

Quality control of the chemical information in the Crystallography Open Database

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Abstract

- Crystallography Open Database (COD, [1]) is the largest open access crystallographic database, housing over 500k crystal structure entries.
- However, with the advent of machine learning methods and the increased reliance on black box approaches [2] quality control becomes vital to ensure:
 - ► integrity and consistency of both chemical and crystallographic descriptions;
 - presence and correctness of chemical representations such as SMILES [3] and chemical names;
 - correctness of chemical connectivity.

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- Solutions show the potential to improve the data quality in the COD:
 - cross-checking of crystallographic structures and chemical representations detects mismatches due to potential errors [4];
 - generated chemical names provide a reference point for name analysis and comparison;
 - derived covalent radii table provides insight into the choice of cutoff values [5].

Overlaying crystallographic and chemical annotations

Unsupervised method to derive a covalent radii table

- Uses Voronoi tessellation to find possible direct neighbours;
- Analyses distance distributions to locate van der Waals gap;
- Solves a system of equations to estimate covalent radii;
- Resulting radii table closely follows general trends of published tables.





- 1. Connectivity is inferred from the coordinates (CIF files);
- 2. Crystal contents are broken down into molecular entities;

Source #2

- 3. Chemical descriptions are extracted from chemical names, CML files and SMILES;
- 4. Molecular entities of compared crystals are matched;
- 5. Corresponding molecular entities are overlaid.

Cross-check results

Source #1

No. of pairs Matches

Validation of the derived covalent radii table

Compared to expert-assigned connectivity [3], the derived radii:

- tend to miss bonds involving I, Mn and N;
- mark false-positive Mo-Mo, Ti-Ti, V-V and W-W bonds, similarly as the table by Cordero et al. [7].

Generation of preferred IUPAC names (PINs)

Attempt to reproduce 3696 PINs from the IUPAC Blue Book [8]

	ChemOnomatopist v0.10.0	STOUT v2.0 [9]
Correct PIN	1321	1132
Alternative name	781	1874
Incorrect	987	690
Refused	607	0
Time	pprox 4 min.	pprox 230 min.

- ChemOnomatopist performs best with:
 - saturated and unsaturated acyclic and cyclic hydrocarbons;
 - bicyclic compounds, including heterocycles;
 - noncarbon acids.
- ChemOnomatopist needs improvement to handle:
 - multiplication nomenclature;
 - amides, amidines, esters and ethers;
 - diazenes, hydrazines, hydrazides and urea compounds;
 - charges and stereochemistry.
- STOUT most likely has been trained on older generation PINs.

Coordinate-derived	Chemical names	39636	88%
Coordinate-derived	CML	1551	89%
Coordinate-derived	Expert-curated [3]	188 137	85%
Chemical names	CML	1533	97%
Chemical names	Expert-curated [3]	34 670	92%

- Analysis of a couple dozens of mismatches identified incomplete or incorrect published chemical annotations [4].
- ► More interesting traits are dominated by differences in notation:
 - aromatic form vs. Kekulé form;
 - marked vs. unmarked metal coordination [6].

This research has received funding from the Research Council of Lithuania under Grant agreement No. MIP-23-87

On-line poster material: https://bit.ly/3X4D8Di



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