import ase

import pycce as pc

seed = 8800
np.set_printoptions(suppress=True, precision=5)
```

First, as always, generate spin bath with BathCell instance. To get parameters we use ase interface. It allows to conveniently read structure files of any type.

```
[2]: # Generate unitcell from ase
from ase import io
s = io.read('si.cif')
s = pc.bath.BathCell.from_ase(s)
```

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```

# Add types of isotopes
s.add_isotopes(('29Si', 0.047))
# set z direction of the defect
s.zdir = [1, 1, 1]
# Generate supercell
atoms = s.gen_supercell(200, remove=[('Si', [0., 0., 0.])], seed=seed)

```

### 3.3.1 Calculations with point dipole hyperfine couplings

Here we compute Hahn-echo decay with point dipole hyperfine couplings. All of the parameters are converged, however it never hurts to check!

```

[3]: # Parameters of CCE calculations engine

# Order of CCE approximation
CCE_order = 2
# Bath cutoff radius
r_bath = 80 # in A
# Cluster cutoff radius
r_dipole = 10 # in A

# position of central spin
position = [0, 0, 0]
# Qubit levels (in Sz basis)
alpha = [0, 1]; beta = [1, 0]
# Mag. Field (Bx By Bz)
B = np.array([0, 0, 1000]) # in G
# Number of pulses in CPMG seq (0 = FID, 1 = HE etc)
pulses = 1

# Setting the runner engine
calc = pc.Simulator(spin=0.5, position=position, alpha=alpha, beta=beta,
                    bath=atoms, r_bath=r_bath, magnetic_field=B, pulses=pulses,
                    r_dipole=r_dipole, order=CCE_order)

```

```

[4]: # Time points
time_space = np.linspace(0, 2, 201) # in ms

```

For comparison, we compute both with generalized CCE and usual CCE coherence. Note a relatively large bath ( $r_{\text{bath}} = 80$ ), so the calculations will take some time.

```

[5]: l_cce = calc.compute(time_space, method='CCE')
l_gen = calc.compute(time_space, method='gCCE')

```

### 3.3.2 Hyperfine couplings of the shallow donor

We compute the hyperfine couplings of the shallow donor, following the formulae by Rogerio de Sousa and S. Das Sarma (*Phys Rev B* 68, 115322 (2003)).

```
[6]: # PHYSICAL REVIEW B 68, 115322 (2003)
n = 0.81
a = 25.09

def factor(x, y, z, n=0.81, a=25.09, b=14.43):
    top = np.exp(-np.sqrt(x**2/(n*b)**2 + (y**2 + z**2)/(n*a)**2))
    bottom = np.sqrt(np.pi * (n * a)**2 * (n * b) )

    return top / bottom

def contact_si(r, gamma_n, gamma_e=pc.ELECTRON_GYRO, a_lattice=5.43, nu=186, n=0.81,
↪a=25.09, b=14.43):
    k0 = 0.85 * 2 * np.pi / a_lattice
    pre = 8 / 9 * gamma_n * gamma_e * pc.HBAR * nu
    xpart = factor(r[0], r[1], r[2], n=n, a=a, b=b) * np.cos(k0 * r[0])
    ypart = factor(r[1], r[2], r[0], n=n, a=a, b=b) * np.cos(k0 * r[1])
    zpart = factor(r[2], r[0], r[1], n=n, a=a, b=b) * np.cos(k0 * r[2])
    return pre * (xpart + ypart + zpart) ** 2
```

We make a copy of the BathArray object, and set up their hyperfines according to the reference above.

```
[7]: newatoms = atoms.copy()

# Generate hyperfine from point dipole
newatoms.from_point_dipole(position)

# Following PRB paper
newatoms['A'][newatoms.dist() < n*a] = 0
newatoms['A'] += np.eye(3)[np.newaxis, :, :] * contact_si(newatoms['xyz'].T, newatoms.
↪types['29Si'].gyro)[: , np.newaxis, np.newaxis]
```

Now we set up a Simulator object. Because hyperfines in newatoms are nonzero, they are **not** approximated as the ones of point dipole.

```
[8]: calc = pc.Simulator(spin=0.5, position=position, alpha=alpha, beta=beta,
                        bath=newatoms, r_bath=r_bath, magnetic_field=B, pulses=pulses,
                        r_dipole=r_dipole, order=CCE_order)
```

```
[9]: shallow_l_cce = calc.compute(time_space, method='CCE')
shallow_l_gen = calc.compute(time_space, method='gCCE')
```

### 3.3.3 Compare the results

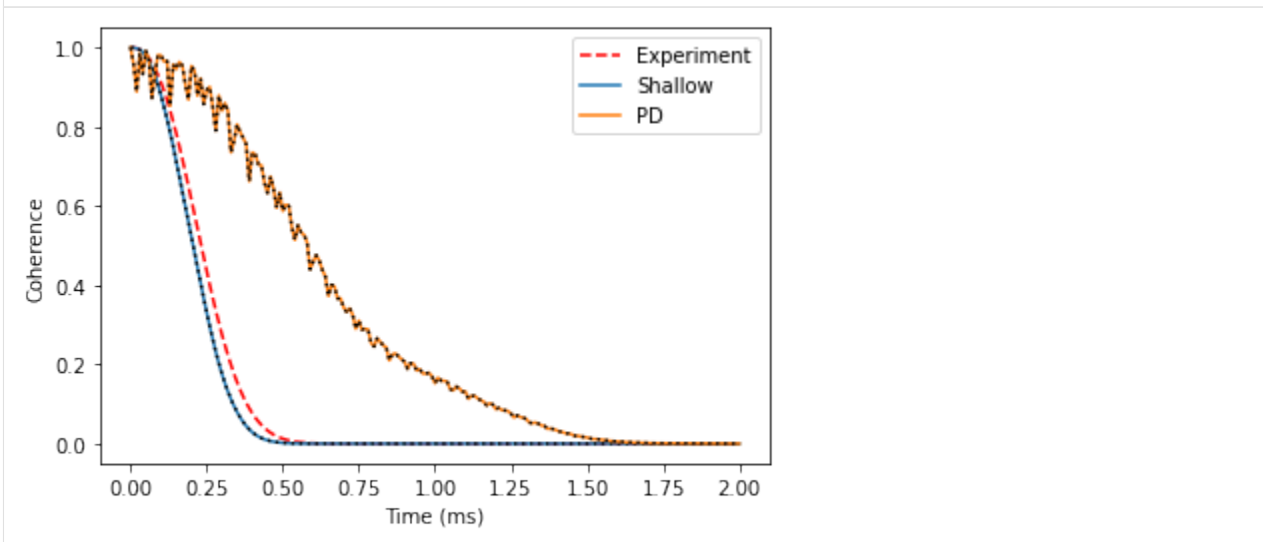
We find that the point dipole gives a poor agreement with the experimental data. Model wavefunction, on the contrary, produces great agreement with the experimental coherence time from work of Eisuke Abe et al. (*Phys Rev B* 82, 121201(R) (2010)).

```
[10]: t2exp = 0.27 # Experimental T2 from PhysRevB.82.121201
decay = lambda t: np.exp(-(t/t2exp)**2.4)
plt.plot(time_space, decay(time_space), color='red', label='Experiment', ls='--')

plt.plot(time_space, shallow_l_cce.real, label='Shallow')
plt.plot(time_space, shallow_l_gen.real, ls=':', c='black')

plt.plot(time_space, l_cce.real, label='PD')
plt.plot(time_space, l_gen.real, ls=':', c='black')
plt.legend();
plt.xlabel('Time (ms)')
plt.ylabel('Coherence')
```

```
[10]: Text(0, 0.5, 'Coherence')
```



Interesting to note - the decay depends significantly on the orientation of the magnetic field. You can check it yourself!

## 3.4 Correlation function

In this tutorial we will compute the coherence function of the NV Center in diamond and then reproduce it from the correlation function of the noise.

The correlation function  $C(t)$  of the effective magnetic field (noise) along the  $z$ -axis can be defined as follows:

$$C(t) = \langle \beta_z(t) \beta_z(0) \rangle$$

With  $\beta_z$  given as:

$$\beta_z(t) = U^\dagger(t) \left( \sum_{\{I\}} A_{zz} I_z \right) U(t)$$

Where  $U(t)$  is time propagator.

Within the CCE formalism, the correlation function is computed as:

$$C(t) = \sum_{\{i\}} \tilde{C}_{\{i\}}(t) + \sum_{\{ij\}} \tilde{C}_{\{ij\}}(t) + \dots$$

With contributions computed as:

$$\tilde{C}_{\nu}(t) = C_{\nu}(t) - \sum_{\nu' \subset \nu} \tilde{C}_{\nu'}(t)$$

```
[1]: import numpy as np
import matplotlib.pyplot as plt
import pandas as pd

import sys

import pycce as pc
import ase

seed = 42055
np.set_printoptions(suppress=True, precision=5)
```

### 3.4.1 Generate nuclear spin bath

Building a BathArray of nuclear spins from the ase.Atoms object.

```
[2]: from ase.build import bulk

# Generate unitcell from ase
diamond = bulk('C', 'diamond', cubic=True)
diamond = pc.bath.BathCell.from_ase(diamond)
# Add types of isotopes
diamond.add_isotopes(('13C', 0.011))
# set z direction of the defect
diamond.zdir = [1, 1, 1]
# Add the defect. remove and add atoms at the positions (in cell coordinates)
atoms = diamond.gen_supercell(200, remove=[('C', [0., 0, 0]),
                                           ('C', [0.5, 0.5, 0.5])],
                              add=('14N', [0.5, 0.5, 0.5]),
                              seed=seed)
```

Next, we define all of the parameters of the simulation. We are interested in the very specific regime, when all nearby nuclear spins are removed. To achieve this goal we define an `inner = 20` parameter, and remove all nuclear spins within this radius.

```
[3]: position = np.array([0, 0, 0])
inner = 20
smallatoms = atoms[atoms.dist(position) >= inner]

parameters = dict(
    order=2, # CCE order
    r_bath=60, # Size of the bath in A
    r_dipole=6, # Cutoff of pairwise clusters in A
    position=position, # Position of central Spin
    alpha=[0, 0, 1], # 0 qubit state
    beta=[0, 1, 0], # 1 qubit state
    magnetic_field = 500, # magnetic field along z-axis
```

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```

    pulses=1 # N pulses in CPMG sequence
) # Qubit levels

ts = np.linspace(0, 2.5, 1001) # Time points in ms

```

### 3.4.2 Coherence calculations

Next, we set up Simulator objects and check convergence with respect to the CCE order.

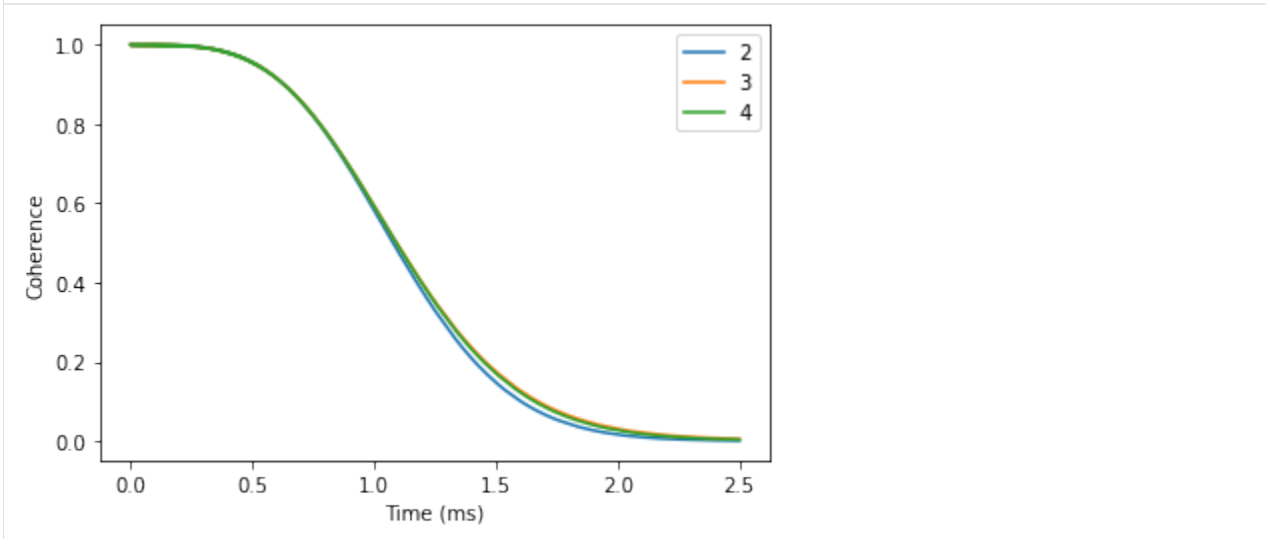
```
[4]: calc = pc.Simulator(spin=1, bath=smallatoms, **parameters)
```

```
[5]: orders = [2, 3, 4]
coh = {}
for o in orders:
    calc.generate_clusters(o)
    coh[o] = calc.compute(ts, method='cce', quantity='coherence')
coh = np.abs(pd.DataFrame(coh, index=ts))
coh.index.name = 'Time (ms)'
```

Visually verify the convergence.

```
[6]: coh.plot()
plt.ylabel('Coherence')
```

```
[6]: Text(0, 0.5, 'Coherence')
```



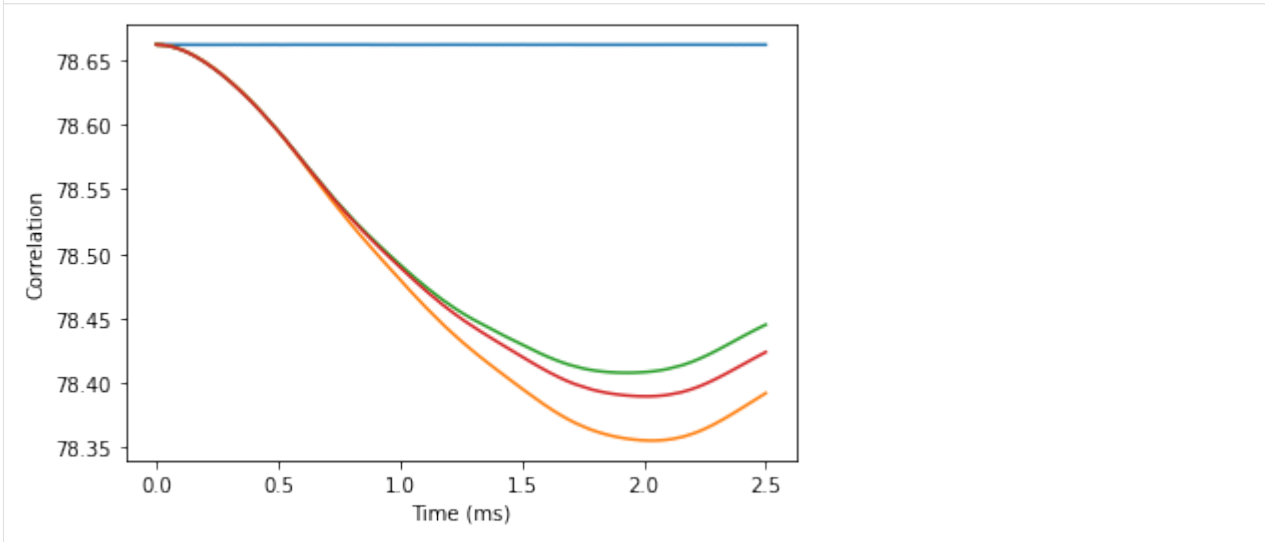
### 3.4.3 Noise calculations

To compute the correlation function of the noise, we call `Simulator.compute` method and specify `quantity = 'noise'`.

First we determine convergence of the correlation function with the CCE order.

```
[7]: for o in [1, 2, 3, 4]:
      calc.generate_clusters(o)
      noise = calc.compute(ts, method='cce', quantity='noise')
      plt.plot(ts, noise.real, label=o)
plt.xlabel('Time (ms)')
plt.ylabel('Correlation')
```

```
[7]: Text(0, 0.5, 'Correlation')
```



The difference between third and fourth order is fairly small, we will use the fourth order for the following calculations.

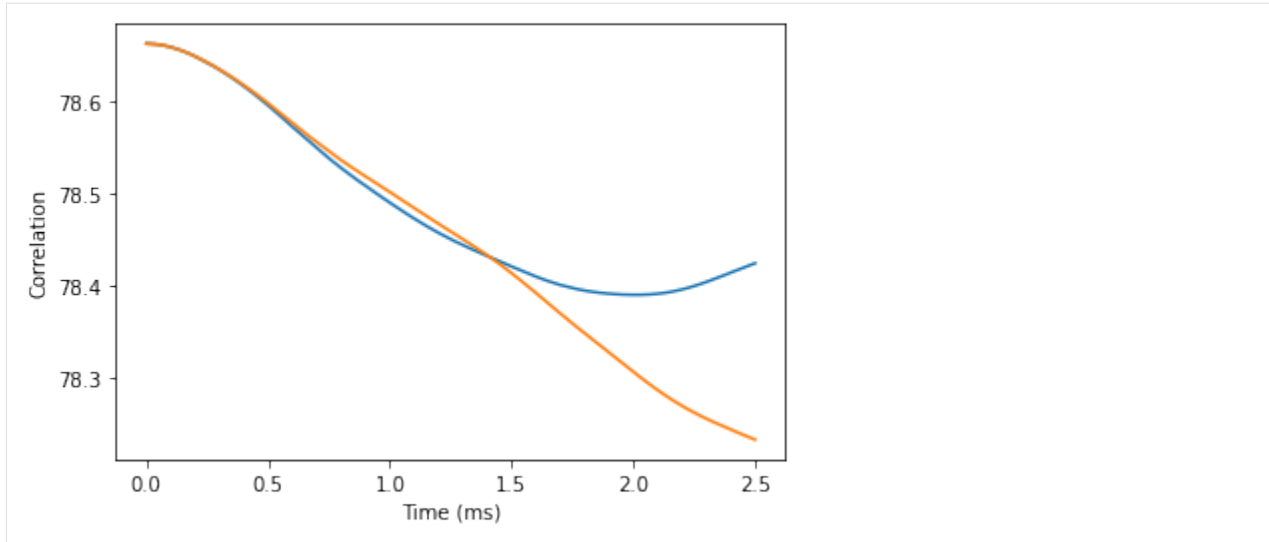
```
[8]: calc.generate_clusters(4)

noise = calc.compute(ts, method='cce', quantity='noise')
genoise = calc.compute(ts, method='gcce', quantity='noise', nbstates=0)
```

Compare the results obtained with CCE and gCCE approaches. Note that they are slightly different. However, as we will see it does not impact the predicted coherence.

