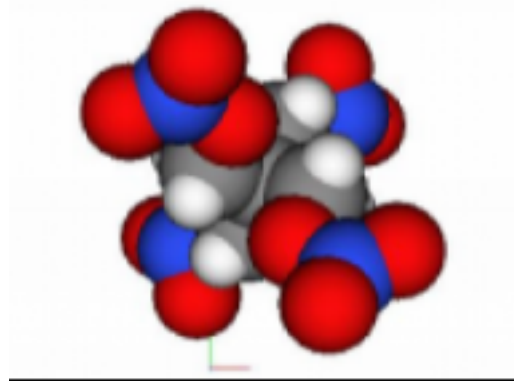

GSAS-2



GSAS-II Developers Documentation

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GSAS-II MAIN MODULE

Main routines for the GSAS-II program

class GSASII.**GSASII** (*parent*)

Define the main GSAS-II frame and its associated menu items

CheckNotebook ()

Make sure the data tree has the minimally expected controls. (BHT) correct?

class **CopyDialog** (*parent, title, text, data*)

Creates a dialog for copying control settings between data tree items

GSASII.**ErrorDialog** (*title, message, parent=None, wtype=4*)

Display an error message

GSASII.**ExitMain** (*event*)

Called if the main window is closed

GSASII.**FillMainMenu** (*menubar*)

Define contents of the main GSAS-II menu for the (main) data tree window in the mac, used also for the data item windows as well.

GSASII.**GetFileList** (*fileType, skip=None*)

Appears unused. Note routine of same name in GSASIIpwdGUI

GSASII.**GetHKLFdatafromTree** (*HKLFname*)

Returns single crystal data from GSASII tree

Parameters **HKLFname** (*str*) – a single crystal histogram name as obtained from
GSASIIstruct.GetHistogramNames ()

Returns HKLFdata = single crystal data list of reflections

GSASII.**GetHistogramNames** (*hType*)

Returns a list of histogram names found in the GSASII data tree Note routine
GSASIIstrIO.GetHistogramNames () also exists to get same info from GPX file.

Parameters **hType** (*str*) – list of histogram types

Returns list of histogram names

GSASII.**GetPWDRdatafromTree** (*PWDRname*)

Returns powder data from GSASII tree

Parameters **PWDRname** (*str*) – a powder histogram name as obtained from
GSASIIstruct.GetHistogramNames ()

Returns PWDRdata = powder data dictionary with Powder data arrays, Limits, Instrument Parameters, Sample Parameters

GSASII.GetPhaseData()

Returns a dict with defined phases. Note routine `GSASIIstrIO.GetPhaseData()` also exists to get same info from GPX file.

GSASII.GetPhaseNames()

Returns a list of defined phases. Note routine `GSASIIstrIO.GetPhaseNames()` also exists to get same info from GPX file.

GSASII.GetPowderIparm(*rd, prevIparm, lastIparmfile, lastdatafile*)

Open and read an instrument parameter file for a data file Returns the list of parameters used in the data tree

Parameters

- **rd** (*obj*) – the raw data (histogram) data object.
- **prevIparm** (*str*) – not used
- **lastIparmfile** (*str*) – Name of last instrument parameter file that was read, or a empty string.
- **lastdatafile** (*str*) – Name of last data file that was read.

Returns a list of two dicts, the first containing instrument parameters and the second used for future TOF datasets (timemaps?)

GSASII.GetUsedHistogramsAndPhasesfromTree()

Returns all histograms that are found in any phase and any phase that uses a histogram. This also assigns numbers to used phases and histograms by the order they appear in the file. Note routine `GSASIIstrIO.GetUsedHistogramsAndPhasesfromTree()` also exists to get same info from GPX file.

Returns

(Histograms,Phases)

- Histograms = dictionary of histograms as {name:data,...}
- Phases = dictionary of phases that use histograms

GSASII.MakeLSParmDict()

Load all parameters used for computation from the tree into a dict of paired values [value, refine flag]. Note that this is different than the parmDict used in the refinement, which only has values.

Returns

(parmDict,varyList) where:

- parmDict is a dict with values and refinement flags for each parameter and
- varyList is a list of variables (refined parameters).

GSASII.OnAddPhase(*event*)

Add a new, empty phase to the tree. Called by Data/Add Phase menu

GSASII.OnDataDelete(*event*)

Delete one or more histograms from data tree. Called by the Data/DeleteData menu

GSASII.OnDeletePhase(*event*)

Delete a phase from the tree. Called by Data/Delete Phase menu

GSASII.OnDummyPowder(*event*)

Called in response to Import/Powder Data/Simulate menu item to create a Dummy powder diffraction data set.

Reads an instrument parameter file and then gets input from the user

GSASII.OnFileClose (*event*)

Clears the data tree in response to the File/New Project menu button. User is given option to save the project.

GSASII.OnFileExit (*event*)

Called in response to the File/Quit menu button

GSASII.OnFileOpen (*event*, *filename=None*)

Reads in a GSAS-II .gpx project file in response to the File/Open Project menu button

GSASII.OnFileSave (*event*)

Save the current project in response to the File/Save Project menu button

GSASII.OnFileSaveas (*event*)

Save the current project in response to the File/Save as menu button

GSASII.OnImageRead (*event*)

Called to read in an image in any known format

GSASII.OnImageSum (*event*)

Sum together image data(?)

GSASII.OnImportGeneric (*reader*, *readerlist*, *label*, *multiple=False*, *usedRanIdList=[]*)

Used to import Phases, powder dataset or single crystal datasets (structure factor tables) using reader objects subclassed from `GSASIIIO.ImportPhase`, `GSASIIIO.ImportStructFactor` or `GSASIIIO.ImportPowderData`. If a reader is specified, only that will be attempted, but if no reader is specified, every one that is potentially compatible (by file extension) will be tried on the selected file(s).

Parameters

- **reader** (*readerobject*) – This will be a reference to a particular object to be used to read a file or None, if every appropriate reader should be used.
- **readerlist** (*list*) – a list of reader objects appropriate for the current read attempt. At present, this will be either `self.ImportPhaseReaderlist`, `self.ImportSfactReaderlist` or `self.ImportPowderReaderlist` (defined in `_init_Imports` from the files found in the path), but in theory this list could be tailored. Used only when reader is None.
- **label** (*str*) – string to place on the open file dialog: Open *label* input file
- **multiple** (*bool*) – True if multiple files can be selected in the file dialog. False is default. At present True is used only for reading of powder data.
- **usedRanIdList** (*list*) – an optional list of random Ids that have been used and should not be reused

Returns a list of reader objects (*rd_list*) that were able to read the specified file(s). This list may be empty.

GSASII.OnImportPhase (*event*)

Called in response to an Import/Phase/... menu item to read phase information. `dict self.ImportMenuId` is used to look up the specific reader item associated with the menu item, which will be None for the last menu item, which is the “guess” option where all appropriate formats will be tried.

GSASII.OnImportPowder (*event*)

Called in response to an Import/Powder Data/... menu item to read a powder diffraction data set. `dict self.ImportMenuId` is used to look up the specific reader item associated with the menu item, which will be None for the last menu item, which is the “guess” option where all appropriate formats will be tried.

Also reads an instrument parameter file for each dataset.

GSASII.OnImportSfact (*event*)

Called in response to an Import/Structure Factor/... menu item to read single crystal datasets. dict self.ImportMenuId is used to look up the specific reader item associated with the menu item, which will be None for the last menu item, which is the “guess” option where all appropriate formats will be tried.

GSASII.OnMakePDFs (*event*)

Calculates PDFs

GSASII.OnPatternTreeItemActivated (*event*)

Called when a tree item is activated

GSASII.OnPatternTreeItemCollapsed (*event*)

Called when a tree item is collapsed

GSASII.OnPatternTreeItemDelete (*event*)

Called when a tree item is deleted – not sure what this does

GSASII.OnPatternTreeItemExpanded (*event*)

Called when a tree item is expanded

GSASII.OnPatternTreeKeyDown (*event*)

Not sure what this does

GSASII.OnPatternTreeSelChanged (*event*)

Called when a data tree item is selected

GSASII.OnPwdrSum (*event*)

Sum together powder data(?)

GSASII.OnReadPowderPeaks (*event*)

Bound to menu Data/Read Powder Peaks – still needed?

GSASII.OnRefine (*event*)

Perform a refinement. Called from the Calculate/Refine menu.

GSASII.OnRenameData (*event*)

Renames an existing phase. Called by Data/Rename Phase menu

GSASII.OnSeqRefine (*event*)

Perform a sequential refinement. Called from the Calculate/Sequential refine menu.

GSASII.OnSize (*event*)

Called when the main window is resized. Not sure why

GSASII.ReadPowderInstprm (*instfile*)

Read a GSAS-II (new) instrument parameter file

Parameters *instfile* (*str*) – name of instrument parameter file

GSASII.ReadPowderIparm (*instfile*, *bank*, *databanks*, *rd*)

Read a GSAS (old) instrument parameter file

Parameters

- **instfile** (*str*) – name of instrument parameter file
- **bank** (*int*) – the bank number read in the raw data file
- **databanks** (*int*) – the number of banks in the raw data file. If the number of banks in the data and instrument parameter files agree, then the sets of banks are assumed to match up and bank is used to select the instrument parameter file. If not, the user is asked to make a selection.
- **rd** (*obj*) – the raw data (histogram) data object. This sets rd.instbank.

GSASII.ShowLSParms (*event*)

Displays a window showing all parameters in the refinement. Called from the Calculate/View LS Parms menu.

class GSASII.SumDialog (*parent, title, text, dataType, data*)

Allows user to supply scale factor(s) when summing data

class GSASII.GSASIImain (*redirect=True, filename=None, useBestVisual=False, clearSigInt=True*)

Defines a wxApp for GSAS-II

Creates a wx frame (self.main) which contains the display of the data tree.

MacOpenFile (*filename*)

Called on Mac every time a file is dropped on the app when it is running, treat this like a File/Open project menu action. Should be ignored on other platforms

OnInit ()

Called automatically when the app is created.

GSASII.main ()

Start up the GSAS-II application

GSASIIOBJ: DATA OBJECTS

This module defines and/or documents the data structures used in GSAS-II, as well as provides misc. support routines.

2.1 Constraints Tree Item

Constraints are stored in a dict, separated into groups. Note that parameter are named in the following pattern, p:h:<var>:n, where p is the phase number, h is the histogram number <var> is a variable name and n is the parameter number. If a parameter does not depend on a histogram or phase or is unnumbered, that number is omitted. Note that the contents of each dict item is a List where each element in the list is a *constraint definition objects*. The constraints in this form are converted in `GSASIIstrIO.ProcessConstraints()` to the form used in `GSASIImapvars`

The keys in the Constraints dict are:

key	explanation
Hist	This specifies a list of constraints on histogram-related parameters, which will be of form :h:<var>:n.
HAP	This specifies a list of constraints on parameters that are defined for every histogram in each phase and are of form p:h:<var>:n.
Phase	This specifies a list of constraints on phase parameters, which will be of form p::<var>:n.
Global	This specifies a list of constraints on parameters that are not tied to a histogram or phase and are of form ::<var>:n

Each constraint is defined as an item in a list. Each constraint is of form:

```
[<mult1>, <var1>], [<mult2>, <var2>], ..., <fixedval>, <varyflag>, <constype>]
```

Where the variable pair list item containing two values [<mult>, <var>], where:

- <mult> is a multiplier for the constraint (float)
- <var> a **G2VarObj** object (previously a str variable name of form 'p:h:name[:at]')

Note that the last three items in the list play a special role:

- <fixedval> is the fixed value for a *constant equation* (constype=c) constraint or is None. For a *New variable* (constype=f) constraint, a variable name can be specified as a str (used for externally generated constraints)
- <varyflag> is True or False for *New variable* (constype=f) constraints or is None. This will be implemented in the future to indicate if these variables should be refined.
- <constype> is one of four letters, 'e', 'c', 'h', 'f' that determines the type of constraint:

- ‘e’ defines a set of equivalent variables. Only the first variable is refined (if the appropriate refine flag is set) and all other equivalent variables in the list are generated from that variable, using the appropriate multipliers.
- ‘c’ defines a constraint equation of form, $m_1 \times var_1 + m_2 \times var_2 + \dots = c$
- ‘h’ defines a variable to hold (not vary). Any variable on this list is not varied, even if its refinement flag is set. Only one [mult,var] pair is allowed in a hold constraint and the mult value is ignored. This is of particular value when needing to hold one or more variables where a single flag controls a set of variables such as, coordinates, the reciprocal metric tensor or anisotropic displacement parameter.
- ‘f’ defines a new variable (function) according to relationship $newvar = m_1 \times var_1 + m_2 \times var_2 + \dots$

2.2 Covariance Tree Item

The Covariance tree item has results from the last least-squares run. They are stored in a dict with these keys:

key	sub-key	explanation
newCellDict		dict with lattice parameters computed by <code>GSASIIstrMath.GetNewCellParms()</code> (dict)
title		Name of gpx file(?) (str)
variables		Values for all N refined variables (list of float values, length N, ordered to match varyList)
sig		Uncertainty values for all N refined variables (list of float values, length N, ordered to match varyList)
varyList		List of directly refined variables (list of str values, length N)
newAtomDict		dict with atom position values computed in <code>GSASIIstrMath.ApplyXYZshifts()</code> (dict)
Rvals		R-factors, GOF, Marquardt value for last refinement cycle (dict)
	Nobs	Number of observed data points (int)
	Rwp	overall weighted profile R-factor (% , float)
	chisq	sum[w*(Iobs-Icalc)**2] for all data note this is not the reduced chi squared (float)
	lamMax	Marquardt value applied to Hessian diagonal (float)
	GOF	The goodness-of-fit, aka square root of the reduced chi squared. (float)
covMatrix		The (NxN) covVariance matrix (np.array)

2.3 Phase Tree Items

Phase information is stored in the GSAS-II data tree as children of the Phases item in a dict with keys:

key	sub-key	explanation
General		Overall information for the phase (dict)
	AtomPtrs	list of four locations to use to pull info from the atom records (list)
	F000X	x-ray F(000) intensity (float)
	F000N	neutron F(000) intensity (float)
	Mydir	directory of current .gpx file (str)
	MCSA controls	Monte Carlo-Simulated Annealing controls (dict)
	Cell	List with 8 items: cell refinement flag (bool) a, b, c, (Angstrom, float) alpha, beta & gamma (degrees, float) volume (A^3, float)
	Type	‘nuclear’ or ‘macromolecular’ for now (str)

Continued on next page

Table 2.1 – continued from previous page

key	sub-key	explanation
	Map	dict of map parameters
	SH Texture	dict of spherical harmonic preferred orientation parameters
	Isotope	dict of isotopes for each atom type
	Isotopes	dict of scattering lengths for each isotope combination for each element in phase
	Name	phase name (str)
	SGData	Space group details as a <i>space group (SGData) object</i> as defined in <code>GSASIIspc.SpcGroup()</code> .
	Pawley neg wt	Restraint value for negative Pawley intensities (float)
	Flip	dict of Charge flip controls
	Data plot type	data plot type ('Mustrain', 'Size' or 'Preferred orientation') for powder data (str)
	Mass	Mass of unit cell contents in g/mol
	POhkl	March-Dollase preferred orientation direction
	Z	dict of atomic numbers for each atom type
	vdWRadii	dict of van der Waals radii for each atom type
	Color	Colors for atoms (list of (r,b,g) triplets)
	AtomTypes	List of atom types
	AtomMass	List of masses for atoms
	doPawley	Flag for Pawley intensity extraction (bool)
	NoAtoms	Number of atoms per unit cell of each type (dict)
	Pawley dmin	maximum Q (as d-space) to use for Pawley extraction (float)
	BondRadii	Default radius for each atom used to compute interatomic distances (list of floats)
	AngleRadii	Default radius for each atom used to compute interatomic angles (list of floats)
	DisAgICtls	Dict with distance/angle search controls, which has keys 'Name', 'AtomTypes', 'BondRadii', 'AngleRadii' which are as above except are possibly edited. Also contains 'Factors', which is a 2 element list with a multiplier for bond and angle search range [typically (0.85,0.85)].
ranId		unique random number Id for phase (int)
pId		Phase Id number for current project (int).
Atoms		Atoms in phase as a list of lists. The outer list is for each atom, the inner list contains varying items depending on the type of phase, see the <i>Atom Records</i> description. (list of lists)
Drawing		Display parameters (dict)
	ballScale	Size of spheres in ball-and-stick display (float)
	bondList	dict with bonds
	contourLevel	map contour level in e/A^3 (float)
	showABC	Flag to show view point triplet (bool). True=show.
	viewDir	cartesian viewing direction (np.array with three elements)
	Zclip	clipping distance in A (float)
	backColor	background for plot as and R,G,B triplet (default = [0, 0, 0], black). (list with three atoms)
	selectedAtoms	List of selected atoms (list of int values)
	showRigidBody	Flag to highlight rigid body placement
	sizeH	Size ratio for H atoms (float)
	bondRadius	Size of binds in A (float)
	atomPtrs	positions of x, type, site sym, ADP flag in Draw Atoms (list)
Continued on next page		

Table 2.1 – continued from previous page

key	sub-key	explanation
	viewPoint	list of lists. First item in list is [x,y,z] in fractional coordinates for the center of the plot. Second item list of previous & current atom number viewed (may be [0,0])
	showHydrogen	Flag to control plotting of H atoms.
	unitCellBox	Flag to control display of the unit cell.
	ellipseProb	Probability limit for display of thermal ellipsoids in % (float).
	vdwScale	Multiplier of van der Waals radius for display of vdW spheres.
	Atoms	A list of lists with an entry for each atom that is plotted.
	Zstep	Step to de/increase Z-clip (float)
	Quaternion	Viewing quaternion (4 element np.array)
	radiusFactor	Distance ratio for searching for bonds. ? Bonds are located that are within $r(R_a+R_b)$ and $(R_a+R_b)/r$ where R_a and R_b are the atomic radii.
	oldxy	previous view point (list with two floats)
	cameraPos	Viewing position in A for plot (float)
	depthFog	True if use depthFog on plot - set currently as False (bool)
RBModels		Rigid body assignments (note Rigid body definitions are stored in their own main top-level tree entry.)
Pawley ref		Pawley reflections
Histograms		A dict of dicts. The key for the outer dict is the histograms tied to this phase. The inner dict contains the combined phase/histogram parameters for items such as scale factors, size and strain parameters. (dict)
MCSA		Monte-Carlo simulated annealing parameters (dict)

2.4 Rigid Body Objects

Rigid body descriptions are available for two types of rigid bodies: ‘Vector’ and ‘Residue’. Vector rigid bodies are developed by a sequence of translations each with a refinable magnitude and Residue rigid bodies are described as Cartesian coordinates with defined refinable torsion angles.

key	sub-key	explanation
Vector	RBIId	vector rigid bodies (dict of dict)
	AtInfo	Drad, Color: atom drawing radius & color for each atom type (dict)
	RBname	Name assigned by user to rigid body (str)
	VectMag	vector magnitudes in Å (list)
	rbXYZ	Cartesian coordinates for Vector rigid body (list of 3 float)
	rbRef	3 assigned reference atom nos. in rigid body for origin definition, use center of atoms flag (list of 3 int & 1 bool)
	VectRef	refinement flags for VectMag values (list of bool)
	rbTypes	Atom types for each atom in rigid body (list of str)
	rbVect	Cartesian vectors for each translation used to build rigid body (list of lists)
	useCount	Number of times rigid body is used in any structure (int)
Residue	RBIId	residue rigid bodies (dict of dict)
	AtInfo	Drad, Color: atom drawing radius & color for each atom type(dict)
	RBname	Name assigned by user to rigid body (str)
	rbXYZ	Cartesian coordinates for Residue rigid body (list of 3 float)
	rbTypes	Atom types for each atom in rigid body (list of str)
	atNames	Names of each atom in rigid body (e.g. C1,N2...) (list of str)
	rbRef	3 assigned reference atom nos. in rigid body for origin definition, use center of atoms flag (list of 3 int & 1 bool)
	rbSeq	Orig,Piv,angle,Riding (list): definition of internal rigid body torsion; origin atom (int), pivot atom (int), torsion angle (float), riding atoms (list of int)
	SelSeq	[int,int] used by SeqSizer to identify objects
	useCount	Number of times rigid body is used in any structure (int)
RBIIds		unique Ids generated upon creation of each rigid body (dict)
	Vector	Ids for each Vector rigid body (list)
	Residue	Ids for each Residue rigid body (list)

2.5 Space Group Objects

Space groups are interpreted by `GSASIIspc.SpcGroup()` and the information is placed in a SGdata object which is a dict with these keys:

key	explanation
SpGrp	space group symbol (str)
Laue	one of the following 14 Laue classes: -1, 2/m, mmm, 4/m, 4/mmm, 3R, 3mR, 3, 3m1, 31m, 6/m, 6/mmm, m3, m3m (str)
SGInv	True if centrosymmetric, False if not (bool)
SGLatt	Lattice centering type. Will be one of P, A, B, C, I, F, R (str)
SGUniq	unique axis if monoclinic. Will be a, b, or c for monoclinic space groups. Will be blank for non-monoclinic. (str)
SGCen	Symmetry cell centering vectors. A (n,3) np.array of centers. Will always have at least one row: <code>np.array([[0, 0, 0]])</code>
SGOps	symmetry operations as a list of form <code>[[M1, T1], [M2, T2], ...]</code> where M_n is a 3x3 np.array and T_n is a length 3 np.array. Atom coordinates are transformed where the Asymmetric unit coordinates [X is (x,y,z)] are transformed using $X' = M_n * X + T_n$
SGSys	symmetry unit cell: type one of 'triclinic', 'monoclinic', 'orthorhombic', 'tetragonal', 'rhombohedral', 'trigonal', 'hexagonal', 'cubic' (str)
SGPolax	Axes for space group polarity. Will be one of '', 'x', 'y', 'x y', 'z', 'x z', 'y z', 'xyz'. In the case where axes are arbitrary '111' is used (P 1, and ?).

2.6 Atom Records

If `phasedict` points to the phase information in the data tree, then atoms are contained in a list of atom records (list) in `phasedict['Atoms']`. Also needed to read atom information are four pointers, `cx, ct, cs, cia = phasedict['General']['atomPtrs']`, which define locations in the atom record, as shown below. Items shown are always present; additional ones for macromolecular phases are marked 'mm'

location	explanation
ct-4	mm - residue number (str)
ct-3	mm - residue name (e.g. ALA) (str)
ct-2	mm - chain label (str)
ct-1	atom label (str)
ct	atom type (str)
ct+1	refinement flags; combination of 'F', 'X', 'U' (str)
cx,cx+1,cx+2	the x,y and z coordinates (3 floats)
cs	site symmetry (str)
cs+1	site multiplicity (int)
cia	ADP flag: Isotropic ('I') or Anisotropic ('A')
cia+1	Uiso (float)
cia+2...cia+6	U11, U22, U33, U12, U13, U23 (6 floats)
atom[-1]	unique atom identifier (int)

2.7 Drawing Atom Records

If `phasedict` points to the phase information in the data tree, then drawing atoms are contained in a list of drawing atom records (list) in `phasedict['Drawing']['Atoms']`. Also needed to read atom information are four pointers, `cx, ct, cs, ci = phasedict['Drawing']['AtomPtrs']`, which define locations in the atom record, as shown below. Items shown are always present; additional ones for macromolecular phases are marked 'mm'

location	explanation
ct-4	mm - residue number (str)
ct-3	mm - residue name (e.g. ALA) (str)
ct-2	mm - chain label (str)
ct-1	atom label (str)
ct	atom type (str)
cx,cx+1,cx+2	the x,y and z coordinates (3 floats)
cs-1	Sym Op symbol; sym. op number + unit cell id (e.g. '1,0,-1') (str)
cs	atom drawing style; e.g. 'balls & sticks' (str)
cs+1	atom label style (e.g. 'name') (str)
cs+2	atom color (RBG triplet) (int)
cs+3	ADP flag: Isotropic ('I') or Anisotropic ('A')
cs+4	Uiso (float)
cs+5...cs+11	U11, U22, U33, U12, U13, U23 (6 floats)
ci	unique atom identifier; matches source atom Id in Atom Records (int)

2.8 Powder Diffraction Tree Items

Every powder diffraction histogram is stored in the GSAS-II data tree with a top-level entry named beginning with the string "PWDR ". The diffraction data for that information are directly associated with that tree item and there are a series of children to that item. The routines `GSASII.GSASII.GetUsedHistogramsAndPhasesfromTree()` and `GSASIIstrIO.GetUsedHistogramsAndPhases()` will load this information into a dictionary where the child tree name is used as a key, and the information in the main entry is assigned a key of `Data`, as outlined below.

