

Championing data standards in chemical crystallography with CIF

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IUCr Committee on Data (CommDat) member



**universität
wien**



IUCr
International Union
of Crystallography



International Union of Crystallography: a scientific union

The **International Union of Crystallography (IUCr)** is a non-profit scientific union serving the worldwide interests of crystallographers and other scientists employing crystallographic methods

- 55 National Committees make up the General Assembly
- The scientific work is conducted through Commissions and Committees
- The **Committee on Data (CommDat)** work with the IUCr's Commissions, including the Commission on Journals, having a coordinating and advisory role regarding data
- CommDat exist alongside, and have a formal relationship with, the Committee for the Maintenance of the CIF Standard (COMCIFS)
- The **Committee for the Maintenance of the CIF Standard (COMCIFS)** oversees the development of the CIF and reports to the Executive Committee of the IUCr
- Drafts of new definitions and official dictionaries are submitted to COMCIFS for technical validation and ultimately for approval

Paul Ewald of the Polytechnic Institute of Brooklyn, President of the International Union of Crystallography (a scientific union, not a labor union)

Elizabeth A. Wood, *Crystals and Light: An Introduction to Optical Crystallography*, D. Van Nostrand, Princeton 1964.



IUCr
International Union
of Crystallography



Simon Coles
CommDat chair



James Hester
COMCIFS chair

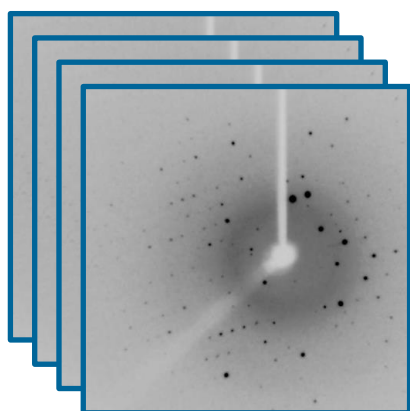




Flavors and data cycle of crystallographic data

Crystallographic data lifecycle

- Raw data images collected in a diffraction experiment
- Processed data (unit-cell parameters and indexed Bragg reflections with associated integrated intensities)
- Crystal structure (symmetry, unit-cell parameters, atomic coordinates, ADPs)



Raw (primary) data

loop_	refln_index_h	refln_index_k	refln_index_l	refln_F_squared_calc	refln_F_squared_meas	refln_F_squared_sigma	refln_observed_status
2	0	0	25185.30	25637.90	369.48	o	
4	0	0	1787.58	1693.17	29.40	o	
6	0	0	59.02	71.69	4.22	o	
8	0	0	27.82	14.56	3.68	o	
10	0	0	483.46	450.47	23.58	o	
1	1	0	7357.92	7392.33	96.02	o	
2	1	0	27.32	30.41	1.75	o	
3	1	0	1335.00	1608.41	31.10	o	
4	1	0	9037.50	8887.28	88.78	o	
5	1	0	4968.33	5067.08	84.21	o	
6	1	0	29.52	35.79	2.56	o	
7	1	0	4.53	6.29	2.14	o	
8	1	0	1.53	3.02	2.39	o	
9	1	0	62.03	58.07	4.39	o	

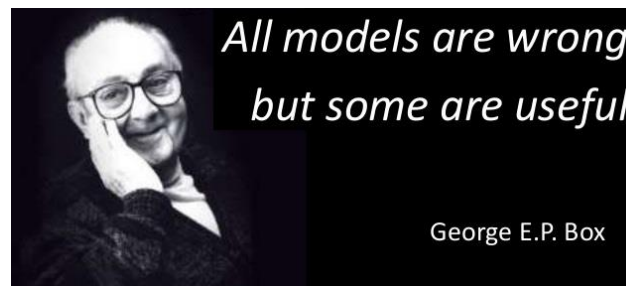
Processed data

```
loop_
_atom_site_label
_atom_site_type_symbol
_atom_site_fract_x
_atom_site_fract_y
_atom_site_fract_z
_atom_site_U_iso_or_equiv
_atom_site_adp_type
_atom_site_occupancy
_atom_site_symmetry_multiplicity
_atom_site_calc_flag
_atom_site_refinement_flags
_atom_site_disorder_assembly
_atom_site_disorder_group
C1 C 0.10650(11) 0.6028(2) 0.15602(9) 0.0265(4) Uani 1 1 d . . .
C2 C 0.07667(12) 0.6156(2) 0.09181(10) 0.0346(5) Uani 1 1 d . . .
H2 H 0.0274 0.6713 0.0837 0.042 Uiso 1 1 calc R . . .
C3 C 0.11916(13) 0.5466(3) 0.03925(10) 0.0414(5) Uani 1 1 d . . .
H3 H 0.0986 0.5546 -0.0049 0.050 Uiso 1 1 calc R . . .
C4 C 0.19043(12) 0.4672(3) 0.05067(10) 0.0390(5) Uani 1 1 d . . .
H4 H 0.2188 0.4203 0.0144 0.047 Uiso 1 1 calc R . . .
C5 C 0.22184(11) 0.4543(2) 0.11478(10) 0.0326(5) Uani 1 1 d . . .
H5 H 0.2715 0.3992 0.1223 0.039 Uiso 1 1 calc R . . .
C6 C 0.18050(12) 0.5223(3) 0.16750(9) 0.0253(4) Uani 1 1 d . . .
```

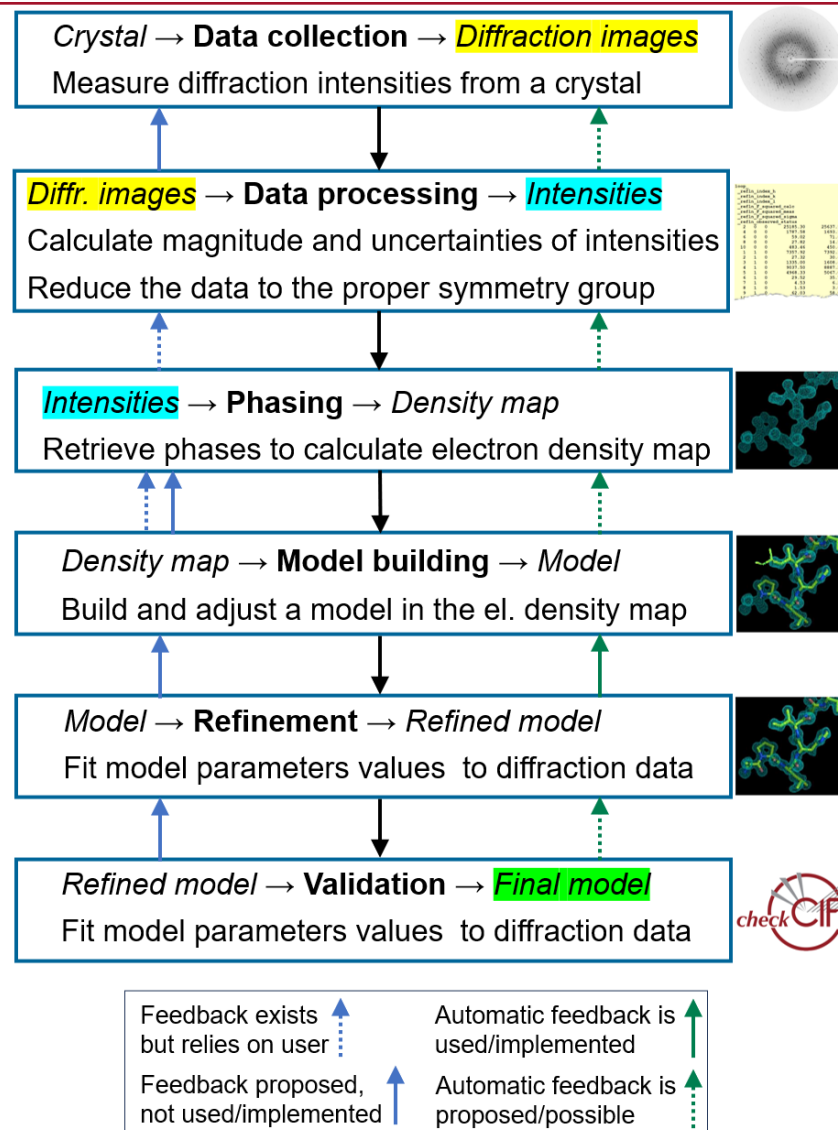
Derived (structural) data

Crystal structures are models!

