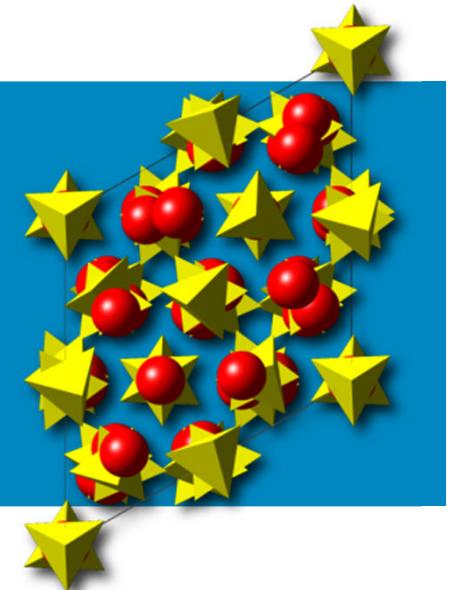


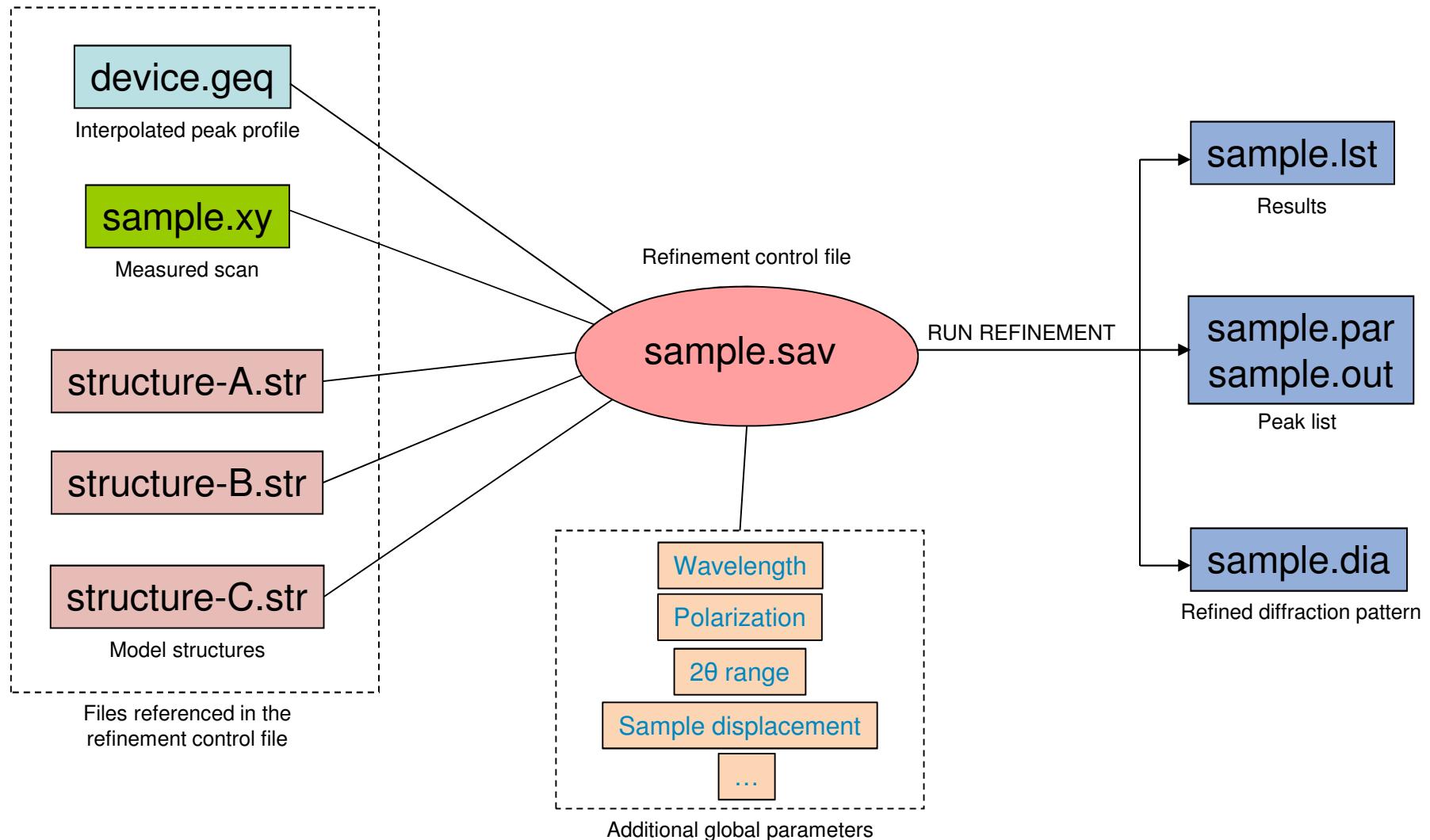
Lesson 8

Crystal Structure Files



Nicola Döbelin
RMS Foundation, Bettlach, Switzerland

Repetition: BGMN Project Structure



Structure Databases

- STR files shipped with Profex (created manually by the Profex developer)
- <http://www.bgmn.de/download-structures.html> (created manually by BGMN team)
- Create manually from:
 - ICSD (<http://www.fiz-karlsruhe.de/icsd.html>)
 - PDF-4+ (<http://www.icdd.com/>)
 - American Mineralogist Structure Database
(<http://rruff.geo.arizona.edu/AMS/amcsd.php>)
 - Crystallography Open Database COD
(<http://www.crystallography.net/>)
 - Cambridge Crystallographic Data Centre
(<http://beta-www.ccdc.cam.ac.uk/pages/Home.aspx>)



Commercial

Open Access

A screenshot of a Firefox browser window showing the AMCSD search interface. The title bar reads "Firefox" and the address bar shows the URL "rruff.geo.arizona.edu/AMS/amcsd.php". The main content area has a black header bar with the text "American Mineralogist Crystal Structure Database". Below this, a paragraph of text describes the database's purpose and maintenance. The search form contains a text input field with "anatase" and several buttons: "Mineral", "Author", "Chemistry Search", "Cell Parameters and Symmetry", "Diffraction Search", "General Search", and "Search Tips". At the bottom of the search form are "Search" and "Reset" buttons. Below the search form is a "Logic interface" section with radio buttons for "AND" and "OR", and a "Viewing" section with options for "amc long form", "amc short form", "cif", "amc", "cif", and "diffraction data".