key	sub-key	explanation
Limits		A list of two two element lists, as <code>[[Ld,Hd],[L,H]]</code> where <code>Ld</code> is the current and default lowest two-theta value to be used and <code>Hd</code> are the current and default highest two-theta value.
Reflection Lists		A dict with an entry for each phase in the histogram. Each dict item is a dict containing reflections, as described in the Reflections description.
Instrument Parameters		A list containing two dicts where the possible keys in each dict are listed below. The value for each item is a list containing three values: the default value, the current value and a refinement flag which can be True, False or 0 where 0 indicates a value that cannot be refined. The first and second values are floats unless otherwise noted. The first dict are noted as [1]
	Lam	Specifies a wavelength in Angstroms [1]
	Lam1	Specifies the primary wavelength in Angstrom, when an external x-ray source is used [1]
	Lam2 I(L2)/I(L1)	Specifies the secondary wavelength in Angstrom, when an external x-ray source is used [1] Ratio of Lam2 to Lam1 [1]
	Type	Histogram type (str) [1]: <ul style="list-style-type: none"> • ‘PXC’ for constant wavelength x-ray • ‘PNC’ for constant wavelength neutron • ‘PNT’ for time of flight neutron
	Zero	Two-theta zero correction in <i>degrees</i> [1]
	Azimuth	Azimuthal setting angle for data recorded with differing detector geometry [1]
	U, V, W	Cagliotti profile coefficients for Gaussian instrument broadening where the FWHM goes as $U \tan^2 \theta + V \tan \theta + W$ [1]
	X, Y	Cauchy (Lorentzian) instrumental broadening coefficients [1]
	SH/L	Variant of the Finger-Cox-Jephcoat asymmetric peak broadening. Note that this is the average between S/L and H/L where S is the peak height, H is the slit height and L is the goniometer diameter [1]
	Polariz.	Polarization coefficient. [1]
wtFactor		A weighting factor to increase or decrease the leverage of the histogram (float). A value of 1.0 weights the data with their uncertainties and a larger value increases the weighting of the data (to decreasing the uncertainties).
Sample Parameters		Specifies a dict with parameters that describe how the sample was collected, as listed below. Refinable parameters are a list of floats and a bool, where the second value specifies if the value is refinable or otherwise the value is a float unless otherwise noted.
	Scale	The histogram scale factor (refinable)
	Absorption	The sample absorption coefficient as μr where r is the radius [1]
	DisplaceX, DisplaceY	Sample displacement from goniometer center where Y is parallel to the direction and X is perpendicular. Units are μm (refinable)
	Phi, Chi, Omega	Goniometer sample setting angles, in degrees.
	Gonio. radius	Radius of the diffractometer in mm
	InstrName	A name for the instrument, used in preparing a CIF (str).
	Force, Temperature, Humidity, Pressure, Voltage	Variables that describe how the measurement was performed. These are not directly in any computations.
	ranId	The random-number Id for the histogram (same value as the histogram key is ranId)
	Type	Type of diffraction data, may be ‘Debye-Scherrer’ or ‘Bragg’ (str).
	Diffuse	not in use?
14 hId		Chapter 2. GSASIIobj: Data objects The number assigned to the histogram when the project was first edited (can change)
ranId		A random number id for the histogram that does not change
Background		The background is stored as a list with where the first value is the

2.9 Powder Reflection Data Structure

For every phase in a histogram, the `Reflection Lists` value is a dict one element of which is `RefList`, which is a np.array containing reflections. The columns in that array are documented below.

index	explanation
0,1,2	h,k,l (float)
3	multiplicity
4	d-space, Angstrom
5	pos, two-theta
6	sig, Gaussian width
7	gam, Lorentzian width
8	F_{obs}^2
9	F_{calc}^2
10	reflection phase, in degrees
11	intensity correction for reflection, this times F_{obs}^2 or F_{calc}^2 gives Iobs or Icalc

2.10 Single Crystal Tree Items

Every single crystal diffraction histogram is stored in the GSAS-II data tree with a top-level entry named beginning with the string “HKL ”. The diffraction data for that information are directly associated with that tree item and there are a series of children to that item. The routines `GSASII.GSASII.GetUsedHistogramsAndPhasesfromTree()` and `GSASIIstrIO.GetUsedHistogramsAndPhases()` will load this information into a dictionary where the child tree name is used as a key, and the information in the main entry is assigned a key of `Data`, as outlined below.

key	sub-key	explanation
Data		A dict that contains the reflection table, as described in the <i>Single Crystal Reflections</i> description.
Instrument Parameters		A list containing two dicts where the possible keys in each dict are listed below. The value for most items is a list containing two values: the initial value, the current value. The first and second values are floats unless otherwise noted.
	Lam	Specifies a wavelength in Angstroms (two floats)
	Type	Histogram type (two str values): <ul style="list-style-type: none"> • ‘SXC’ for constant wavelength x-ray • ‘SNC’ for constant wavelength neutron • ‘SNT’ for time of flight neutron
	InstrName	A name for the instrument, used in preparing a CIF (str).
wtFactor		A weighting factor to increase or decrease the leverage of data in the histogram (float). A value of 1.0 weights the data with their standard uncertainties and a larger value increases the weighting of the data (equivalent to decreasing the uncertainties).
hId		The number assigned to the histogram when the project is loaded or edited (can change)
ranId		A random number id for the histogram that does not change

2.11 Single Crystal Reflection Data Structure

For every single crystal a histogram, the 'Data' item contains the structure factors as an np.array in item 'RefList'. The columns in that array are documented below.

index	explanation
0,1,2	h,k,l (float)
3	multiplicity
4	d-space, Angstrom
5	F_{obs}^2
6	$\sigma(F_{obs}^2)$
7	F_{calc}^2
8	$F_{obs}^2 T$
9	$F_{calc}^2 T$
10	reflection phase, in degrees
11	intensity correction for reflection, this times F_{obs}^2 or F_{calc}^2 gives Iobs or Icalc

2.12 Classes and routines

`GSASIIobj.AtomIdLookup = {}`

dict listing for each phase index as a str, the atom label and atom random Id, keyed by atom sequential index as a str; best to access this using `LookupAtomLabel()`

`GSASIIobj.AtomRanIdLookup = {}`

dict listing for each phase the atom sequential index keyed by atom random Id; best to access this using `LookupAtomId()`

`GSASIIobj.CompileVarDesc()`

Set the values in the variable description lookup table (`VarDesc`) into `reVarDesc`. This is called in `getDescr()` so the initialization is always done before use.

Note that keys may contain regular expressions, where '[xyz]' matches 'x' 'y' or 'z' (equivalently '[x-z]' describes this as range of values). '.' matches any string. For example:

```
'AUiso': 'Atomic isotropic displacement parameter',
```

will match variable 'p:AUiso:a'. If parentheses are used in the key, the contents of those parentheses can be used in the value, such as:

```
'AU([123][123]): 'Atomic anisotropic displacement parameter U\1',
```

will match AU11, AU23,... and U11, U23 etc will be displayed in the value when used.

`GSASIIobj.DefaultControls = {'F**2': True, 'shift factor': 1.0, 'deriv type': 'analytic Hessian', 'max cyc': 3, 'minF/sig':`

Values to be used as defaults for the initial contents of the `Controls` data tree item.

class `GSASIIobj.G2VarObj(*args)`

Defines a GSAS-II variable either using the phase/atom/histogram unique Id numbers or using a character string that specifies variables by phase/atom/histogram number (which can change). Note that `LoadID()` should be used to (re)load the current Ids before creating or later using the `G2VarObj` object.

A `G2VarObj` object can be created with a single parameter:

Parameters `varname (str/tuple) –`

a single value can be used to create a `G2VarObj` object. If a string, it must be of form "p:h:var" or "p:h:var:a", where

- *p* is the phase number (which may be left blank);
- *h* is the histogram number (which may be left blank);
- *a* is the atom number (which may be left blank in which case the third colon is omitted).

Alternately a single tuple of form (Phase,Histogram,VarName,AtomID) can be used, where Phase, Histogram, and AtomID are None or are ranId values and VarName is a string.

If four positional arguments are supplied, they are:

Parameters

- **phasenum** (*str/int*) – The number for the phase
- **histnum** (*str/int*) – The number for the histogram
- **varname** (*str*) – a single value can be used to create a `G2VarObj`
- **atomnum** (*str/int*) – The number for the atom

varname ()

Formats the GSAS-II variable name as a “traditional” GSAS-II variable string (p:h:<var>:a) or (p:h:<var>)

Returns the variable name as a str

`GSASIIobj.HistIdLookup = {}`

dict listing histogram name and random Id, keyed by sequential histogram index as a str; best to access this using `LookupHistName()`

`GSASIIobj.HistRanIdLookup = {}`

dict listing histogram sequential index keyed by histogram random Id; best to access this using `LookupHistId()`

`GSASIIobj.IndexAllIds (Histograms, Phases)`

Scan through the used phases & histograms and create an index to the random numbers of phases, histograms and atoms. While doing this, confirm that assigned random numbers are unique – just in case lightning strikes twice in the same place.

Note: this code assumes that the atom random Id (ranId) is the last element each atom record.

This is called in two places (only) `GSASIIstrIO.GetUsedHistogramsAndPhases()` (which loads the histograms and phases from a GPX file) and `GSASII.GSASII.GetUsedHistogramsAndPhases()` (which loads the histograms and phases from the data tree.)

TODO: do we need a lookup for rigid body variables?

`GSASIIobj.LookupAtomId (pId, ranId)`

Get the atom number from a phase and atom random Id

Parameters

- **pId** (*int/str*) – the sequential number of the phase
- **ranId** (*int*) – the random Id assigned to an atom

Returns the index number of the atom (str)

`GSASIIobj.LookupAtomLabel (pId, index)`

Get the atom label from a phase and atom index number

Parameters

- **pId** (*int/str*) – the sequential number of the phase

- **index** (*int*) – the index of the atom in the list of atoms

Returns the label for the atom (*str*) and the random Id of the atom (*int*)

`GSASIIobj.LookupHistId (ranId)`

Get the histogram number and name from a histogram random Id

Parameters **ranId** (*int*) – the random Id assigned to a histogram

Returns the sequential Id (*hId*) number for the histogram (*str*)

`GSASIIobj.LookupHistName (hId)`

Get the histogram number and name from a histogram Id

Parameters **hId** (*int/str*) – the sequential assigned to a histogram

Returns (*hist,ranId*) where *hist* is the name of the histogram (*str*) and *ranId* is the random # id for the histogram (*int*)

`GSASIIobj.LookupPhaseId (ranId)`

Get the phase number and name from a phase random Id

Parameters **ranId** (*int*) – the random Id assigned to a phase

Returns the sequential Id (*pId*) number for the phase (*str*)

`GSASIIobj.LookupPhaseName (pId)`

Get the phase number and name from a phase Id

Parameters **pId** (*int/str*) – the sequential assigned to a phase

Returns (*phase,ranId*) where *phase* is the name of the phase (*str*) and *ranId* is the random # id for the phase (*int*)

`GSASIIobj.MakeUniqueLabel (lbl, labellist)`

Make sure that every a label is unique against a list by adding digits at the end until it is not found in list.

Parameters

- **lbl** (*str*) – the input label
- **labellist** (*list*) – the labels that have already been encountered

Returns *lbl* if not found in *labellist* or *lbl* with `_1-9`` (or ``_10-99`, etc.) appended at the end

`GSASIIobj.PhaseIdLookup = {}`

dict listing phase name and random Id keyed by sequential phase index as a *str*; best to access this using `LookupPhaseName()`

`GSASIIobj.PhaseRanIdLookup = {}`

dict listing phase sequential index keyed by phase random Id; best to access this using `LookupPhaseId()`

`GSASIIobj.ShortHistNames = {}`

a dict containing a possibly shortened and when non-unique numbered version of the histogram name. Keyed by the histogram sequential index.

`GSASIIobj.ShortPhaseNames = {}`

a dict containing a possibly shortened and when non-unique numbered version of the phase name. Keyed by the phase sequential index.

`GSASIIobj.VarDesc = {}`

This dictionary lists descriptions for GSAS-II variables, as set in `CompileVarDesc()`. See that function for a description for how keys and values are written.

`GSASIIobj.fmtVarDescr (varname)`

Return a string with a more complete description for a GSAS-II variable

TODO: This will not handle rigid body parameters yet

Parameters `name (str)` – A full G2 variable name with 2 or 3 colons (<p>:<h>:name[:<a>])

Returns a string with the description

`GSASIIobj.getDescr (name)`

Return a short description for a GSAS-II variable

Parameters `name (str)` – The descriptive part of the variable name without colons (:)

Returns a short description or None if not found

`GSASIIobj.getVarDescr (varname)`

Return a short description for a GSAS-II variable

Parameters `name (str)` – A full G2 variable name with 2 or 3 colons (<p>:<h>:name[:<a>])

Returns a five element list as [*p*, 'h', 'name', 'a', 'description'], where *p*, *h*, *a* are str values or *None*, for the phase number, the histogram number and the atom number; *name* will always be an str; and *description* is str or *None*. If the variable name is incorrectly formed (for example, wrong number of colons), *None* is returned instead of a list.

`GSASIIobj.reVarDesc = {}`

This dictionary lists descriptions for GSAS-II variables with the same values as `VarDesc` except that keys have been compiled as regular expressions. Initialized in `CompileVarDesc()`.

GSAS-II UTILITY MODULES

3.1 GSASIIdata: Data for computations

At present this module defines one dict, `ramachandranDist`, which contains arrays for All and specific amino acids

3.2 ElementTable: Periodic Table Data

Element table data for building periodic table with valences & JMOL colors. Need these in case we go back to this periodic table coloring scheme.

Defines list `ElTable` which contains all defined oxidation states for each element, the location in the table, an element name, a color, a size and a second color.

3.3 FormFactors: Scattering Data

Contains atomic scattering factors from “New Analytical Scattering Factor Functions for Free Atoms and Ions for Free Atoms and Ions”, D. Waasmaier & A. Kirfel, *Acta Cryst.* (1995). A51, 416-413.

Also, tabulated coefficients for calculation of Compton Cross Section as a function of $\sin(\theta)/\lambda$ from “Analytic Approximations to Incoherently Scattered X-Ray Intensities”, H. H. M. Balyuzi, *Acta Cryst.* (1975). A31, 600.

3.4 ImageCalibrants: Calibration Standards

GSASII powder calibrants as a dictionary `ImageCalibrants.Calibrants` with substances commonly used for powder calibrations for image data.

Each entry in `ImageCalibrants` consists of:

```
'key': ([Bravais num, ], [(a,b,c,alpha,beta,gamma), ], no. lines skipped, (dmin,pixLimit,cutOff))
```

as an example:

```
'LaB6 SRM660a': ([2, ], [(4.1569162,4.1569162,4.1569162,90,90,90), ], 0, (1.0,10,10)),
```

or where “Bravais num” and “(a,b,...)” are repeated in the case of mixtures:

```
'LaB6 & CeO2': ([2, 0], [(4.1569, 4.1569, 4.1569, 90, 90, 90), (5.4117, 5.4117, 5.4117, 90, 90, 90)], 0, (1.0, 2, 1))
```

To expand this list with locally needed additions, do not modify this file, because you may lose these changes during a software update. Instead duplicate the format of this file in a file named *UserCalibrants.py* and there define the material(s) you want:

```
Calibrants={
    'LaB6 skip 2 lines': ([2, ], [(4.1569162, 4.1569162, 4.1569162, 90, 90, 90), ], 2, (1.0, 10, 10)),
}
```

New key values will be added to the list of options. If a key is duplicated, the information in *UserCalibrants.py* will override the information in this file.

Note, some useful Bravais numbers are: F-cubic=0, I-cubic=1, P-cubic=2, R3/m (hex)=3, P6=4, P4mmm=6

3.5 GSASIIpath: locations & updates

Routines for dealing with file locations, etc.

Determines the location of the compiled (.pyd or .so) libraries.

Interfaces with subversion (svn): Determine the subversion release number by determining the highest version number where `SetVersionNumber()` is called (best done in every GSASII file). Other routines will update GSASII from the subversion server if svn can be found.

`GSASIIpath.GetVersionNumber()`

Return the maximum version number seen in `SetVersionNumber()`

`GSASIIpath.SetVersionNumber(RevString)`

Set the subversion version number

Parameters `RevString (str)` – something like “\$Revision: 1047 \$” that is set by subversion when the file is retrieved from subversion.

Place `GSASIIpath.SetVersionNumber("$Revision: 1047 $")` in every python file.

`GSASIIpath.svnFindLocalChanges(fpath='/Users/toby/software/G2/GSASII')`

Returns a list of files that were changed locally. If no files are changed, the list has length 0

Parameters `fpath` – path to repository dictionary, defaults to directory where the current file is located

Returns None if there is a subversion error (likely because the path is not a repository or svn is not found)

`GSASIIpath.svnGetLog(fpath='/Users/toby/software/G2/GSASII', version=None)`

Get the revision log information for a specific version of the

Parameters

- **fpath (str)** – path to repository dictionary, defaults to directory where the current file is located.
- **version (int)** – the version number to be looked up or None (default) for the latest version.

Returns a dictionary with keys (one hopes) ‘author’, ‘date’, ‘msg’, and ‘revision’

`GSASIIpath.svnGetRev(fpath='/Users/toby/software/G2/GSASII', local=True)`

Obtain the version number for the either the last update of the local version or contacts the subversion server to get the latest update version (# of Head).

Parameters

- **fpath** (*str*) – path to repository dictionary, defaults to directory where the current file is located
- **local** (*bool*) – determines the type of version number, where True (default): returns the latest installed update False: returns the version number of Head on the server

Returns the version number as an str or None if there is a subversion error (likely because the path is not a repository or svn is not found)

`GSASIIpath.svnUpdateDir (fpath='/Users/toby/software/G2/GSASII', version=None)`

This performs an update of the files in a local directory from a server.

Parameters

- **fpath** (*str*) – path to repository dictionary, defaults to directory where the current file is located
- **version** – the number of the version to be loaded. Used only cast as a string, but should be an integer or something that corresponds to a string representation of an integer value when cast. A value of None (default) causes the latest version on the server to be used.

`GSASIIpath.svnUpdateProcess (version=None, projectfile=None)`

perform an update of GSAS-II in a separate python process

`GSASIIpath.whichsvn ()`

Returns a path to the subversion exe file, if any is found. Searches the current path as well as subdirectory “svn” and “svn/bin” in the location of the GSASII source files.

Returns None if svn is not found or an absolute path to the subversion executable file.

3.6 GSASIIElem: functions for element types

`GSASIIElem.CheckElement (El)`

Check if element El is in the periodic table

Parameters **El** (*str*) – One or two letter element symbol, capitalization ignored

Returns True if the element is found

`GSASIIElem.ComptonFac (El, SQ)`

compute Compton scattering factor

Parameters

- **El** – element dictionary
- **SQ** – $(\sin\text{-theta}/\lambda)^2$

Returns compton scattering factor

`GSASIIElem.FPcalc (Orbs, KEv)`

Compute real & imaginary resonant X-ray scattering factors

Parameters

- **Orbs** – list of orbital dictionaries as defined in GetXsectionCoeff
- **KEv** – x-ray energy in keV

Returns C: (f',f'',mu): real, imaginary parts of resonant scattering & atomic absorption coeff.

`GSASIIElem.FixValence (El)`

Returns the element symbol, even when a valence is present

`GSASIIElem.GetAtomInfo (El)`

reads element information from file `atmdata.dat`

`GSASIIElem.GetBLtable (General)`

returns a dictionary of neutron scattering length data for atom types & isotopes found in `General`

Parameters `General` (*dict*) – dictionary of phase info.; includes `AtomTypes` & `Isotopes`

Returns `BLtable`, dictionary of scattering length data; key is atom type

`GSASIIElem.GetFFC5 (ElSym)`

Get 5 term form factor and Compton scattering data

Parameters `ElSym` – str(1-2 character element symbol with proper case);

Return `El` dictionary with 5 term form factor & compton coefficients

`GSASIIElem.GetFFtable (atomTypes)`

returns a dictionary of form factor data for atom types found in `atomTypes`

Parameters `atomTypes` (*list*) – list of atom types

Returns `FFtable`, dictionary of form factor data; key is atom type

`GSASIIElem.GetFormFactorCoeff (El)`

Read X-ray form factor coefficients from `atomdata.asc` file

Parameters `El` (*str*) – element 1-2 character symbol, case irrelevant

Returns `FormFactors`: list of form factor dictionaries

Each X-ray form factor dictionary is:

- `Symbol`: 4 character element symbol with valence (e.g. 'NI+2')
- `Z`: atomic number
- `fa`: 4 A coefficients
- `fb`: 4 B coefficients
- `fc`: C coefficient

`GSASIIElem.GetMagFormFacCoeff (El)`

Read magnetic form factor data from `atomdata.asc` file

Parameters `El` – 2 character element symbol

Returns `MagFormFactors`: list of all magnetic form factors dictionaries for element `El`.

each dictionary contains:

- `'Symbol':Symbol`
- `'Z':Z`
- `'mfa': 4 MA coefficients`
- `'nfa': 4 NA coefficients`
- `'mfb': 4 MB coefficients`
- `'nfb': 4 NB coefficients`
- `'mfc': MC coefficient`

- ‘nfc’: NC coefficient

GSASIIElem.**GetXsectionCoeff** (*El*)

Read atom orbital scattering cross sections for fprime calculations via Cromer-Lieberman algorithm

Parameters *El* – 2 character element symbol

Returns Orbs: list of orbitals each a dictionary with detailed orbital information used by FPcalc

each dictionary is:

- ‘OrbName’: Orbital name read from file
- ‘IfBe’ 0/2 depending on orbital
- ‘BindEn’: binding energy
- ‘BB’: BindEn/0.02721
- ‘XSectIP’: 5 cross section inflection points
- ‘ElEterm’: energy correction term
- ‘SEdge’: absorption edge for orbital
- ‘Nval’: 10/11 depending on IfBe
- ‘LEner’: 10/11 values of log(energy)
- ‘LXSect’: 10/11 values of log(cross section)

GSASIIElem.**ScatFac** (*El*, *SQ*)

compute value of form factor

Parameters

- *El* – element dictionary defined in GetFormFactorCoeff
- *SQ* – (sin-theta/lambda)**2

Returns real part of form factor

GSASIIElem.**getBLvalues** (*BLtables*, *ifList=False*)

Needs a doc string

GSASIIElem.**getFFvalues** (*FFtables*, *SQ*, *ifList=False*)

Needs a doc string

3.7 GSASIIlattice: Unit cells

Perform lattice-related computations

Note that *g* is the reciprocal lattice tensor, and *G* is its inverse, $G = g^{-1}$, where

$$G = \begin{pmatrix} a^2 & ab \cos \gamma & ac \cos \beta \\ ab \cos \gamma & b^2 & bc \cos \alpha \\ ac \cos \beta & bc \cos \alpha & c^2 \end{pmatrix}$$

The “A tensor” terms are defined as $A = (G_{11} \ G_{22} \ G_{33} \ 2G_{12} \ 2G_{13} \ 2G_{23})$ and *A* can be used in this fashion: $d^* = \sqrt{A_1 h^2 + A_2 k^2 + A_3 l^2 + A_4 hk + A_5 hl + A_6 kl}$, where *d* is the d-spacing, and *d** is the reciprocal lattice spacing, $Q = 2\pi d^* = 2\pi/d$

`GSASIIlattice.A2Gmat (A, inverse=True)`

Fill real & reciprocal metric tensor (G) from A.

Parameters

- **A** – reciprocal metric tensor elements as [G11,G22,G33,2*G12,2*G13,2*G23]
- **inverse** (*bool*) – if True return both G and g; else just G

Returns reciprocal (G) & real (g) metric tensors (list of two numpy 3x3 arrays)

`GSASIIlattice.A2cell (A)`

Compute unit cell constants from A

Parameters **A** – [G11,G22,G33,2*G12,2*G13,2*G23] G - reciprocal metric tensor

Returns a,b,c,alpha, beta, gamma (degrees) - lattice parameters

`GSASIIlattice.A2invcell (A)`

Compute reciprocal unit cell constants from A returns tuple with a*,b*,c*,alpha*, beta*, gamma* (degrees)

`GSASIIlattice.CellAbsorption (ElList, Volume)`

Compute unit cell absorption

Parameters

- **ElList** (*dict*) – dictionary of element contents including mu and number of atoms be cell
- **Volume** (*float*) – unit cell volume

Returns mu-total/Volume

`GSASIIlattice.CellBlock (nCells)`

Generate block of unit cells n*n*n on a side; [0,0,0] centered, n = 2*nCells+1 currently only works for nCells = 0 or 1 (not >1)

`GSASIIlattice.CentCheck (Cent, H)`

needs doc string

`GSASIIlattice.CosAngle (U, V, G)`

calculate cos of angle between U & V in generalized coordinates defined by metric tensor G

Parameters

- **U** – 3-vectors assume numpy arrays, can be multiple reflections as (N,3) array
- **V** – 3-vectors assume numpy arrays, only as (3) vector
- **G** – metric tensor for U & V defined space assume numpy array

Returns cos(phi)

`GSASIIlattice.CosSinAngle (U, V, G)`

calculate sin & cos of angle between U & V in generalized coordinates defined by metric tensor G

Parameters

- **U** – 3-vectors assume numpy arrays
- **V** – 3-vectors assume numpy arrays
- **G** – metric tensor for U & V defined space assume numpy array

Returns cos(phi) & sin(phi)

`GSASIIlattice.CrsAng (H, cell, SGData)`

needs doc string

`GSASIIlattice.F1nh` (*Start, SHCoef, phi, beta, SGData*)

needs doc string

`GSASIIlattice.GenHBravais` (*dmin, Bravais, A*)

Generate the positionally unique powder diffraction reflections

Parameters

- **dmin** – minimum d-spacing in Å
- **Bravais** – lattice type (see `GetBraviasNum`). Bravais is one of: 0 F cubic 1 I cubic 2 P cubic 3 R hexagonal (trigonal not rhombohedral) 4 P hexagonal 5 I tetragonal 6 P tetragonal 7 F orthorhombic 8 I orthorhombic 9 C orthorhombic 10 P orthorhombic 11 C monoclinic 12 P monoclinic 13 P triclinic
- **A** – reciprocal metric tensor elements as [G11,G22,G33,2*G12,2*G13,2*G23]

Returns HKL unique d list of [h,k,l,d,-1] sorted with largest d first

`GSASIIlattice.GenHLaue` (*dmin, SGData, A*)

Generate the crystallographically unique powder diffraction reflections for a lattice and Bravais type

Parameters

- **dmin** – minimum d-spacing
- **SGData** – space group dictionary with at least
 - ‘SGLaue’: Laue group symbol: one of ‘-1’, ‘2/m’, ‘mmm’, ‘4/m’, ‘6/m’, ‘4/mmm’, ‘6/mmm’, ‘3m1’, ‘31m’, ‘3’, ‘3R’, ‘3mR’, ‘m3’, ‘m3m’
 - ‘SGLatt’: lattice centering: one of ‘P’, ‘A’, ‘B’, ‘C’, ‘I’, ‘F’
 - ‘SGUniq’: code for unique monoclinic axis one of ‘a’, ‘b’, ‘c’ (only if ‘SGLaue’ is ‘2/m’) otherwise an empty string
- **A** – reciprocal metric tensor elements as [G11,G22,G33,2*G12,2*G13,2*G23]

Returns HKL = list of [h,k,l,d] sorted with largest d first and is unique part of reciprocal space ignoring anomalous dispersion

`GSASIIlattice.GenSHCoeff` (*SGLaue, SamSym, L, IfLMN=True*)

needs doc string

`GSASIIlattice.GetBraviasNum` (*center, system*)

Determine the Bravais lattice number, as used in `GenHBravais`

Parameters

- **center** – one of: ‘P’, ‘C’, ‘I’, ‘F’, ‘R’ (see `SGLatt` from `GSASIIspc.SpcGroup`)
- **system** – one of ‘cubic’, ‘hexagonal’, ‘tetragonal’, ‘orthorhombic’, ‘trigonal’ (for R) ‘monoclinic’, ‘triclinic’ (see `SGSys` from `GSASIIspc.SpcGroup`)

Returns a number between 0 and 13 or throws a `ValueError` exception if the combination of center, system is not found (i.e. non-standard)

`GSASIIlattice.GetKcl` (*L, N, SGLaue, phi, beta*)

needs doc string

`GSASIIlattice.GetKclKs1` (*L, N, SGLaue, psi, phi, beta*)

This is used for spherical harmonics description of preferred orientation; cylindrical symmetry only (M=0) and no sample angle derivatives returned

`GSASIIlattice.GetKs1` (*L, M, SamSym, psi, gam*)

needs doc string

`GSASIIlattice.G1nh` (*Start, SHCoef, psi, gam, SamSym*)

needs doc string

`GSASIIlattice.Gmat2A` (*G*)

Extract A from reciprocal metric tensor (G)

Parameters *G* – reciprocal metric tensor (3x3 numpy array)

Returns *A* = [G11,G22,G33,2*G12,2*G13,2*G23]

`GSASIIlattice.Gmat2AB` (*G*)

Computes orthogonalization matrix from reciprocal metric tensor G

Returns

tuple of two 3x3 numpy arrays (A,B)

- A for crystal to Cartesian transformations $A \cdot x = \text{np.inner}(A, x) = X$
- B (= inverse of A) for Cartesian to crystal transformation $B \cdot X = \text{np.inner}(B, X) = x$

`GSASIIlattice.Gmat2cell` (*g*)

Compute real/reciprocal lattice parameters from real/reciprocal metric tensor (g/G) The math works the same either way.

