Spglib for Python

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Installation

Source codes, examples, and the test are downloaded SourceForge (https://sourceforge.net/project/showfiles.php?group_id=215020) or GitHub (https://github.com/atztogo/spglib/releases).

Using package distribution service

The easiest way to install python-spglib is to use the pypi package, for which numpy is required to be installed before the installation. A command to install spglib is:

% pip install spglib

Previous versions < 1.8.x are installed by:

% pip install pyspglib

Conda is another choice:

% conda install -c https://conda.anaconda.org/jochym spglib

These packages are made by Paweł T. Jochym.

Building using setup.py (distutils)

To manually install python-spglib using setup.py, python header files (python-dev), gcc, and numpy are required before the build. The installation steps are shown as follows:

- 1. Go to the python directory
- 2. Type the command:

% python setup.py install --home=<my-directory>

The <my-directory> is possibly current directory, . or maybe:

% python setup.py install --user

to use the user scheme (see the python document.)

3. Put ./lib/python path into \$PYTHONPATH, e.g., in your .washrag.

Test

The test script test_spglib.py is found in python/test directory. Got to this directory and run this script. It will be like below:

How to import spglib module

Change in version 1.9.0!

For versions 1.9.x or later:

```
import spglib
```

For versions 1.8.x or before:

```
from pyspglib import spglib
```

If the version is not sure:

```
try:
    import spglib as spg
except ImportError:
    from pyspglib import spglib as spg
```

Version number

In version 1.8.3 or later, the version number is obtained by spglib.__version__ or get_version.

Example

Examples are found in examples (https://github.com/atztogo/spglib/tree/master/python/examples) directory.

Variables

Crystal structure (cell)

A crystal structure is given by a **tuple**. This tuple format is supported at version 1.9.1 or later. Optionally, an **ASE Atoms-like object** is also supported. An alternative Atoms class (atoms.py (https://github.com/atztogo/spglib/blob/master/python/examples/atoms.py)) that contains minimum set of methods is prepared in the examples (https://github.com/atztogo/spglib/tree/master/python/examples) directory. When using ASE Atoms-like object, get_symmetry with collinear polarizations is not supported.

The tuple format is shown as follows. There are three or four elements in the tuple: cell = (lattice, positions, numbers) or cell = (lattice, positions, numbers, magmoms) where magmoms represents collinear polarizations on atoms and is optional.

Lattice parameters lattice are given by a 3x3 matrix with floating point values, where $\mathbf{a}, \mathbf{b}, \mathbf{c}$ are given as rows, which results in the transpose of the definition for C-API (latti (variable.html#variables-lattice)). Fractional atomic positions positions are given by a Nx3 matrix with floating point values, where N is the number of atoms. Numbers to distinguish atomic species numbers are given by a list of N integers. The collinear polarizations magmoms only work with get_symmetry and are given as a list of N floating | values.

Version 1.9.5 or later: When a crystal structure is not properly given, the methods that use the crsytal structure will return None.

Symmetry tolerance (symprec)

Distance tolerance in Cartesian coordinates to find crystal symmetry.

Methods

get_version

New in version 1.8.3

```
version = get_version()
```

This returns version number of spglib by tuple with three numbers.

get_spacegroup

```
spacegroup = get_spacegroup(cell, symprec=1e-5)
```

International space group short symbol and number are obtained as a string. With symbol_type=1, Schoenflies symbol is given instead of international symbol.

get_symmetry

```
symmetry = get_symmetry(cell, symprec=1e-5)
```

Symmetry operations are obtained as a dictionary. The key rotation contains a numpy array of integer, which is "number of symmetry operations" x "3x3 matrices". The key translation contains a numpy array of float, which is "number of symmetry operations" x "vectors". The orders of the rotation matrices and the translation vectors correspon with each other, e.g., the second symmetry operation is organized by the set of the second rotation matrix and second translation vector in the respective arrays. Therefore a set symmetry operations may obtained by:

```
[(r, t) for r, t in zip(dataset['rotations'], dataset['translations'])]
```

The operations are given with respect to the fractional coordinates (not for Cartesian coordinates). The rotation matrix and translation vector are used as follows:

```
new\_vector[3x1] = rotation[3x3] * vector[3x1] + translation[3x1]
```

The three values in the vector are given for the a, b, and c axes, respectively. The key equivalent_atoms gives a mapping table of atoms to symmetrically independent atoms is used to find symmetrically equivalent atoms. The numbers contained are the indices of atoms starting from 0, i.e., the first atom is numbered as 0, and then 1, 2, 3, ... np.unique(equivalent_atoms) gives representative symmetrically independent atoms. A list of atoms that are symmetrically euivalent to some independent atom (here feexample 1 is in equivalent_atom) is found by np.where(equivalent_atom=1) [0].

If cell is given as a tuple and collinear polarizations are given as the fourth element of this tuple, symmetry operations are searched considering this freedome. In ASE Atoms-c object, this is not supported.

refine cell

Behaviour changed in version 1.8.x

```
lattice, scaled_positions, numbers = refine_cell(cell, symprec=1e-5)
```

Bravais lattice (3x3 numpy array), atomic scaled positions (a numpy array of [number_of_atoms,3]), and atomic numbers (a 1D numpy array) that are symmetrized following spagroup type are returned. When the search failed, None is returned.

The detailed control of standardization of unit cell may be done using $\verb|standardize_cell|.$

find primitive

Behaviour changed in version 1.8.x

```
lattice, scaled_positions, numbers = find_primitive(cell, symprec=1e-5)
```

When a primitive cell is found, lattice parameters (3x3 numpy array), scaled positions (a numpy array of [number_of_atoms,3]), and atomic numbers (a 1D numpy array) is return When the search failed, None is returned.

