# AFLOWpi Developers Guide Release

**Andrew Supka** 

Contents

Contents:

Contents 1

2 Contents

# plot module

plot.bands (calcs, yLim=[-10, 10], DOSPlot='', LSDA=False, runlocal=False, postfix='', tight banding=False)

Generates electronic band structure plots for the calculations in the dictionary of dictionaries of calculations with the option to have a DOS or PDOS plot to accompany it.

**Parameters** calcs (dict) – dictionary of dictionaries representing the set of calculations

#### **Keyword Arguments**

- yLim (list) List or tuple of two integers for max and min range in horizontal axis of DOS plot
- **LSDA** (*bool*) To plot DOS as spin polarized or not (calculation must have been done as spin polarized)
- **DOSPlot** (*str*) DOS or the PDOS plots next to the eletronic band structure plot (assuming you ran either ppDOS or ppPDOS) (default: NONE)
- **postfix** (str) Postfix to the filename of the plot
- **tight\_banding** (*bool*) Whether to treat the input data as from Quantum Espresso or WanT bands x

#### Returns None

plot.distortionEnergy (calcs1, xaxis, yaxis, zaxis='Energy', calcs=None, colorbarUnits=None, titleArray=None, plotTitle=None, xAxisStr=None, yAxisStr=None, file-Name='distortionEnergy.pdf', percentage=False, runlocal=True)

plot.dos (calcs, yLim=[-10, 10], LSDA=False, runlocal=False)

Generates DOS plots for the calculations in the dictionary of dictionaries of calculations

Parameters calcs (dict) – dictionary of dictionaries representing the set of calculations

# **Keyword Arguments**

- yLim (list) List or tuple of two integers for max and min range in horizontal axis of DOS plot
- LSDA (*bool*) To plot DOS as spin polarized or not (calculation must have been done as spin polarized)
- runlocal (bool) Do the plotting right now or if False do it when the calculations are running
- **postfix** (*str*) Postfix to the filename of the plot
- **tight\_banding** (*bool*) Whether to treat the input data as from Quantum Espresso or WanT bands.x

#### **Returns** None

- plot.grid\_plot (calcs, xaxis, yaxis, zaxis='Energy', colorbarUnits=None, zaxis\_title=None, plot\_title=None, xAxisStr=None, yAxisStr=None, fileName='grid\_plot.pdf', runlo-cal=True)
- plot.interpolatePlot (calcs, variable1, variable2, zaxis='Energy', xaxisTitle=None, yaxisTitle=None, zaxisTitle=None, title=None, fileName='interpolatePlot.pdf', delta=False, text\_min=False, vhline\_min=False, circle\_min=False, delta\_min=True, rel\_center=False, plot\_color='jet', find\_min=False)

Takes a list of calculations and plots the energy of the calculations as a function of two input variables the first value is the baseline for the energy value and the energy plotted is the difference between that energy and the other energies in the grid

#### **Parameters**

- calcs (dict) dictionary of dictionaries of calculations
- variable1 (str) a string of the variable in the calculations that you want as your x axis
- variable2 (str) a string of the variable in the calculations that you want to your y axis

#### **Keyword Arguments**

- **title** (*str*) Title of plot (default: None)
- zaxis (str) Choice out of the keywords in each calc to plot in the Z axis (default: Energy)
- xaxisTitle (str) title of xaxis (default: same as variable1)
- **yaxisTitle** (*str*) title of yaxis (default: same as variable2)
- zaxisTitle (str) title of zaxis (default: same as zaxis)
- **fileName** (*str*) Name (and path where default is directory where script is run from ) of the output file (default: 'interpolatePlot.pdf')
- **delta** (bool) Z-axis scale relative to its first value
- **delta\_min** (bool) Z-axis scale relative to its lowest value
- **text\_min** (*bool*) Display text of minimum value next to the minimum value if find min=True
- **vhline\_min** (*bool*) Display text of minimum value next to the minimum value if find\_min=True
- **circle\_min** (*bool*) Display text of minimum value next to the minimum value if find\_min=True
- **delta\_min** (*bool*) Display text of minimum value next to the minimum value if find min=True
- rel\_center (bool) Display text of minimum value next to the minimum value if find min=True
- **plot\_color** (*str*) string of the matplotlib colormap to be used
- find\_min (bool) Interpolate between points and find value of variable1 and variable2 for the minimum value of Z axis

#### Returns None

plot.interpolatePlot1D (calcs, variable1, yaxis='Energy', xaxisTitle=None, yaxisTitle=None, title=None, fileName='interpolatePlot.pdf', delta=False, circle\_min=False)
Takes a list of calculations and plots the energy of the calculations as a function of two input variables the first value is the baseline for the energy value and the energy plotted is the difference between that energy and the other energies in the grid

#### **Parameters**

- calcs (dict) dictionary of dictionaries of calculations
- variable1 (dict) a string of the variable in the calculations that you want as your x axis

#### **Keyword Arguments**

- yaxis (str) Choice out of the keywords in each calc to plot in the Z axis (default: Energy)
- **xaxisTitle** (*str*) title of xaxis (default: same as variable1)
- yaxisTitle (str) title of yaxis (default: same as yaxis)
- **title** (*str*) Title of plot (default: None)
- **fileName** (*str*) Name (and path where default is directory where script is run from ) of the output file (default: 'interpolatePlot.pdf')
- **delta** (bool) Z-axis scale relative to its first value
- circle\_min (bool) Display text of minimum value next to the minimum value
- plot.opdos (calcs, yLim=[-10, 10], LSDA=False, runlocal=False, postfix='', scale=False)

Generates electronic band structure plots for the calculations in the dictionary of dictionaries of calculations with the option to have a DOS or PDOS plot to accompany it.

Parameters calcs (dict) – dictionary of dictionaries representing the set of calculations

#### **Keyword Arguments**

- yLim (list) List or tuple of two integers for max and min range in horizontal axis of DOS plot
- LSDA (*bool*) To plot DOS as spin polarized or not (calculation must have been done as spin polarized)
- runlocal (bool) Do the plotting right now or if False do it when the calculations are running
- **postfix** (str) Postfix to the filename of the plot
- **tight\_banding** (*bool*) Whether to treat the input data as from Quantum Espresso or WanT bands.x

#### Returns None

```
plot.phonon (calcs, runlocal=False, postfix='', THz=True)
plot.radialPDF (calcs, atomNum, filterElement=None, runlocal=False, title='', **kwargs)
    kwargs get passed to pyplot.hist
plot.read_transport_datafile(ep_data_file, mult_x=1.0, mult_y=1.0)
    Arguments:
    Keyword Arguments:
    Returns:
```

plot.transport\_plots (calcs, runlocal=False, postfix='')

# prep module

```
prep.ConfigSectionMap (section, option, configFile=None)
prep.__null__(*args, **kwargs)
prep.addToBlockWrapper (oneCalc, ID, block, addition)
```

Wraps AFLOWpi.prep.\_\_addToBlock for use inside \_calcs\_container methods

#### **Parameters**

- oneCalc (dict) a dictionary containing properties about the AFLOWpi calculation
- ID (str) ID string for the particular calculation and step
- **block** (*str*) string of the block in the \_ID.py that the addition is to be added to for that step of workflow
- addition (str) a string containing code to be written to the specific block in the \_ID.py for each calculation

#### Returns None

# prep.askAFLOWpiVars (refAFLOWpiVars)

Cycle on the keys of the refAFLOWpiVars dictionary and ask to define them

**Parameters refAFLOWpiVars** (*dict*) – the variables in the ref files that you need to input to run the calculation

#### Returns None

```
prep.bands (calcs, dk=None, nk=None)
```

Wrapper function to write the function AFLOWpi.prep.\_\_oneBands to the \_ID.py

**Parameters** calcs (*dict*) – a dictionary of dicionaries representing the set of calculations

#### **Keyword Arguments**

- dk (float) distance between points for Electronic Band Structure calculation
- **nk** (*int*) approximate number of k points to be calculated along the path

**Returns** The identical "calcs" input variable

# prep.bandsAflow (dk, LAT)

Query aflow for band structure path and generate the path for band structure calculation

#### **Parameters**

- dk (float) distance between k points along path in Brillouin Zone
- LAT (int) bravais lattice number from Quantum Espresso convention

#### **Keyword Arguments None**

**Returns info** – some information nks (str): number of k points in path stringk (str): kpoint path string

Return type str

```
prep.build_calcs (PARAM_VARS, build_type='product')
```

prep.calcFromFile (aflowkeys, fileList, reffile=None, pseudodir=None, build=True, workdir=None, keep name=False, clean input=True)

Reads in a string of an QE input file path, a string of an QE input, a file object of a QE input or a list of them and attempts to fill create a calculation from them. If they are missing things such as k\_points card, they are automtically generated.