Parameters (or *G*) (*g*) – real (or reciprocal) metric tensor 3x3 array

Returns a,b,c,alpha, beta, gamma (degrees) (or a*,b*,c*,alpha*,beta*,gamma* degrees)

`GSASIIlattice.Hx2Rh` (*Hx*)

needs doc string

`GSASIIlattice.MaxIndex` (*dmin, A*)

needs doc string

`GSASIIlattice.OdfChk` (*SGLaue, L, M*)

needs doc string

`GSASIIlattice.Rh2Hx` (*Rh*)

needs doc string

`GSASIIlattice.SamAng` (*Tth, Gangls, Sangl, IFCoup*)

Compute sample orientation angles vs laboratory coord. system

Parameters

- **Tth** – Signed theta
- **Gangls** – Sample goniometer angles phi,chi,omega,azimuth
- **Sangl** – Sample angle zeros om-0, chi-0, phi-0
- **IFCoup** – True if omega & 2-theta coupled in CW scan

Returns psi,gam: Sample odf angles dPSdA,dGMdA: Angle zero derivatives

`GSASIIlattice.SwapIndx` (*Axis, H*)

needs doc string

`GSASIIlattice.U6toUij` (*U6*)

Fill matrix (Uij) from U6 = [U11,U22,U33,U12,U13,U23] NB: there is a non numpy version in GSASIIspc: U2Uij

Parameters *U6* (*list*) – 6 terms of u11,u22,...

Returns Uij - numpy [3][3] array of uij

`GSASIIlattice.Uij2Ueqv(Uij, GS, Amat)`

returns 1/3 trace of diagonalized U matrix

`GSASIIlattice.Uij2betaij(Uij, G)`

Convert Uij to beta-ij tensors – stub for eventual completion

Parameters

- **Uij** – numpy array [Uij]
- **G** – reciprocal metric tensor

Returns beta-ij - numpy array [beta-ij]

`GSASIIlattice.UijtoU6(U)`

Fill vector [U11,U22,U33,U12,U13,U23] from Uij NB: there is a non numpy version in GSASIIspc: Uij2U

`GSASIIlattice.calc_V(A)`

Compute the real lattice volume (V) from A

`GSASIIlattice.calc_rDsqr(H, A)`

needs doc string

`GSASIIlattice.calc_rDsqr2(H, G)`

needs doc string

`GSASIIlattice.calc_rDsqrZ(H, A, Z, tth, lam)`

needs doc string

`GSASIIlattice.calc_rV(A)`

Compute the reciprocal lattice volume (V*) from A

`GSASIIlattice.calc_rVsqr(A)`

Compute the square of the reciprocal lattice volume (1/V**2) from A'

`GSASIIlattice.cell2A(cell)`

Obtain A = [G11,G22,G33,2*G12,2*G13,2*G23] from lattice parameters

Parameters **cell** – [a,b,c,alpha,beta,gamma] (degrees)

Returns G reciprocal metric tensor as 3x3 numpy array

`GSASIIlattice.cell2AB(cell)`

Computes orthogonalization matrix from unit cell constants

Parameters **cell** (*tuple*) – a,b,c, alpha, beta, gamma (degrees)

Returns tuple of two 3x3 numpy arrays (A,B) A for crystal to Cartesian transformations $A^*x = \text{np.inner}(A,x) = X B$ (= inverse of A) for Cartesian to crystal transformation $B^*X = \text{np.inner}(B,X) = x$

`GSASIIlattice.cell2GS(cell)`

returns Uij to betaij conversion matrix

`GSASIIlattice.cell2Gmat(cell)`

Compute real and reciprocal lattice metric tensor from unit cell constants

Parameters **cell** – tuple with a,b,c,alpha, beta, gamma (degrees)

Returns reciprocal (G) & real (g) metric tensors (list of two numpy 3x3 arrays)

`GSASIIlattice.combinations(items, n)`

take n distinct items, order matters

`GSASIIlattice.criticalEllipse(prob)`

Calculate critical values for probability ellipsoids from probability

`GSASIIlattice.fillgmat (cell)`

Compute lattice metric tensor from unit cell constants

Parameters `cell` – tuple with a,b,c,alpha, beta, gamma (degrees)

Returns 3x3 numpy array

`GSASIIlattice.getHKLmax (dmin, SGData, A)`

finds maximum allowed hkl for given A within dmin

`GSASIIlattice.invcell2Gmat (invcell)`

Compute real and reciprocal lattice metric tensor from reciprocal unit cell constants

Parameters `invcell` – [a*,b*,c*,alpha*, beta*, gamma*] (degrees)

Returns reciprocal (G) & real (g) metric tensors (list of two 3x3 arrays)

`GSASIIlattice.invpolfcal (ODFln, SGData, phi, beta)`

needs doc string

`GSASIIlattice.permutations (items)`

take all items, order matters

`GSASIIlattice.polfcal (ODFln, SamSym, psi, gam)`

needs doc string

`GSASIIlattice.rotMat (angle, axis=0)`

Prepare rotation matrix for angle in degrees about axis(=0,1,2)

Parameters

- **angle** – angle in degrees
- **axis** – axis (0,1,2 = x,y,z) about which for the rotation

Returns rotation matrix - 3x3 numpy array

`GSASIIlattice.rotMat4 (angle, axis=0)`

Prepare rotation matrix for angle in degrees about axis(=0,1,2) with scaling for OpenGL

Parameters

- **angle** – angle in degrees
- **axis** – axis (0,1,2 = x,y,z) about which for the rotation

Returns rotation matrix - 4x4 numpy array (last row/column for openGL scaling)

`GSASIIlattice.sec2HMS (sec)`

Convert time in sec to H:M:S string

Parameters `sec` – time in seconds

Returns H:M:S string (to nearest 100th second)

`GSASIIlattice.selections (items, n)`

take n (not necessarily distinct) items, order matters

`GSASIIlattice.selftestlist = [<function test0 at 0x5baaa30>, <function test1 at 0x5baaa70>, <function test2 at 0x5baa...`

Defines a list of self-tests

`GSASIIlattice.sortHKLd (HKLd, ifreverse, ifdup)`

needs doc string

Parameters

- **HKLd** – a list of [h,k,l,d,...];

- **ifreverse** – True for largest d first
- **ifdup** – True if duplicate d-spacings allowed

```
GSASIIlattice.test1()
    test cell2A and A2Gmat

GSASIIlattice.test2()
    test Gmat2A, A2cell, A2Gmat, Gmat2cell

GSASIIlattice.test3()
    test invcell2Gmat

GSASIIlattice.test4()
    test calc_rVsq, calc_rV, calc_V

GSASIIlattice.test5()
    test A2invcell

GSASIIlattice.test6()
    test cell2AB

GSASIIlattice.test7()
    test GetBraviasNum(...) and GenHBravais(...)

GSASIIlattice.test8()
    test GenHLaue

GSASIIlattice.test9()
    test GenHLaue

GSASIIlattice.textureIndex(SHCoef)
    needs doc string

GSASIIlattice.uniqueCombinations(items, n)
    take n distinct items, order is irrelevant
```

3.8 GSASIIspc: Space group module

Space group interpretation routines. Note that space group information is stored in a *Space Group (SGData)* object.

GSASIIspc.AllOps(*SGData*)

Returns a list of all operators for a space group, including those for centering and a center of symmetry

Parameters **SGData** – from `SpcGroup()`

Returns

(SGTextList,offsetList,symOpList,G2oprList) where

- SGTextList: a list of strings with formatted and normalized symmetry operators.
- offsetList: a tuple of (dx,dy,dz) offsets that relate the GSAS-II symmetry operation to the operator in SGTextList and symOpList. these dx (etc.) values are added to the GSAS-II generated positions to provide the positions that are generated by the normalized symmetry operators.
- symOpList: a list of tuples with the normalized symmetry operations as (M,T) values (see SGOps in the *Space Group object*)
- G2oprList: The GSAS-II operations for each symmetry operation as a tuple with (center,mult,opnum), where center is (0,0,0), (0.5,0,0), (0.5,0.5,0.5),...; where mult is 1 or -1 for

the center of symmetry and opnum is the number for the symmetry operation, in *SGOps* (starting with 0).

`GSASIIspc.ApplyStringOps (A, SGData, X, Uij=[])`

Needs a doc string

`GSASIIspc.ElementPosition (SGData)`

Under development. Object here is to return a list of symmetry element types and locations suitable for say drawing them. So far I have the element type... getting all possible locations without lookup may be impossible!

`GSASIIspc.GenAtom (XYZ, SGData, All=False, Uij=[], Move=True)`

Generates the equivalent positions for a specified coordinate and space group

Parameters

- **XYZ** – an array, tuple or list containing 3 elements: x, y & z
- **SGData** – from `SpcGroup()`
- **All** – True return all equivalent positions including duplicates; False return only unique positions
- **Uij** – [U11,U22,U33,U12,U13,U23] or [] if no Uij
- **Move** – True move generated atom positions to be inside cell False do not move atoms

Returns

[[XYZEquiv],Idup,[UijEquiv]]

- [XYZEquiv] is list of equivalent positions (XYZ is first entry)
- Idup = [-][C]SS where SS is the symmetry operator number (1-24), C (if not 0,0,0)
- is centering operator number (1-4) and - is for inversion Cell = unit cell translations needed to put new positions inside cell [UijEquiv] - equivalent Uij; absent if no Uij given

`GSASIIspc.GenHKLf (HKL, SGData)`

Uses old GSAS Fortran routine genhkl.for

Parameters

- **HKL** – [h,k,l]
- **SGData** – space group data obtained from `SpcGroup`

Returns

iabsnt,mulp,Uniq,phi

- iabsnt = True if reflection is forbidden by symmetry
- mulp = reflection multiplicity including Friedel pairs
- Uniq = numpy array of equivalent hkl in descending order of h,k,l

`GSASIIspc.GetCSuinel (siteSym)`

returns Uij terms, multipliers, GUI flags & Uiso2Uij multipliers

`GSASIIspc.GetCSxinel (siteSym)`

Needs a doc string

`GSASIIspc.GetKNsym (key)`

Needs a doc string

`GSASIIspc.GetNXUPQsym (siteSym)`

Needs a doc string

`GSASIIspc.GetOprPtrName (key)`

Needs a doc string

`GSASIIspc.HStrainNames (SGData)`

Needs a doc string

`GSASIIspc.Latt2text (Latt)`

From lattice type ('P', 'A', etc.) returns ';' delimited cell centering vectors

`GSASIIspc.MT2text (M, T)`

From space group matrix/translation operator returns text version

`GSASIIspc.MoveToUnitCell (xyz)`

Translates a set of coordinates so that all values are ≥ 0 and < 1

Parameters `xyz` – a list or numpy array of fractional coordinates

Returns `XYZ` - numpy array of new coordinates now 0 or greater and less than 1

`GSASIIspc.Muiso2Shkl (muiso, SGData, cell)`

this is to convert isotropic mustrain to generalized Shkls - doesn't work just now

`GSASIIspc.MustrainCoeff (HKL, SGData)`

Needs a doc string

`GSASIIspc.MustrainNames (SGData)`

Needs a doc string

`GSASIIspc.Opposite (XYZ, toler=0.0002)`

Gives opposite corner, edge or face of unit cell for position within tolerance. Result may be just outside the cell within tolerance

Parameters

- `XYZ` – $0 \leq \text{np.array}[x,y,z] < 1$ as by `MoveToUnitCell`
- `toler` – unit cell fraction tolerance making opposite

Returns `XYZ`: array of opposite positions; always contains `XYZ`

`GSASIIspc.SGErrors (IErr)`

Interprets the error message code from `SpcGroup`. Used in `SpaceGroup`.

Parameters `IErr` – see `SGError` in `SpcGroup()`

Returns `ErrString` - a string with the error message or "Unknown error"

`GSASIIspc.SGPrint (SGData)`

Print the output of `SpcGroup` in a nicely formatted way. Used in `SpaceGroup`

Parameters `SGData` – from `SpcGroup()`

Returns `SGText` - list of strings with the space group details

`GSASIIspc.SGpolar (SGData)`

Determine identity of polar axes if any

`GSASIIspc.SpaceGroup (SGSymbol)`

Print the output of `SpcGroup` in a nicely formatted way.

Parameters `SGSymbol` – space group symbol (string) with spaces between axial fields

Returns nothing

GSASIIspc.SpcGroup (*SGSymbol*)

Determines cell and symmetry information from a short H-M space group name

Parameters **SGSymbol** – space group symbol (string) with spaces between axial fields

Returns

(SGError,SGData) * SGError = 0 for no errors; >0 for errors (see SGErrors below for details) *

SGData - is a dict (see *Space Group object*) with entries:

- 'SpGrp': space group symbol, slightly cleaned up
- 'Laue': one of '-1', '2/m', 'mmm', '4/m', '4/mmm', '3R', '3mR', '3', '3m1', '31m', '6/m', '6/mmm', 'm3', 'm3m'
- 'SGInv': boolean; True if centrosymmetric, False if not
- 'SGLatt': one of 'P', 'A', 'B', 'C', 'I', 'F', 'R'
- 'SGUniq': one of 'a', 'b', 'c' if monoclinic, '' otherwise
- 'SGCen': cell centering vectors [0,0,0] at least
- 'SGOps': symmetry operations as [M,T] so that $M*x+T = x$
- 'SGSys': one of 'triclinic', 'monoclinic', 'orthorhombic', 'tetragonal', 'rhombohedral', 'trigonal', 'hexagonal', 'cubic'
- 'SGPolax': one of '', 'x', 'y', 'x y', 'z', 'x z', 'y z', 'xyz', '111' for arbitrary axes

GSASIIspc.StandardizeSpcName (*spcgroup*)

Accept a spacegroup name where spaces may have not been used in the names according to the GSAS convention (spaces between symmetry for each axis) and return the space group name as used in GSAS

GSASIIspc.StringOpsProd (*A, B, SGData*)

Find $A*B$ where A & B are in strings '- ' + '100*c+n' + '+ijk' where '-' indicates inversion, c(>0) is the cell centering operator, n is operator number from SgOps and ijk are unit cell translations (each may be <0). Should return resultant string - C. SGData - dictionary using entries:

- 'SGCen': cell centering vectors [0,0,0] at least
- 'SGOps': symmetry operations as [M,T] so that $M*x+T = x$

GSASIIspc.SytSym (*XYZ, SGData*)

Generates the number of equivalent positions and a site symmetry code for a specified coordinate and space group

Parameters

- **XYZ** – an array, tuple or list containing 3 elements: x, y & z
- **SGData** – from SpcGroup

Returns a two element tuple:

- The 1st element is a code for the site symmetry (see GetKNSym)
- The 2nd element is the site multiplicity

GSASIIspc.selftestlist = [<function test0 at 0x1e305f0>, <function test1 at 0x1e30630>, <function test2 at 0x1e30670>]

Defines a list of self-tests

GSASIIspc.test0 ()

self-test #0: exercise MoveToUnitCell

GSASIIspc.test1 ()

self-test #1: SpcGroup and SGPrint against previous results

```

GSASIIspc.test2()
    self-test #2: SpcGroup against cctbx (sgtbx) computations

GSASIIspc.test3()
    self-test #3: exercise SytSym (includes GetOprPtrName, GenAtom, GetKNSym) for selected space groups
    against info in IT Volume A

```

3.9 *gltext: draw OpenGL text*

Routines that render text on OpenGL without use of GLUT.

Code written by Christian Brugger & Stefan Hacker and distributed under GNU General Public License.

```

class gltext.Text (text='Text', font=None, font_size=8, foreground=wx.Colour(), centered=False)
    A simple class for using System Fonts to display text in an OpenGL scene. The Text adds a global Cache of
    already created text elements to TextElement's base functionality so you can save some memory and increase
    speed

    centered
        Display the text centered

    draw_text (position=wx.Point(0, 0), scale=1.0, rotation=0)
        position (wx.Point) - x/y Position to draw in scene scale (float) - Scale rotation (int) - Rotation in degree

        Draws the text to the scene

    font
        Font of the object

    font_size
        Font size

    foreground
        Color/Overlay bitmap of the text

    getTextElement ()
        Returns the text element bound to the Text class

    getTexture ()
        Returns the texture of the bound TextElement

    getTexture_size ()
        Returns a texture size tuple

    setCentered (value, reinit=True)
        value (bool) - New centered value reinit (bool) - Create a new texture

        Sets a new value for 'centered'

    setFont (value, reinit=True)
        value (bool) - New Font reinit (bool) - Create a new texture

        Sets a new font

    setFont_size (value, reinit=True)
        value (bool) - New font size reinit (bool) - Create a new texture

        Sets a new font size

    setForeground (value, reinit=True)
        value (bool) - New centered value reinit (bool) - Create a new texture

```

Sets a new value for 'centered'

setText (*value*, *reinit=True*)

value (bool) - New Text reinit (bool) - Create a new texture

Sets a new text

text

Text of the object

text_element

TextElement bound to this class

texture

Texture of bound TextElement

texture_size

Size of the used texture

class `gltext.TextElement` (*text='', font=None, foreground=wx.Colour(), centered=False*)

A simple class for using system Fonts to display text in an OpenGL scene

bind ()

Increase refcount

centered

Is text centered

createTexture ()

Creates a texture from the settings saved in TextElement, to be able to use normal system fonts conveniently a wx.MemoryDC is used to draw on a wx.Bitmap. As wxwidgets device contexts don't support alpha at all it is necessary to apply a little hack to preserve antialiasing without sticking to a fixed background color:

We draw the bmp in b/w mode so we can use its data as a alpha channel for a solid color bitmap which after GL_ALPHA_TEST and GL_BLEND will show a nicely antialiased text on any surface.

To access the raw pixel data the bmp gets converted to a wx.Image. Now we just have to merge our foreground color with the alpha data we just created and push it all into a OpenGL texture and we are DONE *inhalesdelpy*

DRAWBACK of the whole conversion thing is a really long time for creating the texture. If you see any optimizations that could save time PLEASE CREATE A PATCH!!!

deleteTexture ()

Deletes the OpenGL texture object

draw_text (*position=wx.Point(0, 0), scale=1.0, rotation=0*)

position (wx.Point) - x/y Position to draw in scene *scale* (float) - Scale *rotation* (int) - Rotation in degree

Draws the text to the scene

font

Font of the object

foreground

Color of the text

isBound ()

Return refcount

owner_cnt

Owner count

release()
Decrease refcount

text
Text of the object

texture
Used texture

texture_size
Size of the used texture

GSAS-II GUI ROUTINES

4.1 GSASIIgrid: Basic GUI routines

class GSASIIgrid.**ASCIIValidator** (*result=None, key=None*)

A validator to be used with a TextCtrl to prevent entering characters other than ASCII characters.

The value is checked for validity after every keystroke If an invalid number is entered, the box is highlighted. If the number is valid, it is saved in result[key]

Parameters

- **result** (*dict/list*) – List or dict where value should be placed when valid
- **key** (*any*) – key to use for result (int for list)

Clone ()

Create a copy of the validator, a strange, but required component

OnChar (*event*)

Called each type a key is pressed ignores keys that are not allowed for int and float types

TestValid (*tc*)

Check if the value is valid by casting the input string into ASCII.

Save it in the dict/list where the initial value was stored

Parameters *tc* (*wx.TextCtrl*) – A reference to the TextCtrl that the validator is associated with.

TransferFromWindow ()

Needed by validator, strange, but required component

TransferToWindow ()

Needed by validator, strange, but required component

class GSASIIgrid.**AddHelp** (*frame, helpType, helpLbl=None, title=''*)

For the Mac: creates an entry to the help menu of type 'Help on <helpType>': where helpType is a reference to an HTML page to be opened.

NOTE: when appending this menu (menu.Append) be sure to set the title to '&Help' so that wx handles it correctly.

OnHelpById (*event*)

Called when Help on... is pressed in a menu. Brings up a web page for documentation.

GSASIIgrid.**CallScrolledMultiEditor** (*parent, dictlst, elem1st, prelbl=[], postlbl=[], title='Edit items', header='', size=(300, 250), CopyButton=False*)

Shell routine to call a ScrolledMultiEditor dialog. See [ScrolledMultiEditor](#) for parameter definitions.

Returns True if the OK button is pressed; False if the window is closed with the system menu or the Cancel button.

class GSASIIgrid.**DataFrame** (*parent, frame, data=None, name=None, size=None, pos=None*)

Create the data item window and all the entries in menus used in that window. For Linux and windows, the menu entries are created for the current data item window, but in the Mac the menu is accessed from all windows. This means that a different menu is posted depending on which data item is posted. On the Mac, all the menus contain the data tree menu items, but additional menus are added specific to the data item.

Note that while the menus are created here, the binding for the menus is done later in various GSASII*GUI modules, where the functions to be called are defined.

Bind (**args, **kwargs*)

Override the Bind() function: on the Mac the binding is to the main window, so that menus operate with any window on top. For other platforms, call the default wx.Frame Bind()

PostfillDataMenu (*empty=False*)

Create the “standard” part of data frame menus. Note that on Linux and Windows, this is the standard help Menu. On Mac, this menu duplicates the tree menu, but adds an extra help command for the data item and a separator.

PrefillDataMenu (*menu, helpType, helpLbl=None, empty=False*)

Create the “standard” part of data frame menus. Note that on Linux and Windows nothing happens here. On Mac, this menu duplicates the tree menu, but adds an extra help command for the data item and a separator.

class GSASIIgrid.**DisAglDialog** (*parent, data, default*)

Distance/Angle Controls input dialog. After ShowModal () returns, the results are found in dict self.data, which is accessed using GetData ().

Parameters

- **parent** (*wx.Frame*) – reference to parent frame (or None)
- **data** (*dict*) – a dict containing the current search ranges or an empty dict, which causes default values to be used. Will be used to set element *DisAglCtrls* in *Phase Tree Item*
- **default** (*dict*) – A dict containing the default search ranges for each element.

Draw (*data*)

Creates the contents of the dialog. Normally called by __init__ ().

GetData ()

Returns the values from the dialog

OnOk (*event*)

Called when the OK button is pressed

OnReset (*event*)

Called when the Reset button is pressed

class GSASIIgrid.**EnumSelector** (*parent, dct, item, choices, values=None, **kw*)

A customized wxpython.ComboBox that selects items from a list of choices, but sets a dict (list) entry to the corresponding entry from the input list of values.

Parameters

- **parent** (*wx.Panel*) – the parent to the ComboBox (usually a frame or panel)
- **dct** (*dict*) – a dict (or list) to contain the value set for the ComboBox.
- **item** – the dict key (or list index) where dct[item] will be set to the value selected in the ComboBox. Also, dct[item] contains the starting value shown in the widget. If the value

does not match an entry in `values`, the first value in `choices` is used as the default, but `dct[item]` is not changed.

- **choices** (*list*) – a list of choices to be displayed to the user such as

```
["default", "option 1", "option 2", ]
```

Note that these options will correspond to the entries in `values` (if specified) item by item.

- **values** (*list*) – a list of values that correspond to the options in `choices`, such as

```
[0, 1, 2]
```

The default for `values` is to use the same list as specified for `choices`.

- **(other)** – additional keyword arguments accepted by `ComboBox` can be specified.

class GSASIIgrid.**G2CheckBox** (*parent, label, loc, key*)

A customized version of a `CheckBox` that automatically initializes the control to match a supplied

Parameters

- **parent** (*wx.Panel*) – name of panel or frame that will be the parent to the `TextCtrl`. Can be `None`.
- **label** (*str*) – text to put on check button
- **loc** (*dict/list*) – the dict or list with the initial value to be placed in the `CheckBox`.
- **key** (*int/str*) – the dict key or the list index for the value to be edited by the `CheckBox`. The `loc[key]` element must exist. The `CheckBox` will be initialized from this value. If the value is anything other than `True` (or `1`), it will be taken as `False`.

class GSASIIgrid.**G2HtmlWindow** (*parent, *args, **kwargs*)

Displays help information in a primitive HTML browser type window

class GSASIIgrid.**G2MultiChoiceDialog** (*parent, title, header, ChoiceList, toggle=True, **kw*)

A dialog similar to `MultiChoiceDialog` except that buttons are added to set all choices and to toggle all choices.