- The actual raw data in X-ray crystallography are diffraction data leading to electron density
- Atom positions that make up a crystal structure are a model that attempts to explain the raw data in as complete a fashion as possible



George E.P. Box

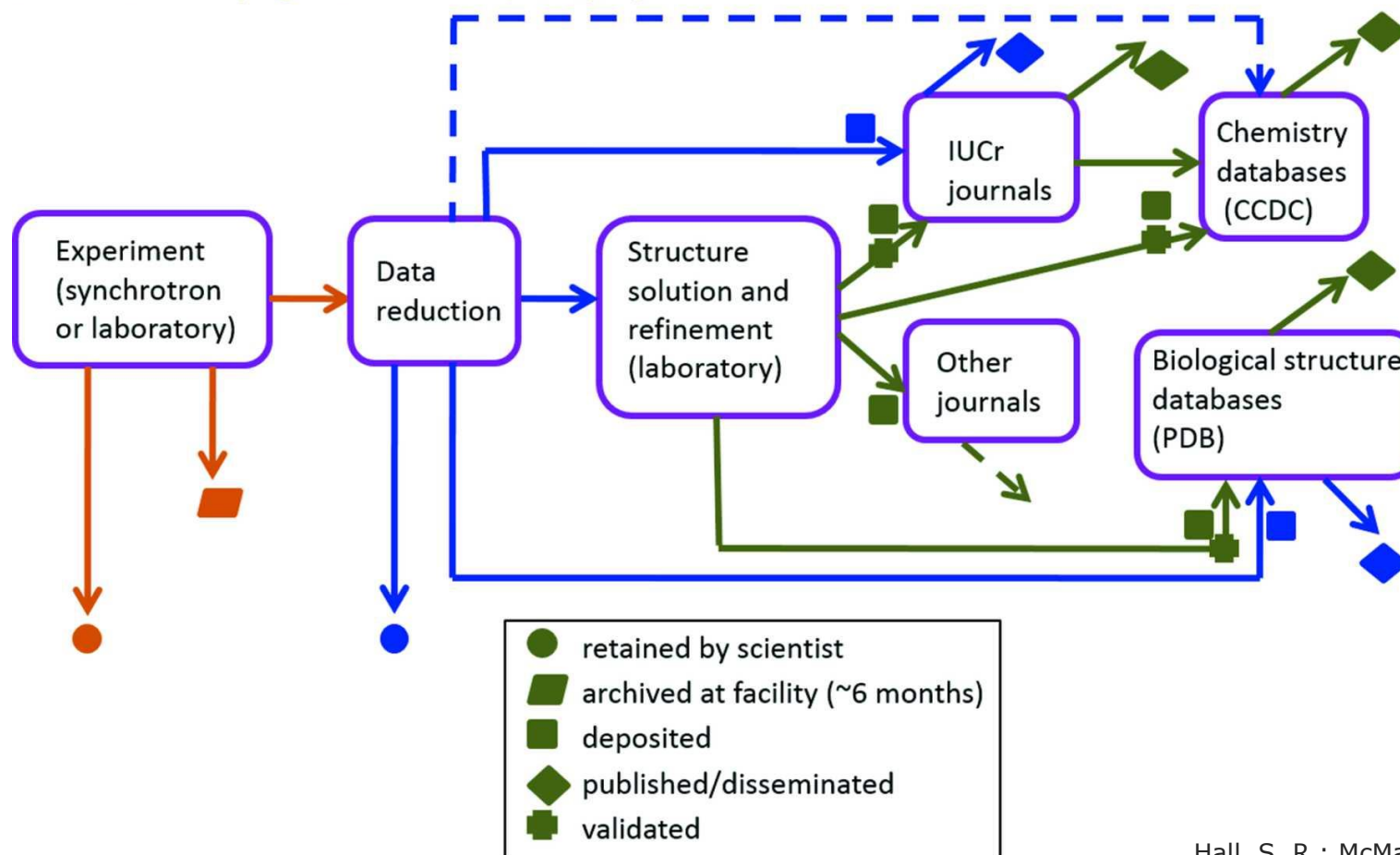


Coherent information flow in crystallography

Raw experimental data (e.g. diffraction images)

Reduced/processed data (e.g. structure factors)

Derived data (e.g. coordinates, a.d.p.s)

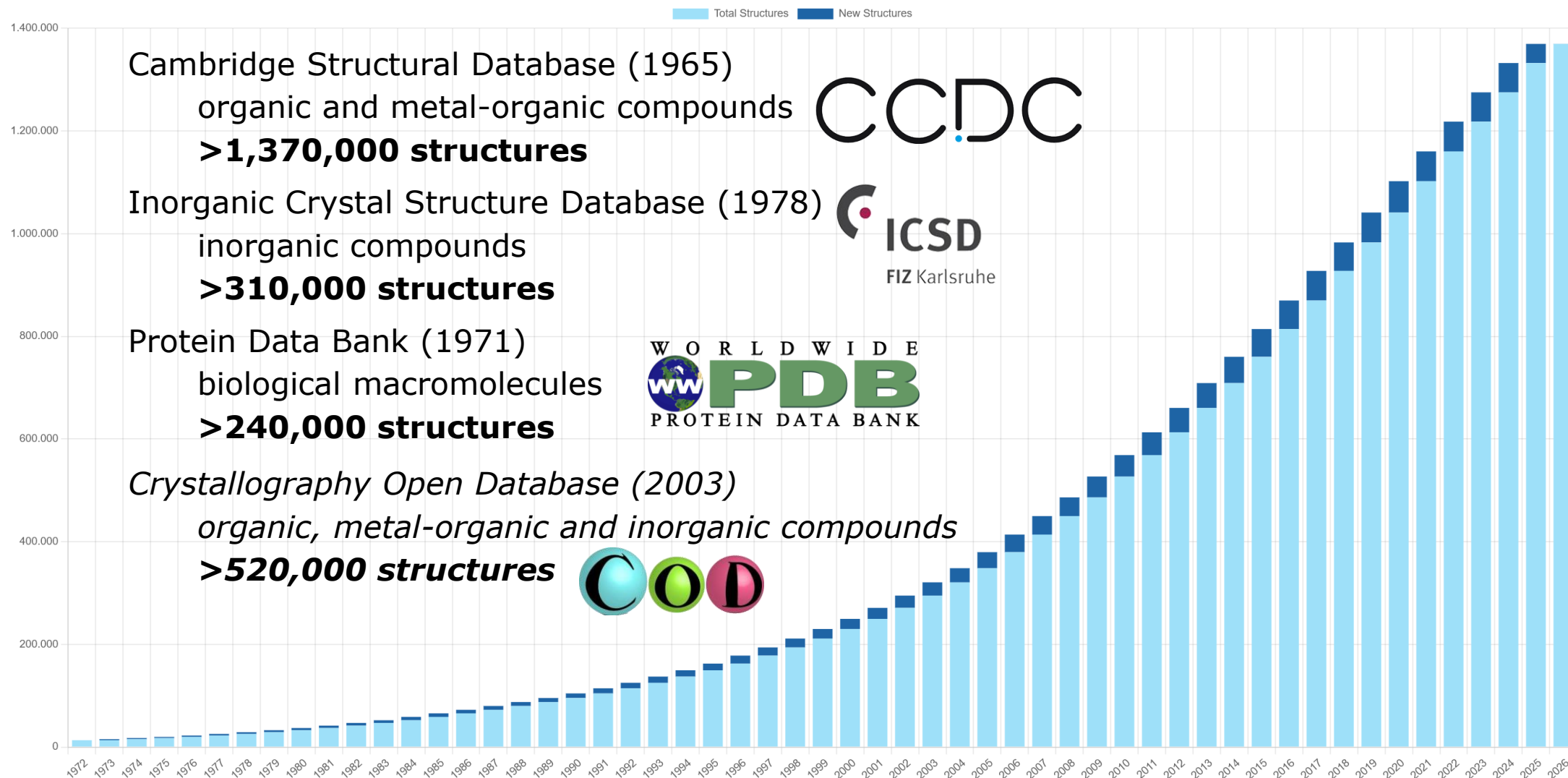


Hall, S. R.; McMahon, B. *Data Sci. J.* **2016**, *15*, 3.



