The Crystallographic Information File (CIF) Description and Usage

Ton Spek, Bijvoet Center for Biomolecular Research Utrecht University Leiden. 27-Jan.-2009.



Overview of this Lecture

- An Overview of the Status of Single Crystal X-Ray Structure Determination
- The What, Why and How of CIF
- CIF Usage and Applications

A Single Crystal X-Ray Study Involves

- A Structure Query (What did I make?)
- Needed: A Single Crystal (0.2 mm)
- Collection of X-Ray Diffraction Data
- Solution of the Phase Problem (To get a Preliminary Model)
- Structure Model Parameter Refinement
- Interpretation of the Result/Geometry Analysis
- Validation of the Analysis
- Report and (Co)Publication

Crystal Requirements

- Preferably a single crystal (Sharp Extinctions under Polarized Light)
- Block rather than slim long needle
- Fresh from mother liquor
- Unstable crystals covered with inert oil Data collection under cold N₂ stream.
- Twins and split crystals possible but best avoided when possible.







Short History of Data Collection

- 1912 von Laue et al. Experiment X-ray Film
- 1913 Bragg Diffractometer + Detector
- Tot ~ 1965 Film (Weissenberg, etc.)
- ~1960-1995 Serial Detector Diffractometer
 ~ 50 datasets / year
- 1995 present Image plate/ CCD Detectors a few hours Collection Time for a Routine Structure
- ~2007 Digital Detectors (AXIOM, Pilatus etc.) new options: shutterless, low noise etc.





Central Formulae

- Diffraction Spots: $2d_{hkl} \sin \theta_{hkl} = n\lambda$
- Electron Density Map (3D Fourier Map)
- $\rho_{x,y,z} = 1/V \Sigma_{hkl} F_{hkl} \exp\{-2\pi i(hx + ky + lz)\}$
- Structure Factor (Model)
- $$\begin{split} F_{hkl}(calc) &= \Sigma_j \, f_j \, T_{j,hkl} \, exp\{2\pi i (hx_j + ky_j + lz_j)\} \\ \bullet \ Least \ Squares \ Model \ Refinement \end{split}$$
- $\begin{array}{l} \text{Minimize: } \boldsymbol{\Sigma}_{hkl} \left[\mathbf{w}_{hkl} (|\mathbf{F}_{hkl}(\mathbf{obs})|^2 |\mathbf{F}_{hkl}(\mathbf{calc})|^2) \right]^2 \\ \text{Gradients of the set of th$
- Convergence Criteria: R1, wR2, S

Solution of the Phase Problem

- Early Method: Trial & Error (Salts such as NaCl, Silicates etc.)
- Patterson Methods (Heavy Atom)
- Direct Methods (SHELXS, DIRDIF, SIR)
- New: Charge Flipping (Ab-initio)
- Phase Problem Solved! Given reasonable data.







Interpretation in Terms of Atoms

- Position of highest density => Position x,y,z
- Deviation of the density shape from the ideal atomic electron density => Thermal motion parameters:
- Isotropic: U(iso) or
- Anisotropic: U¹¹,U¹²,U¹³,U²²,U²³,U³³
 (Displacement Parameters) => ORTEP
 Note: ORTEP does NOT represent the electron distribution.

Interpretation in Terms of Bonds

- Bonds between atoms of type A and B are assigned on the basis of atomic covalent radii with: d < R(A) + R(B) + 0.4
- 'Crystallographic Bonds' are not necessarily Chemical Bonds.
- Van der Waals Radii are used to detect isolated molecular species or short contacts.

Display Options

Ball-and-Stick

Simpel but may hide problems with a structure.

• ORTEP

Often preferred because it visualizes most model parameters and possible problems.

• **CPK** Spacefilling PLOT illustrating the shape etc







Parameter Model Refinement

- Translate the 3D electron density in terms of position and temperature parameters
- Non- Lineair Least Squares Method Min: $\Sigma_{hkl} \; [w_{hkl} (|F_{hkl} (obs)|^2 - |F_{hkl} (calc)|^2)]^2$
- Time consuming: disorder, weak data

Analysis

Interpretation (geometry, intermolecular interactions etc.)

- Structure Validation
- (Co)Publikation
- Results to Cambridge Crystallographic Database CSD (~500000)

Computing in the Past

- University Mainframe
- Mainly Batch Mode (Long Waits)
- Storage on Magnetic Tape
- Piles of Lineprinter Output
- Primitive Computer Graphics
- No Internet









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Computing Today

- Personal Workstation (MS-Windows or Linux)
- No queues for Computing Facilities
- Good graphics & Hardcopy
- Unlimited Disc Storage
- Internet (exchange, information)
- Automation



Problems Around 1990

- Multiple Data Storage Types
- No Standard Computer Readable Format
- Data Entry of Published data via Retyping.
- No easy numerical checking for referees etc.
- CSD Database Archival by Retyping from the paper
- Multiple typo's in Published Data

Solution

- CIF-Standard Proposal
- S.R. Hall, F.H. Allen, I.D. Brown (1991). Acta Cryst. A47, 655-685.
- Pioneered and Adopted by the International Union for Crystallography
- Adopted by the author of the most used software package SHELXL (G.M.Sheldrick)

What is CIF ?

