

GSAS-II Developers Documentation *Release 0.2.0*

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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 ConstraintDialog (parent, title, text, data, separator='*')

Window to edit Constraint values

class GSASII. 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.GetPWDRdatafromTree(PWDRname)

Returns powder data from GSASII tree

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

GSASII.GetPhaseData()

Returns a list of defined phases. Used only in GSASIIgrid Note routine GSASIIstruct.GetPhaseData also exists.

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 :returns: two dicts:

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

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.OnFileClose (event)

Clears the data tree in response to the File/Close 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)

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.

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.OnViewLSParms (event)

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

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.

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

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

${\bf class} \; {\tt GSASII.ViewParmDialog} \; (parent, \, title, \, parmDict)$

Window to show all parameters in the refinement. Called from OnViewLSParms

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.

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

Each constraint is defined as a list using a series of terms of form

```
[[<mult1>, <var1>], [<mult2>, <var2>],..., <fixed val>, <vary flag>, <cons type>]
```

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

- <mult> is a multiplier for the constraint (float)
- <var> is the name of the variable (str) (or to be implemented a VarName object.)

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

- <fixed val> is the fixed value for a constraint equation or is None
- <vary flag> is True, False or None and is intended to use to indicate if new variables should be refined.
- <cons type> 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. The vary flag for those variables is ignored.
 - 'c' defines a constraint equation of form, $m_1 \times var_1 + m_2 \times var_2 + ... = c$

- 'h' defines a variable to hold (not vary). Any variable on this list is not varied, even if its refinement flag is set. This is of particular value when needing to hold one or more variables in a set such as the reciprocal metric tensor or anisotropic displacement parameter.
- 'f' defines a relationship to define a new variable 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	explan	ation					
newCellDict		dict	with	lattice	paran	neters	computed	by
		GSASI	IstrMat	th.GetNe	ewCellPar	ms() (dict	t)	
title		Name o	f gpx file((?) (str)				
variables		Values	for all N 1	efined var	iables (list of	f float value	es, length N, or	dered
		to matcl	h varyList	:)				
sig		Uncerta	inty value	es for all N	refined varia	ables (list c	of float values, l	ength
		N, orde	N, ordered to match varyList)					
varyList		List of o	List of directly refined variables (list of str values, length N)					
newAtomDict		dict	dict with atom position values computed in					
		GSASIIstrMath.ApplyXYZshifts() (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 (N	(N) covVa	ariance mat	trix (np.array	7)		

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	?
	Cell	List with 7 items: cell refinement flag (bool) a, b, c, (Angstrom, float)
		alpha, beta & gamma (degrees, float)
	Type	for now 'nuclear' (str)
	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
		Continued on next page