AMCSD: Retrieving CIF Files

The screenshot shows a Firefox browser window displaying the American Mineralogist Crystal Structure Database (AMCSD) search results. The URL in the address bar is rruff.geo.arizona.edu/AMS/result.php. The page title is "American Mineralogist Crystal Structure Database". A message at the top says "7 matching records for this search." Below this, there are two entries:

[Anatase](#)
Howard C J, Sabine T M, Dickson F
Acta Crystallographica B47 (1991) 462-468
Structural and thermal parameters for rutile and anatase
Locality: synthetic
_database_code_amcsd 0019093
3.7845 3.7845 9.5143 90 90 90 I4_1/amd
atom x y z U(1,1) U(2,2) U(3,3) U(1,2) U(1,3) U(2,3)
Ti 0 0 0 .0052 0 0 .0052 0 .0070
O 0 0 .20806 .0117 0 0 .0027 0 .0072

[Download AMC data \(View Text File\)](#)
[Download CIF data \(View Text File\)](#)
[Download diffraction data \(View Text File\)](#)
[View JMOL 3-D Structure](#)

[Anatase](#)
Parker R
Zeitschrift fur Kristallographie 59 (1924) 1-54
Zur Kristallstruktur von Anatass und Rutil. (II. Teil. Die Anatassstruktur).

A red box highlights the "Download CIF data (View Text File)" link. A green arrow points from this link to a green box containing the text "Download CIF file". A red box also highlights the year "1924" in the second entry.

Watch out for high-temp and high-pressure datasets

Download CIF file

The screenshot shows a web browser window for the COD website. The URL in the address bar is www.crystallography.net/search.html. The page title is "Crystallography Open Database". On the left, there is a sidebar with several sections:

- COD Home**: Home, What's new?
- Accessing COD Data**: Browse, Search, Search by structural formula
- Add Your Data**: Deposit your data, Manage depositions, Manage/release prepublications
- Documentation**: COD Wiki, Obtaining COD, Citing COD, COD Mirrors, Advices to donators, Useful links

The main content area is titled "Search" and includes a note: "(For more information on search see the [hints and tips](#))". It features two search boxes: "Search by COD ID:" and "Enter SMILES:". Below these is a note: "Note: substructure search by SMILES is currently available in a subset of COD containing 70 000 structures." A large form follows, containing fields for text (1 or 2 words), journal, year, volume, issue, Z (min, max), Z' (min, max), 1 to 8 elements, NOT these elements, volume min and max, number of distinct elements min and max, filters (with a checkbox for "has F_{obs}"), and buttons for "Reset" and "Send".

COD: Retrieving CIF Files

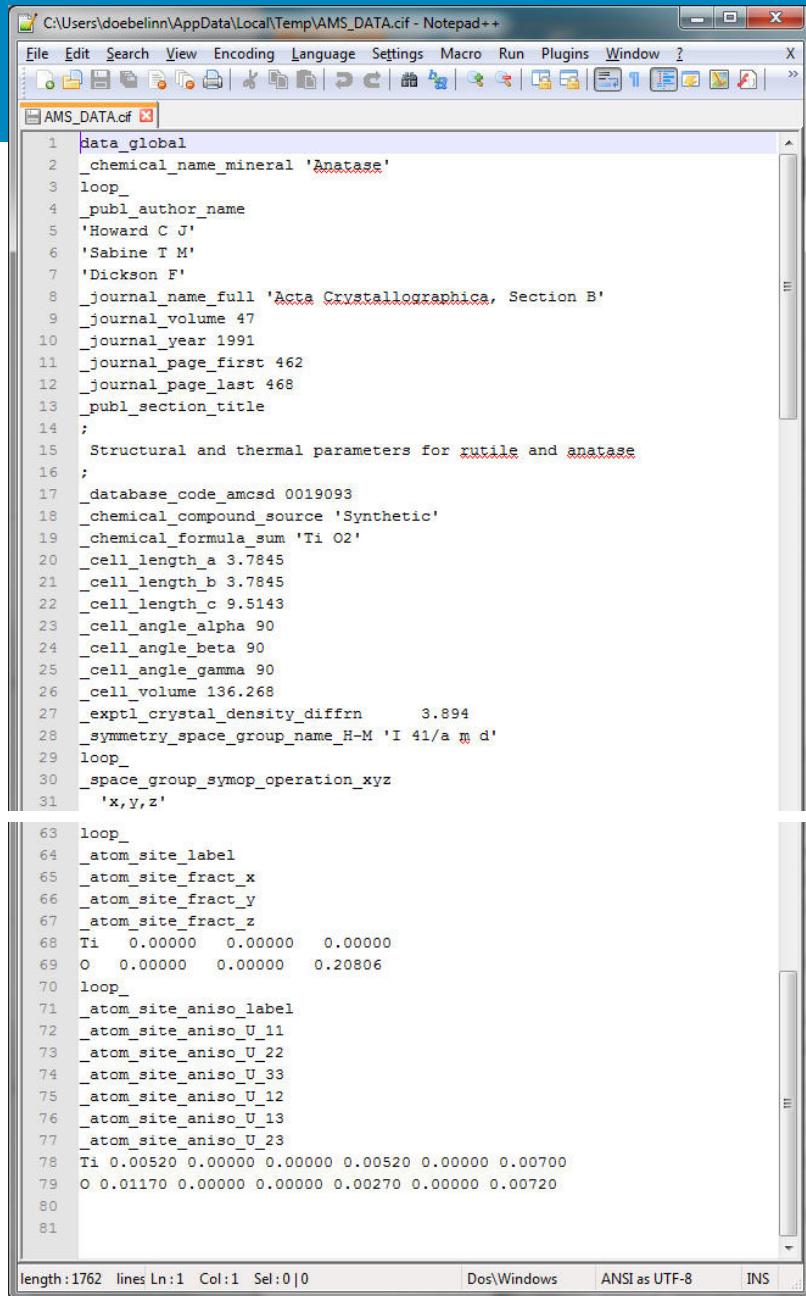
The screenshot shows a web browser displaying the Crystallography Open Database (COD) search results. The URL in the address bar is www.crystallography.net/result.php?CODSESSION=of6r7lhhh8n7a7k5fksld7pbldta1lh&order_by=file&order_desc. The main content area displays a table of search results with 9 entries. A green box highlights the 'CIF' link for the first entry (COD ID 9015929). A green arrow points from this link to a green box containing the text 'Download CIF file'. The table columns include COD ID, Links, Formula, Space group, Cell parameters, Cell volume, and Bibliography.

