Resource Summary Report

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Critical Assessment of Techniques for Protein Structure Prediction

RRID:SCR_008434
Type: Tool

Proper Citation

Critical Assessment of Techniques for Protein Structure Prediction (RRID:SCR_008434)

Resource Information

URL: http://predictioncenter.org/casp7/Casp7.html

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Description: The main goal of CASP is to obtain an in-depth and objective assessment of our current abilities and inabilities in the area of protein structure prediction. To this end, participants will predict as much as possible about a set of soon to be known structures. These will be true predictions, not "post-dictions" made on already known structures. Sponsors: This resource is supported by the US National Library of Medicine (NIH/NLM), National Institute of General Medical Sciences (NIH/NIGMS), BioSapiens Network of Excellence, and HP Invent. Keywords: Clinical, Technique, Protein, Prediction, Structure,

Synonyms: CASP7

Resource Type: portal, topical portal, data or information resource

Funding:

Resource Name: Critical Assessment of Techniques for Protein Structure Prediction

Resource ID: SCR_008434

Alternate IDs: nif-0000-30234

Old URLs: http://predictioncenter.gc.ucdavis.edu/casp7/

Record Creation Time: 20220129T080247+0000

Record Last Update: 20250417T065330+0000

Ratings and Alerts

No rating or validation information has been found for Critical Assessment of Techniques for Protein Structure Prediction.

No alerts have been found for Critical Assessment of Techniques for Protein Structure Prediction.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Kirillova S, et al. (2009) Protein domain boundary predictions: a structural biology perspective. The open biochemistry journal, 3, 1.