Parameters

- **ParentFrame** (*wx.Frame*) – reference to parent frame
- **title** (*str*) – heading above list of choices
- **header** (*str*) – Title to place on window frame
- **ChoiceList** (*list*) – a list of choices where one will be selected
- **toggle** (*bool*) – If `True` (default) the toggle and select all buttons are displayed
- **kw** – optional keyword parameters for the `wx.Dialog` may be included such as `size` [which defaults to `(320,310)`] and `style` (which defaults to `wx.DEFAULT_DIALOG_STYLE|wx.RESIZE_BORDER|wx.CENTRE`) `wx.OK` | `wx.CANCEL`); note that `wx.OK` and `wx.CANCEL` controls the presence of the eponymous buttons in the dialog.

Returns the name of the created dialog

GetSelections ()

Returns a list of the indices for the selected choices

class GSASIIgrid.**GSGrid** (*parent, name=''*)

Basic `wx.Grid` implementation

class GSASIIgrid.**GSNoteBook** (*parent, name='', size=None*)

Notebook used in various locations; implemented with wx.aui extension

GSASIIgrid.**GetPatternTreeDataNames** (*G2frame, dataTypes*)

Needs a doc string

GSASIIgrid.**GetPatternTreeItemId** (*G2frame, parentId, itemText*)

Needs a doc string

class GSASIIgrid.**GridFractionEditor** (*grid*)

A grid cell editor class that allows entry of values as fractions as well as sine and cosine values [as s() and c()]

class GSASIIgrid.**HelpButton** (*parent, msg*)

Create a help button that displays help information. The text is displayed in a modal message window.

TODO: it might be nice if it were non-modal: e.g. it stays around until the parent is deleted or the user closes it, but this did not work for me.

Parameters

- **parent** – the panel which will be the parent of the button
- **msg** (*str*) – the help text to be displayed

GSASIIgrid.**HorizontalLine** (*sizer, parent*)

Draws a horizontal line as wide as the window. This shows up on the Mac as a very thin line, no matter what I do

GSASIIgrid.**ItemSelector** (*ChoiceList, ParentFrame=None, title='Select an item', size=None, header='Item Selector', useCancel=True, multiple=False*)

Provide a wx dialog to select a single item from list of choices

Parameters

- **ChoiceList** (*list*) – a list of choices where one will be selected
- **ParentFrame** (*wx.Frame*) – Name of parent frame (default None)
- **title** (*str*) – heading above list of choices (default 'Select an item')
- **size** (*wx.Size*) – Size for dialog to be created (default None – size as needed)
- **header** (*str*) – Title to place on window frame (default 'Item Selector')
- **useCancel** (*bool*) – If True (default) both the OK and Cancel buttons are offered
- **multiple** (*bool*) – If True then multiple items can be selected (default False)

Returns the selection index or None or a selection list if multiple is true

GSASIIgrid.**MovePatternTreeToGrid** (*G2frame, item*)

Needs a doc string

class GSASIIgrid.**MyHelp** (*frame, helpType=None, helpLbl=None, morehelpitems=[], title=''*)

A class that creates the contents of a help menu. The menu will start with two entries:

- 'Help on <helpType>': where helpType is a reference to an HTML page to be opened
- About: opens an About dialog using OnHelpAbout. N.B. on the Mac this gets moved to the App menu to be consistent with Apple style.

NOTE: for this to work properly with respect to system menus, the title for the menu must be &Help, or it will not be processed properly:

```
menu.Append(menu=MyHelp(self,...),title("&Help"))
```

OnCheckUpdates (*event*)

Check if the GSAS-II repository has an update for the current source files and perform that update if requested.

OnHelpAbout (*event*)

Display an 'About GSAS-II' box

OnHelpById (*event*)

Called when Help on... is pressed in a menu. Brings up a web page for documentation.

OnSelectVersion (*event*)

Allow the user to select a specific version of GSAS-II

class GSASIIgrid.**MyHtmlPanel** (*frame, id*)

Defines a panel to display HTML help information, as an alternative to displaying help information in a web browser.

class GSASIIgrid.**NumberValidator** (*typ, positiveonly=False, min=None, max=None, result=None, key=None, OKcontrol=None, CIFinput=False*)

A validator to be used with a TextCtrl to prevent entering characters other than digits, signs, and for float input, a period and exponents.

The value is checked for validity after every keystroke If an invalid number is entered, the box is highlighted. If the number is valid, it is saved in result[key]

Parameters

- **typ** (*type*) – the base data type. Must be int or float.
- **positiveonly** (*bool*) – If True, negative integers are not allowed (default False). This prevents the + or - keys from being pressed. Used with typ=int; ignored for typ=float.
- **min** (*number*) – Minimum allowed value. If None (default) the lower limit is unbounded
- **max** (*number*) – Maximum allowed value. If None (default) the upper limit is unbounded
- **result** (*dict/list*) – List or dict where value should be placed when valid
- **key** (*any*) – key to use for result (int for list)
- **OKcontrol** (*function*) – function or class method to control an OK button for a window. Ignored if None (default)
- **CIFinput** (*bool*) – allows use of a single '?' or '.' character as valid input.

CheckInput (*previousInvalid*)

called to test every change to the TextCtrl for validity and to change the appearance of the TextCtrl

Anytime the input is invalid, call self.OKcontrol (if defined) because it is fast. If valid, check for any other invalid entries only when changing from invalid to valid, since that is slower.

Parameters **previousInvalid** (*bool*) – True if the TextCtrl contents were invalid prior to the current change.

Clone ()

Create a copy of the validator, a strange, but required component

OnChar (*event*)

Called each type a key is pressed ignores keys that are not allowed for int and float types

ShowValidity (*tc*)

Set the control colors to show invalid input

Parameters **tc** (*wx.TextCtrl*) – A reference to the TextCtrl that the validator is associated with.

TestValid (*tc*)

Check if the value is valid by casting the input string into the current type.

Set the invalid variable in the TextCtrl object accordingly.

If the value is valid, save it in the dict/list where the initial value was stored, if appropriate.

Parameters *tc* (*wx.TextCtrl*) – A reference to the TextCtrl that the validator is associated with.

TransferFromWindow ()

Needed by validator, strange, but required component

TransferToWindow ()

Needed by validator, strange, but required component

class GSASIIgrid.**PickTwoDialog** (*parent, title, prompt, names, choices*)

This does not seem to be in use

class GSASIIgrid.**ScrolledMultiEditor** (*parent, dictlst, elemlst, prelbl=[], postlbl=[], title='Edit items', header='', size=(300, 250), CopyButton=False, minvals=[], maxvals=[], sizevals=[]*)

Define a window for editing a potentially large number of dict- or list-contained values with validation for each item. Edited values are automatically placed in their source location. If invalid entries are provided, the TextCtrl is turned yellow and the OK button is disabled.

The type for each TextCtrl validation is determined by the initial value of the entry (int, float or string). Float values can be entered in the TextCtrl as numbers or also as algebraic expressions using operators + - / * () and **, in addition pi, sind(), cosd(), tand(), and sqrt() can be used, as well as appreviations s(), sin(), c(), cos(), t(), tan() and sq().

Parameters

- **parent** (*wx.Frame*) – name of parent window, or may be None
- **dictlst** (*tuple*) – a list of dicts or lists containing values to edit
- **elemlst** (*tuple*) – a list of keys for each item in a dictlst. Must have the same length as dictlst.
- **parent** – name of parent window, or may be None
- **prelbl** (*tuple*) – a list of labels placed before the TextCtrl for each item (optional)
- **postlbl** (*tuple*) – a list of labels placed after the TextCtrl for each item (optional)
- **title** (*str*) – a title to place in the frame of the dialog
- **header** (*str*) – text to place at the top of the window. May contain new line characters.
- **size** (*wx.Size*) – a size parameter that dictates the size for the scrolled region of the dialog. The default is (300,250).
- **CopyButton** (*bool*) – if True adds a small button that copies the value for the current row to all fields below (default is False)
- **minvals** (*list*) – optional list of minimum values for validation of float or int values. Ignored if value is None.
- **maxvals** (*list*) – optional list of maximum values for validation of float or int values. Ignored if value is None.
- **sizevals** (*list*) – optional list of wx.Size values for each input widget. Ignored if value is None.

Returns the wx.Dialog created here. Use method .ShowModal() to display it.

Example for use of ScrolledMultiEditor:


```
dlg = <pkg>.ScrolledMultiEditor(frame,dictlst,elemlst,prelbl,postlbl,
                                header=header)
if dlg.ShowModal() == wx.ID_OK:
    for d,k in zip(dictlst,elemlst):
        print d[k]
```

Example definitions for dictlst and elemlst:

```
dictlst = (dict1,list1,dict1,list1)
elemlst = ('a', 1, 2, 3)
```

This causes items dict1['a'], list1[1], dict1[2] and list1[3] to be edited.

Note that these items must have int, float or str values assigned to them. The dialog will force these types to be retained. String values that are blank are marked as invalid.

ControlOKButton (*setvalue*)

Enable or Disable the OK button for the dialog. Note that this is passed into the ValidatedTxtCtrl for use by validators.

Parameters *setvalue* (*bool*) – if True, all entries in the dialog are checked for validity. if False then the OK button is disabled.

GSASIIgrid.SetDataMenuBar (*G2frame, menu=None*)

Set the menu for the data frame. On the Mac put this menu for the data tree window instead.

Note that data frame items do not have menus, for these (*menu=None*) display a blank menu or on the Mac display the standard menu for the data tree window.

GSASIIgrid.ShowHelp (*helpType, frame*)

Called to bring up a web page for documentation.

class GSASIIgrid.ShowLSParms (*parent, title, parmDict, varyList, fullVaryList, size=(300, 430)*)

Create frame to show least-squares parameters

class GSASIIgrid.SingleFloatDialog (*parent, title, prompt, value, limits=[0.0, 1.0], format='%.5g'*)

Dialog to obtain a single float value from user

class GSASIIgrid.SingleStringDialog (*parent, title, prompt, value='', size=(200, -1)*)

Dialog to obtain a single string value from user

Parameters

- **parent** (*wx.Frame*) – name of parent frame
- **title** (*str*) – title string for dialog
- **prompt** (*str*) – string to tell use what they are inputting
- **value** (*str*) – default input value, if any

GetValue ()

Use this method to get the value entered by the user :returns: string entered by user

Show ()

Use this method after creating the dialog to post it :returns: True if the user pressed OK; False if the User pressed Cancel

class GSASIIgrid.SymOpDialog (*parent, SGData, New=True, ForceUnit=False*)

Class to select a symmetry operator

class GSASIIgrid.Table (*data=[], rowLabels=None, colLabels=None, types=None*)

Basic data table for use with GSgrid

`GSASIIgrid.UpdateControls (G2frame, data)`

Edit overall GSAS-II controls in main Controls data tree entry

`GSASIIgrid.UpdateHKLControls (G2frame, data)`

Needs a doc string

`GSASIIgrid.UpdateNotebook (G2frame, data)`

Called when the data tree notebook entry is selected. Allows for editing of the text in that tree entry

`GSASIIgrid.UpdatePWHKPlot (G2frame, kind, item)`

Called when the histogram main tree entry is called. Displays the histogram weight factor, refinement statistics for the histogram and the range of data for a simulation.

Also invokes a plot of the histogram.

`GSASIIgrid.UpdateSeqResults (G2frame, data)`

Called when the Sequential Results data tree entry is selected to show results from a sequential refinement.

Parameters

- **G2frame** (*wx.Frame*) – main GSAS-II data tree windows
- **data** (*dict*) – a dictionary containing the following items:
 - ‘histNames’ - list of histogram names in order as processed by Sequential Refinement
 - ‘varyList’ - list of variables - identical over all refinements in sequence
 - ‘histName’ - dictionaries for all data sets processed, which contains:
 - * ‘variables’ - result[0] from leastsq call
 - * ‘varyList’ - list of variables; same as above
 - * ‘sig’ - esds for variables
 - * ‘covMatrix’ - covariance matrix from individual refinement
 - * ‘title’ - histogram name; same as dict item name
 - * ‘newAtomDict’ - new atom parameters after shifts applied
 - * ‘newCellDict’ - new cell parameters after shifts to A0-A5 applied

`class GSASIIgrid.ValidatedTextCtrl (parent, loc, key, notBlank=True, min=None, max=None, OKcontrol=None, OnLeave=None, typeHint=None, CIFinput=False, **kw)`

Create a TextCtrl widget that uses a validator to prevent the entry of inappropriate characters and changes color to highlight when invalid input is supplied. As valid values are typed, they are placed into the dict or list where the initial value came from. The type of the initial value must be int, float or str or None (see *key* and *typeHint*); this type (or the one in *typeHint*) is preserved.

Float values can be entered in the TextCtrl as numbers or also as algebraic expressions using operators + - / * () and **, in addition pi, sind(), cosd(), tand(), and sqrt() can be used, as well as abbreviations s, sin, c, cos, t, tan and sq.

Parameters

- **parent** (*wx.Panel*) – name of panel or frame that will be the parent to the TextCtrl. Can be None.
- **loc** (*dict/list*) – the dict or list with the initial value to be placed in the TextCtrl.
- **key** (*int/str*) – the dict key or the list index for the value to be edited by the TextCtrl. The *loc[key]* element must exist, but may have value None. If None, the type for the element is taken from *typeHint* and the value for the control is set initially blank (and thus invalid.)

This is a way to specify a field without a default value: a user must set a valid value. If the value is not None, it must have a base type of int, float, str or unicode; the TextCtrl will be initialized from this value.

- **notBlank** (*bool*) – if True (default) blank values are invalid for str inputs.
- **min** (*number*) – minimum allowed valid value. If None (default) the lower limit is unbounded.
- **max** (*number*) – maximum allowed valid value. If None (default) the upper limit is unbounded
- **OKcontrol** (*function*) – specifies a function or method that will be called when the input is validated. The called function is supplied with one argument which is False if the TextCtrl contains an invalid value and True if the value is valid. Note that this function should check all values in the dialog when True, since other entries might be invalid. The default for this is None, which indicates no function should be called.
- **OnLeave** (*function*) – specifies a function or method that will be called when the focus for the control is lost. The called function is supplied with (at present) three keyword arguments:
 - **invalid**: (*bool*) True if the value for the TextCtrl is invalid
 - **value**: (*int/float/str*) the value contained in the TextCtrl
 - **tc**: (*wx.TextCtrl*) the TextCtrl name

The number of keyword arguments may be increased in the future, if needs arise, so it is best to code these functions with a `**kwargs` argument so they will continue to run without errors

The default for OnLeave is None, which indicates no function should be called.

- **typeHint** (*type*) – the value of typeHint is overrides the initial value for the dict/list element `loc[key]`, if set to int or float, which specifies the type for input to the TextCtrl. Defaults as None, which is ignored.
- **CIFinput** (*bool*) – for str input, indicates that only printable ASCII characters may be entered into the TextCtrl. Forces output to be ASCII rather than Unicode. For float and int input, allows use of a single '?' or '.' character as valid input.
- **(other)** – other optional keyword parameters for the wx.TextCtrl widget such as Size or Style may be specified.

EvaluateExpression ()

Show the computed value when an expression is entered to the TextCtrl Make sure that the number fits by truncating decimal places and switching to scientific notation, as needed. Called on loss of focus.

ShowStringValidity (*previousInvalid=True*)

Check if input is valid. Anytime the input is invalid, call self.OKcontrol (if defined) because it is fast. If valid, check for any other invalid entries only when changing from invalid to valid, since that is slower.

Parameters **previousInvalid** (*bool*) – True if the TextCtrl contents were invalid prior to the current change.

class GSASIIgrid.**downdate** (*parent=None*)

Dialog to allow a user to select a version of GSAS-II to install

getVersion ()

Get the version number in the dialog

4.2 GSASIIIO: Misc I/O routines

Module with miscellaneous routines for input and output. Many are GUI routines to interact with user.

Includes support for image reading.

Also includes base classes for data import routines.

GSASIIIO.CheckImageFile (*G2frame, imagefile*)

Get an new image file name if the specified one does not exist

Parameters

- **G2frame** (*wx.Frame*) – main GSAS-II Frame and data object
- **imagefile** (*str*) – name of image file

Returns imagefile, if it exists, or the name of a file that does exist or False if the user presses Cancel

class GSASIIIO.ExportBaseclass (*G2frame, formatName, extension, longFormatName=None*)

Defines a base class for the exporting of GSAS-II results

CloseFile (*fp=None*)

Close a file opened in OpenFile

Parameters **fp** (*file*) – the file object to be closed. If None (default) file object self.fp is closed.

ExportSelect (*AskFile=True*)

Selects histograms or phases when needed. Sets a default file name.

Parameters **AskFile** (*bool*) – if AskFile is True (default) get the name of the file in a dialog

Returns True in case of an error

GetAtoms (*phasenam*)

Gets the atoms associated with a phase. Can be used with standard or macromolecular phases

Parameters **phasenam** (*str*) – the name for the selected phase

Returns

a list of items for eac atom where each item is a list containing: label, typ, mult, xyz, and td, where

- label and typ are the atom label and the scattering factor type (str)
- mult is the site multiplicity (int)
- xyz is contains a list with four pairs of numbers: x, y, z and fractional occupancy and their standard uncertainty (or a negative value)
- td is contains a list with either one or six pairs of numbers: if one number it is U_{iso} and with six numbers it is U_{11} , U_{22} , U_{33} , U_{12} , U_{13} & U_{23} paired with their standard uncertainty (or a negative value)

GetCell (*phasenam*)

Gets the unit cell parameters and their s.u.'s for a selected phase

Parameters **phasenam** (*str*) – the name for the selected phase

Returns *cellList, cellSig* where each is a 7 element list corresponding to a, b, c, alpha, beta, gamma, volume where *cellList* has the cell values and *cellSig* has their uncertainties.

InitExport (*event*)

Determines the type of menu that called the Exporter.

OpenFile (*fil=None*)

Open the output file

Parameters *fil* (*str*) – The name of the file to open. If None (default) the name defaults to `self.filename`.

Returns the file object opened by the routine which is also saved as `self.fp`

Write (*line*)

write a line of output, attaching a line-end character

Parameters *line* (*str*) – the text to be written.

askSaveFile ()

Ask the user to supply a file name

Returns a file name (*str*)

dumpTree (*mode='type'*)

Print out information on the data tree dicts loaded in `loadTree`

loadParmDict ()

Load the GSAS-II refinable parameters from the tree into a dict (`self.parmDict`). Update refined values to those from the last cycle and set the uncertainties for the refined parameters in another dict (`self.sigDict`).

Expands the `parm` & `sig` dicts to include values derived from constraints.

loadTree ()

Load the contents of the data tree into a set of dicts (`self.OverallParms`, `self.Phases` and `self.Histogram` as well as `self.powderDict` & `self.xtalDict`)

- The childrenless data tree items are overall parameters/controls for the entire project and are placed in `self.OverallParms`
- Phase items are placed in `self.Phases`
- Data items are placed in `self.Histogram`. The key for these data items begin with a keyword, such as `PWDR`, `IMG`, `HKLF`,... that identifies the data type.

GSASIIIO.ExtractFileFromZip (*filename*, *selection=None*, *confirmread=True*, *confirmoverwrite=True*, *parent=None*, *multipleselect=False*)

If the filename is a zip file, extract a file from that archive.

Parameters

- **Selection** (*list*) – used to predefine the name of the file to be extracted. Filename case and zip directory name are ignored in selection; the first matching file is used.
- **confirmread** (*bool*) – if True asks the user to confirm before expanding the only file in a zip
- **confirmoverwrite** (*bool*) – if True asks the user to confirm before overwriting if the extracted file already exists
- **multipleselect** (*bool*) – if True allows more than one zip file to be extracted, a list of file(s) is returned. If only one file is present, do not ask which one, otherwise offer a list of choices (unless selection is used).

Returns the name of the file that has been created or a list of files (see `multipleselect`)

If the file is not a zipfile, return the name of the input file. If the zipfile is empty or no file has been selected, return None

GSASIIIO.FileDlgFixExt (*dlg, file*)

this is needed to fix a problem in linux `wx.FileDialog`

GSASIIIO.GetEdfData (*filename, imageOnly=False*)

Read European detector data edf file

GSASIIIO.GetG2Image (*filename*)

Read an image as a python pickle

GSASIIIO.GetGESumData (*filename, imageOnly=False*)

Read SUM file as produced at 1-ID from G.E. images

GSASIIIO.GetImageData (*G2frame, imagefile, imageOnly=False*)

Read an image with the file reader keyed by the file extension

Parameters

- **G2frame** (*wx.Frame*) – main GSAS-II Frame and data object. Note: not used!
- **imagefile** (*str*) – name of image file
- **imageOnly** (*bool*) – If True return only the image, otherwise (default) return more (see below)

Returns an image as a numpy array or a list of four items: Comments, Data, Npix and the Image, as selected by imageOnly

GSASIIIO.GetImgData (*filename, imageOnly=False*)

Read an ADSC image file

GSASIIIO.GetMAR345Data (*filename, imageOnly=False*)

Read a MAR-345 image plate image

GSASIIIO.GetPowderPeaks (*fileName*)

Read powder peaks from a file

GSASIIIO.GetTifData (*filename, imageOnly=False*)

Read an image in a pseudo-tif format, as produced by a wide variety of software, almost always incorrectly in some way.

class GSASIIIO.ImportBaseclass (*formatName, longFormatName=None, extensionlist=[], strictExtension=False*)

Defines a base class for the reading of input files (diffraction data, coordinates,...). See [Writing a Import Routine](#) for an explanation on how to use a subclass of this class.

BlockSelector (*ChoiceList, ParentFrame=None, title='Select a block', size=None, header='Block Selector', useCancel=True*)

Provide a wx dialog to select a block if the file contains more than one set of data and one must be selected

CIFValidator (*filepointer*)

A `ContentsValidator()` for use to validate CIF files.

ContentsValidator (*filepointer*)

This routine will attempt to determine if the file can be read with the current format. This will typically be overridden with a method that takes a quick scan of [some of] the file contents to do a “sanity” check if the file appears to match the selected format. Expected to be called via `self.Validator()`

ExtensionValidator (*filename*)

This methods checks if the file has the correct extension Return False if this filename will not be supported by this reader Return True if the extension matches the list supplied by the reader Return None if the reader allows un-registered extensions

exception ImportError

Defines an Exception that is used when an import routine hits an expected error, usually in `.Reader`.

Good practice is that the Reader should define a value in `self.errors` that tells the user some information about what is wrong with their file.

`ImportBaseclass.MultipleBlockSelector` (*ChoiceList*, *ParentFrame=None*, *title='Select a block'*, *size=None*, *header='Block Selector'*)

Provide a wx dialog to select a block of data if the file contains more than one set of data and one must be selected.

Returns a list of the selected blocks

`ImportBaseclass.MultipleChoicesDialog` (*choicelist*, *headinglist*, *ParentFrame=None*, ***kwargs*)

A modal dialog that offers a series of choices, each with a title and a wx.Choice widget. Typical input:

- choicelist*=[('a','b','c'), ('test1','test2'), ('no choice',)]

- headinglist* = ['select a, b or c', 'select 1 of 2', 'No option here']

optional keyword parameters are: *head* (window title) and *title* returns a list of selected indices for each choice (or None)

class `GSASIIIO.ImportPhase` (*formatName*, *longFormatName=None*, *extensionlist=[]*, *strictExtension=False*)

Defines a base class for the reading of files with coordinates

Objects constructed that subclass this (in `import/G2phase_*.py`) will be used in `GSASII.GSASII.OnImportPhase()`. See [Writing a Import Routine](#) for an explanation on how to use this class.

PhaseSelector (*ChoiceList*, *ParentFrame=None*, *title='Select a phase'*, *size=None*, *header='Phase Selector'*)

Provide a wx dialog to select a phase if the file contains more than one phase

class `GSASIIIO.ImportPowderData` (*formatName*, *longFormatName=None*, *extensionlist=[]*, *strictExtension=False*)

Defines a base class for the reading of files with powder data. See [Writing a Import Routine](#) for an explanation on how to use this class.

powderdata = None

A powder data set is a list with items [x,y,w,yc,yb,yd]: `np.array(x)`, # x-axis values `np.array(y)`, # powder pattern intensities `np.array(w)`, # $1/\text{sig}(\text{intensity})^2$ values (weights) `np.array(yc)`, # calc. intensities (zero) `np.array(yb)`, # calc. background (zero) `np.array(yd)`, # obs-calc profiles

class `GSASIIIO.ImportStructFactor` (*formatName*, *longFormatName=None*, *extensionlist=[]*, *strictExtension=False*)

Defines a base class for the reading of files with tables of structure factors.

Structure factors are read with a call to `GSASII.GSASII.OnImportSfact()` which in turn calls `GSASII.GSASII.OnImportGeneric()`, which calls methods `ExtensionValidator()`, `ContentsValidator()` and `Reader()`.

See [Writing a Import Routine](#) for an explanation on how to use import classes in general. The specifics for reading a structure factor histogram require that the `Reader()` routine in the import class need to do only a few things: It should load `RefDict` item 'RefList' with the reflection list, and set `Parameters` with the instrument parameters (initialized with `InitParameters()` and set with `UpdateParameters()`). Also, set `Controls`, which specifies how the histogram is plotted (initialized with `InitControls()` and set with `UpdateControls()`).

Controls = None

`self.Controls` is a dict with plotting controls

InitControls()

initialize the controls structure

InitParameters()

initialize the instrument parameters structure

Parameters = None

self.Parameters is a list with two dicts for data parameter settings

RefDict = None

self.RefDict is a dict containing the reflection information, as read from the file. Item 'RefList' contains the reflection information. See the [Single Crystal Reflection Data Structure](#) for the contents of each row. Dict element 'FF' contains the form factor values for each element type; if this entry is left as initialized (an empty list) it will be initialized as needed later.

