
easyInterface

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EXAMPLE GALLERIES

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easyInterface is a library to interface crystallographic calculators to front end applications, jupyter notebooks and scripting interfaces.

The code of the project is on Github: [easyInterface](#)

FEATURES OF EASYINTERFACE

easyInterface is a way of storing information about crystal structures, providing commonly used functions in an easy to use package. The data structure interfaces to crystallographic libraries, making a common way to calculate observable phenomena regardless of your choice of backend calculator. Currently we support:

- [Cryspy](#) - a crystallographic library for neutron data analysis.

With more interfaces coming.

PROJECTS USING EASYINTERFACE

easyInterface is currently being used in the following projects:

- [easyDiffraction](#) - Scientific software for modelling and analysis of neutron diffraction data

INSTALLATION

3.1 Install via pip

You can do a direct install via pip by using:

```
$ pip install easyInterface
```

3.2 Install as an easyInterface developer

You can get the latest development source from our [Github repository](#). You need `setuptools` installed in your system to install `easyInterface`. For example, you can do:

```
$ git clone https://github.com/easyDiffraction/easyInterface
$ cd easyInterface
$ pip install -r requirements.txt
$ pip install -e .
```

3.3 Main Contents

3.3.1 Introduction to easyInterface

Here we can see some examples of `easyInterface` in action

Note: Click [here](#) to download the full example code or to run this example in your browser via Binder

Creating a QT interface

This demonstrates an example of how to load an example and create a QT interface to a `crispy` calculator. Information about the project is then displayed.

```
# import os
#
# from easyInterface.Utils.Helpers import getExamplesDir
# from easyInterface.Diffraction.Calculators import CrispyCalculator
# from easyInterface.Diffraction.QtInterface import QtCalculatorInterface
```

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```
#
# data_dir = getExamplesDir()
# main_rcif = os.path.join(data_dir, 'Fe3O4_powder-1d_neutrons-pol_5C1(LLB)', 'main.
↪cif')
# calculator = CryspyCalculator(main_rcif)
#
# interface = QtCalculatorInterface(calculator, None)
#
# print(interface.project_dict)
#
# print(interface.phasesIds())
#
# print(interface.getPhase(interface.phasesIds()[0]))
```

Total running time of the script: (0 minutes 0.000 seconds)

3.3.2 Scripted Examples

This section gathers examples which don't produce any figures. These examples show the basic features of easyInterface.

Note: Click [here](#) to download the full example code or to run this example in your browser via Binder

Creating a interface

This demonstrates an example of how to load an example and create a interface to a cryspy calculator. Information about the project is then displayed.

```
import os
from easyInterface.Uutils.Helpers import getExamplesDir
from easyInterface.Diffraction.Calculators import CryspyCalculator
from easyInterface.Diffraction.Interface import CalculatorInterface

data_dir = getExamplesDir()
main_rcif = os.path.join(data_dir, 'Fe3O4_powder-1d_neutrons-pol_5C1(LLB)', 'main.cif
↪')
calculator = CryspyCalculator(main_rcif)

interface = CalculatorInterface(calculator)

print(interface.project_dict)

print(interface.phasesIds())

print(interface.getPhase(interface.phasesIds()[0]))
```

Total running time of the script: (0 minutes 0.000 seconds)

Note: Click [here](#) to download the full example code or to run this example in your browser via Binder

Performing a fit

This demonstrates an example of how to load an example and create an interface to a crspsy calculator and then fit a value.