COD ID	Links	Formula	Space group	Cell parameters	Cell volume	Bibliography
9015929	CIF	O ₂ Ti	I ₄₁ /a m d :1	3.7845; 3.7845; 9.5143	136.268	Howard, C. J.; Sabine, T. M.; Dickson, F. Structural and thermal properties of anatase <i>Acta Crystallographica</i> , Wyckoff, R. W. G. Second edition. Interscience <i>Crystal Structures</i> , 1963, 1, 239-444
9009086	CIF	O ₂ Ti	I ₄₁ /a m d :1	3.785; 3.785; 9.514	136.3	Horn, M.; Schwerdtfeger, C. F.; Meagher, E. P. Refinement of the structure of anatase at several temperatures Sample: T = 800 C Locality: Legenbach quarry, Binnatal, Switzerland <i>Zeitschrift für Kristallographie</i> , 1972, 136, 273-281
9008216	CIF	O ₂ Ti	I ₄₁ /a m d :1	3.804; 3.804; 9.614	139.119	Horn, M.; Schwerdtfeger, C. F.; Meagher, E. P. Refinement of the structure of anatase at several temperatures Sample: T = 800 C Locality: Legenbach quarry, Binnatal, Switzerland <i>Zeitschrift für Kristallographie</i> , 1972, 136, 273-281
9008215	CIF	O ₂ Ti	I ₄₁ /a m d :1	3.7971; 3.7971; 9.579	138.11	Horn, M.; Schwerdtfeger, C. F.; Meagher, E. P. Refinement of the structure of anatase at several temperatures Sample: T = 600 C Locality: Legenbach quarry, Binnatal, Switzerland <i>Zeitschrift für Kristallographie</i> , 1972, 136, 273-281
9008214	CIF	O ₂ Ti	I ₄₁ /a m d :1	3.7892; 3.7892; 9.537	136.933	Horn, M.; Schwerdtfeger, C. F.; Meagher, E. P. Refinement of the structure of anatase at several temperatures Sample: T = 300 C Locality: Legenbach quarry, Binnatal, Switzerland <i>Zeitschrift für Kristallographie</i> , 1972, 136, 273-281
9008213	CIF	O ₂ Ti	I ₄₁ /a m d :1	3.7842; 3.7842; 9.5146	136.251	Horn, M.; Schwerdtfeger, C. F.; Meagher, E. P. Refinement of the structure of anatase at several temperatures Sample: T = 25 C Locality: Legenbach quarry, Binnatal <i>Zeitschrift für Kristallographie</i> , 1972, 136, 273-281

CIF File Format

CIF (Crystallographic Information File) Format:

- Standardized file format
- Contains structural information
- Contains references
- Specification available at <http://www.iucr.org>