Table 2.1 – continued from previous page

Name SGData Pawley neg wt Flip Data plot type Mass POhkl Z vdWRadii Color AtomTypes AtomMass doPawley NoAtoms Pawley dmin BondRadii AngleRadii	phase name (str) Space group details as a space group (SGData) object as defined in GSASIIspc.SpcGroup(). Restraint value for negative Pawley intensities (float) Charge flip controls dict? ? Mass of unit cell contents in g/mol March-Dollase preferred orientation direction ? ? Colors for atoms (list of (r,b,g) triplets) List of atom types List of masses for atoms Flag for Pawley intensity extraction (bool) Number of atoms per unit cell of each type (dict) maximum Q (as d-space) to use for Pawley extraction (float) Radius for each atom used to compute interatomic distances (list of
Pawley neg wt Flip Data plot type Mass POhkl Z vdWRadii Color AtomTypes AtomMass doPawley NoAtoms Pawley dmin BondRadii	GSASIIspc.SpcGroup(). Restraint value for negative Pawley intensities (float) Charge flip controls dict? ? Mass of unit cell contents in g/mol March-Dollase preferred orientation direction ? ? Colors for atoms (list of (r,b,g) triplets) List of atom types List of masses for atoms Flag for Pawley intensity extraction (bool) Number of atoms per unit cell of each type (dict) maximum Q (as d-space) to use for Pawley extraction (float)
Flip Data plot type Mass POhkl Z vdWRadii Color AtomTypes AtomMass doPawley NoAtoms Pawley dmin BondRadii	Restraint value for negative Pawley intensities (float) Charge flip controls dict? ? Mass of unit cell contents in g/mol March-Dollase preferred orientation direction ? ? Colors for atoms (list of (r,b,g) triplets) List of atom types List of masses for atoms Flag for Pawley intensity extraction (bool) Number of atoms per unit cell of each type (dict) maximum Q (as d-space) to use for Pawley extraction (float)
Flip Data plot type Mass POhkl Z vdWRadii Color AtomTypes AtomMass doPawley NoAtoms Pawley dmin BondRadii	Charge flip controls dict? ? Mass of unit cell contents in g/mol March-Dollase preferred orientation direction ? ? Colors for atoms (list of (r,b,g) triplets) List of atom types List of masses for atoms Flag for Pawley intensity extraction (bool) Number of atoms per unit cell of each type (dict) maximum Q (as d-space) to use for Pawley extraction (float)
Data plot type Mass POhkl Z vdWRadii Color AtomTypes AtomMass doPawley NoAtoms Pawley dmin BondRadii	? Mass of unit cell contents in g/mol March-Dollase preferred orientation direction ? ? Colors for atoms (list of (r,b,g) triplets) List of atom types List of masses for atoms Flag for Pawley intensity extraction (bool) Number of atoms per unit cell of each type (dict) maximum Q (as d-space) to use for Pawley extraction (float)
Mass POhkl Z vdWRadii Color AtomTypes AtomMass doPawley NoAtoms Pawley dmin BondRadii	Mass of unit cell contents in g/mol March-Dollase preferred orientation direction ? ? Colors for atoms (list of (r,b,g) triplets) List of atom types List of masses for atoms Flag for Pawley intensity extraction (bool) Number of atoms per unit cell of each type (dict) maximum Q (as d-space) to use for Pawley extraction (float)
POhkl Z vdWRadii Color AtomTypes AtomMass doPawley NoAtoms Pawley dmin BondRadii	March-Dollase preferred orientation direction ? ? Colors for atoms (list of (r,b,g) triplets) List of atom types List of masses for atoms Flag for Pawley intensity extraction (bool) Number of atoms per unit cell of each type (dict) maximum Q (as d-space) to use for Pawley extraction (float)
