Cif2x Documentation

Release 1.0-alpha

ISSP, University of Tokyo

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Introduction

1.1 What is cif2x?

In recent years, the use of machine learning for predicting material properties and designing substances (known as materials informatics) has gained considerable attention. The accuracy of machine learning depends heavily on the preparation of appropriate training data. Therefore, the development of tools and environments for the rapid generation of training data is expected to contribute significantly to the advancement of research in materials informatics.

Cif2x is a tool that generates input files for first-principles calculations from cif files. It constructs parts that vary depending on the type of material and computational conditions from crystal structure data, using input parameters as a template. It is capable of generating multiple input files tailored to specific computational conditions. Currently, it supports VASP https://www.vasp.at, Quantum ESPRESSO https://www.quantum-espresso.org, and OpenMX http://www.openmx-square.org, with plans to support AkaiKKR http://kkr.issp.u-tokyo.ac.jp in the future.

1.2 License

The distribution of the program package and the source codes for cif2x follow GNU General Public License version 3 (GPL v3) or later.

1.3 Contributors

This software was developed by the following contributors.

- ver.1.0-alpha (Released on 2023/12/28)
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1.4 Copyright

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This software was developed with the support of "Project for advancement of software usability in materials science" of The Institute for Solid State Physics, The University of Tokyo.

1.5 Operating environment

This tool was tested on the following platforms:

• Ubuntu Linux + python3

Installation and basic usage

Prerequisite

Input file generator for first-principles calculation cif2x included in HTP-tools requires the following programs and libraries:

- python 3.x
- pymatgen module
- ruamel.yaml module
- f90nml module
- · qe-tools module
- numpy module
- · pandas module
- · monty module
- OpenBabel module (optional)

Official pages

• GitHub repository

Downloads

cif2x can be downloaded by the following command with git:

```
$ git clone https://github.com/issp-center-dev/cif2x.git
```

Installation

Once the source files are obtained, you can install cif2x by running the following command. The required libraries will also be installed automatically at the same time.

```
$ cd ./cif2x
$ python3 -m pip install .
```

The executable file cif2x will be installed. You may need to add --user option next to install keyword above in case you are not allowed to install packages system-wide.

Directory structure

```
|-- LICENSE
|-- README.md
|-- pyproject.toml
|-- docs/
   |-- ja/
    |-- en/
   |-- tutorial/
|-- src/
   |-- cif2x/
        |-- __init__.py
        |-- main.py
        |-- cif2struct.py
        |-- struct2qe.py
        |-- qe/
        | |-- __init__.py
          |-- calc_mode.py
           |-- cards.py
        | |-- content.py
          |-- qeutils.py
          |-- tools.py
        |-- struct2vasp.pv
        |-- struct2openmx.py
        |-- openmx/
            |-- __init__.py
            |-- vps_table.py
        |-- utils.py
   sample/
```

Basic usage

cif2x is a tool to generate a set of input files for first-principles calculation software. It takes an input parameter file as a template, and generates parameter items that may vary by materials and calculation conditions from crystallographic data. In the present version, cif2x supports Quantum ESPRESSO, VASP, and OpenMX.

1. Prepare input parameter file

First, you need to create an input parameter file in YAML format that describes contents of the input file to be generated for the first-principles calculation software.

2. Prepare crystal structure files and pseudo-potential files

The crystal structure data need to be prepared for the target materials. The file format is CIF, POSCAR, xfs, or those supported by pymatgen.

For Quantum ESPRESSO, the pseudo-potential files and the index file in CSV format need to be placed. Their locations are specified in the input parameter file.

For VASP, the location of the pseudo-potential files will be specified in a file ~/.config/.pmgrc.yaml or by an environment variable. It may be specified in the input parameter file.

3. Run command

Run cif2x command with the input parameter file and the crystal structure data as arguments. To generate input files for Quantum ESPRESSO, the target option -t QE should be

specified. The option turns to -t VASP for VASP, and -t OpenMX for OpenMX.