Databases in the data deluge era

CSD Growth





In 1991...



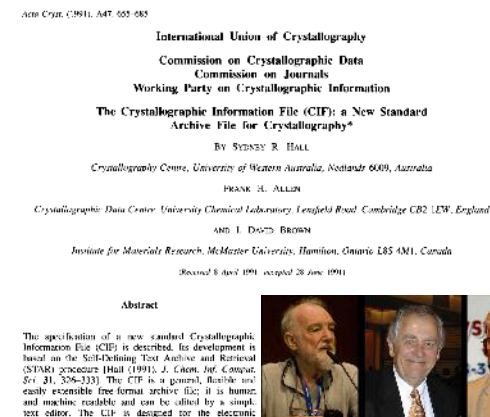
...USSR passes into history



...Operation Desert Storm began



...World Wide Web publicly debuts
(HTML, XML and Java were yet to come!)



...IUCr introduced the Crystallographic Information File



What is CIF and how is it used?

Crystallographic Information Framework (CIF) is a data exchange format for crystallographic and related structural science data

- The CIF format and extensible CIF dictionaries are **defined, adopted and curated by the IUCr**
- Until 2005 the acronym stood for 'Crystallographic Information File', but the name was modified in recognition of its application across crystallography and all related structural science fields
- CIF offers domain-specific ontology (collection of data identifiers, attributes and relationships)
- Simple syntax well suited to archive and exchange purposes (human- and machine-readable)
- Adopted by major databases** (CSD, ICSD, COD, RRUFF, Mindat) and **required as ESI in most journals**
- Can also be adopted for the **hosting structure factors / reflection intensity data**
- imgCIF/CBF** can handle binary data (including raw data images) and is isomorphous to CIF
- Canonical information on CIF contained in International Tables Vol. G

```
loop_
  _atom_site_label
  _atom_site_type_symbol
  _atom_site_fract_x
  _atom_site_fract_y
  _atom_site_fract_z
  _atom_site_U_iso_or_equiv
  _atom_site_adp_type
  _atom_site_occupancy
  _atom_site_symmetry_multiplicity
  _atom_site_calc_flag
  _atom_site_refinement_flags
  _atom_site_disorder_assembly
  _atom_site_disorder_group
C1 C 0.10650(11) 0.6028(2) 0.1560(2) 0.0265(4) Uani 1 1 d . . .
C2 C 0.07667(12) 0.6156(2) 0.09181(10) 0.0346(5) Uani 1 1 d . . .
H2 H 0.0274 0.6713 0.0837 0.042 Uiso 1 1 calc R . .
C3 C 0.11916(13) 0.5466(3) 0.03925(10) 0.0414(5) Uani 1 1 d . . .
H3 H 0.0986 0.5546 -0.0049 0.050 Uiso 1 1 calc R . .
C4 C 0.19043(12) 0.4672(3) 0.05067(10) 0.0390(5) Uani 1 1 d . . .
H4 H 0.2188 0.4203 0.0144 0.047 Uiso 1 1 calc R . .
C5 C 0.22184(11) 0.4543(2) 0.11478(10) 0.0326(5) Uani 1 1 d . . .
H5 H 0.2715 0.3992 0.1223 0.039 Uiso 1 1 calc R . .
C6 C 0.1802(10) 0.5223(2) 0.16750(9) 0.0253(4) Uani 1 1 d . . .
```

Acta Cryst. (1991). A47, 655–685

International Union of Crystallography
Commission on Crystallographic Data
Commission on Journals
Working Party on Crystallographic Information

The Crystallographic Information File (CIF): a New Standard
Archive File for Crystallography*

By SYDNEY R. HALL,

Crystallography Centre, University of Western Australia, Nedlands 6009, Australia

FRANK H. ALLEN

Crystallographic Data Centre, University Chemical Laboratory, Lensfield Road, Cambridge CB2 1EW, England

AND I. DAVID BROWN

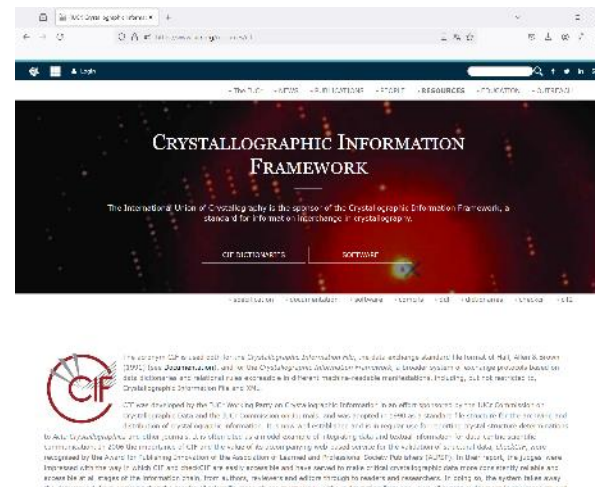
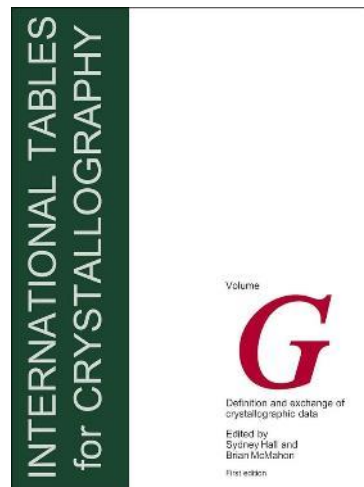
Institute for Materials Research, McMaster University, Hamilton, Ontario L8S 4M1, Canada

(Received 8 April 1991; accepted 28 June 1991)

Abstract

The specification of a new standard Crystallographic Information File (CIF) is described. Its development is based on the Self-Defining Text Archive and Retrieval (STAR) procedure (Hall (1991), *J. Chem. Inf. Comput. Sci.* 31, 326–333). The CIF is a general, flexible and easily extensible free-format archive file; it is human and machine readable and can be edited by a simple text editor. The CIF is designed for the electronic

There is an increasing need in many branches of science for a uniform but flexible method of archiving and exchanging data in electronic form. Rapid advances in computer technology, coupled with the expansion of local, national and international networks, have fuelled the need for such a facility. The variety and relative inflexibility of existing data exchange formats have inhibited their effective use. This is true even in fields where the basic data requirements are well defined. Problems of data ex-



```
loop_
  _refln_index_h
  _refln_index_k
  _refln_index_l
  _refln_F_squared_calc
  _refln_F_squared_meas
  _refln_F_squared_sigma
  _refln_observed_status
2 0 0 25185.30 25637.90 369.48 o
4 0 0 1787.58 1693.17 29.40 o
6 0 0 59.02 71.69 4.22 o
8 0 0 27.82 14.56 3.68 o
10 0 0 483.46 450.47 23.58 o
1 1 0 7357.92 7392.33 96.02 o
2 1 0 27.32 30.41 1.75 o
3 1 0 1335.00 1608.41 31.10 o
4 1 0 9037.50 8887.28 88.78 o
5 1 0 4968.33 5067.08 84.21 o
6 1 0 29.52 35.79 2.56 o
7 1 0 4.53 6.29 2.14 o
8 1 0 1.53 3.02 2.39 o
9 1 0 62.03 58.07 4.39 o
```



