

# Resource Summary Report

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## RSRef

RRID:SCR\_017211

Type: Tool

### Proper Citation

RSRef (RRID:SCR\_017211)

### Resource Information

**URL:** <https://biochem.missouri.edu/chapman/software.htm>

**Proper Citation:** RSRef (RRID:SCR\_017211)

**Description:** Software for fitting of atomic models into density maps derived from x-ray crystallography or electron microscopy.

**Resource Type:** software resource, software application, data processing software

**Defining Citation:** [PMID:23376441](#)

**Keywords:** Fitting, atomic, model, density, map, x ray, crystallography, electron, microscopy

**Funding:** NIGMS R01 GM66875;  
NIGMS R01 GM78538

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

**Resource Name:** RSRef

**Resource ID:** SCR\_017211

**Alternate URLs:** <http://xtal.ohsu.edu/software/rsref/readme.txt>

**Record Creation Time:** 20220129T080334+0000

**Record Last Update:** 20260731T043011+0000

### Ratings and Alerts

No rating or validation information has been found for RSRef.

No alerts have been found for RSRef.

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## Data and Source Information

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

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## Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Meyer NL, et al. (2019) Structure of the gene therapy vector, adeno-associated virus with its cell receptor, AAVR. eLife, 8.