Resource Summary Report

Generated by RRID on Aug 5, 2026

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: data processing software, software application, software resource

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: 20260805T044406+0000

Ratings and Alerts

No rating or validation information has been found for RSRef.

No alerts have been found for RSRef.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 1 mentions in open access literature.

Listed below are recent publications. The full list is available at RRID .

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