\$ cif2x -t QE input.yaml material.cif

Tutorial

The procedure to use the input file generator cif2x for first-principles calculation software consists of preparing an input parameter file, crystal structure data, and pseudo-potential files, and running the program cif2x. In the current version, the supported software includes Quantum ESPRESSO, VASP, and OpenMX. In this tutorial, we will explain the steps along a sample for Quantum ESPRESSO in docs/tutorial/cif2x.

3.1 Prepare an input parameter file

An input parameter file describes the content of input files for the first-principles calculation software. An example is presented below. It is a text file in YAML format that contains options to crystal structure data, and contents of the input file used as an input for the first-principle calculation. See *file format* section for the details of specification.

In YAML format, parameters are given in dictionary form as keyword: value, where value is a scalar such as a number or a string, or a set of values enclosed in [] or listed in itemized form, or a nested dictionary.

```
structure:
 use_ibrav: false
  tolerance: 0.05
optional:
 pseudo_dir: ./pseudo
 pp_file: ./pseudo/pp_psl_pbe_rrkjus.csv
tasks:
  - mode: scf
    output_file: scf.in
    output_dir: scf
    template: scf.in_tmpl
    content:
      namelist:
        control:
          prefix: pwscf
          pseudo_dir:
          outdir: ./work
        system:
          ecutwfc:
```

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```
ecutrho:
CELL_PARAMETERS:
ATOMIC_SPECIES:
ATOMIC_POSITIONS:
option: crystal
K_POINTS:
option: automatic
grid: [8,8,8]
```

The input parameter file consists of structure, optional, and tasks sections. The structure section specifies options to the crystal structure data. The optional section holds global settings concerning the pseudo-potentials.

The tasks section describes inputs for the first-principles calculations. In case of generating multiple files for a series of calculations, the tasks section takes a list of parameter sets. For each set, the calculation type is specified by the mode parameter: scf and nscf are supported as modes, as well as arbitrary modes for generic output.

The content of the output is given in content section. The input files of Quantum ESPRESSO are composed of the parts in namelist format of Fortran90 starting from &keyword, and the blocks called cards that start with keywords such as K_POINTS and end with blank lines. The content block holds namelist and cards in a form of nested dictionary. Basically, the specified items are written to the input files as-is, except for several cases. If a keyword is left blank, its value will be obtained form the crystal structure data or other sources.

Besides, templates of the input files can be used. The content of the file given by the template keyword is considered as input data along with the entries in content block. When the entries of the same keywords appear both, those of the input parameter files will be used. Therefore, it is possible to use template files and overwrite some entries by the input parameter file as needed. In the present example, the file (scf.in_tmpl) shown below is read as a template, and the entries on cutoff parameters as well as cards of CELL_PARAMETER, ATOMIC_SPECIES, ATOMIC_POSITIONS, K_POINTS are generated from the crystal structure data and pseudo-potential files. It is noted that the values of ecutwfc and ecutrho are overwritten by the empty lines.

```
&control
    calculation = 'scf'
    prefix = 'pwscf'
    pseudo_dir = './pseudo'
    outdir = './work'
    tstress = .true.
    tprnfor = .true.
&system
    ibrav = 0
    nat = 7
   ntyp = 3
    ecutwfc = 36.0
    ecutrho = 180.0
    occupations = 'smearing'
    smearing = 'm-p'
    degauss = 0.01
    noncolin = .true.
    nspin = 2
&electrons
```

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```
missing_beta = 0.1
conv_thr = 1e-08
/
```

3.2 Generating input files

The program cif2x is executed with the input parameter file (input.yaml) and crystal structure data (Co3SnS2_nosym.cif) as follows.

```
$ cif2x -t QE input.yaml Co3SnS2_nosym.cif
```

The required pseudo-potential files should be placed in the directory ./pseudo, and the index file for the pseudo-potential should be prepared as ./psudo/pp_psl_pbe_rrkjus.csv.

Run cif2x and a set of input files for Quantum ESPRESSO will be created. The output file is specified by output_file parameter of the input parameter file, and stored in the directory given by output_dir. In this example, the input file for SCF calculation is created as ./scf/scf.in.