#### **Parameters**

- aflowkeys (dict) a dictionary generated by AFLOWpi.prep.init
- **fileList** (*list*) a string of an QE input file path, a string of an QE input, a file object of a QE input or a list of them

# **Keyword Arguments**

- **reffile** (*str*) a partially filled QE input file used in case the input(s) in fileList are missing. i.e. wfc cutoff. If the names of the Pseudopotential files are not included in the input(s) in fileList, they are chosen depending on the pseudodir chosen and included when the calculation set is formed.
- workdir (*str*) a string of the workdir path that be used to override what is in the config file used when initating the AFLOWpi session
- **pseudodir** (*str*) a string of the pseudodir path that be used to override what is in the config file used when initating the AFLOWpi session
- build (bool) < DEFUNCT OPTION. NEEDS REMOVAL>

class prep.calcs\_container (dictionary)

```
addToAll (block=None, addition=None)
```

bands (dk=None, nk=100)

Wrapper method to write call AFLOWpi.prep.bands for calculating the Electronic Band Structure.

**Parameters** calcs (dict) – a dictionary of dicionaries representing the set of calculations

# **Keyword Arguments**

- **dk** (*float*) the density in the Brillouin zone of the k point sampling along the entirety of the path between high symmetry points.
- **nk** (*int*) the approximate number of sampling points in the Brillouin Zone along the entirety of the path between high symmetry points. Points are chosen so that they are equidistant along the entirety of the path. The actual number of points will be slightly different than the inputted value of nk. nk!=None will override any value for dk.

#### Returns None

```
change_input (namelist=None, parameter=None, value=None)
change_pseudos (directory)
conventional_cell_input()
```

Wrapper method to call AFLOWpi.pseudo.crawlingMinimization in the high level user interface. Adds a new step to the workflow.

Parameters self - the \_calcs\_container object

#### **Keyword Arguments**

- mult\_jobs (*bool*) if True split the individual scf jobs into separate cluster jobs if False run them serially
- **grid\_density** (*int*) controls the number of calculations to generate for the minimization num\_{calcs}=grid\_density^{num\_{parameters}-num\_{constraints}}
- initial\_variance (float) amount to vary the values of the parameters from the initial value. i.e. (0.02 = +/-2% variance)
- **thresh** (*float*) threshold for \$DeltaX\$ of the lattice parameters between brute force minimization iterations.
- **constraint** (*list*) a list or tuple containing two entry long list or tuples with the first being the constraint type and the second the free parameter in params that its constraining for example in a orthorhombic cell: constraint=(["volume",'c'],) allows for A and B to move freely but C is such that it keeps the cell volume the same in all calculations generated by the input oneCalc calculation.
- **final\_minimization** (*str*) calculation to be run at the end of the brute force minimization options include "scf", "relax", and "vcrelax"

#### Returns None

```
dos (kpFactor=2, project=True)
```

Wrapper method to call AFLOWpi.prep.doss in the high level user interface. Adds a new step to the workflow.

Parameters self - the \_calcs\_container object

# **Keyword Arguments**

- kpFactor (float) factor to which the k-point grid is made denser in each direction
- project (bool) if True: do the projected DOS after completing the DOS

# Returns None

```
evCurve_min (pThresh=25, final_minimization='relax')
get_initial_inputs()
increase_step(func)
items()
items()
keys()
new_step (update_positions=True, update_structure=True, new_job=True, ext='')
phonon (nrx1=2, nrx2=2, nrx3=2, innx=2, de=0.01, mult_jobs=False, raman=False, LOTO=False, disp_sym=True, atom_sym=True, field_strength=0.01)
pseudo_test_brute(ecutwfc, dual=[], sampling=[], conv_thresh=0.01, constraint=None, initial_relax=None, min_thresh=0.01, initial_variance=0.05, grid_density=7, mult_jobs=False, options=None)
```

```
relax()
resubmit (reset=True)
scf()
scfuj (thresh=0.1, nIters=20, paodir=None, relax='scf', mixing=0.0)
     Wrapper method to call AFLOWpi.scfuj.scfPrep and AFLOWpi.scfuj.run in the high level user interface.
     Adds a new step to the workflow.
         Parameters self - the _calcs_container object
         Keyword Arguments
              • thresh (float) – threshold for self consistent hubbard U convergence
              • niters (int) – max number of iterations of the acbn0 cycle
              • paodir (string) – the path of the PAO directory. This will override an entry of the paodir
                in the AFLOWpi config file used for the session
              • mixing (float) – the amount of the previous acbn0 U iteration to mix into the current (only
                needed when there is U val oscillation)
         Returns None
split_chain (update_positions=True, update_structure=True)
submit()
transport (temperature=[300], epsilon=True, run_bands=True)
     Wrapper method to call AFLOWpi.scfuj.prep_transport and AFLOWpi.scfuj.run_transport in the high
     level user interface. Adds a new step to the workflow.
```

Parameters self - the \_calcs\_container object

#### **Keyword Arguments**

- epsilon (bool) if True episilon tensor will be computed
- temperature (list) list of temperature(s) at which to calculate transport properties

#### Returns None

```
values()
     vcrelax()
prep.changeCalcs (calcs, keyword='calculation', value='scf')
     A Wrapper function that writes the function AFLOWpi.prep.__changeCalcs to the __ID.py
```

**Parameters** calcs (dict) – a dictionary of dicionaries representing the set of calculations

#### **Keyword Arguments**

- **keyword** (str) a string which signifies the type of change that is to be made
- value the value of the choice.

**Returns** The identical set of calculations as the input to this function

```
prep.cleanCalcs (calcs, runlocal=False)
```

Wrapper function for AFLOWpi.prep.\_\_cleanCalcs

**Parameters** calcs (dict) – a dictionary of dicionaries representing the set of calculations

**Keyword Arguments runlocal** (bool) – a flag to choose whether or not to run the wrapped function now or write it to the \_ID.py to run during the workflow

#### Returns None

prep.construct\_and\_run (\_\_submitNodeName\_\_, oneCalc, ID, build\_command='', subset\_tasks=[], fault\_tolerant=False, mult\_jobs=True, subset\_name='SUBSET', keep\_file\_names=False, clean\_input=True)

this is a check to see if we're restarting when mult\_jobs==True

prep.doss(calcs, kpFactor=1.5)

Wrapper function to write the function AFLOWpi.prep.\_\_oneDoss to the \_ID.py

**Parameters** calcs (dict) – a dictionary of dicionaries representing the set of calculations

**Keyword Arguments kpFactor** (*float*) – the factor to which we make each direction in the kpoint grid denser

**Returns** The identical "calcs" input variable

#### prep.extractvars(refFile)

Read refFile and return an empty dictionary with all the keys and None values

Parameters refFile (str) – filename of the reference input file for the frame

**Returns** A dictionary containing the keyword extracted from the reference input file as keys with None for values..

#### prep.generateAnotherCalc (old, new, calcs)

Modify the calculation in each subdir and update the master dictionary

#### **Parameters**

- old (str) string to replace
- **new** (*str*) replacement string
- calcs (dict) dictionary of dictionaries of calculations

Returns A new set of calculations with a new ID of the hash of the new input strings

```
prep.getMPGrid (primLatVec, offset=True, string=True)
```

from\_file (fileList, reffile=None, pseudodir=None, build=True, workdir=None)

Reads in a string of an QE input file path, a string of an QE input, a file object of a QE input or a list of them and attempts to fill create a calculation from them. If they are missing things such as k\_points card, they are automtically generated.

