

# Resource Summary Report

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## RCSB PDB Software Tools

RRID:SCR\_000035

Type: Tool

### Proper Citation

RCSB PDB Software Tools (RRID:SCR\_000035)

### Resource Information

**URL:** <http://sw-tools.pdb.org/index.html>

**Proper Citation:** RCSB PDB Software Tools (RRID:SCR\_000035)

**Description:** Information Portal to Biological Macromolecular Structures provides variety of software tools made available through the RCSB. These tools include: data extraction and deposition preparation tools, data format conversion and validation tools, data parsing tools, dictionary and data management tools, visualization tools that support PDBx/mmCIF, and other PDBx/mmCIF software library tools.

**Synonyms:** RCSB Software Tools

**Resource Type:** topical portal, data or information resource, software resource, portal

**Keywords:** Information Portal, Biological Macromolecular Structure, RCSB, RCSB software tools, data extraction, format conversion, data parsing, data management, visualization, pdbx/mmCIF, pdbx, mmCIF

**Funding:**

**Availability:** Free, Available for download, Freely available

**Resource Name:** RCSB PDB Software Tools

**Resource ID:** SCR\_000035

**Alternate IDs:** nif-0000-31399

**License:** <http://sw-tools.pdb.org/license.txt>

**Record Creation Time:** 20220129T080159+0000

**Record Last Update:** 20250409T055948+0000

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## Ratings and Alerts

No rating or validation information has been found for RCSB PDB Software Tools.

No alerts have been found for RCSB PDB Software Tools.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We found 2 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [NIF](#).

Vallat B, et al. (2018) Development of a Prototype System for Archiving Integrative/Hybrid Structure Models of Biological Macromolecules. *Structure (London, England : 1993)*, 26(6), 894.

Williams SM, et al. (2017) A Mutation Directs the Structural Switch of DNA Binding Proteins under Starvation to a Ferritin-like Protein Cage. *Structure (London, England : 1993)*, 25(9), 1449.