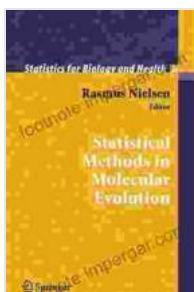


Statistical Methods In Molecular Evolution: Statistics For Biology And Health

Statistical methods play a vital role in the field of molecular evolution, providing researchers with the tools to analyze and interpret genetic data. This book provides a comprehensive overview of statistical methods in molecular evolution, with a focus on applications and real-world examples. The book is divided into three parts:



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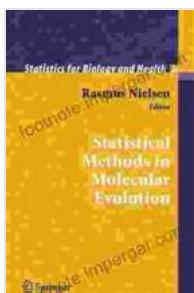
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- Bayesian statistics
- Machine learning
- Statistical genomics

This book provides a comprehensive overview of statistical methods in molecular evolution. The book is written in a clear and concise style, and it is packed with real-world examples. This book is an essential resource for researchers and students in the fields of biology and health.

About the author

Dr. Ziheng Yang is a professor of statistical genetics at the University of California, Berkeley. He is the author of several books and papers on statistical methods in molecular evolution. Dr. Yang is a Fellow of the American Statistical Association and the Royal Statistical Society.



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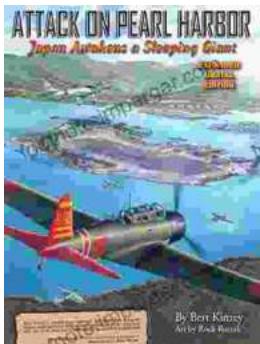
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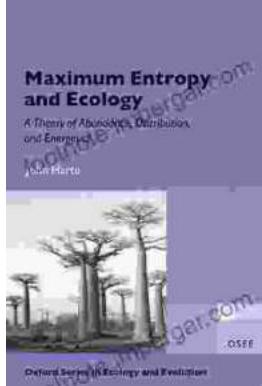
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