

Resource Summary Report

Generated by [Kravitz](#) on Sep 10, 2026

ModFOLD

RRID:SCR_024518

Type: Tool

Proper Citation

ModFOLD (RRID:SCR_024518)

Resource Information

URL: <https://www.reading.ac.uk/bioinf/ModFOLD/>

Proper Citation: ModFOLD (RRID:SCR_024518)

Description: Web server for the global and local quality estimation of 3D protein models

Synonyms: ModFOLD6, ModFOLD8

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

Defining Citation: [PMID:28460136](#)

Keywords: quality estimation, 3D protein models,

Availability: Restricted

Resource Name: ModFOLD

Resource ID: SCR_024518

Record Creation Time: 20231002T161337+0000

Record Last Update: 20260905T133032+0000

Ratings and Alerts

No rating or validation information has been found for ModFOLD.

No alerts have been found for ModFOLD.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 5 mentions in open access literature.

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

De'Ath C, et al. (2026) YdbL directly modulates YdbH-YnbE bridge formation to maintain Escherichia coli outer membrane homeostasis. bioRxiv : the preprint server for biology.

Edmunds NS, et al. (2024) Benchmarking of AlphaFold2 accuracy self-estimates as indicators of empirical model quality and ranking: a comparison with independent model quality assessment programmes. Bioinformatics (Oxford, England), 40(8).

Fu H, et al. (2024) EsigPBP3 Was the Important Pheromone-Binding Protein to Recognize Male Pheromones and Key Eucalyptus Volatiles. International journal of molecular sciences, 25(5).

Toth JM, et al. (2021) ResiRole: residue-level functional site predictions to gauge the accuracies of protein structure prediction techniques. Bioinformatics (Oxford, England), 37(3), 351.

Rodríguez-García Á, et al. (2021) Molecular Analysis of Streptomycin Resistance Genes in Clinical Strains of Mycobacterium tuberculosis and Biocomputational Analysis of the MtGidB L101F Variant. Antibiotics (Basel, Switzerland), 10(7).