#### **Parameters**

- **aflowkeys** (*dict*) a dictionary generated by AFLOWpi.prep.init
- **fileList** (*str*) a string of an QE input file path, a string of an QE input, a file object of a QE input or a list of them

# **Keyword Arguments**

- **reffile** (*str*) a partially filled QE input file used in case the input(s) in fileList are missing. i.e. wfc cutoff. If the names of the Pseudopotential files are not included in the input(s) in fileList, they are chosen depending on the pseudodir chosen and included when the calculation set is formed.
- workdir (str) a string of the workdir path that be used to override what is in the config file used when initating the AFLOWpi session

- **pseudodir** (*str*) a string of the pseudodir path that be used to override what is in the config file used when initating the AFLOWpi session
- build (bool) <DEFUNCT OPTION. NEEDS REMOVAL>

```
items()
iteritems()
keys()
load(step=1)
    Loads the calc logs from a given step
```

**Parameters** step (*int*) – the step of the calculation for whose calclogs are to be loaded

**Returns** calcs – the loaded calc logs

Return type dict

scfs (allAFLOWpiVars, refFile, name='first', pseudodir=None, build\_type='product', build=True,
run=True)

A wrapper method to call AFLOWpi.prep.scfs to form the calculation set. This will also create directory within the set directory for every calculation in the set.

#### **Parameters**

- **allAFLOWpiVars** (*dict*) a dictionary whose keys correspond to the keywords in the reference input file and whose values will be used to construct the set of calculations
- **refFile** (*str*) a filename as a string, a file object, or a string of the file that contains keywords to construct the inputs to the different calculations in the set

#### **Keyword Arguments**

- **pseudodir** (*str*) path of the directory that contains your Pseudopotential files The value in the AFLOWpi config file used will override this.
- build\_type (str) how to construct the calculation set from allAFLOWpiVars dictionary:

#### zip | The first calculation takes the first entry from the list of

each of the keywords. The second calculation takes the second and so on. The keywords for all lists in allAFLOWpiVars must be the same length for this method.

# product | Calculation set is formed via a "cartesian product" with

the values the list of each keyword combined. (i.e if allAFLOWpiVars has one keyword with a list of 5 entires and another with 4 and a third with 10, there would be 2000 calculations in the set formed from them via product mode.

- build (bool) < DEFUNCT OPTION. NEEDS REMOVAL>
- run (bool) <DEFUNCT OPTION. NEEDS REMOVAL>

**Returns** A dictionary of dictionaries containing the set of calculations.

```
status (status={}, step=0, negate_status=False)
Loads the calc logs from a given step
```

**Parameters** step (*int*) – The step of the calculation for whose calclogs are to be loaded. If no step is specified then it will default to load calculations from all steps with the chosen status.

# **Keyword Arguments**

- status (dict) key, value pairs for status type and their value to filter on. i.e. status={'Finished':False}
- **negate\_status** (*bool*) filter on the opposite of the status filters

**Returns** calcs – the loaded calcs for one or more steps with the given status

Return type dict

#### values()

# Parameters

- **PROJECT** (str) Name of project
- **SET** (*str*) Name of set
- author (str) Name of author
- **CORRESPONDING** (*str*) Name of corresponding
- **SPONSOR** (*str*) Name of sponsor

e.g. initFrame('LNTYPE','', 'MF', 'marco.fornari@cmich.edu','DOD-MURI'). Return the AFLOKEYS dictionary.

#### prep.line\_prepender (filename, new\_text)

prepends a file with a new line containing the contents of the string new\_text.

#### **Parameters**

- **filename** (*str*) string of the filename that is to be prepended
- new\_text (str) a string that is one line long to be prepended to the file

**Returns** None

```
prep.loadlogs (PROJECT='', SET='', logname='', config=None)
```

#### prep.lockAtomMovement (calcs)

A Wrapper function that writes the function AFLOWpi.prep.\_\_freezeAtoms to the \_\_ID.py

Parameters calcs (dict) – a dictionary of dicionaries representing the set of calculations

Returns None

#### prep.maketree (calcs, pseudodir=None, build=True, workdir=None)

Make the directoy tree and place in the input file there

#### Parameters calcs (dict) -

Dictionary of dictionaries of calculations

#### **Keyword Arguments**

- **pseudodir** (str) path of pseudopotential files directory
- workdir (*str*) a string of the workdir path that be used to override what is in the config file used when initating the AFLOWpi session
- build (bool) <DEFUNCT OPTION. NEEDS REMOVAL>

#### Returns None

#### prep.modifyCalcs (old, new, calcs)

Modify the calculation in each subdir and update the master dictionary

#### **Parameters**

- old (str) string to replace
- **new** (*str*) replacement string
- calcs (dict) dictionary of dictionaries of calculations

#### prep.modifyInputPrefixPW(calcs, pre)

A Wrapper function that is used to write a function to the \_\_ID.py

**Parameters** calcs (*dict*) – a dictionary of dicionaries representing the set of calculations

**Returns** None

# prep.modifyNamelistPW(calcs, namelist, parameter, value, runlocal=False)

A Wrapper function that is used to write the function AFLOWpi.prep.\_\_modifyNameListPW to the \_ID.py. If the value is intended to be a string in the QE input file, it must be dually quoted i.e. value=""scf" will become 'scf' in the input file.

#### **Parameters**

- calcs (dict) a dictionary of dicionaries representing the set of calculations
- namelist (str) a string of the fortran namelist that the parameter is in
- parameter (str) a string of the parameter name
- **value** the value of that parameter

**Keyword Arguments runlocal** (*bool*) – a flag to choose whether or not to run the wrapped function now or write it to the \_ID.py to run during the workflow.

**Returns** Either the identical set of calculations if runlocal == False or the set of calculations with the parameter's value changed in their oneCalc['\_AFLOWPI\_INPUT\_'] if runlocal==True

# prep.newstepWrapper(pre)

A function that wraps another function that is to be run before the a certain function runs. Its use must be in the form:

@newstepwrapper(func) def being\_wrapped(oneCalc,ID,\*args,\*\*kwargs)

where func is the function that is to be run before and being\_wrapped is the function being wrapped. oneCalc and ID must be the first two arguments in the function being wrapped. additional arguments and keyword arguments can follow.

**Parameters** pre (func) – function object that is to be wrapped before another function runs

**Returns** the returned values of function being wrapped or the execution of the function is skipped entirely.

# class prep.plotter(calcs)

Class for adding common plotting functions from AFLOWpi.plot module to the high level user interface.

bands (yLim=[-10, 10], DOSPlot='', LSDA=False, runlocal=False, postfix='')

Wrapper method to call AFLOWpi.plot.bands in the high level user interface.

Parameters self - the plotter object

# **Keyword Arguments**

• yLim (*list*) – a tuple or list of the range of energy around the fermi/Highest occupied level energy that is to be included in the plot.

• **DOSPlot** (*str*) – a string that flags for the option to have either a DOS plot share the Y-axis of the band structure plot.