3.3 Specifying parameter sets

In some cases, a series of input files should be generated with varying their parameter values. For example, the convergence is examined by modifying the cutoff values or grid resolution of k points. The input parameter can be given a list or a range of values, and the input files for every combination from the choices of parameter values are generated and stored in separate directories. To specify parameter set, a special syntax \${...} is adopted.

```
content:
    K_POINTS:
        option: automatic
        grid: ${ [ [4,4,4], [8,8,8], [12,12,12] ] }
```

When K_POINTS is given as above, the input files having the grid value to be [4,4,4], [8,8,8], [12,12,12] will be generated in the sub-directories, 4x4x4/, 8x8x8/, 12x12x12/, respectively.

8 Chapter 3. Tutorial

Command reference

4.1 cif2x

Generate input files for first-principles calculation software

SYNOPSIS:

```
cif2x [-v][-q] -t target input_yaml material.cif
cif2x -h
cif2x --version
```

DESCRIPTION:

This program reads an input parameter file specified by input_yaml and a crystal data file specified by material.cif, and generates a set of input files for first-principles calculation software. In the current version, the supported software includes Quantum ESPRESSO, VASP, and OpenMX. It takes the following command line options.

• -v

increases verbosity of the runtime messages. When specified multiple times, the program becomes more verbose.

• -q

decreases verbosity of the runtime messages. It cancels the effect of -v option, and when specified multiple times, the program becomes more quiet.

-t target

specifies the target first-principles calculation software. The supported software for *target* is listed as follows:

- QE, espresso, quantum_espresso: generates input files for Quantum ESPRESSO.
- VASP: generates input files for VASP.
- OpenMX: generates input files for OpenMX
- input_yaml

specifies an input parameter file in YAML format.

- material.cif
 - specifies crystal structure data file. It is in CIF (Crystallographic Information Framework) format, or other format supported by pymatgen.
- -h
 displays help and exits.
- --version displays version information.

File format

5.1 Input parameter file

An input parameter file describes information necessary to generate input files for first-principles calculation software by cif2x. It should be given in YAML format, and consist of the following sections.

- 1. structure section: describes how to handle crystal structure data.
- 2. optional section: describes pseudo-potential files, and symbol definitions for reference feature of YAML.
- 3. tasks section: describes contents of input files.

5.1.1 structure

use_ibrav (default value: false)

This parameter specifies whether ibrav parameter is used for Quantum ESPRESSO as the input of the crystal structure. When it is set to true, the lattice is transformed to match the convention of Quantum ESPRESSO, and the lattice parameters a, b, c, cosab, cosac, and cosbc are written to the input file as needed.

tolerance (default value: 0.01)

This parameter specifies the tolerance in the difference between the reconstructed Structure data and the original data when use_ibrav is set to true.

supercell (default value: none)

This parameter specifies the size of supercell, when it is adopted, in the form of $[n_x, n_y, n_z]$.

5.1.2 optional

This section contains global settings needed for the first-principles calculation software. The available parameters are described in the corresponding sections below.

5.1.3 tasks

This section defines contents of the input files. It is organized as a list of blocks, each corresponding to an input file, to allow for generating a set of input files for an input. The terms described in each block are explained in the following.

```
mode (Quantum ESPRESSO)
```

This parameter specifies the type of calculation. In the current version, the supported mode includes scf and nscf for pw.x of Quantum ESPRESSO. If an unsupported mode is specified, the settings in content will be exported as is.

```
output_file (Quantum ESPRESSO)
```

This parameter specifies the file name of the output.

```
output_dir
```

This parameter specifies the directory name of the output. The default value is the current directory.

content

This parameter describes the content of the output. For Quantum ESPRESSO, it contains the namelist data (blocks starting from &system, &control, etc.) in namelist block, and other card data (such as K_POINTS) as individual blocks. Some card data may take parameters.

```
template (Quantum ESPRESSO)
```

```
template_dir (VASP)
```

These parameters specifies the template file and the template directory for the input files, respectively. If they are not given, templates will not be used. The content of the template file is merged with those of content. The entries in the template file will be superseded by those of content if the entries of the same keys appear both.