Wide applicability of STAR

CIF is based on the Self-Defining Text Archive and Retrieval (STAR) procedure

- STAR language has the capacity to handle the more complex data representations found in various disciplines such as imaging, quantum chemistry, botany, etc.
- Non-CIF STAR applications include
 - NMR imaging field (Biological Magnetic Resonance Data Bank)
 - Molecular Information File (MIF)
 - QCHEM ontology for quantum chemistry
 - Botanical ontologies (Florabase, Western Australian Herbarium), e.g. plant naming hierarchies lend themselves to the use of 'definition' methods to connect these hierarchies

326

J. Chem. Inf. Comput. Sci. **1991**, *31*, 326–333

The STAR File: A New Format for Electronic Data Transfer and Archiving

SYDNEY R. HALL

Crystallography Centre, University of Western Australia, Nedlands 6009, Australia

Received October 2, 1990

A new type of format is proposed for the computer archiving and electronic transmission of text and numerical data. The Self-defining Text Archive and Retrieval (STAR) File uses standard ASCII text to specify both the data structure and the information. The syntax of this file is simple, and it may be easily interpreted visually or by computer. The STAR format is the basis for the Crystallographic Information File (CIF), which has been adopted by the International Union of Crystallography for the submission of data and text to crystallographic journals and data bases.



Sydney R. Hall



Extensibility of CIF: dictionaries

CIF can host not only structural data

- Formally, the CIF approach makes **no distinction between 'data' and 'metadata'** and is adaptable and extensible to any domain of structural science
- CIF can host more than just data (symmetry, unit-cell parameters, atomic coordinates, ADPs); it can also host structural, experimental, and processing details, as well as provenance, authorship, journal details, etc.
- CIF dictionaries** provide a formal taxonomy of crystallographic terms and ideas that can be **discipline-specific**
 - Core dictionary (coreCIF) „set of data names designed to cover the requirements of archiving and exchanging raw and processed data and derived structural results“
 - Powder dictionary (pdCIF)
 - Modulated and composite structures dictionary (msCIF)
 - Electron density dictionary (rhoCIF)
 - Twinning dictionary
 - Magnetic dictionary (magCIF)
 - Topology dictionary (topoCIF)
 - Macromolecular dictionary (mmCIF)
 - Symmetry dictionary (symCIF)
 - Image dictionary (imgCIF)
 - Restraints dictionary





Data flavors: the raw, the cooked and the medium-rare

- Q: Which of the data types – (1) raw, (2) processed or (3) derived data – can be described using the CIF format?
- A: All of them! However, raw data requires imgCIF/CBF ontologies, the particular formats describing diffraction images.
- Q: Do the major structural databases accept raw diffraction data?
- A: Currently not, but most large-scale photon facilities host and make raw diffraction data publicly available after the embargo period. The in-house diffractometer data can be uploaded to free data-sharing platforms (e.g. Zenodo).
- Q: And how about the processed data (structure factors)?
- A: From 2011 the Cambridge Structural Database (CSD) and the Inorganic Crystal Structure Database (ICSD) strongly encourage the inclusion of structure factor data along with CIFs, in line with the recommendations by the IUCr. This service is also available in the Crystallography Open Database (COD). Besides, many journals accept CIF files with structure factors as Electronic Supplementary Information.



Raw diffraction data reuse: the good, the bad and the challenging A Satellite Workshop to the XXVI IUCr Congress

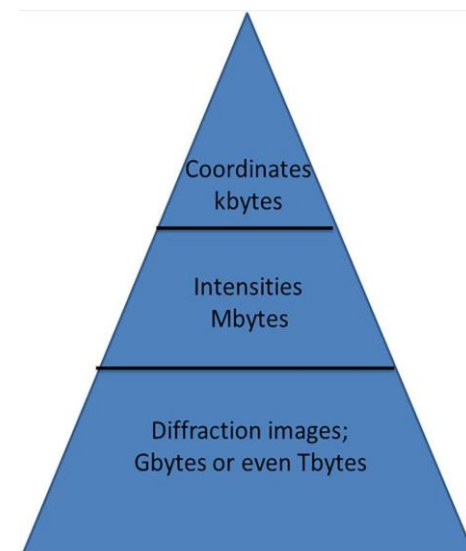
Organised by Loes Kroon-Batenburg, Selina Storm, John R. Helliwell and Brian McMahon under the auspices of the IUCr Committee on Data

The raw, the cooked and the medium-rare: unmerged diffraction data as a rich source of opportunities for data re-use and improvements in methods and results

G. Bricogne, C. Flensburg, R. H. Fogh, P. A. Keller, I. J. Tickle and C. Vonrhein

Tuesday 22 August 2023

International Convention Centre, Melbourne, Victoria, Australia



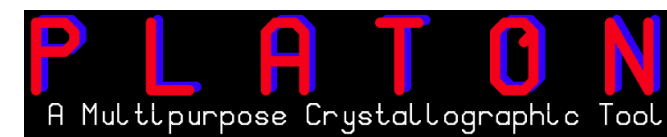
The Big Data pyramid, after: J. R. Helliwell, *Struct. Dyn.* **2019**, 6, 054306.



The element of trust: data validation

checkCIF is the IUCr service that reports on the completeness, consistency and integrity of CIF files

- Began as IUCr journals in-house suite for validation and consistency checks
- Consolidated with Ton Spek's PLATON (tool for analyzing geometrical calculations, e.g. bond lengths, angles, torsions and various structural tests, e.g. for missing symmetry, voids in the lattice) and now offered as public service
- Adopted by many journal publishers and databases (checkCIF reports required!)



Anthony L. Spek

checkCIF is sponsored by IUCr Journals, Elsevier, Wiley, Royal Society of Chemistry, and Crystals.

A service of the International Union of Crystallography

checkCIF reports on the consistency and integrity of crystal structure determinations reported in CIF format.

Please upload your CIF using the form below.

File name: No file selected.

Select form of checkCIF report

☒ HTML

☐ PDF

☐ PDF (recommended for CIFs that might take a long time to check)

Select validation type

☒ Full validation of CIF and structure factors

☐ Full IUCr publication validation of CIF and structure factors

☐ Validation of CIF only (no structure factors)

Output Validation Response Form

☐ Level A alerts only

☐ Level A and B alerts

☐ Level A, B and C alerts

☒ None

Information about this version of checkCIF ...

Useful links

Prepublication check for submissions to IUCr journals

Details of checkCIF/PLATON tests

CIF dictionary

Download CIF editor (pubCIF) from the IUCr

Download CIF editor (enCIFer) from the CCDC

checkcif.iucr.org/cgi-bin/checkcif_hkl.pl

Datablock: I

Bond precision: = 0.0000 Å Wavelength=0.71073

Cell: a=1 b=1 c=1
alpha=90 beta=90 gamma=90

Temperature: 0 K

Volume: 1 Calculated 738.55(7) Reported 738.55(7)

Space group: P 21 21 21

Hall group: P 2ac 2ab

Moiety formula: ?

Sum formula: ?

Mr: 0.00 152.15

Dx, g cm⁻³: 0.000 1.368

Z: 1 4

Mu (mm⁻¹): 0.000 0.105

F000: 0.0 0.0

F000': 0.00

h, k, l max: 6, 11, 18

Nref: 871

Tmin, Tmax: 0.978, 0.983

Tmin': 0.950

Correction method= Not given

Data completeness= Theta(max)= 26.000

R(reflections)= 0.0438(706) wR2(reflections)= 0.1113(871)

S = 1.085 Npar= 101

The following ALERTS were generated. Each ALERT has the format test-name_ALERT-alert-type_alert-level.

