

University of Reading

HTML Templates

The ModFOLD9 Server Results

Please cite the ModFOLD9 paper:

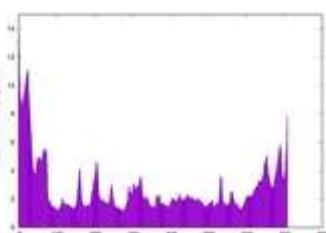
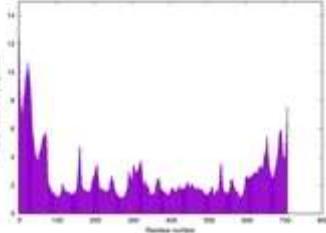
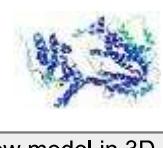
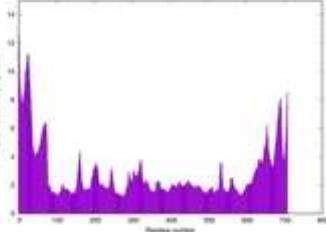
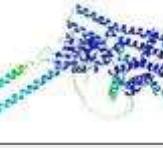
McGuffin, L.J. and Alharbi, S.M.A. (2024) ModFOLD9: a web server for independent estimates of 3D protein model quality. Journal of Molecular Biology, In press. [DOI](#) [PubMed](#)

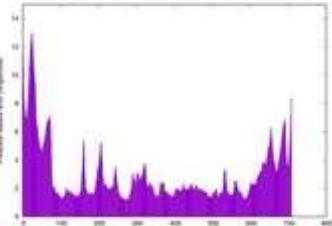
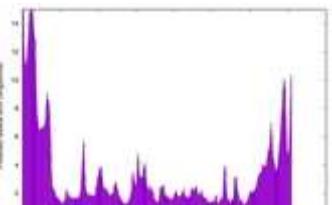
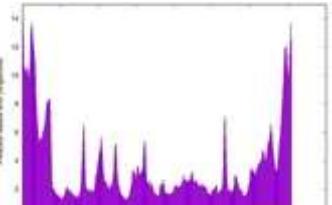
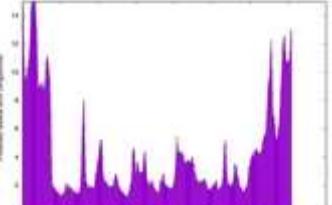
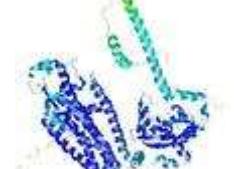
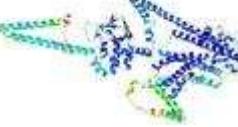
Click on the links below to download machine readable results files:

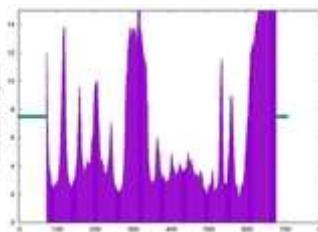
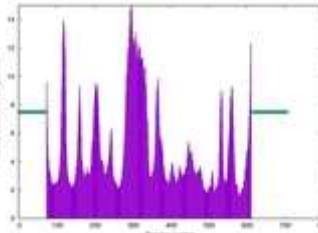
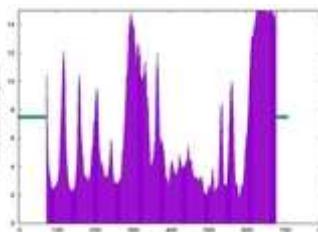
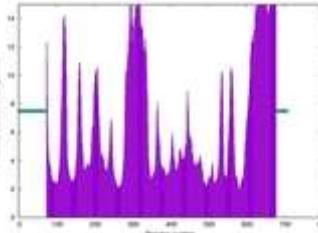
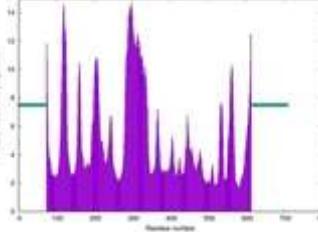
[Download Quality Assessment file](#) (raw model quality prediction scores in CASP QA format)

