



Computational Methods for Protein Structure Prediction and Modeling: 2 (Biological and Medical Physics, Biomedical Engineering)

Ying Xu, Dong Xu, Jie Liang

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Volume two of this two volume sequence focuses on protein structure prediction and includes protein threading, De novo methods, applications to membrane proteins and protein complexes, structure-based drug design, as well as structure prediction as a systems problem. A series of appendices review the biological and chemical basics related to protein structure, computer science for structural informatics, and prerequisite mathematics and statistics. Key Features Presents a comprehensive overview of protein structure prediction methods Each chapter is a self contained review Coverage includes De novo methods of protein structure prediction, applications to membrane proteins and protein complexes and more 2nd volume of a two-volume sequence

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