UpdateControls (*Type='Fosq', FcalcPresent=False*)

Scan through the reflections to update the Controls dictionary

UpdateParameters (*Type=None, Wave=None*)

Revise the instrument parameters

GSASIIIO.IndexPeakListSave (*G2frame, peaks*)

Save powder peaks from the indexing list

class GSASIIIO.MultipleChoicesDialog (*choicelist, headinglist, head='Select options', title='Please select from options below', parent=None*)

A dialog that offers a series of choices, each with a title and a wx.Choice widget. Intended to be used Modally. typical input:

- choicelist=[('a','b','c'), ('test1','test2'),('no choice',)]
- headinglist = ['select a, b or c', 'select 1 of 2', 'No option here']

selections are placed in self.chosen when OK is pressed

GSASIIIO.PDFSave (*G2frame, exports*)

Save a PDF G(r) and S(Q) in column formats

GSASIIIO.PeakListSave (*G2frame, file, peaks*)

Save powder peaks to a data file

GSASIIIO.ProjFileOpen (*G2frame*)

Read a GSAS-II project file and load into the G2 data tree

GSASIIIO.ProjFileSave (*G2frame*)

Save a GSAS-II project file

GSASIIIO.PutG2Image (*filename, Comments, Data, Npix, image*)

Write an image as a python pickle

GSASIIIO.ReadCIF (*URLorFile*)

Open a CIF, which may be specified as a file name or as a URL using PyCifRW (from James Hester). The open routine gets confused with DOS names that begin with a letter and colon "C:dir" so this routine will try to open the passed name as a file and if that fails, try it as a URL

Parameters **URLorFile** (*str*) – string containing a URL or a file name. Code will try first to open it as a file and then as a URL.

Returns a PyCifRW CIF object.

GSASIIIO.SaveIntegration (*G2frame, PickId, data*)

Save image integration results as powder pattern(s)

GSASIIIO.SetNewPhase (*Name='New Phase', SGData=None, cell=None*)

Create a new phase dict with default values for various parameters

Parameters

- **Name** (*str*) – Name for new Phase

- **SGData** (*dict*) – space group data from `GSASIIspc:SpcGroup()`; defaults to data for P 1
- **cell** (*list*) – unit cell parameter list; defaults to [1.0,1.0,1.0,90.,90,90.,1.]

`GSASIIIO.powderFxyeSave` (*G2frame, exports, powderfile*)
Save a powder histogram as a GSAS FXYE file

`GSASIIIO.powderXyeSave` (*G2frame, exports, powderfile*)
Save a powder histogram as a Topas XYE file

`GSASIIIO.sfloat` (*S*)
Convert a string to float. An empty field is treated as zero

`GSASIIIO.sint` (*S*)
Convert a string to int. An empty field is treated as zero

`GSASIIIO.trim` (*val*)
Simplify a string containing leading and trailing spaces as well as newlines, tabs, repeated spaces etc. into a shorter and more simple string, by replacing all ranges of whitespace characters with a single space.

Parameters *val* (*str*) – the string to be simplified

Returns the (usually) shortened version of the string

4.3 ReadMarCCDFrame: Read Mar Files

`class ReadMarCCDFrame.marFrame` (*File, byteOrd='<', IFD={}*)
A class to extract correct mar header and image info from a MarCCD file

Parameters

- **File** (*str*) – file object [from `open()`]
- **byteOrd** – ‘<’ (default) or ‘>’
- **IFD** (*dict*) – ?

4.4 GSASIIpy3: Python 3.x Routines

Module to hold python 3-compatible code, to keep it separate from code that will break with `__future__` options.

`GSASIIpy3.FormatValue` (*val, maxdigits=10*)
Format a float to fit in `maxdigits` spaces, showing as much precision as possible, more or less.

Parameters

- **val** (*float*) – number to be formatted.
- **maxdigits** (*int*) – the number of digits to be used for display of the number (defaults to 10).

Returns a string with \leq `maxdigits` characters (I hope).

`GSASIIpy3.FormulaEval` (*string*)
Evaluates a algebraic formula into a float, if possible. Works properly on fractions e.g. $2/3$ only with python 3.0+ division.

Expressions such as $2/3$, $3*\pi$, $\sin(45)/2$, $2*\sqrt{2}$, $2**10$ can all be evaluated.

Parameters *string* (*str*) – Character string containing a Python expression to be evaluated.

Returns the value for the expression as a float or None if the expression does not evaluate to a valid number.

GSAS-II GUI SUBMODULES

5.1 GSASIIphsGUI: Phase GUI

Module to create the GUI for display of phase information in the data display window when a phase is selected. Phase information is stored in one or more *Phase Tree Item* objects. Note that there are functions that respond to some tabs in the phase GUI in other modules (such as GSASIIddata).

GSASIIphsGUI.**UpdatePhaseData** (*G2frame*, *Item*, *data*, *oldPage*)

Create the data display window contents when a phase is clicked on in the main (data tree) window. Called only from `GSASIIgrid.MovePatternTreeToGrid()`, which in turn is called from `GSASII.GSASII.OnPatternTreeSelChanged()` when a tree item is selected.

Parameters

- **G2frame** (*wx.frame*) – the main GSAS-II frame object
- **Item** (*wx.TreeItemId*) – the tree item that was selected
- **data** (*dict*) – all the information on the phase in a dictionary
- **oldPage** (*int*) – This sets a tab to select when moving from one phase to another, in which case the same tab is selected to display first. This is set only when the previous data tree selection is a phase, if not the value is None. The default action is to bring up the General tab.

5.2 GSASIIddataGUI: Phase Diffraction Data GUI

Module to create the GUI for display of diffraction data * phase information that is shown in the data display window (when a phase is selected.)

GSASIIddataGUI.**UpdateDDData** (*G2frame*, *DDData*, *data*)

Display the Diffraction Data associated with a phase (items where there is a value for each histogram and phase)

Parameters

- **G2frame** (*wx.frame*) – the main GSAS-II frame object
- **DDData** (*wx.ScrolledWindow*) – notebook page to be used for the display
- **data** (*dict*) – all the information on the phase in a dictionary

5.3 GSASIIElemGUI: GUI to select and delete element lists

Module to select elements from a periodic table and to delete an element from a list of selected elements.

```
class GSASIIElemGUI.DeleteElement (parent, choice)
    Delete element from selected set widget

    ElButton (id, name, pos)
        Needs a doc string

class GSASIIElemGUI.PickElement (parent, oneOnly=False, ifNone=False)
    Makes periodic table widget for picking element - caller maintains element list

    ElButton (name, pos, tip, color)
        Needs a doc string
```

5.4 GSASIIconstrGUI: Constraint GUI routines

Used to define constraints and rigid bodies.

```
class GSASIIconstrGUI.ConstraintDialog (parent, title, text, data, separator='*', varname='',
                                         varyflag=False)
    Window to edit Constraint values

class GSASIIconstrGUI.MultiIntegerDialog (parent, title, prompts, values)
    Input a series of integers based on prompts

GSASIIconstrGUI.UpdateConstraints (G2frame, data)
    Called when Constraints tree item is selected. Displays the constraints in the data window

GSASIIconstrGUI.UpdateRigidBodies (G2frame, data)
    Called when Rigid bodies tree item is selected. Displays the rigid bodies in the data window
```

5.5 GSASIIimgGUI: Image GUI

Control image display and processing

```
GSASIIimgGUI.UpdateImageControls (G2frame, data, masks)
    Shows and handles the controls on the "Image Controls" data tree entry

GSASIIimgGUI.UpdateMasks (G2frame, data)
    Shows and handles the controls on the "Masks" data tree entry

GSASIIimgGUI.UpdateStressStrain (G2frame, data)
    Shows and handles the controls on the "Stress/Strain" data tree entry
```

5.6 GSASIIpwdGUI: Powder Pattern GUI routines

Used to define GUI controls for the routines that interact with the powder histogram (PWDR) data tree items.

```
GSASIIpwdGUI.IsHistogramInAnyPhase (G2frame, histoName)
    Needs a doc string

GSASIIpwdGUI.SetDefaultSample ()
    Needs a doc string
```

`GSASIIpwdGUI.UpdateBackground (G2frame, data)`
respond to selection of PWDR background data tree item.

`GSASIIpwdGUI.UpdateIndexPeaksGrid (G2frame, data)`
respond to selection of PWDR Index Peak List data tree item.

`GSASIIpwdGUI.UpdateInstrumentGrid (G2frame, data)`
respond to selection of PWDR Instrument Parameters data tree item.

`GSASIIpwdGUI.UpdateLimitsGrid (G2frame, data)`
respond to selection of PWDR Limits data tree item.

`GSASIIpwdGUI.UpdatePDFGrid (G2frame, data)`
respond to selection of PWDR PDF data tree item.

`GSASIIpwdGUI.UpdatePeakGrid (G2frame, data)`
respond to selection of PWDR powder peaks data tree item.

`GSASIIpwdGUI.UpdateReflectionGrid (G2frame, data, HKLF=False, Name='')`
respond to selection of PWDR Reflections data tree item.

`GSASIIpwdGUI.UpdateSampleGrid (G2frame, data)`
respond to selection of PWDR Sample Parameters data tree item.

`GSASIIpwdGUI.UpdateUnitCellsGrid (G2frame, data)`
respond to selection of PWDR Unit Cells data tree item.

5.7 GSASIIrestrGUI: Restraint GUI routines

Used to define restraints.

`GSASIIrestrGUI.UpdateRestraints (G2frame, data, Phases, phaseName)`
Respond to selection of the Restraints item on the data tree

GSAS-II STRUCTURE SUBMODULES

6.1 GSASIIstrMain: main structure routine

GSASIIstrMain.**BestPlane** (*PlaneData*)

Needs a doc string

GSASIIstrMain.**DisAglTor** (*DATData*)

Needs a doc string

GSASIIstrMain.**PrintDistAngle** (*DisAglCtls*, *DisAglData*, *out*=<open file '<stdout>', mode 'w' at 0x240078>)

Print distances and angles

Parameters

- **DisAglCtls** (*dict*) – contains distance/angle radii usually defined using `GSASIIgrid.DisAglDialog()`
- **DisAglData** (*dict*) – contains phase data: Items 'OrigAtoms' and 'TargAtoms' contain the atoms to be used for distance/angle origins and atoms to be used as targets. Item 'SGData' has the space group information (see *Space Group object*)
- **out** (*file*) – file object for output. Defaults to sys.stdout.

GSASIIstrMain.**Refine** (*GPXfile*, *dlg*)

Needs a doc string

GSASIIstrMain.**RetDistAngle** (*DisAglCtls*, *DisAglData*)

Compute and return distances and angles

Parameters

- **DisAglCtls** (*dict*) – contains distance/angle radii usually defined using `GSASIIgrid.DisAglDialog()`
- **DisAglData** (*dict*) – contains phase data: Items 'OrigAtoms' and 'TargAtoms' contain the atoms to be used for distance/angle origins and atoms to be used as targets. Item 'SGData' has the space group information (see *Space Group object*)

Returns

AtomLabels, DistArray, AngArray where:

AtomLabels is a dict of atom labels, keys are the atom number

DistArray is a dict keyed by the origin atom number where the value is a list of distance entries. The value for each distance is a list containing:

0. the target atom number (int);
1. the unit cell offsets added to x,y & z (tuple of int values)
2. the symmetry operator number (which may be modified to indicate centering and center of symmetry)
3. an interatomic distance in A (float)
4. an uncertainty on the distance in A or 0.0 (float)

AngArray is a dict keyed by the origin (central) atom number where the value is a list of angle entries. The value for each angle entry consists of three values:

0. a distance item reference for one neighbor (int)
1. a distance item reference for a second neighbor (int)
2. a angle, uncertainty pair; the s.u. may be zero (tuple of two floats)

The AngArray distance reference items refer directly to the index of the items in the DistArray item for the list of distances for the central atom.

`GSASIIstrMain.SeqRefine (GPXfile, dlg)`

Needs a doc string

`GSASIIstrMain.main ()`

Needs a doc string

6.2 GSASIIstrMath - structure math routines

`GSASIIstrMath.ApplyRBModelDerivs (dFdvDict, parmDict, rigidbodyDict, Phase)`

Needs a doc string

`GSASIIstrMath.ApplyRBModels (parmDict, Phases, rigidbodyDict, Update=False)`

Takes RB info from RBModels in Phase and RB data in rigidbodyDict along with current RB values in parmDict & modifies atom contents (xyz & Uij) of parmDict

`GSASIIstrMath.ApplyXYZshifts (parmDict, varyList)`

takes atom x,y,z shift and applies it to corresponding atom x,y,z value

Parameters

- **parmDict** (*dict*) – parameter dictionary
- **varyList** (*list*) – list of variables

Returns newAtomDict - dictionary of new atomic coordinate names & values; key is parameter shift name

`GSASIIstrMath.Dict2Values (parmdict, varylist)`

Use before call to leastsq to setup list of values for the parameters in parmdict, as selected by key in varylist

`GSASIIstrMath.GetAbsorb (refl, hfx, calcControls, parmDict)`

Needs a doc string

`GSASIIstrMath.GetAbsorbDerv (refl, hfx, calcControls, parmDict)`

Needs a doc string

`GSASIIstrMath.GetAtomFXU (pfx, calcControls, parmDict)`

Needs a doc string

`GSASIIstrMath.GetFobsSq` (*Histograms, Phases, parmDict, calcControls*)

Needs a doc string

`GSASIIstrMath.GetHStrainShift` (*refl, SGData, phfx, parmDict*)

Needs a doc string

`GSASIIstrMath.GetHStrainShiftDerv` (*refl, SGData, phfx*)

Needs a doc string

`GSASIIstrMath.GetIntensityCorr` (*refl, uniq, G, g, pfx, phfx, hfx, SGData, calcControls, parmDict*)

Needs a doc string

`GSASIIstrMath.GetIntensityDerv` (*refl, uniq, G, g, pfx, phfx, hfx, SGData, calcControls, parmDict*)

Needs a doc string

`GSASIIstrMath.GetNewCellParms` (*parmDict, varyList*)

Needs a doc string

`GSASIIstrMath.GetPrefOri` (*refl, uniq, G, g, phfx, hfx, SGData, calcControls, parmDict*)

Needs a doc string

`GSASIIstrMath.GetPrefOriDerv` (*refl, uniq, G, g, phfx, hfx, SGData, calcControls, parmDict*)

Needs a doc string

`GSASIIstrMath.GetReflPos` (*refl, wave, G, hfx, calcControls, parmDict*)

Needs a doc string

`GSASIIstrMath.GetReflPosDerv` (*refl, wave, A, hfx, calcControls, parmDict*)

Needs a doc string

`GSASIIstrMath.GetSampleSigGam` (*refl, wave, G, GB, phfx, calcControls, parmDict*)

Needs a doc string

`GSASIIstrMath.GetSampleSigGamDerv` (*refl, wave, G, GB, phfx, calcControls, parmDict*)

Needs a doc string

`GSASIIstrMath.HessRefine` (*values, HistoPhases, parmDict, varylist, calcControls, pawleyLookup, dlg*)

Loop over histograms and compute derivatives of the fitting model (M) with respect to all parameters. For each histogram, the Jacobian matrix, dMdv, with dimensions (n by m) where n is the number of parameters and m is the number of data points *in the histogram*. The (n by n) Hessian is computed from each Jacobian and it is returned. This routine is used when refinement derivatives are selected as “analytic Hessian” in Controls.

Returns Vec,Hess where Vec is the least-squares vector and Hess is the Hessian

`GSASIIstrMath.SCExtinction` (*ref, phfx, hfx, pfx, calcControls, parmDict, varyList*)

Single crystal extinction function; puts correction in ref[13] and returns corrections needed for derivatives

`GSASIIstrMath.SHPOcal` (*refl, g, phfx, hfx, SGData, calcControls, parmDict*)

spherical harmonics preferred orientation (cylindrical symmetry only)

`GSASIIstrMath.SHPOcalDerv` (*refl, g, phfx, hfx, SGData, calcControls, parmDict*)

spherical harmonics preferred orientation derivatives (cylindrical symmetry only)

`GSASIIstrMath.SHTXcal` (*refl, g, pfx, hfx, SGData, calcControls, parmDict*)

Spherical harmonics texture

`GSASIIstrMath.SHTXcalDerv` (*refl, g, pfx, hfx, SGData, calcControls, parmDict*)

Spherical harmonics texture derivatives

`GSASIIstrMath.StructureFactor` (*refDict, G, hfx, pfx, SGData, calcControls, parmDict*)

Compute structure factors for all h,k,l for phase puts the result, F^2, in each ref[8] in refList input:

Parameters

- **refDict** (*dict*) – where ‘RefList’ list where each ref = h,k,l,m,d,... ‘FF’ dict of form factors - filed in below
- **G** (*np.array*) – reciprocal metric tensor
- **pfx** (*str*) – phase id string
- **SGData** (*dict*) – space group info. dictionary output from SpcGroup
- **calcControls** (*dict*) –
- **ParmDict** (*dict*) –

GSASIIstrMath.**StructureFactor2** (*refDict, G, hfx, pfx, SGData, calcControls, parmDict*)

Compute structure factors for all h,k,l for phase puts the result, F^2 , in each ref[8] in refList input:

Parameters

- **refDict** (*dict*) – where ‘RefList’ list where each ref = h,k,l,m,d,... ‘FF’ dict of form factors - filed in below
- **G** (*np.array*) – reciprocal metric tensor
- **pfx** (*str*) – phase id string
- **SGData** (*dict*) – space group info. dictionary output from SpcGroup
- **calcControls** (*dict*) –
- **ParmDict** (*dict*) –

GSASIIstrMath.**StructureFactorDerv** (*refDict, G, hfx, pfx, SGData, calcControls, parmDict*)

Needs a doc string

GSASIIstrMath.**Values2Dict** (*parmdict, varylist, values*)

Use after call to leastsq to update the parameter dictionary with values corresponding to keys in varylist

GSASIIstrMath.**dervRefine** (*values, HistoPhases, parmDict, varylist, calcControls, pawleyLookup, dlg*)

Loop over histograms and compute derivatives of the fitting model (M) with respect to all parameters. Results are returned in a Jacobian matrix (aka design matrix) of dimensions (n by m) where n is the number of parameters and m is the number of data points. This can exceed memory when m gets large. This routine is used when refinement derivatives are selected as “analytic Jacobian” in Controls.

Returns Jacobian numpy.array dMdv for all histograms concatenated

GSASIIstrMath.**errRefine** (*values, HistoPhases, parmDict, varylist, calcControls, pawleyLookup, dlg*)

Needs a doc string

GSASIIstrMath.**getPowderProfile** (*parmDict, x, varylist, Histogram, Phases, calcControls, pawleyLookup*)

Needs a doc string

GSASIIstrMath.**getPowderProfileDerv** (*parmDict, x, varylist, Histogram, Phases, rigidbodyDict, calcControls, pawleyLookup*)

Needs a doc string

GSASIIstrMath.**penaltyDeriv** (*pNames, pVal, HistoPhases, parmDict, varyList*)

Needs a doc string

GSASIIstrMath.**penaltyFxn** (*HistoPhases, parmDict, varyList*)

Needs a doc string

6.3 GSASIIstrIO: structure I/O routines

GSASIIstrIO.CheckConstraints (*GPXfile*)

Load constraints and related info and return any error or warning messages

GSASIIstrIO.GPXBackup (*GPXfile*, *makeBack=True*)

makes a backup of the current .gpx file (?)

Parameters

- **GPXfile** (*str*) – full .gpx file name
- **makeBack** (*bool*) – if True (default), the backup is written to a new file; if False, the last backup is overwritten

Returns the name of the backup file that was written

GSASIIstrIO.GetAllPhaseData (*GPXfile*, *PhaseName*)

Returns the entire dictionary for PhaseName from GSASII gpx file

Parameters

- **GPXfile** (*str*) – full .gpx file name
- **PhaseName** (*str*) – phase name

Returns phase dictionary

GSASIIstrIO.GetConstraints (*GPXfile*)

Read the constraints from the GPX file and interpret them

called in `CheckConstraints()`, `GSASIIstrMain.Refine()` and `GSASIIstrMain.SeqRefine()`.

GSASIIstrIO.GetControls (*GPXfile*)

Returns dictionary of control items found in GSASII gpx file

Parameters **GPXfile** (*str*) – full .gpx file name

Returns dictionary of control items

GSASIIstrIO.GetFprime (*controlDict*, *Histograms*)

Needs a doc string

GSASIIstrIO.GetHistogramData (*Histograms*, *Print=True*, *pFile=None*)

needs a doc string

GSASIIstrIO.GetHistogramNames (*GPXfile*, *hType*)

Returns a list of histogram names found in GSASII gpx file

Parameters

- **GPXfile** (*str*) – full .gpx file name
- **hType** (*str*) – list of histogram types

Returns list of histogram names (types = PWDR & HKLF)

GSASIIstrIO.GetHistogramPhaseData (*Phases*, *Histograms*, *Print=True*, *pFile=None*, *resetRefList=True*)

Loads the HAP histogram/phase information into dicts

Parameters

- **Phases** (*dict*) – phase information

- **Histograms** (*dict*) – Histogram information
- **Print** (*bool*) – prints information as it is read
- **pFile** (*file*) – file object to print to (the default, None causes printing to the console)
- **resetRefList** (*bool*) – Should the contents of the Reflection List be initialized on loading. The default, True, initializes the Reflection List as it is loaded.

Returns (hapVary,hapDict,controlDict) * hapVary: list of refined variables * hapDict: dict with refined variables and their values * controlDict: dict with computation controls (?)

GSASIIstrIO.**GetHistograms** (*GPXfile, hNames*)

Returns a dictionary of histograms found in GSASII gpx file

Parameters

- **GPXfile** (*str*) – full .gpx file name
- **hNames** (*str*) – list of histogram names

Returns dictionary of histograms (types = PWDR & HKLF)

GSASIIstrIO.**GetPawleyConstr** (*SGLaue, PawleyRef, pawleyVary*)

needs a doc string

GSASIIstrIO.**GetPhaseData** (*PhaseData, RestraintDict={}, rIds={}, Print=True, pFile=None*)

needs a doc string

GSASIIstrIO.**GetPhaseNames** (*GPXfile*)

Returns a list of phase names found under 'Phases' in GSASII gpx file

Parameters **GPXfile** (*str*) – full .gpx file name

Returns list of phase names

GSASIIstrIO.**GetRestrains** (*GPXfile*)

Read the restrains from the GPX file. Throws an exception if not found in the .GPX file

GSASIIstrIO.**GetRigidBodies** (*GPXfile*)

Read the rigid body models from the GPX file

GSASIIstrIO.**GetRigidBodyModels** (*rigidbodyDict, Print=True, pFile=None*)

needs a doc string

GSASIIstrIO.**GetUsedHistogramsAndPhases** (*GPXfile*)

Returns all histograms that are found in any phase and any phase that uses a histogram. This also assigns numbers to used phases and histograms by the order they appear in the file.

Parameters **GPXfile** (*str*) – full .gpx file name

Returns

(Histograms,Phases)

- Histograms = dictionary of histograms as {name:data,...}
- Phases = dictionary of phases that use histograms

GSASIIstrIO.**PrintRestrains** (*cell, SGData, AtPtrs, Atoms, AtLookup, textureData, phaseRest, pFile*)

needs a doc string

GSASIIstrIO.**ProcessConstraints** (*constList*)

Interpret the constraints in the constList input into a dictionary, etc. All GSASIIobj.G2VarObj objects are mapped to the appropriate phase/hist/atoms based on the object internals (random Ids). If this can't be done (if

a phase has been deleted, etc.), the variable is ignored. If the constraint cannot be used due to too many dropped variables, it is counted as ignored.

Parameters **constList** (*list*) – a list of lists where each item in the outer list specifies a constraint of some form, as described in the [GSASIIobj Constraint definition](#).

Returns

a tuple of (constDict, fixedList, ignored) where:

- **constDict** (list of dicts) contains the constraint relationships
- **fixedList** (list) contains the fixed values for each type of constraint.
- **ignored** (int) counts the number of invalid constraint items (should always be zero!)

GSASIIstrIO.SetHistogramData (*parmDict, sigDict, Histograms, Print=True, pFile=None*)
needs a doc string

GSASIIstrIO.SetHistogramPhaseData (*parmDict, sigDict, Phases, Histograms, Print=True, pFile=None*)
needs a doc string

GSASIIstrIO.SetPhaseData (*parmDict, sigDict, Phases, RBIds, covData, RestraintDict=None, pFile=None*)
needs a doc string

GSASIIstrIO.SetRigidBodyModels (*parmDict, sigDict, rigidbodyDict, pFile=None*)
needs a doc string

GSASIIstrIO.SetSeqResult (*GPXfile, Histograms, SeqResult*)
Needs doc string

Parameters **GPXfile** (*str*) – full .gpx file name

GSASIIstrIO.SetUsedHistogramsAndPhases (*GPXfile, Histograms, Phases, RigidBodyes, CovData, makeBack=True*)

Updates gpxfile from all histograms that are found in any phase and any phase that used a histogram. Also updates rigid body definitions.

Parameters

- **GPXfile** (*str*) – full .gpx file name
- **Histograms** (*dict*) – dictionary of histograms as {name:data,...}
- **Phases** (*dict*) – dictionary of phases that use histograms
- **RigidBodyes** (*dict*) – dictionary of rigid bodies
- **CovData** (*dict*) – dictionary of refined variables, varyList, & covariance matrix
- **makeBack** (*bool*) – True if new backup of .gpx file is to be made; else use the last one made

GSASIIstrIO.ShowBanner (*pFile=None*)
Print authorship, copyright and citation notice

GSASIIstrIO.ShowControls (*Controls, pFile=None*)
Print controls information

GSASIIstrIO.cellFill (*pfx, SGData, parmDict, sigDict*)
Returns the filled-out reciprocal cell (A) terms and their uncertainties from the parameter and sig dictionaries.

Parameters

- **pfx** (*str*) – parameter prefix (“n:”, where n is a phase number)

- **SGdata** (*dict*) – a symmetry object
- **parmDict** (*dict*) – a dictionary of parameters
- **sigDict** (*dict*) – a dictionary of uncertainties on parameters

Returns A,sigA where each is a list of six terms with the A terms

GSASIIstrIO.**cellVary** (*pfx, SGData*)
needs a doc string

GSASIIstrIO.**getBackupName** (*GPXfile, makeBack*)
Get the name for the backup .gpx file name

Parameters

- **GPXfile** (*str*) – full .gpx file name
- **makeBack** (*bool*) – if True the name of a new file is returned, if False the name of the last file that exists is returned

Returns the name of a backup file

GSASIIstrIO.**getCellEsd** (*pfx, SGData, A, covData*)
needs a doc string

GSASIIMAPVARS: PARAMETER CONSTRAINTS

Module to implements algebraic constraints, parameter redefinition and parameter simplification constraints.