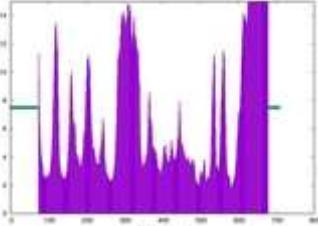
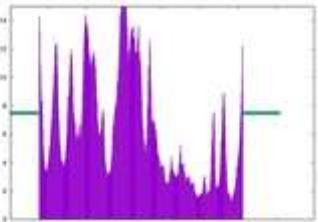
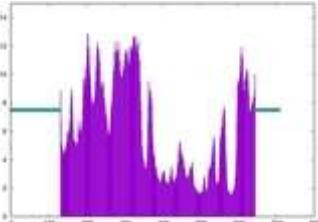
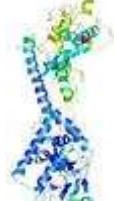
[Make a compressed file containing all evaluated 3D models](#) (models are in PDB format with predicted per-residue errors in the B-factor column)

Results will be available for **21 days** after job completion (subject to server capacity)

Graphical ModFOLD9 results for NoName Help					
Rank	Model ID	Confidence and P-value	Global model quality score	Residue quality plots	3D view of local quality
1	IntFOLD701.pdb	CERT: 3.159E-4	0.6200	 View and download	 View model in 3D and download Fix errors using ReFOLD3
2	CNGC6AlphaFold.pdb	CERT: 3.53E-4	0.6141	 View and download	 View model in 3D and download Fix errors using ReFOLD3
3	IntFOLD702.pdb	CERT: 4.2E-4	0.6048	 View and download	 View model in 3D and download Fix errors using ReFOLD3

4	IntFOLD703.pdb	CERT: 4.829E-4	0.5974	 View and download	 View model in 3D and download Fix errors using ReFOLD3
5	swissmodel02.pdb	CERT: 5.541E-4	0.5901	 View and download	 View model in 3D and download Fix errors using ReFOLD3
6	ESMFold.pdb	CERT: 8.915E-4	0.5648	 View and download	 View model in 3D and download Fix errors using ReFOLD3
7	RoseTTAFold2.pdb	HIGH: 1.296E-3	0.5449	 View and download	 View model in 3D and download Fix errors using ReFOLD3
8	OmegaFold.pdb	HIGH: 2.204E-3	0.5166	 View and download	 View model in 3D and download Fix errors using ReFOLD3

9	modeller04.pdb	UNCERT: 1.017E-1	0.3120	 View and download	 View model in 3D and download Fix errors using ReFOLD3
10	modeller07.pdb	UNCERT: 1.071E-1	0.3090	 View and download	 View model in 3D and download Fix errors using ReFOLD3
11	modeller06.pdb	UNCERT: 1.188E-1	0.3029	 View and download	 View model in 3D and download Fix errors using ReFOLD3
12	modeller05.pdb	UNCERT: 1.349E-1	0.2954	 View and download	 View model in 3D and download Fix errors using ReFOLD3
13	modeller02.pdb	UNCERT: 1.403E-1	0.2931	 View and download	 View model in 3D and download Fix errors using ReFOLD3

14	modeller01.pdb	UNCERT: 1.531E-1	0.2879	 View and download	 View model in 3D and download Fix errors using ReFOLD3
15	modeller03.pdb	UNCERT: 2.46E-1	0.2580	 View and download	 View model in 3D and download Fix errors using ReFOLD3
16	swissmodel01.pdb	UNCERT: 3.346E-1	0.2354	 View and download	 View model in 3D and download Fix errors using ReFOLD3

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[@reading.ac.uk](https://www.reading.ac.uk/bioinf/services/nFOLD/ModFOLDResultsv9_0.jsp?time=1_12_14_112_25-6-2024_QA_single2&md5=oo45edtpiv3mo5jj&targ...)

[Full contact details](#)

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