Z vdWRadii Color AtomTypes AtomMass doPawley NoAtoms Pawley dmin BondRadii	? ? Colors for atoms (list of (r,b,g) triplets) List of atom types List of masses for atoms Flag for Pawley intensity extraction (bool) Number of atoms per unit cell of each type (dict) maximum Q (as d-space) to use for Pawley extraction (float)
vdWRadii Color AtomTypes AtomMass doPawley NoAtoms Pawley dmin BondRadii	? Colors for atoms (list of (r,b,g) triplets) List of atom types List of masses for atoms Flag for Pawley intensity extraction (bool) Number of atoms per unit cell of each type (dict) maximum Q (as d-space) to use for Pawley extraction (float)
Color AtomTypes AtomMass doPawley NoAtoms Pawley dmin BondRadii	Colors for atoms (list of (r,b,g) triplets) List of atom types List of masses for atoms Flag for Pawley intensity extraction (bool) Number of atoms per unit cell of each type (dict) maximum Q (as d-space) to use for Pawley extraction (float)
AtomTypes AtomMass doPawley NoAtoms Pawley dmin BondRadii	List of atom types List of masses for atoms Flag for Pawley intensity extraction (bool) Number of atoms per unit cell of each type (dict) maximum Q (as d-space) to use for Pawley extraction (float)
AtomMass doPawley NoAtoms Pawley dmin BondRadii	List of atom types List of masses for atoms Flag for Pawley intensity extraction (bool) Number of atoms per unit cell of each type (dict) maximum Q (as d-space) to use for Pawley extraction (float)
AtomMass doPawley NoAtoms Pawley dmin BondRadii	List of masses for atoms Flag for Pawley intensity extraction (bool) Number of atoms per unit cell of each type (dict) maximum Q (as d-space) to use for Pawley extraction (float)
doPawley NoAtoms Pawley dmin BondRadii	Flag for Pawley intensity extraction (bool) Number of atoms per unit cell of each type (dict) maximum Q (as d-space) to use for Pawley extraction (float)
NoAtoms Pawley dmin BondRadii	Number of atoms per unit cell of each type (dict) maximum Q (as d-space) to use for Pawley extraction (float)
Pawley dmin BondRadii	maximum Q (as d-space) to use for Pawley extraction (float)
BondRadii	
AngleRadii	floats)
11118101144011	Radius for each atom used to compute interatomic angles (list of floats)
	unique random number Id for phase (int)
	Phase Id number for current project (int).
	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</i>
	Records description. (list of lists)
	Display parameters (dict)
hallScale	Size of spheres in ball-and-stick display (float)
	dict with bonds
	? (float)
	Flag to show view point triplet (bool). True=show.
	cartesian viewing direction (np.array with three elements)
	clipping distance in A (float)
*	background for plot as and R,G,B triplet (default = $[0, 0, 0]$, black). (list
DackColor	with three atoms)
selected A toms	List of selected atoms (list of int values)
	Flag to highlight rigid body placement
_	Size ratio for H atoms (float)
	Size of binds in A (float)
	? (list)
	list of lists. First item in list is $[x,y,z]$ in fractional coordinates for the
A 1CM L OIIII	center of the plot. Second item ?.
chow Hydrogon	Flag to control plotting of H atoms.
	Flag to control display of the unit cell.
	Probability limit for display of thermal ellipsoids in % (float).
	Multiplier of van der Waals radius for display of vdW spheres.
	A list of lists with an entry for each atom that is plotted.
_	Step to de/increase Z-clip (float)
	Viewing quaternion (4 element np.array)
radius Eactor	Distance ratio for searching for bonds. ? Bonds are located that are within r(Ra+Rb) and (Ra+Rb)/r where Ra and Rb are the atomic radii.
1441401 40101	
	ballScale bondList contourLevel showABC viewDir Zclip backColor selectedAtoms showRigidBodies sizeH bondRadius atomPtrs viewPoint showHydrogen unitCellBox ellipseProb vdwScale Atoms Zstep Quaternion radiusFactor

