

# Using COD and TCOD, searching and getting data

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USNC/Cr Database Workshop 2022



BY SA

# Family of open-access structural databases

Database	Records	License	URL	Est.
COD	490 000	public domain	<a href="https://www.crystallography.net/cod/">https://www.crystallography.net/cod/</a>	2003
PCOD	1 000 000	public domain	<a href="https://www.crystallography.net/pcod/">https://www.crystallography.net/pcod/</a>	2003
MPOD	300	public domain	<a href="http://mpod.cimav.edu.mx/">http://mpod.cimav.edu.mx/</a>	2010
TCOD	2 600	public domain	<a href="https://www.crystallography.net/tcod/">https://www.crystallography.net/tcod/</a>	2013
ROD	1 100	public domain	<a href="https://solsa.crystallography.net/rod/">https://solsa.crystallography.net/rod/</a>	2017

## COD entry

- ▶ Describes a single crystal structure
- ▶ Has provenance record
- ▶ May be accompanied by diffraction data
- ▶ Is assigned permanent 7-digit identifier

# Deposition types in the COD

- ▶ Published material:
  - ▶ harvested journal supplements
  - ▶ donated collections
  - ▶ individual depositors
- ▶ Prepublication material
- ▶ Personal communications

# Entry information card

## Contents:

- ▶ 3D structure preview
- ▶ links to download files
- ▶ links to other databases
- ▶ bibliographic data
- ▶ cell parameters
- ▶ contents (name, formula, SMILES)
- ▶ space group information
- ▶ experiment details
- ▶ version history

## Example:

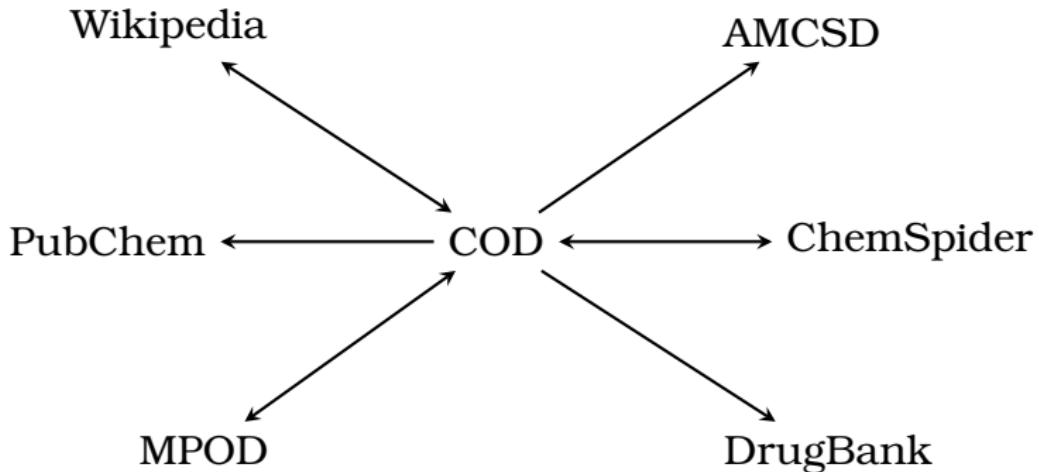
<https://www.crystallography.net/cod/4308000.html>

# Crystallographic Information File/Framework (CIF)

COD and TCOD uses CIF v1.1

- ▶ ASCII-based human-readable text
- ▶ Data storage:
  - ▶ key-value pairs
  - ▶ tables
- ▶ 3D viewers:
  - ▶ Avogadro
  - ▶ Jmol
- ▶ I/O software libraries:
  - ▶ codcif, CIF API (C)
  - ▶ ciftools-java (Java)
  - ▶ PyCifRW, pycodcif (Python)

# Links with other databases



# Data versioning

## Changes:

- ▶ are recorded
- ▶ get stable sequential identifiers
- ▶ are accompanied by log messages
- ▶ can be viewed

# Curation

## Curation

- ▶ Deposition time fixes
- ▶ (Semi-)automated fixes
- ▶ Manual curation

### Deposition time fixes

- ▶ Fix CIF syntax errors
- ▶ Reformat or estimate space group
- ▶ Reformat summary chemical formula
- ▶ Calculate cell volume
- ▶ Exclude unknown or placeholder CIF data items
- ▶ Convert temperature values
- ▶ Perform various other fixes

# Unusual entries

COD entries regarded unusual:

- ▶ duplicate entries ( $\approx 4500$ , e.g., 1000018)
- ▶ theoretical structures ( $\approx 600$ , e.g., 2100167)
- ▶ entries without coordinates ( $\approx 240$ , e.g., 1000195)
- ▶ retracted structures ( $\approx 150$ , e.g., 2015946)

# Validation database

- ▶ Uses CIF dictionaries to check data conformity
- ▶ Updated periodically
- ▶ Provides insight into most frequent issues

## Table of validation issues:

[https://sql.crystallography.net/db/cod\\_validation/validation\\_issue](https://sql.crystallography.net/db/cod_validation/validation_issue)