Click on the hyperlinks for more details of the test.

Alert level A

SYMM001_ALERT_1_A Symmetry cell setting is missing

The cell setting should be one of the following

- * monoclinic
- * orthorhombic
- * tetragonal
- * rhombohedral
- * trigonal
- * hexagonal
- * cubic

The following tests will not be performed.

SYMM01_ALERT_1_G

PLAT198_ALERT_1_A Missing _diffm_ambient_temperature Datum Please Add

PLAT801_ALERT_4_A Cell Data Missing, Incomplete or Out-of-Order Please Check

Alert level G

PLAT005_ALERT_5_G No Embedded Refinement Details Found in the CIF Please Do I

PLAT104_ALERT_1_G The Reported Crystal System is Inconsistent with P212121 Check

Completeness check

Alert level A

PLAT150_ALERT_1_A Volume as Calculated Differs from that Given ... 738.55 Ang-3

PLAT198_ALERT_1_A Missing _diffm_ambient_temperature Datum Please Add

PLAT701_ALERT_1_A Bond Calc 1.384(5), Rep 1.368(5), Dev... 3.20 Sigma

PLAT701_ALERT_1_A Bond Calc 1.555 1.555 # 1 Check

PLAT701_ALERT_1_A Bond Calc 1.527(5), Rep 1.496(4), Dev... 6.20 Sigma

PLAT702_ALERT_1_A Angle Calc 121.9(3), Rep 122.9(3), Dev... 3.33 Sigma

PLAT702_ALERT_1_A Angle Calc 1.555 1.555 # 13 Check

PLAT707_ALERT_1_A D...A Calc 2.732(4), Rep 2.676(4), Dev... 14.00 Sigma

PLAT707_ALERT_1_A D...A Calc 1.555 4.745 # 19 Check

PLAT707_ALERT_1_A D...A Calc 2.747(3), Rep 2.689(3), Dev... 19.33 Sigma

PLAT707_ALERT_1_A D...A Calc 1.555 2.665 # 19 Check

Alert level B

PLAT701_ALERT_1_B Bond Calc 1.382(4), Rep 1.371(4), Dev... 2.75 Sigma

PLAT701_ALERT_1_B Bond Calc 1.555 1.555 # 4 Check

PLAT701_ALERT_1_B Bond Calc 1.383(5), Rep 1.368(5), Dev... 3.00 Sigma

PLAT701_ALERT_1_B Bond Calc 1.555 1.555 # 6 Check

PLAT701_ALERT_1_B Bond Calc 1.395(4), Rep 1.385(5), Dev... 2.50 Sigma

PLAT702_ALERT_1_B Angle Calc 123.6(3), Rep 122.9(3), Dev... 2.33 Sigma

PLAT702_ALERT_1_B Angle Calc 1.555 1.555 # 14 Check

Alert level C

PLAT046_ALERT_1_C Reported Z, MW and D(calc) are Inconsistent 1.329 Check

PLAT068_ALERT_1_C Reported F000 Differs from Calc'd (or Missing)... Please Check

PLAT242_ALERT_2_C Low 'MainMol' Ueq as Compared to Neighbors of ... C8 Check

PLAT340_ALERT_2_C Low Bond Precision on: C-C Bonds 0.0045 Ang.

PLAT355_ALERT_3_C Long O-H(X0.82,N0.98A) O3 -H3A ... 1.01 Ang.

PLAT701_ALERT_1_C Bond Calc 1.195(5), Rep 1.186(4), Dev... 1.80 Sigma

PLAT702_ALERT_1_C Angle Calc 1.555 1.555 # 11 Check

PLAT702_ALERT_1_C Angle Calc 120.9(3), Rep 120.3(3), Dev... 2.00 Sigma

PLAT702_ALERT_1_C Angle Calc 1.555 1.555 # 4 Check

PLAT702_ALERT_1_C Angle Calc 118.5(3), Rep 117.9(3), Dev... 2.00 Sigma

PLAT702_ALERT_1_C Angle Calc 1.555 1.555 # 10 Check

PLAT702_ALERT_1_C Angle Calc 120.0(3), Rep 120.5(3), Dev... 1.67 Sigma

PLAT702_ALERT_1_C Angle Calc 1.555 1.555 # 12 Check

PLAT702_ALERT_1_C Angle Calc 119.1(3), Rep 119.5(3), Dev... 1.33 Sigma

PLAT702_ALERT_1_C Angle Calc 1.555 1.555 # 16 Check

PLAT702_ALERT_1_C Angle Calc 120.7(3), Rep 121.1(3), Dev... 1.33 Sigma

PLAT702_ALERT_1_C Angle Calc 1.555 1.555 # 19 Check

Alert level G

PLAT005_ALERT_5_G No Embedded Refinement Details Found in the CIF Please Do I

PLAT007_ALERT_5_G Number of Unrefined Donor-H Atoms 2 Report

PLAT721_ALERT_1_G Bond Calc 1.04000, Rep 1.02990 Dev... 0.01 Ang.

PLAT721_ALERT_1_G Bond Calc 1.555 1.555 # 14 Check

PLAT721_ALERT_1_G Bond Calc 0.92000, Rep 0.89950 Dev... 0.02 Ang.

PLAT721_ALERT_1_G Bond Calc 1.555 1.555 # 17 Check

PLAT721_ALERT_1_G Bond Calc 1.01000, Rep 0.98500 Dev... 0.02 Ang.

PLAT721_ALERT_1_G Bond Calc 1.555 1.555 # 18 Check

PLAT721_ALERT_1_G Bond Calc 0.97000, Rep 0.95040 Dev... 0.02 Ang.

PLAT722_ALERT_1_G Angle Calc 1.555 1.555 # 19 Check

PLAT722_ALERT_1_G Angle Calc 119.00, Rep 117.90 Dev... 1.10 Degree

PLAT722_ALERT_1_G Angle Calc 1.555 1.555 # 17 Check

PLAT722_ALERT_1_G Angle Calc 119.00, Rep 117.40 Dev... 1.60 Degree

PLAT722_ALERT_1_G Angle Calc 1.555 1.555 # 25 Check

PLAT725_ALERT_2_G D-H Calc 1.01000, Rep 0.98000 Dev... 0.03 Ang.

PLAT725_ALERT_2_G D-H Calc 1.555 1.555 # 19 Check

PLAT725_ALERT_2_G D-H Calc 0.97000, Rep 0.95000 Dev... 0.02 Ang.

PLAT725_ALERT_2_G D-H Calc 1.555 1.555 # 19 Check

PLAT726_ALERT_2_G H...A Calc 1.74000, Rep 1.71000 Dev... 0.03 Ang.

PLAT726_ALERT_2_G H...A Calc 1.555 4.745 # 19 Check

PLAT726_ALERT_2_G H...A Calc 1.78000, Rep 1.74000 Dev... 0.04 Ang.