2.3. Phase Tree Items 7

Table 2.1 – continued from previous page

key	sub-key	explanation
	oldxy	? (list with two floats)
	cameraPos	Viewing position in A for plot (float)
	depthFog	? (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

2.4 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: np.array([[0, 0, 0]])
SGOps	symmetry operations as a list of form [[M1,T1], [M2,T2],] 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 $X' = \begin{bmatrix} 1 & 1 & 1 \\ 1 & 1 & 1 \end{bmatrix}$
	$M_n * X + T_n$
SGSys	symmetry unit cell: type one of 'triclinic', 'monoclinic', 'orthorhombic', 'tetrag-
	onal', '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 ?).

2.5 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.

location	explanation
cx,cx+1,cx+2	the x,y and z coordinates
cx+3	fractional occupancy (also cs-1)
ct-1	atom label
ct	atom type
ct+1	refinement flags
cs	site symmetry string
cs+1	site multiplicity
cia	ADP flag: Isotropic ('I') or Anisotropic ('A')
cia+1	Uiso
cia+2cia+6	U11, U22, U33, U12, U13, U23

2.6 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 routine <code>GetUsedHistogramsAndPhasesfromTree()</code> 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 <code>Data</code>, as outlined below.

Background

key	sub-key	explanation
Limits		A list of two two element lists, as [[Ld,Hd],[L,H]] whe
		the current and default lowest two-theta value to be us
		and Hd are the current and default highest two-theta val
Reflection Lists		A dict with an entry for each phase in the histogram. The
		dict item is a list or reflections as described in the <i>Pov</i>
		description.
Instrument Parameters		A list containing two dicts where the possible keys in each
		below. The value for each item is a list containing three v
		value, the current value and a refinement flag which can
		True, False or 0 where 0 indicates a value that cannot
		first and second values are floats unless otherwise note
		first dict are noted as [1]
	Lam	Specifies a wavelength in Angstroms [1]
	Lam1	Specifies the primary wavelength in Angstrom, when ar
	Baini	source is used [1]
	Lam2 I(L2)/I(L1)	Specifies the secondary wavelength in Angstrom, when a
		source is used [1] Ratio of Lam2 to Lam1 [1]
	Туре	
	-JP~	Histogram type (str) [1]:
		 '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 differin
		[1]
	U, V, W	Cagliotti profile coefficients for Gaussian instrumen
		where the FWHM goes as $U \tan^2 \theta + V \tan \theta + W$ [1]
	X, Y	Cauchy (Lorentzian) instrumental broadening coefficien
	SH/L	Variant of the Finger-Cox-Jephcoat asymmetric peak b
		Note that this is the average between S/L and H/L wh
		height, H is the slit height and L is the goniometer diame
	Polariz.	Polarization coefficient. [1]
wtFactor		A weighting factor to increase or decrease the leverage o
		togram (float). A value of 1.0 weights the data with their
		tainties and a larger value increases the weighting of the
		to decreasing the uncertainties).
Sample Parameters		Specifies a dict with parameters that describe how the
		lected, as listed below. Refinable parameters are a list co
		and a bool, where the second value specifies if the value
		erwise 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 ra
	DisplaceX, DisplaceY	Sample displacement from goniometer center where Y is
		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 perfo
		directly in any computations.
	ranId	The random-number Id for the histogram (same value as
		key is ranId)
	Туре	Type of diffraction data, may be 'Debye-Scherrer' or 'E
		(str).
10	Diffuse	not in use?
10 hId		not in use? hapter 2. GSASIlobj: Data objects The number assigned to the histogram when the project
		edited (can change)
ranId		A random number id for the histogram that does not cha
Doolcaround		The healtground is stored as a list with where the first

The background is stored as a list with where the first

2.7 Powder Reflection Data Structure

For every phase in a histogram, the Reflection Lists value is a list of reflections. The items in that list are documented below.

in-	explanation
dex	
0,1,2	h,k,l (float)
3	multiplicity
4	d-space, Angstrom
5	pos, two-theta
6	sig, Gaussian width
7	gam, Lorenzian width
8	Fobs**2
9	Fcalc**2
10	reflection phase, in degrees
11	the equivalent reflections as a (m x 3) np.array, where m is 0.5 * multiplicity. Note that Freidel pairs,
	(-h,-k-,l), are not included.
12	phase shift for each of the equivalent reflections as a length (m) array
13	intensity correction for reflection, this times Fobs**2 or Fcalc**2 gives Iobs or Icalc
14	dict with the form factor (f or b) by atom type symbol at the reflection position.

2.8 Classes and routines

GSASIIobj.LoadHistogramIDs (histList, idList)

Save the Id values for a series of histograms

class GSASIIobj.VarName (*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 VarName object.

A VarName object can be created with a single parameter:

Parameters varname (str) -

- a single value can be used to create a VarName object. The string 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 VarName object can be created with exactly four positional parameters:

Parameters

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

fullDescr()

Return a longer description for a GSAS-II variable

Returns a short description or 'no definition' if not found

getDescr()

Return a short description for a GSAS-II variable

Returns a short description or 'no definition' if not found

name()

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

Returns the variable name as a str

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", 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 of substances commonly used for powder calibrations for image data.

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: 989 \$" that is set by subversion when the file is retrieved from subversion.

Place GSASIIpath.SetVersionNumber("\$Revision: 989 \$") 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.

Returns A dictionary with the files that have been changed/added and a code describing how they have been updated (see changetype) or None if there is a subversion error (likely because the path is not a repository or svn is not found)

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-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 irrevelant 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

${\tt GSASIIElem.GetMagFormFacCoeff}~(El)$

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

${\tt GSASIIElem.GetXsectionCoeff}~(El)$

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)

Needs a doc string

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

Needs a doc string

3.7 GSASIllattice: 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: $d^*=\sqrt{A_1h^2+A_2k^2+A_3l^2+A_4hk+A_5hl+A_6kl}$, 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.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

 ${\tt GSASIIIattice.Flnh}~(Start, SHCoef, phi, beta, SGData)$

needs doc string

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

Generate the positionally unique powder diffraction reflections

Parameters

- dmin minimum d-spacing in A
- **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. GetKclKsl (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

```
GSASIIIattice.Glnh (Start, SHCoef, psi, gam, SamSym) needs doc string
```

```
GSASIIlattice. Gmat2A(G)
```

