

# 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](#)

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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 [RRID](#) .

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