

Resource Summary Report

Generated by [PRECISE-TBI](#) on Sep 9, 2026

MADOKA

RRID:SCR_024522

Type: Tool

Proper Citation

MADOKA (RRID:SCR_024522)

Resource Information

URL: <http://madoka.denglab.org/>

Proper Citation: MADOKA (RRID:SCR_024522)

Description: Web server as ultra fast approach for large scale protein structure similarity searching. The upload protein structure file should be in PDB format. Used for searching similar protein structures by aligning input structure with the whole PDB library.

Resource Type: data access protocol, software resource, web service

Defining Citation: [PMID:31870277](#)

Keywords: large scale protein structure similarity searching, searching similar protein structures, aligning input structure with whole PDB library,

Availability: Free, Freely available

Resource Name: MADOKA

Resource ID: SCR_024522

Record Creation Time: 20231002T161337+0000

Record Last Update: 20260905T133032+0000

Ratings and Alerts

No rating or validation information has been found for MADOKA.

No alerts have been found for MADOKA.

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 [PRECISE-TBI](#) .

Wegrzyn K, et al. (2021) Defining a novel domain that provides an essential contribution to site-specific interaction of Rep protein with DNA. Nucleic acids research, 49(6), 3394.