Resource Summary Report

Generated by SPARC SAWG on Jul 28, 2026

O

RRID:SCR_018394
Type: Tool

Proper Citation

O (RRID:SCR_018394)

Resource Information

URL: http://xray.bmc.uu.se/~alwyn/TAJ/Home.html

Proper Citation: O (RRID:SCR_018394)

Description: Software for visualization of macromolecular structures, atomic coordinates, and 3D intensity maps, and for building macromolecular structures.

Resource Type: data visualization software, 3d visualization software, image, software resource, software application, 3d spatial image, data processing software, data or information resource

Defining Citation: DOI:10.1107/s0108767390010224

Keywords: Macromolecular structure, atomic coordinate, protein structure, 3D visualization, model building, atomic modelling

Availability: Free, Available for download, Freely available

Resource Name: O

Resource ID: SCR_018394

Record Creation Time: 20220129T080340+0000

Record Last Update: 20260728T043544+0000

Ratings and Alerts

No rating or validation information has been found for O.

No alerts have been found for O.

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 SPARC SAWG .

Ahel J, et al. (2020) Moyamoya disease factor RNF213 is a giant E3 ligase with a dynein-like core and a distinct ubiquitin-transfer mechanism. eLife, 9.