

Resource Summary Report

Generated by [SPARC SAWG](#) on Jul 29, 2026

Multiple Mapping Method Server

RRID:SCR_018015

Type: Tool

Proper Citation

Multiple Mapping Method Server (RRID:SCR_018015)

Resource Information

URL: <http://manaslu.fiserlab.org/MMM/>

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Description: Web server for modeling protein structure by using Multiple Mapping Method. Approach to sequence-to-structure alignment in comparative protein structure modeling.

Abbreviations: MMM server

Synonyms: Multiple Mapping Method server

Resource Type: software application, service resource, data access protocol, software resource, data processing software, alignment software, web service, image analysis software

Defining Citation: [PMID:16437570](#)

Keywords: Modeling protein structure, multiple mapping method, sequence, structure, alignment, comparative protein structure

Availability: Free, Freely available

Resource Name: Multiple Mapping Method Server

Resource ID: SCR_018015

Record Creation Time: 20220129T080338+0000

Record Last Update: 20260729T044449+0000

Ratings and Alerts

No rating or validation information has been found for Multiple Mapping Method Server.

No alerts have been found for Multiple Mapping Method Server.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

Listed below are recent publications. The full list is available at [SPARC SAWG](#) .

Ahmad J, et al. (2021) Structures of synthetic nanobody-SARS-CoV-2-RBD complexes reveal distinct sites of interaction and recognition of variants. Research square.

Rathkey JK, et al. (2020) Human polymorphisms in GSDMD alter the inflammatory response. The Journal of biological chemistry, 295(10), 3228.