

# Resource Summary Report

Generated by [dkNET](#) on Apr 27, 2025

## Critical Assessment of Techniques for Protein Structure Prediction

RRID:SCR\_008434

Type: Tool

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### Proper Citation

Critical Assessment of Techniques for Protein Structure Prediction (RRID:SCR\_008434)

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### 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:** data or information resource, topical portal, portal

**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:** 20250426T060032+0000

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## 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.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We found 1 mentions in open access literature.

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

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