Extract A from reciprocal metric tensor (G)

Parameters G – reciprocal maetric 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*x = np.inner(A,x) = X
- B (= inverse of A) for Cartesian to crystal transformation B*X = 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)

```
GSASIIIattice.Hx2Rh (Hx)
needs doc string

GSASIIIattice.MaxIndex (dmin, A)
needs doc string

GSASIIIattice.OdfChk (SGLaue, L, M)
needs doc string

GSASIIIattice.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,azmuth
                • 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_rDsq(H, A)
     needs doc string
GSASIIlattice.calc_rDsq2 (H, G)
     needs doc string
GSASIIlattice.calc_rDsqZ(H, A, Z, tth, lam)
     needs doc string
GSASIIlattice.calc_rV(A)
     Compute the reciprocal lattice volume (V*) from A
GSASIIlattice.calc_rVsq(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 =
               np.inner(A,x) = X B (= inverse of A) for Cartesian to crystal transformation B*X = 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.rotdMat(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.rotdMat4 (angle, axis=0)
     Prepare rotation matrix for angle in degrees about axis(=0,1,2) with scaling for OpenGL
```

3.7. GSASIllattice: Unit cells

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 0x279ea70>, <function test1 at 0x279eab0>, <function test2 at 0x279e
     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. ElemPosition (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

 ${\tt GSASIIspc.Muiso2Shkl}~(\textit{muiso}, \textit{SGData}, \textit{cell})$

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 >= 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 0x27b6630>, <function test1 at 0x27b6670>, <function test2 at 0x27b66b0> 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 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 GUI ROUTINES

4.1 GSASIIgrid: Basic GUI routines

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, elemlst, prelbl=[], postlbl=[], title='Edit items', header='', size=(300, 250))

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 Close 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 dialog

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

Displays help information in a primitive HTML browser type window

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()]

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)

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

Returns the selection index or None

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.

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 high-lighted. 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)

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.

${\tt TestValid}(tc)$

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))
```

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
- elemIst (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).

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

Example for use of ScrolledMultiEditor:

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. **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

 ${\bf class} \ {\tt GSASIIgrid.SymOpDialog} \ (parent, SGData, New=True, ForceUnit=False)$

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)

Needs a doc string

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'

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 appreviations 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 TextCrtl 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
- **size** (*wx.Size*) an optional size parameter that dictates the size for the TextCtrl. None (the default) indicates that the default size should be used.
- **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 used if the initial value for the dict/list element loc[key] is None. In this case typeHint must be int or float, which specifies the type for input to the TextCtrl. Defaults as None.

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=None)

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

${\bf class} \; {\tt GSASIIIO.ExportBaseclass} \; ({\it G2frame, formatName, longFormatName=None}) \\$

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

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)

- •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
- **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 importing of data files (diffraction data, coordinates,...

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

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

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

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 indicies for each choice (or None)

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

 $\textbf{PhaseSelector} \ (\textit{ChoiceList}, \textit{ParentFrame} = \textit{None}, \textit{title} = \textit{`Select a phase'}, \textit{size} = \textit{None}, \textit{header} = \textit{`Phase'}, \textit{size} = \textit{None}, \textit{size} = \textit{N$

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

Defines a base class for the reading of files with coordinates

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/sig(intensity)^2 values (weights) np.array(yc), # calc. intensities (zero) np.array(yb), # calc. background (zero) np.array(yd), # obs-calc profiles

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

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

Scan through the reflections to update the Controls dictionary

