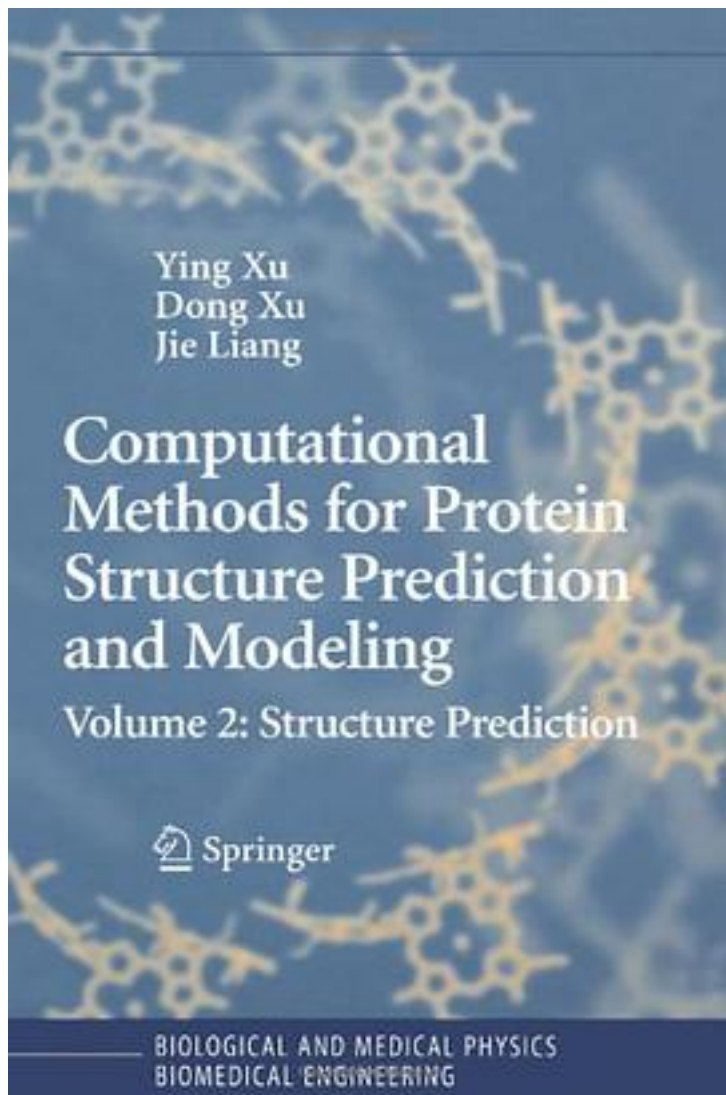


Computational Methods for Protein Structure Prediction and Modeling



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出版者: Springer Verlag

出版时间: 2006-12

装帧: HRD

isbn: 9780387333212

Volume Two of this two-volume sequence presents a comprehensive overview of protein structure prediction methods 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.

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