Bioinformatics: Sequence Alignment and Markov Models PDF
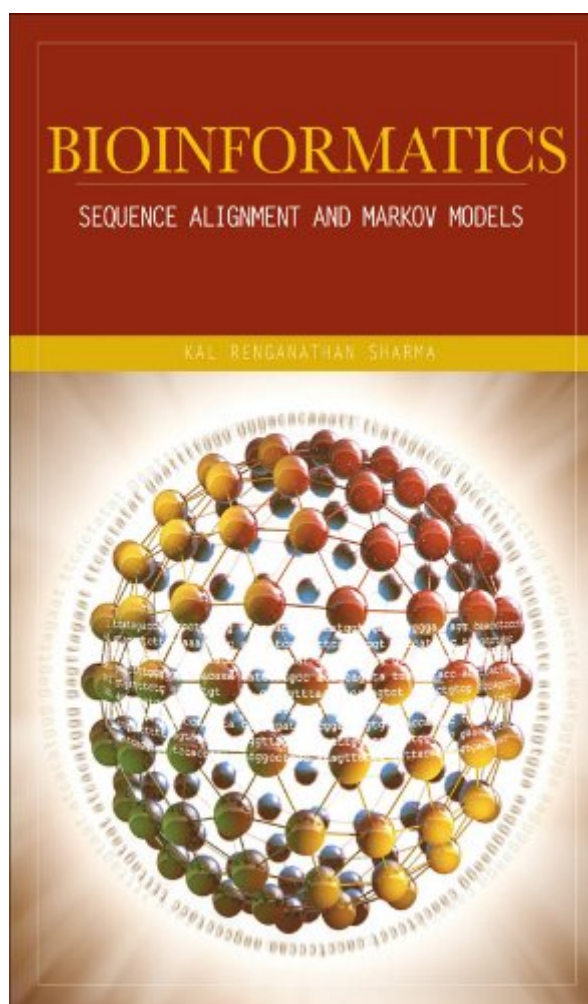

Bioinformatics: Sequence Alignment and Markov Models by Kal Sharma ISBN B001FR2TEM

GET FULLY UP-TO-DATE ON BIOINFORMATICS-THE TECHNOLOGY OF THE 21ST CENTURY

Bioinformatics showcases the latest developments in the field along with all the foundational information you'll need. It provides in-depth coverage of a wide range of autoimmune disorders and detailed analyses of suffix trees, plus late-breaking advances regarding biochips and genomes.

Featuring helpful gene-finding algorithms, *Bioinformatics* offers key information on sequence alignment, HMMs, HMM applications, protein secondary structure, microarray techniques, and drug discovery and development. Helpful diagrams accompany mathematical equations

throughout, and exercises appear at the end of each chapter to facilitate self-evaluation.

This thorough, up-to-date resource features:

- Worked-out problems illustrating concepts and models
- End-of-chapter exercises for self-evaluation
- Material based on student feedback
- Illustrations that clarify difficult math problems
- A list of bioinformatics-related websites

Bioinformatics covers:

- Sequence representation and alignment
- Hidden Markov models
- Applications of HMMs
- Gene finding
- Protein secondary structure prediction
- Microarray techniques
- Drug discovery and development
- Internet resources and public domain databases

Bioinformatics: Sequence Alignment and Markov Models Review

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