```
GSASIIIO. IndexPeakListSave (G2frame, peaks)
     Save powder peaks from the indexing list
class GSASIIIO.MultipleChoicesDialog(choicelist,
                                                            headinglist,
                                                                         head='Select options',
                                               tle='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
GSASIIIO.ProjFileSave (G2frame)
     Save a GSAS-II project file
GSASIIIO.PutG2Image (filename, Comments, Data, Npix, image)
     Write an image as a python pickle
GSASIIIO.ReadEXPPhase (G2frame, filename)
     Read a phase from a GSAS .EXP file. Called in the GSAS phase import routine (see imports/G2phase.py)
GSASIIIO.ReadPDBPhase (filename)
     Read a phase from a PDB file. Called in the PDB phase import routine (see imports/G2phase.py)
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 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
```

4.3 ReadMarCCDFrame: Read Mar Files

class ReadMarCCDFrame.marFrame (File, byteOrd='<', IFD={})</pre>

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 <= 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 man (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. **UpdateDData** (*G2frame*, *DData*, *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
- **DData** (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.

5.5 GSASIlimgGUI: 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.DistAngle (DisAglCtls, DisAglData)
Needs a doc string

GSASIIstrMain.Refine (GPXfile, dlg)
Needs a doc string

GSASIIstrMain.SeqRefine (GPXfile, dlg)
Needs a doc string

GSASIIstrMain.main ()
Needs a doc string
```

6.2 GSASIIstrMath - structure math routines

```
GSASIIstrMath. ApplyRBModelDervs (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, G, g, pfx, phfx, hfx, SGData, calcControls, parmDict) Needs a doc string
- GSASIIstrMath.**GetIntensityDerv** (refl, G, g, pfx, phfx, hfx, SGData, calcControls, parmDict) Needs a doc string
- GSASIIstrMath.**GetNewCellParms** (parmDict, varyList)
 Needs a doc string
- GSASIIstrMath.**GetPrefOri** (refl, G, g, phfx, hfx, SGData, calcControls, parmDict)
 Needs a doc string
- GSASIIstrMath.**GetPrefOriDerv** (*refl*, *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
- $\texttt{GSASIIstrMath.GetSampleSigGam} \ (\textit{refl}, \textit{wave}, \textit{G}, \textit{GB}, \textit{phfx}, \textit{calcControls}, \textit{parmDict}) \\ \textbf{Needs a doc string}$
- ${\tt GSASIIstrMath.GetSampleSigGamDerv} \ (\textit{refl}, \textit{wave}, \textit{G}, \textit{GB}, \textit{phfx}, \textit{calcControls}, \textit{parmDict}) \\ {\tt Needs a doc string}$
- 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 "analtytic 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** (refList, 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

- refList (*list*) [ref] where each ref = h,k,l,m,d,...,[equiv h,k,l],phase[equiv]
- **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** (refList, 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

 ${\tt GSASIIstrMath.dervRefine} \ (\textit{values}, \ \textit{HistoPhases}, \ \textit{parmDict}, \ \textit{varylist}, \ \textit{calcControls}, \ \textit{pawleyLookup}, \\ \textit{dlg})$

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 "analtytic Jacobian" in Controls.

Returns Jacobian numpy.array dMdv for all histograms concatinated

- GSASIIstrMath.errRefine (values, HistoPhases, parmDict, varylist, calcControls, pawleyLookup, dlg) Needs a doc string
- GSASIIstrMath.getPowderProfile (parmDict, x, varylist, Histogram, Phases, calcControls, pawley-Lookup)

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

${\tt GSASIIstrIO.CheckConstraints} \, (\textit{GPXfile})$

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

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
- hNames (str) list of histogram names

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

GSASIIstrIO. GetHistogramPhaseData(Phases, Histograms, Print=True, pFile=None, resetRe-fList=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={}*, *rbIds={}*, *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.**GetRestraints** (*GPXfile*)

Read the restraints 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

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.**PrintRestraints** (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.

Parameters constList (*list*) – a list of lists where each item in the outer list specifies a constraint of some form. The last item in each inner list determines which of the four constraints types has been input:

- h (hold): a single variable that will not be varied. It will be removed from the varyList later.
- c (constraint): specifies a linear relationship that can be varied as a new grouped variable a fixed value.
- f (fixed): specifies a linear relationship that is assigned a fixed value.
- e (equivalence): specifies a series of variables where the first variable in the last can be used to generate the values for all the remaining variables.

Returns

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

- constDict (list) contains the constraint relationships
- fixedList (list) contains the fixed values for 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

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, RigidBodies, Cov-Data, 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
- **RigidBodies** (*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 contraints, parameter redefinition and parameter simplification contraints.

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 Pj is a parameter name and Mjk is a constant.

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

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

where Cn is a constant. These equations define an algebraic constrant.

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:

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 P0, P1 and P2,... 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 Group-Constraints.) 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 contraint 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 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).

- If all parameters in a parameter redefinition (new var) relationship are varied, the parameter assigned to this expression (::constr:n, see paramPrefix) newly generated parameter is varied. Note that any generated "missing" relations are not varied. Only the input relations are 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 in internally consistent, use CheckConstraints

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

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

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 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 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

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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 contraints where each constraint grouped containts 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 1st 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

