



# Bioinformatics and Molecular Evolution

*Paul G. Higgs, Teresa K. Attwood*

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In the current era of complete genome sequencing, *Bioinformatics and Molecular Evolution* provides an up-to-date and comprehensive introduction to bioinformatics in the context of evolutionary biology.

This accessible text:

- provides a thorough examination of sequence analysis, biological databases, pattern recognition, and applications to genomics, microarrays, and proteomics
- emphasizes the theoretical and statistical methods used in bioinformatics programs in a way that is accessible to biological science students
- places bioinformatics in the context of evolutionary biology, including population genetics, molecular evolution, molecular phylogenetics, and their applications
- features end-of-chapter problems and self-tests to help students synthesize the materials and apply their understanding
- is accompanied by a dedicated website - [www.blackwellpublishing.com/higgs](http://www.blackwellpublishing.com/higgs) - containing downloadable sequences, links to web resources, answers to self-test questions, and all artwork in downloadable format (artwork also available to instructors on CD-ROM).

This important textbook will equip readers with a thorough understanding of the quantitative methods used in the analysis of molecular evolution, and will be essential reading for advanced undergraduates, graduates,

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