



Protein Folding Protocols (Methods in Molecular Biology)

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Covering experiment and theory, bioinformatics approaches, and state-of-the-art simulation protocols for better sampling of the conformational space, this volume describes a broad range of techniques to study, predict, and analyze the protein folding process. Protein Folding Protocols also provides sample approaches toward the prediction of protein structure starting from the amino acid sequence, in the absence of overall homologous sequences.

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