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Bayesian methods are central to bioinformatics, but also play an increasing role in structural bioinformatics. This book serves as an excellent entry point into the applications of probabilistic modeling methods in structural biology. There are two main themes in the book. One is the use of Bayesian methods to infer structures of biopolymers by analyzing experimental data (from NMR, SAXS etc) in a probabilistic framework. The second main theme is the use of Bayesian methods to analyze PDF Bayesian Methods In Structural Bioinformatics Free ...

This book is an edited volume, the goal of which is to provide an overview of the current state-of-the-art in statistical methods applied to problems in structural bioinformatics (and in particular protein structure prediction, simulation, experimental structure determination and analysis).

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The Statistical Structural Biology group focuses on Bayesian, probabilistic models of protein structure and their application to protein structure prediction, protein design and protein structure determination from experimental data (NMR, SAXS), including data obtained from protein ensembles.