Resource Summary Report

Generated by RRID on Aug 14, 2026

iMol

RRID:SCR_018735
Type: Tool

Proper Citation

iMol (RRID:SCR_018735)

Resource Information

URL: http://www.pirx.com/iMol/index.shtml

Proper Citation: iMol (RRID:SCR_018735)

Description: Software molecular visualization application for Mac OS X operating system. Tool for chemists and molecular biologists used for handling small and large molecules, loading multiple molecules, moving and rotating them independently, or displaying molecular dynamics trajectory. Allows loading molecules using file formats: PDB, XYZ, MOL2, HIN, CAR, ALC, BIO.

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

Keywords: Molecular visualization, Mac OS X operating system, molecule, loading molecule, moving molecule, rotating molecule, displaying dynamics trajectory

Availability: Free, Available for download, Freely available

Resource Name: iMol

Resource ID: SCR_018735

Record Creation Time: 20220129T080341+0000

Record Last Update: 20260814T043015+0000

Ratings and Alerts

No rating or validation information has been found for iMol.

No alerts have been found for iMol.

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 .

Kang HE, et al. (2020) Incomplete glycosylation during prion infection unmasks a prion protein epitope that facilitates prion detection and strain discrimination. The Journal of biological chemistry, 295(30), 10420.