Resource Summary Report

Generated by RRID on Aug 31, 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/

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

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: 20260829T182552+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 3 mentions in open access literature.

Listed below are recent publications. The full list is available at RRID .

Ahmad J, et al. (2021) Structures of synthetic nanobody-SARS-CoV-2 receptor-binding domain complexes reveal distinct sites of interaction. The Journal of biological chemistry, 297(4), 101202.

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.