PLAT726_ALERT_2_G H...A Calc 1.555 2.665 # 19 Check

A typo in a cell parameter ($b = 9.3092 \text{ \AA}$ instead of $9.0392 \text{ \AA}$) generates inconsistencies and anomalies in bond geometry

Consistency check



Three prongs of the *checkCIF* approach

FAIR principles do not, in themselves, cover the crucial aspects of intrinsic data quality and FAIR data are not *per se* high quality data! Three C's of *checkCIF**:

- **Completeness:** Including minimal metadata (complete crystal description, details of the diffraction experiment, structure solution and refinement strategy), required for reproducibility
- **Correctness:** Integrity and internal self-consistency (e.g. space group must agree with lattice parameters)
- **Context:** Comparison with similar structures in the wider universe of knowledge (e.g. tests for missed symmetry, missed twinning, solvent accessible voids, and mis-assigned atom types)



*McMahon, B. *The vital role of Crystallographic Information Files in chemical and biological crystallography to underpin the databases' validation reports*, Data Science Skills in Publishing Workshop, 2019, Vienna.



checkCIF reports and alerts

The automated *checkCIF* report contains three types of alerts:

- **ALERT level A** = In general: serious problem
- **ALERT level B** = Potentially serious problem
- **ALERT level C** = Check and explain

Try to eliminate the problems, but do not change the CIF or alter your refinement just to make *checkCIF* alerts vanish! Sometimes **alerts** cannot be eliminated, but instead **have valid reasons for their presence**. In these cases **validation reply form (vrf)** statements can be inserted into a CIF file, which acknowledge and account for *checkCIF* alerts.

```
# start Validation Reply Form
_vrf_ATOM007_Sb3N5_35_GPa
;
PROBLEM: _atom_site_aniso_label is missing
RESPONSE: Due to incompleteness of the high pressure dataset for a sample
in a diamond anvil cell ADPs of all atoms were refined
in isotropic approximation.
;
_vrf_PLAT029_Sb3N5_35_GPa
;
PROBLEM: _diffrn_measured_fraction_theta_full value Low .      0.462 Why?
RESPONSE: This is a high pressure data set, a substantial part of
a reciprocal space was shaded by the diamond anvil cell.
;
_vrf_PLAT911_Sb3N5_35_GPa
;
PROBLEM: Missing FCF Refl Between Thmin & STh/L=      0.600      80
Report
RESPONSE: Certain part of the reflections is missing due to shading
by the diamond anvil cell.
;
# end Validation Reply Form
```



research communications

checkCIF validation ALERTS: what they mean and how to respond

Anthony L. Spek*

Acta Cryst. (2020). E76, 1–11

<https://doi.org/10.1107/S2056989019016244> 1

A. L. Spek, *Acta Crystallogr. E* **2020**, 76, 1–11.

The following ALERTS were generated. Each ALERT has the format
test-name_ALERT_alert-type_alert-level.
Click on the hyperlinks for more details of the test.

Alert level A

ATOM007_ALERT_1_A _atom_site_aniso_label is missing
Unique label identifying the atom site.

Author Response: Due to incompleteness of the high pressure dataset for a sample in a diamond anvil cell ADPs of all atoms were refined in isotropic approximation.

PLAT029_ALERT_3_A _diffrn_measured_fraction_theta_full value Low . 0.462 Why?

Author Response: This is a high pressure data set, a substantial part of a reciprocal space was shaded by the diamond anvil cell.

Alert level B

PLAT911_ALERT_3_B Missing FCF Refl Between Thmin & STh/L= 0.600 80 Report

Author Response: Certain part of the reflections is missing due to shading by the diamond anvil cell.



CIF and *checkCIF*: Editors' requirements



Membership ▼ Publishing ▼ Policy and campaigning ▼ Standards and recognition ▼ Fundi

[Home](#) > [Publishing](#) > [Publish with us](#) > [Publish a journal article](#) > [Data sharing](#)

Data sharing

Guidance for best practice and reproducibility of experimental data.

Recommended repositories

A data repository is an external storage space for researchers to deposit datasets associated with their research. Data should be submitted to a discipline-specific, community-recognised repository where possible, or alternatively to an institutional repository, or a generalist repository if no subject discipline repository is available for the given data type.

The choice of repository is the author's decision, provided it is in line with institutional or funder guidelines. The exception to this is small molecule crystal data, which must be deposited with the Cambridge Crystallographic Data Centre (CCDC).

Data availability statements

- Crystallographic data for [compound number] has been deposited at the [name of repository, such as CCDC / ICSD / PBD] under [accession number] and can be obtained from [URL of data record, format <https://doi.org/DOI>].

CIF and *check*CIF: Editors' requirements

X-Ray crystallography

These guidelines provide details for the presentation of single crystal and powder diffraction data; they apply to submissions to any of our journals.

— Small molecule single crystal data

Authors should present their crystal data in a CIF (Crystallographic Information File) format and deposit this with the [Cambridge Crystallographic Data Centre](#) (CCDC) before submission. Data will be held in the CCDC's confidential archive until publication of the article, but it will be made accessible to reviewers and the publisher assigned to review the data.

At the point of publication, any deposited data will be made publicly available through the CCDC Access Structures service. In addition, organic and metal-organic structures will be curated into the Cambridge Structural Database, and inorganic structures will be curated into the Inorganic Crystal Structures Database (FIZ Karlsruhe).

Upon deposition, each data set is assigned a Digital Object Identifier (DOI), so that the crystal structure is unambiguously identified and registered.

Include CCDC or ICSD numbers in the manuscript prior to submission as part of a Data Availability Statement. During submission authors will be asked to cite CCDC or ICSD reference numbers; CIFs should not be submitted with the manuscript. Any revised CIFs should be deposited directly with the CCDC before the revised manuscript is submitted to us.



— CheckCIF

In addition, authors are required to provide a checkCIF report for their reported crystal data. The checkCIF report can be obtained via the International Union of Crystallography's (IUCr) [free checkCIF service](#), or as part of the CCDC deposition process. Any 'level A' alerts in the report should be explained in the submission details for the article or an explanation provided within the submitted CIF. Authors should submit the checkCIF reports to the Royal Society of Chemistry along with the manuscript files.

If the editor deems it necessary during the peer-review process, the crystallography associated with the manuscript may undergo specialist crystallographic assessment, in which case a report will be provided along with the other reports from reviewers. Any points raised in this assessment should be attended to and all revised CIFs should be deposited with the CCDC prior to uploading the revised manuscript.

For recommended information to include in your CIF, please see the CCDC CIF [deposition guidelines](#). If SQUEEZE or MASK procedures are used, this should be noted in the CIF file.

We encourage authors to include hkl data in the deposited CIF file. Alternatively authors can submit hkl data and the structure files (.fcf) separately during deposition with the CCDC. Raw data accompanying a structure should be made available by the authors for the review process, on request.

Raw data: cross-referencing between CCDC and external databases (e.g. Zenodo)

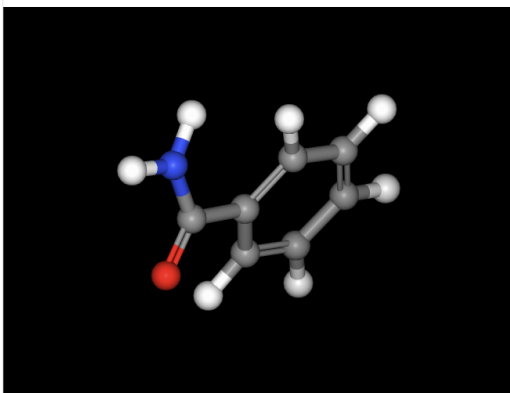
DOI: [10.5517/ccdc.csd.cc2jmntn](https://doi.org/10.5517/ccdc.csd.cc2jmntn)

LOHZER : New Structure undergoing enhancement

Space Group: $P 2_1/c$ (14), Cell: a 5.56(4)Å b 5.04(3)Å c 21.72(4)Å, α 90° β 90.100(11)° γ 90°