From: http://ww1.iucr.org/cif/index.html

CIF

The acronym CIF is used both for the Crystallographic Information File, the data exchange standard file format of Hall, Allen & Brown (1991) (see Documentation), and for the Crystallographic Information Framework, a broader system of exchange protocols based on data dictionaries and relational rules expressible in different machine-readable manifestations, including, but not restricted to, Crystallographic Information File and XML.

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A selection of papers and articles.
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CIE Die

 Provder dictionary (pdCIF) [ASCII [ITTML [PDF] More information]
 Modulated and commute structure

Practical Approach

- We ignore here the scary details that are not relevant in the current context
- We will Discuss the File structure
- We will look at its relevance for publication
- We will discuss software to edit and check the CIF file
- We will look at software that uses CIF as Input.

File Structure

- Both Computer and Human Readable Ascii encoded file
- Free Format
- Mostly 80 colums wide
- Parsable in units
- Data Order Flexible
- Dataname and Value associations

Constructs

• data_*name*

- where *name* the choosen identifier of the data
- Data associations e.g. _ cell_length_a 16.6392(2)
- Repetition (loop)
- loop_ ___**symmetry_equiv_pos_as_xyz**
 - · x, y, z'
 - '-x, y+1/2, -z'



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Note on Editing the CIF

- The Idea of editing the CIF is to add missing information to the CIF.
- Some Acta Cryst. authors have been found to polish away less nice numerical values. This leaves traces and is generally detected by the validation software and not good for the career of the culprit...

CIF Applications

- Data Archival
- Deposition to the CSD (=> CSD number)
- Supplementary Material for Publication
- Input for Geometry and Graphics Software e,g. Mercury (from CCDC) and PLATON
- Standard Format for publications (Structure Communications) in Acta Cryst. Sections C & E.
- Structure Validation

Reflection CIF (FCF)
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Structure Validation

- Pioneered by the IUCr
- Currently most journals have implemented a validation scheme.
- Papers:
- A.L.Spek (2003). J. Appl. Cryst. 36, 7-13.
- A.L.Spek (2009). Acta Cryst. D65, 148-155.

Why Crystal Structure Validation

- · The explosion of structure determinations
- An analyses of the nearly 500000 structures in the CSD learns that a significant number are in error
- Many analyses are done today by non-specialists
- Limited number of experts to detect pitfalls
- Validation provides a list of issues that need special attention of the analyst, specialist and referee.
- · Validation sets quality standards.

VALIDATION QUESTIONS

Single crystal validation addresses three simple but important questions:

- 1 Is the reported information complete?
- 2 What is the quality of the analysis?
- 3 Is the Structure Correct?

How is Validation Implemented

- Computer readable structure analysis results in CIF format (Syd Hall & George Sheldrick)
- A file (Check.def) defines the issues that are tested with levels of severity and associated explanation and advise.
- The tests are executed by the program PLATON
- The tests can be executed both in-house or through the WEB-based IUCr CHECKCIF server.

ALERT LEVELS

CheckCif Report in terms of a list of ALERTS

- ALERT A Serious Problem
- ALERT B Potentially Serious Problem
- ALERT C Check & Explain
- ALERT G Verify or Take Notice

ALERT TYPES

- 1 CIF Construction/Syntax errors, Missing or Inconsistent Data.
- 2 Indicators that the Structure Model may be Wrong or Deficient.
- 3 Indicators that the quality of the results may be low.
- 4 Cosmetic Improvements, Queries and Suggestions.

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Bood precision:	C-C + 0.0157 A	Wavelength-0.71073	
Cell: a-7.63	396(15) b=27.725(6) c-	-12.051(2)	
alpha	-90 beta-98.80(3) ga	4883-90	
Temperature: 153 K			
	Calculated	Reported	
Volume	2520.5(9)	2520.5(9)	
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Hall group	-P 2yn	-P 2yn	
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Which Key Issues are Addressed

- Missed symmetry ("being Marshed")
- Wrong chemistry (Misassigned atom types)
- Too many, too few or misplaced H-atoms
- · Missed solvent accessible voids in the structure
- Missed Twinning
- Absolute structure
- Data quality and completeness

FCF-VALIDATION

Forthcoming:

Automatic twinning detection as part of the IUCr CheckCif procedure

- Detection of ignored twinning
- Detection of Applied Twinning Correction without being reported
 - (Already available via PLATON/Check)

Examples

• Following are some examples of the type of problems addressed.

Enthusiastic Last Paragraph of C&E-News

... The Northwestern chemists are now exploring the reactivity of C5Me5+. "You can sit down and write lots of interesting reactions on paper," Lambert says, and it will be interesting to see if the molecule reacts as expected. But Considering this cation's track record, it might be safer to expect more surprises. [!!]

NOT SO HOT AFTER ALL !!

Editors Note in the next issue of Angewandte Chemie

CORRIGENDUM

Note from the Editors: unfortunately the results reported in the communication "The Stable Pentamethylcyclopentadienyl Cation" by Joseph B. Lambert et al. in issue 82002 (pp. 1429–1431) must be corrected. Guy Bertrand et al. quickly discovered that not the pentamethylcyclopentadienyl cation but the pentamethylcyclopentenyl cation was prepared and characterized (the corresponding communication will be published in issue 13, and will appear earlier on the Angewandte Chemie homepage).

Concluding Remarks

- The CIF standard makes it possible to easily do follow-up calculations for published structures
- The available information is more complete
- <u>http://www.cryst.chem.uu.nl</u> for more information