```
import os
from easyInterface.Utils.Helpers import getExamplesDir
from easyInterface.Diffraction.Calculators import CrspsyCalculator
from easyInterface.Diffraction.Interface import CalculatorInterface

data_dir = getExamplesDir()
main_rcif = os.path.join(data_dir, 'PbSO4_powder-1d_neutrons-unpol_D1A(ILL)', 'main.
↪cif')

calculator = CrspsyCalculator(main_rcif)

interface = CalculatorInterface(calculator)

print(interface.project_dict)

phase_ids = interface.phasesIds()

print(phase_ids)

phase = interface.getPhase(phase_ids[0])
phase['phasename'] = 'PbSO5'
interface.addPhase(phase)
interface.removePhase('PbSO5')

phase = interface.getPhase(phase_ids[0])
interface.setPhaseValue(phase_ids[0], ['atoms', 'Pb', 'fract_x'], 0.18)
interface.setPhases(phase)
print(phase)

interface.setPhaseRefine(phase_ids[0], ['atoms', 'Pb', 'fract_x'], True)

calc = interface.getCalculations()
print(calc)

res = interface.refine()

print(res)
```

Total running time of the script: (0 minutes 0.000 seconds)

3.3.3 Interface and Calculators

easyInterface Interface

class easyInterface.Diffraction.Interface.CalculatorInterface (*calculator*)
Interface to calculators in the *easyInterface.Diffraction.Calculator* class.

addExperiment (*experiment*)

Add an experiment to the list of experiments in both the project dict and the calculator.

Parameters **experiment** (*Experiment*) – Experiment object to be added to the system.

Return type NoReturn

addExperimentDefinition (*exp_path*)

Add an experiment to be simulated from a cif file. Note that this will not have any crystallographic phases associated with it.

Parameters **exp_path** (*str*) – Path to a experiment file (*.cif*)

Return type NoReturn

addPhase (*phase*)

Add a new phase from an easyInterface phase object to the list of existing crystal phases in the calculator.

Parameters **phase** (*Phase*) – New phase to be added to the phase list.

Return type NoReturn

addPhaseDefinition (*phase_path*)

Add new phases from a cif file to the list of existing crystal phases in the calculator.

Parameters **phase_path** (*str*) – Path to a phase definition file (*.cif*)

Example:

```
interface = CalculatorInterface(calculator)
phase_path = '~/Experiments/new_phase.cif'
interface.addPhaseDefinition(phase_path)
```

Return type NoReturn

addPhaseToExp (*exp_name, phase_name, scale=0.0*)

Link a phase in the project dictionary to an experiment in the project dictionary. Links in the calculator will also be made.

Parameters

- **exp_name** (*str*) – The name of the experiment
- **phase_name** (*str*) – The name of the phase to be associated with the experiment
- **scale** (*float*) – The scale of the crystallographic phase in the experimental system.

Raises **KeyError** – If the exp_name or phase_name are unknown

Return type NoReturn

asCifDict ()

Converts the project dictionary into a *cif* structure.

Return type *str*

Returns Project dictionary as a string encoded to the cif specification.

asDict ()

Converts the project dictionary info a standard python dictionary. If there is an error then an empty dictionary is returned.

Return type *dict*

Returns Python dictionary of the project dictionary.

canRedo ()

Informs on if the project dictionary can have redo() called. Typically called after an undo function call.

Return type *bool*

Returns Can or Can't redo the project dictionary.

canUndo()

Inform on if the project dictionary can have undo() called.

Return type bool

Returns Can or Can't undo the project dictionary.

clearUndoStack()

Resets the Undo/Redo stack of the project dictionary.

ALL PREVIOUS UNDO/REDO EDITS WILL BE LOST

Return type NoReturn

experimentsCount()

Get the number of experiments in the project dictionary.

Return type int

Returns number of experiments in the project dictionary.

experimentsIds()

Returns labels of the experiments in the project dictionary.

Return type List[str]

property final_chi_square

Calculates the final chi squared of the simulation. Where the final chi squared is the chi squared divided by the number of data points.

Return type float

Returns Final chi squared

getCalculation(calculation_name)

Returns a specified calculation from the project dictionary.

Parameters **calculation_name** (str) – Name of the calculation to be returned.

Raises **KeyError** – If the calculation_name is not known.

Return type Calculation

Returns Calculation requested.

getCalculations()

Returns all calculations in the project dictionary. Calculations will be updated if members of the phases or experiments section of the project dictionary has been modified.

Return type Calculations

Returns Calculations object containing all calculations.

getDictByPath(keys)

Returns an object in the project dictionary by the path to the object.

Parameters **keys** (List[str]) – Path to the object in the project dictionary

Return type Any

Returns Object from the project dictionary.

Raises **KeyError** – The supplied keys do not return an object in the project dictionary

getExperiment (*experiment_name*)

Returns a experiment from the project dictionary by name if one is supplied. If the experiment name is None then all experiments are returned. If the experiment name does not exist KeyError is thrown.

Parameters **experiment_name** (Optional[str]) – Name of the experiment to be returned or None for all experiments

Return type *Experiment*

Returns Copy of the project dictionaries phase object with name experiment_name

Raises **KeyError** – The supplied key is not a valid experiment name

getPhase (*phase_name*)

Returns a phase from the project dictionary by name if one is supplied. If the phase name is None then all phases are returned. If the phase name does not exist KeyError is thrown.

Parameters **phase_name** (Optional[str]) – Name of the phase to be returned or None for all phases

Return type *Phase*

Returns Copy of the project dictionaries phase object with name phase_name

Raises **KeyError** – The supplied key is not a valid phase name

name ()

Returns the name of the current project.

Return type str

Returns Name of the current project

phasesCount ()

Get the number of phases in the project dictionary.

Return type int

Returns number of phases in the project dictionary.

phasesIds ()

Get the labels of the phases in the project dictionary.

Return type List[str]

Returns labels of the phases in the project dictionary.

redo ()

Perform an redo operation on the project dictionary.

Return type NoReturn

refine ()