Vaitkus et al., 2021

# Querying COD

Search methods:

- ▶ bibliography
- ▶ cell parameters
- ▶ cell contents
- ▶ substructure by SMILES
- ▶ substructure by drawn fragment

Response formats:

- ▶ human-readable HTML
- ▶ database ID list
- ▶ CIF URL list
- ▶ CSV
- ▶ archive of matching files

# Download the whole COD. Quarterly releases

## Links

- ▶ Link for the most recent release:

<https://www.crystallography.net/cod/archives/cod-cifs-mysql.tgz>

- ▶ Older releases:

<https://www.crystallography.net/cod/archives/<YEAR>/data/>

## Contents

- ▶ coordinate files
- ▶ SQL table data

## Sizes of 2021-10-16 release

- ▶ TAR GZ – 17 GB
- ▶ TAR XZ – 12 GB
- ▶ ZIP – 18 GB

# Individual entry access

Whole file tree for browsing:

<https://www.crystallography.net/cod/cif>

Download individual entry

▶ coordinate file

<https://www.crystallography.net/cod/<ID>.cif>

▶ diffraction data

<https://www.crystallography.net/cod/<ID>.hkl>

▶ coordinate file at specified revision

<https://www.crystallography.net/cod/<ID>.cif@<revision>>

# Download the whole COD. *Rsync*

## *Rsync* access

```
$ mkdir cif hkl  
$ rsync -aq --delete rsync://www.crystallography.net/cif/ cif/  
$ rsync -aq --delete rsync://www.crystallography.net/hkl/ hkl/
```

(for verbose mode, replace `-q` with `-v`)

## Advantages:

- ▶ Subsequent `rsync` executions will fetch updates
- ▶ Network traffic is kept minimal

# Access with *Subversion*

Checkout (downloads uncompressed COD CIF collection)

```
$ export LC_TIME=en_US.UTF-8 # Fix locale  
$ svn checkout svn://www.crystallography.net/cod/cif
```

Fetch updates

```
$ svn up cif/
```

Go back to specific revision (here 1234)

```
$ svn up cif/ -r1234
```

# Examining changes with Subversion

Get contents of CIF file in revision 12345

```
$ svn cat svn://www.crystallography.net/cod/cif/1/00/00/1000000.cif -r12345
```

Get change of CIF file in revision 91932

```
$ svn diff svn://www.crystallography.net/cod/cif/1/00/00/1000000.cif -c91932
```

Index: 1000000.cif

```
=====
--- 1000000.cif (revision 91931)
+++ 1000000.cif (revision 91932)
@@ -348,3 +348,4 @@
 N2 H2C O6 1_655 .89 1.88 2.750(5) 164.8
 N2 H2D O4 2_645 .89 2.05 2.869(5) 153.3
 N2 H2E O7 1_655 .89 1.98 2.861(6) 170.9
+_journal_paper_doi 10.1107/S0108270100008532
```

# RESTful API for querying the COD

## Documentation:

[https://wiki.crystallography.net/RESTful\\_API/](https://wiki.crystallography.net/RESTful_API/)

Fetch entries describing cucurbiturils from year 2017, in CSV format

```
$ curl https://www.crystallography.net/cod/result -F text=cucurbituril  
-F format=csv -F year=2017
```

# SQL access

Connection details:

- ▶ **Server:** sql.crystallography.net
- ▶ **Database:** cod
- ▶ **User:** cod\_reader

**Explanation of fields in data table:**

[https://wiki.crystallography.net/cod\\_mysql\\_schema/](https://wiki.crystallography.net/cod_mysql_schema/)

## SQL access (*cont.*)

Find five most voluminous MOFs

```
$ mysql -h sql.crystallography.net cod -u cod_reader -e "select file,
    commonname, vol from data where commonname like '\"%MOF%\" order by
    vol desc limit 5"
+-----+-----+-----+
| file      | commonname   | vol      |
+-----+-----+-----+
| 4120255  | bio-MOF-102 | 425902  |
| 4120254  | bio-MOF-101 | 238778  |
| 4109100  | MOF-HTB'     | 148818  |
| 4111295  | mesoMOF-1   | 122163  |
| 4109099  | MOF-HTB      | 110988  |
+-----+-----+-----+
```

# OPTIMADE interface

Select ternary structures having C, Si, Ge or Sn, but not having Pb

```
$ curl -L https://www.crystallography.net/cod/optimade/structures -F 'filter=elements HAS ANY "C", "Si", "Ge", "Sn" AND NOT elements HAS "Pb" AND elements LENGTH 3'
```

Andersen et al., 2021

# Theoretical Crystallography Open Database (TCOD)

- ▶ Has 8-digit identifiers
- ▶ Contains computational details
- ▶ Embeds files used in computations

Merkys et al., 2017

# Acknowledgments

## **COD Advisory Board**

Saulius Gražulis  
Andrius Merkys  
Daniel Chateigner  
Robert T. Downs  
Werner Kaminsky  
Armel Le Bail  
Luca Lutterotti  
Peter Moeck  
Peter Murray-Rust  
Miguel Quirós