The screenshot shows a Notepad++ window displaying a CIF (Crystallographic Information File) document. The file path is C:\Users\doebelin\My Documents\Temp\AMS_DATA.cif. The window title is "AMS_DATA.cif". The text content is a structured data file with various sections and loops, primarily describing crystallographic parameters for Anatase.

```
1 data_global
2 _chemical_name_mineral 'Anatase'
3 loop_
4 _publ_author_name
5 'Howard C J'
6 'Sabine T M'
7 'Dickson F'
8 _journal_name_full 'Acta Crystallographica, Section B'
9 _journal_volume 47
10 _journal_year 1991
11 _journal_page_first 462
12 _journal_page_last 468
13 _publ_section_title
14 ;
15 Structural and thermal parameters for rutile and anatase
16 ;
17 _database_code_amcsd 0019093
18 _chemical_compound_source 'Synthetic'
19 _chemical_formula_sum 'Ti O2'
20 _cell_length_a 3.7845
21 _cell_length_b 3.7845
22 _cell_length_c 9.5143
23 _cell_angle_alpha 90
24 _cell_angle_beta 90
25 _cell_angle_gamma 90
26 _cell_volume 136.268
27 _exptl_crystal_density_diffrn      3.894
28 _symmetry_space_group_name_H-M 'I 41/a m d'
29 loop_
30 _space_group_symop_operation_xyz
31 'x,y,z'

32
33 loop_
34 _atom_site_label
35 _atom_site_fract_x
36 _atom_site_fract_y
37 _atom_site_fract_z
38 Ti  0.00000   0.00000   0.00000
39 O   0.00000   0.00000   0.20806
40
41 loop_
42 _atom_site_aniso_label
43 _atom_site_aniso_U_11
44 _atom_site_aniso_U_22
45 _atom_site_aniso_U_33
46 _atom_site_aniso_U_12
47 _atom_site_aniso_U_13
48 _atom_site_aniso_U_23
49 Ti 0.00520 0.00000 0.00000 0.00520 0.00000 0.00700
50 O 0.01170 0.00000 0.00000 0.00270 0.00000 0.00720
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```

CIF Format Specification

<http://www.iucr.org/resources/cif/dictionaries>

The screenshot shows the IUCr website's "CIF dictionaries" page. At the top, there is a navigation bar with links for iucr, journals, books, news, education, people, resources, iycr2014, world directory, other directories, data, cif, lists, blogs, forums, commissions, nexus, and symmetry font. Below the navigation bar, the breadcrumb trail shows Home > resources > cif > dictionaries. On the left, there is a sidebar with a search bar and a list of CIF dictionary types: Search for CIF data name, Core CIF dictionary, Restraints dictionary, Powder CIF dictionary, Modulated structures, CIF dictionary, Electron density CIF dictionary, Twinning CIF dictionary, Macromolecular CIF dictionary, Image CIF dictionary, Symmetry CIF dictionary, and Suggested new CIF data items.

CIF dictionaries

CIF dictionaries provide a formal taxonomy of crystallographic terms and ideas. Dictionary entries are constructed in a structured machine-readable manner that facilitates validation and structuring of data. New entries may be devised for public or private dictionaries. A candidate data-name definition should fulfil the following conditions: (i) describe a specific and well defined concept: precision of definition is essential for an effective interchange mechanism; (ii) have appropriate granularity: data names can define a very small piece of information (a standard uncertainty on a particular physical measurable) or a very large amount (the text of a scientific paper). An appropriate choice should be made (and for DDL2 formalized through membership of subcategories, category and category groups, as appropriate); (iii) have well defined relationships with other data items (through its assigned category membership and parent/child links); for DDL2 the prior construction of a formal entity/relationship schema may be helpful; (iv) constraints on the data type and permissible values should be provided where applicable; (v) the name chosen should be globally unique; this is achieved through monitoring of names in public dictionaries by a regulatory committee (COMCIFS) and by registering of prefix strings for exclusive use in local dictionaries. Some thought may need to be applied to the choice of DDL appropriate for a candidate dictionary.

Current CIF Dictionaries

Canonical data descriptors and their attributes are presented in machine-readable data dictionaries.

Descriptors for small-molecule, inorganic and other small-unit-cell structures

- Core dictionary (coreCIF)
[ASCII | HTML | PDF | More information]
- Restraints dictionary
[ASCII | HTML | PDF | More information]
- Powder dictionary (pdCIF)
[ASCII | HTML | PDF | More information]
- Modulated and composite structures dictionary (msCIF)
[ASCII | HTML | PDF | More information]
- Electron density dictionary (rhoCIF)
[ASCII | HTML | PDF | More information]
- Twinning dictionary
[ASCII | HTML | PDF | More information]

Problems with CIF Files

Problems with CIF files:

1. May be (and very often ARE) incomplete
2. May use non-standard atomic settings
3. Must be converted to STR format for BGMN



Creating Structure Files

Official Documentation:

<http://www.bgmn.de/variables.html>

Control File (*.sav)

Structure File (*.str)

Device Conf. File (*.sav)

Variables for use with BGMN

The expression interpreter

- [standard functions](#)
- [multiple equation symbols in assignments](#)
- [user-defined string functions](#)

Variables in the task description SAV file for BGMN

NTHREADS, VAL, VERZERR, STRUC, STRUCOUT, SimpleSTRUOUT, PDBOUT, RESOUT, FCFOUT, OUTPUT, LIST, RU, UNT, UNTC, DDM, PARAM, LAMBDA, SYNCHROTRON, EPS1, EPS2, EPS3, EPS4, POL, PROTOKOLL, ONLYISO, ITMAX, DIAGRAMM, PLAN, STANDALONEPLAN, GOAL, WMIN, WMAX, CUT, LIMIT2, LIMIT4, LIMIT6, LIMIT8, LIMIT10, ANISOLIMIT, ANISO4LIMIT

Items in the structure description *.str-file

Definition of the lattice

SpacegroupNo, HermannMauguin, GeneralCondition, A, B, C, ALPHA, BETA, GAMMA, UNIT

Anisotropic variables

ANISO, ANISOLIN, ANISOSQR, ANISO4

Scaling factor/preferred orientation

GEWICHT, SPHAR0, SPHAR2, SPHAR4, SPHAR6, SPHAR8, SPHAR10

Real structure

B1, B2, k1, k2, k3, sk, H, h, k, l, zweiTheta, RP

Peak deselection

GeneralCondition

Atomic positions

E=, Wyckoff, x, y, z, TDS, B, betaij, U, Uij

GOAL's, quant analysis

GOAL, GEWICHT

BGMN-specific functions

PHASE, sk, B1, k2, ANISO, GEWICHT(h,k,l), GrainSize, TDS,

Advanced feature: subphases

RefMult, iref, GEWICHT[i], B1[i], B2[i], k1[i], k2[i], k3[i], DELTAask, DELTAzweiTheta, LeBail, FPARAM

Advanced feature: Structure amplitudes etc

F, Finv, H

Advanced feature: User calculated structure amplitudes

FMult, F[i], phi[i], Finv[i], phiinv[i]

Advanced feature: molecules

set, setgitter, cross, diffvec, normvec, skalpro, distance, angle, cpXYZ, T, D, X, Y, Z, WW, WWalt, Straf, Theory, Bondings, BondLevel

Micro absorption correction according to Brindley

my, my[i]

X ray density

density, density[i]

Variables in the task description SAV file for computation of standard profiles

NTHREADS, VERZERR, TubeTails, R, FocusH, FocusW, HSlitR, HSlitW, RoundSlitR, RoundSlitD PColl, PCollA, VSlitR, VSlitH, SampID, SampIw, SampIH, DeltaOmega, SCollA, SSlitW, SSlitR, SColl, SCollA, DetW, DetH, DetArrayW, MonR, MonH, EPSG, zweiTheta[i], GSUM, TSlitR, TSlitH, FocusS, FocusA, GEOMETRY, WMIN, WMAX, WSTEP, GEQ, D, T, STANDARDPAR, VAL,



Creating Structure Files

BGMN_manual_2005.pdf

Manual

Rietveld Analysis Program

BGMN[®]

Jörg Bergmann, Dresden, Germany

Rietveld Analysis Program BGMN	
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From CIF to BGMN Structure Files (I)

CIF File

```
_chemical_name_mineral 'Anatase'  
  
_symmetry_space_group_name_H-M 'I 41/a m d'  
  