The detailed control of standardization of unit cell can be done using $\verb|standardize_cell|.$

standardize_cell

New in version 1.8.x

```
lattice, scaled_positions, numbers = standardize_cell(bulk, to_primitive=False, no_idealize=False, symprec=1e-5)
```

to_primitive=True is used to create the standardized primitive cell, and no_idealize=True disables to idealize lengths and angles of basis vectors and positions of ator according to crystal symmetry. Now refine_cell and find_primitive are shorthands of this method with combinations of these options. When the search failed, None returned. Is returned. More detailed explanation is shown in the spglib (C-API) document.

get_symmetry_dataset

At version 1.9.4, the member 'choice' is added.

```
dataset = get_symmetry_dataset(cell, symprec=1e-5)
```

dataset is a dictionary. The keys are:

- number: International space group number
- international: International short symbol

- hall: Hall symbol
- choice: Centring, origin, basis vector setting
- transformation_matrix: Transformation matrix from lattice of input cell to Bravais lattice $L^{bravais} \equiv L^{original}*T$
- origin shift: Origin shift choice in the setting of Bravais lattice
- wyckoffs: Wyckoff letters
- equivalent atoms: Mapping table to equivalent atoms
- rotations and translations: Rotation matrices and translation vectors. See get_symmetry for more details
- pointgroup_symbol: Symbol of the crystallographic point group in the Hermann-Mauguin notation.
- std_lattice, std_positions, std_types: Standardized crystal structure corresponding to a Hall symbol found. These are equivalently given in the array formats lattice, positions, and numbers presented at Crystal structure (cell), respectively.

When the search failed, None is returned.

get_symmetry_from_database

```
symmetry = get_symmetry_from_database(hall_number)
```

A set of crystallographic symmetry operations corresponding to hall_number is returned by a dictionary where rotation parts and translation parts are accessed by the keys rotations and translations, respectively. The definition of hall_number is found at Space group type (api.html#api-spg-get-dataset-spacegroup-type).

When something wrong happened, None is returned.

get_spacegroup_type

New at version 1.9.4

```
spacegroup_type = get_spacegroup_type(hall_number)
```

This function allows to directly access to the space-group-type database in spglib (spg_database.c). A dictionary is returned. To specify the space group type with a specific choic hall_number is used. The definition of hall_number is found at Space group type (api.html#api-spg-get-dataset-spacegroup-type). The keys of the returned dictionary is as follows:

number
international_short
international_full
international
schoenflies
hall_symbol
choice
pointgroup_schoenflies
pointgroup_international
arithmetic_crystal_class_number
arithmetic_crystal_class_symbol

Here spacegroup_type['international_short'] is equivalent to dataset['international'] of get_symmetry_dataset, spacegroup_type['hall_symbolis equivalent to dataset['hall'] of get_symmetry_dataset, and spacegroup_type['pointgroup_international'] is equivalent to dataset['pointgroup_symbol'] of get_symmetry_dataset.

When something wrong happened, None is returned.

niggli reduce

New at version 1.9.4

```
niggli_lattice = niggli_reduce(lattice, eps=1e-5)
```

Niggli reduction is achieved using this method. The algorithm detail is found at https://atztogo.github.io/niggli/ (https://atztogo.github.io/niggli/) and the references are there in vectors are stored in lattice and niggli_lattice as shown in Crystal structure (cell). esp is the tolerance parameter, but unlike symprec the unit is not a length. This i to check if difference of norms of two basis vectors is close to zero or not and if two basis vectors are orthogonal by the value of dot product being close to zero or not. The detail shown at https://atztogo.github.io/niggli/ (https://atztogo.github.io/niggli/).

When the search failed, None is returned.

delaunay_reduce

New at version 1.9.4

```
delaunay_lattice = delaunay_reduce(lattice, eps=1e-5)
```

Delaunay reduction is achieved using this method. The algorithm is found in the international tables for crystallography volume A. Basis vectors are stored in lattice and niggli_lattice as shown in Crystal structure (cell). esp is the tolerance parameter, but unlike symprec the unit is not a length. This is used as the criterion if volume is closero or not and if two basis vectors are orthogonal by the value of dot product being close to zero or not.

When the search failed, None is returned.

get_ir_reciprocal_mesh

```
mapping, grid = get_ir_reciprocal_mesh(mesh, cell, is_shift=[0, 0, 0])
```

Irreducible k-points are obtained from a sampling mesh of k-points. mesh is given by three integers by array and specifies mesh numbers along reciprocal primitive axis. is_sh is given by the three integers by array. When is_shift is set for each reciprocal primitive axis, the mesh is shifted along the axis in half of adjacent mesh points irrespective of t mesh numbers. When the value is not 0, is_shift is set.

mapping and grid are returned. grid gives the mesh points in fractional coordinates in reciprocal space. mapping gives mapping to the irreducible k-point indices that ar obtained by

```
np.unique(mapping)
```

Here np is the imported numpy module. The grid point is accessed by grid[index].

When the sesarch failed, None is returned.

For example, the irreducible k-points in fractional coordinates are obtained by

```
ir_grid = []
mapping, grid = get_ir_reciprocal_mesh([ 8, 8, 8 ], cell, [1, 1, 1])
for i in np.unique(mapping):
    ir_grid.append(grid[i])
```

Source (_sources/python-spglib.txt)

Back

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