Options include: "" | A blank string (default) will cause No Density of

States plotted alongside the Band Structure

"APDOS" | Atom Projected Density of States "DOS" | Normal Density of States

- LSDA (bool) Plot the up and down of a spin polarized orbital projected DOS calculation.
- **runlocal** (*bool*) a flag to choose whether or not to run the wrapped function now or write it to the \_ID.py to run during the workflow
- **postfix** (*str*) a string of an optional postfix to the plot filename for every calculation.

#### Returns None

dos (yLim=[-10, 10], LSDA=False, runlocal=False, postfix='')

Wrapper method to call AFLOWpi.plot.dos in the high level user interface.

Parameters self – the plotter object

#### **Keyword Arguments**

- yLim (*list*) a tuple or list of the range of energy around the fermi/Highest occupied level energy that is to be included in the plot.
- LSDA (bool) Plot the up and down of a spin polarized DOS calculation.
- **runlocal** (*bool*) a flag to choose whether or not to run the wrapped function now or write it to the \_ID.py to run during the workflow
- postfix (str) a string of an optional postfix to the plot filename for every calculation.

#### Returns None

opdos (yLim=[-10, 10], LSDA=False, runlocal=False, postfix='')

Wrapper method to call AFLOWpi.plot.opdos in the high level user interface.

Parameters self - the plotter object

#### **Keyword Arguments**

- yLim (*list*) a tuple or list of the range of energy around the fermi/Highest occupied level energy that is to be included in the plot.
- LSDA (bool) Plot the up and down of a spin polarized orbital projected DOS calculation.
- **runlocal** (*bool*) a flag to choose whether or not to run the wrapped function now or write it to the \_ID.py to run during the workflow
- **postfix** (*string*) a string of an optional postfix to the plot filename for every calculation.

#### Returns None

phonon (runlocal=False, postfix='', THz=True)

transport (runlocal=False, postfix='')

Wrapper method to call AFLOWpi.plot.epsilon in the high level user interface.

Parameters self - the plotter object

**Keyword Arguments** 

- **nm** (bool) whether to plot in nanometers for spectrum or eV for energy
- **runlocal** (*bool*) a flag to choose whether or not to run the wrapped function now or write it to the \_ID.py to run during the workflow

#### Returns None

```
prep.prep_split_step(calcs, subset_creator, subset_tasks=[], mult_jobs=False, sub-
step_name='SUBSET', keep_file_names=False, clean_input=True)
```

#### prep.remove\_blank\_lines(inp\_str)

Removes whitespace lines of text

**Parameters** inp\_str (str) – input string of text

Returns the same string with blank lines removed

#### prep.runAfterAllDone (calcs, command, faultTolerant=True)

Adds a command to the BATCH command block at the end of each calculation's \_ID.py for all calculations in the set. Used to execute a command over all calculations in particular step have completed.

**Parameters** calcs (dict) – a dictionary of dicionaries representing the set of calculations

# **Keyword Arguments**

- **command** (*str*) the text to be added to the BATCH block
- **faultTolerant** (*bool*) a flag to choose if we return True if some of the calculations ran but did not complete successful

#### Returns None

prep.scfs (aflowkeys, allAFLOWpiVars, refFile, pseudodir=None, build\_type='product', build=True)

Read a reference input file, and construct a set of calculations from the allAFLOWpiVars dictionary defining values for the keywords in the reference input file. This will also create directory within the set directory for every calculation in the set.

# **Parameters**

- **allAFLOWpiVars** (*dict*) a dictionary whose keys correspond to the keywords in the reference input file and whose values will be used to construct the set of calculations
- **refFile** (*str*) a filename as a string, a file object, or a string of the file that contains keywords to construct the inputs to the different calculations in the set

#### **Keyword Arguments**

- **pseudodir** (*str*) path of the directory that contains your Pseudopotential files The value in the AFLOWpi config file used will override this.
- **build\_type** (*str*) how to construct the calculation set from allAFLOWpiVars dictionary:

# zip | The first calculation takes the first entry from the list of

each of the keywords. The second calculation takes the second and so on. The keywords for all lists in allAFLOWpiVars must be the same length for this method.

#### product | Calculation set is formed via a "cartesian product" with

the values the list of each keyword combined. (i.e if allAFLOWpiVars has one keyword with a list of 5 entires and another with 4 and a third with 10, there would be 2000 calculations in the set formed from them via product mode.

• build (bool) – <DEFUNCT OPTION. NEEDS REMOVAL>

**Returns** A dictionary of dictionaries containing the set of calculations.

prep.totree (tobecopied, calcs, rename=None, symlink=False)

Populate all the subdirectories for the calculation with the file in input

#### **Parameters**

- **tobecopied** (*str*) filepath to be copied to the AFLOWpi directory tree
- calcs (dict) Dictionary of dictionaries of calculations

#### **Keyword Arguments**

- **rename** (*bool*) option to rename the file/directory being moves into the AFLOWpi directory tree
- **symlink** (*bool*) whether to copy the data to the AFLOWpi directory tree or to use symbolic links

#### Returns None

#### prep.unlockAtomMovement (calcs)

A Wrapper function that writes the function AFLOWpi.prep.\_unfreezeAtoms to the \_\_ID.py

**Parameters** calcs (*dict*) – a dictionary of dicionaries representing the set of calculations

Returns None

#### prep.updateStructs (calcs, update\_structure=True, update\_positions=True)

A Wrapper function that writes the function AFLOWpi.prep.\_oneUpdateStructs to the \_\_ID.py

Parameters calcs (dict) – a dictionary of dicionaries representing the set of calculations

# **Keyword Arguments**

- **update\_structure** (*bool*) if True update the cell parameter if possible from the output of previous calculations in the workflow.
- **update\_positions** (*bool*) if True update the atomic positions if possible from the output of previous calculations in the workflow.

**Returns** The identical set of calculations as the input to this function

prep.updatelogs (calcs, logname, runlocal=False)

# prep.varyCellParams (oneCalc, ID, param=(), amount=0.15, steps=8, constraint=None)

Forms and returns a set of calcs with varied cell params must be in A,B,C, and, in degrees,alpha,beta,gamma and then returns it.

#### **Parameters**

- oneCalc (dict) a dictionary containing properties about the AFLOWpi calculation
- ID (str) ID string for the particular calculation and step

# **Keyword Arguments**

- param (tuple) the params assoc. with the amount and step. i.e. ('celldm(1)','celldm(3))
- amount (float) percentage amount to be varied up and down. i.e (0.04,0.02,0.01)
- steps (int) how many steps to within each range. i.e (4,5,7)
- **constraint** (*list*) a list or tuple containing two entry long list or tuples with the first being the constraint type and the second the free parameter in params that its constraining for example in a orthorhombic cell: constraint=(["volume", 'c'],) allows for A and B to move

freely but C is such that it keeps the cell volume the same in all calculations generated by the input oneCalc calculation.

**Returns** A dictionary of dictionaries representing a new calculation set

prep.writeToScript (executable, calcs, from\_step=0)

Generates calls on several functions to set up everything that is needed for a new step in the workflow. The mechanics of the \_ID.py are written to it here.