Parameter redefinition (new vars) is done by creating one or more relationships between a set of parameters

```
Mx1 * Px + My1 * Py + ...  
Mx2 * Px + Mz2 * Pz + ...
```

where P_j is a parameter name and M_{jk} is a constant.

Constant constraint Relations can also be supplied in the form of an equation:

```
nx1 * Px + ny1 * Py + ... = C1
```

where C_n is a constant. These equations define an algebraic constraint.

Parameters can also be “fixed” (held), which prevents them from being refined.

All of the above three cases are input using routines GroupConstraints and GenerateConstraints. The input consists of a list of relationship dictionaries:

```
constrDict = [  
    {'0:12:Scale': 2.0, '0:14:Scale': 4.0, '0:13:Scale': 3.0, '0:0:Scale': 0.5},  
    {'2::C(10,6,1)': 1.0, '1::C(10,6,1)': 1.0},  
    {'0::A0': 0.0}]  
fixedList = ['5.0', None, '0']
```

Where the dictionary defines the first part of an expression and the corresponding fixedList item is either None (for parameter redefinition) or the constant value, for a constant constraint equation. A dictionary that contains a single term defines a variable that will be fixed (held). The multiplier and the fixedList value in this case are ignored.

Parameters can also be equivalenced or “slaved” to another parameter, such that one (independent) parameter is equated to several (now dependent) parameters. In algebraic form this is:

```
P0 = M1 * P1 = M2 * P2 = ...
```

Thus parameters P_0, P_1 and P_2, \dots are linearly equivalent. Routine StoreEquivalence is used to specify these equivalences.

Parameter redefinition (new vars) describes a new, independent, parameter, which is defined in terms of dependent parameters that are defined in the Model, while fixed constrained relations effectively reduce the complexity of the Model by removing a degree of freedom. It is possible for a parameter to appear in both a parameter redefinition expression and a fixed constraint equation, but a parameter cannot be used a parameter equivalence cannot be used elsewhere (not fixed, constrained or redefined). Likewise a fixed parameter cannot be used elsewhere (not equivalenced, constrained or redefined).

Relationships are grouped so that a set of dependent parameters appear in only one group (done in routine GroupConstraints.) Note that if a group contains relationships/equations that involve N dependent parameters, there must exist N-C new parameters, where C is the number of constraint equations in the group. Routine GenerateConstraints takes the output from GroupConstraints and generates the “missing” relationships and saves that information in the module’s global variables. Each generated parameter is named sequentially using paramPrefix.

A list of parameters that will be varied is specified as input to GenerateConstraints (varyList). A fixed parameter will simply be removed from this list preventing that parameter from being varied. Note that all parameters in a constraint relationship must specified as varied (appear in varyList) or none can be varied. This is checked in GenerateConstraints (as well as for generated relationships in SetVaryFlags).

- When a new variable is created, the variable is assigned the name associated in the constraint definition or it is assigned a default name of form `: : constr<n>` (see paramPrefix). The vary setting for variables used in the constraint are ignored. Note that any generated “missing” relations are not varied. Only the input relations can be varied.
- If all parameters in a fixed constraint equation are varied, the generated “missing” relations in the group are all varied. This provides the N-C degrees of freedom.

7.1 External Routines

To define a set of constrained and unconstrained relations, one defines a list of dictionary defining constraint parameters and their values, a list of fixed values for each constraint and a list of parameters to be varied. In addition, one uses `StoreEquivalence()` to define parameters that are equivalent. One can then use `CheckConstraints()` to check that the input is internally consistent and finally `GroupConstraints()` and `GenerateConstraints()` to generate the internally used tables. Routines `Map2Dict()` is used to initialize the parameter dictionary and `Dict2Map()`, `Dict2Deriv()`, and `ComputeDepESD()` are used to apply constraints. Routine `VarRemapShow()` is used to print out the constraint information, as stored by `GenerateConstraints()`.

InitVars() This is optionally used to clear out all defined previously defined constraint information

StoreEquivalence() To implement parameter redefinition, one calls StoreEquivalence. This should be called for every set of equivalence relationships. There is no harm in using StoreEquivalence with the same independent variable:

```
StoreEquivalence('x', ('y',))
StoreEquivalence('x', ('z',))
```

or equivalently

```
StoreEquivalence('x', ('y', 'z'))
```

The latter will run more efficiently. Note that mixing independent and dependent variables is problematic. This is not allowed:

```
StoreEquivalence('x', ('y',))
StoreEquivalence('y', ('z',))
```

Use StoreEquivalence before calling GenerateConstraints or CheckConstraints

CheckConstraints() To check that input is internally consistent, use CheckConstraints

Map2Dict() To determine values for the parameters created in this module, one calls Map2Dict. This will not apply constraints.

Dict2Map() To take values from the new independent parameters and constraints, one calls Dict2Map. This will apply constraints.

Dict2Deriv() Use Dict2Deriv to determine derivatives on independent parameters from those on dependent ones

ComputeDepESD() Use ComputeDepESD to compute uncertainties on dependent variables

VarRemapShow() To show a summary of the parameter remapping, one calls VarRemapShow

7.2 Global Variables

dependentParmList: contains a list by group of lists of parameters used in the group. Note that parameters listed in dependentParmList should not be refined as they will not affect the model

indParmList: a list of groups of Independent parameters defined in each group. This contains both parameters used in parameter redefinitions as well as names of generated new parameters.

fixedVarList: a list of variables that have been ‘fixed’ by defining them as equal to a constant (`::var: = 0`). Note that the constant value is ignored at present. These variables are later removed from varyList which prevents them from being refined. Unlikely to be used externally.

arrayList: a list by group of relationship matrices to relate parameters in dependentParmList to those in indParmList. Unlikely to be used externally.

invarrayList: a list by group of relationship matrices to relate parameters in indParmList to those in dependentParmList. Unlikely to be used externally.

fixedDict: a dictionary containing the fixed values corresponding to parameter equations. The dict key is an ascii string, but the dict value is a float. Unlikely to be used externally.

7.3 Routines

Note that parameter names in GSAS-II are strings of form `<ph>:<hst>:<nam>`

`GSASIImapvars.CheckConstraints (varyList, constrDict, fixedList)`

Takes a list of relationship entries comprising a group of constraints and checks for inconsistencies such as conflicts in parameter/variable definitions and or inconsistently varied parameters.

Parameters

- **varyList** (*list*) – a list of parameters names that will be varied
- **constrDict** (*dict*) – a list of dicts defining relationships/constraints (as created in `GSASIIstrIO.ProcessConstraints()` and documented in `GroupConstraints()`)
- **fixedList** (*list*) – a list of values specifying a fixed value for each dict in constrDict. Values are either strings that can be converted to floats or `None` if the constraint defines a new parameter rather than a constant.

Returns

two strings:

- the first lists conflicts internal to the specified constraints
- the second lists conflicts where the varyList specifies some parameters in a constraint, but not all

If there are no errors, both strings will be empty

`GSASIImapvars.ComputeDepESD(covMatrix, varyList, parmDict)`

Compute uncertainties for dependent parameters from independent ones returns a dictionary containing the esd values for dependent parameters

`GSASIImapvars.Dict2Deriv(varyList, derivDict, dMdv)`

Compute derivatives for Independent Parameters from the derivatives for the original parameters

Parameters

- **varyList** (*list*) – a list of parameters names that will be varied
- **derivDict** (*dict*) – a dict containing derivatives for parameter values keyed by the parameter names.
- **dMdv** (*list*) – a Jacobian, as a list of np.array containing derivatives for dependent parameter computed from derivDict

`GSASIImapvars.Dict2Map(parmDict, varyList)`

Applies the constraints defined using `StoreEquivalence()`, `GroupConstraints()` and `GenerateConstraints()` by changing values in a dict containing the parameters. This should be done before the parameters are used for any computations

Parameters

- **parmDict** (*dict*) – a dict containing parameter values keyed by the parameter names. This will contain updated values for both dependent and independent parameters after Dict2Map is called. It will also contain some unexpected entries of every constant value {'0':0.0} & {'1.0':1.0}, which do not cause any problems.
- **varyList** (*list*) – a list of parameters names that will be varied

`GSASIImapvars.GenerateConstraints(groups, parmlist, varyList, constrDict, fixedList)`

Takes a list of relationship entries comprising a group of constraints and builds the relationship lists and their inverse and stores them in global variables Also checks for internal conflicts or inconsistencies in parameter/variable definitions.

Parameters

- **groups** (*list*) – a list of grouped constraints where each constraint grouped contains a list of indices for constraint constrDict entries, created in `GroupConstraints()` (returned as 1st value)
- **parmlist** (*list*) – a list containing lists of parameter names contained in each group, created in `GroupConstraints()` (returned as 2nd value)
- **varyList** (*list*) – a list of parameters names (strings of form <ph>:<hst>:<nam>) that will be varied
- **constrDict** (*dict*) – a list of dicts defining relationships/constraints (as defined in `GroupConstraints()`)
- **fixedList** (*list*) – a list of values specifying a fixed value for each dict in constrDict. Values are either strings that can be converted to floats, float values or None if the constraint defines a new parameter.
- **constrDict** – a list of dicts defining relationships/constraints.

`GSASIImapvars.GetDependentVars()`

Return a list of dependent variables: e.g. variables that are constrained in terms of other variables

Returns a list of variable names

`GSASIImapvars.GetIndependentVars()`

Return a list of independent variables: e.g. variables that are created by constraints of other variables

Returns a list of variable names

`GSASIImpvars.GramSchmidtOrtho(a, nkeep=0)`

Use the Gram-Schmidt process (<http://en.wikipedia.org/wiki/Gram-Schmidt>) to find orthonormal unit vectors relative to first row.

If nkeep is non-zero, the first nkeep rows in the array are not changed

input: arrayin: a 2-D non-singular square array

returns: a orthonormal set of unit vectors as a square array

`GSASIImpvars.GroupConstraints(constrDict)`

divide the constraints into groups that share no parameters.

Parameters `constrDict` (*dict*) – a list of dicts defining relationships/constraints

`constrDict = [{<constr1>}, {<constr2>}, ...]`

where {<constr1>} is {‘var1’: mult1, ‘var2’: mult2,... }

Returns

two lists of lists:

- a list of grouped constraints where each constraint grouped contains a list of indices for constraint `constrDict` entries
- a list containing lists of parameter names contained in each group

`GSASIImpvars.InitVars()`

Initializes all constraint information

`GSASIImpvars.Map2Dict(parmDict, varyList)`

Create (or update) the Independent Parameters from the original set of Parameters

Removes dependent variables from the `varyList`

This should be done once, after the constraints have been defined using `StoreEquivalence()`, `GroupConstraints()` and `GenerateConstraints()` and before any variable refinement is done to complete the parameter dictionary by defining independent parameters and satisfying the constraint equations.

Parameters

- **parmDict** (*dict*) – a dict containing parameter values keyed by the parameter names. This will contain updated values for both dependent and independent parameters after `Dict2Map` is called. It will also contain some unexpected entries of every constant value {‘0’:0.0} & {‘1.0’:1.0}, which do not cause any problems.
- **varyList** (*list*) – a list of parameters names that will be varied

`GSASIImpvars.PrintIndependentVars(parmDict, varyList, sigDict, PrintAll=False, pFile=None)`

Print the values and uncertainties on the independent variables

`GSASIImpvars.StoreEquivalence(independentVar, dependentList)`

Takes a list of dependent parameter(s) and stores their relationship to a single independent parameter (`independentVar`)

Parameters

- **independentVar** (*str*) – name of master parameter that will be used to determine the value to set the dependent variables
- **dependentList** (*list*) – a list of parameters that will set from `independentVar`. Each item in the list can be a string with the parameter name or a tuple containing a name and multiplier: `['parm1', ('parm2', .5),]`

`GSASIImapvars.VarKeys` (*constr*)

Finds the keys in a constraint that represent variables e.g. eliminates any that start with ‘_’

Parameters `constr` (*dict*) – a single constraint entry of form:

```
{'var1': mult1, 'var2': mult2, ... '_notVar': val, ...}
```

(see `GroupConstraints()`)

Returns a list of keys where any keys beginning with ‘_’ are removed.

`GSASIImapvars.VarRemapShow` (*varyList*)

List out the saved relationships. This should be done after the constraints have been defined using `StoreEquivalence()`, `GroupConstraints()` and `GenerateConstraints()`.

Returns a string containing the details of the constraint relationships

GSASIIIMAGE: IMAGE CALC MODULE

Ellipse fitting & image integration

GSASIIimage.**EdgeFinder** (*image, data*)

this makes list of all x,y where I>edgeMin suitable for an ellipse search? Not currently used but might be useful in future?

GSASIIimage.**Fill2ThetaAzimuthMap** (*masks, TA, tam, image*)

Needs a doc string

GSASIIimage.**FitCircle** (*ring*)

Needs a doc string

GSASIIimage.**FitDetector** (*rings, varyList, parmDict, Print=True*)

Needs a doc string

GSASIIimage.**FitEllipse** (*ring*)

Needs a doc string

GSASIIimage.**FitRing** (*ring, delta*)

Needs a doc string

GSASIIimage.**FitStrSta** (*Image, StrSta, Controls, Masks*)

Needs a doc string

GSASIIimage.**FitStrain** (*rings, p0, wave, phi*)

Needs a doc string

GSASIIimage.**GetAzm** (*x, y, data*)

Give azimuth value for detector x,y position; calibration info in data

GSASIIimage.**GetDetXYfromThAzm** (*Th, Azm, data*)

Needs a doc string

GSASIIimage.**GetDetectorXY** (*dsp, azm, data*)

Needs a doc string

GSASIIimage.**GetDsp** (*x, y, data*)

Give d-spacing value for detector x,y position; calibration info in data

GSASIIimage.**GetEllipse** (*dsp, data*)

uses Dandelin spheres to find ellipse or hyperbola parameters from detector geometry as given in image controls dictionary (data) and a d-spacing (dsp)

GSASIIimage.**GetEllipse2** (*tth, dxy, dist, cent, tilt, phi*)

uses Dandelin spheres to find ellipse or hyperbola parameters from detector geometry on output radii[0] (b-minor axis) set < 0. for hyperbola

`GSASIIimage.GetTth(x, y, data)`

Give 2-theta value for detector x,y position; calibration info in data

`GSASIIimage.GetTthAzm(x, y, data)`

Give 2-theta, azimuth values for detector x,y position; calibration info in data

`GSASIIimage.GetTthAzmD(x, y, data)`

Give 2-theta, azimuth & geometric correction values for detector x,y position; calibration info in data

`GSASIIimage.GetTthAzmDsp(x, y, data)`

Needs a doc string - checked OK for ellipses dont know about hyperbola

`GSASIIimage.ImageCalibrate(self, data)`

Needs a doc string

`GSASIIimage.ImageCompress(image, scale)`

Needs a doc string

`GSASIIimage.ImageIntegrate(image, data, masks, blkSize=128, dlq=None)`

Needs a doc string

`GSASIIimage.ImageLocalMax(image, w, Xpix, Ypix)`

Needs a doc string

`GSASIIimage.ImageRecalibrate(self, data, masks)`

Needs a doc string

`GSASIIimage.Make2ThetaAzimuthMap(data, masks, iLim, jLim)`

Needs a doc string

`GSASIIimage.calcFij(omg, phi, azm, th)`

Does something...

Uses parameters as defined by Bob He & Kingsley Smith, Adv. in X-Ray Anal. 41, 501 (1997)

Parameters

- **omg** – his omega = sample omega rotation; 0 when incident beam || sample surface, 90 when perp. to sample surface
- **phi** – his phi = sample phi rotation; usually = 0, axis rotates with omg.
- **azm** – his chi = azimuth around incident beam
- **th** – his theta = theta

`GSASIIimage.checkEllipse(Zsum, distSum, xSum, ySum, dist, x, y)`

Needs a doc string

`GSASIIimage.makeMat(Angle, Axis)`

Make rotation matrix from Angle and Axis

Parameters

- **Angle** (*float*) – in degrees
- **Axis** (*int*) – 0 for rotation about x, 1 for about y, etc.

`GSASIIimage.makeRing(dsp, ellipse, pix, reject, scalex, scaley, image)`

Needs a doc string

`GSASIIimage.peneCorr(tth, dep)`

Needs a doc string

`GSASIIimage.pointInPolygon(pXY, xy)`

Needs a doc string

GSASIIIMATH: COMPUTATION MODULE

Routines for least-squares minimization and other stuff

GSASIIImath.**AV2Q** (*A*, *V*)
convert angle (radians) & vector to quaternion $q=r+ai+bj+ck$

GSASIIImath.**AVdeg2Q** (*A*, *V*)
convert angle (degrees) & vector to quaternion $q=r+ai+bj+ck$

GSASIIImath.**AtomTLS2UIJ** (*atomData*, *atPtrs*, *Amat*, *rbObj*)
default doc string

Parameters *name* (*type*) – description

Returns type name: description

GSASIIImath.**AtomUIj2TLS** (*atomData*, *atPtrs*, *Amat*, *Bmat*, *rbObj*)
default doc string

Parameters *name* (*type*) – description

Returns type name: description

GSASIIImath.**ChargeFlip** (*data*, *reflDict*, *pgbar*)
default doc string

Parameters *name* (*type*) – description

Returns type name: description

GSASIIImath.**FillAtomLookUp** (*atomData*)
create a dictionary of atom indexes with atom IDs as keys

Parameters *atomData* (*list*) – Atom table to be used

Returns dict atomLookUp: dictionary of atom indexes with atom IDs as keys

GSASIIImath.**FindAtomIndexByIDs** (*atomData*, *IDs*, *Draw=True*)
finds the set of atom array indices for a list of atom IDs. Will search either the Atom table or the drawAtom table.

Parameters

- **atomData** (*list*) – Atom or drawAtom table containing coordinates, etc.
- **IDs** (*list*) – atom IDs to be found
- **Draw** (*bool*) – True if drawAtom table to be searched; False if Atom table is searched

Returns list indx: atom (or drawAtom) indices

`GSASIImath.FourierMap (data, reflDict)`

default doc string

Parameters *name (type)* – description

Returns type name: description

`GSASIImath.GetAngleSig (Oatoms, Atoms, Amat, SGData, covData={})`

default doc string

Parameters *name (type)* – description

Returns type name: description

`GSASIImath.GetAtomCoordsByID (pId, parmDict, AtLookup, indx)`

default doc string

Parameters *name (type)* – description

Returns type name: description

`GSASIImath.GetAtomItemsById (atomData, atomLookUp, IdList, itemLoc, numItems=1)`

gets atom parameters for atoms using atom IDs

Parameters

- **atomData** (*list*) – Atom table to be used
- **atomLookUp** (*dict*) – dictionary of atom indexes with atom IDs as keys
- **IdList** (*list*) – atom IDs to be found
- **itemLoc** (*int*) – pointer to desired 1st item in an atom table entry
- **numItems** (*int*) – number of items to be retrieved

Returns type name: description

`GSASIImath.GetAtomsById (atomData, atomLookUp, IdList)`

gets a list of atoms from Atom table that match a set of atom IDs

Parameters

- **atomData** (*list*) – Atom table to be used
- **atomLookUp** (*dict*) – dictionary of atom indexes with atom IDs as keys
- **IdList** (*list*) – atom IDs to be found

Returns list atoms: list of atoms found

`GSASIImath.GetDATSig (Oatoms, Atoms, Amat, SGData, covData={})`

default doc string

Parameters *name (type)* – description

Returns type name: description

`GSASIImath.GetDistSig (Oatoms, Atoms, Amat, SGData, covData={})`

default doc string

Parameters *name (type)* – description

Returns type name: description

`GSASIImath.GetSHCoeff (pId, parmDict, SHkeys)`

default doc string

Parameters *name (type)* – description

Returns type name: description

GSASIImath.**GetTorsionSig** (*Oatoms, Atoms, Amat, SGData, covData={}*)
default doc string

Parameters *name (type)* – description

Returns type name: description

GSASIImath.**GetXYZDist** (*xyz, XYZ, Amat*)

gets distance from position xyz to all XYZ, xyz & XYZ are np.array and are in crystal coordinates; Amat is crystal to Cart matrix

Parameters *name (type)* – description

Returns type name: description

GSASIImath.**HessianLSQ** (*func, x0, Hess, args=(), ftol=1.49012e-08, xtol=1.49012e-08, maxcyc=0, Print=False*)

Minimize the sum of squares of a function (*f*) evaluated on a series of values (*y*): $\sum_{y=0}^{N_{obs}} f(y, args)$

Nobs
x = arg min (sum(func(y)**2, axis=0))
 y=0

Parameters

- **func** (*function*) – callable method or function should take at least one (possibly length N vector) argument and returns M floating point numbers.
- **x0** (*np.ndarray*) – The starting estimate for the minimization of length N
- **Hess** (*function*) – callable method or function A required function or method to compute the weighted vector and Hessian for func. It must be a symmetric NxN array
- **args** (*tuple*) – Any extra arguments to func are placed in this tuple.
- **ftol** (*float*) – Relative error desired in the sum of squares.
- **xtol** (*float*) – Relative error desired in the approximate solution.
- **maxcyc** (*int*) – The maximum number of cycles of refinement to execute, if -1 refine until other limits are met (ftol, xtol)
- **Print** (*bool*) – True for printing results (residuals & times) by cycle

Returns

(x,cov_x,infodict) where

- **x** : ndarray The solution (or the result of the last iteration for an unsuccessful call).
- **cov_x** : ndarray Uses the fjac and ipvt optional outputs to construct an estimate of the jacobian around the solution. None if a singular matrix encountered (indicates very flat curvature in some direction). This matrix must be multiplied by the residual standard deviation to get the covariance of the parameter estimates – see curve_fit.
- **infodict** : dict a dictionary of optional outputs with the keys:
 - ‘fvec’ : the function evaluated at the output
 - ‘num cyc’:
 - ‘nfev’:
 - ‘lamMax’:

– ‘psing’:

`GSASIImath.OmitMap (data, reflDict, pgsbar=None)`
default doc string

Parameters *name (type)* – description

Returns type name: description

`GSASIImath.PeaksEquiv (data, Ind)`

Find the equivalent map peaks for those selected. Works on the contents of data[‘Map Peaks’].

Parameters

- **data** – the phase data structure
- **Ind** (*list*) – list of selected peak indices

Returns augmented list of peaks including those related by symmetry to the ones in Ind

`GSASIImath.PeaksUnique (data, Ind)`

Finds the symmetry unique set of peaks from those selected. Works on the contents of data[‘Map Peaks’].

Parameters

- **data** – the phase data structure
- **Ind** (*list*) – list of selected peak indices

Returns the list of symmetry unique peaks from among those given in Ind

`GSASIImath.Q2AV (Q)`

convert quaternion to angle (radians 0-2pi) & normalized vector $q=r+ai+bj+ck$

`GSASIImath.Q2AVdeg (Q)`

convert quaternion to angle (degrees 0-360) & normalized vector $q=r+ai+bj+ck$

`GSASIImath.Q2Mat (Q)`

make rotation matrix from quaternion $q=r+ai+bj+ck$

`GSASIImath.RotateRBXYZ (Bmat, Cart, oriQ)`

rotate & transform cartesian coordinates to crystallographic ones no translation applied. To be used for numerical derivatives

Parameters *name (type)* – description

Returns type name: description

`GSASIImath.SearchMap (data)`

Does a search of a density map for peaks meeting the criterion of peak height is greater than mapData[‘cutOff’]/100 of mapData[‘rhoMax’] where mapData is data[‘General’][‘mapData’]; the map is also in mapData.

Parameters **data** – the phase data structure

Returns

(peaks,mags,dzeros) where

- **peaks** : ndarray x,y,z positions of the peaks found in the map
- **mags** : ndarray the magnitudes of the peaks
- **dzeros** : ndarray the distance of the peaks from the unit cell origin

`GSASIImath.SetMolCent (model, RBDData)`

default doc string

Parameters *name (type)* – description

Returns type name: description

GSASIImath.**TLS2Uij** (*xyz, g, Amat, rbObj*)
default doc string

Parameters *name (type)* – description

Returns type name: description

GSASIImath.**UpdateMCSAxyz** (*Bmat, MCSA*)
default doc string

Parameters *name (type)* – description

Returns type name: description

GSASIImath.**UpdateRBUIJ** (*Bmat, Cart, RBObj*)
default doc string

Parameters *name (type)* – description

Returns type name: description

GSASIImath.**UpdateRXYZ** (*Bmat, RBObj, RBDData, RBType*)
default doc string

Parameters *name (type)* – description

Returns type name: description

GSASIImath.**ValEsd** (*value, esd=0, nTZ=False*)

Format a floating point number with a given level of precision or with in crystallographic format with a “esd”, as value(esd). If esd is negative the number is formatted with the level of significant figures appropriate if abs(esd) were the esd, but the esd is not included. if the esd is zero, approximately 6 significant figures are printed. nTZ=True causes “extra” zeros to be removed after the decimal place. for example:

- “1.235(3)” for value=1.2346 & esd=0.003
- “1.235(3)e4” for value=12346. & esd=30
- “1.235(3)e6” for value=0.12346e7 & esd=3000
- “1.235” for value=1.2346 & esd=-0.003
- “1.240” for value=1.2395 & esd=-0.003
- “1.24” for value=1.2395 & esd=-0.003 with nTZ=True
- “1.23460” for value=1.2346 & esd=0.0

Parameters

- **value** (*float*) – number to be formatted
- **esd** (*float*) – uncertainty or if esd < 0, specifies level of precision to be shown e.g. esd=-0.01 gives 2 places beyond decimal
- **nTZ** (*bool*) – True to remove trailing zeros (default is False)

Returns value(esd) or value as a string

GSASIImath.**adjHKLmax** (*SGData, Hmax*)
default doc string

Parameters *name (type)* – description

Returns type name: description

`GSASIImath.anneal` (*func*, *x0*, *args=()*, *schedule='fast'*, *full_output=0*, *T0=None*, *Tf=1e-12*, *maxeval=None*, *maxaccept=None*, *maxiter=400*, *boltzmann=1.0*, *learn_rate=0.5*, *feqs=1e-06*, *quench=1.0*, *m=1.0*, *n=1.0*, *lower=-100*, *upper=100*, *dwell=50*, *slope=0.9*, *ranStart=False*, *ranRange=0.1*, *autoRan=False*, *dlg=None*)

Minimize a function using simulated annealing.

