

# Resource Summary Report

Generated by [NIF](#) on Apr 20, 2025

## MADOKA

RRID:SCR\_024522

Type: Tool

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### Proper Citation

MADOKA (RRID:SCR\_024522)

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### 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, web service, software resource

**Defining Citation:** [PMID:31870277](#)

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

**Funding:**

**Availability:** Free, Freely available

**Resource Name:** MADOKA

**Resource ID:** SCR\_024522

**Record Creation Time:** 20231002T161337+0000

**Record Last Update:** 20250420T015308+0000

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### Ratings and Alerts

No rating or validation information has been found for MADOKA.

No alerts have been found for MADOKA.

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## 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 [NIF](#).

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.