#### **Parameters**

- calcs (dict) dictionary of dictionaries of calculations
- executable (str) < DEFUNCT OPTION: HERE FOR LEGACY SUPPORT>
- \*args <DEFUNCT OPTION: HERE FOR LEGACY SUPPORT>

Keyword Arguments \*\*kwargs - <DEFUNCT OPTION: HERE FOR LEGACY SUPPORT>

**Returns** A set of calculations for a new step in the workflow

# pseudo module

```
pseudo.brute_test(calcs, ecutwfc, dual=None, sampling=None, constraint=None, thresh=0.001, initial_variance=0.05, grid_density=10, mult_jobs=False, conv_thresh=0.01, calc_type='relax')
```

pseudo.crawlingMinimization (calcs, options=None, faultTolerant=True, constraint=None, thresh=0.001, initial\_variance=0.05, grid\_density=10, mult\_jobs=False, final\_minimization='relax')

pseudo.getMinimization(origCalcs, fitVars=None, options=None, runlocal=False, faultTolerant=True, minimize\_var='Energy')

pseudo.plot (resultList, xaxis='', xtitle=None, ytitle=None, title=None, rename=None, file\_name='') takes in a list of dictionaries of dictionaries of calculations and generates a plot with the x axis being some value in the list 'key' and splits the calculations and plots them with each plot being a unique combination of the items in key that are not 'xaxis'

#### **Parameters**

- resultList (list) list of dictionaries of dictionaries of calculations
- xaxis (str) the keyword in oneCalc that you choose to be the x axis in your plots

# **Keyword Arguments**

- **xtitle** (*str*) title of the x axis of the plot (default: None)
- ytitle (str) title of the y axis of the plot (default: None)
- **plotTitle** (*str*) title of the plots (default: None)
- **rename** (*dict*) a mapping of the names of the keywords of whose values used to generate the plot to some other name. ex. { 'AFLOWPI ECUTW ':'wavefunction cutoff'}
- file\_name (str) use this instead of "PT\_RESULTS.pdf" as filename of plot

# Returns None

# retr module

 $\verb|retr.abc2celldm| (a=None, b=None, c=None, alpha=None, beta=None, gamma=None, ibrav=None)| \\ Convert a,b,c,alpha,beta,gamma into celldm for QE| \\$ 

#### Parameters None -

#### **Keyword Arguments**

- a (float) length of a
- **b** (*float*) length of b
- c (float) length of c
- alpha (float) Angle between axis b and c
- beta (float) Angle between axis a and c
- gamma (*float*) Angle between axis a and b
- **ibrav** (*int*) ibrav to be used to convert for defaults

**Returns** celldm\_array – an array of (ibrav,celldm(1),celldm(2),celldm(3),celldm(4),celldm(5),celldm(6))

# Return type array

 $\texttt{retr.abc2free} \ (a=None, b=None, c=None, alpha=None, beta=None, gamma=None, ibrav=None, return-String=True)$ 

Converts a,b,c,alpha,beta,gamma into QE convention primitive lattice vectors

#### Parameters None -

#### **Keyword Arguments**

- a (float) length of a
- **b** (*float*) length of b
- $\mathbf{c}$  (float) length of  $\mathbf{c}$
- alpha (float) Angle between axis b and c
- beta (float) Angle between axis a and c
- gamma (float) Angle between axis a and b
- **ibrav** (*int*) bravais lattice type by QE convention
- returnString (bool) return vectors as a string or as a numpy matrix

**Returns** Either a numpy.matrix or a string of the primitive lattice vectors generated from the celldm input

retr.abcVol (a=None, b=None, c=None, alpha=None, beta=None, gamma=None, ibrav=None)
Get volume from a,b,c,alpha,beta,gamma

#### Parameters None -

#### **Keyword Arguments**

- **a** (*float*) length of a
- $\mathbf{b}$  (float) length of b
- $\mathbf{c}$  (float) length of  $\mathbf{c}$
- alpha (*float*) Angle between axis b and c
- beta (float) Angle between axis a and c
- gamma (*float*) Angle between axis a and b
- **ibrav** (*int*) ibrav to be used to convert for defaults

**Returns** cell\_vol – volume of the cell

Return type float

#### retr.attachPosFlags (positionString, flagString)

Reattaches the flags to the end of the atomic position

#### **Parameters**

- **positionString** (*str*) string of the atomic positions
- **flagString** (*str*) string of the flags

#### **Keyword Arguments None**

Returns positionString – Positions with flags attached

Return type str

retr.celldm2abc (ibrav=None, celldm1=None, celldm2=None, celldm3=None, celldm4=None, celldm5=None, celldm6=None, cosine=True, degrees=False)

Convert celldm for espresso into A,B,C,and angles alpha,beta,gamma

Parameters cellparamatrix (numpy.matrix) - matrix of cell vectors

#### **Keyword Arguments**

- **cosine** (*bool*) If True alpha,beta,gamma are cos(alpha),cos(beta),cos(gamma),
- degrees (bool) If True return alpha, beta, gamma in degrees; radians if False

**Returns paramArray** – a list of the parameters a,b,c,alpha,beta,gamma generated from the input matrix

#### Return type list

retr.celldm2free (ibrav=None, celldm1=None, celldm2=None, celldm3=None, celldm4=None, celldm5=None, celldm6=None, returnString=True)

Converts QE's celldm format into QE convention primitive lattice vectors

# Parameters None -

#### **Keyword Arguments**

- **ibrav** (*int*) bravais lattice type by QE convention
- **celldm1** (*float*) a
- celldm2 (float) b/a

- **celldm3** (*float*) c/a
- celldm4 (float) Cosine of the angle between axis b and c
- celldm5 (float) Cosine of the angle between axis a and c
- celldm6 (float) Cosine of the angle between axis a and b
- returnString (bool) return vectors as a string or as a numpy matrix

**Returns** Either a numpy.matrix or a string of the primitive lattice vectors generated from the celldm input

retr.celldm2params (a, b, c, alpha, beta, gamma, k, v)

DEFUNCT <Marked for removal>

Arguments:

**Keyword Arguments:** 

Returns:

retr.checkStatus(PROJECT, SET='', config='', step=0, status={}, negate\_status=False)

function that loads the calclogs of each of the calculations in your run and displays status information about them.

**Parameters PROJECT** (*str*) – Project name

**Keyword Arguments** 

- **SET** (*str*) Set name
- config (str) Config file used
- step (int) Which number step to seee (default all of them)
- **status** (*dict*) dictionary containing the status and the value you want to see (ex. {'Error':True})
- negate\_status (bool) Whether to take the calculations with that status or without it

**Returns** calcsList (list) list of each step of the calculations you selected that satisfay the status criteria

#### retr.chemAsKeys(calcs)

For sets of calcs that only differ by chemistry you can replace the hash by the chemical name. May not always work. Especially if there are two compounds with swapped positions of atoms of different elements

Parameters calcs (dict) – dictionary of dictionaries of calculations

**Keyword Arguments None** 

**Returns calcsCopy** – returns new dictionay with keys as the chemical stoiciometry

Return type dict

retr.compMatrices (matrix1, matrix2)

Compare two numpy matrices to see if they are equal or not

#### **Parameters**

- matrix1 (numpy.matrix) first matrix
- matrix2 (numpy.matrix) second matrix

**Keyword Arguments None** 

**Returns** equalBool – True if they're equal False if they're not

#### Return type bool

#### retr.convertFCC(oneCalc, ID)

#### **Parameters**

- oneCalc (dict) a dictionary containing properties about the AFLOWpi calculation
- ID (str) ID string for the particular calculation and step

**Keyword Arguments:** 

Returns:

#### retr.detachPosFlags (positionString)

Detach the position flags from the string of the atomic positions

**Parameters** positionString (str) – string of the atomic positions

**Keyword Arguments None** 

Returns positions – atomic positions as a list flags (list): a list of the flags for by each position

Return type list

retr.free2abc(cellparamatrix, cosine=True, degrees=True, string=True, bohr=False)

Convert lattice vectors to a,b,c,alpha,beta,gamma of the primitive lattice

Parameters cellparamatrix (numpy.matrix) - matrix of cell vectors

#### **Keyword Arguments**

- **cosine** (*bool*) If True alpha,beta,gamma are cos(alpha),cos(beta),cos(gamma),
- degrees (bool) If True return alpha, beta, gamma in degrees; radians if False
- string (bool) If True return a,b,c,alpha,beta,gamma as a string; if False return as a list
- bohr (bool) If True return a,b,c in bohr radii; if False return in angstrom

**Returns paramArray** – a list of the parameters a,b,c,alpha,beta,gamma generated from the input matrix

Return type list

retr.free2ibrav(cellparamatrix, ibrav=0, primitive=True)

Convert lattice vectors to celldm

Parameters cellparamatrix (numpy.matrix) - matrix of cell vectors

#### **Keyword Arguments**

- **ibrav** (*int*) Overrides the bravais lattice automatically detected (must be in QE convention if primitive)
- **primitive** (bool) If True it will treat cellparamatrix as primitive lattice vectors

Returns ibravStr - string of the celldm used for QE pwscf input

Return type str

#### retr.getBravaisLatticeName (bravaisLatticeNumber)

Returns the name of the crystal system from the bravias lattice number

**Parameters** braviasLatticeNumber (int) – the number of the bravais lattice in QE convention

**Keyword Arguments None** 

**Returns** A string of the name of the crystal system

#### retr.getCellMatrixFromInput(inputString, string=False, scale=True)

Get the primitive cell vectors from the input.