```
[9]: plt.plot(ts, noise.real, label='CCE')
plt.plot(ts, genoise.real, label='gCCE')
plt.xlabel('Time (ms)')
plt.ylabel('Correlation')
```

```
[9]: Text(0, 0.5, 'Correlation')
```



Assuming that the noise is Gaussian, we can reproduce the coherence from the average phase squared  $\langle \phi^2 \rangle$ , accumulated by the spin qubit:

$$L(t) = e^{-\langle \phi^2(t) \rangle}$$

The average phase is obtained from the autocorrelation function as:

$$\langle \phi^2(t) \rangle = \int_0^t d\tau C(\tau) F(\tau)$$

Where  $F(\tau)$  is the correlation filter function (see [Phys. Rev. A 86, 012314 \(2012\)](#) for details).

PyCCE code already has implemented calculations of the phase in the `pycce.filter` module:

`pycce.filter.gaussian_phase` takes three positional arguments: - `timespace` - time points at which correlation function was computed; - `corr` - noise autocorrelation function; - `npulses` - number of pulses in CPMG sequence.

Here we compute the phase for the Hahn-echo experiment. Note that the implementation of `gaussian_phase` is not heavily optimized and can take a hot second.

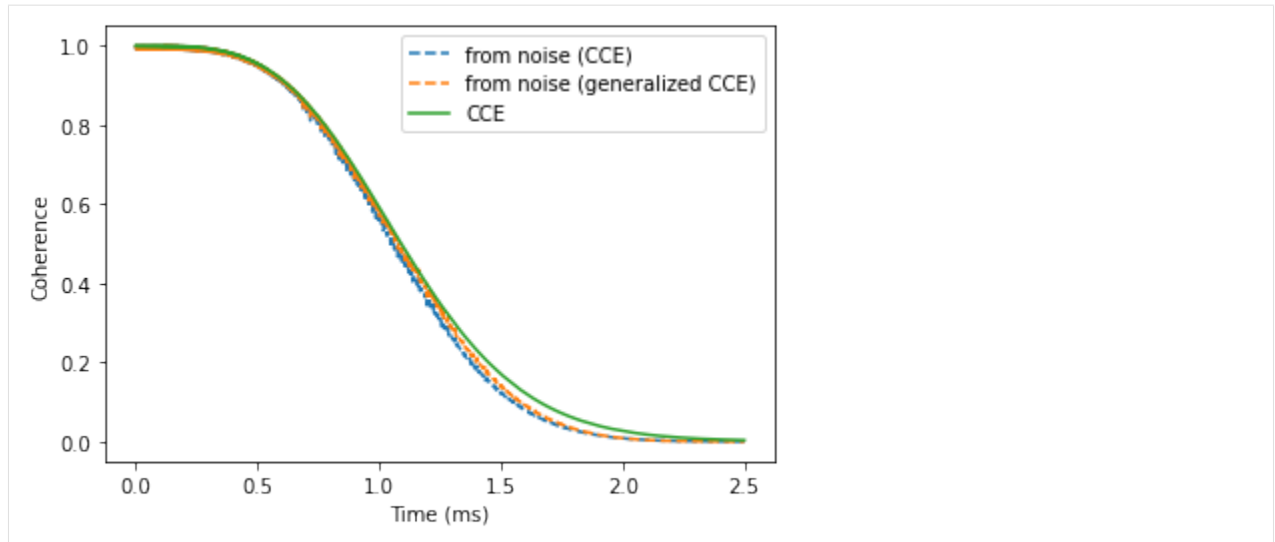
```
[10]: import pycce.filter
```

```
chis = pycce.filter.gaussian_phase(ts, np.abs(noise), 1)
gchis = pycce.filter.gaussian_phase(ts, np.abs(genoise), 1)
```

Now compare results from direct calculations of the coherence function, and the one reconstructed from the noise autocorrelation:

```
[11]: plt.plot(ts, np.exp(-chis), ls='--', label='from noise (CCE)')
plt.plot(ts, np.exp(-gchis), ls='--', marker='', label='from noise (generalized CCE)')
plt.plot(ts, coh[4], label='CCE')
plt.legend()
plt.xlabel('Time (ms)')
plt.ylabel('Coherence')
```

```
[11]: Text(0, 0.5, 'Coherence')
```



The recommended order of the tutorials is from the top to bottom:

- *NV Center in Diamond* example goes through the *Quick Start* example in more details.
- *VV in SiC* tutorial explores the difference between generalized CCE with and without random bath state sampling. Also, in this example we introduce the way to work with DFT output of hyperfine tensors.
- *Shallow donor in Si* example shows the way to include the custom hyperfine couplings for more delocalized defects in semiconductors.
- *Correlation function* example explains the way to use autocorrelation function of the noise to predict the decay of the coherence of the NV center in diamond.



## GENERATING THE SPIN BATH

### 4.1 Random bath

Documentation for the `pycce.random_bath` function, used to generate random bath.

**random\_bath**(*names*, *size*, *number=1000*, *density=None*, *types=None*, *density\_units='cm-3'*, *center=None*, *seed=None*)

Generate random bath containing spins with names provided with argument `name` in the box of size `size`. By default generates coordinates in range  $(-size/2; +size/2)$  but this behavior can be changed by providing `center` keyword.

#### Examples

Generate 2000  $^{13}\text{C}$  nuclear spins in the cubic box with the side of 100 angstrom:

```
>>> atoms = random_bath('13C', 100, number=2000, seed=10)
>>> print(atoms.size)
2000
>>> print(round(atoms.x.min()), round(atoms.x.max()))
-50.0 50.0
```

Generate electron spin bath with density  $10^{17}\text{cm}^{-3}$  in the cuboid box:

```
>>> electrons = random_bath('e', [1e3, 2e3, 3e3], density=1e17,
>>>                          density_units='cm-3', seed=10)
>>> print(electrons.size, round(electrons.x.min()), round(electrons.x.max()))
600 -494.0 500.0
>>> print(electrons.types)
SpinDict(e: (e, 0.5, -17608.59705))
```

#### Parameters

- **names** (*str or array-like with length n*) – Name of the bath spin or array with the names of the bath spins,
- **size** (*float or ndarray with shape (3,)*) – Size of the box. If float is given, assumes 3D cube with the edge = `size`. Otherwise the size specifies the dimensions of the box. Dimensionality is controlled by setting entries of the size array to 0.
- **number** (*int or array-like with length n*) – Number of the bath spins in the box or array with the numbers of the bath spins. Has to have the same length as the name array.

- **density** (*float or array-like with length n*) – Concentration of the bath spin or array with the concentrations. Has to have the same length as the **name** array.
- **types** (*SpinDict*) – Dictionary with SpinTypes or input to create one.
- **density\_units** (*str*) – If number of spins provided as density, defines units. Values are accepted in the format *m*, or *m<sup>x</sup>* or *m-x* where *m* is the length unit, *x* is dimensionality of the bath (e.g. *x* = 1 for 1D, 2 for 2D etc). If only *m* is provided the dimensions are inferred from **size** argument. Accepted length units:
  - *m* meters;
  - *cm* centimeters;
  - *a* angstroms.
- **center** (*ndarray with shape (3,)*) – Coordinates of the (0, 0, 0) point of the final coordinate system in the initial coordinates. Default is **size** / 2 - center is in the middle of the box.

**Returns** Array of the bath spins with random positions.

**Return type** BathArray with shape (np.prod(number))

## 4.2 BathCell

Documentation for the `pycce.BathCell` - class for convenient generation of `BathArray` and the necessary helper functions.

**class BathCell** (*a=None, b=None, c=None, alpha=None, beta=None, gamma=None, angle='rad', cell=None*)  
 Generator of the bath spins positions from the unit cell of the material.

### Parameters

- **a** (*float*) – *a* parameter of the primitive cell.
- **b** (*float*) – *b* parameter of the primitive cell.
- **c** (*float*) – *c* parameter of the primitive cell.
- **alpha** (*float*) –  $\alpha$  angle of the primitive cell.
- **beta** (*float*) –  $\beta$  angle of the primitive cell.
- **gamma** (*float*) –  $\gamma$  angle of the primitive cell.
- **angle** (*str*) – units of the  $\alpha, \beta, \gamma$  angles. Can be either radians ('rad'), or degrees ('deg').
- **cell** (*ndarray with shape (3, 3)*) – Parameters of the cell.

`cell` is 3x3 matrix with columns of coordinates of crystallographic vectors in the cartesian reference frame. See `cell` attribute.

If provided, overrides *a*, *b*, and *c*.

### cell

Parameters of the cell. `cell` is 3x3 matrix with entries:

$$\begin{bmatrix} a_x & b_x & c_x \\ a_y & b_y & c_y \\ a_z & b_z & c_z \end{bmatrix}$$

where *a*, *b*, *c* are crystallographic vectors and *x*, *y*, *z* are their coordinates in the cartesian reference frame.

**Type** ndarray with shape (3, 3)

**atoms**

Dictionary containing coordinates and occupancy of each lattice site:

```
{atom_1: [array([x1, y1, z1]), array([x2, y2, z2])],
 atom_2: [array([x3, y3, z3]), ...]}
```

**Type** dict

**isotopes**

Dictionary containing spin types and their concentration for each lattice site type:

```
{atom_1: {spin_1: concentration, spin_2: concentration},
 atom_2: {spin_3: concentration ...}}
```

where atom\_i are lattice site types, and spin\_i are spin types.

**Type** dict

**property zdir**

z-direction of the reference cartesian coordinate frame in cell coordinates.

**Type** ndarray

**rotate**(*rotation\_matrix*)

Rotate the BathCell using the rotation matrix provided.

**Parameters** **rotation\_matrix** (*ndarray with shape (3,)*) – Rotation matrix R which rotates the old basis of the cartesian reference frame to the new basis.

**set\_zdir**(*direction, type='cell'*)

Set z-direction of the cell.

**Parameters**

- **direction** (*ndarray with shape (3,)*) – Direction of the z axis.
- **type** (*str*) – How coordinates in **direction** are stored. If **type**="cell", assumes crystallographic coordinates. If **type**="angstrom" assumes that z direction is given in the cartesian reference frame.

**add\_atoms**(\**args, type='cell'*)

Add coordinates of the lattice sites to the unit cell.

**Parameters**

- **\*args** (*tuple*) – List of tuples, each containing the type of atom N (*str*), and the xyz coordinates in the format (*float, float, float*): (N, [x, y, z]).
- **type** (*str*) – Type of coordinates. Can take values of ['cell', 'angstrom'].

If **type**="cell", assumes crystallographic coordinates.

If **type**="angstrom" assumes that coordinates are given in the cartesian reference frame.

**Returns** View of `cell.atoms` dictionary, where each key is the type of lattice site, and each value is the list of coordinates in crystallographic frame.

**Return type** dict

## Examples

```
>>> cell = BathCell(10)
>>> cell.add_atoms(('C', [0, 0, 0]), ('C', [5, 5, 5]), type='angstrom')
>>> cell.add_atoms(('Si', [0, 0.5, 0.]), type='cell')
>>> print(cell.atoms)
{'C': [array([0., 0., 0.]), array([0.5, 0.5, 0.5])], 'Si': [array([0. , 0.5, 0.5
↵])]}
```

### add\_isotopes(\*args)

Add spins that can populate each lattice site type.

**Parameters** *\*args* (*tuple or list of tuples*) – Each tuple can have any of the following formats:

- Name of the lattice site *N* (*str*), name of the spin *X* (*str*), concentration *c* (*float*, in decimal): (*N*, *X*, *c*).
- Isotope name *X* and concentration *c*: (*X*, *c*).

In this case, the name of the isotope is given in the format "{*digits*}\_{*atom\_name*}" where *digits* is any set of digits 0-9, *atom\_name* is the name of the corresponding lattice site. Convenient when generating nuclear spin bath.

### Returns

View of `cell.isotopes` dictionary which contains information about lattice site types, spin types, and their concentrations:

```
{atom_1: {spin_1: concentration, spin_2: concentration},
 atom_2: {spin_3: concentration ...}}
```

**Return type** dict

## Examples

```
>>> cell = BathCell(10)
>>> cell.add_atoms(('C', [0, 0, 0]), ('C', [5, 5, 5]), type='angstrom')
>>> cell.add_isotopes(('C', 'X', 0.001), ('13C', 0.0107))
>>> print(cell.isotopes)
{'C': {'X': 0.001, '13C': 0.0107}}
```

### gen\_supercell(size, add=None, remove=None, seed=None)

Generate supercell populated with spins.

---

**Note:** If `isotopes` were not provided, assumes the natural concentration of nuclear spin isotopes for each lattice site type. However, if any isotope concentration is provided, then uses only user-defined ones.

---

### Parameters

- **size** (*float*) – Approximate linear size of the supercell. The generated supercell will have minimal distance between opposite sides larger than this parameter.

- **add** (*tuple or list of tuples*) – Tuple or list of tuples containing `common_isotopes` to add as a defect. Each tuple contains name of the new isotope and its coordinates in the cell basis: (`isotope_name`, `x_cell`, `y_cell`, `z_cell`).
- **remove** (*tuple or list of tuples*) – Tuple or list of tuples containing bath to remove in the defect. Each tuple contains name of the atom to remove and its coordinates in the cell basis: (`atom_name`, `x_cell`, `y_cell`, `z_cell`).
- **seed** (*int*) – Seed for random number generator.

---

**Note:** While `add` takes the `spin` name as an argument, `remove` takes the lattice site name.

---

**Returns** Array of the spins in the given supercell.

**Return type** *BathArray*

**to\_cartesian**(*coord*)

Transform coordinates from crystallographic basis to the cartesian reference frame.

**Parameters** `coord` (*ndarray with shape (3,) or (n, 3)*) – Coordinates in crystallographic basis or array of coordinates.

**Returns** Cartesian coordinates in angstrom.

**Return type** ndarray with shape (3,) or (n, 3)

**to\_cell**(*coord*)

Transform coordinates from the cartesian coordinates of the reference frame to the cell coordinates.

**Parameters** `coord` (*ndarray with shape (3,) or (n, 3)*) – Cartesian coordinates in angstrom or array of coordinates.

**Returns** Coordinates in the cell basis.

**Return type** ndarray with shape (3,) or (n, 3)

**classmethod from\_ase**(*atoms\_object*)

Generate `BathCell` instance from `ase.Atoms` object of Atomic Simulations Environment (ASE) package.

**Parameters** `atoms_object` (*Atoms*) – Atoms object, used to generate new `BathCell` instance.

**Returns** New instance of the `BathCell` with atoms read from `ase.Atoms`.

**Return type** *BathCell*

**defect**(*cell, atoms, add=None, remove=None*)

Generate a defect in the given supercell.

The defect will be located in the unit cell, located roughly in the middle of the supercell, generated by `BathCell`, such that (0, 0, 0) of cartesian reference frame is located at (0, 0, 0) position of this unit cell.

**Parameters**

- **cell** (*ndarray with shape (3, 3)*) – parameters of the unit cell.
- **atoms** (*BathArray*) – Array of spins in the supercell.
- **add** (*tuple or list of tuples*) – Add spin type(s) to the supercell at specified positions to create point defect. Each tuple contains name of the new isotope and its coordinates in the cell basis: (`isotope_name`, `x_cell`, `y_cell`, `z_cell`).

- **remove** (*tuple or list of tuples*) – Remove lattice site from the supercell at specified position to create point defect. Each tuple contains name of the atom to remove and its coordinates in the cell basis: (*atom\_name*, *x\_cell*, *y\_cell*, *z\_cell*).

**Returns** Array of spins with the defect added.

**Return type** *BathArray*

## 4.3 BathArray

Documentation for the `pycce.BathArray` - central class, containing properties of the bath spins.

**class BathArray** (*shape=None, array=None, names=None, hyperfines=None, quadrupoles=None, types=None, imap=None, ca=None, sn=None, hf=None, q=None, efg=None*)

Subclass of ndarray containing information about the bath spins.

The subclass has fixed structured datatype:

```
_dtype_bath = np.dtype([('N', np.unicode_, 16),
                        ('xyz', np.float64, (3,)),
                        ('A', np.float64, (3, 3)),
                        ('Q', np.float64, (3, 3))])
```

Accessing different fields results in the ndarray view.

Each of the fields can be accessed as the attribute of the `BathArray` instance and modified accordingly. In addition to the name fields, the information of the bath spin types is stored in the `types` attribute. All of the items in `types` can be accessed as attributes of the `BathArray` itself.

### Examples

Generate empty `BathArray` instance.

```
>>> ba = BathArray((3,))
>>> print(ba)
[(' ', [0., 0., 0.], [[0., 0., 0.], [0., 0., 0.], [0., 0., 0.]], [[0., 0., 0.], [0., 0., 0.],
↪0., 0.], [0., 0., 0.])]
(' ', [0., 0., 0.], [[0., 0., 0.], [0., 0., 0.], [0., 0., 0.]], [[0., 0., 0.], [0., 0., 0.],
↪0., 0.], [0., 0., 0.])]
(' ', [0., 0., 0.], [[0., 0., 0.], [0., 0., 0.], [0., 0., 0.]], [[0., 0., 0.], [0., 0., 0.],
↪0., 0.], [0., 0., 0.])]
```

Generate `BathArray` from the set of arrays:

```
>>> import numpy as np
>>> ca = np.random.random((3, 3))
>>> sn = ['1H', '2H', '3H']
>>> hf = np.random.random((3, 3, 3))
>>> ba = BathArray(ca=ca, hf=hf, sn=sn)
>>> print(ba.N, ba.types)
['1H' '2H' '3H'] SpinDict(1H: (1H, 0.5, 26.7519), 2H: (2H, 1, 4.1066, 0.00286), 3H:
↪(3H, 0.5, 28.535))
```

**Warning:** Due to how structured arrays work, if one uses a boolean array to access a subarray, and then access the name field, the initial array *will not change*.

Example:

```
>>> ba = BathArray((10,), sn='1H')
>>> print(ba.N)
['1H' '1H' '1H' '1H' '1H' '1H' '1H' '1H' '1H' '1H']
>>> bool_mask = np.arange(10) % 2 == 0
>>> ba[bool_mask]['N'] = 'e'
>>> print(ba.N)
['1H' '1H' '1H' '1H' '1H' '1H' '1H' '1H' '1H' '1H']
```

To achieve the desired result, one should first access the name field and only then apply the boolean mask:

```
>>> ba['N'][bool_mask] = 'e'
>>> print(ba.N)
['e' '1H' 'e' '1H' 'e' '1H' 'e' '1H' 'e' '1H']
```

### Parameters

- **shape** (*tuple*) – Shape of the array.
- **array** (*array-like*) – Either an unstructured array with shape (n, 3) containing coordinates of bath spins as rows OR structured ndarray with the same fields as the datatype of the bath.
- **name** (*array-like*) – Array of the bath spin name.
- **hyperfines** (*array-like*) – Array of the hyperfine tensors with shape (n, 3, 3).
- **quadrupoles** (*array-like*) – Array of the quadrupole tensors with shape (n, 3, 3).
- **efg** (*array-like*) – Array of the electric field gradients with shape (n, 3, 3) for each bath spin. Used to compute Quadrupole tensors for spins  $\geq 1$ . Requires the spin types either be found in `common_isotopes` or specified with `types` argument.
- **types** (`SpinDict`) – `SpinDict` or input to create one. Contains either `SpinTypes` of the bath spins or tuples which will initialize those. See `pycce.bath.SpinDict` documentation for details.
- **imap** (`InteractionMap`) – Instance of `InteractionMap` containing user defined interaction tensors between bath spins stored in the array.
- **ca** (*array-like*) – Shorthand notation for `array` argument.
- **sn** (*array-like*) – Shorthand notation for `name` argument.
- **hf** (*array-like*) – Shorthand notation for `hyperfines` argument.
- **q** (*array-like*) – Shorthand notation for `quadrupoles` argument.

**sort** (*axis=-1, kind=None, order=None*)

Sort array in-place. Is implemented only when `imap` is `None`. Otherwise use `np.sort`.

**property name**

Array of the name attribute for each spin in the array from `types` dictionary.

---

**Note:** While the value of this attribute should be the same as the `N` field of the `BathArray` instance, `.name` *should not* be used for production as it creates a *new* array from `types` dictionary.

---

**Type** ndarray

**property s**

Array of the `spin` (spin value) attribute for each spin in the array from `types` dictionary.

**Type** ndarray

**property dim**

Array of the `dim` (dimensions of the spin) attribute for each spin in the array from `types` dictionary.

**Type** ndarray

**property gyro**

Array of the `gyro` (gyromagnetic ratio) attribute for each spin in the array from `types` dictionary.

**Type** ndarray

**property q**

Array of the `q` (quadrupole moment) attribute for each spin in the array from `types` dictionary.

**Type** ndarray

**property detuning**

Array of the `detuning` attribute for each spin in the array from `types` dictionary.

**Type** ndarray

**property x**

Array of x coordinates for each spin in the array (`bath['xyz'][:, 0]`).

**Type** ndarray

**property y**

Array of y coordinates for each spin in the array (`bath['xyz'][:, 1]`).

**Type** ndarray

**property z**

Array of z coordinates for each spin in the array (`bath['xyz'][:, 2]`).

**Type** ndarray

**property N**

Array of name for each spin in the array (`bath['N']`).

**Type** ndarray

**property xyz**

Array of coordinates for each spin in the array (`bath['xyz']`).

**Type** ndarray

**property A**

Array of hyperfine tensors for each spin in the array (`bath['A']`).

**Type** ndarray

**property Q**

Array of quadrupole tensors for each spin in the array (`bath['Q']`).



Type ndarray

**add\_type**(\*args, \*\*kwargs)

Add spin type to the types dictionary.

**Parameters**

- **\*args** – Any number of positional inputs to create SpinDict entries. E.g. the tuples of form (name str, spin float, gyro float, q float).
- **\*\*kwargs** – Any number of keyword inputs to create SpinDict entries. E.g. name = (spin, gyro, q).

For details and allowed inputs see SpinDict documentation.

**Returns** A view of self.types instance.

**Return type** SpinDict

**add\_interaction**(i, j, tensor)

Add interactions tensor between bath spins with indexes i and j.

---

**Note:** If called from the subarray this method **does not** change the tensors of the total BathArray.

---

**Parameters**

- **i** (*int or ndarray (n,)*) – Index of the first spin in the pair or array of the indexes of the first spins in n pairs.
- **j** (*int or ndarray with shape (n,)*) – Index of the second spin in the pair or array of the indexes of the second spins in n pairs.
- **tensor** (*ndarray with shape (3,3) or (n, 3,3)*) – Interaction tensor between the spins i and j or array of tensors.

**update**(ext\_bath, error\_range=0.2, ignore\_isotopes=True, inplace=True)

Update the properties of the spins in the array using data from other BathArray instance. For each spin in ext\_bath check whether there is such spin in the array that has the same position within allowed error range given by error\_range and has the same name. If such spins is found in the array, then it's coordinates, hyperfine tensor and quadrupole tensor are updated using the values of the spin in the ext\_bath object.

If ignore\_isotopes is true, then the name check ignores numbers in the name of the spins.

**Parameters**

- **ext\_bath** (BathArray) – Array of the new spins.
- **error\_range** (float) – +- distance in Angstrom within which two positions are considered to be the same. Default is 0.2 A.
- **ignore\_isotopes** (bool) – True if ignore numbers in the name of the spins. Default True.
- **inplace** (bool) – True if changes parameters of the array in place. If False, returns copy of the array.

**Returns** updated BathArray instance.

**Return type** BathArray

**from\_point\_dipole**(*position*, *gyro\_e*=- 17608.59705, *inplace*=True)

Generate hyperfine couplings, assuming that bath spins interaction with central spin is the same as the one between two magnetic point dipoles.

**Parameters**

- **position** (*ndarray with shape (3,)*) – position of the central spin
- **gyro\_e** (*float or ndarray with shape (3,3)*) – gyromagnetic ratio of the central spin

**OR**

tensor corresponding to interaction between magnetic field and central spin.

- **inplace** (*bool*) – True if changes parameters of the array in place. If False, returns copy of the array.

**Returns** updated BathArray instance with changed hyperfine couplings.

**Return type** *BathArray*

**from\_cube**(*cube*, *gyro\_e*=- 17608.59705, *inplace*=True)

Generate hyperfine couplings, assuming that bath spins interaction with central spin can be approximated as a point dipole, interacting with given spin density distribution.

**Parameters**

- **cube** (*Cube*) – An instance of *Cube* object, which contains spatial distribution of spin density. For details see documentation of *Cube* class.
- **gyro\_e** (*float*) – Gyromagnetic ratio of the central spin.
- **inplace** (*bool*) – True if changes parameters of the array in place. If False, returns copy of the array.

**Returns** Updated BathArray instance with changed hyperfine couplings.

**Return type** *BathArray*

**from\_func**(*func*, *gyro\_e*=- 17608.59705, *vectorized*=False, *inplace*=True)

Generate hyperfine couplings from user-defined function.

**Parameters**

- **func** (*func*) – Callable with signature:

```
func(coord, gyro, central_gyro)
```

where *coord* is array of the bath spin coordinate, *gyro* is the gyromagnetic ratio of bath spin, *central\_gyro* is the gyromagnetic ratio of the central bath spin.

- **gyro\_e** (*float*) – gyromagnetic ratio of the central spin to be used in the function.
- **vectorized** (*bool*) – If True, assume that *func* takes arrays of all bath spin coordinates and array of gyromagnetic ratios as arguments.
- **inplace** (*bool*) – True if changes parameters of the array in place. If False, returns copy of the array.

**Returns** Updated BathArray instance with changed hyperfine couplings.

**Return type** *BathArray*

**from\_efg**(*efg*, *inplace*=True)

Generate quadrupole splittings from electric field gradient tensors for spins  $\geq 1$ .

**Parameters**

- **efg** (*array-like*) – Array of the electric field gradients for each bath spin. The data for spins-1/2 should be included but can be any value.
- **inplace** (*bool*) – True if changes parameters of the array in place. If False, returns copy of the array.

**Returns** Updated BathArray instance with changed quadrupole tensors.

**Return type** *BathArray*

**dist**(*position=None*)

Compute the distance of the bath spins from the given position.

**Parameters** **position** (*ndarray with shape (3,)*) – Cartesian coordinates of the position from which to compute the distance. Default is (0, 0, 0).

**Returns** Array of distances of each bath spin from the given position in angstrom.

**Return type** ndarray with shape (n,)

**savetxt**(*filename, fmt='%18.8f', strip\_isotopes=False, \*\*kwargs*)

Save name of the isotopes and their coordinates to the txt file of xyz format.

**Parameters**

- **filename** (*str or file*) – Filename or file handle.
- **fmt** (*str*) – Format of the coordinate entry.
- **strip\_isotopes** (*bool*) – True if remove numbers from the name of bath spins. Default False.
- **\*\*kwargs** – Additional keywords of the `numpy.savetxt` function.

**sort**(*a, \*args, \*\*kwargs*)

Return a indexes of an sorted array. Overrides `numpy.argsort` function.

**same\_bath\_indexes**(*barray\_1, barray\_2, error\_range=0.2, ignore\_isotopes=True*)

Find indexes of the same array elements in two BathArray instances.

**Parameters**

- **barray\_1** (*BathArray*) – First array.
- **barray\_2** (*BathArray*) – Second array.
- **error\_range** (*float*) – If distance between positions in two arrays is smaller than `error_range` they are assumed to be the same.
- **ignore\_isotopes** (*bool*) – True if ignore numbers in the name of the spins. Default True.

**Returns**

tuple containing:

- **ndarray**: Indexes of the elements in the first array found in the second.
- **ndarray**: Indexes of the elements in the second array found in the first.

**Return type** tuple

**utilities.rotmatrix**(*final\_vector*)

Generate 3D rotation matrix which applied on initial vector will produce vector, aligned with final vector.

## Examples

```
>>> R = rotmatrix([0,0,1], [1,1,1])
>>> R @ np.array([0,0,1])
array([0.577, 0.577, 0.577])
```

### Parameters

- **initial\_vector** (*ndarray with shape (3, )*) – Initial vector.
- **final\_vector** (*ndarray with shape (3, )*) – Final vector.

**Returns** Rotation matrix.

**Return type** ndarray with shape (3, 3)

## 4.3.1 InteractionMap

**class InteractionMap**(*rows=None, columns=None, tensors=None*)

Dict-like object containing information about tensor interactions between two spins.

Each key is a tuple of two bath spin indexes.

### Parameters

- **rows** (*array-like with shape (n,)*) – Indexes of the bath spins, appearing on the left in the pairwise interaction.
- **columns** (*array-like with shape (n,)*) – Indexes of the bath spins, appearing on the right in the pairwise interaction.
- **tensors** (*array-like with shape (n, 3, 3)*) – Tensors of pairwise interactions between two spins with the indexes in rows and columns.

### mapping

Actual dictionary storing the data.

**Type** dict

### property indexes

Array with the indexes of pairs of bath spins, for which the tensors are stored.

**Type** ndarray with shape (n, 2)

### shift(*start, inplace=True*)

Add an offset *start* to the indexes. If *inplace* is False, returns the copy of InteractionMap.

### Parameters

- **start** (*int*) – Offset in indexes.
- **inplace** (*bool*) – If True, makes changes inplace. Otherwise returns copy of the map.

**Returns** Map with shifted indexes.

**Return type** *InteractionMap*

**keys()** → a set-like object providing a view on D's keys

**items()** → a set-like object providing a view on D's items

### subspace(*array*)

Get new InteractionMap with indexes readressed from array. Within the subspace indexes are renumbered.

## Examples

The subspace of [3,4,7] indexes will contain InteractionMap only within [3,4,7] elements with new indexes [0, 1, 2].

```
>>> import numpy as np
>>> im = InteractionMap()
>>> im[0, 3] = np.eye(3)
>>> im[3, 7] = np.ones(3)
>>> for k in im: print(k, '\n', im[k],)
(0, 3)
[[1. 0. 0.]
 [0. 1. 0.]
 [0. 0. 1.]]
(3, 7)
[[1. 1. 1.]
 [1. 1. 1.]
 [1. 1. 1.]]
>>> array = [3, 4, 7]
>>> sim = im.subspace(array)
>>> for k in sim: print(k, '\n', sim[k])
(0, 2)
[[1. 1. 1.]
 [1. 1. 1.]
 [1. 1. 1.]]
```

**Parameters** **array** (*ndarray*) – Either bool array containing True for elements within the subspace or array of indexes presented in the subspace.

**Returns** The map for the subspace.

**Return type** *InteractionMap*

**classmethod** **from\_dict**(*dictionary*, *presorted=False*)

Generate InteractionMap from the dictionary. :param dictionary: Dictionary with tensors. :type dictionary: dict :param presorted: If true, assumes that the keys in the dictionary were already presorted. :type presorted: bool

**Returns** New instance generated from the dictionary.

**Return type** *InteractionMap*

### 4.3.2 Cube

**class** **Cube**(*filename*)

Class to process the .cube datafiles with spin polarization.

**Parameters** **filename** (*str*) – Name of the .cube file.

**comments**

First two lines of the .cube file.

**Type** *str*

**origin**

Coordinates of the origin in angstrom.

**Type** ndarray with shape (3,)

**voxel**

Parameters of the voxel - unit of the 3D grid in angstrom.

**Type** ndarray with shape (3,3)

**size**

Size of the cube.

**Type** ndarray with shape (3,)

**atoms**

Array of atoms in the cube.

**Type** BathArray with shape (n)

**data**

Data stored in cube.

**Type** ndarray with shape (size[0], size[1], size[2])

**grid**

Coordinates of the points at which data is computed.

**Type** ndarray with shape (size[0], size[1], size[2], 3)

**integral**

Data integrated over cube.

**Type** float

**spin**

integral / 2 - total spin.

**Type** float

**transform**(*rotmatrix=None, shift=None*)

Changes coordinates of the grid. DOES NOT ASSUME PERIODICITY.

**Parameters**

- **rotmatrix** (ndarray with shape (3, 3)) – Rotation matrix  $R$ :

$$R = \begin{bmatrix} n_1^{(1)} & n_1^{(2)} & n_1^{(3)} \\ n_2^{(1)} & n_2^{(2)} & n_2^{(3)} \\ n_3^{(1)} & n_3^{(2)} & n_3^{(3)} \end{bmatrix}$$

where  $n_i^{(j)}$  corresponds to the coefficient of initial basis vector  $i$  for  $j$  new basis vector:

$$e'_j = n_1^{(j)} \vec{e}_1 + n_2^{(j)} \vec{e}_2 + n_3^{(j)} \vec{e}_3$$

In other words, columns of  $R$  are coordinates of the new basis in the old basis.

Given vector in initial basis  $v = [v_1, v_2, v_3]$ , vector in new basis is given as  $v' = R.T @ v$ .

- **shift** (ndarray with shape (3,)) – Shift in the origin of coordinates (in the old basis).

**integrate**(*position, gyro\_n, gyro\_e=-17608.59705, spin=None*)

Integrate over polarization data, stored in Cube object, to obtain hyperfine dipolar-dipolar tensor.

**Parameters**

- **position** (*ndarray with shape (3,) or (n, 3)*) – Position of the bath spin at which to compute hyperfine tensor or array of positions.
- **gyro\_n** (*float or ndarray with shape (n,)*) – Gyromagnetic ratio of the bath spin or array of the ratios.
- **gyro\_e** (*float*) – Gyromagnetic ratio of central spin.
- **spin** (*float*) – Total spin of the central spin. If not given, taken from the integral of the polarization.

**Returns** Hyperfine tensor or array of hyperfine tensors.

**Return type** ndarray with shape (3, 3) or (n, 3, 3)

## 4.4 SpinDict and SpinType

Documentation for the `SpinDict` - dict-like class which describes the properties of the different types of the spins in the bath.

**class SpinDict**(\*args, \*\*kwargs)

Wrapper class for dictionary tailored for containing properties of the spin types. Can take `np.void` or `BathArray` instances as keys. Every entry is instance of the `SpinType`.

Each entry of the `SpinDict` can be initialized as follows:

- As a Tuple containing name (optional), spin, gyromagnetic ratio, quadrupole constant (optional) and de-tuning (optional).
- As a `SpinType` instance.

### Examples

```
>>> types = SpinDict()
>>> types['1H'] = ('1H', 1 / 2, 26.7519)
>>> types['2H'] = 1, 4.1066, 0.00286
>>> types['3H'] = SpinType('3H', 1 / 2, 28.535, 0)
>>> print(types)
SpinDict({'1H': (1H, 0.5, 26.7519, 0.0), '2H': (2H, 1, 4.1066, 0.00286), '3H': (3H, ↵
↵0.5, 28.535, 0)})
```

If `SpinType` of the given bath spin is not provided, when requested `SpinDict` will try to find information about the bath spins in the `common_isotopes`.

If found, adds an entry to the given `SpinDict` instance and returns it. Otherwise `KeyError` is raised.

To initialize several `SpinType` entries one can use `add_types` method.

#### Parameters

- **\*args** – Any numbers of arguments which could initialize `SpinType` instances.
- **\*\*kwargs** – Any numbers of keyword arguments which could initialize `SpinType` instances. For details see `SpinDict.add_type` method.

**add\_type**(\*args, \*\*kwargs)

Add one or several spin types to the spin dictionary.

#### Parameters

- **\*args** – Any numbers of arguments which could initialize `SpinType` instances. Accepted arguments:
  - Tuple containing name, spin, gyromagnetic ratio, quadrupole constant (optional) and detuning (optional).
  - `SpinType` instance.

Can also initialize one instance of `SpinType` if each argument corresponds to each positional argument necessary to initialize.

- **\*\*kwargs** – Any numbers of keyword arguments which could initialize `SpinType` instances. Usefull as an alternative for updating the dictionary. for each keyword argument adds an entry to the `SpinDict` with the same name as keyword.

## Examples

```
>>> types = SpinDict()
>>> types.add_type('1H', 1 / 2, 26.7519)
>>> types.add_type(('1H_det', 1 / 2, 26.7519, 10), ('2H', 1, 4.1066, 0.00286),
>>>               SpinType('3H', 1 / 2, 28.535, 0), e=(1 / 2, 6.7283, 0))
>>> print(types)
SpinDict(1H: (1H, 0.5, 26.7519), 1H_det: (1H_det, 0.5, 26.7519, 10),
2H: (2H, 1, 4.1066, 0.00286), 3H: (3H, 0.5, 28.535), e: (e, 0.5, 6.7283))
```

**class SpinType**(name, s=0.0, gyro=0.0, q=0.0, detuning=0.0)  
Class which contains properties of each spin type in the bath.

### Parameters

- **name** (*str*) – Name of the bath spin.
- **s** (*float*) – Total spin of the bath spin.  
Default 0.
- **gyro** (*float*) – Gyromagnetic ratio in rad \* kHz / G.  
Default 0.
- **q** (*float*) – Quadrupole moment in barn (for s > 1/2).  
Default 0.
- **detuning** (*float*) – Energy detuning from the zeeman splitting in kHz, included as an extra  $+\omega\hat{S}_z$  term in the Hamiltonian, where  $\omega$  is the detuning.  
Default 0.

### name

Name of the bath spin.

**Type** str

### s

Total spin of the bath spin.

**Type** float

### dim

Spin dimensionality = 2s + 1.

**Type** int



**gyro**

Gyromagnetic ratio in rad/(ms \* G).

**Type** float

**q**

Quadrupole moment in barn (for  $s > 1/2$ ).

**Type** float

**detuning**

Energy detuning from the zeeman splitting in kHz.

**Type** float

**common\_isotopes** = `SpinDict(1H: (0.5, 26.7522), 2H: (1.0, 4.1066, 0.0029), 3He: (0.5, -20.3789), ...)`

An instance of the `SpinDict` dictionary, containing properties for the most of the common isotopes with nonzero spin. The isotope is considered common if it is stable and has nonzero concentration in nature.

**Type** *SpinDict*

**common\_concentrations** = `{element ('H', 'He', ...) : { isotope ('1H', '2H', ..) : concentration}}`

Nested dict containing natural concentrations of the stable nuclear isotopes.

**Type** dict



## RUNNING THE SIMULATIONS

### 5.1 Setting up the Simulator Object

Documentation for the `pycce.Simulator` - main class for conducting CCE Simulations.

```
class Simulator(spin, position=None, alpha=None, beta=None, gyro=- 17608.59705, magnetic_field=None,  
D=0.0, E=0.0, r_dipole=None, order=None, bath=None, pulses=None, as_delay=False,  
n_clusters=None, **bath_kw)
```

The main class for CCE calculations.

The typical usage includes:

1. Read array of the bath spins. This is done with `Simulator.read_bath` method which accepts either reading from `.xyz` file or from the `BathArray` instance with defined positions and names of the bath spins. In the process, the subset of the array within the distance of `r_dipole` from the central spin is taken and for this subset the Hyperfine couplings can be generated.

If no `hyperfine` keyword is provided and there are some hyperfine couplings already, then no changes are done to the hyperfine tensors. If `hyperfine='pd'`, the hyperfine couplings are computed assuming point dipole approximation. For all accepted arguments, see `Simulator.read_bath`.

2. Generate set of clusters with `Simulator.generate_clusters`, determined by the maximum connectivity radius `r_dipole` and the maximum size of the cluster order.
3. Compute the desired property with `Simulator.compute` method.

---

**Note:** Directly setting up the attribute values will rerun `Simulator.read_bath` and/or `Simulator.generate_clusters` to reflect updated value of the given attribute.

E.g. If `Simulator.r_bath` is set to some new value after initialization, then `Simulator.read_bath` and `Simulator.generate_clusters` are called with the increased bath.

---

First two steps are usually done during the initialization of the `Simulator` object by providing the necessary arguments.

## Notes

Depending on the number of provided arguments, in the initialization process will call the following methods to setup the calculation engine.

- If `bath` is provided, `Simulator.read_bath` is called with additional keywords in `**bath_kw`.
- If both `r_dipole` and `order` are provided and `bath` is not `None`, the `Simulator.generate_clusters` is called.

See the corresponding method documentation for details.

Examples:

```
>>> atoms = random_bath('13C', 100, number=2000, seed=10)
>>> calc = Simulator(1, bath=atoms, r_bath=40, r_dipole=6,
>>>                  order=2, D=2.88 * 2 * np.pi * 1e6,
>>>                  magnetic_field=500, pulses=1)
>>> print(calc)
Simulator for spin-1.
alpha: [0.+0.j 1.+0.j 0.+0.j]
beta: [0.+0.j 0.+0.j 1.+0.j]
gyromagnetic ratio: -17608.59705 kHz * rad / G
zero field splitting:
array([[ -6031857.895,    0.    ,    0.    ],
       [    0.    , -6031857.895,    0.    ],
       [    0.    ,    0.    , 12063715.79 ]])
magnetic field:
array([ 0.,  0., 500.])

Parameters of cluster expansion:
r_bath: 40
r_dipole: 6
order: 2

Bath consists of 549 spins.