3D viewer

Ball and Stick ▼ No Labels ▼



▶ No Packing ▼ H DISORDER

Chemical diagram



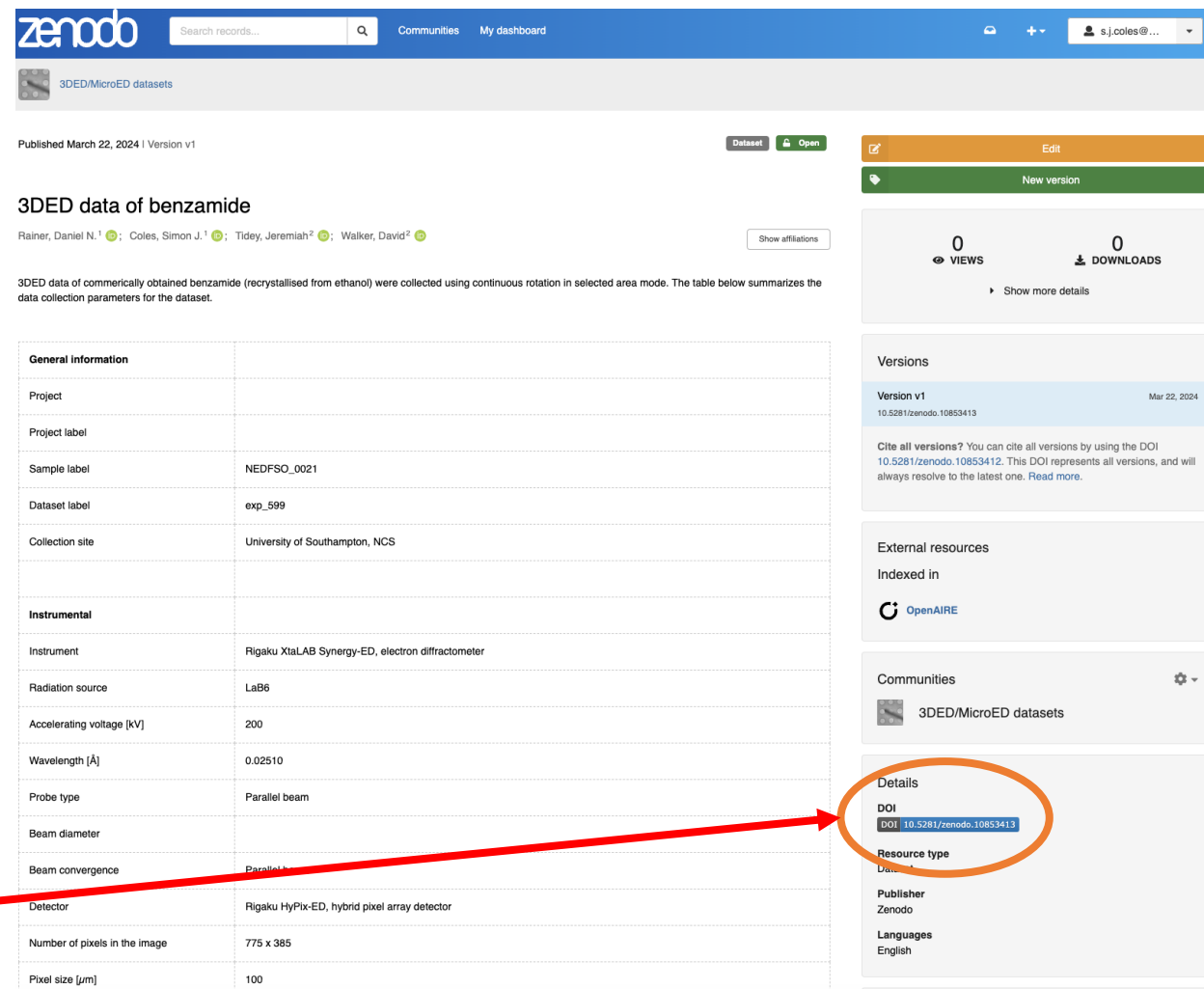
View group symbols key

Additional details

Deposition Number	2342602
Data Citation	Daniel N. Rainer, Simon J. Coles, Mark E. Light, Jeremiah P. Tidey, David Walker CCDC 2342602: Experimental Crystal Structure Determination, 2024, DOI: 10.5517/ccdc.csd.cc2jmntn
Deposited on	22/03/2024

Raw data DOI(s)

DOI [10.5281/zenodo.10853412](https://doi.org/10.5281/zenodo.10853412)



zenodo Search records... Communities My dashboard s.j.coles@...

3DED/MicroED datasets

Published March 22, 2024 | Version v1 Dataset Open Edit New version

3DED data of benzamide

Rainer, Daniel N.¹ ; Coles, Simon J.¹ ; Tidey, Jeremiah² ; Walker, David² Show affiliations

3DED data of commercially obtained benzamide (recrystallised from ethanol) were collected using continuous rotation in selected area mode. The table below summarizes the data collection parameters for the dataset.

General information	
Project	
Project label	
Sample label	NEDFSO_0021
Dataset label	exp_599
Collection site	University of Southampton, NCS
Instrumental	
Instrument	Rigaku XtaLAB Synergy-ED, electron diffractometer
Radiation source	LaB6
Accelerating voltage [kV]	200
Wavelength [Å]	0.02510
Probe type	Parallel beam
Beam diameter	
Beam convergence	Parallel beam
Detector	Rigaku HyPix-ED, hybrid pixel array detector
Number of pixels in the image	775 x 385
Pixel size [µm]	100

0 VIEWS 0 DOWNLOADS Show more details

Versions

Version v1 Mar 22, 2024
[10.5281/zenodo.10853413](https://doi.org/10.5281/zenodo.10853413)

Cite all versions? You can cite all versions by using the DOI [10.5281/zenodo.10853412](https://doi.org/10.5281/zenodo.10853412). This DOI represents all versions, and will always resolve to the latest one. Read more.

External resources

Indexed in OpenAIRE

Communities

3DED/MicroED datasets

Details

DOI [10.5281/zenodo.10853413](https://doi.org/10.5281/zenodo.10853413)

Resource type Dataset

Publisher Zenodo

Languages English

Courtesy of Simon Coles

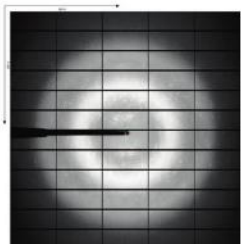
Continuing FAIRification of raw data within CIF

A brief history of raw data deposition

- Launch of *IUCrData* for peer-reviewed short structure reports (2016)
- A Gold Standard for metadata to describe an MX raw data (2020)
- Raw Data Letters**, a new section of *IUCrData* for authors to describe unprocessed diffraction images (2022)
- checkCIF for raw data** (2022)



Beam centre check image

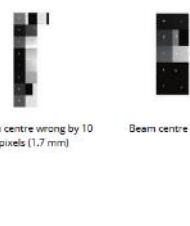


Predicted peak positions



Reversed axis of rotation

Predicted peak positions



Beam centre wrong by 10 pixels (1.7 mm)

Beam centre correct

research papers

IUCrJ

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BIOLOGY | MEDICINE

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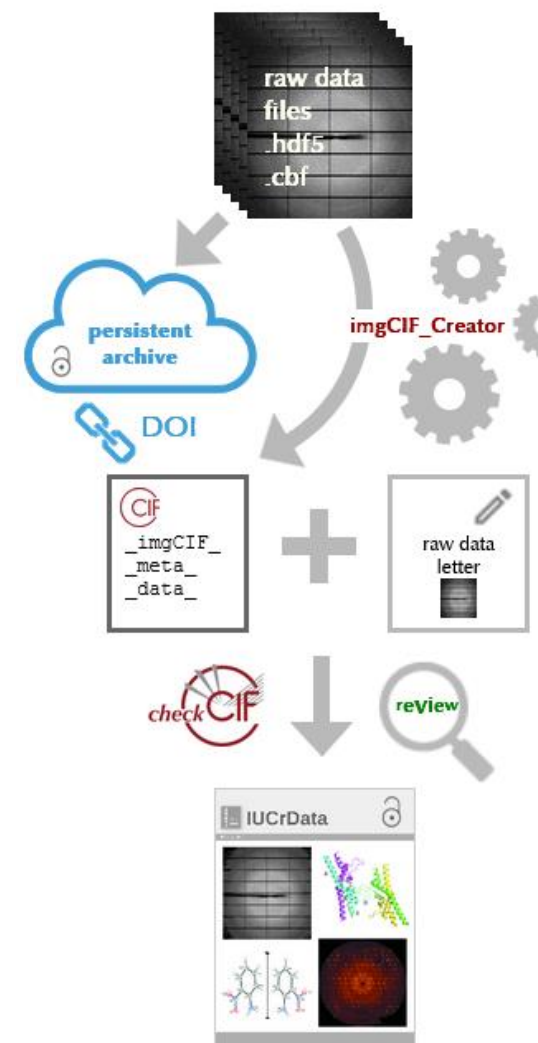
Gold Standard for macromolecular crystallography diffraction data