Perform a refinement on parameters which are marked in the project dictionary. If the refinement fails then only the “refinement_message” will be returned in the results dictionary with an explanation of the error.

Return type dict

Returns Refinement results of the following fields: “num_refined_parameters”, “refinement_message”, “nfev”, “nit”, “njev”, “final_chi_sq”

removeExperiment (*experiment_name*)

Remove a experiment from both the project dictionary and the calculator.

Parameters **experiment_name** (str) – Name of the experiment to be removed.

Return type NoReturn

removePhase (*phase_name*)

Remove a phase of a given name from the dictionary and the calculator object.

Parameters **phase_name** (*str*) – name of the phase to be removed.

Return type NoReturn

removePhaseFromExp (*exp_name, phase_name*)

Remove the link between an experiment and a crystallographic phase. Links in the calculator will also be removed.

Parameters

- **exp_name** (*str*) – The name of the experiment.
- **phase_name** (*str*) – The name of the phase to be removed.

Raises **KeyError** – If the *exp_name* or *phase_name* are unknown

Return type NoReturn

saveCifs (*save_dir*)

Write project cif files (*main.cif*, *experiments.cif* and *phases.cif*) to a user supplied directory. This contains all information needed to recreate the project dictionary.

Parameters **save_dir** (*str*) – Directory to where the project cif files should be saved.

Return type NoReturn

setCalculatorFromProject ()

Resets the project phases and experiments fields of the project dictionary from the calculator.

Return type NoReturn

setDictByPath (*keys, value*)

Set an object in the project dictionary by a key path.

Parameters

- **keys** (*List[str]*) – Path to the object to be modified/created
- **value** (*Any*) – Value to be set at the key path

Return type NoReturn

setExperiment (*experiment*)

Set an experiment to the project dictionary. If an experiment by the same name exists, the necessary changes will be propagated. If it does not exist, then it will be added to the project dictionary.

Parameters **experiment** (*Experiment*) – Experiment container with experimental information

Raises **TypeError** – If the input isn't an *Experiment*.

Return type NoReturn

setExperimentDefinition (*exp_path*)

Set an experiment/s to be simulated from a cif file. Note that this will not have any crystallographic phases associated with it.

Parameters **exp_path** (*str*) – Path to a experiment file (*.cif*)

Return type NoReturn

setExperimentDefinitionFromString (*exp_cif_string*)

Set an experiment/s to be simulated from a string. Note that this will not have any crystallographic phases associated with it.

Parameters `exp_cif_string` (`str`) – String containing the contents of an experiment file (.cif)

Return type `NoReturn`

setExperimentRefine (`experiment`, `key`, `value=True`)

Shortcut for setting the refinement key for items in the experiment list.

Parameters

- **experiment** (`str`) – Name of experiment to be modified
- **key** (`List[str]`) – Location of element to be modified in the named experiment.
- **value** (`bool`) – Should the parameter specified by above be refined

Raises **KeyError** – If experiment is unknown

Return type `NoReturn`

setExperimentValue (`experiment`, `key`, `value`)

Shortcut for setting the value key for items in the experiment list.

Parameters

- **experiment** (`str`) – Name of experiment to be modified
- **key** (`List[str]`) – Location of element to be modified in the named experiment.
- **value** – New value of the parameter specified by above

Raises **KeyError** – If experiment is unknown

setExperiments (`experiments`)

Overwrite all experiments in the project dictionary with supplied experiments.

Parameters **experiments** (`Union[Experiment, Experiments]`) – Experiments container with experimental information

Raises **TypeError** – If the input isn't an *Experiments* or *Experiment*.

setPhase (`phase`)

Modify a phase in the calculator. The phase will be added if it does not currently exist.

Parameters **phase** (*Phase*) – easyInterface phase object to be added.

Raises **TypeError** – If the phase object is not a easyInterface phase object.

Return type `NoReturn`

setPhaseDefinition (`phase_path`)

Parse a phases cif file and replace existing crystal phases

Parameters **phase_path** (`str`) – Path to new phase definition file (.cif)

Example:

```
interface = CalculatorInterface(calculator)
phase_path = '~/Experiments/phases.cif'
interface.setPhaseDefinition(phase_path)
```

Return type `NoReturn`

setPhaseRefine (`phase`, `key`, `value=True`)

Shortcut for setting the refinement key for items in the phase list.

Parameters

- **phase** (`str`) – Name of phase to be modified
- **key** (`List[str]`) – Location of element to be modified in the named phase.
- **value** (`bool`) – Should the parameter specified by above be refined

Raises **KeyError** – If phase is unknown

Return type `NoReturn`

setPhaseValue (*phase, key, value*)

Shortcut for setting the value key for items in the phase list.

Parameters

- **phase** (`str`) – Name of phase to be modified
- **key** (`List[str]`) – Location of element to be modified in the named phase.
- **value** – New value of the parameter specified by above

Raises **KeyError** – If phase is unknown

Return type `NoReturn`

setPhases (*phases*)

Set the phases in the calculator to an easyInterface phases object. If a phase in the supplied phases exists then the phase will be modified, if not, it will be added.

Parameters **phases** (`Union[Phase, Phases]`) – phases to be added to the calculator.

Raises **TypeError** – If the phase object is not a easyInterface phase/phases object or dictionary object.

Return type `NoReturn`

setProjectFromCalculator ()

Sets the project dictionary from the calculator given on initialisation. Calling this function will regenerate the project dictionary and changes may be lost.

Return type `NoReturn`

undo ()

Perform an undo operation on the project dictionary.

Return type `NoReturn`

updateCalculations ()

Calculate all experiments and populate the calculations field in the project dictionary. Note that this will only occur if a member of the phases or experiments section of the project dictionary has been modified since the last call to *updateCalculations*.

Return type `NoReturn`

updateExperiments ()

Synchronise the experiments portion of the project dictionary from the calculator.

Return type `NoReturn`

updatePhases ()

Synchronise the phases in project dictionary by queering the calculator object.

Return type `NoReturn`

writeExpCif (*save_dir*)

Write the *experiments.cif* where all experiments in the project dictionary are saved to file. This includes the instrumental parameters and which phases are in the experiment/s

Parameters **save_dir** (*str*) – Directory to where the experiment cif file should be saved.

Return type NoReturn

writeMainCif (*save_dir*)

Write the *main.cif* where links to the experiments and phases are stored and other generalised project information.

Parameters **save_dir** (*str*) – Directory to where the main cif file should be saved.

Return type NoReturn

writePhaseCif (*save_dir*)

Write the *phases.cif* where all phases in the project dictionary are saved to file. This cif file should be compatible with other crystallographic software.

Parameters **save_dir** (*str*) – Directory to where the phases cif file should be saved.

Return type NoReturn

easyInterfaces Project Dictionary

class easyInterface.Diffraction.Interface.**ProjectDict** (*interface, app, calculator, info, phases, experiments, calculations*)

This class deals with the creation and modification of the main project dictionary.

classmethod **default** ()

Create a default and empty project dictionary

Return type *ProjectDict*

Returns Default project dictionary with undo/redo functionality

classmethod **fromPars** (*experiments, phases, calculations={}*)

Create a main project dictionary from phases and experiments.

Parameters

- **calculations** (Union[*Calculations, Calculation, List[Calculation], None*]) –
- **experiments** (Union[*Experiments, Experiment, List[Experiment]*]) – A collection of experiments to be compared to calculations
- **phases** (Union[*Phases, Phase, List[Phase]*]) – A Collection of crystallographic phases to be calculated

Return type *ProjectDict*

Returns Project dictionary with undo/redo

easyInterface Crspsy Calculator

class easyInterface.Diffraction.Calculators.**CrspsyCalculator** (*main_rcif_path=None*)

addExpDefinitionFromString (*exp_rcif_content*)