#### **Parameters**

- oneCalc (dict) a dictionary containing properties about the AFLOWpi calculation
- ID (str) ID string for the particular calculation and step

# **Keyword Arguments**

- string (bool) Return a string or matrix
- scale (bool) scale the vectors by alat

**Returns cellParamMatrix** – primitive lattice params

**Return type** numpy.matrix

#### retr.getCellOutput (oneCalc, ID)

retreives information about the structure and its chemistry and prints it into a file in the AFLOWpi folder inside the project/set directories

#### **Parameters**

- oneCalc (dict) a dictionary containing properties about the AFLOWpi calculation
- ID (str) ID string for the particular calculation and step

#### **Keyword Arguments None**

**Returns outputStr** – Output of the report for the calculation

Return type str

#### retr.getCellVolume (oneCalc, ID, conventional=True, string=True)

Gets the cell volume from output if avaiable and if not gets it from the input

#### **Parameters**

- oneCalc (dict) a dictionary containing properties about the AFLOWpi calculation
- ID (str) ID string for the particular calculation and step

#### **Keyword Arguments**

- conventional (bool) return the volume of the primitive or conventional lattice
- string (bool) to return as a string or as a float

Returns vol – Volume of the cell

**Return type** str

# retr.getCellVolumeFromVectors (cellInput)

Calculates cell volume from basis vectors

**Parameters** cellInput (numpy.matrix) – basis set defining the cell

#### **Keyword Arguments None**

**Returns vol** – volume of the cell (may be not scaled. it depends on your input matrix)

Return type float

# $\verb"retr.getForce" (one Calc, ID")$

Gets last entry of the total force in the calculation from the engine output.

#### **Parameters**

- oneCalc (dict) a dictionary containing properties about the AFLOWpi calculation
- ID (str) ID string for the particular calculation and step

# **Keyword Arguments None**

**Returns force\_string** – Total force from engine output

Return type str

# retr.getIbravFromVectors(cellVectors)

Arguments:

**Keyword Arguments:** 

Returns:

# retr.getPointGroup (oneCalc, ID, source='input')

NOT COMPLETED.

#### **Parameters**

- oneCalc (dict) a dictionary containing properties about the AFLOWpi calculation
- ID (str) ID string for the particular calculation and step

**Keyword Arguments source** (*string*) – either 'input' or 'output' to get point group from input or output atomic positions

Returns None

# retr.getPositionsFromInput(oneCalc, ID)

#### **Parameters**

- oneCalc (dict) a dictionary containing properties about the AFLOWpi calculation
- ID (str) ID string for the particular calculation and step

**Keyword Arguments:** 

Returns:

#### retr.getPositionsFromOutput (oneCalc, ID)

Get the atomic positions from the output. If atomic positions can not be read from output then the positions from the input are returned.

#### **Parameters**

- oneCalc (dict) a dictionary containing properties about the AFLOWpi calculation
- ID (str) ID string for the particular calculation and step

#### **Keyword Arguments None**

**Returns pos** – atomic positions

Return type numpy.matrix

#### retr.getRecipParams(oneCalc)

reads the output from the SCF or relax calculation to get the reciprocal cell parameters produced by the calculation.

Parameters oneCalc (dict) – a dictionary of a single calculation

**Keyword Arguments None** 

```
Returns alat – alat multiplier paramMatrix (numpy.matrix): matrix of reciprocal lattice vectors Return type float
```

```
retr.get_parameters (oneCalc, ID, conventional=True)
retr.glideXshiftY (symMatrix, cellMatrix, shift)
retr.glideXshiftZ (symMatrix, cellMatrix, shift)
retr.glideYshiftX (symMatrix, cellMatrix, shift)
retr.glideYshiftZ (symMatrix, cellMatrix, shift)
retr.glideZshiftX (symMatrix, cellMatrix, shift)
retr.glideZshiftY (symMatrix, cellMatrix, shift)
```

retr.grabEnergy (oneCalc, ID)

Grabs energy for oneCalc and adds the keyword 'Energy' with the value of the energy grabbed from the output

#### **Parameters**

- oneCalc (dict) a dictionary containing properties about the AFLOWpi calculation
- ID (str) ID string for the particular calculation and step

# **Keyword Arguments None**

#### Returns

oneCalc – a dictionary containing properties about the AFLOWpi calculation with the 'Energy' keyword added

Return type dict

# retr.grabEnergyOut (calcs)

Goes in every subdirectory of the calculation and searches for the final energy of the calculation and returns a new copy of the input dictionary that includes the final energy.

Parameters calcs (dict) - dictionary of dictionaries of calculations

#### **Keyword Arguments None**

**Returns** calcs – dictionary of dictionaries of calculations with energy added

Return type dict

```
retr.ibrav2String(ibrav=None, celldm1=None, celldm2=None, celldm3=None, celldm4=None, celldm5=None, celldm6=None)
```

DEFUNCT < CONSIDER FOR REMOVAL>

Arguments:

**Keyword Arguments:** 

Returns:

# retr.inputDict2params (inputDict)

Arguments:

**Keyword Arguments:** 

Returns:

retr.invertX (symMatrix, cellMatrix)

#### **Parameters**

- **symMatrix** (*numpy.matrix*) atomic positions in matrix form
- cellMatrix (numpy.matrix) primitive lattice vectors in matrix form

**Keyword Arguments:** 

Returns:

retr.invertXYZ (symMatrix, cellMatrix)

#### **Parameters**

- **symMatrix** (*numpy.matrix*) atomic positions in matrix form
- **cellMatrix** (*numpy.matrix*) primitive lattice vectors in matrix form

**Keyword Arguments:** 

Returns:

retr.invertY (symMatrix, cellMatrix)

#### **Parameters**

- **symMatrix** (*numpy.matrix*) atomic positions in matrix form
- cellMatrix (numpy.matrix) primitive lattice vectors in matrix form

**Keyword Arguments:** 

Returns:

retr.invertZ (symMatrix, cellMatrix)

#### **Parameters**

- **symMatrix** (*numpy.matrix*) atomic positions in matrix form
- cellMatrix (numpy.matrix) primitive lattice vectors in matrix form

**Keyword Arguments:** 

Returns:

retr.pw2cif (calcs, inpt=True, outp=True, runlocal=False, outputFolder=None, filePrefix='')
Writes a simple CIF file for engine input and outputs for viewing the structure

**Parameters** calcs (*dict*) – dictionary of dictionaries of calculations

#### **Keyword Arguments**

- inpt (bool) Do CIF for input
- **outp** (*bool*) Do CIF for output
- **outputFolder** (str) Output directory for the CIF
- **filePrefix** (*str*) Optional prefix to the CIF filenames

**Returns** None

retr.rotateAlpha(symMatrix, cellMatrix, angle)

