Critical Assessment of Methods of Protein Structure Prediction (CASP) - Round XIV.

Title: Critical Assessment of Methods of Protein Structure Prediction (CASP) - Round XIV.
Publication Type: Journal Article
Year of Publication: 2021
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Journal: Proteins
Date Published: 2021 Sep 17
ISSN: 1097-0134
Abstract: CASP is a community experiment to advance methods of computing three-dimensional protein structures from amino acid sequences. The methods evaluated include model building, refinement, estimation of model accuracy, and prediction of inter-residue contacts and distances, are also described.
DOI: 10.1002/prot.26237
Alternate Journal: Proteins
PubMed ID: 34533838