Set an experiment/s to be simulated from a string. Note that this will not have any crystallographic phases associated with it.

Parameters **exp_rcif_content** (*str*) – String containing the contents of an experiment file (*.cif*)

Return type NoReturn

addExpsDefinition (*exp_path*)

Add an experiment to be simulated from a cif file. Note that this will not have any crystallographic phases associated with it.

Parameters **exp_path** (*str*) – Path to a experiment file (*.cif*)

Return type NoReturn

addPhaseDefinition (*phases_path*)

Add new phases from a cif file to the list of existing crystal phases in the calculator.

Parameters **phases_path** (*str*) – Path to a phase definition file (*.cif*)

Return type NoReturn

asCifDict ()

...

Return type dict

getCalculations ()

Returns all calculations from the calculator object.

Return type *Calculations*

getExperiments ()

Returns all experiments from the calculator object.

Return type *Experiments*

getPhases ()

Returns all phases from the calculator object.

Return type *Phases*

readPhaseDefinition (*phases_path*)

Parse the relevant phases file and update the corresponding model

Return type Optional[Tuple[*Phase*, Phase]]

refine ()

refinement ...

Return type Tuple[dict, dict]

removeExpsDefinition (*experiment_name*)

Remove a experiment from both the project dictionary and the calculator.

Parameters **experiment_name** (*str*) – Name of the experiment to be removed.

Return type NoReturn

removePhaseDefinition (*phase_name*)

Remove a phase of a given name from the calculator object.

Parameters **phase_name** (*str*) – name of the phase to be removed.

Return type NoReturn

saveCifs (*save_dir*, *filename*='main.cif', *exp_name*='experiments.cif', *phase_name*='phases.cif')

Write project cif files (*main.cif*, *experiments.cif* and *phases.cif*) to a user supplied directory. This contains all information needed to recreate the calculator object.

Parameters

- **phase_name** (*str*) – What to call the phases file.
- **exp_name** (*str*) – What to call the experiments file.
- **filename** (*str*) – What to call the main file.
- **save_dir** (*str*) – Directory to where the main cif file should be saved.

Return type NoReturn

setExperiments (*experiments*)

Set experiments (Experimental data tab in GUI)

Return type NoReturn

setExpsDefinition (*exp_path*)

Set an experiment/s to be simulated from a cif file. Note that this will not have any crystallographic phases associated with it.

Parameters **exp_path** (*str*) – Path to a experiment file (.cif)

Return type NoReturn

setObjFromProjectDicts (*phases*, *experiments*)

Set all the cryspy parameters from project dictionary

Return type NoReturn

setPhaseDefinition (*phases_path*)

Parse a phases cif file and replace existing crystal phases

Parameters **phases_path** (*str*) – Path to new phase definition file (.cif)

Return type NoReturn

setPhases (*phases*)

Set phases (sample model tab in GUI)

Return type NoReturn

writeExpCif (*save_dir*, *exp_name*='experiments.cif')

Write the *experiments.cif* where all experiments in the calculator are saved to file. This includes the instrumental parameters and which phases are in the experiment/s

Parameters

- **exp_name** (*str*) – What to call the experiments file.
- **save_dir** (*str*) – Directory to where the experiment cif file should be saved.

Return type NoReturn

writeMainCif (*save_dir*, *filename*='main.cif', *exp_filename*='experiments.cif',
phase_filename='phases.cif')

Write the *main.cif* where links to the experiments and phases are stored and other generalised project information.

Parameters

- **phase_filename** (*str*) – What to call the phases file.
- **exp_filename** (*str*) – What to call the experiments file.

- **filename** (str) – What to call the main file.
- **save_dir** (str) – Directory to where the main cif file should be saved.

Return type NoReturn

writePhaseCif (save_dir, phase_name='phases.cif')

Write the *phases.cif* where all phases in the calculator are saved to file. This cif file should be compatible with other crystallographic software.

Parameters

- **phase_name** (str) – What to call the phases file.
- **save_dir** (str) – Directory to where the phases cif file should be saved.

Return type NoReturn

3.3.4 Container Classes

Phase Classes