_cell_length_a 3.7845  
_cell_length_c 9.5143  
  
Ti  
0.00000  
0.00000  
0.00000  
0.0052 0.0000 0.0000  
0.0052 0.0000 0.0070  
  
O  
0.00000  
0.00000  
0.20806  
0.0117 0.0000 0.0000  
0.0027 0.0000 0.0072
```

OK

Not available

Wrong format

Wrong unit

OK (sort of...)

Not available

OK

Wrong format

OK (sort of...)

Not available

OK

Wrong format

BGMN STR File

PHASE=Anatase

SpacegroupNo=141

HermannMauguin=I4_1/a2/m2/d

A=0.3784

C=0.9514

E=Ti+4

Wyckoff=a

x=0.00000

y=0.00000

z=0.00000

TDS=0.0045795

E=O-2

Wyckoff=e

x=0.00000

y=0.00000

z=0.20806

TDS=0.0056849

From CIF to BGMN Structure Files (II)

Category	Solution
OK →	Can be mapped from CIF to STR
OK (sort of...) →	Can be mapped / or fixed easily
Wrong format / unit →	Need conversion
Not available →	PROBLEM!!!

PROBLEM!!!

=

gather more information from other sources
and fix manually

From CIF to BGMN Structure Files (III)

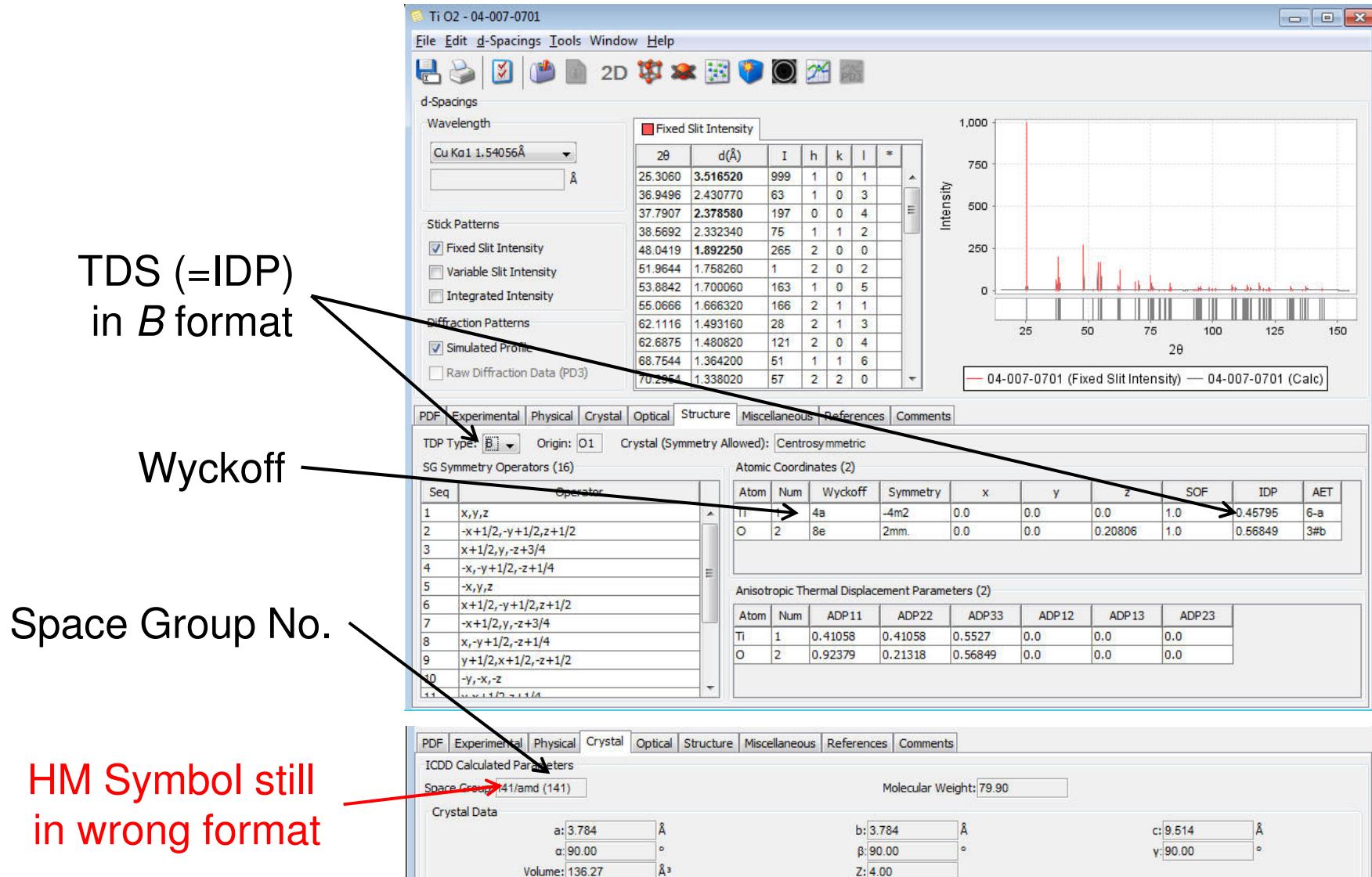
PROBLEM!!!

=

gather more information from other sources
and fix manually

- Find a better, more complete, CIF file
(e.g. from another structure refinement)
- Search for the same structure in a different database
- Find missing information elsewhere and add to CIF file
(original publication, different database, different publication)

PDF-4+ Database



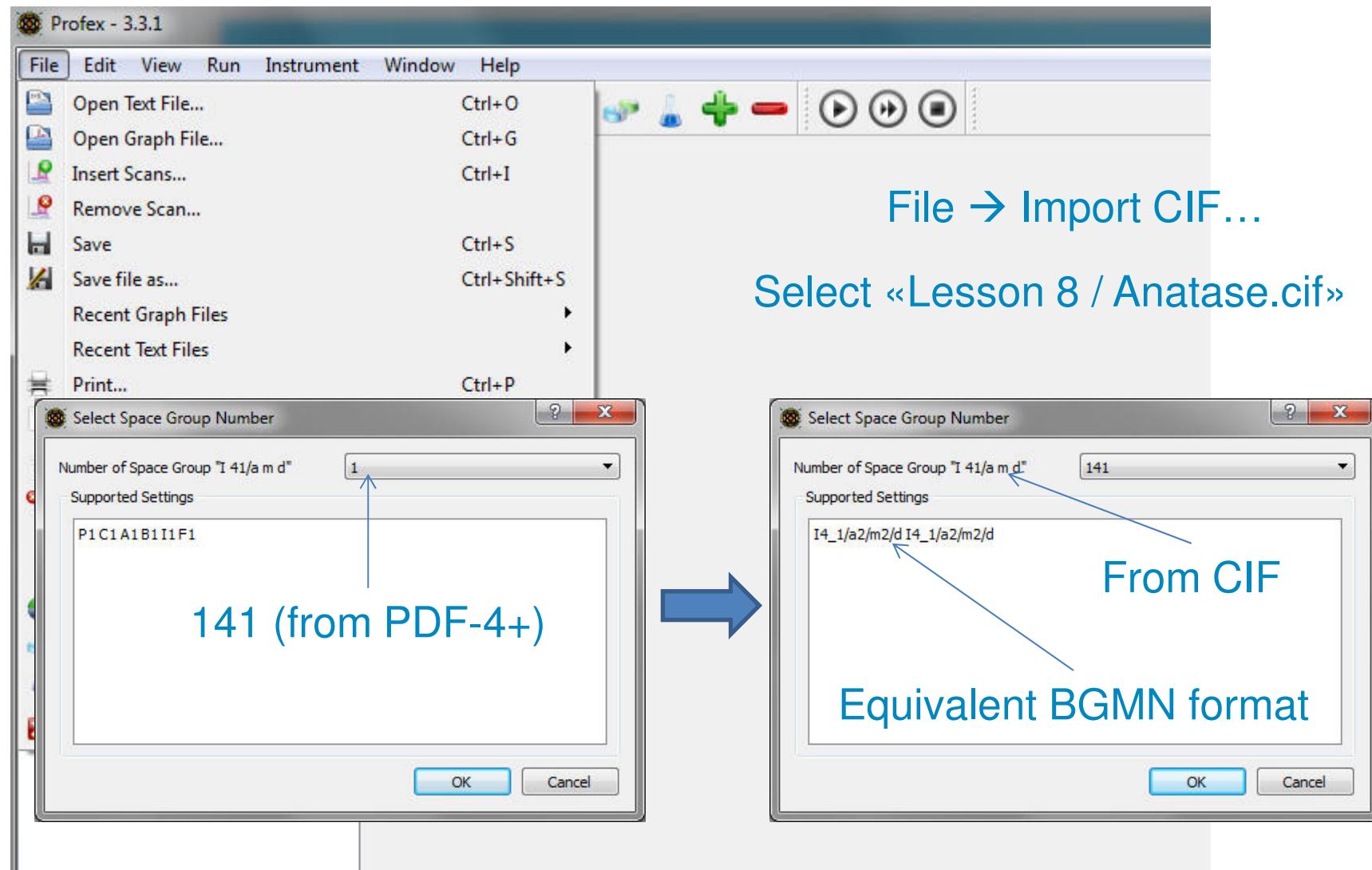
From CIF to BGMN Structure Files



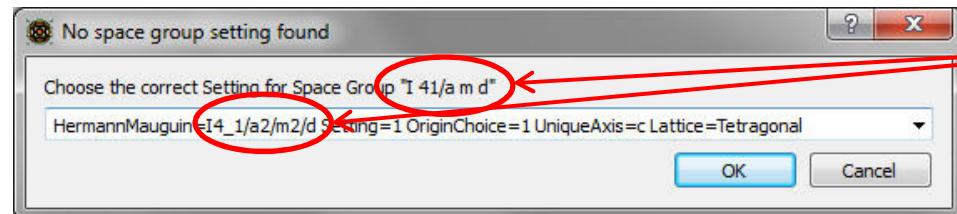
Profex CIF Import



CIF to STR with Profex (I)

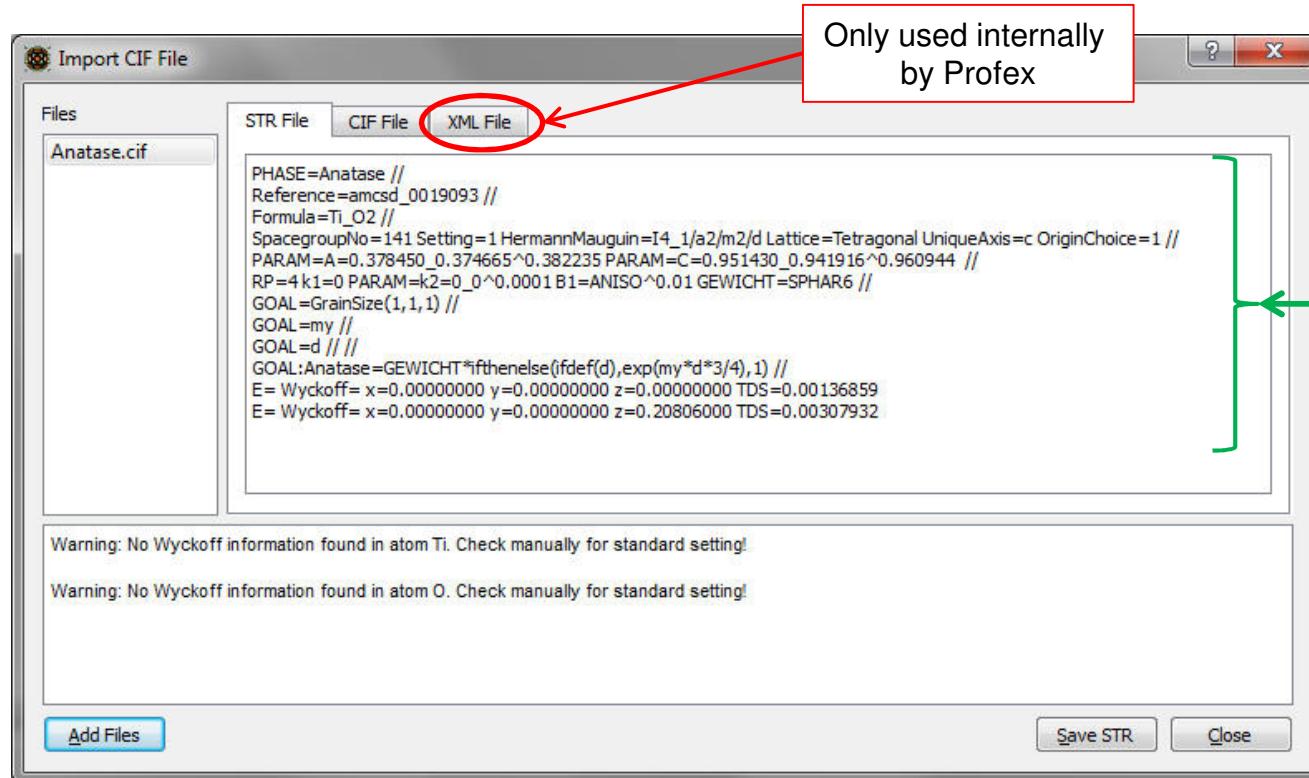