Clusters include:
549 clusters of order 1.
457 clusters of order 2.
```

## Parameters

- **spin** (*float*) – Total spin of the central spin.
- **position** (*ndarray*) – Cartesian coordinates in Angstrom of the central spin. Default (0., 0., 0.).
- **alpha** (*float or ndarray with shape (2s+1, )*) – 0 state of the qubit in  $S_z$  basis or the index of eigenstate to be used as one.  
Default: state with  $m_s = +s$  where  $m_s$  is the z-projection of the spin and  $s$  is the total spin if no information of central spin Hamiltonian is provided. Otherwise lowest energy eigenstate of the central spin Hamiltonian.
- **beta** (*float or ndarray with shape (2s+1, )*) – 1 state of the qubit in  $S_z$  basis or the index of the eigenstate to be used as one.

Default: state with  $m_s = +s - 1$  where  $m_s$  is the z-projection of the spin and  $s$  is the total spin if no information of central spin Hamiltonian is provided. Otherwise second lowest energy eigenstate of the central spin Hamiltonian.

- **gyro** (*float or ndarray with shape (3, 3)*) – Gyromagnetic ratio of central spin in rad / ms / G.

OR

Tensor describing central spin interactions with the magnetic field.

Default -17608.597050 kHz \* rad / G - gyromagnetic ratio of the free electron spin.

- **D** (*float or ndarray with shape (3, 3)*) – D (longitudinal splitting) parameter of central spin in ZFS tensor of central spin in kHz.

OR

Total ZFS tensor. Default 0.

- **E** (*float*) – E (transverse splitting) parameter of central spin in ZFS tensor of central spin in kHz. Default 0. Ignored if D is None or tensor.

- **bath** (*ndarray or str*) – First positional argument of the `Simulator.read_bath` method.

Either:

– Instance of `BathArray` class;

– ndarray with `dtype([('N', np.unicode_, 16), ('xyz', np.float64, (3, ))])` containing names of bath spins (same ones as stored in `self.n_type`) and positions of the spins in angstroms;

– the name of the xyz text file containing 4 cols: name of the bath spin and xyz coordinates in A.

- **r\_dipole** (*float*) – Maximum connectivity distance between two bath spins.
- **order** (*int*) – Maximum size of the cluster to be considered in CCE expansion.
- **n\_clusters** (*dict*) – Dictionary which contain maximum number of clusters of the given size. Has the form `n_clusters = {order: number}`, where `order` is the size of the cluster, `number` is the maximum number of clusters with this size.

If provided, sort the clusters by the strength of cluster interaction, equal to the lowest pairwise interaction in the cluster. Then the strongest number of clusters is taken.

- **pulses** (*list or int or Sequence*) – Number of pulses in CPMG sequence or list with pulses.
- **\*\*bath\_kw** – Additional keyword arguments for the `Simulator.read_bath` method.

#### **position**

Position of the central spin in Cartesian coordinates.

**Type** ndarray with shape (3, )

#### **spin**

Value of the central spin  $s$ .

**Type** float

**gyro**

Gyromagnetic ratio of central spin in rad / ms / G.

*OR*

Tensor describing central spin interactions with the magnetic field.

Default -17608.597050 kHz \* rad / G - gyromagnetic ratio of the free electron spin.

**Type** gyro (float or ndarray with shape (3,3))

**zfs**

Zero field splitting tensor of the central spin

**Type** ndarray with shape (3,3)

**clusters**

Dictionary containing information about cluster structure of the bath.

Each keys n correspond to the size of the cluster. Each `Simulator.clusters[n]` contains ndarray of shape (m, n), where m is the number of clusters of given size, n is the size of the cluster. Each row of this array contains indexes of the bath spins included in the given cluster. Generated during `.generate_clusters` call.

**Type** dict

**as\_delay**

True if time points are delay between pulses (for equispaced pulses), False if time points are total time. Ignored if `pulses` contains the time delays.

**Type** bool

**state**

Initial state of the qubit in gCCE simulations. Assumed to be  $1/\sqrt{2}(0 + 1)$  unless provided during `Simulator.compute` call.

**Type** ndarray

**interlaced**

True if use hybrid CCE approach - for each cluster sample over states of the supercluster.

**Type** bool

**seed**

Seed for random number generator, used in random bath states sampling.

**Type** int

**nbstates**

Number of random bath states to sample over.

**Type** int

**fixstates**

If not None, shows which bath states to fix in random bath states.

Each key is the index of bath spin, value - fixed  $\hat{S}_z$  projection of the mixed state of nuclear spin.

**Type** dict

**masked**

True if mask numerically unstable points (with coherence > 1) in the averaging over bath states.

---

**Note:** It is up to user to check whether the possible instability is due to numerical error or unphysical assumptions of the calculations.

---

**Type** bool

**second\_order**

True if add second order perturbation theory correction to the cluster Hamiltonian in conventional CCE. Relevant only for conventional CCE calculations.

**Type** bool

**level\_confidence**

Maximum fidelity of the qubit state to be considered eigenstate of the central spin Hamiltonian when `second_order` set to True.

**Type** float

**projected\_bath\_state**

Array with z-projections of the bath spins states.

**Type** ndarray with shape (n,)

**bath\_state**

Array of bath states.

**Type** bath\_state (ndarray)

**timespace**

Time points at which compute the desired property.

**Type** timespace (ndarray with shape (n,))

**property\_alpha**

0 qubit state of the central spin in Sz basis **OR** index of the energy state to be considered as one.

**Type** ndarray or int

**property\_beta**

1 qubit state of the central spin in Sz basis **OR** index of the energy state to be considered as one.

**Type** ndarray or int

**property\_magnetic\_field**

Array containing external magnetic field as (Bx, By, Bz). Default (0, 0, 0).

**Type** ndarray

**property\_order**

Maximum size of the cluster.

**Type** int

**property\_n\_clusters**

Dictionary which contain maximum number of clusters of the given size. Has the form `n_clusters = {order: number}`, where `order` is the size of the cluster, `number` is the maximum number of clusters with this size.

If provided, sorts the clusters by the strength of cluster interaction, equal to the lowest pairwise interaction in the cluster. Then the strongest `number` of clusters is taken.

**Type** dict

**property r\_dipole**

Maximum connectivity distance.

**Type** float

**property pulses**

List-like object, containing the sequence of the instantaneous ideal control pulses.

Each item is Pulse object, containing the following attributes:

- **axis** (*str*): Axis of rotation of the central spin. Can be 'x', 'y', or 'z'.
- **angle** (*float or str*): Angle of rotation of central spin. Can be provided in rad, or as a string, containing fraction of pi: 'pi', 'pi/2', '2\*pi' etc. Default is None.
- **delay** (*float or ndarray*): Delay before the pulse or array of delays with the same shape as time points.
- **bath\_names** (*str or array-like of str*): Name or array of names of bath spin types, impacted by the bath pulse.
- **bath\_axes** (*str or array-like of str*): Axis of rotation or array of axes of the bath spins. If **bath\_names** is provided, but **bath\_axes** and **bath\_angles** are not, assumes the same axis and angle as the one of the central spin.
- **bath\_angles** (*float or str or array-like*): Angle of rotation or array of axes of rotations of the bath spins.

If delay is not provided in **all** pulses, assumes even delay of CPMG sequence.

If only **some** delays are provided, assumes 0 delay in the pulses without delay provided.

**Type** *Sequence*

**set\_zfs**(*D=None, E=0*)

Set Zero Field Splitting of the central spin from longitudinal ZFS *D* and transverse ZFS *E*.

**Parameters**

- **D** (*float or ndarray with shape (3, 3)*) – D (longitudinal splitting) parameter of central spin in ZFS tensor of central spin in kHz.

**OR**

Total ZFS tensor. Default 0.

- **E** (*float*) – E (transverse splitting) parameter of central spin in ZFS tensor of central spin in kHz. Default 0. Ignored if D is None or tensor.

**set\_magnetic\_field**(*magnetic\_field=None*)

Set magnetic field from either value of the magnetic field along z-direction or full magnetic field vector.

**Parameters magnetic\_field** (*float or array-like*) – Magnetic field along z-axis.

**OR**

Array containing external magnetic field as (Bx, By, Bz). Default (0, 0, 0).

**set\_states**(*alpha=None, beta=None*)

Set 0 and 1 Qubit states of the Simulator object.

**Parameters**

- **alpha** (*int or ndarray with shape (2s+1, )*) – 0 state of the qubit in  $S_z$  basis or the index of eigenstate to be used as one.



Default: Lowest energy eigenstate of the central spin Hamiltonian. Otherwise state with  $m_s = +s$  where  $m_s$  is the z-projection of the spin and  $s$  is the total spin if no information of central spin Hamiltonian is provided.

- **beta** (*int* or *ndarray with shape (2s+1, )*) – 1 state of the qubit in  $S_z$  basis or the index of the eigenstate to be used as one.

Default: Second lowest energy eigenstate of the central spin Hamiltonian. Otherwise state with  $m_s = +s - 1$  where  $m_s$  is the z-projection of the spin and  $s$  is the total spin if no information of central spin Hamiltonian is provided.

**eigenstates**(*alpha=None, beta=None, magnetic\_field=None, D=None, E=0, return\_eigen=True*)

Compute eigenstates of the central spin Hamiltonian.

If **alpha** is provided, set alpha state as eigenstate. Similarly, if **beta** is provided, set beta state as eigenstate

#### Parameters

- **alpha** (*int*) – Index of the state to be considered as 0 (alpha) qubit state in order of increasing energy (0 - lowest energy).
- **beta** (*int*) – Index of the state to be considered as 1 (beta) qubit state.
- **magnetic\_field** (*ndarray with shape (3,)*) – Array containing external magnetic field as (Sx, By, Bz).
- **D** (*float* or *ndarray with shape (3, 3)*) – D (longitudinal splitting) parameter of central spin in kHz OR total ZFS tensor.
- **E** (*float*) – E (transverse splitting) parameter of central spin in kHz. Ignored if D is None or tensor.
- **return\_eigen** (*bool*) – If true, returns eigenvalues and eigenvectors of the central spin Hamiltonian.

#### Returns

*tuple* containing:

- **ndarray with shape (2s+1,)**: Array with eigenvalues of the central spin Hamiltonian.
- **ndarray with shape (2s+1, 2s+1)**: Array with eigenvectors of the central spin Hamiltonian. Each column of the array is eigenvector.

**Return type** tuple

## 5.2 Reading the Bath

Documentation for the `Simulator.read_bath` and `Simulator.generate_clusters` method. These methods are called automatically on the initialization of the `Simulator` object if the necessary keywords are provided. Otherwise they can also be called by themselves to update the properties of the spin bath in `Simulator` object.

`Simulator.read_bath`(*bath=None, r\_bath=None, skiprows=1, external\_bath=None, hyperfine=None, types=None, error\_range=None, ext\_r\_bath=None, imap=None*)

Read spin bath from the file or from the `BathArray`.

#### Parameters

- **bath** (*ndarray, BathArray* or *str*) – Either:
  - Instance of `BathArray` class;

- ndarray with dtype([('N', np.unicode\_, 16), ('xyz', np.float64, (3, ))]) containing names of bath spins (same ones as stored in self.n\_type) and positions of the spins in angstroms;
- the name of the xyz text file containing 4 cols: name of the bath spin and xyz coordinates in A.
- **r\_bath** (*float*) – Cutoff size of the spin bath.
- **skiprows** (*int*, *optional*) – If bath is name of the file, this argument gives number of rows to skip while reading the .xyz file (default 1).
- **external\_bath** (*BathArray*, *optional*) – BathArray containing spins read from DFT output (see `pycce.io`).
- **hyperfine** (*str*, *func*, or *Cube instance*, *optional*) – This argument tells the code how to generate hyperfine couplings.

If (`hyperfine = None` and all A in provided bath are 0) or (`hyperfine = 'pd'`), use point dipole approximation.

Otherwise can be an instance of `Cube` object, or callable with signature: `func(coord, gyro, central_gyro)`, where `coord` is array of the bath spin coordinate, `gyro` is the gyromagnetic ratio of bath spin, `central_gyro` is the gyromagnetic ratio of the central bath spin.

- **types** (*SpinDict*) – `SpinDict` or input to create one. Contains either `SpinTypes` of the bath spins or tuples which will initialize those.

See `pycce.bath.SpinDict` documentation for details.

- **error\_range** (*float*, *optional*) – Maximum distance between positions in bath and external bath to consider two positions the same (default 0.2).
- **ext\_r\_bath** (*float*, *optional*) – Maximum distance from the central spins of the bath spins for which to use the DFT positions.
- **imap** (*InteractionMap*) – Instance of `InteractionMap` class, containing interaction tensors for bath spins. Each key of the `InteractionMap` is a tuple with indexes of two bath spins. The value is the 3x3 tensor describing the interaction between two spins in a format:

$$I^i J I^j = I_x^i J_{xx} I_x^j + I_x^i J_{xy} I_y^j \dots$$

---

**Note:** For each bath spin pair without interaction tensor in `imap`, coupling is approximated assuming magnetic point dipole–dipole interaction. If `imap = None` all interactions between bath spins are approximated in this way. Then interaction tensor between spins  $i$  and  $j$  is computed as:

$$\mathbf{J}_{ij} = -\gamma_i \gamma_j \frac{\hbar^2}{4\pi\mu_0} \left[ \frac{3\vec{r}_{ij} \otimes \vec{r}_{ij} - |\vec{r}_{ij}|^2 \mathbf{I}}{|\vec{r}_{ij}|^5} \right]$$

Where  $\gamma_i$  is gyromagnetic ratio of  $i$  spin,  $\mathbf{I}$  is 3x3 identity matrix, and  $\vec{r}_{ij}$  is distance between two spins.

---

**Returns** The view of `Simulator.bath` attribute, generated by the method.

**Return type** *BathArray*

`Simulator.generate_clusters`(*order=None*, *r\_dipole=None*, *r\_inner=0*, *strong=False*, *ignore=None*, *n\_clusters=None*)

Generate set of clusters used in CCE calculations.

The clusters are generated from the following procedure:

- Each bath spin  $i$  forms a cluster of one.
- Bath spins  $i$  and  $j$  form cluster of two if there is an edge between them (distance  $d_{ij} \leq r_{\text{dipole}}$ ).
- Bath spins  $i$ ,  $j$ , and  $k$  form a cluster of three if enough edges connect them (e.g., there are two edges  $ij$  and  $jk$ )
- And so on.

In general, we assume that spins  $\{i..n\}$  form clusters if they form a connected graph. Only clusters up to the size imposed by the order parameter (equal to CCE order) are included.

#### Parameters

- **order** (*int*) – Maximum size of the cluster.
- **r\_dipole** (*float*) – Maximum connectivity distance.
- **r\_inner** (*float*) – Minimum connectivity distance.
- **strong** (*bool*) – True - generate only clusters with “strong” connectivity (all nodes should be interconnected). Default False.
- **ignore** (*list or str, optional*) – If not None, includes the names of bath spins which are ignored in the cluster generation.
- **n\_clusters** (*dict*) – Dictionary which contain maximum number of clusters of the given size. Has the form `n_clusters = {order: number}`, where `order` is the size of the cluster, `number` is the maximum number of clusters with this size.

If provided, sort the clusters by the strength of cluster interaction, equal to the lowest pairwise interaction in the cluster. Then the strongest number of clusters is taken.

#### Returns

**View of Simulator.clusters.** `Simulator.clusters` is a dictionary with keys corresponding to size of the cluster.

I.e. `Simulator.clusters[n]` contains ndarray of shape (m, n), where m is the number of clusters of given size, n is the size of the cluster. Each row contains indexes of the bath spins included in the given cluster.

**Return type** dict

## 5.3 Calculate Properties with Simulator

Documentation for the `Simulator.compute` method - the interface to run calculations with PyCCE.

`Simulator.compute(timespace, quantity='coherence', method='cce', **kwargs)`

General function for computing properties with CCE.

The dynamics are simulated using the Hamiltonian:

$$\begin{aligned}\hat{H}_S &= \mathbf{SDS} + \mathbf{B}\gamma_S\mathbf{S} \\ \hat{H}_{SB} &= \sum_i \mathbf{S}\mathbf{A}_i\mathbf{I}_i \\ \hat{H}_B &= \sum_i \mathbf{I}_i\mathbf{P}_i\mathbf{I}_i + \mathbf{B}\gamma_i\mathbf{I}_i + \sum_{i>j} \mathbf{I}_i\mathbf{J}_{ij}\mathbf{I}_j\end{aligned}$$

Here  $\hat{H}_S$  is the central spin Hamiltonian with Zero Field splitting tensor  $\mathbf{D}$  and gyromagnetic ratio tensor  $\gamma_S = \mu_S g$  are read from `Simulator.zfs` and `Simulator.gyro` respectively.

The  $\hat{H}_{SB}$  is the Hamiltonian describing interactions between central spin and the bath. The hyperfine coupling tensors  $\mathbf{A}_i$  are read from the `BathArray` stored in `Simulator.bath['A']`. They can be generated using point dipole approximation or provided by the user (see `Simulator.read_bath` for details).

The  $\hat{H}_B$  is the Hamiltonian describing interactions between the bath spins. The self interaction tensors  $\mathbf{P}_i$  are read from the `BathArray` stored in `Simulator.bath['Q']` and have to be provided by the user.

The gyromagnetic ratios  $\gamma_i$  are read from the `BathArray.gyros` attribute, which is generated from the properties of the types of bath spins, stored in `BathArray.types`. They can either be provided by user or read from the `pycce.common_isotopes` object.

The interaction tensors  $\mathbf{J}_{ij}$  are assumed from point dipole approximation or can be provided in `BathArray.imap` attribute.

---

**Note:** The `compute` method takes two keyword arguments to determine which quantity to compute and how:

- *method* can take 'cce' or 'gcce' values, and determines which method to use - conventional or generalized CCE.
- *quantity* can take 'coherence' or 'noise' values, and determines which quantity to compute - coherence function or autocorrelation function of the noise.

Each of the methods can be performed with monte carlo bath state sampling (if `nbstates` keyword is non zero) and with interlaced averaging (If `interlaced` keyword is set to `True`).

---

## Examples

First set up `Simulator` object using random bath of 1000 <sup>13</sup>C nuclear spins.

```
>>> import pycce as pc
>>> import numpy as np
>>> atoms = pc.random_bath('13C', 100, number=2000, seed=10) # Random spin bath
>>> calc = pc.Simulator(1, bath=atoms, r_bath=40, r_dipole=6,
>>>                    order=2, D=2.88 * 1e6, # D of NV in GHz -> kHz
>>>                    magnetic_field=500, pulses=1)
>>> ts = np.linspace(0, 2, 101) # timesteps
```

We set magnetic field to 500 G along z-axis and chose 1 decoupling pulse (Hahn-echo) in this example. The zero field splitting is set to the one of NV center in diamond.

Run conventional CCE calculation at time points `timespace` to obtain coherence without second order effects:

```
>>> calc.compute(ts)
```

This will call `Simulator.cce_coherence` method with default keyword values.

Compute the coherence conventional CCE coherence with second order interactions between bath spins:

```
>>> calc.compute(ts, second_order=True)
```

Compute the coherence with conventional CCE with bath state sampling (over 10 states):

```
>>> calc.compute(ts, nbstates=10)
```

Compute the coherence with generalized CCE:

```
>>> calc.compute(ts, method='gcce')
```

This will call `Simulator.gcce_dm` method with default keyword values and obtain off diagonal element as  $0\hat{\rho}_C1$ , where  $\hat{\rho}_C$  is the density matrix of the qubit.

Compute the coherence with generalized CCE with bath state sampling (over 10 states):

```
>>> calc.compute(ts, method='gcce', nbstates=10)
```

### Parameters

- **timespace** (*ndarray with shape (n,)*) – Time points at which compute the desired property.
- **quantity** (*str*) – Which quantity to compute. Case insensitive.  
Possible values:
  - 'coherence': compute coherence function.
  - 'noise': compute noise autocorrelation function.
- **method** (*str*) – Which implementation of CCE to use. Case insensitive.  
Possible values:
  - 'cce': conventional CCE, where interactions are mapped on 2 level pseudospin.
  - 'gcce': Generalized CCE where central spin is included in each cluster.
- **magnetic\_field** (*ndarray with shape (3,)*) – Magnetic field vector of form (Bx, By, Bz).  
Default is **None**. Overrides `Simulator.magnetic_field` if provided.
- **D** (*float or ndarray with shape (3,3)*) – D (longitudinal splitting) parameter of central spin in ZFS tensor of central spin in kHz.

### OR

Total ZFS tensor.

Default is **None**. Overrides `Simulator.zfs` if provided.

- **E** (*float*) – E (transverse splitting) parameter of central spin in ZFS tensor of central spin in kHz. Ignored if D is None or tensor.  
Default is 0.
- **pulses** (*list or int or Sequence*) – Number of pulses in CPMG sequence.

### OR

Sequence of the instantaneous ideal control pulses. It can be provided as an instance of `Sequence` class (See documentation for `pycce.Sequence`).

`pulses` can be provided as a list with tuples or dictionaries, each tuple or dictionary is used to initialize `Pulse` class instance.

For example, for only central spin pulses the `pulses` argument can be provided as a list of tuples, containing:

1. axis the rotation is about;
2. angle of rotation;

3. (optional) Time before the pulse. Can be as fixed, as well as varied. If varied, it should be provided as an array with the same length as `timespace`.

E.g. for Hahn-Echo the pulses can be defined as `[('x', np.pi)]` or `[('x', np.pi, timespace / 2)]`.

---

**Note:** If delay is not provided in **all** pulses, assumes even delay of CPMG sequence. If only **some** delays are provided, assumes `delay = 0` in the pulses without delay.

Then total experiment is assumed to be:

$$\text{tau} - \text{pulse} - 2\text{tau} - \text{pulse} - \dots - 2\text{tau} - \text{pulse} - \text{tau}$$

Where tau is the delay between pulses.

The sum of delays at each time point should be less or equal to the total time of the experiment at the same time point, provided in `timespace` argument.

---

**Warning:** In conventional CCE calculations, only *pi* pulses on the central spin are allowed.

In the calculations of noise autocorrelation this parameter is ignored.

Default is **None**. Overrides ```Simulator.pulses``` if provided.

- **alpha** (*int or ndarray with shape (2s+1, )*) – 0 state of the qubit in  $S_z$  basis or the index of eigenstate to be used as one.

Default is **None**. Overrides ```Simulator.alpha``` if provided.

- **beta** (*int or ndarray with shape (2s+1, )*) – 1 state of the qubit in  $S_z$  basis or the index of the eigenstate to be used as one.

Default is **None**. Overrides ```Simulator.beta``` if provided.

- **as\_delay** (*bool*) – True if time points are delay between pulses (for equispaced pulses), False if time points are total time. Ignored in gCCE if `pulses` contains the time fractions. Conventional CCE calculations do not support custom time fractions.

Default is **False**.

- **interlaced** (*bool*) – True if use hybrid CCE approach - for each cluster sample over states of the supercluster.

Default is **False**.

- **state** (*ndarray with shape (2s+1,)*) – Initial state of the central spin, used in gCCE and noise autocorrelation calculations.

Defaults to  $\frac{1}{N}(0 + 1)$  if not set.

- **bath\_state** (*array-like*) – List of bath spin states. If `len(shape) == 1`, contains  $I_z$  projections of  $I_z$  eigenstates. Otherwise, contains array of initial density matrices of bath spins.

Default is **None**. If not set, the code assumes completely random spin bath (density matrix of each nuclear spin is proportional to identity,  $\hat{I}^{\otimes N}/N$ ).

- **nbstates** (*int*) – Number of random bath states to sample over.

If provided, sampling of random states is carried and `bath_states` values are ignored.

Default is 0.

- **seed** (*int*) – Seed for random number generator, used in random bath states sampling.

Default is **None**.

- **masked** (*bool*) – True if mask numerically unstable points (with coherence > 1) in the averaging over bath states.

---

**Note:** It is up to user to check whether the possible instability is due to numerical error or unphysical assumptions of the calculations.

---

Default is **True** for coherence calculations, **False** for noise calculations.

- **parallel\_states** (*bool*) – True if to use MPI to parallelize the calculations of density matrix equally over present mpi processes for random bath state sampling calculations.

Compared to `parallel` keyword, when this argument is True each process is given a fraction of random bath states. This makes the implementation faster. Works best when the number of bath states is divisible by the number of processes, `nbstates % size == 0`.

Default is **False**.

- **fixstates** (*dict*) – If not None, shows which bath states to fix in random bath states. Each key is the index of bath spin, value - fixed  $\hat{S}_z$  projection of the mixed state of nuclear spin.

Default is **None**.

- **second\_order** (*bool*) – True if add second order perturbation theory correction to the cluster Hamiltonian in conventional CCE. Relevant only for conventional CCE calculations.

If set to True sets the qubit states as eigenstates of central spin Hamiltonian from the following procedure. If qubit states are provided as vectors in  $S_z$  basis, for each qubit state compute the fidelity of the qubit state and all eigenstates of the central spin and chose the one with fidelity higher than `level_confidence`. If such state is not found, raises an error.

**Warning:** Second order corrections are not implemented as mean field.

I.e., setting `second_order=True` and `nbstates != 0` leads to the calculation, when mean field effect is accounted only from dipolar interactions within the bath.

Default is **False**.

- **level\_confidence** (*float*) – Maximum fidelity of the qubit state to be considered eigenstate of the central spin Hamiltonian.

Default is 0.95.

- **direct** (*bool*) – True if use direct approach (requires way more memory but might be more numerically stable). False if use memory efficient approach.

Default is **False**.

- **parallel** (*bool*) – True if parallelize calculation of cluster contributions over different mpi processes.

Default is **False**.

- **Returns** – ndarray: Computed property.

## 5.4 Pulse sequences

Documentation of the Pulse and Sequence classes, used in definition of the complicated pulse sequences.

**class Pulse**(*axis=None, angle=None, delay=None, bath\_names=None, bath\_axes=None, bath\_angles=None*)  
Class containing properties of each control pulse, applied to the system.

### Parameters

- **axis** (*str*) – Axis of rotation of the central spin. Can be ‘x’, ‘y’, or ‘z’. Default is None.
- **angle** (*float or str*) – Angle of rotation of central spin. Can be provided in rad, or as a string, containing fraction of pi: ‘pi’, ‘pi/2’, ‘2\*pi’ etc. Default is None.
- **delay** (*float or ndarray*) – Delay before the pulse or array of delays with the same shape as time points. Default is None.
- **bath\_names** (*str or array-like of str*) – Name or array of names of bath spin types, impacted by the bath pulse. Default is None.
- **bath\_axes** (*str or array-like of str*) – Axis of rotation or array of axes of the bath spins. Default is None. If **bath\_names** is provided, but **bath\_axes** and **bath\_angles** are not, assumes the same axis and angle as the one of the central spin
- **bath\_angles** (*float or str or array-like*) – Angle of rotation or array of axes of rotations of the bath spins.

### Examples