```
class easyInterface.Diffraction.DataClasses.PhaseObj.Atom.ADP (u_11,      u_22,
                                                             u_33, u_12, u_13,
                                                             u_23)
```

Data store for Atom site anisotropic displacement parameters

```
class easyInterface.Diffraction.DataClasses.PhaseObj.Atom.Atom (atom_site_label,
                                                                type_symbol,
                                                                scat_length_neutron,
                                                                fract_x, fract_y,
                                                                fract_z,      oc-
                                                                cupancy,
                                                                adp_type,
                                                                U_iso_or_equiv,
                                                                ADp, MSp)
```

Storage for details about an atom

```
classmethod default (atom_site_label)
```

Default constructor for an atom given a unique name in the phase

Parameters **atom_site_label** (str) – The atoms unique name in the phase

Return type *Atom*

Returns Default atom with a given name

```
classmethod fromPars (atom_site_label, type_symbol, scat_length_neutron, fract_x, fract_y,
                                                                fract_z, occupancy, adp_type, U_iso_or_equiv, ADp=None, MSp=None)
```

Atom constructor from parameters

Parameters

- **atom_site_label** (str) – The unique name of the atom in the phase
- **type_symbol** (str) – The type of atom
- **scat_length_neutron** (float) – Neutron scattering length
- **fract_x** (float) – X position
- **fract_y** (float) – Y position

- **fract_z** (float) – Z position
- **occupancy** (float) – Site occupancy
- **adp_type** (str) – ADP type code
- **U_iso_or_equiv** (float) – Isotropic atomic displacement parameter

Return type *Atom*

Returns Fully formed atom data store

classmethod **fromXYZ** (*atom_site_label*, *type_symbol*, *x*, *y*, *z*)

Construct an atom from name, type and position

Parameters

- **atom_site_label** (str) – The atoms unique name in the phase
- **type_symbol** (str) – The type of atom
- **x** (float) – X position
- **y** (float) – Y position
- **z** (float) – Z position

Return type *Atom*

Returns Atom with name type and position filled in

class `easyInterface.Diffraction.DataClasses.PhaseObj.Atom.Atoms` (*atoms*)

Container for multiple atoms

class `easyInterface.Diffraction.DataClasses.PhaseObj.Atom.MSP` (*MSPtype*, *chi_11*,
chi_22, *chi_33*,
chi_12, *chi_13*,
chi_23)

Data store for Atom site magnetic susceptibility parameters

class `easyInterface.Diffraction.DataClasses.PhaseObj.Cell.Cell` (*length_a*,
length_b,
length_c, *angle_alpha*,
angle_beta,
angle_gamma)

Container for crystallographic unit cell parameters

classmethod **default** ()

Default constructor for a crystallographic unit cell

Return type *Cell*

Returns Default crystallographic unit cell container

classmethod **fromPars** (*length_a*, *length_b*, *length_c*, *angle_alpha*, *angle_beta*, *angle_gamma*)

Constructor of a crystallographic unit cell when parameters are known

Parameters

- **length_a** (float) – Unit cell length a
- **length_b** (float) – Unit cell length b
- **length_c** (float) – Unit cell length c
- **angle_alpha** (float) – Unit cell angle alpha

- **angle_beta** (float) – Unit cell angle beta
- **angle_gamma** (float) – Unit cell angle gamma

Return type *Cell*

Returns

class easyInterface.Diffraction.DataClasses.PhaseObj.Phase.**Phase** (*name, space-group, cell, atoms, sites*)

Container for crystallographic phase information

classmethod default (*name*)

Default constructor for a crystallographic phase with a given name

Return type *Phase*

Returns Default empty phase with a name

class easyInterface.Diffraction.DataClasses.PhaseObj.Phase.**Phases** (*phases*)

Container for multiple phases

renamePhase (*old_phase_name, new_phase_name*)

Easy method of renaming a phase

Parameters

- **old_phase_name** (str) – phase name to be changed
- **new_phase_name** (str) – new phase name

Return type NoReturn

class easyInterface.Diffraction.DataClasses.PhaseObj.SpaceGroup.**SpaceGroup** (*crystal_system, space_group_name_Hi, space_group_IT_number, origin_choice*)

Data Classes

class easyInterface.Diffraction.DataClasses.DataObj.Calculation.**BraggPeaks** (*bragg_peaks*)

Container for multiple calculations

class easyInterface.Diffraction.DataClasses.DataObj.Calculation.**CalculatedPattern** (*x, y_calc, y_diff_lower, y_diff_upper*)

Storage container for a calculated pattern

class easyInterface.Diffraction.DataClasses.DataObj.Calculation.**Calculation** (*name, bragg_peaks, calculated_pattern, limits*)

Storage container for calculations

class easyInterface.Diffraction.DataClasses.DataObj.Calculation.**Calculations** (*calculations*)

Container for multiple calculations

```
class easyInterface.Diffraction.DataClasses.DataObj.Calculation.CrystalBraggPeaks (name,  
                                                                    h,  
                                                                    k,  
                                                                    l,  
                                                                    ttheta)
```

Generator for HKL reflections and corresponding two theta.