Herbert J. Bernstein,^{a,*} Andreas Förster,^b Asmit Bhowmick,^c Aaron S. Brewster,^c Sandor Brockhauser,^{d,e,f} Luca Gelisio,^g David R. Hall,^h Filip Leonarski,ⁱ Valerio Mariani,^g Gianluca Santoni,^j Clemens Vonrhein^k and Graeme Winter^h

784 <https://doi.org/10.1107/S2052252520008672>

IUCrJ (2020). 7, 784–792

raw data → archive → publish



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*chemical crystallography,
macromolecular crystallography,
raw data archiving*

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structural chemistry, databases*



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data science*



New challenge: Crystal Structure Prediction standards and CIF dictionary

Number of theoretically calculated crystal structures far exceeds that of experimentally determined structures!

- In one year a draft of the final dictionary will be communicated to the IUCr and the finalized version will be officially published
- In the meantime feedback, input and approval from community is required
- Available at: github.com/COMCIFS/Structure_Prediction_Dictionary

CSP Data Standards

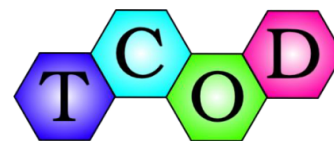
Scope of the project:

- An inclusive initiative that establishes **community standards** for predicted crystal structures
- Enables effective publication of results and easy **search and retrieval** across resources
- Supports levels of **reproducibility** deemed appropriate by different communities
- **Flexible structure** for future expansion as CSP methods are continuing to evolve



IUCr
International Union
of Crystallography

CCDC



Crystal Structure
Prediction

Inorganic CSP

Organic CSP

MOFs CSP



James Hester



Nicholas Francia



Ian Bruno

Potential of CIF to host non-crystallographic (spectroscopic) data

research papers



Received 23 November 2018
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Edited by A. Borbély, Ecole National Supérieure
des Mines, Saint-Etienne, France

Keywords: Raman spectroscopy; open
databases; combined Raman–X-ray diffraction;
DDLm dictionary; CIF2.

Raman Open Database: first interconnected Raman–X-ray diffraction open-access resource for material identification

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Detailed crystallographic information provided by X-ray diffraction (XRD) is complementary to molecular information provided by Raman spectroscopy. Accordingly, the combined use of these techniques allows the identification of an unknown compound without ambiguity. However, a full combination of Raman and XRD results requires an appropriate and reliable reference database with complete information. This is already available for XRD. The main objective of this paper is to introduce and describe the recently developed Raman Open Database (ROD, <http://solsa.crystallography.net/rod>). It comprises a collection of high-quality uncorrected Raman spectra. The novelty of this database is its interconnectedness with other open databases like the Crystallography Open Database (<http://www.crystallography.net/cod>) and Theoretical Crystallography Open Database (<http://www.crystallography.net/tcod>). The syntax adopted to format entries in the ROD is based on the worldwide recognized and used CIF format, which offers a simple way for data exchange, writing and description. ROD also uses JCAMP-DX files as an alternative format for submitted spectra. JCAMP-DX files are compatible to varying degrees with most commercial Raman software and can be read and edited using standard text editors.



Journal of
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Radiation

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J. Synchrotron Rad. (2012). **19**, 869–874

q2xafs workshop

Towards data format standardization for X-ray absorption spectroscopy

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A working group on data format standardization for X-ray absorption spectroscopy (XAS) has recently formed under the auspices of the International X-ray Absorption Society and the XAFS Commission of the International Union of Crystallography. This group of beamline scientists and XAS practitioners has been tasked to propose data format standards to meet the needs of the world-wide XAS community. In this report, concepts for addressing three XAS data storage needs are presented: a single spectrum interchange format, a hierarchical format for multispectral X-ray experiment, and a relational database format for XAS data libraries.

Keywords: XAFS; standardization; data formats.



Synergies between IUCr and other scientific standard organizations

CIF integrates standards developed within IUPAC

- Wavelength description uses the IUPAC nomenclature
- Enantioexcess is as per IUPAC recommendation
- InChi and InChIKey can be embedded in CIF
- Systematic chemical name and chemical formula follow IUPAC standards
- Topological CIF dictionary includes recommendations of IUPAC on nomenclature for coordination polymers

Liaisons between CIF and CDIF

- Physical Sciences Data Infrastructure (PDSI) at Southampton managed by Simon Coles strives to introduce CDIF (the CODATA Cross Domain Interoperability Framework) that integrates a wide range of resources across chemistry (with crystallography and CIF at the core)

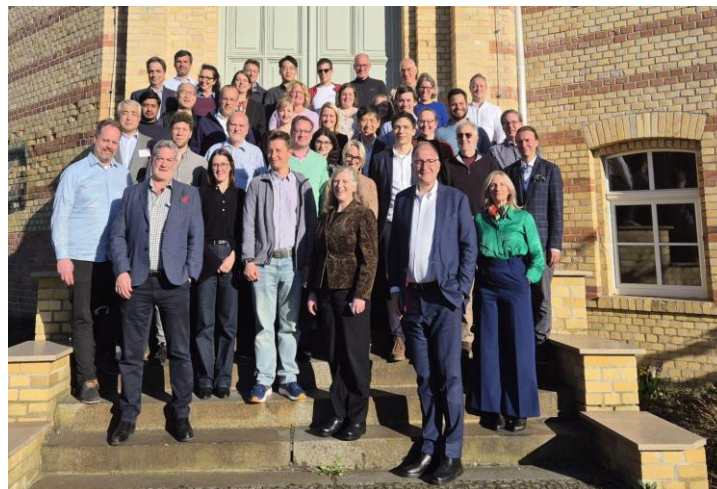


Coalition for the Sustainability of Digital Data Standards in the Chemical Sciences

IUCr (representing the CIF standard) actively participates in the DigSustain meetings from the beginning



DigSustain1: Sustainable Business Modeling for Digital Standards Development, Cambridge, UK, March 2024



DigSustain2: Digital Data Standards Sustainability in Chemical Sciences, Delitzsch, Germany, April 2025



DigSustain3: Coalition for the Sustainability of Digital Data Standards in the Chemical Sciences Planning Meeting, London, UK, November 2025

Stay tuned for more details in the talk of Wendy Patterson!



Where do we go from here?

Impact of CIF can be assessed by its popularity

- *'The single most important impact of CIF has been the ability to easily **access data globally across journals, databases and laboratories**'**
- CIF is now one of Universal Data Languages
- Peter Murray-Rust called CIF *'**the best working data system in the scientific world**'* (2014)
- 'Technology lock-in': foundational technology choice can become so embedded that it stifles future alternatives

Progress and adoption can be measured only indirectly

- Efficiency to submission–deposition processes
- Better understanding of particular data items across laboratories and databases
- Adoption in programming libraries and end-user software
- *'Anyone can place their own tag–value items into a CIF, but until their definitions are accepted into the global ontology they can only be for local use'**
- Official global ontology is the responsibility of the IUCr COMCIFS group, who moderates and maintain the standard
- Important to keep momentum and train new generations
- **The art of understanding and being understood**



*S. R. Hall, B. McMahon, *Data Science Journal* **2016**, 15, 1–15.



Thank you for listening!

Acknowledgments



Simon Coles



James Hester



John R. Helliwell



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