```
>>> Pulse('x', 'pi')
Pulse((x, 3.14))
>>> Pulse('x', 'pi', bath_names=['13C', '14C'])
Pulse((x, 3.14), {13C: (x, 3.14), 14C: (x, 3.14)})
>>> import numpy as np
>>> Pulse('x', 'pi', delay=np.linspace(0, 1, 5), bath_names=['13C', '14C'], bath_
↪axes='x', bath_angles='pi/2')
Pulse((x, 3.14), {13C: (x, 1.57), 14C: (x, 1.57)}, t = [0.  0.25 0.5  0.75 1.  ])
```

#### axis

Axis of rotation of the central spin

**Type** str

#### angle

Angle of rotation of central spin in rad.

**Type** float

#### flip

True if the angle == pi.

**Type** bool

#### bath\_names

Array of names of bath spin types, impacted by the bath pulse.

**Type** ndarray

#### bath\_axes

Array of axes of rotation of the bath spins.



**Type** ndarray

**bath\_angles**

Array of angles of rotation of the bath spins.

**Type** ndarray

**rotation**

Matrix representation of the pulse for the given cluster. Generated by Sequence object.

**Type** ndarray

**property delay**

Delay before the pulse or array of delays with the same shape as time points.

**Type** float or ndarray

**class Sequence** (*t=None*)

List-like object, which contains the sequence of the pulses.

Each element is a Pulse instance, which can be generated from either the tuple with positional arguments or from the dictionary, or set manually.

If delay is not provided in **all** pulses in the sequence, assume equispaced pulse sequence:

t - pulse - 2t - pulse - 2t - ... - pulse - t

If only **some** delays are provided, assumes 0 delay in the pulses without delay provided.

**Examples**

```
>>> import numpy as np
>>> Sequence([('x', np.pi, 0),
>>>           {'axis': 'y', 'angle': 'pi', 'delay': np.linspace(0, 1, 3), 'bath_
↳names': '13C'},
>>>           Pulse('x', 'pi', 1)])
[Pulse((x, 3.14), t = 0), Pulse((y, 3.14), {13C: (y, 3.14)}, t = [0. 0.5 1. ]),
↳Pulse((x, 3.14), t = 1)]
```

**small\_sigma**

Dictionary with Pauli matrices of the central spin

**Type** dict

**delays**

List with delays before each pulse or None if equispaced. Generated by .generate\_pulses method.

**Type** list or None

**rotations**

List with matrix representations of the rotation from each pulse. Generated by .generate\_pulses method.

**Type** list

**append**(*item*)

S.append(value) – append value to the end of the sequence

**set\_central\_spin**(*alpha, beta*)

Set Pauli matrices of the central spin.

**Parameters**

- **alpha** (*ndarray*) – 0 state of the qubit in  $S_z$  basis.
- **beta** (*ndarray*) – 1 state of the qubit in  $S_z$  basis.

**generate\_pulses**(*dimensions=None, bath=None, vectors=None, central\_spin=True*)

Generate list of matrix representations of the rotations, induced by the sequence of the pulses.

The rotations are stored in the `.rotation` attribute of the each `Pulse` object and in `Sequence.rotations`.

#### Parameters

- **dimensions** (*ndarray with shape (N,)*) – Array of spin dimensions in the system.
- **bath** (*BathArray with shape (n,)*) – Array of bath spins in the system.
- **vectors** (*ndarray with shape (N, 3, prod(dimensions), prod(dimensions))*) – Array with vectors of spin matrices for each spin in the system.
- **central\_spin** (*bool*) – True if generate the rotations including central spin rotations. Default is True.

#### Returns

*tuple* containing:

- **list** or **None**: List with delays before each pulse or None if equispaced.
- **list**: List with matrix representations of the rotation from each pulse.

**Return type** *tuple*

**bath\_rotation**(*vectors, axis, angle*)

Generate rotation of the bath spins with given spin vectors.

#### Parameters

- **vectors** (*ndarray with shape (n, 3, x, x)*) – Array of  $n$  bath spin vectors.
- **axis** (*str*) – Axis of rotation.
- **angle** (*float*) – Angle of rotation.

**Returns** Matrix representation of the spin rotation.

**Return type** *ndarray with shape (x, x)*

## HAMILTONIAN PARAMETERS INPUT

The total Hamiltonian of the system is set as:

$$\hat{H} = \hat{H}_S + \hat{H}_{SB} + \hat{H}_B$$

with

$$\begin{aligned}\hat{H}_S &= \mathbf{SDS} + \mathbf{B}\gamma_S\mathbf{S} \\ \hat{H}_{SB} &= \sum_i \mathbf{SA}_i\mathbf{I}_i \\ \hat{H}_B &= \sum_i \mathbf{I}_i\mathbf{P}_i\mathbf{I}_i + \mathbf{B}\gamma_i\mathbf{I}_i + \sum_{i>j} \mathbf{I}_i\mathbf{J}_{ij}\mathbf{I}_j\end{aligned}$$

Where  $\hat{H}_S$  is the Hamiltonian of the free central spin,  $\hat{H}_{SB}$  denotes interactions between central spin and bath spin, and  $\hat{H}_B$  are intrinsic bath spin interactions:

- $\mathbf{D}(\mathbf{P})$  is the self interaction tensor of the central spin (bath spin). For the electron spin, corresponds to the Zero field splitting (ZFS) tensor. For nuclear spins corresponds to the quadrupole interactions tensor.
- $\gamma_i$  is the magnetic field interaction tensor of the  $i$ -spin describing the interaction of the spin and the external magnetic field.
- $\mathbf{A}$  is the interaction tensor between central and bath spins. In the case of nuclear spin bath, corresponds to the hyperfine couplings.
- $\mathbf{J}$  is the interaction tensor between bath spins.

Each of this terms can be defined within **PyCCE** framework as following.

In general, central spin properties are stored in the `Simulator` instance, bath properties are stored in the `BathArray` instance.

### 6.1 Central Spin Hamiltonian

The central spin Hamiltonian is provided as attributes of the `Simulator` object:

- $\mathbf{D}$  is set with `Simulator.set_zfs` method or during the initialization of the `Simulator` object either from observables  $D$  and  $E$  of the zero field splitting **OR** directly as tensor for the interaction `SDS` in kHz. By default is zero.

Examples:

```

>>> c = Simulator(1)
>>> print(c.zfs)
[[ 0.  0.  0.]
 [ 0. -0.  0.]
 [ 0.  0.  0.]]
>>> c.set_zfs(D=1e6)
>>> print(c.zfs)
[[-333333.333  0.  0. ]
 [  0. -333333.333  0. ]
 [  0.  0.  666666.667]]

```

- $\gamma_S$ , the tensor describing the interaction of the spin and the external magnetic field in units of gyromagnetic ratio  $\text{rad} \cdot \text{kHz} \cdot \text{G}^{-1}$ . By default is equal to the gyromagnetic ratio of the free electron spin,  $-17609 \text{ rad} \cdot \text{ms}^{-1} \cdot \text{G}^{-1}$ .

For the electron spin, it is proportional to g-tensor  $\mathbf{g}$  as:

$$\gamma_S = \mathbf{g} \frac{\mu_B}{\hbar},$$

where  $\mu_B$  is Bohr magneton.

For the nuclear central spin, it is proportional to the chemical shift tensor  $\sigma$  and gyromagnetic ratio  $\gamma$  as:

$$\gamma_S = \gamma(1 - \sigma)$$

Examples:

```

>>> c = Simulator(1)
>>> print(c.gyro)
-17608.59705

```

**Note:** While all other coupling parameters are given in the units of frequency, the gyromagnetic ratio (and therefore tensors coupling magnetic field with the spin) are conventionally given in the units of **angular** frequency and differ by  $2\pi$ .

The magnetic field is set with with `Simulator.set_magnetic_field` method or during the initialization of the `Simulator` object in G.

## 6.2 Spin-Bath Hamiltonian

The interactions between central spin and bath spins and are provided in the ['A'] namefield of the `BathArray` object in kHz.

Interaction tensors can be either:

- Directly provided by setting the values of `bath['A']` in kHz for each bath spin.
- Approximated from magnetic point dipole–dipole interactions by calling `BathArray.from_point_dipole` method. Then the tensors are computed as:

$$\mathbf{A}_j = -\gamma_S \gamma_j \frac{\hbar^2}{4\pi\mu_0} \left[ \frac{3\vec{r}_j^\top \otimes \vec{r}_j - |\vec{r}_j|^2 \mathbf{I}}{|\vec{r}_j|^5} \right]$$

Where  $\gamma_j$  is gyromagnetic ratio of  $j$  spin,  $\vec{r}_j$  is position of the bath spin, and  $\mathbf{I}$  is 3x3 identity matrix. The default option when reading the bath by `Simulator` object.

- Approximated from the spin density distribution of the central spin by calling `BathArray.from_cube` method.

Examples:

```
>>> bath = random_bath('13C', size=100, number=5, seed=1)
>>> print(bath)
(['13C', [ 1.182, 45.046, -35.584], [[0., 0., 0.], [0., 0., 0.], [0., 0., 0.]],
→ [[0., 0., 0.], [0., 0., 0.], [0., 0., 0.]])
('13C', [ 44.865, -18.817, -7.667], [[0., 0., 0.], [0., 0., 0.], [0., 0., 0.]],
→ [[0., 0., 0.], [0., 0., 0.], [0., 0., 0.]])
('13C', [ 32.77 , -9.08 , 4.959], [[0., 0., 0.], [0., 0., 0.], [0., 0., 0.]],
→ [[0., 0., 0.], [0., 0., 0.], [0., 0., 0.]])
('13C', [-47.244, 25.351, 3.814], [[0., 0., 0.], [0., 0., 0.], [0., 0., 0.]],
→ [[0., 0., 0.], [0., 0., 0.], [0., 0., 0.]])
('13C', [-17.027, 28.843, -19.681], [[0., 0., 0.], [0., 0., 0.], [0., 0., 0.]],
→ [[0., 0., 0.], [0., 0., 0.], [0., 0., 0.]])
>>> bath['A'] = 1
>>> print(bath)
(['13C', [ 1.182, 45.046, -35.584], [[1., 1., 1.], [1., 1., 1.], [1., 1., 1.]],
→ [[0., 0., 0.], [0., 0., 0.], [0., 0., 0.]])
('13C', [ 44.865, -18.817, -7.667], [[1., 1., 1.], [1., 1., 1.], [1., 1., 1.]],
→ [[0., 0., 0.], [0., 0., 0.], [0., 0., 0.]])
('13C', [ 32.77 , -9.08 , 4.959], [[1., 1., 1.], [1., 1., 1.], [1., 1., 1.]],
→ [[0., 0., 0.], [0., 0., 0.], [0., 0., 0.]])
('13C', [-47.244, 25.351, 3.814], [[1., 1., 1.], [1., 1., 1.], [1., 1., 1.]],
→ [[0., 0., 0.], [0., 0., 0.], [0., 0., 0.]])
('13C', [-17.027, 28.843, -19.681], [[1., 1., 1.], [1., 1., 1.], [1., 1., 1.]],
→ [[0., 0., 0.], [0., 0., 0.], [0., 0., 0.]])
>>> bath.from_point_dipole([0, 0, 0])
>>> print(bath)
(['13C', [ 1.182, 45.046, -35.584], [[-0.659, 0.032, -0.025], [ 0.032, 0.559, -
→ 0.963], [-0.025, -0.963, 0.1 ]], [[0., 0., 0.], [0., 0., 0.], [0., 0., 0.]])
('13C', [ 44.865, -18.817, -7.667], [[ 1.558, -1.092, -0.445], [-1.092, -0.588,
→ 0.187], [-0.445, 0.187, -0.97 ]], [[0., 0., 0.], [0., 0., 0.], [0., 0., 0.]])
('13C', [ 32.77 , -9.08 , 4.959], [[ 5.32 , -2.327, 1.271], [-2.327, -2.434, -
→ 0.352], [ 1.271, -0.352, -2.886]], [[0., 0., 0.], [0., 0., 0.], [0., 0., 0.]])
('13C', [-47.244, 25.351, 3.814], [[ 1.06 , -1. , -0.151], [-1. , -0.268,
→ 0.081], [-0.151, 0.081, -0.792]], [[0., 0., 0.], [0., 0., 0.], [0., 0., 0.]])
('13C', [-17.027, 28.843, -19.681], [[-0.903, -2.081, 1.42 ], [-2.081, 1.393, -
→ 2.405], [ 1.42 , -2.405, -0.49 ]], [[0., 0., 0.], [0., 0., 0.], [0., 0., 0.]])
```

### 6.3 Bath Hamiltonian

The self interaction tensors of the bath spins are stored in the ['Q'] namefield of the `BathArray` object. By default they are set to 0. They can be either:

- Directly provided by setting the values of `bath['Q']` in kHz for each bath spin.
- Computed from the electric field gradient (EFG) tensors at each bath spin position, using `BathArray.from_efg` method.

The gyromagnetic ratio  $\gamma_j$  of each bath spin type is stored in the `BathArray.types`.

The couplings between bath spins are assumed to follow point dipole-dipole interactions as:

$$\mathbf{P}_{ij} = -\gamma_i \gamma_j \frac{\hbar^2}{4\pi\mu_0} \left[ \frac{3\vec{r}_{ij} \otimes \vec{r}_{ij} - |\vec{r}_{ij}|^2 \mathbf{I}}{|\vec{r}_{ij}|^5} \right]$$

Where  $\gamma_i$  is gyromagnetic ratio of  $i$  tensor,  $\mathbf{I}$  is 3x3 identity matrix, and  $\vec{r}_{ij}$  is distance between two vectors.

However, user can define the interaction tensors for specific bath spin pairs stored in the `BathArray` instance. This can be achieved by:

- Calling `BathArray.add_interaction` method of the `BathArray` instance.
- Providing `InteractionsMap` instance as `imap` keyword to the `Simulator.read_bath`.

Examples:

```
>>> import numpy as np
>>> bath = random_bath('13C', size=100, number=5, seed=1)
>>> print(bath.types)
SpinDict(13C: (13C, 0.5, 6.7283))
>>> test_tensor = np.random.random((3, 3))
>>> bath.add_interaction(0, 1, (test_tensor + test_tensor.T) / 2)
>>> print(bath.imap[0, 1])
[[0.786 0.53  0.404]
 [0.53  0.821 0.366]
 [0.404 0.366 0.655]]
>>> print(bath.imap[0, 1])
[[0.786 0.53  0.404]
 [0.53  0.821 0.366]
 [0.404 0.366 0.655]]
```

## ELECTRONIC STRUCTURE OUTPUT

Each of the interfaces includes the function that should be used to read electronic structure calculations output into `BathArray` instance.

### 7.1 Quantum Espresso interface

**read\_qe**(*pwfile*, *hyperfine*=None, *efg*=None, *s*=1, *pwtype*=None, *types*=None, *isotopes*=None, *center*=None, *center\_type*=None, *rotation\_matrix*=None, *rm\_style*='col', *find\_isotopes*=True)

Function to read PW/GIPAW output from Quantum Espresso into `BathArray`.

Changes the names of the atoms to the most abundant isotopes if `find_isotopes` set to True. If that is not the desired outcome, user can define which isotopes to use using keyword `isotopes`. If `find_isotopes` is False, then keep the original names even when `isotopes` argument is provided.

#### Parameters

- **pwfile** (*str*) – Name of PW input or output file. If the file doesn't have proper extension, parameter `pw_type` should indicate the type.
- **hyperfine** (*str*) – name of the GIPAW hyperfine output.
- **efg** (*str*) – Name of the gipaw electric field tensor output.
- **s** (*float*) – Spin of the central spin. Default 1.
- **pwtype** (*str*) – Type of the `pwfile`. if not listed, will be inferred from extension of `pwfile`.
- **types** (`SpinDict` or *list of tuples*) – `SpinDict` containing `SpinTypes` of isotopes or input to make one.
- **isotopes** (*dict*) – Optional. Dictionary with entries: {"element": "isotope"}, where "element" is the name of the element in DFT output, "isotope" is the name of the isotope.
- **center** (*ndarray of shape (3,)*) – Position of (0, 0, 0) point in input coordinates.
- **center\_type** (*str*) – Type of the coordinates provided in center argument. Possible value include: 'bohr', 'angstrom', 'crystal', 'alat'. Default assumes the same as in PW file.
- **rotation\_matrix** (*ndarray of shape (3,3)*) – Rotation matrix to rotate basis. For details see `utilities.transform`.
- **rm\_style** (*str*) – Indicates how rotation matrix should be interpreted. Can take values "col" or "row". Default "col"
- **find\_isotopes** (*bool*) – If true, sets isotopes instead of names of the atoms.

**Returns** `BathArray` containing atoms with hyperfine couplings and quadrupole tensors from QE output.

**Return type** *BathArray*

## 7.2 ORCA interface

**read\_orca**(*fname*, *isotopes=None*, *types=None*, *center=None*, *find\_isotopes=True*, *rotation\_matrix=None*, *rm\_style='col'*)

Function to read ORCA output containing the hyperfines couplings and EFG tensors.

if *find\_isotopes* is set to True changes the names of the atoms to the most abundant isotopes. If that is not the desired outcome, user can define which isotopes to use using keyword *isotopes*.

### Parameters

- **fname** (*str*) – file name of the ORCA output.
- **isotopes** (*dict*) – Optional. Dictionary with entries:

```
{"element" : "isotope"}
```

where “element” is the name of the element in DFT output, “isotope” is the name of the isotope.

- **types** (*SpinDict* or *list of tuples*) – *SpinDict* containing *SpinTypes* of isotopes or input to make one.
- **center** (*ndarray of shape (3,)*) – position of (0, 0, 0) point in the DFT coordinates.
- **rotation\_matrix** (*ndarray of shape (3,3)*) – Rotation matrix to rotate basis. For details see *utilities.transform*.
- **rm\_style** (*str*) – Indicates how rotation matrix should be interpreted. Can take values “col” or “row”. Default “col”
- **find\_isotopes** (*bool*) – If true, sets isotopes instead of names of the atoms.

**Returns** Array of bath spins with hyperfine couplings and quadrupole tensors from Orca output.

**Return type** *BathArray*



## CCE CALCULATORS

Documentation for the calculator objects called by Simulator object.

### 8.1 Base class