5.1.4 Specifying parameter set

An input parameter may be given a list or range of parameters. In this case, a separate directory is created for every combination of parameters to store the generated input files. A special syntax \${...} is used to specify the parameter set as follows:

```
• a list: ${[ A, B, ... ]}
```

a set of parameter values is described as a Python list. Each entry may be a scalar value, or a list of values.

```
• a range: ${range(N)}, ${range(start, end, step)}
```

a range of parameter is given by the keyword range. The former specifies the values from 0 to N-1, and the latter from start to end with every step. (If step is omitted, it is assumed to be 1.)

5.2 Parameters for Quantum ESPRESSO

The entries of optional section and content part of the tasks section specific to Quantum ESPRESSO are explained below. In the current version, scf mode and nscf mode of pw.x are supported.

5.2.1 optional section

pp_file

This parameter specifies the index file in CSV format that relates the element type and the pseudo-potential file. This file contains the following columns: element name, type of pseudo-potential, nexclude, orbitals. An example line is given as:

```
Fe,pbe-spn-rrkjus_psl.0.2.1,4,spd
```

The name of the pseudo-potential file corresponding to the above example reads Fe.pbe-spn-rrkjus_psl.0.2.1.UPF.

cutoff file

This parameter specifies the index file in CSV format that relates the pseudo-potential file and the cutoff values. This file contains the following columns: name of pseudo-potential file, ecutwfc value, ecutrho value.

pseudo_dir

This parameter specifies the name of the directory that holds pseudo-potential files. It is used when the cutoff values are obtained from the pseudo-potential files. It is indenepent from the pseudo_dir parameter in the input files for Quantum ESPRESSO.

5.2.2 content

namelist

- The lattice specifications in &system block will be superseded according to use_ibrav parameter in the structure section.
 - use_ibrav = false: ibrav is set to 0, and the lattice parameters including a, b, c, cosab, cosac, cosbc, celldm are removed.
 - use_ibrav = true: ibrab is set to the index of Bravais lattices obtained from the crystal structure data. The Structure data will be reconstructed to match the convention of Quantum ESPRESSO.
- nat (the number of atoms) and ntyp (the number of element types) will be superseded by the values obtained from the crystal structure data.
- The cutoff values ecutwfc and ecutrho are obtained from the pseudo-potential files if these parameters are left blank.

CELL PARAMETERS

- This block will not be generated if use_ibrav is set to true. Otherwise, the lattice vectors are exported in units of angstrom.
- The information of the lattice vectors are obtained from the crystal structure data. When the data field is defined and contains a 3x3 matrix, that value will be used for the set of lattice vectors instead.

ATOMIC SPECIES

- This block exports a list of atom species, atomic mass, and the file name of the pseudo-potential data.
- The information of the atoms are obtained from the crystal structure data. The file names of the pseudo-potential data are referred from the CSV-formatted index file specified by pp_list parameter.
- When the data field is defined and contains the required data, these values will be used instead.

ATOMIC POSITIONS

- This block exports the atomic species and their fractional coordinates.
- When ignore_species is given to specify an atomic species or a list of species, the values of if_pos for these species will be set to 0. It is used for MD or structure relaxations.
- When the data field is defined and contains the required data, these values will be used instead.

K POINTS

- This block exports the information of k points. The type of the output is specified by the option parameter that takes one of the following:
 - gamma: uses Γ point.
 - crystal: generates a list of k points in mesh pattern. The mesh width is given by the grid parameter, or derived from the vol_density or k_resolution parameters.
 - automatic: generates a mesh of k points. It is given by the grid parameter, or derived from the vol_density or k_resolution parameters. The shift is obtained from the kshifts parameter.
- The mesh width is determined in the following order:
 - the grid parameter, specified by a list of n_x, n_y, n_z , or a scalar value n. For the latter, $n_x = n_y = n_z = n$ is assumed.
 - derived from the vol_density parameter.
 - derived from the k_resolution parameter, whose default value is 0.15.
- When the data field is defined and contains the required data, these values will be used.

5.3 Parameters for VASP

The entries of optional section and content part of the tasks section specific to VASP are explained below.

5.3.1 optional

The type and the location of pseudo-potential files are specified.

