

# Resource Summary Report

Generated by [PRECISE-TBI](#) on Jul 31, 2026

## Multiple Mapping Method Server

RRID:SCR\_018015

Type: Tool

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### Proper Citation

Multiple Mapping Method Server (RRID:SCR\_018015)

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### Resource Information

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

**Proper Citation:** Multiple Mapping Method Server (RRID:SCR\_018015)

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

**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:** 20260731T043018+0000

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### 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.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We found 2 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [PRECISE-TBI](#) .

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.