```
class RunObject(timespace, clusters, bath, magnetic_field, alpha, beta, state, spin, zfs, gyro, nbstates=None,  
               bath_state=None, seed=None, masked=True, fixstates=None, projected_bath_state=None,  
               parallel=False, direct=False, parallel_states=False, **kwargs)
```

Abstract class of the CCE simulation runner.

Implements cluster correlation expansion, interlaced averaging, and sampling over random bath states. Requires definition of the following methods, from which the kernel will be automatically created:

- `.generate_hamiltonian(self)` method which, using the attributes of the `self` object, computes cluster hamiltonian stored in `self.cluster_hamiltonian`.
- `.compute_result(self)` method which, using the attributes of the `self`, computes the resulting quantity for the given cluster.

Alternatively, user can define the kernel manually. Then the following methods have to be overridden:

- `.kernel(self, cluster, *args, **kwargs)` method which takes indexes of the bath spins in the given cluster as a first positional argument. This method is required for usual CCE runs.
- `.interlaced_kernel(self, cluster, supercluster, *args, **kwargs)` method which takes indexes of the bath spins in the given cluster as a first positional argument, indexes of the supercluster as a second positional argument. This method is required for interlaced CCE runs.

#### Parameters

- **timespace** (*ndarray with shape (t, )*) – Time delay values at which to compute propagators.
- **clusters** (*dict*) – Clusters included in different CCE orders of structure `{int order: ndarray([[i, j], [i, j]])}`.
- **bath** (*BathArray with shape (n,)*) – Array of  $n$  bath spins.
- **magnetic\_field** (*ndarray*) – Magnetic field of type `magnetic_field = np.array([Bx, By, Bz])`.
- **alpha** (*int or ndarray with shape (2s+1, )*) – 0 state of the qubit in  $S_z$  basis or the index of eigenstate to be used as one.
- **beta** (*int or ndarray with shape (2s+1, )*) – 1 state of the qubit in  $S_z$  basis or the index of the eigenstate to be used as one.

- **state** (*ndarray with shape (2s+1, )*) – Initial state of the central spin, used in gCCE and noise autocorrelation calculations. Defaults to  $\frac{1}{N}(0 + 1)$  if not set **OR** if alpha and beta are provided as indexes.
- **spin** (*float*) – Value of the central spin.
- **zfs** (*ndarray with shape (3, 3)*) – Zero Field Splitting tensor of the central spin.
- **gyro** (*float or ndarray with shape (3, 3)*) – Gyromagnetic ratio of the central spin

**OR**

tensor corresponding to interaction between magnetic field and central spin.

- **nbstates** (*int*) – Number of random bath states to sample over in bath state sampling runs.
- **bath\_state** (*ndarray*) – Array of bath states in any accepted format.
- **seed** (*int*) – Seed for the random number generator in bath states sampling.
- **masked** (*bool*) – True if mask numerically unstable points (with `result > result[0]`) in the sampling over bath states False if not. Default True.
- **fixstates** (*dict*) – If provided, contains indexes of bath spins with fixed pure state for sampling runs and interlaced runs.

Each key is the index of bath spin, value - fixed  $\hat{I}_z$  projection of the **pure**  $\hat{I}_z$  eigenstate of bath spin.

- **projected\_bath\_state** (*ndarray with shape (n, )*) – Array with z-projections of the bath spins states. Overridden in runs with random bath state sampling.
- **parallel** (*bool*) – True if parallelize calculation of cluster contributions over different mpi processes. Default False.
- **direct** (*bool*) – True if use direct approach in run (requires way more memory but might be more numerically stable). False if use memory efficient approach. Default False.
- **parallel\_states** (*bool*) – True if use MPI to parallelize the calculations of density matrix for each random bath state.
- **\*\*kwargs** – Additional keyword arguments to be set as the attributes of the given object.

**result\_operator** (*b, /*)

Operator which will combine the result of expansion,.

Default: `operator.imul`.

**contribution\_operator** (*b, /*)

Operator which will combine multiple contributions of the same cluster in the optimized approach.

Default: `operator.ipow`.

**removal\_operator** (*b, /*)

Operator which will remove subcluster contribution from the given cluster contribution. First argument cluster contribution, second - subcluster contribution.

Default: `operator.itruediv`.

**addition\_operator** (*axis=None, dtype=None, out=None, keepdims=<no value>, initial=<no value>, where=<no value>*)

Group operation which will combine contributions from the different clusters into one contribution in the direct approach.

Default: `numpy.prod`.

**nbstates**

Number of random bath states to sample over in bath state sampling runs.

**Type** int

**timespace**

Time points at which result will be computed.

**Type** ndarray with shape (t, )

**clusters**

Clusters included in different CCE orders of structure `{int order: ndarray([[i,j],[i,j]])}`.

**Type** dict

**bath**

Array of  $n$  bath spins.

**Type** BathArray with shape (n,)

**spin**

Value of the central spin

**Type** float

**zfs**

Zero Field Splitting tensor of the central spin.

**Type** ndarray with shape (3, 3)

**gyro**

float or ndarray with shape (3, 3): Gyromagnetic ratio of the central spin

**OR**

tensor corresponding to interaction between magnetic field and central spin.

**bath\_state**

Array of bath states in any accepted format.

**Type** ndarray

**projected\_bath\_state**

Array with z-projections of the bath spins states. Overridden in runs with random bath state sampling.

**Type** ndarray with shape (n,)

**magnetic\_field**

Magnetic field of type `magnetic_field = np.array([Bx, By, Bz])`.

**Type** ndarray

**initial\_alpha**

0 state of the qubit in  $S_z$  basis or the index of eigenstate to be used as one.

**Type** ndarray

**initial\_beta**

1 state of the qubit in  $S_z$  basis or the index of eigenstate to be used as one.

**Type** ndarray

**alpha**

0 state of the qubit in  $S_z$  basis. If initially provided as index, generated as a state during the run.

**Type** ndarray

**beta**

1 state of the qubit in  $S_z$  basis. If initially provided as index, generated as a state during the run.

**Type** ndarray

**state**

Initial state of the central spin, used in gCCE and noise autocorrelation calculations.

Defaults to  $\frac{1}{N}(0 + 1)$  if not set **OR** if alpha and beta are provided as indexes.

**Type** ndarray

**parallel**

True if parallelize calculation of cluster contributions over different mpi processes. Default False.

**Type** bool

**parallel\_states**

True if use MPI to parallelize the calculations of density matrix for each random bath state.

**Type** bool

**direct**

True if use direct approach in run (requires way more memory but might be more numerically stable). False if use memory efficient approach. Default False.

**Type** bool

**seed**

Seed for the random number generator in bath states sampling.

**Type** int

**masked**

True if mask numerically unstable points (with result > result[0]) in the sampling over bath states False if not. Default True.

**Type** bool

**fixstates**

If provided, contains indexes of bath spins with fixed pure state for sampling runs and interlaced runs.

Each key is the index of bath spin, value - fixed  $\hat{I}_z$  projection of the **pure**  $\hat{I}_z$  eigenstate of bath spin.

**Type** dict

**hamiltonian**

central spin hamiltonian.

**Type** ndarray

**energies**

Eigen energies of the central spin hamiltonian.

**Type** ndarray

**eigenvectors**

Eigen states of the central spin hamiltonian.

**Type** ndarray

**cluster**

Array of the bath spins inside the given cluster.

**Type** *BathArray*

**states**

Array of the states of bath spins inside the given cluster.

**Type** ndarray

**others**

Array of the bath spins outside the given cluster.

**Type** *BathArray*

**other\_states**

Array of the z-projections of the states of bath spins outside the given cluster.

**Type** ndarray

**cluster\_hamiltonian**

Full hamiltonian of the given cluster. In conventional CCE, tuple with two projected hamiltonians.

**Type** *Hamiltonian* or tuple

**result**

Result of the calculation.

**Type** ndarray

**preprocess()**

Method which will be called before cluster-expanded run.

**postprocess()**

Method which will be called after cluster-expanded run.

**kernel**(*cluster*, \**args*, \*\**kwargs*)

Central kernel that will be called in the cluster-expanded calculations

**Parameters**

- **cluster** (*ndarray*) – Indexes of the bath spins in the given cluster.
- **\*args** – Positional arguments of the kernel.
- **\*\*kwargs** – Keyword arguments of the kernel.

**Returns** Results of the calculations.

**Return type** ndarray

**run**(\**args*, \*\**kwargs*)

Method that runs cluster-expanded single calculation.

**Parameters**

- **\*args** – Positional arguments of the kernel.
- **\*\*kwargs** – Keyword arguments of the kernel.

**Returns** Results of the calculations.

**Return type** ndarray

**sampling\_run**(\**args*, \*\**kwargs*)

Method that runs bath sampling calculations.

**Parameters**

- **\*args** – Positional arguments of the kernel.
- **\*\*kwargs** – Keyword arguments of the kernel.

**Returns** Results of the calculations.

**Return type** ndarray

**interlaced\_kernel**(*cluster, supercluster, \*args, \*\*kwargs*)

Central kernel that will be called in the cluster-expanded calculations with interlaced averaging of bath spin states.

**Parameters**

- **cluster** (ndarray) – Indexes of the bath spins in the given cluster.
- **supercluster** (ndarray) – Indexes of the bath spins in the supercluster of the given cluster. Supercluster is the union of all clusters in `.clusters` attribute, for which given cluster is a subset.
- **\*args** – Positional arguments of the kernel.
- **\*\*kwargs** – Keyword arguments of the kernel.

**Returns** Results of the calculations.

**Return type** ndarray

**interlaced\_run**(\*args, \*\*kwargs)

Method that runs cluster-expanded single calculation with interlaced averaging of bath spin states.

**Parameters**

- **\*args** – Positional arguments of the interlaced kernel.
- **\*\*kwargs** – Keyword arguments of the interlaced kernel.

**Returns** Results of the calculations.

**Return type** ndarray

**sampling\_interlaced\_run**(\*args, \*\*kwargs)

Method that runs bath sampling calculations with interlaced averaging of bath spin states.

**Parameters**

- **\*args** – Positional arguments of the interlaced kernel.
- **\*\*kwargs** – Keyword arguments of the interlaced kernel.

**Returns** Results of the calculations.

**Return type** ndarray

**classmethod from\_simulator**(*sim, \*\*kwargs*)

Class method to generate RunObject from the properties of Simulator object.

**Parameters**

- **sim** (Simulator) – Object, whose properties will be used to initialize RunObject instance.
- **\*\*kwargs** – Additional keyword arguments that will replace ones, recovered from the Simulator object.

**Returns** New instance of RunObject class.

**Return type** RunObject

**generate\_supercluster\_states**(*self, supercluster*)

Helper function to generate all possible pure states of the given supercluster.

**Parameters**

- **self** (`RunObject`) – Instance of the `RunObject` class, used in the calculation.
- **supercluster** (`ndarray with shape (n, )`) – Indexes of the bath spins in the supercluster.

**Yields** `ndarray with shape (n, )` – Pure state of the given supercluster.

## 8.2 Conventional CCE

**propagators**(`timespace, H0, H1, pulses, as_delay=False`)

Function to compute propagators U0 and U1 in conventional CCE.

**Parameters**

- **timespace** (`ndarray with shape (t, )`) – Time delay values at which to compute propagators.
- **H0** (`ndarray with shape (n, n)`) – Hamiltonian projected on alpha qubit state.
- **H1** (`ndarray with shape (n, n)`) – Hamiltonian projected on beta qubit state.
- **pulses** (`int or Sequence`) – Sequence of pulses.
- **as\_delay** (`bool`) – True if time points are delay between pulses. False if time points are total time.

**Returns**

*tuple* containing:

- **ndarray with shape (t, n, n)**: Matrix representation of the propagator conditioned on the alpha qubit state for each time point.
- **ndarray with shape (t, n, n)**: Matrix representation of the propagator conditioned on the beta qubit state for each time point.

**Return type** `tuple`

**compute\_coherence**(`H0, H1, timespace, N, as_delay=False, states=None`)

Function to compute cluster coherence function in conventional CCE.

**Parameters**

- **H0** (`ndarray`) – Hamiltonian projected on alpha qubit state.
- **H1** (`ndarray`) – Hamiltonian projected on beta qubit state.
- **timespace** (`ndarray`) – Time points at which to compute coherence function.
- **N** (`int`) – Number of pulses in CPMG.
- **as\_delay** (`bool`) – True if time points are delay between pulses, False if time points are total time.
- **states** (`ndarray`) – `ndarray` of bath states in any accepted format.

**Returns** Coherence function of the central spin.

**Return type** `ndarray`

**class CCE**(\*args, pulses=0, as\_delay=False, second\_order=False, level\_confidence=0.95, \*\*kwargs)  
Class for running conventional CCE simulations.

---

**Note:** Subclass of the RunObject abstract class.

---

### Parameters

- **\*args** – Positional arguments of the RunObject.
- **pulses** (*int* or *Sequence*) – number of pulses in CPMG sequence or instance of Sequence object. For now, only CPMG sequences are supported in conventional CCE simulations.
- **as\_delay** (*bool*) – True if time points are delay between pulses, False if time points are total time.
- **second\_order** (*bool*) – True if add second order perturbation theory correction to the cluster Hamiltonian. If set to True sets the qubit states as eigenstates of central spin Hamiltonian from the following procedure. If qubit states are provided as vectors in  $S_z$  basis, for each qubit state compute the fidelity of the qubit state and all eigenstates of the central spin and chose the one with fidelity higher than `level_confidence`. If such state is not found, raises an error.
- **level\_confidence** (*float*) – Maximum fidelity of the qubit state to be considered eigenstate of the central spin Hamiltonian. Default 0.95.
- **\*\*kwargs** – Keyword arguments of the RunObject.

### **initial\_pulses**

Input pulses

**Type** *int* or *Sequence*

### **pulses**

If input Sequence contains only pi pulses at even delay, stores number of pulses. Otherwise stores full Sequence.

**Type** *int* or *Sequence*

### **as\_delay**

True if time points are delay between pulses, False if time points are total time.

**Type** *bool*

### **second\_order**

True if add second order perturbation theory correction to the cluster hamiltonian.

**Type** *bool*

### **level\_confidence**

Maximum fidelity of the qubit state to be considered eigenstate of the central spin hamiltonian.

**Type** *float*

### **energy\_alpha**

Eigen energy of the alpha state in the central spin Hamiltonian.

**Type** *float*

### **energy\_beta**

Eigen energy of the beta state in the central spin Hamiltonian.



**Type** float

**energies**

All eigen energies of the central spin Hamiltonian.

**Type** ndarray with shape (2s+1, )

**projections\_alpha\_all**

Array of vectors with spin operator matrix elements of type  $[0\hat{S}_x i, 0\hat{S}_y i, 0\hat{S}_z i]$ , where 0 is the alpha qubit state,  $i$  are all eigenstates of the central spin hamiltonian.

**Type** ndarray with shape (2s+1, 3)

**projections\_beta\_all**

Array of vectors with spin operator matrix elements of type  $[1\hat{S}_x i, 1\hat{S}_y i, 1\hat{S}_z i]$ , where 1 is the beta qubit state,  $i$  are all eigenstates of the central spin hamiltonian.

**Type** ndarray with shape (2s+1, 3)

**projections\_alpha**

Vector with spin operator matrix elements of type  $[0\hat{S}_x 0, 0\hat{S}_y 0, 0\hat{S}_z 0]$ , where 0 is the alpha qubit state

**Type** ndarray with shape (3,)

**projections\_beta**

Vectors with spin operator matrix elements of type  $[1\hat{S}_x 1, 1\hat{S}_y 1, 1\hat{S}_z 1]$ , where 1 is the beta qubit state.

**Type** ndarray with shape (3,)

**use\_pulses**

True if use full Sequence. False if use only number of pulses.

**Type** bool

**preprocess()**

Method which will be called before cluster-expanded run.

**postprocess()**

Method which will be called after cluster-expanded run.

**generate\_hamiltonian()**

Using the attributes of the `self` object, compute the two projected cluster hamiltonians.

**Returns**

Tuple containing:

- **Hamiltonian**: Cluster hamiltonian when qubit in the alpha state.
- **Hamiltonian**: Cluster hamiltonian when qubit in the alpha state.

**Return type** tuple

**compute\_result()**

Using the attributes of the `self` object, compute the coherence function as overlap in the bath evolution.

**Returns** Computed coherence.

**Return type** ndarray

## 8.3 Generalized CCE

**propagator**(*timespace, hamiltonian, pulses=None, as\_delay=False*)

Function to compute time propagator U.

### Parameters

- **timespace** (*ndarray with shape (t, )*) – Time points at which to compute propagators.
- **hamiltonian** (*ndarray with shape (n, n)*) – Matrix representation of the cluster hamiltonian.
- **pulses** ([Sequence](#)) – Pulses as an instance of Sequence class with rotations already generated. Default is None.
- **as\_delay** (*bool*) – True if time points are delay between pulses, False if time points are total time. Default is False.

**Returns** Array of propagators, evaluated at each time point in timespace.

**Return type** ndarray with shape (t, n, n)

**compute\_dm**(*dm0, H, timespace, pulse\_sequence=None, as\_delay=False, states=None*)

Function to compute density matrix of the central spin, given Hamiltonian H.

### Parameters

- **dm0** (*ndarray*) – Initial density matrix of central spin.
- **H** (*ndarray*) – Cluster Hamiltonian.
- **timespace** (*ndarray*) – Time points at which to compute density matrix.
- **pulse\_sequence** ([Sequence](#)) – Sequence of pulses.
- **as\_delay** (*bool*) – True if time points are delay between pulses, False if time points are total time.
- **states** (*ndarray*) – ndarray of bath states in any accepted format.

**Returns** Array of density matrices evaluated at all time points in timespace.

**Return type** ndarray

**full\_dm**(*dm0, H, timespace, pulse\_sequence=None, as\_delay=False*)

A function to compute density matrix of the cluster, using hamiltonian H from the initial density matrix of the cluster.

### Parameters

- **dm0** (*ndarray*) – Initial density matrix of the cluster
- **H** (*ndarray*) – Cluster Hamiltonian
- **timespace** (*ndarray*) – Time points at which to compute coherence function.
- **pulse\_sequence** ([Sequence](#)) – Sequence of pulses.
- **as\_delay** (*bool*) – True if time points are delay between pulses, False if time points are total time. Ignored if delays are provided in the pulse\_sequence.

**Returns** Array of density matrices of the cluster, evaluated at the time points from timespace.

**Return type** ndarray

**generate\_dm0**(*dm0, dimensions, states=None*)

A function to generate initial density matrix or statevector of the cluster. :param dm0: Initial density matrix of the central spin. :type dm0: ndarray :param dimensions: ndarray of bath spin dimensions. Last entry - electron spin dimensions. :type dimensions: ndarray :param states: ndarray of bath states in any accepted format. :type states: ndarray

**Returns** Initial density matrix of the cluster **OR** statevector if dm0 is vector and states are provided as list of pure states.

**Return type** ndarray

**generate\_pure\_initial\_state**(*state0, dimensions, states*)

A function to generate initial state vector of the cluster with central spin.

**Parameters**

- **state0** (*ndarray*) – Initial state of the central spin.
- **dimensions** (*ndarray*) – ndarray of bath spin dimensions. Last entry - electron spin dimensions.
- **states** (*ndarray*) – ndarray of bath states in any accepted format.

**Returns** Initial state vector of the cluster.

**Return type** ndarray

**gen\_density\_matrix**(*states=None, dimensions=None*)

Generate density matrix from the ndarray of states.

**Parameters**

- **states** (*ndarray*) – Array of bath spin states. If None, assume completely random state. Can have the following forms:
  - array of the  $\hat{I}_z$  projections for each spin. Assumes that each bath spin is in the pure eigenstate of  $\hat{I}_z$ .
  - array of the diagonal elements of the density matrix for each spin. Assumes mixed state and the density matrix for each bath spin is diagonal in  $\hat{I}_z$  basis.
  - array of the density matrices of the bath spins.
- **dimensions** (*ndarray*) – array of bath spin dimensions. Last entry - electron spin dimensions.

**Returns** Density matrix of the system.

**Return type** ndarray

**class gCCE**(*\*args, as\_delay=False, pulses=None, \*\*kwargs*)

Class for running generalized CCE simulations.

---

**Note:** Subclass of the RunObject abstract class.

---

**Parameters**

- **\*args** – Positional arguments of the RunObject.
- **pulses** (*Sequence*) – Sequence object, containing series of pulses, applied to the system.
- **as\_delay** (*bool*) – True if time points are delay between pulses, False if time points are total time.

- **\*\*kwargs** – Keyword arguments of the RunObject.

**pulses**

Sequence object, containing series of pulses, applied to the system.

**Type** *Sequence*

**as\_delay**

True if time points are delay between pulses, False if time points are total time.

**Type** bool

**dm0**

Initial density matrix of the central spin.

**Type** ndarray with shape (2s+1, 2s+1)

**normalization**

Coherence at time 0.

**Type** float

**zero\_cluster**

Coherence computed for the isolated central spin.

**Type** ndarray with shape (n,)

**preprocess()**

Method which will be called before cluster-expanded run.

**postprocess()**

Method which will be called after cluster-expanded run.

**generate\_hamiltonian()**

Using the attributes of the `self` object, compute the cluster hamiltonian including the central spin.

**Returns** Cluster hamiltonian.

**Return type** *Hamiltonian*

**compute\_result()**

Using the attributes of the `self` object, compute the coherence function of the central spin.

**Returns** Computed coherence.

**Return type** ndarray

## 8.4 Noise Autocorrelation

**correlation\_it\_j0**(*operator\_i, operator\_j, dm0\_expanded, U*)

Function to compute correlation function of the operator *i* at time *t* and operator *j* at time 0

**Parameters**

- **operator\_i** (*ndarray with shape (n, n)*) – Matrix representation of operator *i*.
- **operator\_j** (*ndarray with shape (n, n)*) – Matrix representation of operator *j*.
- **dm0\_expanded** (*ndarray with shape (n, n)*) – Initial density matrix of the cluster.
- **U** (*ndarray with shape (t, n, n)*) – Time evolution propagator, evaluated over *t* time points.

**Returns** Autocorrelation of the z-noise at each time point.

**Return type** ndarray with shape (t,)

**compute\_correlations**(*nspin*, *dm0\_expanded*, *U*, *central\_spin=None*)

Function to compute correlations for the given cluster, given time propagator U.

**Parameters**

- **nspin** (*BathArray*) – BathArray of the given cluster of bath spins.
- **dm0\_expanded** (*ndarray with shape (n, n)*) – Initial density matrix of the cluster.
- **U** (*ndarray with shape (t, n, n)*) – Time evolution propagator, evaluated over t time points.
- **central\_spin** (*float*) – Value of the central spin.

**Returns** correlation of the Overhauser field, induced by the given cluster at each time point.

**Return type** ndarray with shape (t,)

**class gCCENoise**(\*args, \*\*kwargs)

Class for running generalized CCE simulations of the noise autocorrelation function.

---

**Note:** Subclass of the RunObject abstract class.

---

**Parameters**

- **\*args** – Positional arguments of the RunObject.
- **\*\*kwargs** – Keyword arguments of the RunObject.

**result\_operator**(*b*, /)

Overridden operator which will combine the result of expansion: `operator.iadd`.

**contribution\_operator**(*b*, /)

Overridden operator which will combine multiple contributions of the same cluster in the optimized approach: `operator.imul`.

**removal\_operator**(*b*, /)

Overridden operator which remove subcluster contribution from the given cluster contribution: `operator.isub`.

**addition\_operator**(*axis=None*, *dtype=None*, *out=None*, *keepdims=<no value>*, *initial=<no value>*, *where=<no value>*)

Overridden group operation which will combine contributions from the different clusters into one contribution in the direct approach: `numpy.sum`.

**preprocess**()

Method which will be called before cluster-expanded run.

**postprocess**()

Method which will be called after cluster-expanded run.

**generate\_hamiltonian**()

Using the attributes of the `self` object, compute the cluster hamiltonian including the central spin.

**Returns** Cluster hamiltonian.

**Return type** *Hamiltonian*

**compute\_result()**

Using the attributes of the `self` object, compute autocorrelation function of the noise from bath spins in the given cluster.

**Returns** Computed autocorrelation function.

**Return type** ndarray

**class CCENoise(\*args, \*\*kwargs)**

Class for running conventional CCE simulations of the noise autocorrelation function.

---

**Note:** Subclass of the `RunObject` abstract class.

---

**Warning:** In general, for calculations of the autocorrelation function, better results are achieved with generalized CCE, which accounts for the evolution of the entangled state of the central spin.

Second order couplings between nuclear spins are not implemented.

**Parameters**

- **\*args** – Positional arguments of the `RunObject`.
- **\*\*kwargs** – Keyword arguments of the `RunObject`.

**result\_operator(b, /)**

Overridden operator which will combine the result of expansion: `operator.iadd`.

**contribution\_operator(b, /)**

Overridden operator which will combine multiple contributions of the same cluster in the optimized approach: `operator.imul`.

**removal\_operator(b, /)**

Overridden operator which remove subcluster contribution from the given cluster contribution: `operator.isub`.

**addition\_operator(axis=None, dtype=None, out=None, keepdims=<no value>, initial=<no value>, where=<no value>)**

Overridden group operation which will combine contributions from the different clusters into one contribution in the direct approach: `numpy.sum`.

**preprocess()**

Method which will be called before cluster-expanded run.

**postprocess()**

Method which will be called after cluster-expanded run.

**generate\_hamiltonian()**

Using the attributes of the `self` object, compute the projected cluster hamiltonian, averaged for two qubit states.

**Returns** Cluster hamiltonian.

**Return type** *Hamiltonian*

**compute\_result()**

Using the attributes of the `self` object, compute autocorrelation function of the noise from bath spins in the given cluster.

**Returns** Computed autocorrelation function.

**Return type** ndarray

## 8.5 Cluster-correlation Expansion Decorators

The way we find cluster in the code.

**generate\_clusters**(*bath, r\_dipole, order, r\_inner=0, ignore=None, strong=False, nclusters=None*)

Generate clusters for the bath spins.

### Parameters

- **bath** (*BathArray*) – Array of bath spins.
- **r\_dipole** (*float*) – Maximum connectivity distance.
- **order** (*int*) – Maximum size of the clusters to find.
- **r\_inner** (*float*) – Minimum connectivity distance.
- **ignore** (*list or str, optional*) – If not None, includes the names of bath spins which are ignored in the cluster generation.
- **strong** (*bool*) – Whether to find only completely interconnected clusters (default False).
- **nclusters** (*dict*) – Dictionary which contain maximum number of clusters of the given size. Has the form `n_clusters = {order: number}`, where `order` is the size of the cluster, `number` is the maximum number of clusters with this size.

If provided, sorts the clusters by the strength of cluster interaction, equal to the lowest pairwise interaction in the cluster. Then the strongest number of clusters is taken.

**Returns** Dictionary with keys corresponding to size of the cluster, and value corresponds to ndarray of shape (matrix, N). Here matrix is the number of clusters of given size, N is the size of the cluster. Each row contains indexes of the bath spins included in the given cluster.

**Return type** dict

**make\_graph**(*bath, r\_dipole, r\_inner=0, ignore=None, max\_size=5000*)

Make a connectivity matrix for bath spins.

### Parameters

- **bath** (*BathArray*) – Array of bath spins.
- **r\_dipole** (*float*) – Maximum connectivity distance.
- **r\_inner** (*float*) – Minimum connectivity distance.
- **ignore** (*list or str, optional*) – If not None, includes the names of bath spins which are ignored in the cluster generation.
- **max\_size** (*int*) – Maximum size of the bath before less optimal (but less memory intensive) approach is used.

**Returns** Connectivity matrix.

**Return type** crs\_matrix

**connected\_components**(*csgraph, directed=False, connection='weak', return\_labels=True*)

Find connected components using `scipy.sparse.csgraph`. See documentation of `scipy.sparse.csgraph.connected_components`

**find\_subclusters**(*maximum\_order, graph, labels, n\_components, strong=False*)

Find subclusters from connectivity matrix.

**Parameters**

- **maximum\_order** (*int*) – Maximum size of the clusters to find.
- **graph** (*csr\_matrix*) – Connectivity matrix.
- **labels** (*ndarray with shape (n,)*) – Array of labels of the connected components.
- **n\_components** (*int*) – The number of connected components *n*.
- **strong** (*bool*) – Whether to find only completely interconnected clusters (default *False*).

**Returns** Dictionary with keys corresponding to size of the cluster, and value corresponds to ndarray of shape (matrix, N). Here matrix is the number of clusters of given size, N is the size of the cluster. Each row contains indexes of the bath spins included in the given cluster.

**Return type** dict

**combine\_clusters**(*cs1, cs2*)

Combine two dictionaries with clusters.

**Parameters**

- **cs1** (*dict*) – First cluster dictionary with keys corresponding to size of the cluster, and value corresponds to ndarray of shape (matrix, N).
- **cs2** (*dict*) – Second cluster dictionary with the same structure.

**Returns** Combined dictionary with unique clusters from both dictionaries.

**Return type** dict

**expand\_clusters**(*sc*)

Expand dict so each new cluster will include all possible additions of one more bath spin. This increases maximum size of the cluster by one.

**Parameters** **sc** (*dict*) – Initial clusters dictionary.

**Returns** Dictionary with expanded clusters.

**Return type** dict

**find\_valid\_subclusters**(*graph, maximum\_order, nclusters=None, bath=None, strong=False*)

Find subclusters from connectivity matrix.

**Parameters**

- **maximum\_order** (*int*) – Maximum size of the clusters to find.
- **graph** (*csr\_matrix*) – Connectivity matrix.
- **nclusters** (*dict*) – Dictionary which contain maximum number of clusters of the given size.
- **bath** (*BathArray*) – Array of bath spins.
- **strong** (*bool*) – Whether to find only completely interconnected clusters (default *False*).

**Returns** Dictionary with keys corresponding to size of the cluster, and value corresponds to ndarray of shape (matrix, N). Here matrix is the number of clusters of given size, N is the size of the cluster. Each row contains indexes of the bath spins included in the given cluster.

**Return type** dict

General decorators that are used to expand kernel of the `RunObject` class or subclasses to the whole bath *via* CCE.

This module contains information about the way the cluster expansion is implemented in the package.



**cluster\_expansion\_decorator**(*\_func=None*, \*, *result\_operator=<built-in function imul>*,  
*contribution\_operator=<built-in function ipow>*, *removal\_operator=<built-in  
function itruediv>*, *addition\_operator=<function prod>*)

Decorator for creating cluster correlation expansion of the method of RunObject class.

#### Parameters

- **\_func** (*func*) – Function to expand.
- **result\_operator** (*func*) – Operator which will combine the result of expansion (default: operator.imul).
- **contribution\_operator** (*func*) – Operator which will combine multiple contributions of the same cluster (default: operator.ipow) in the optimized approach.
- **removal\_operator** – Operator which will remove subcluster contribution from the given cluster contribution. First argument cluster contribution, second - subcluster contribution (default: operator.itruediv).
- **addition\_operator** (*func*) – Group operation which will combine contributions from the different clusters into one contribution (default: np.prod).

**Returns** Expanded function.

**Return type** func

**optimized\_approach**(*function, self, \*arg, result\_operator=<built-in function imul>*,  
*contribution\_operator=<built-in function ipow>*, *\*\*kwarg*)

Optimized approach to compute cluster correlation expansion.

#### Parameters

- **function** (*func*) – Function to expand.
- **self** (*RunObject*) – Object whose method is expanded.
- **\*arg** – list of positional arguments of the expanded function.
- **result\_operator** (*func*) – Operator which will combine the result of expansion (default: operator.imul).
- **contribution\_operator** (*func*) – Operator which will combine multiple contributions of the same cluster (default: operator.ipow).
- **\*\*kwarg** – Dictionary containing all keyword arguments of the expanded function.

**Returns** Expanded function.

**Return type** func

**direct\_approach**(*function, self, \*arg, result\_operator=<built-in function imul>*, *removal\_operator=<built-in  
function itruediv>*, *addition\_operator=<function prod>*, *\*\*kwarg*)

Direct approach to compute cluster correlation expansion.

#### Parameters

- **function** (*func*) – Function to expand.
- **self** (*RunObject*) – Object whose method is expanded.
- **result\_operator** (*func*) – Operator which will combine the result of expansion (default: operator.imul).

- **removal\_operator** (*func*) – Operator which will remove subcluster contribution from the given cluster contribution. First argument cluster contribution, second - subcluster contribution (default: `operator.itruediv`).
- **addition\_operator** (*func*) – Group operation which will combine contributions from the different clusters into one contribution (default: `np.prod`).
- **\*\*kwarg** – Dictionary containing all keyword arguments of the expanded function.

**Returns** Expanded method.

**Return type** `func`

**interlaced\_decorator**(*\_func=None*, \*, *result\_operator=<built-in function imul>*,  
*contribution\_operator=<built-in function ipow>*)

Decorator for creating interlaced cluster correlation expansion of the method of `RunObject` class.

**Parameters**

- **\_func** (*func*) – Function to expand.
- **result\_operator** (*func*) – Operator which will combine the result of expansion (default: `operator.imul`).
- **contribution\_operator** (*func*) – Operator which will combine multiple contributions of the same cluster (default: `operator.ipow`) in the optimized approach.

**Returns** Expanded method.

**Return type** `func`

Decorators that are used to perform bath state sampling over the kernel of `RunObject`.

**generate\_bath\_state**(*bath*, *nbstates*, *seed=None*, *fixstates=None*, *parallel=False*)

Generator of the random *pure*  $\hat{I}_z$  bath eigenstates.

**Parameters**

- **bath** (`BathArray`) – Array of bath spins.
- **nbstates** (*int*) – Number of random bath states to generate.
- **seed** (*int*) – Optional. Seed for RNG.
- **fixstates** (*dict*) – Optional. dict of which bath states to fix. Each key is the index of bath spin, value - fixed  $\hat{I}_z$  projection of the mixed state of nuclear spin.

**Yields** *ndarray* – Array of shape = `len(bath)` containing z-projections of the bath spins states.

**monte\_carlo\_method\_decorator**(*func*)

Decorator to sample over random bath states given function.

## HAMILTONIAN FUNCTIONS

### 9.1 Base Class

**class** `Hamiltonian`(*dimensions*, *vectors=None*)

Class containing properties of the Hamiltonian.

Essentially wrapper for ndarray with additional attributes of `dimensions` and `spins`.

Usual methods (e.g. `__getitem__` or `__setitem__`) access the `data` attribute.

---

**Note:** Algebraic operations with `Hamiltonian` will return ndarray instance.

---

**Parameters** `dimensions` (*array-like*) – array of the dimensions for each spin in the Hilbert space of the Hamiltonian.

**dimensions**

array of the dimensions for each spin in the Hilbert space of the Hamiltonian.

**Type** ndarray

**spins**

array of the spins, spanning the Hilbert space of the Hamiltonian.

**Type** ndarray

**vectors**

list with spin vectors of form `[[Ix, Iy, Iz], [Ix, Iy, Iz], ...]`.

**Type** list

**data**

matrix representation of the Hamiltonian.

**Type** ndarray

## 9.2 Total Hamiltonian

**hamiltonian\_wrapper**(*func=None, \*, projected=False*)

Wrapper with the general structure for the cluster Hamiltonian. Adds several additional arguments to the wrapped function.

**Additional parameters:**

- **central\_spin** (*float*) – value of the central spin.

**Parameters**

- **\_func** (*func*) – Function to be wrapped.
- **projected** (*bool*) – True if return two projected Hamiltonians, False if remove single not projected one.

**Returns** Wrapped function.

**Return type** *func*

**projected\_hamiltonian**(*bath, vectors, projections\_alpha, projections\_beta, mfield, others=None, other\_states=None, energy\_alpha=None, energy\_beta=None, energies=None, projections\_alpha\_all=None, projections\_beta\_all=None*)

Compute projected hamiltonian on state and beta qubit states. Wrapped function so the actual call does not follow the one above!

**Parameters**

- **bath** (*BathArray*) – array of all bath spins in the cluster.
- **projections\_alpha** (*ndarray with shape (3,)*) – Projections of the central spin level alpha  $[\hat{S}_x, \hat{S}_y, \hat{S}_z]$ .
- **projections\_beta** (*ndarray with shape (3,)*) – Projections of the central spin level beta.  $[\hat{S}_x, \hat{S}_y, \hat{S}_z]$
- **mfield** (*ndarray with shape (3,)*) – Magnetic field of type *mfield* = *np.array*([Bx, By, Bz]).
- **others** (*BathArray with shape (m,)*) – array of all bath spins outside the cluster
- **other\_states** (*ndarray with shape (m,) or (m, 3)*) – Array of Iz projections for each bath spin outside of the given cluster.
- **energy\_alpha** (*float*) – Energy of the alpha state
- **energy\_beta** (*float*) – Energy of the beta state
- **energies** (*ndarray with shape (2s-1,)*) – Array of energies of all states of the central spin.
- **projections\_alpha\_all** (*ndarray with shape (2s-1, 3)*) – Array of vectors of the central spin matrix elements of form:

$$[\alpha \hat{S}_{xj}, \alpha \hat{S}_{yj}, \alpha \hat{S}_{zj}],$$

where  $\alpha$  is the alpha qubit state, and  $\alpha$  are all states.

- **projections\_beta\_all** (*ndarray with shape (2s-1, 3)*) – Array of vectors of the central spin matrix elements of form:

$$[\beta \hat{S}_{xj}, \beta \hat{S}_{yj}, \beta \hat{S}_{zj}],$$

where  $\beta$  is the beta qubit state, and  $\alpha$  are all states.

### Returns

*tuple* containing:

- **Hamiltonian**: Hamiltonian of the given cluster, conditioned on the alpha qubit state.
- **Hamiltonian**: Hamiltonian of the given cluster, conditioned on the beta qubit state.

**Return type** *tuple*

**total\_hamiltonian**(*bath, vectors, mfield, zfs=None, others=None, other\_states=None, central\_gyro=-17608.59705*)

Compute total Hamiltonian for the given cluster including mean field effect of all bath spins. Wrapped function so the actual call does not follow the one above!

### Parameters

- **bath** (*BathArray*) – array of all bath spins.
- **mfield** (*ndarray with shape (3,)*) – Magnetic field of type `mfield = np.array([Bx, By, Bz])`.
- **others** (*BathArray with shape (m,)*) – array of all bath spins outside the cluster
- **other\_states** (*ndarray with shape (m,) or (m, 3)*) – Array of Iz projections for each bath spin outside of the given cluster.
- **zfs** (*ndarray with shape (3, 3)*) – Zero Field Splitting tensor of the central spin.
- **central\_gyro** (*float or ndarray with shape (3, 3)*) – gyromagnetic ratio of the central spin OR tensor corresponding to interaction between magnetic field and central spin.
- **central\_spin** (*float*) – value of the central spin.

**Returns** hamiltonian of the given cluster, including central spin.

**Return type** *Hamiltonian*

## 9.3 Separate Terms

Documentation for the functions used to generate spin Hamiltonian for each cluster.

**expanded\_single**(*ivec, gyro, mfield, self\_tensor, detuning=0.0*)

Function to compute the single bath spin term.

### Parameters

- **ivec** (*ndarray with shape (3, n, n)*) – Spin vector of the bath spin in the full Hilbert space of the cluster.
- **gyro** (*float or ndarray with shape (3, 3)*) –
- **mfield** (*ndarray with shape (3,)*) – Magnetic field of type `mfield = np.array([Bx, By, Bz])`.
- **self\_tensor** (*ndarray with shape (3, 3)*) – tensor of self-interaction of type IPI where I is bath spin.
- **detuning** (*float*) – Additional term of  $d \cdot I_z$  allowing to simulate different energy splittings of bath spins.

**Returns** Single bath spin term.

**Return type** ndarray with shape (n, n)

**dipole\_dipole**(*coord\_1, coord\_2, g1, g2, ivec\_1, ivec\_2*)

Compute dipole\_dipole interactions between two bath spins.

**Parameters**

- **coord\_1** (ndarray with shape (3,)) – Coordinates of the first spin.
- **coord\_2** (ndarray with shape (3,)) – Coordinates of the second spin.
- **g1** (float) – Gyromagnetic ratio of the first spin.
- **g2** (float) – Gyromagnetic ratio of the second spin.
- **ivec\_1** (ndarray with shape (3, n, n)) – Spin vector of the first spin in the full Hilbert space of the cluster.
- **ivec\_2** (ndarray with shape (3, n, n)) – Spin vector of the second spin in the full Hilbert space of the cluster.

**Returns** Dipole-dipole interactions.

**Return type** ndarray with shape (n, n)

**bath\_interactions**(*nspin, ivectors*)

Compute interactions between bath spins.

**Parameters**

- **nspin** (BathArray) – Array of the bath spins in the given cluster.
- **ivectors** (array-like) – array of expanded spin vectors, each with shape (3, n, n).

**Returns** All intrabath interactions of bath spins in the cluster.

**Return type** ndarray with shape (n, n)

**bath\_mediated**(*nspin, ivectors, energy\_state, energies, projections*)

Compute all hyperfine-mediated interactions between bath spins.

**Parameters**

- **nspin** (BathArray) – Array of the bath spins in the given cluster.
- **ivectors** (array-like) – array of expanded spin vectors, each with shape (3,n,n).
- **energy\_state** (float) – Energy of the qubit state on which the interaction is conditioned.
- **energies** (ndarray with shape (2s-1,)) – Array of energies of all states of the central spin.
- **projections** (ndarray with shape (2s-1, 3)) – Array of vectors of the central spin matrix elements of form:

$$[i\hat{S}_x j, i\hat{S}_y j, i\hat{S}_z j],$$

where *i* are different states of the central spin.

**Returns** Hyperfine-mediated interactions.

**Return type** ndarray with shape (n, n)

**conditional\_hyperfine**(*hyperfine\_tensor, ivec, projections*)

Compute conditional hyperfine Hamiltonian.

**Parameters**

- **hyperfine\_tensor** (*ndarray with shape (3, 3)*) – Tensor of hyperfine interactions of the bath spin.
- **ivec** (*ndarray with shape (3, n, n)*) – Spin vector of the bath spin in the full Hilbert space of the cluster.
- **projections** (*ndarray with shape (3,)*) – Array of vectors of the central spin matrix elements of form:

$$[i\hat{S}_{xj}, i\hat{S}_{yj}, i\hat{S}_{zj}],$$

where  $j$  are different states of the central spin. If  $i = j$ , produces the usual conditioned hyperfine interactions and just equal to projections of  $\hat{S}_z$  of the central spin state  $[\hat{S}_x, \hat{S}_y, \hat{S}_z]$ .

If  $i \neq j$ , gives second order perturbation.

**Returns** Conditional hyperfine interaction.

**Return type** ndarray with shape (n, n)

**hyperfine**(*hyperfine\_tensor, svec, ivec*)

Compute hyperfine interactions between central spin and bath spin.

**Parameters**

- **hyperfine\_tensor** (*ndarray with shape (3, 3)*) – Tensor of hyperfine interactions of the bath spin.
- **svec** (*ndarray with shape (3, n, n)*) – Spin vector of the central spin in the full Hilbert space of the cluster.
- **ivec** (*ndarray with shape (3, n, n)*) – Spin vector of the bath spin in the full Hilbert space of the cluster.

**Returns** Hyperfine interaction.

**Return type** ndarray with shape (n, n)

**self\_central**(*svec, mfield, zfs=None, gyro=- 17608.59705*)

Function to compute the central spin term in the Hamiltonian.

**Parameters**

- **svec** (*ndarray with shape (3, n, n)*) – Spin vector of the central spin in the full Hilbert space of the cluster.
- **mfield** (*ndarray with shape (3,)*) – Magnetic field of type `mfield = np.array([Bx, By, Bz])`.
- **zfs** (*ndarray with shape (3, 3)*) – Zero Field Splitting tensor of the central spin.
- **gyro** (*float or ndarray with shape (3, 3)*) – gyromagnetic ratio of the central spin OR tensor corresponding to interaction between magnetic field and central spin.

**Returns** Central spin term.

**Return type** ndarray with shape (n, n)

**overhauser\_central**(*svec, others\_hyperfines, others\_state*)

Compute Overhauser field term on the central spin from all other spins, not included in the cluster.

**Parameters**

- **svec** (*ndarray with shape (3, n, n)*) – Spin vector of the central spin in the full Hilbert space of the cluster.

- **others\_hyperfines** (*ndarray with shape (m, 3, 3)*) – Array of hyperfine tensors for all bath spins not included in the cluster.
- **others\_state** (*ndarray with shape (m,) or (m, 3)*) – Array of Iz projections for each bath spin outside of the given cluster.

**Returns** Central spin Overhauser term.

**Return type** ndarray with shape (n, n)

**overhauser\_bath**(*ivec, position, gyro, other\_gyros, others\_position, others\_state*)

Compute Overhauser field term on the bath spin in the cluster from all other spins, not included in the cluster.

**Parameters**

- **ivec** (*ndarray with shape (3, n, n)*) – Spin vector of the bath spin in the full Hilbert space of the cluster.
- **position** (*ndarray with shape (3,)*) – Position of the bath spin.
- **gyro** (*float*) – Gyromagnetic ratio of the bath spin.
- **other\_gyros** (*ndarray with shape (m,)*) – Array of the gyromagnetic ratios of the bath spins, not included in the cluster.
- **others\_position** (*ndarray with shape (m, 3)*) – Array of the positions of the bath spins, not included in the cluster.
- **others\_state** (*ndarray with shape (m,) or (m, 3)*) – Array of Iz projections for each bath spin outside of the given cluster.

**Returns** Bath spin Overhauser term.

**Return type** ndarray with shape (n, n)

**eta\_hamiltonian**(*nspin, central\_spin, alpha, beta, eta*)

EXPERIMENTAL. Compute hamiltonian with eta-term - gradually turn off or turn on the secular interactions for alpha and beta qubit states.

**Parameters**

- **nspin** (*BathArray*) – Array of the bath spins in the given cluster.
- **central\_spin** (*float*) – central spin.
- **alpha** (*ndarray with shape (2s+1,)*) – Vector representation of the alpha qubit state in Sz basis.
- **beta** (*ndarray with shape (2s+1,)*) – Vector representation of the beta qubit state in Sz basis.
- **eta** (*float*) – Value of dimensionless parameter eta (from 0 to 1).

**Returns** Eta term.

**Return type** ndarray with shape (n, n)



## UTILITY FUNCTIONS

Here are the various functions used throughout the **PyCCE** code. There is no real structure in this section.

Module with helper functions to obtain CPMG coherence from the noise autocorrelation function.

**filterfunc**(*ts, tau, npulses*)

Time-domain filter function for the given CPMG sequence.

**Parameters**

- **ts** (*ndarray with shape (n,)*) – Time points at which filter function will be computed.
- **tau** (*float*) – Delay between pulses.
- **npulses** (*int*) – Number of pulses in CPMG sequence.

**Returns** Filter function for the given CPMG sequence

**Return type** ndarray with shape (n,)

**gaussian\_phase**(*timespace, corr, npulses, units='khz'*)

Compute average random phase squared assuming Gaussian noise.

**Parameters**

- **timespace** (*ndarray with shape (n,)*) – Time points at which correlation function was computed.
- **corr** (*ndarray with shape (n,)*) – Noise autocorrelation function.
- **npulses** (*int*) – Number of pulses in CPMG sequence.
- **units** (*str*) – If units contain frequency or angular frequency ('rad' in units).

**Returns** Random phase accumulated by the qubit.

**Return type** ndarray with shape (n,)

**rotmatrix**(*initial\_vector, final\_vector*)

Generate 3D rotation matrix which applied on initial vector will produce vector, aligned with final vector.

## Examples

