Predictions for Entire Proteomes

RRID:SCR_002803
Type: Tool

Proper Citation

Predictions for Entire Proteomes (RRID:SCR_002803)

Resource Information

URL: http://www.predictprotein.org/

Description: Web application for sequence analysis and the prediction of protein structure and function. The user interface intakes protein sequences or alignments and returned multiple sequence alignments, motifs, and nuclear localization signals.

Resource Name: Predictions for Entire Proteomes

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Resource Type: Resource, data analysis software, data processing software, software application, sequence analysis software, web application, software resource

Keywords: sequence analysis database, protein structure prediction, protein structure, protein function

Resource ID: SCR_002803

Parent Organization: Columbia University; New York; USA

Funding Agency: BMBF

References: PMID:24799431

Availability: Open for academic use, Acknowledgement requested

Website Status: Last checked up

Alternate IDs: nif-0000-00136
Old URLs: http://cubic.bioc.columbia.edu/pep/

Mentions Count: 229

Ratings and Alerts

No rating or validation information has been found for Predictions for Entire Proteomes.

No alerts have been found for Predictions for Entire Proteomes.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 229 mentions in open access literature.

Listed below are recent publications. The full list is available at FDI Lab - SciCrunch Infrastructure.