#### **Parameters**

- **symMatrix** (*numpy.matrix*) atomic positions in matrix form
- cellMatrix (numpy.matrix) primitive lattice vectors in matrix form

**Keyword Arguments:** 

Returns:

retr.rotateBeta(symMatrix, cellMatrix, angle)

#### **Parameters**

- **symMatrix** (*numpy.matrix*) atomic positions in matrix form
- cellMatrix (numpy.matrix) primitive lattice vectors in matrix form

**Keyword Arguments:** 

Returns:

retr.rotateGamma (symMatrix, cellMatrix, angle)

#### **Parameters**

- **symMatrix** (*numpy.matrix*) atomic positions in matrix form
- cellMatrix (numpy.matrix) primitive lattice vectors in matrix form

**Keyword Arguments:** 

Returns:

retr.shiftCell(symMatrix)

Parameters symMatrix (numpy.matrix) – atomic positions in matrix form

**Keyword Arguments:** 

Returns:

retr.shiftX(symMatrix, cellMatrix, shift)

#### **Parameters**

- **symMatrix** (*numpy.matrix*) atomic positions in matrix form
- cellMatrix (numpy.matrix) primitive lattice vectors in matrix form

**Keyword Arguments:** 

Returns:

retr.shiftY(symMatrix, cellMatrix, shift)

#### **Parameters**

- **symMatrix** (*numpy.matrix*) atomic positions in matrix form
- cellMatrix (numpy.matrix) primitive lattice vectors in matrix form

**Keyword Arguments:** 

Returns:

 $\verb"retr.shiftZ" (symMatrix, cellMatrix, shift)$ 

#### **Parameters**

- **symMatrix** (*numpy.matrix*) atomic positions in matrix form
- cellMatrix (numpy.matrix) primitive lattice vectors in matrix form

**Keyword Arguments:** 

Returns:

retr.transform\_input\_conv(oneCalc, ID)

#### **Parameters**

- oneCalc (dict) a dictionary containing properties about the AFLOWpi calculation
- ID (str) ID string for the particular calculation and step

**Keyword Arguments:** 

Returns:

retr.writeInputFromOutput (calcs, replace=False, runlocal=False)

Parameters calcs (dict) – dictionary of dictionaries of calculations

# **Keyword Arguments**

- replace (bool) If true replace the input with updated atomic positions and cell parameters
- runlocal (bool) run local or write to <ID>.py

Returns None

# run module

run.addatexit\_\_\_(command, \*args, \*\*kwargs)

Wrapper to add function to be run at exit

#### **Parameters**

- **command** (*func*) function to be run
- \*args arguments for command

Keyword Arguments \*\*kwargs – Keyword arguments for command

Returns None

run . bands (calcs, engine='', execPrefix=None, execPostfix=' ', holdFlag=True, config=None)
Wrapper to set up Electronic Band Structure calculation

**Parameters** calcs (*dict*) – Dictionary of dictionaries of calculations

#### **Keyword Arguments**

- **engine** (*str*) executable that you are calling to run the calculations
- **execPrefix** (*str*) commands to go before the executable when run (ex. mpiexec nice -n 19 <executable>) (default = None)
- **execPostfix** (*str*) commands to go after the executable when run (ex. <execPrefix> <executable> -ndiag 12 -nimage 2) (default = None)
- holdFlag (bool) DEFUNCT. NEEDS REMOVAL
- config (str) DEFUNCT. NEEDS REMOVAL

#### **Returns** None

# run.clean\_cell\_params(output)

Parses the atomic shifts in a supercell from fd.x outputted pw.x input files to correct for formatting issues when they are imported to AFLOWpi

**Parameters** output (*str*) – pw.x input files generated by fd.x

**Keyword Arguments None** 

**Returns output** – pw.x input files generated by fd.x (cleaned by AFLOWpi)

Return type str

#### run.cleanup(calcs)

Deletes all files a calculation set's directory tree that are prepended with '\_'

Parameters calcs (dict) – Dictionary of dictionaries of calculations

#### **Keyword Arguments None**

Returns None

run. dos (calcs, engine='', execPrefix=None, execPostfix=None, holdFlag=True, config=None)
Wrapper to set up DOS nscf calculation

**Parameters** calcs (*dict*) – Dictionary of dictionaries of calculations

#### **Keyword Arguments**

- engine (str) executable that you are calling to run the calculations
- **execPrefix** (*str*) commands to go before the executable when run (ex. mpiexec nice -n 19 < executable>) (default = None)
- **execPostfix** (*str*) commands to go after the executable when run (ex. <execPrefix> <executable> -ndiag 12 -nimage 2) (default = None)
- holdFlag (bool) DEFUNCT. NEEDS REMOVAL
- config (str) DEFUNCT. NEEDS REMOVAL

#### Returns None

run . emr (calcs, engine='', execPrefix=None, execPostfix=None, holdFlag=True, config=None) Wrapper to set up GIPAW EMR calculation

**Parameters** calcs (dict) – Dictionary of dictionaries of calculations

#### **Keyword Arguments**

- engine (str) executable that you are calling to run the calculations
- **execPrefix** (*str*) commands to go before the executable when run (ex. mpiexec nice -n 19 <executable>) (default = None)
- **execPostfix** (*str*) commands to go after the executable when run (ex. <execPrefix> <executable> -ndiag 12 -nimage 2) (default = None)
- holdFlag (bool) DEFUNCT. NEEDS REMOVAL
- config (str) DEFUNCT. NEEDS REMOVAL

# Returns None

#### run.generateSubRef (qsubRefFileString, oneCalc, ID)

Reads in the reference cluster submission file specified in "jobreffile" in the config used. Tries to insert a few parameters.

OBSOLETE PLANNED FOR REMOVAL

#### **Parameters**

- qsubRefFileString (str) string of the "reference" cluster submission file
- oneCalc (dict) dictionary of one of the calculations
- **ID** (*str*) **ID** of calculation

#### **Keyword Arguments None**

Returns cluster TypeDict - string cluster submission file

# Return type dict

run.gvectors (calcs, engine='', execPrefix=None, execPostfix=None, holdFlag=True, config=None) Wrapper to set up GIPAW gvectors calculation

**Parameters** calcs (*dict*) – Dictionary of dictionaries of calculations

#### **Keyword Arguments**

- engine (str) executable that you are calling to run the calculations
- **execPrefix** (*str*) commands to go before the executable when run (ex. mpiexec nice -n 19 < executable>) (default = None)
- **execPostfix** (*str*) commands to go after the executable when run (ex. <execPrefix> <executable> -ndiag 12 -nimage 2) (default = None)
- holdFlag (bool) DEFUNCT. NEEDS REMOVAL
- config (str) DEFUNCT. NEEDS REMOVAL

#### **Returns** None

run.hyperfine (calcs, engine='', execPrefix=None, execPostfix=None, holdFlag=True, config=None, isotope=())

Wrapper to set up GIPAW hyperfine calculation

Parameters calcs (dict) - Dictionary of dictionaries of calculations

#### **Keyword Arguments**

- engine (str) executable that you are calling to run the calculations
- **execPrefix** (*str*) commands to go before the executable when run (ex. mpiexec nice -n 19 < executable >) (default = None)
- **execPostfix** (*str*) commands to go after the executable when run (ex. <execPrefix> <executable> -ndiag 12 -nimage 2) (default = None)
- holdFlag (bool) DEFUNCT. NEEDS REMOVAL
- config (str) DEFUNCT. NEEDS REMOVAL

#### Returns None

run.nmr (calcs, engine='', execPrefix=None, execPostfix=None, holdFlag=True, config=None)
Wrapper to set up GIPAW NMR calculation