According to pymatgen, the pseudo-potential files are obtained from PMG_VASP_PSP_DIR/functional/POTCAR. {element}(.gz) or PMG_VASP_PSP_DIR/functional/{element}/POTCAR, where PMG_VASP_PSP_DIR points to the directory and it is given in the configuration file ~/.config/.pmgrc.yaml or by the environment variable of the same name. functional refers to the type of the pseudo-potential, whose value is predefined as POT_GGA_PAW_PBE, POT_LDA_PAW, etc.

pseudo_functional

This parameter specifies the type of the pseudo-potential. The relation to the *functional* value above is defined in the table of pymatgen, for example, by PBE to POT_GGA_PAW_PBE, or by LDA to POT_LDA_PAW, or in a similar manner.

When the pseudo_dir parameter is specified, it is used as the directory that holds the pseudo-potential files, ignoring the convention of pymatgen.

```
psuedo_dir
```

This parameter specifies the directory that holds the pseudo-potential files. The paths to the pseudo-potential file turn to pseudo_dir/POTCAR.{element}(.gz), or pseudo_dir/{element}/POTCAR.

5.3.2 tasks

The template files are assumed to be placed in the directory specified by the template_dir parameter by the names INCAR, KPOINTS, POSCAR, and POTCAR. The missing files will be ignored.

5.3.3 content

incar

• This block contains parameters described in the INCAR file

kpoints

• type

The type parameter describes how KPOINTS are specified. The following values are allowed, with some types accepting parameters. See pymatgen.io.vasp manual for further details.

```
    automatic
        parameter: grid
    gamma_automatic
        parameter: grid, shift
    monkhorst_automatic
        parameter: grid, shift
    automatic_density
        parameter: kppa, force_gamma
    automatic_gamma_density
        parameter: grid_density
    automatic_density_by_vol
        parameter: grid_density, force_gamma
    automatic_density_by_lengths
        parameter: length_density, force_gamma
```

- automatic_linemode

parameter: division, path_type (corresponding to the path_type parameter of High-SymmKpath.)

5.4 Parameters for OpenMX

The entries of optional section and content part of the tasks section specific to OpenMX are explained below.

5.4.1 optional

data_path

This parameter specifies the name of directory that holds files for pseudo-atomic orbitals and pseudo-potentials. It corresponds to the DATA.PATH parameter.

5.4.2 content

precision

This parameter specifies the set of pseudo-atomic orbitals listed in Tables 1 and 2 of Section 10.6 of the OpenMX manual. It is one of quick, standard, or precise. The default value is quick.

Extension guide

6.1 Adding modes of Quantum ESPRESSO

In order to add supports to modes of Quantum ESPRESSO, the mapping between the modes and the transformation classes should be added to create_modeproc() function in src/cif2x/qe/calc_mode.py.

```
def create_modeproc(mode, qe):
   if mode in ["scf", "nscf"]:
       modeproc = QEmode_pw(qe)
   else:
       modeproc = QEmode_generic(qe)
   return modeproc
```

The transformation functionality for each mode is provided as a derived class of QEmode_base class. This class implements methods update_namelist() for updating the namelist block, and update_cards() for generating data of card blocks. In the current version, two classes are provided: QEmode_pw class for scf and nscf calculations of pw.x, and QEmode_generic class for generating output as-is.

```
class QEmode_base:
    def __init__(self, qe):
    def update_namelist(self, content):
    def update_cards(self, content):
```

For the namelist, the transformation class generates values for blank entries from crystal structure data and other sources. It may also force to set values such as the lattice parameters that are determined from the crystal structure data, or those that must be specified consistently with other parameters. The functions are provided for each mode separately.

For card blocks, a function is provided for each card, and the mapping between the card type and the function is given in the card_table variable. The method update_cards() in the base class picks up and runs the function associated to the card, and updates the content of the card. Of course, a new update_cards() function may be defined.

```
self.card_table = {
   'CELL_PARAMETERS': generate_cell_parameters,
   'ATOMIC_SPECIES': generate_atomic_species,
   'ATOMIC_POSITIONS': generate_atomic_positions,
   'K_POINTS': generate_k_points,
}
```

The functions for cards are gathered in src/cif2x/qe/cards.py with the function names as generate_{card name}. These functions takes parameters for card blocks as argument, and returns a dictionary containing the card name, the options, and the data field.