```
GSASIImapvars.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

GSASIImapvars.**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 {'param1': mult1, 'param2': mult2,...}
```

Returns

two lists of lists:

- a list of grouped contraints where each constraint grouped containts a list of indices for constraint constrDict entries
- a list containing lists of parameter names contained in each group

```
GSASIImapvars.InitVars()
```

Initializes all constraint information

```
GSASIImapvars.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

GSASIImapvars.**PrintIndependentVars** (parmDict, varyList, sigDict, PrintAll=False, pFile=None) Print the values and uncertainties on the independent variables

```
GSASIImapvars. 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),]

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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 contraint relationships

GSASIIIMAGE: IMAGE CALC MODULE

```
Ellipse fitting & image integration
needs some minor refactoring to remove wx code
GSASIIimage.EdgeFinder(image, data)
     this makes list of all x,y where I>edgeMin suitable for an ellipse search?
GSASIIimage.Fill2ThetaAzimuthMap (masks, TA, tam, image)
     Needs a doc string
GSASIIimage.FitCircle(ring)
     Needs a doc string
GSASIIimage.FitDetector (rings, varyList, parmDict)
     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)
     Needs a doc string
GSASIIimage.GetDetXYfromThAzm(Th, Azm, data)
     Needs a doc string
GSASIIimage. GetDetectorXY (dsp, azm, data)
     Needs a doc string
GSASIIimage.GetDsp (x, y, data)
     Needs a doc string
GSASIIimage.GetEllipse (dsp, data)
     Needs a doc string
GSASIIimage.GetTth(x, y, data)
     Needs a doc string
GSASIIimage.GetTthAzm(x, y, data)
     Needs a doc string
```

```
GSASIIimage.GetTthAzmDsp(x, y, data)
     Needs a doc string
GSASIIimage. ImageCalibrate (self, data)
     Needs a doc string
GSASIIimage. ImageCompress (image, scale)
     Needs a doc string
GSASIIimage. ImageIntegrate (image, data, masks)
     Needs a doc string
GSASIIimage.ImageLocalMax(image, w, Xpix, Ypix)
     Needs a doc string
GSASIIimage.ImageRecalibrate (self, data)
     Needs a doc string
GSASIIimage.Make2ThetaAzimuthMap(data, masks, iLim, jLim)
     Needs a doc string
GSASIIimage.calcDist(radii, tth)
     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
                • \mathbf{phi} - his \mathbf{phi} = sample \mathbf{phi} rotation; usually = 0, axis rotates with omg.
                • azm – his chi = azimuth around incident beam
                • th - his theta = theta
GSASIIimage.calcZdisCosB(radius, tth, radii)
     Needs a doc string
GSASIIimage.checkEllipse(Zsum, distSum, xSum, ySum, dist, x, y)
     Needs a doc string
GSASIIimage.makeIdealRing(ellipse, azm=None)
     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
```

GSASIIMATH: COMPUTATION MODULE

Routines for least-squares minimization and other stuff

GSASIImath.AV2Q(A, V)

convert angle (radians) & vector to quaternion q=r+ai+bj+ck

GSASIImath. AVdeg2Q(A, V)

convert angle (degrees) & vector to quaternion q=r+ai+bj+ck

GSASIImath. **AtomTLS2UIJ** (atomData, atPtrs, Amat, rbObj) default doc string

Parameters name (*type*) – description

Returns type name: description

GSASIImath. AtomUij2TLS (atomData, atPtrs, Amat, Bmat, rbObj) default doc string

Parameters name (type) – description

Returns type name: description

GSASIImath. ChargeFlip (data, reflData, pgbar) default doc string

Parameters name (*type*) – description

Returns type name: description

GSASIImath.**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

GSASIImath.**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 containting 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, reflData)
     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 (pld, 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 (pld, 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, reflData)

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 map-Data['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, RBData)

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. UpdateRBXYZ (Bmat, RBObj, RBData, 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, max-eval=None, maxaccept=None, maxiter=400, boltzmann=1.0, learn_rate=0.5, feps=1e-06, quench=1.0, m=1.0, n=1.0, lower=-100, upper=100, dwell=50, slope=0.9, dlg=None)

Minimize a function using simulated annealing.

Schedule is a schedule class implementing the annealing schedule. Available ones are 'fast', 'cauchy', 'boltz-mann'

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).
- **feps** (*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

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 (\psi)
                • 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.getRamaDeriv(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 (nparray) – crystallographic coordinates of 5 atoms
                 • Amat (nparray) – crystal to cartesian transformation matrix
                 • Coeff (list) – pseudo potential coefficients
          Returns list (deriv) derivatives of pseudopotential with respect to 5 atom crystallographic xyz coor-
               dinates.
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.getRestDeriv(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 (*nparray*) crystallographic coordinates of 5 atoms
- Amat (nparray) 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

 ${\tt GSASIImath.getTOFalpha}$ (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

 ${\tt GSASIImath.getTOFbetaDeriv}\,(\mathit{dsp})$

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 matric for
                 • varyList (list) – full list of all variables in v-cov matrix
                 • covMatrix (nparray) – full variance-covariance matrix from the last least squares refine-
                   ment
          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, pgbar)
     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.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. Coehlo 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
GSASII index. 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
GSASII index. Values 2A (ibrav, values)
     needs a doc string
GSASIIindex.calc_M20 (peaks, HKL)
     needs a doc string
GSASIIindex.findBestCell(dlg, ncMax, A, Ntries, ibrav, peaks, V1)
     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
GSASII index.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
\texttt{GSASII} \texttt{index.scaleAbyV} (A, V)
     needs a doc string
GSASIIindex.sortM20 (cells)
     needs a doc string
GSASIIindex.swapMonoA(A)
     needs a doc string
```