Schedule is a schedule class implementing the annealing schedule. Available ones are 'fast', 'cauchy', 'boltzmann'

Parameters

- **func** (*callable*) – $f(x, *args)$ Function to be optimized.
- **x0** (*ndarray*) – Initial guess.
- **args** (*tuple*) – Extra parameters to *func*.
- **schedule** (*base_schedule*) – Annealing schedule to use (a class).
- **full_output** (*bool*) – Whether to return optional outputs.
- **T0** (*float*) – Initial Temperature (estimated as 1.2 times the largest cost-function deviation over random points in the range).
- **Tf** (*float*) – Final goal temperature.
- **maxeval** (*int*) – Maximum function evaluations.
- **maxaccept** (*int*) – Maximum changes to accept.
- **maxiter** (*int*) – Maximum cooling iterations.
- **learn_rate** (*float*) – Scale constant for adjusting guesses.
- **boltzmann** (*float*) – Boltzmann constant in acceptance test (increase for less stringent test at each temperature).
- **feqs** (*float*) – Stopping relative error tolerance for the function value in last four coolings.
- **quench,m,n** (*float*) – Parameters to alter fast_sa schedule.
- **lower,upper** (*float/ndarray*) – Lower and upper bounds on x .
- **dwell** (*int*) – The number of times to search the space at each temperature.
- **slope** (*float*) – Parameter for log schedule
- **ranStart=False** (*bool*) – True for set 10% of ranges about x

Returns

(*xmin*, *Jmin*, *T*, *feval*, *iters*, *accept*, *retval*) where

- *xmin* (*ndarray*): Point giving smallest value found.
- *Jmin* (*float*): Minimum value of function found.
- *T* (*float*): Final temperature.
- *feval* (*int*): Number of function evaluations.
- *iters* (*int*): Number of cooling iterations.
- *accept* (*int*): Number of tests accepted.
- *retval* (*int*): Flag indicating stopping condition:
 - 0: Points no longer changing

- 1: Cooled to final temperature
- 2: Maximum function evaluations
- 3: Maximum cooling iterations reached
- 4: Maximum accepted query locations reached
- 5: Final point not the minimum amongst encountered points

Notes: Simulated annealing is a random algorithm which uses no derivative information from the function being optimized. In practice it has been more useful in discrete optimization than continuous optimization, as there are usually better algorithms for continuous optimization problems.

Some experimentation by trying the difference temperature schedules and altering their parameters is likely required to obtain good performance.

The randomness in the algorithm comes from random sampling in numpy. To obtain the same results you can call `numpy.random.seed` with the same seed immediately before calling `scipy.optimize.anneal`.

We give a brief description of how the three temperature schedules generate new points and vary their temperature. Temperatures are only updated with iterations in the outer loop. The inner loop is over `xrange(dwell)`, and new points are generated for every iteration in the inner loop. (Though whether the proposed new points are accepted is probabilistic.)

For readability, let `d` denote the dimension of the inputs to `func`. Also, let `x_old` denote the previous state, and `k` denote the iteration number of the outer loop. All other variables not defined below are input variables to `scipy.optimize.anneal` itself.

In the ‘fast’ schedule the updates are

```
u ~ Uniform(0, 1, size=d)
y = sgn(u - 0.5) * T * ((1 + 1/T)**abs(2u-1) - 1.0)
xc = y * (upper - lower)
x_new = x_old + xc

c = n * exp(-n * quench)
T_new = T0 * exp(-c * k**quench)
```

In the ‘cauchy’ schedule the updates are

```
u ~ Uniform(-pi/2, pi/2, size=d)
xc = learn_rate * T * tan(u)
x_new = x_old + xc

T_new = T0 / (1+k)
```

In the ‘boltzmann’ schedule the updates are

```
std = minimum( sqrt(T) * ones(d), (upper-lower) / (3*learn_rate) )
y ~ Normal(0, std, size=d)
x_new = x_old + learn_rate * y

T_new = T0 / log(1+k)
```

`GSASIImath.calcRamaEnergy(phi, psi, Coeff=[])`

Computes pseudo potential energy from a pair of torsion angles and a numerical description of the potential energy surface. Used to create penalty function in LS refinement: $Eval(\phi, \psi) = C[0] * \exp(-V/1000)$

where $V = -C[3] * (\phi - C[1])^2 - C[4] * (\psi - C[2])^2 - 2 * (\phi - C[1]) * (\psi - C[2])$

Parameters

- **phi** (*float*) – first torsion angle (ϕ)
- **psi** (*float*) – second torsion angle (ψ)
- **Coeff** (*list*) – pseudo potential coefficients

Returns list (sum, Eval): pseudo-potential difference from minimum & value; sum is used for penalty function.

`GSASIImath.calcTorsionEnergy (TOR, Coeff=[])`
default doc string

Parameters *name* (*type*) – description

Returns type name: description

`GSASIImath.findOffset (SGData, A, Fhkl)`
default doc string

Parameters *name* (*type*) – description

Returns type name: description

`GSASIImath.getAngSig (VA, VB, Amat, SGData, covData={})`
default doc string

Parameters *name* (*type*) – description

Returns type name: description

`GSASIImath.getAtomXYZ (atoms, cx)`
default doc string

Parameters *name* (*type*) – description

Returns type name: description

`GSASIImath.getCWgam (ins, pos)`
default doc string

Parameters *name* (*type*) – description

Returns type name: description

`GSASIImath.getCWgamDeriv (pos)`
default doc string

Parameters *name* (*type*) – description

Returns type name: description

`GSASIImath.getCWsig (ins, pos)`
default doc string

Parameters *name* (*type*) – description

Returns type name: description

`GSASIImath.getCWsigDeriv (pos)`
default doc string

Parameters *name* (*type*) – description

Returns type name: description

`GSASIImath.getDensity (generalData)`
default doc string

Parameters *name (type)* – description

Returns type name: description

`GSASIImath.getDistDerv (Oxyz, Txyz, Amat, Tunit, Top, SGData)`
default doc string

Parameters *name (type)* – description

Returns type name: description

`GSASIImath.getMass (generalData)`
default doc string

Parameters *name (type)* – description

Returns type name: description

`GSASIImath.getRamaDerv (XYZ, Amat, Coeff)`

Computes numerical derivatives of torsion angle pair pseudo potential with respect of crystallographic atom coordinates of the 5 atom sequence

Parameters

- **XYZ** (*narray*) – crystallographic coordinates of 5 atoms
- **Amat** (*narray*) – crystal to cartesian transformation matrix
- **Coeff** (*list*) – pseudo potential coefficients

Returns list (deriv) derivatives of pseudopotential with respect to 5 atom crystallographic xyz coordinates.

`GSASIImath.getRestAngle (XYZ, Amat)`
default doc string

Parameters *name (type)* – description

Returns type name: description

`GSASIImath.getRestChiral (XYZ, Amat)`
default doc string

Parameters *name (type)* – description

Returns type name: description

`GSASIImath.getRestDerv (Func, XYZ, Amat, ops, SGData)`
default doc string

Parameters *name (type)* – description

Returns type name: description

`GSASIImath.getRestDist (XYZ, Amat)`
default doc string

Parameters *name (type)* – description

Returns type name: description

`GSASIImath.getRestPlane (XYZ, Amat)`
default doc string

Parameters *name (type)* – description

Returns type name: description

`GSASIImath.getRestPolefig (ODFln, SamSym, Grid)`

default doc string

Parameters *name (type)* – description

Returns type name: description

`GSASIImath.getRestPolefigDerv (HKL, Grid, SHCoeff)`

default doc string

Parameters *name (type)* – description

Returns type name: description

`GSASIImath.getRestRama (XYZ, Amat)`

Computes a pair of torsion angles in a 5 atom string

Parameters

- **XYZ** (*narray*) – crystallographic coordinates of 5 atoms
- **Amat** (*narray*) – crystal to cartesian transformation matrix

Returns list (phi,psi) two torsion angles in degrees

`GSASIImath.getRestTorsion (XYZ, Amat)`

default doc string

Parameters *name (type)* – description

Returns type name: description

`GSASIImath.getSyXYZ (XYZ, ops, SGData)`

default doc string

Parameters *name (type)* – description

Returns type name: description

`GSASIImath.getTOFalpha (ins, dsp)`

default doc string

Parameters *name (type)* – description

Returns type name: description

`GSASIImath.getTOFalphaDeriv (dsp)`

default doc string

Parameters *name (type)* – description

Returns type name: description

`GSASIImath.getTOFbeta (ins, dsp)`

default doc string

Parameters *name (type)* – description

Returns type name: description

`GSASIImath.getTOFbetaDeriv (dsp)`

default doc string

Parameters *name (type)* – description

Returns type name: description

`GSASIImath.getTOFgamma (ins, dsp)`
default doc string

Parameters *name (type)* – description

Returns type name: description

`GSASIImath.getTOFgammaDeriv (dsp)`
default doc string

Parameters *name (type)* – description

Returns type name: description

`GSASIImath.getTOFsig (ins, dsp)`
default doc string

Parameters *name (type)* – description

Returns type name: description

`GSASIImath.getTOFsigDeriv (dsp)`
default doc string

Parameters *name (type)* – description

Returns type name: description

`GSASIImath.getTorsionDeriv (XYZ, Amat, Coeff)`
default doc string

Parameters *name (type)* – description

Returns type name: description

`GSASIImath.getVCov (varyNames, varyList, covMatrix)`
obtain variance-covariance terms for a set of variables. NB: the varyList and covMatrix were saved by the last least squares refinement so they must match.

Parameters

- **varyNames** (*list*) – variable names to find v-cov matrix for
- **varyList** (*list*) – full list of all variables in v-cov matrix
- **covMatrix** (*nparray*) – full variance-covariance matrix from the last least squares refinement

Returns nparray vcov: variance-covariance matrix for the variables given in varyNames

`GSASIImath.getWave (Parms)`
default doc string

Parameters *name (type)* – description

Returns type name: description

`GSASIImath.invQ (Q)`
get inverse of quaternion $q=r+ai+bj+ck$; $q^*=r-ai-bj-ck$

`GSASIImath.makeQuat (A, B, C)`
Make quaternion from rotation of A vector to B vector about C axis

Parameters **A,B,C** (*np.array*) – Cartesian 3-vectors

Returns quaternion & rotation angle in radians $q=r+ai+bj+ck$

`GSASIImath.mcsaSearch (data, RBdata, reflType, reflData, covData, pgbars)`
 default doc string

Parameters *name (type)* – description

Returns type name: description

`GSASIImath.normQ (QA)`
 get length of quaternion & normalize it $q=r+ai+bj+ck$

`GSASIImath.printRho (SGLaue, rho, rhoMax)`
 default doc string

Parameters *name (type)* – description

Returns type name: description

`GSASIImath.prodQQ (QA, QB)`
 Grassman quaternion product QA, QB quaternions; $q=r+ai+bj+ck$

`GSASIImath.prodQVQ (Q, V)`
 compute the quaternion vector rotation $qvq^{-1} = v'$ $q=r+ai+bj+ck$

`GSASIImath.randomAVdeg (r0, r1, r2, r3)`
 create random angle (deg), vector from 4 random number in range (-1,1)

`GSASIImath.randomQ (r0, r1, r2, r3)`
 create random quaternion from 4 random numbers in range (-1,1)

`GSASIImath.setPeakparms (Parms, Parms2, pos, mag, ifQ=False, useFit=False)`
 default doc string

Parameters *name (type)* – description

Returns type name: description

`GSASIImath.sortArray (data, pos, reverse=False)`
 data is a list of items sort by pos in list; reverse if True

GSASIIINDEX: CELL INDEXING MODULE

Cell indexing program: variation on that of A. Coelho includes cell refinement from peak positions (not zero as yet)

This needs a bit of refactoring to remove the little bit of GUI code referencing wx

GSASIIindex.**A2values** (*ibrav, A*)
needs a doc string

GSASIIindex.**DoIndexPeaks** (*peaks, wave, controls, bravais*)
needs a doc string

GSASIIindex.**FitHKL** (*ibrav, peaks, A, Pwr*)
needs a doc string

GSASIIindex.**FitHKLZ** (*wave, ibrav, peaks, A, Z, Zref, Pwr*)
needs a doc string

GSASIIindex.**IndexPeaks** (*peaks, HKL*)
needs a doc string

GSASIIindex.**TestData** ()
needs a doc string

GSASIIindex.**Values2A** (*ibrav, values*)
needs a doc string

GSASIIindex.**calc_M20** (*peaks, HKL*)
needs a doc string

GSASIIindex.**findBestCell** (*dlg, ncMax, A, Ntries, ibrav, peaks, VI*)
needs a doc string

GSASIIindex.**getDmax** (*peaks*)
needs a doc string

GSASIIindex.**getDmin** (*peaks*)
needs a doc string

GSASIIindex.**halfCell** (*ibrav, A, peaks*)
needs a doc string

GSASIIindex.**monoCellReduce** (*ibrav, A*)
needs a doc string

GSASIIindex.**oddPeak** (*indx, peaks*)
needs a doc string

`GSASIIindex.ran2axis` (k, N)
needs a doc string

`GSASIIindex.ranAbyR` ($Bravais, A, k, N, ranFunc$)
needs a doc string

`GSASIIindex.ranAbyV` ($Bravais, dmin, dmax, V$)
needs a doc string

`GSASIIindex.ranaxis` ($dmin, dmax$)
needs a doc string

`GSASIIindex.rancell` ($Bravais, dmin, dmax$)
needs a doc string

`GSASIIindex.refinePeaks` ($peaks, ibrav, A$)
needs a doc string

`GSASIIindex.refinePeaksZ` ($peaks, wave, ibrav, A, Zero, ZeroRef$)
needs a doc string

`GSASIIindex.rotOrthoA` (A)
needs a doc string

`GSASIIindex.scaleAbyV` (A, V)
needs a doc string

`GSASIIindex.sortM20` ($cells$)
needs a doc string

`GSASIIindex.swapMonoA` (A)
needs a doc string

GSASII PLOT: PLOTTING ROUTINES

```
class GSASIIplot.G2Plot3D (parent, id=-1, dpi=None, **kwargs)
    needs a doc string

class GSASIIplot.G2PlotMpl (parent, id=-1, dpi=None, **kwargs)
    needs a doc string

class GSASIIplot.G2PlotNoteBook (parent, id=-1)
    create a tabbed window for plotting

    Delete (name)
        delete a tabbed page

    OnNotebookKey (event)
        Called when a keystroke event gets picked up by the notebook window rather the child. This is not
        expected, but somehow it does sometimes on the Mac and perhaps Linux.

        Assume that the page associated with the currently displayed tab has a child, .canvas; give that child the
        focus and pass it the event.

    OnPageChanged (event)
        respond to someone pressing a tab on the plot window

    Rename (oldName, newName)
        rename a tab

    add3D (name='')
        Add a tabbed page with a 3D plot

    addMpl (name='')
        Add a tabbed page with a matplotlib plot

    addOgl (name='')
        Add a tabbed page with an openGL plot

    clear ()
        clear all pages from plot window

class GSASIIplot.G2PlotOgl (parent, id=-1, dpi=None, **kwargs)
    needs a doc string

class GSASIIplot.GSASIItoolbar (plotCanvas)
    needs a doc string

    OnHelp (event)
        needs a doc string
```

OnKey (*event*)

needs a doc string

`GSASIIplot.OnStartMask` (*G2frame*)

Initiate the start of a Frame or Polygon map

Parameters

- **G2frame** (*wx.Frame*) – The main GSAS-II tree “window”
- **eventkey** (*str*) – a single letter (‘f’ or ‘p’) that determines what type of mask is created.

`GSASIIplot.PlotCovariance` (*G2frame, Data*)

needs a doc string

`GSASIIplot.PlotDeltSig` (*G2frame, kind*)

needs a doc string

`GSASIIplot.PlotExposedImage` (*G2frame, newPlot=False, event=None*)

General access module for 2D image plotting

`GSASIIplot.PlotISFG` (*G2frame, newPlot=False, type=''*)

Plotting package for PDF analysis; displays I(q), S(q), F(q) and G(r) as single or multiple plots with waterfall and contour plots as options

`GSASIIplot.PlotImage` (*G2frame, newPlot=False, event=None, newImage=True*)

Plot of 2D detector images as contoured plot. Also plot calibration ellipses, masks, etc.

`GSASIIplot.PlotIntegration` (*G2frame, newPlot=False, event=None*)

Plot of 2D image after image integration with 2-theta and azimuth as coordinates

`GSASIIplot.PlotPatterns` (*G2frame, newPlot=False*)

Powder pattern plotting package - displays single or multiple powder patterns as intensity vs 2-theta, q or TOF. Can display multiple patterns as “waterfall plots” or contour plots. Log I plotting available.

`GSASIIplot.PlotPeakWidths` (*G2frame*)

Plotting of instrument broadening terms as function of 2-theta. Seen when “Instrument Parameters” chosen from powder pattern data tree

`GSASIIplot.PlotPowderLines` (*G2frame*)

plotting of powder lines (i.e. no powder pattern) as sticks

`GSASIIplot.PlotRama` (*G2frame, phaseName, Rama, RamaName, Names=[], PhiPsi=[], Coeff=[]*)

needs a doc string

`GSASIIplot.PlotRigidBody` (*G2frame, rbType, AtInfo, rbData, defaults*)

RB plotting package. Can show rigid body structures as balls & sticks

`GSASIIplot.PlotSeq` (*G2frame, SeqData, SeqSig, SeqNames, sampleParm*)

needs a doc string

`GSASIIplot.PlotSizeStrainPO` (*G2frame, data, Start=False*)

Plot 3D mustrain/size/preferred orientation figure. In this instance data is for a phase

`GSASIIplot.PlotSngl` (*self, newPlot=False*)

Single crystal structure factor plotting package - displays zone of reflections as rings proportional to F, F**2, etc. as requested

`GSASIIplot.PlotStructure` (*G2frame, data*)

Crystal structure plotting package. Can show structures as balls, sticks, lines, thermal motion ellipsoids and polyhedra

`GSASIIplot.PlotTRImage` (*G2frame, tax, tay, taz, newPlot=False*)

a test plot routine - not normally used

GSASIIplot.**PlotTexture** (*G2frame, data, Start=False*)

Pole figure, inverse pole figure, 3D pole distribution and 3D inverse pole distribution plotting. dict generalData contains all phase info needed which is in data

GSASIIplot.**PlotTorsion** (*G2frame, phaseName, Torsion, TorName, Names=[], Angles=[], Coeff=[]*)

needs a doc string

GSASIIplot.**PlotXY** (*G2frame, XY, newPlot=False, type=''*)

simple plot of xy data, used for diagnostic purposes

GSASII POWDER CALCULATION MODULE

GSASIIpwr.**Absorb** (*Geometry, MuR, Tth, Phi=0, Psi=0*)

Calculate sample absorption :param str Geometry: one of 'Cylinder','Bragg-Brentano','Tilting Flat Plate in transmission','Fixed flat plate' :param float MuR: absorption coeff * sample thickness/2 or radius :param Tth: 2-theta scattering angle - can be numpy array :param float Phi: flat plate tilt angle - future :param float Psi: flat plate tilt axis - future

GSASIIpwr.**AbsorbDerv** (*Geometry, MuR, Tth, Phi=0, Psi=0*)

needs a doc string

GSASIIpwr.**CalcPDF** (*data, inst, xydata*)

needs a doc string

GSASIIpwr.**Dict2Values** (*parmdict, varylist*)

Use before call to leastsq to setup list of values for the parameters in parmdict, as selected by key in varylist

GSASIIpwr.**DoPeakFit** (*FitPgm, Peaks, Background, Limits, Inst, Inst2, data, oneCycle=False, controls=None, dIlg=None*)

needs a doc string

GSASIIpwr.**GetAsfMean** (*ElList, Sthl2*)

Calculate various scattering factor terms for PDF calcs

Parameters

- **ElList** (*dict*) – element dictionary contains scattering factor coefficients, etc.
- **Sthl2** (*np.array*) – numpy array of sin theta/lambda squared values

Returns mean(f^2), mean(f^2), mean(compton)

GSASIIpwr.**GetNumDensity** (*ElList, Vol*)

needs a doc string

GSASIIpwr.**LorchWeight** (*Q*)

needs a doc string

GSASIIpwr.**Oblique** (*ObCoeff, Tth*)

currently assumes detector is normal to beam

GSASIIpwr.**Polarization** (*Pola, Tth, Azm=0.0*)

Calculate angle dependent x-ray polarization correction (not scaled correctly!)

Parameters

- **Pola** – polarization coefficient e.g 1.0 fully polarized, 0.5 unpolarized

- **Azm** – azimuthal angle e.g. 0.0 in plane of polarization
- **Tth** – 2-theta scattering angle - can be numpy array which (if either) of these is “right”?

Returns (pola, dpdPola) * pola = $((1-Pola)*np\cos d(Azm)**2 + Pola*np\sin d(Azm)**2)*np\cos d(Tth)**2 + (1-Pola)*np\sin d(Azm)**2 + Pola*np\cos d(Azm)**2$ * dpdPola: derivative needed for least squares

GSASIIpwd.**Ruland** (*RulCoff, wave, Q, Compton*)
needs a doc string

GSASIIpwd.**SetBackgroundParms** (*Background*)
needs a doc string

GSASIIpwd.**TestData** ()
needs a doc string

GSASIIpwd.**Transmission** (*Geometry, Abs, Diam*)
Calculate sample transmission

Parameters

- **Geometry** (*str*) – one of ‘Cylinder’, ‘Bragg-Brentano’, ‘Tilting flat plate in transmission’, ‘Fixed flat plate’
- **Abs** (*float*) – absorption coeff in cm-1
- **Diam** (*float*) – sample thickness/diameter in mm

GSASIIpwd.**Values2Dict** (*parmdict, varylist, values*)
Use after call to leastsq to update the parameter dictionary with values corresponding to keys in varylist

GSASIIpwd.**calcIncident** (*Iparm, xdata*)
needs a doc string

class GSASIIpwd.**cauchy_gen** (*momtype=1, a=None, b=None, xa=-10.0, xb=10.0, xtol=1e-14, badvalue=None, name=None, longname=None, shapes=None, extradoc=None*)
needs a doc string

GSASIIpwd.**ellipseSize** (*H, Sij, GB*)
needs a doc string

GSASIIpwd.**ellipseSizeDerv** (*H, Sij, GB*)
needs a doc string

GSASIIpwd.**factorize** (*num*)
Provide prime number factors for integer num Returns dictionary of prime factors (keys) & power for each (data)

class GSASIIpwd.**fcjde_gen** (*momtype=1, a=None, b=None, xa=-10.0, xb=10.0, xtol=1e-14, badvalue=None, name=None, longname=None, shapes=None, extradoc=None*)

Finger-Cox-Jephcoat D(2phi,2th) function for S/L = H/L Ref: J. Appl. Cryst. (1994) 27, 892-900.

Parameters

- **x** – array -1 to 1
- **t** – 2-theta position of peak
- **s** – sum(S/L,H/L); S: sample height, H: detector opening, L: sample to detector opening distance
- **dx** – 2-theta step size in deg

Returns

for `fcj.pdf`

- $T = x \cdot dx + t$
- $s = S/L + H/L$
- if $x < 0$:

$$fcj.pdf = [1/\sqrt{(\cos(T)^2/\cos(t)^2)-1} - 1/s] / |\cos(T)|$$

- if $x \geq 0$: `fcj.pdf = 0`

`GSASIIpwd.getBackground` (*pfx, parmDict, bakType, xdata*)

needs a doc string

`GSASIIpwd.getBackgroundDerv` (*pfx, parmDict, bakType, xdata*)

needs a doc string

`GSASIIpwd.getEpsVoigt` (*pos, alp, bet, sig, gam, xdata*)

needs a doc string

`GSASIIpwd.getFCJVoigt` (*pos, intens, sig, gam, shl, xdata*)

needs a doc string

`GSASIIpwd.getFCJVoigt3` (*pos, sig, gam, shl, xdata*)

needs a doc string

`GSASIIpwd.getFWHM` (*pos, Inst*)

needs a doc string

`GSASIIpwd.getHKLpeak` (*dmin, SGData, A*)

needs a doc string

`GSASIIpwd.getPeakProfile` (*dataType, parmDict, xdata, varyList, bakType*)

needs a doc string

`GSASIIpwd.getPeakProfileDerv` (*dataType, parmDict, xdata, varyList, bakType*)

needs a doc string

`GSASIIpwd.getWidthsCW` (*pos, sig, gam, shl*)

needs a doc string

`GSASIIpwd.getWidthsTOF` (*pos, alp, bet, sig, gam*)

needs a doc string

`GSASIIpwd.getdEpsVoigt` (*pos, alp, bet, sig, gam, xdata*)

needs a doc string

`GSASIIpwd.getdFCJVoigt3` (*pos, sig, gam, shl, xdata*)

needs a doc string

`GSASIIpwd.getgamFW` (*g, s*)

needs a doc string

`GSASIIpwd.makeFFTsizeList` (*nmin=1, nmax=1023, thresh=15*)

Provide list of optimal data sizes for FFT calculations

Parameters

- **nmin** (*int*) – minimum data size ≥ 1
- **nmax** (*int*) – maximum data size $> nmin$

- **thresh** (*int*) – maximum prime factor allowed

Returns list of data sizes where the maximum prime factor is < thresh

class GSASIIpwd.**norm_gen** (*momtype=1, a=None, b=None, xa=-10.0, xb=10.0, xtol=1e-14, bad-value=None, name=None, longname=None, shapes=None, extradoc=None*)
needs a doc string

GSAS-II SCRIPTS

13.1 *testDeriv: Check derivative computation*

Use this to check derivatives used in structure least squares refinement against numerical values computed in this script.

To use set `DEBUG=True` in `GSASIIstrMain.py` (line 22, as of version 1110); run the least squares - one cycle is sufficient. Do the “Save Results”; this will write the file `testDeriv.dat` in the local directory.