CIF to STR with Profex (II)



Normally: Match HM Symbols

Here: No difference, just click OK



CIF to STR with Profex (III)

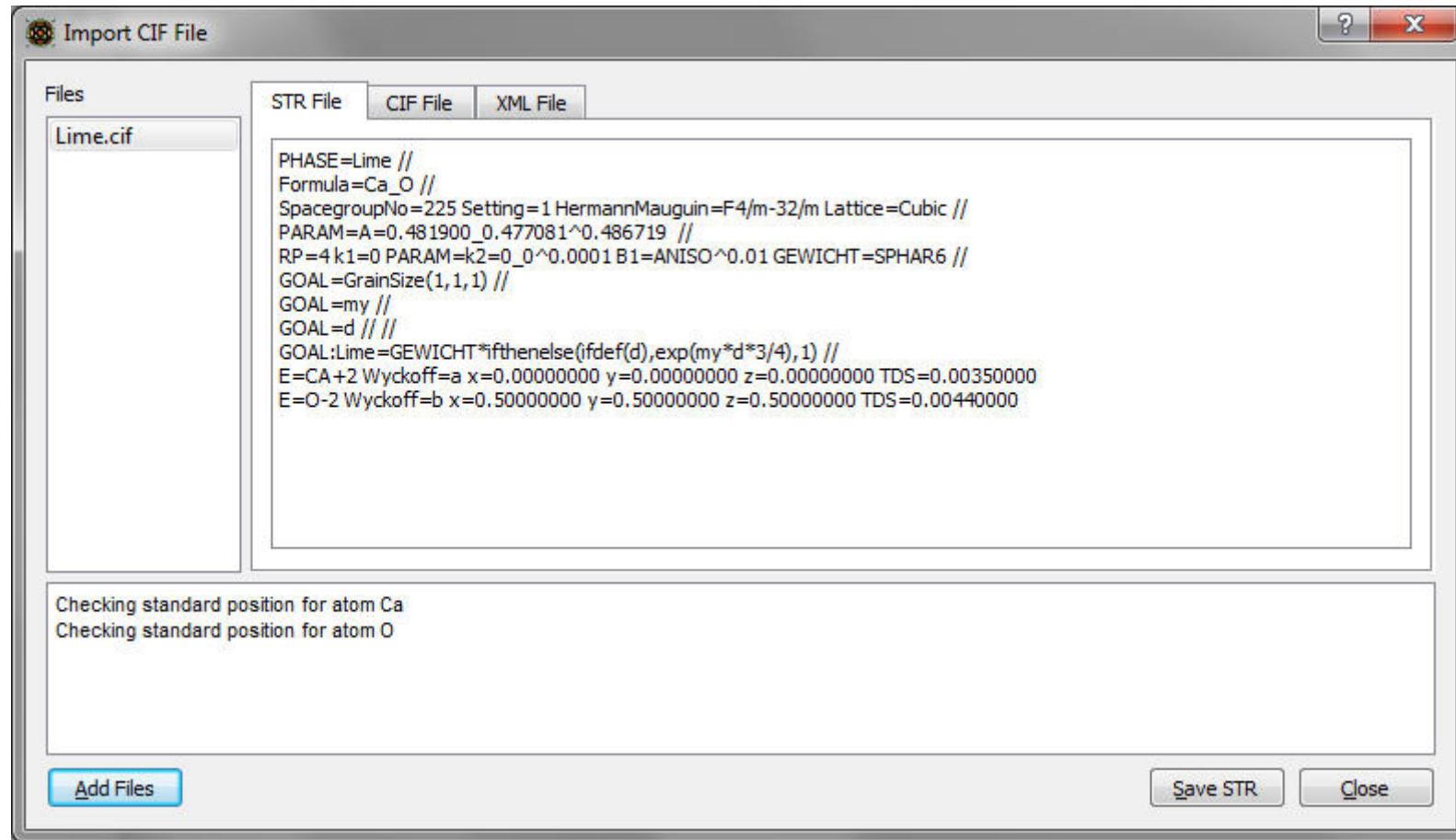
```
PHASE=Anatase //
Reference=amcsd_0019093 //
Formula=Ti_O2 //
SpacegroupNo=141 Setting=1 HermannMauguin=I4_1/a2/m2/d Lattice=Tetragonal
UniqueAxis=c OriginChoice=1 //
PARAM=A=0.378450_0.374665^0.382235 PARAM=C=0.951430_0.941916^0.960944 //
RP=4 k1=0 PARAM=k2=0_0^0.0001 B1=ANISO^0.01 GEWICHT=SPHAR6 //
GOAL=GrainSize(1,1,1) //
GOAL=my //
GOAL=d // //
GOAL:Anatase=GEWICHT*ifthenelse(ifdef(d),exp(my*d*3/4),1) //
E=TI+4 Wyckoff=a x=0.00000000 y=0.00000000 z=0.00000000 TDS=0.00136859
E=O-2 Wyckoff=e x=0.00000000 y=0.00000000 z=0.20806000 TDS=0.00307932
```