GSASIIPLOT: 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
     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.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

GSASIIpwd. 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

GSASIIpwd. **AbsorbDerv** (*Geometry*, *MuR*, *Tth*, *Phi=0*, *Psi=0*) needs a doc string

GSASIIpwd. CalcPDF (data, inst, xydata) needs a doc string

GSASIIpwd.Dict2Values (parmdict, varylist)

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

GSASIIpwd.DoPeakFit (FitPgm, Peaks, Background, Limits, Inst, Inst2, data, oneCycle=False, controls=None, dlg=None)

needs a doc string

GSASIIpwd.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)

GSASIIpwd.GetNumDensity(ElList, Vol)

needs a doc string

 ${ t GSASIIpwd.LorchWeight}(Q)$

needs a doc string

 $\texttt{GSASIIpwd.Oblique}\,(\textit{ObCoeff},\textit{Tth})$

needs a doc string

GSASIIpwd.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)*npcosd(Azm)**2+Pola*npsind(Azm)**2)*npcosd(Tth)**2+ (1-Pola)*npsind(Azm)**2+Pola*npcosd(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

 ${\tt GSASIIpwd.calcIncident}~(\textit{Iparm}, \textit{xdata})$

needs a doc string

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)

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*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 \ge 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(TTh, 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.qetWidthsCW (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.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

 $\textbf{class} \ \texttt{GSASIIpwd.norm_gen} \ (\textit{momtype=1}, \ a=None, \ b=None, \ xa=-10.0, \ xb=10.0, \ xtol=1e-14, \ badvalue=None, name=None, longname=None, shapes=None, extradoc=None) \\ \text{needs a doc string}$

GSASIISOLVE - STRUCTURE SOLVING ROUTINES

```
GSASIIsolve.ShowBanner()
Print authorship, copyright and citation notice

GSASIIsolve.ShowControls(Controls)
Print controls information

GSASIIsolve.Solve(GPXfile)
perform the computation

GSASIIsolve.main()
needs doc string
```

Note that this program requires the Python wxPython, NumPy, SciPy, matplotlib packages, plus the PyOpenGL package (which is installed by GSAS-II if not found).

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