Then run this program to see plots of derivatives for all parameters refined in the last least squares. Shown will be numerical derivatives generated over all observations (including penalty terms) and the corresponding analytical ones produced in the least squares. They should match.

```
testDeriv.main()
```

Starts main application to compute and plot derivatives

13.2 *GSASIItestplot: Plotting for testDeriv*

Plotting module used for script `testDeriv`.

```
class GSASIItestplot.Plot (parent, id=-1, dpi=None, **kwargs)
    Creates a plotting window

class GSASIItestplot.PlotNotebook (id=-1)
    creates a Wx application and a plotting notebook
```

13.3 *scanCCD: reduce data from scanning CCD*

Quickly prototyped routine for reduction of data from detector described in B.H. Toby, T.J. Madden, M.R. Suchomel, J.D. Baldwin, and R.B. Von Dreele, “A Scanning CCD Detector for Powder Diffraction Measurements”. *Journal of Applied Crystallography*. 46(4): p. 1058-63 (2013).

```
scanCCD.main()
```

starts main application to merge data from scanning CCD

13.4 *makeMacApp: Create Mac Applet*

This script creates an AppleScript app to launch GSAS-II. The app is created in the directory where the GSAS-II script is located. A softlink to Python is created, but is named GSAS-II, so that GSAS-II shows up as the name of the app rather than Python in the menu bar, etc. Note that this requires finding an app version of Python (expected name `.../Resources/Python.app/Contents/MacOS/Python` in directory tree of the calling python interpreter).

Run this script with one optional argument, the path to the GSASII.py. The script path may be specified relative to the current path or given an absolute path, but will be accessed via an absolute path. If no arguments are supplied, the GSASII.py script is assumed to be in the same directory as this file.

13.5 *unit_tests: Self-test Module*

A script that can be run to test a series of self-tests in GSAS-II. At present, only modules `GSASIIspc` and `GSASIIlattice` have self-tests.

`unit_tests.test_GSASIIlattice()`

Test registered self-tests in `GSASIIlattice`. Takes no input and returns nothing. Throws an Exception if a test fails.

`unit_tests.test_GSASIIspc()`

Test registered self-tests in `GSASIIspc`. Takes no input and returns nothing. Throws an Exception if a test fails.

GSAS-II EXPORT MODULES

Exports are implemented by deriving a class from `GSASIIIO.ExportBaseClass`. Initialization of `self.exporttype` determines the type of export that will be performed ('project', 'phase', 'single', 'powder', 'image', 'map' or (someday) 'pdf') and of `self.multiple` determines if only a single phase, data set, etc. can be exported at a time (when False) or more than one can be selected.

14.1 Module `G2export_examples`: Examples

Code to demonstrate how GSAS-II data export routines are created. The classes defined here, `ExportPhaseText`, `ExportSingleText`, `ExportPowderReflText`, and `ExportPowderText` each demonstrate a different type of export. Also see `G2export_map.ExportMapASCII` for an example of a map export.

class `G2export_examples.ExportPhaseText (G2frame)`

Used to create a text file for a phase

Parameters `G2frame` (`wx.Frame`) – reference to main GSAS-II frame

Exporter (`event=None`)

Export a phase as a text file

class `G2export_examples.ExportPowderReflText (G2frame)`

Used to create a text file of reflections from a powder data set

Parameters `G2frame` (`wx.Frame`) – reference to main GSAS-II frame

Exporter (`event=None`)

Export a set of powder reflections as a text file

class `G2export_examples.ExportPowderText (G2frame)`

Used to create a text file for a powder data set

Parameters `G2frame` (`wx.Frame`) – reference to main GSAS-II frame

Exporter (`event=None`)

Export a set of powder data as a text file

class `G2export_examples.ExportSingleText (G2frame)`

Used to create a text file with single crystal reflection data

Parameters `G2frame` (`wx.Frame`) – reference to main GSAS-II frame

Exporter (`event=None`)

Export a set of single crystal data as a text file

14.2 Module *G2export_csv*: Spreadsheet export

Code to create .csv (comma-separated variable) files for GSAS-II data export to a spreadsheet program, etc.

class `G2export_csv.ExportPhaseCSV (G2frame)`

Used to create a csv file for a phase

Parameters `G2frame` (*wx.Frame*) – reference to main GSAS-II frame

Exporter (*event=None*)

Export a phase as a csv file

class `G2export_csv.ExportPowderCSV (G2frame)`

Used to create a csv file for a powder data set

Parameters `G2frame` (*wx.Frame*) – reference to main GSAS-II frame

Exporter (*event=None*)

Export a set of powder data as a csv file

class `G2export_csv.ExportPowderRef1CSV (G2frame)`

Used to create a csv file of reflections from a powder data set

Parameters `G2frame` (*wx.Frame*) – reference to main GSAS-II frame

Exporter (*event=None*)

Export a set of powder reflections as a csv file

class `G2export_csv.ExportSingleCSV (G2frame)`

Used to create a csv file with single crystal reflection data

Parameters `G2frame` (*wx.Frame*) – reference to main GSAS-II frame

Exporter (*event=None*)

Export a set of single crystal data as a csv file

`G2export_csv.WriteList (obj, headerItems)`

Write a CSV header

Parameters

- **obj** (*object*) – Exporter object
- **headerItems** (*list*) – items to write as a header

14.3 Module *G2export_PDB*: Macromolecular export

Code to export a phase into the venerated/obsolete (pick one) ASCII PDB format. Also defines exporter `ExportPhaseCartXYZ` which writes atom positions in orthogonal coordinates for a phase.

class `G2export_PDB.ExportPhaseCartXYZ (G2frame)`

Used to create a Cartesian XYZ file for a phase

Parameters `G2frame` (*wx.Frame*) – reference to main GSAS-II frame

Exporter (*event=None*)

Export as a XYZ file

class `G2export_PDB.ExportPhasePDB (G2frame)`

Used to create a PDB file for a phase

Parameters `G2frame` (*wx.Frame*) – reference to main GSAS-II frame

Exporter (*event=None*)
Export as a PDB file

14.4 Module *G2export_image*: 2D Image data export

Demonstrates how an image is retrieved and written. Uses a SciPy routine to write a PNG format file.

class *G2export_image.ExportImagePNG* (*G2frame*)
Used to create a PNG file for a GSAS-II image

Parameters *G2frame* (*wx.Frame*) – reference to main GSAS-II frame

Exporter (*event=None*)
Export an image

14.5 Module *G2export_map*: Map export

Code to write Fourier/Charge-Flip atomic density maps out in formats that can be read by external programs. At present a GSAS format that is supported by FOX and DrawXTL (*ExportMapASCII*) and the CCP4 format that is used by COOT (*ExportMapCCP4*) are implemented.

class *G2export_map.ExportMapASCII* (*G2frame*)
Used to create a text file for a phase

Parameters *G2frame* (*wx.Frame*) – reference to main GSAS-II frame

Exporter (*event=None*)
Export a map as a text file

class *G2export_map.ExportMapCCP4* (*G2frame*)
Used to create a text file for a phase

Parameters *G2frame* (*wx.Frame*) – reference to main GSAS-II frame

Exporter (*event=None*)
Export a map as a text file

OpenFile (*fil=None*)
Open the output file as binary

Parameters *fil* (*str*) – The name of the file to open. If None (default) the name defaults to self.filename.

Returns the file object opened by the routine which is also saved as self.fp

14.6 Module *G2export_shelx*: Examples

Code to export coordinates in the SHELX .ins format (as best as I can makes sense of it).

class *G2export_shelx.ExportPhaseShelx* (*G2frame*)
Used to create a SHELX .ins file for a phase

Parameters *G2frame* (*wx.Frame*) – reference to main GSAS-II frame

Exporter (*event=None*)
Export as a SHELX .ins file

14.7 Module `G2export_CIF`: CIF Exports

This implements a complex exporter `ExportCIF` that can implement an entire project in a complete CIF intended for submission as a publication. In addition, there are two subclasses of `ExportCIF`: `ExportPhaseCIF` and `ExportDataCIF` that export a single phase or data set. Note that `self.mode` determines what is written:

- `self.mode="simple"` creates a simple CIF with only coordinates or data, while
- `self.mode="full"` creates a complete CIF of project.

`G2export_CIF.CIF2dict (cf)`

copy the contents of a CIF out from a `PyCifRW` block object into a dict

Returns `cifblk`, `loopstructure` where `cifblk` is a dict with CIF items and `loopstructure` is a list of lists that defines which items are in which loops.

class `G2export_CIF.CIFdefHelp (parent, msg, helpwin, helptxt)`

Create a help button that displays help information on the current data item

Parameters

- **parent** – the panel which will be the parent of the button
- **msg** (*str*) – the help text to be displayed
- **helpwin** (*wx.Dialog*) – Frame for CIF editing dialog
- **helptxt** (*wx.TextCtrl*) – `TextCtrl` where help text is placed

class `G2export_CIF.CIFtemplateSelect (frame, panel, tplate, G2dict, repaint, title, defaultname='')`

Create a set of buttons to show, select and edit a CIF template

Parameters

- **frame** – *wx.Frame* object of parent
- **panel** – *wx.Panel* object where widgets should be placed
- **tplate** (*str*) – one of 'publ', 'phase', or 'instrument' to determine the type of template
- **G2dict** (*dict*) – GSAS-II dict where CIF should be placed. The key "CIF_template" will be used to store either a list or a string. If a list, it will contain a dict and a list defining loops. If an str, it will contain a file name.
- **repaint** (*function*) – reference to a routine to be called to repaint the frame after a change has been made
- **title** (*str*) – A line of text to show at the top of the window
- **defaultname** (*str*) – specifies the default file name to be used for saving the CIF.

class `G2export_CIF.EditCIFpanel (parent, cifblk, loopstructure, cifdic={}, OKbuttons=[], **kw)`

Creates a scrolled panel for editing CIF template items

Parameters

- **parent** (*wx.Frame*) – parent frame where panel will be placed
- **cifblk** – dict or `PyCifRW` block containing values for each CIF item
- **loopstructure** (*list*) – a list of lists containing the contents of each loop, as an example:

```
[ ["_a", "_b"], ["_c"], ["_d_1", "_d_2", "_d_3"] ]
```

this describes a CIF with this type of structure:

```
loop_ _a _b <a1> <b1> <a2> ...
loop_ _c <c1> <c2>...
loop _d_1 _d_2 _d_3 ...
```

Note that the values for each looped CIF item, such as `_a`, are contained in a list, for example as `cifblk["_a"]`

- **cifdic** (*dict*) – optional CIF dictionary definitions
- **OKbuttons** (*list*) – A list of `wx.Button` objects that should be disabled when information in the CIF is invalid
- **(other)** – optional keyword parameters for `wx.ScrolledPanel`

CIFEntryWidget (*dct, item, dataname*)

Create an entry widget for a CIF item. Use a validated entry for numb values where int is required when limits are integers and floats otherwise. At present this does not allow entry of the special CIF values of `."` and `?"` for numerical values and highlights them as invalid. Use a selection widget when there are specific enumerated values for a string.

ControlOKButton (*setvalue*)

Enable or Disable the OK button(s) for the dialog. Note that this is passed into the `ValidatedTxtCtrl` for use by validators.

Parameters *setvalue* (*bool*) – if True, all entries in the dialog are checked for validity. The first invalid control triggers disabling of buttons. If False then the OK button(s) are disabled with no checking of the invalid flag for each control.

DoLayout ()

Update the Layout and scroll bars for the Panel. Clears `self.LayoutCalled` so that next change to panel can request a new update

OnAddRow (*event*)

add a row to a loop

OnLayoutNeeded (*event*)

Called when an update of the panel layout is needed. Calls `self.DoLayout` after the current operations are complete using `CallAfter`. This is called only once, according to flag `self.LayoutCalled`, which is cleared in `self.DoLayout`.

class `G2export_CIF.EditCIFtemplate` (*parent, cifblk, loopstructure, defaultname*)

Create a dialog for editing a CIF template. The edited information is placed in `cifblk`. If the CIF is saved as a file, the name of that file is saved as `self.newfile`.

Parameters

- **parent** (*wx.Frame*) – parent frame or None
- **cifblk** – dict or `PyCifRW` block containing values for each CIF item
- **loopstructure** (*list*) – a list of lists containing the contents of each loop, as an example:

```
[ ["_a", "_b"], ["_c"], ["_d_1", "_d_2", "_d_3"]]
```

this describes a CIF with this type of structure:

```
loop_ _a _b <a1> <b1> <a2> ...
loop_ _c <c1> <c2>...
loop _d_1 _d_2 _d_3 ...
```

Note that the values for each looped CIF item, such as `_a`, are contained in a list, for example as `cifblk["_a"]`

- **defaultname** (*str*) – specifies the default file name to be used for saving the CIF.

Post ()

Display the dialog

Returns True unless Cancel has been pressed.

class `G2export_CIF.ExportCIF (G2frame)`

Used to create a CIF of an entire project

Parameters **G2frame** (*wx.Frame*) – reference to main GSAS-II frame

Exporter (*event=None*)

Export a CIF. Export can be full or simple (as set by `self.mode`). “simple” skips data, distances & angles, etc. and can only include a single phase while “full” is intended for publication submission.

class `G2export_CIF.ExportDataCIF (G2frame)`

Used to create a simple CIF containing diffraction data only. Uses exact same code as `ExportCIF` except that `self.mode` is set to “simple” and `self.currentExportType` is set to “single” or “powder” in `self.InitExport`. Shows up in menus as Data-only CIF.

Parameters **G2frame** (*wx.Frame*) – reference to main GSAS-II frame

class `G2export_CIF.ExportPhaseCIF (G2frame)`

Used to create a simple CIF of at most one phase. Uses exact same code as `ExportCIF` except that `self.mode` is set to “simple” in `self.InitExport`. Shows up in menu as Quick CIF.

Parameters **G2frame** (*wx.Frame*) – reference to main GSAS-II frame

`G2export_CIF.LoadCIFdic ()`

Create a composite core+powder CIF lookup dict containing information about all items in the CIF dictionaries, loading pickled files if possible. The routine looks for files named `cif_core.cpickle` and `cif_pd.cpickle` in every directory in the path and if they are not found, files `cif_core.dic` and/or `cif_pd.dic` are read.

Returns the dict with the definitions

`G2export_CIF.PickleCIFdict (fil)`

Loads a CIF dictionary, cherry picks out the items needed by local code and sticks them into a python dict and writes that dict out as a cPickle file for later reuse. If the write fails a warning message is printed, but no exception occurs.

Parameters **fil** (*str*) – file name of CIF dictionary, will usually end in `.dic`

Returns the dict with the definitions

`G2export_CIF.dict2CIF (dblk, loopstructure, blockname='Template')`

Create a `PyCifRW` CIF object containing a single CIF block object from a dict and loop structure list.

Parameters

- **dblk** – a dict containing values for each CIF item
- **loopstructure** (*list*) – a list of lists containing the contents of each loop, as an example:

```
[ ["_a", "_b"], ["_c"], ["_d_1", "_d_2", "_d_3"] ]
```

this describes a CIF with this type of structure:

```
loop_ _a _b <a1> <b1> <a2> ...  
loop_ _c <c1> <c2>...  
loop _d_1 _d_2 _d_3 ...
```

Note that the values for each looped CIF item, such as `_a`, are contained in a list, for example as `cifblk['_a']`

- **blockname** (*str*) – an optional name for the CIF block. Defaults to ‘Template’

Returns the newly created PyCifRW CIF object

GSAS-II IMPORT MODULES

Imports are implemented by deriving a class from `GSASIIIO.ImportPhase`, `GSASIIIO.ImportStructFactor` or `GSASIIIO.ImportPowderData` (which are in turn derived from `GSASIIIO.ImportBaseclass`) to implement import of a phase, a single crystal or a powder dataset, respectively. Module file names (*G2phase_*, *G2pwd_* and *G2sfact_*) are used to determine which menu an import routine should be placed into. (N.B. this was an unnecessary choice; this could be done from the class used.)

15.1 Writing an Import Routine

When writing a import routine, one should create a new class derived from `GSASIIIO.ImportPhase`, `GSASIIIO.ImportStructFactor` or `GSASIIIO.ImportPowderData`. As described below, all these classes will implement an `__init__()` and a `Reader()` method, and most will supply a `ContentsValidator()` method, too. See the `ImportPhase`, `ImportStructFactor` or `ImportPowderData` class documentation for details on what values each type of `Reader()` should set.

15.1.1 `__init__()`

The class should supply a `__init__` method which calls the parent `__init__` method and specifies the following parameters:

- *extensionlist*: a list of extensions that may be used for this type of file.
- *strictExtension*: Should be True if only files with extensions in *extensionlist* are allowed; False if all file types should be offered in the file browser. Also if False, the import class will be used on all files when “guess from format” is tried, though readers with matching extensions will be tried first.
- *formatName*: a string to be used in the menu. Should be short.
- *longFormatName*: a longer string to be used to describe the format in help.

15.1.2 `Reader()`

The class must supply a `Reader` method that actually performs the reading. All readers must have at a minimum these arguments:

```
def Reader(self, filename, filepointer, ParentFrame, **unused):
```

where the arguments have the following uses:

- *filename*: a string with the name of the file being read

- *filepointer*: a file object (created by `open()`) that accesses the file and is points to the beginning of the file when `Reader` is called.
- *ParentFrame*: a reference to the main GSAS-II (tree) windows, for the unusual `Reader` routines that will create GUI windows to ask questions.

In addition, the following keyword parameters are defined that `Reader` routines may optionally use:

- *buffer*: a dict that can be used to retain information between repeated calls of the routine
- *blocknum*: counts the number of times that a reader is called
- *usedRanIdList*: a list of previously used random Id values that can be checked to determine that a value is unique.

As an example, the *buffer* dict is used for CIF reading to hold the parsed CIF file so that it can be reused without having to reread the file from scratch.

Reader return values

The `Reader` routine should return the value of `True` if the file has been read successfully. Optionally, use *self.warnings* to indicate any problems.

If the file cannot be read, the `Reader` routine should return `False` or raise an `GSASIIIO.ImportBaseclass.ImportException` exception. (Why either? Sometimes an exception is the easiest way to bail out of a called routine.) Place text in *self.errors* and/or use:

```
ImportException('Error message')
```

to give the user information on what went wrong during the reading.

self.warnings

Use *self.warnings* to indicate any information that should be displayed to the user if the file is read successfully, but perhaps not completely or additional settings will need to be made.

self.errors

Use *self.errors* to give the user information on where and why a read error occurs in the file. Note that text supplied with the `raise` statement will be appended to *self.errors*.

self.repeat

Set *self.repeat* to `True` (the default is `False`) if a `Reader` should be called again to read a second block from a file. Most commonly (only?) used for reading multiple powder histograms from a single file. Variable *self.repeatcount* is used to keep track of the block numbers.

support routines

Note that the base class (`GSASIIIO.ImportBaseclass`) supplies two routines, `BlockSelector()` and `MultipleBlockSelector()` that are useful for selecting amongst one or more datasets (and perhaps phases) for `Reader()` routines that may encounter more than one set of information in a file. Likewise, when an operation will take some time to complete, use `ShowBusy()` and `DoneBusy()` to show the user that something is happening.

15.1.3 ContentsValidator()

Defining a `ContentsValidator` method is optional, but is usually a good idea, particularly if the file extension is not a reliable identifier for the file type. The intent of this routine is to take a superficial look at the file to see if it has the expected characteristics of the expected file type. For example, are there numbers in the expected places?

This routine is passed a single argument:

- *filepointer*: a file object (created by `open()`) that accesses the file and is points to the beginning of the file when `ContentsValidator` is called.

Note that `GSASIIIO.ImportBaseclass.CIFValidator()` is a `ContentsValidator` for validating CIF files.

ContentsValidator return values

The `ContentsValidator` routine should return the value of `True` if the file appears to match the type expected for the class.

If the file cannot be read by this class, the routine should return `False`. Preferably one will also place text in *self.errors* to give the user information on what went wrong during the reading.

15.2 Currently Defined Phase Import Routines

Phase import routines are classes derived from `GSASIIIO.ImportPhase`. They must be found in files named *G2phase*.py* that are in the Python path and the class must override the `__init__` method and add a `Reader` method. The distributed routines are:

15.2.1 Module G2phase: PDB and .EXP

A set of short routines to read in phases using routines that were previously implemented in GSAS-II: PDB and GSAS .EXP file formats

```
class G2phase.EXP_ReaderClass
    Routine to import Phase information from GSAS .EXP files

    ContentsValidator (filepointer)
        Look for a VERSION tag in 1st line

    ReadEXPPhase (G2frame, filepointer)
        Read a phase from a GSAS .EXP file.

    Reader (filename, filepointer, ParentFrame=None, **unused)
        Read a phase from a GSAS .EXP file using ReadEXPPhase ()

class G2phase.PDB_ReaderClass
    Routine to import Phase information from a PDB file

    ContentsValidator (filepointer)
        Taking a stab a validating a PDB file (look for cell & at least one atom)

    ReadPDBPhase (filename, parent=None)
        Read a phase from a PDB file.

    Reader (filename, filepointer, ParentFrame=None, **unused)
        Read a PDF file using ReadPDBPhase ()
```

15.2.2 Module *G2phase_GPX: Import phase from GSAS-II project*

Copies a phase from another GSAS-II project file into the current project.

class `G2phase_GPX.PhaseReaderClass`

Opens a .GPX file and pulls out a selected phase

ContentsValidator (*filepointer*)

Test if the 1st section can be read as a cPickle block, if not it can't be .GPX!

Reader (*filename, filepointer, ParentFrame=None, **unused*)

Read a phase from a .GPX file. Does not (yet?) support selecting and reading more than one phase at a time.

15.2.3 Module *G2phase_CIF: Coordinates from CIF*

Parses a CIF using PyCifRW from James Hester and pulls out the structural information.

If a CIF generated by ISODISTORT is encountered, extra information is added to the phase entry and constraints are generated.

class `G2phase_CIF.CIFPhaseReader`

Implements a phase importer from a possibly multi-block CIF file

ISODISTORT_proc (*blk, atomlbllist, ranIdlookup*)

Process ISODISTORT items to create constraints etc.

15.3 Currently Defined Powder Data Import Routines

Powder data import routines are classes derived from `GSASIIIO.ImportPowderData`. They must be found in files named *G2pwd*.py* that are in the Python path and the class must override the `__init__` method and add a `Reader` method. The distributed routines are:

15.3.1 Module *G2pwd_GPX: GSAS-II projects*

Routine to import powder data from GSAS-II .gpx files

class `G2pwd_GPX.GSAS2_ReaderClass`

Routines to import powder data from a GSAS-II file This should work to pull data out from a out of date .GPX file as long as the details of the histogram data itself don't change

ContentsValidator (*filepointer*)

Test if the 1st section can be read as a cPickle block, if not it can't be .GPX!

Reader (*filename, filepointer, ParentFrame=None, **kwarg*)

Read a dataset from a .GPX file. If multiple datasets are requested, use self.repeat and buffer caching.

15.3.2 Module *G2pwd_fxyc: GSAS data files*

Routine to read in powder data in a variety of formats that are defined for GSAS.

class `G2pwd_fxyc.GSAS_ReaderClass`

Routines to import powder data from a GSAS files

ContentsValidator (*filepointer*)

Validate by checking to see if the file has BANK lines

Reader (*filename, filepointer, ParentFrame=None, **kwarg*)

Read a GSAS (old formats) file of type FXY, FXYE, ESD or STD types. If multiple datasets are requested, use self.repeat and buffer caching.

G2pwd_fxye.**sfloat** (*S*)

convert a string to a float, treating an all-blank string as zero

G2pwd_fxye.**sint** (*S*)

convert a string to an integer, treating an all-blank string as zero

15.3.3 Module G2pwd_xye: Topas .xye data

Routine to read in powder data from a Topas-compatible .xye file

class G2pwd_xye.**xye_ReaderClass**

Routines to import powder data from a .xye file

ContentsValidator (*filepointer*)

Look through the file for expected types of lines in a valid Topas file

Reader (*filename, filepointer, ParentFrame=None, **unused*)

Read a Topas file

15.3.4 Module G2pwd_CIF: CIF powder data

Routine to read in powder data from a CIF.

class G2pwd_CIF.**CIFpwdReader**

Routines to import powder data from a CIF file

ContentsValidator (*filepointer*)

Use standard CIF validator

Reader (*filename, filepointer, ParentFrame=None, **kwarg*)

Read powder data from a CIF. If multiple datasets are requested, use self.repeat and buffer caching.

15.4 Currently Defined Single Crystal Data Import Routines

Single crystal data import routines are classes derived from , `GSASIIIO.ImportStructFactor`. They must be found in files named *G2sfact*.py* that are in the Python path and the class must override the `__init__` method and add a `Reader` method. The distributed routines are:

15.4.1 Module G2sfact: simple HKL import

Read structure factors from a simple hkl file. Two routines are provided to read from files containing F or F² values.

G2sfact.**ColumnValidator** (*parent, filepointer*)

Validate a file to check that it contains columns of numbers

class G2sfact.**HKLF2_ReaderClass**

Routines to import F**2, sig(F**2) reflections from a HKLF file

ContentsValidator (*filepointer*)

Make sure file contains the expected columns on numbers

Reader (*filename, filepointer, ParentFrame=None, **unused*)

Read the file

class G2sfact.**HKLF_ReaderClass**

Routines to import F, sig(F) reflections from a HKLF file

ContentsValidator (*filepointer*)

Make sure file contains the expected columns on numbers

Reader (*filename, filepointer, ParentFrame=None, **unused*)

Read the file

15.4.2 Module G2sfact_CIF: CIF import

Read structure factors from a CIF reflection table.

class G2sfact_CIF.**CIFhklReader**

Routines to import Phase information from a CIF file

ContentsValidator (*filepointer*)

Use standard CIF validator

Reader (*filename, filepointer, ParentFrame=None, **kwarg*)

Read single crystal data from a CIF. If multiple datasets are requested, use self.repeat and buffer caching.

REQUIRED PACKAGES

Note that GSAS-II requires the Python extension packages

- wxPython (<http://wxpython.org/docs/api/>),
- NumPy (<http://docs.scipy.org/doc/numpy/reference/>),
- SciPy (<http://docs.scipy.org/doc/scipy/reference/>),
- matplotlib (<http://matplotlib.org/contents.html>) and
- PyOpenGL (<http://pyopengl.sourceforge.net/documentation>)

These are not distributed as part of the Python standard library and must be obtained separately (or in a bundled Python package such as the Enthought Python Distribution/Canopy). The PyOpenGL package is installed into Python by GSAS-II if not found, so it does not need to be included in the Python bundle.

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