- Add missing information (Element Types and Wyckoff Sequences)
- «Save STR»
- Save file in Structure Database Directory:
...\\Profex-BGMN-Bundle-3.3.1\\Profex\\Structures

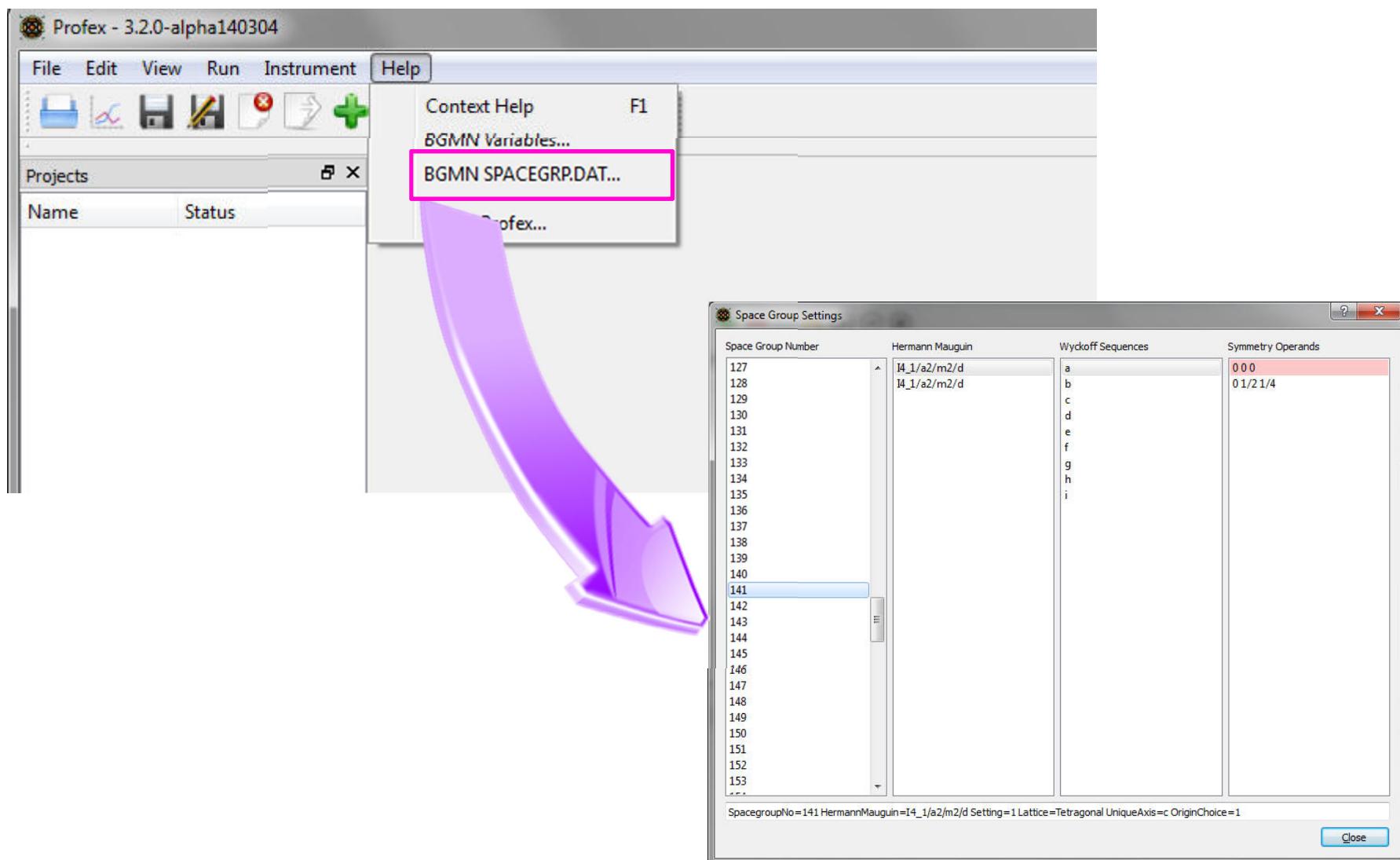
CIF to STR with Profex (IV)

With correct and complete CIF files: No / minimal user input required

Try: «Lesson 8 / Lime.cif»



SPACEGRP.DAT Browser

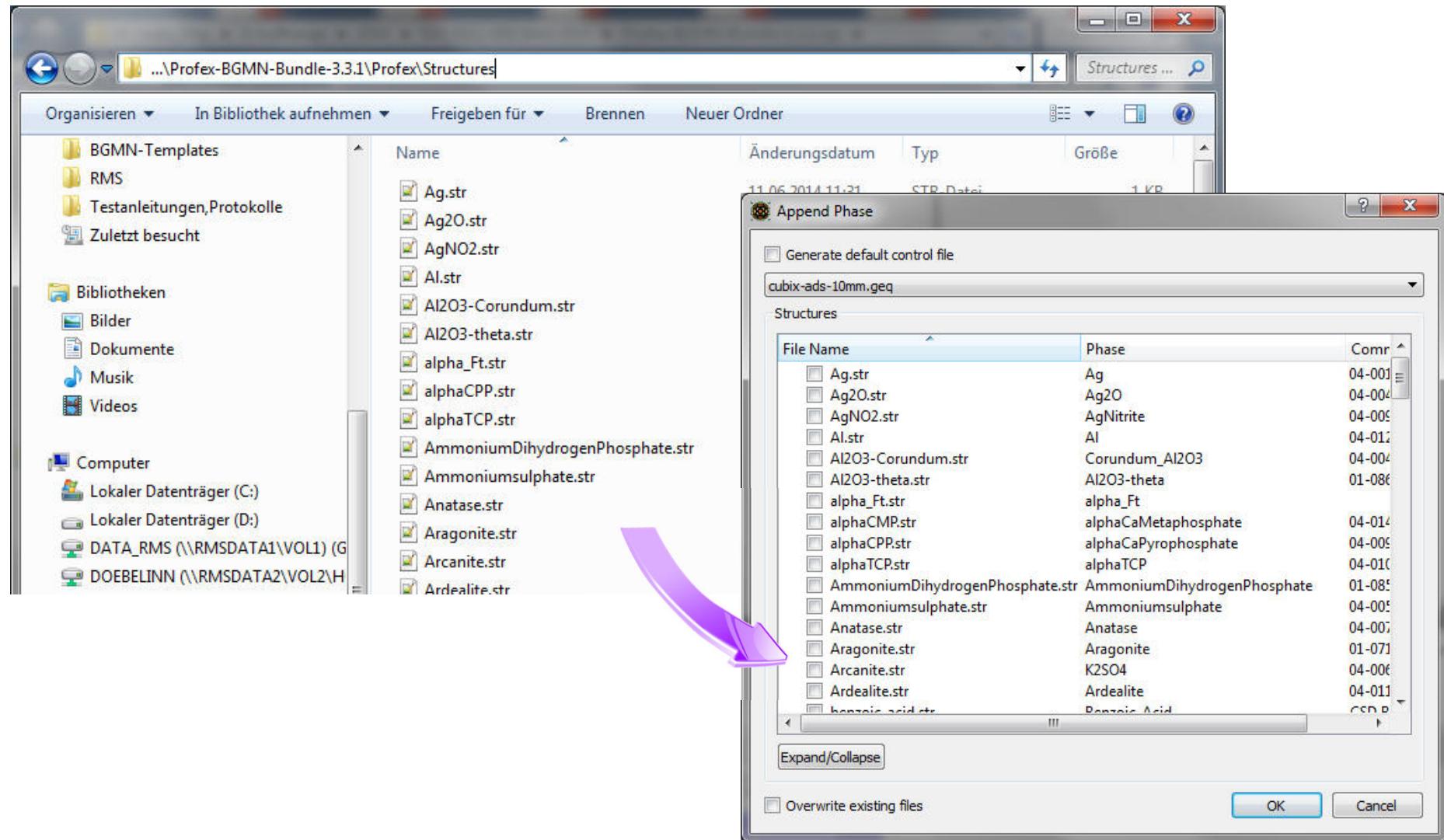


CIF to STR with Profex (VII)

Summary:

- Crystal structure data available from various databases (free and commercial)
- Provided in CIF format
- Usually additional information required
- Semi-automatic conversion CIF → STR with Profex
- Manual fixing / verification required

Profex Structure and Device Database



Profex Structure and Device Database