```
class easyInterface.Diffraction.DataClasses.DataObj.Calculation.Limits (y_obs_lower=-  
                                                                    inf,  
                                                                    y_obs_upper=inf,  
                                                                    y_diff_upper=inf,  
                                                                    y_diff_lower=-  
                                                                    inf,  
                                                                    x_calc=None,  
                                                                    y_calc=None)
```

Generator for limits of a dataset

```
class easyInterface.Diffraction.DataClasses.DataObj.Experiment.Background (ttheta,  
                                                                    in-  
                                                                    ten-  
                                                                    sity)
```

Data store for the background data parameters

classmethod default ()

Default constructor for a background point

Return type *Background*

Returns Default background data object

classmethod fromPars (*ttheta*, *intensity*)

Constructor for background when two theta and intensity are known

Parameters

- **ttheta** (float) – Two Theta angle in degrees
- **intensity** (float) – Value for intensity

Return type *Background*

Returns Background data dict

```
class easyInterface.Diffraction.DataClasses.DataObj.Experiment.Backgrounds (backgrounds)  
Store for a collection of background points
```

```
class easyInterface.Diffraction.DataClasses.DataObj.Experiment.Experiment (name,  
                                                                    wave-  
                                                                    length,  
                                                                    off-  
                                                                    set,  
                                                                    phase,  
                                                                    back-  
                                                                    ground,  
                                                                    res-  
                                                                    o-  
                                                                    lu-  
                                                                    tion,  
                                                                    mea-  
                                                                    sured_pattern)
```

Experimental details data container

classmethod default (*name*)

Default constructor for an Experiment

Parameters *name* (*str*) – What the experiment should be called

Return type *Experiment*

Returns Default empty experiment

classmethod fromPars (*name*, *wavelength*, *offset*, *scale*, *background*, *resolution*, *measured_pattern*)

Constructor of experiment from parameters

Parameters

- **name** (*str*) – What the experiment should be called
- **wavelength** (*float*) – Experimental wavelength
- **offset** (*float*) – Experimental offset
- **scale** (*float*) – Scale parameter
- **background** (*Backgrounds*) – Background model
- **resolution** (*Resolution*) – Resolution model
- **measured_pattern** (*MeasuredPattern*) – The Measured data

Return type *Experiment*

Returns Experiment from parameters

class `easyInterface.Diffraction.DataClasses.DataObj.Experiment.ExperimentPhase` (*name*, *scale*)

Storage container for the Experimental Phase details

classmethod default (*name*)

Default experimental phase data container

Return type *ExperimentPhase*

Returns Default experimental phase data container

classmethod fromPars (*name*, *scale*)

Parameter initialised experimental phase data container

Return type *ExperimentPhase*

Returns Set experimental phase data container

class `easyInterface.Diffraction.DataClasses.DataObj.Experiment.ExperimentPhases` (*experiment_phases*)

Storage of multiple phase markers associated with experiments

class `easyInterface.Diffraction.DataClasses.DataObj.Experiment.Experiments` (*experiments*)

Container for multiple experiments

class `easyInterface.Diffraction.DataClasses.DataObj.Experiment.MeasuredPattern` (*x*, *y_obs*, *sy_obs*, *y_obs_up=None*, *sy_obs_up=None*, *y_obs_down=None*, *sy_obs_down=None*)

Storage container for measured patterns

classmethod default (*polarised=False*)

Default constructor for measured data container.

Parameters **polarised** (bool) – Should the container be initialised as a polarised data container?

Returns Empty data container

property isPolarised

Is the measured data of a polarised type?

Return type bool

Returns True if it is from a polarised measurement, false otherwise

property y_obs_lower

Lower data confidence bound.

Return type list

Returns value of lower confidence bound

property y_obs_upper

Upper data confidence bound.

Return type list

Returns value of upper confidence bound

class easyInterface.Diffraction.DataClasses.DataObj.Experiment.**Resolution** (*u*,
v,
w,
x,
y)

Data store for the resolution parameters

classmethod default ()

Default constructor for the resolution dict

Return type *Resolution*

Returns Default resolution dict

classmethod fromPars (*u, v, w, x, y*)

Constructor when resolution parameters are known

Parameters

- **u** (float) – resolution parameter u
- **v** (float) – resolution parameter v
- **w** (float) – resolution parameter w
- **x** (float) – resolution parameter x
- **y** (float) – resolution parameter y

Return type *Resolution*

Returns Resolution dictionary with values set

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