

Resource Summary Report

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

Defining Citation: [DOI:10.1107/s0108767390010224](https://doi.org/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: 20260729T044441+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 [nidm-terms](#) .

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.