

Resource Summary Report

Generated by [PRECISE-TBI](#) on Sep 19, 2026

MEGA-MD

RRID:SCR_006403

Type: Tool

Proper Citation

MEGA-MD (RRID:SCR_006403)

Resource Information

URL: <http://www.megasoftware.net/mega-md/mega-md.php>

Proper Citation: MEGA-MD (RRID:SCR_006403)

Description: Software to forecast the deleteriousness of non-synonymous single nucleotide variants (nsSNVs) using multiple methods and explore them in the context of the variability permitted in the long-term evolution of the affected positions. A web version (MEGA-MDW) is also available.

Abbreviations: MEGA-MD

Synonyms: Molecular Evolutionary Genetics Analysis - Mutation Diagnosis, Molecular Evolutionary Genetics Analysis software with mutational diagnosis

Resource Type: analysis service resource, data analysis service, production service resource, service resource, software resource

Defining Citation: [PMID:24413669](#)

Keywords: windows, non-synonymous single nucleotide variant

Availability: Free for academic use, Registration required, Acknowledgement requested

Resource Name: MEGA-MD

Resource ID: SCR_006403

Alternate IDs: OMICS_02211

Record Creation Time: 20220129T080236+0000

Record Last Update: 20260919T195110+0000

Ratings and Alerts

No rating or validation information has been found for MEGA-MD.

No alerts have been found for MEGA-MD.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.