Parameters calcs (dict) – Dictionary of dictionaries of calculations

#### **Keyword Arguments**

- engine (str) executable that you are calling to run the calculations
- **execPrefix** (*str*) commands to go before the executable when run (ex. mpiexec nice -n 19 < executable>) (default = None)
- **execPostfix** (*str*) commands to go after the executable when run (ex. <execPrefix> <executable> -ndiag 12 -nimage 2) (default = None)
- holdFlag (bool) DEFUNCT. NEEDS REMOVAL
- **config** (*str*) DEFUNCT. NEEDS REMOVAL

#### Returns None

run.**pdos** (calcs, engine='', execPrefix=None, execPostfix='', holdFlag=True, config=None) Wrapper to set up DOS projection calculation

**Parameters** calcs (dict) – Dictionary of dictionaries of calculations

# **Keyword Arguments**

• **engine** (*str*) – executable that you are calling to run the calculations

- **execPrefix** (*str*) commands to go before the executable when run (ex. mpiexec nice -n 19 < executable>) (default = None)
- **execPostfix** (*str*) commands to go after the executable when run (ex. <execPrefix> <executable> -ndiag 12 -nimage 2) (default = None)
- holdFlag (bool) DEFUNCT. NEEDS REMOVAL
- config (str) DEFUNCT. NEEDS REMOVAL

#### Returns None

run.prep\_fd(\_\_submitNodeName\_\_, oneCalc, ID, nrx1=2, nrx2=2, nrx3=2, innx=2, de=0.01, atom\_sym=True, disp\_sym=True)

Generates input files for fd.x, fd\_ifc.x, and matdyn.x for the finite difference phonon calculations.

#### **Parameters**

- \_\_submitNodeName\_\_ (str) String of hostname that cluster jobs should be submitted from
- oneCalc (dict) dictionary of one of the calculations
- **ID** (*str*) **ID** of calculation

#### **Keyword Arguments**

- **nrx1** (*int*) supercell size for first primitive lattice vector
- nrx2 (int) supercell size for second primitive lattice vector
- nrx3 (int) supercell size for third primitive lattice vector
- innx (int) how many different shifts in each direction for finite difference phonon calculation
- de (float) amount to shift the atoms for finite differences

#### Returns None

#### run.reset\_logs (calcs)

Removes log files from AFLOWpi directory

Parameters calcs (dict) – Dictionary of dictionaries of calculations

#### **Keyword Arguments None**

Returns None

# run.resubmit(calcs)

Stages loaded calculation set to be resubmitted on a cluster

**Parameters** calcs (*dict*) – Dictionary of dictionaries of calculations

#### **Keyword Arguments None**

Returns None

run.**scf** (calcs, engine='', execPrefix=None, execPostfix=None, holdFlag=True, config=None) Wrapper to set up self-consitent calculation

**Parameters** calcs (*dict*) – Dictionary of dictionaries of calculations

#### **Keyword Arguments**

- engine (str) executable that you are calling to run the calculations
- **execPrefix** (*str*) commands to go before the executable when run (ex. mpiexec nice -n 19 <executable>) (default = None)

- **execPostfix** (*str*) commands to go after the executable when run (ex. <execPrefix> <executable> -ndiag 12 -nimage 2) (default = None)
- holdFlag (bool) DEFUNCT. NEEDS REMOVAL
- config (str) DEFUNCT. NEEDS REMOVAL

#### Returns:

#### run.submit()

sets global \_\_submit\_\_flag\_\_ so calculations will start when user script completes

#### Parameters None -

#### **Keyword Arguments None**

Returns None

# run.submitFirstCalcs\_\_\_(calcs)

Submits the first step of a calculation's pipeline

**Parameters** calcs (dict) – Dictionary of dictionaries of calculations

#### **Keyword Arguments None**

Returns None

run.testOne (calcs, calcType='scf', engine='', execPrefix=None, execPostfix=None, holdFlag=True, config=None)

Run all the calculation in the dictionary with a specific engine

**Parameters** calcs (*dict*) – Dictionary of dictionaries of calculations

#### **Keyword Arguments**

- engine (str) executable that you are calling to run the calculations
- **execPrefix** (*str*) commands to go before the executable when run (ex. mpiexec nice -n 19 <executable>) (default = None)
- **execPostfix** (*str*) commands to go after the executable when run (ex. <execPrefix> <executable> -ndiag 12 -nimage 2) (default = None)

#### Returns None

run.write\_fdx\_template(oneCalc, ID, nrx1=2, nrx2=2, nrx3=2, innx=2, de=0.01, atom\_sym=True, disp\_sym=True)

Generates input files for fd.x, fd\_ifc.x, and matdyn.x for the finite difference phonon calculations.

#### **Parameters**

- oneCalc (dict) dictionary of one of the calculations
- **ID** (*str*) **ID** of calculation

#### **Keyword Arguments**

- nrx1 (int) supercell size for first primitive lattice vector
- nrx2 (int) supercell size for second primitive lattice vector
- nrx3 (int) supercell size for third primitive lattice vector
- innx (int) how many different shifts in each direction for finite difference phonon calculation
- **de** (*float*) amount to shift the atoms for finite differences

#### Returns None

# scfuj module

```
scfuj.WanT_bands (oneCalc, ID=None, eShift=10.0, nbnd=None)
     Make input files for WanT bands calculation
          Parameters calc_copy -- dictionary of dictionaries of calculations (-
scfuj.WanT_dos (oneCalc, ID=None, eShift=10.0, temperature=300.0, energy_range=[-10.0, 10.0])
     Make input files for WanT bands calculation
          Parameters calc_copy -- dictionary of dictionaries of calculations (-
scfuj.WanT_epsilon (oneCalc, ID=None, eShift=10.0, temperature=300.0, energy_range=[0.1, 12.5])
     Make input files for WanT bands calculation
          Parameters calc_copy -- dictionary of dictionaries of calculations (-
scfuj.acbn0 (oneCalc, projCalcID, byAtom=False)
scfuj.checkOscillation(ID, oneCalc, uThresh=0.001)
scfuj.chkSpinCalc(oneCalc, ID=None)
     Check whether an calculation is spin polarized or not.
     Arguments:
     -oneCalc : dictionary of a single calculation.
scfuj.chk_species(elm)
scfuj.evCurveMinimize (calcs, config=None, pThresh=10.0, final_minimization='vc-relax')
scfuj.getU_frmACBN0out (oneCalc, ID, byAtom=False)
scfuj.maketree (oneCalc, ID, paodir=None)
     Make the directoy tree and place in the input file there
          Parameters calcs -- Dictionary of dictionaries of calculations (-) -
          Keyword Arguments
                 • - pseudodir - path of pseudopotential files directory
                 · - paodir - path of pseudoatomic orbital basis set
scfuj.nscf_nosym_noinv (oneCalc, ID=None, kpFactor=1.5)
     Add the nesf input to each subdir and update the master dictionary
```

#### **Parameters**

- calc\_copy -- dictionary of one calculation (-) -
- kpFactor -- multiplicative factor for kpoints from SCF to DOS (default  $(\mbox{-})\,-$

2.

scfuj.projwfc(oneCalc, ID=None)

Run projwfc on each calculation

Parameters oneCalc -- dictionary of a single calculation (-) -

```
scfuj.run (calcs, uThresh=0.001, nIters=20, mixing=0.0)
```

scfuj.scfprep(calcs, paodir=None)

scfuj.transport\_prep(oneCalc, ID)

sets up the environment do do scf->nscf->projwfc to get overlap for transport calcs

scfuj.updateUvals (oneCalc, Uvals, ID=None)

Modify scf input file to do a lda+u calculation.

#### **Parameters**

- oneCalc -- Dictionary of one calculation (-) -
- Uvals -- Dictionary of Uvals (-) -

# CHAPTER 7

# Indices and tables

- genindex
- modindex
- search

# plot, ?? prep, ?? pseudo, ?? retr, ?? run, ?? S scfuj, ??