```
>>> R = rotmatrix([0,0,1], [1,1,1])
>>> R @ np.array([0,0,1])
array([0.577, 0.577, 0.577])
```

### Parameters

- **initial\_vector** (*ndarray with shape (3, )*) – Initial vector.
- **final\_vector** (*ndarray with shape (3, )*) – Final vector.

**Returns** Rotation matrix.

**Return type** ndarray with shape (3, 3)

**expand**(*matrix, i, dim*)

Expand matrix M from it's own dimensions to the total Hilbert space.

### Parameters

- **matrix** (*ndarray with shape (dim[i], dim[i])*) – Initial matrix.
- **i** (*int*) – Index of the spin dimensions in dim parameter.
- **dim** (*ndarray*) – Array of dimensions of all spins present in the cluster.

**Returns** Expanded matrix.

**Return type** ndarray with shape (prod(dim), prod(dim))

**dimensions\_spinvectors**(*nspin, central\_spin=None*)

Generate two arrays, containing dimensions of the spins in the cluster and the vectors with spin matrices.

### Parameters

- **nspin** (*BathArray with shape (n,)*) – Array of the n spins within cluster.
- **central\_spin** (*float, optional*) – If provided, include dimensions of the central spin with the total spin s.

### Returns

*tuple* containing:

- **ndarray with shape (n,)**: Array with dimensions for each spin.
- **list**: List with vectors of spin matrices for each spin in the cluster (Including central spin if central\_spin is not None). Each with shape (3, N, N) where N = prod(dimensions).

**Return type** tuple

**spinvec**(*s, j, dimensions*)

Generate spin vector for the particle, containing 3 spin matrices in the total basis of the system.

### Parameters

- **s** (*float*) – Spin of the particle.
- **j** (*j*) – Particle index in dimensions array.
- **dimensions** (*ndarray*) – Array with dimensions of all spins in the cluster.

**Returns** Vector of spin matrices for the given spin in the cluster.

**Return type** ndarray with shape (3, prod(dimensions), prod(dimensions))

**generate\_projections**(*state\_a*, *state\_b=None*)

Generate vector with the spin projections of the given spin states:

$$[a\hat{S}_x b, a\hat{S}_y b, a\hat{S}_z b],$$

where *a* and *b* are the given spin states.

**Parameters**

- **state\_a** (*ndarray*) – state *a* of the central spin in  $\hat{S}_z$  basis.
- **state\_b** (*ndarray*) – state *b* of the central spin in  $\hat{S}_z$  basis.

**Returns** [ $\hat{S}_x, \hat{S}_y, \hat{S}_z$ ] projections.

**Return type** *ndarray* with shape (3,)

**zfs\_tensor**(*D*, *E=0*)

Generate (3, 3) ZFS tensor from observable parameters D and E.

**Parameters**

- **D** (*float or ndarray with shape (3, 3)*) – Longitudinal splitting (D) in ZFS **OR** total ZFS tensor.
- **E** (*float*) – Transverse splitting (E) in ZFS.

**Returns** Total ZFS tensor.

**Return type** *ndarray* with shape (3, 3)

**project\_bath\_states**(*states*)

Generate projections of bath states on  $S_z$  axis from any type of states input. :param states: Array of bath spin states. :type states: array-like

**Returns** Array of  $S_z$  projections of the bath states

**Return type** *ndarray*

**partial\_inner\_product**(*avec*, *total*, *dimensions*, *index=-1*)

Returns partial inner product  $b = a\psi$ , where *a* provided by *avec* contains degrees of freedom to be “traced out” and  $\psi$  provided by *total* is the total statevector.

**Parameters**

- **avec** (*ndarray with shape (a,)*) –
- **total** (*ndarray with shape (a\*b,)*) –
- **dimensions** (*ndarray with shape (n,)*) –
- **O** (*index*) –

Returns:

**class SpinMatrix**(*s*)

Class containing the spin matrices in  $S_z$  basis.

**Parameters** **s** (*float*) – Total spin.

**class MatrixDict**(\**spins*)

Class for storing the SpinMatrix objects.

**keys**() → a set-like object providing a view on D’s keys



## ES INTERFACE

Each of the interfaces uses subclass of the `DFTCoordinates` class to parse the output.

---

**Note:** The interfaces are in beta stage. Please let us know if you encounter any errors.

---

### 11.1 Quantum Espresso

**class** `PWCoordinates`(*filename*, *pwtype=None*, *to\_angstrom=False*)

Coordinates of the system from the PW data of Quantum Espresso. Subclass of the `DFTCoordinates`.

With initialization reads either output or input of PW module of QE.

#### Parameters

- **filename** (*str*) – name of the PW input or output.
- **pwfile** (*str*) – Name of PW input or output file. If the file doesn't have proper extension, parameter `pw_type` should indicate the type.
- **pwtype** (*str*) – Type of the `coord_f`. if not listed, will be inferred from extension of `pwfile`.
- **to\_angstrom** (*bool*) – True if automatically convert the units of `cell` and `coordinates` to Angstrom.

**parse\_output**(*filename*, *to\_angstrom=False*)

Method to read coordinates of atoms from PW output into the `PWCoordinates` instance.

#### Parameters

- **filename** (*str*) – the name of the output file.
- **to\_angstrom** (*bool*) – True if automatically convert the units of `cell` and `coordinates` to Angstrom.

**Returns** None

**parse\_input**(*filename*, *to\_angstrom=False*)

Method to read coordinates of atoms from PW input into the `PWCoordinates` instance.

#### Parameters

- **filename** (*str*) – the name of the output file.
- **to\_angstrom** (*bool*) – True if automatically convert the units of `cell` and `coordinates` to Angstrom.

**cell\_from\_system**(*sdict*)

Function to obtain cell from namelist SYSTEM read from PW input.

**Parameters** *sdict* (*dict*) – Dictionary generated from namelist SYSTEM of PW input.

**Returns**

Cell is 3x3 matrix with entries:

```
[[a_x b_x c_x]
 [a_y b_y c_y]
 [a_z b_z c_z]],
```

where a, b, c are crystallographic vectors, and x, y, z are their coordinates in the cartesian reference frame.

**Return type** ndarray with shape (3,3)

**celldms\_from\_abc**(*ibrav*, *abc\_list*)

Obtain celldms from ibrav value and a, b, c, cosab, cosac, cosbc parameters.

Using ibrav value and abc parameters from PW input generate celldm array, necessary to construct cell parameters. For details about abc and ibrav values see PW input documentation.

**Parameters**

- **ibrav** (*int*) – ibrav parameter of PW input.
- **abc\_list** (*list*) – List, of 6 parameters: a, b, c, cosab, cosac, cosbc

**Returns** list of 6 values, from which cell can be generated.

**Return type** celldm (list)

**read\_gipaw\_tensors**(*lines*, *keyword=None*, *start=None*, *conversion=1*)

Helper function to read GIPAW tensors from the list of lines.

**Parameters**

- **lines** (*list of str*) – List of strings containing lines from the file. Output of `open(file).readlines()`.
- **keyword** (*str*) – Keyword in the line which indicates the beginning of the tensor data block.
- **start** (*int*) – Index of the line which indicates the beginning of the tensor data block.
- **conversion** (*float*) – Conversion factor from GIPAW units to the ones, used in this package.

**Returns** Array of tensors.

**Return type** ndarray with shape (n, 3, 3)

**read\_hyperfine**(*filename*, *spin=1*)

Function to read hyperfine couplings from GIPAW output.

**Parameters**

- **filename** (*str*) – Name of the GIPAW hyperfine output.
- **spin** (*float*) – Spin of the central spin. Default 1.

**Returns**

Tuple containing:

- *ndarray with shape (n,):* Array of Fermi contact terms.

- *ndarray with shape (n, 3,3)*: Array of spin dipolar hyperfine tensors.

**Return type** tuple

**read\_efg**(*filename*)

Function to read electric field gradient tensors from GIPAW output.

**Parameters** **filename** (*str*) – Name of the GIPAW EFG-containing output.

**Returns** Array of EFG tensors.

**Return type** ndarray with shape (n, 3,3)

**read\_qe\_namelists**(*input\_string*)

Read Fortran-like namelists from the large string.

**Parameters** **input\_string** (*str*) – String representation of the QE input file.

**Returns** Dictionary, containing dicts for each namelist found in the input string.

**Return type** dict

**get\_ctype**(*lin*)

Get coordinates type from the line of QE input/output.

**Parameters** **str** – Line from QE input/output containing string with coordinates type.

**Returns** type of the coordinates.

**Return type** str

## 11.2 ORCA

**class ORCACoordinates**(*orca\_output*)

Coordinates of the system from the ORCA output. Subclass of the DFTCoordinates.

With initialization reads output of the ORCA.

**Parameters** **orca\_output** (*str or list of str*) – either name of the output file or list of lines read from that file.

**alat**

The lattice parameter in angstrom.

**Type** float

**cell**

cell is 3x3 matrix with entries:

$$\begin{bmatrix} a_x & b_x & c_x \\ a_y & b_y & c_y \\ a_z & b_z & c_z \end{bmatrix}$$

where a, b, c are crystallographic vectors, and x, y, z are their coordinates in the cartesian reference frame.

**Type** ndarray with shape (3, 3)

**coordinates**

array with the coordinates of atoms in the cell.

**Type** ndarray with shape (n, 3)

**names**

array with the names of atoms in the cell.

**Type** ndarray with shape (n,)

**cell\_units**

Units of cell coordinates: 'bohr', 'angstrom', 'alat'.

**Type** str

**coordinates\_units**

Units of atom coordinates: 'crystal', 'bohr', 'angstrom', 'alat'.

**Type** str

**read\_output**(*orca\_output*)

Method to read coordinates of atoms from ORCA output into the ORCACoordinates instance.

**Parameters** *orca\_output* (*str* or *list of str*) – either name of the output file or list of lines read from that file.

## 11.3 Base class

### class DFTCoordinates

Abstract class of a container of the DFT output coordinates.

**alat**

The lattice parameter in angstrom.

**Type** float

**cell**

cell is 3x3 matrix with entries:

$$\begin{bmatrix} a_x & b_x & c_x \\ a_y & b_y & c_y \\ a_z & b_z & c_z \end{bmatrix}$$

where a, b, c are crystallographic vectors and x, y, z are their coordinates in the cartesian reference frame.

**Type** ndarray with shape (3, 3)

**coordinates**

Array with the coordinates of atoms in the cell.

**Type** ndarray with shape (n, 3)

**names**

Array with the names of atoms in the cell.

**Type** ndarray with shape (n,)

**cell\_units**

Units of cell coordinates: 'bohr', 'angstrom', 'alat'.

**Type** str

**coordinates\_units**

Units of atom coordinates: 'crystal', 'bohr', 'angstrom', 'alat'.

**Type** str

**to\_angstrom**(*inplace=False*)

Method to transform cell and coordinates units to angstroms.



**Parameters** **inplace** (*bool*) – if True changes attributes inplace. Otherwise returns copy.

**Returns** Instance of the subclass with units of coordinates and cell of Angstroms.

**Return type** *DFTCoordinates* or subclass

**get\_angstrom**(*coordinate, units*)

Change given coordinates to angstrom.

**Parameters**

- **coordinates** (*ndarray with shape (n, 3) or (3,)*) – Coordinates to change.
- **units** (*str*) – Initial units of the coordinates.

**Returns** Coordinates in angstrom.

**Return type** *ndarray (n, 3)*

**change\_to\_angstrom**(*coordinates, units, alat=None, cell=None*)

Change coordinates to angstrom.

**Parameters**

- **coordinates** (*ndarray with shape (n, 3) or (3,)*) – Coordinates to change.
- **units** (*str*) – Initial units of the coordinates.
- **alat** (*float*) – The lattice parameter in angstrom.
- **cell** (*ndarray with shape (3, 3)*) – cell is 3x3 matrix with entries:

$$\begin{bmatrix} a_x & b_x & c_x \\ a_y & b_y & c_y \\ a_z & b_z & c_z \end{bmatrix}$$

where a, b, c are crystallographic vectors, and x, y, z are their coordinates in the cartesian reference frame.

**Returns** Coordinates in angstrom.

**Return type** *ndarray with shape (n, 3)*

**fortran\_value**(*value*)

Get value from Fortran-type variable.

**Parameters** **value** (*str*) – Value read from Fortran-type input.

**Returns** value in Python format.

**Return type** value (*bool, str, float*)

**yield\_index**(*word, lines, start=0, case\_sensitive=False*)

Generator which yields indexes of the lines containing specific word.

**Parameters**

- **word** (*str*) – Word to find in the line.
- **lines** (*list of str*) – List of strings containing lines from the file. Output of `open(file).readlines()`.
- **start** (*int*) – First index from which to start search.
- **case\_sensitive** (*bool*) – If True looks for the exact match. Otherwise the search is case insensitive.

**Yields** *i (int)* – Index of the line containing word.

**find\_first\_index**(*word*, *lines*, *start=0*, *case\_sensitive=False*)

Function to find first appearance of the index in the list of lines.

**Parameters**

- **word** (*str*) – Word to find in the line.
- **lines** (*list of str*) – List of strings containing lines from the file. Output of `open(file).readlines()`.
- **start** (*int*) – First index from which to start search.
- **case\_sensitive** (*bool*) – If True looks for the exact match. Otherwise the search is case insensitive.

**Returns** Index of the first line from the start containing word.

**Return type** *i* (int)

**set\_isotopes**(*array*, *isotopes=None*, *inplace=True*, *spin\_types=None*)

Function to set the most common isotopes for the array containing DFT output. If some other isotope is specified, the A tensors are scaled accordingly.

**Parameters**

- **array** (*BathArray*) – Array with DFT spins.
- **isotopes** (*dict*) – Dictionary with chosen isotopes.
- **inplace** (*bool*) – True if change the array inplace.
- **spin\_types** (*SpinDict*) – If provided, allows for custom defined *SpinType* instances.

**Returns** Array with DFT spins with correct isotopes.

**Return type** array (*BathArray*)

PyCCE is an open source Python library to simulate the dynamics of a spin qubit interacting with a spin bath using the cluster-correlation expansion (CCE) method.



## INSTALLATION

The recommended way to install **PyCCE** is to use **pip**:

```
$ pip install pycce
```

Otherwise you can install **PyCCE** directly using the source code. First copy the repository to the desired folder:

```
$ git clone https://github.com/foxfifax/pycce.git
```

Then, execute **pip** in the folder containing **setup.py**:

```
$ pip install .
```

or run the python install command:

```
$ python setup.py install
```



## REQUIREMENTS

The following modules are required to run **PyCCE**.

- Python (version  $\geq 3.6$ ).
- NumPy (version  $\geq 1.16$ ).
- SciPy.
- Numba (version  $\geq 0.50$ ).
- Atomic Simulation Environment (ASE).
- Pandas.

**PyCCE** inherently supports parallelization with the **mpi4py** package, which requires the installation of MPI. However, for serial implementation the **mpi4py** is not required.



## HOW TO CITE

If you make use of **PyCCE** in a scientific publication, please cite the following paper:

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