

Basant K. Tiwary

Bioinformatics and Computational Biology

A Primer for Biologists

Bioinformatics An Introduction Computational Biology

Röbbe Wünschiers



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Bioinformatics Jeremy Ramsden, 2010-10-22 The field of bioinformatics continues to develop energetically This second edition covers new findings using the formula of the original text It is self contained bringing together the multiple disciplines necessary for a profound grasp of the field

Introduction to Bioinformatics Anna Tramontano, 2006-12-06 Guiding readers from the elucidation and analysis of a genomic sequence to the prediction of a protein structure and the identification of the molecular function Introduction to Bioinformatics describes the rationale and limitations of the bioinformatics methods and tools that can help solve biological problems Requiring only a limited mathematical and statistical background the book shows how to efficiently apply these approaches to biological data and evaluate the resulting information The author an expert bioinformatics researcher first addresses the ways of storing and retrieving the enormous amount of biological data produced every day and the methods of decrypting the information encoded by a genome She then covers the tools that can detect and exploit the evolutionary and functional relationships among biological elements Subsequent chapters illustrate how to predict the three dimensional structure of a protein The book concludes with a discussion of the future of bioinformatics Even though the future will undoubtedly offer new tools for tackling problems most of the fundamental aspects of bioinformatics will not change This resource provides the essential information to understand bioinformatics methods ultimately facilitating in the solution of biological problems

Bioinformatics: An Introduction Jeremy J. Ramsden, 2012-12-06 An Introduction to Bioinformatics is intended to be a complete study companion for the advanced undergraduate or beginning graduate student It is self contained in the sense that whatever the starting point may be the reader will gain insight into bioinformatics Underlying the work is the belief that bioinformatics is a kind of metaphoric lens through which the entire field of biology can be brought into focus admittedly as yet imperfect and understood in a unified way Reflecting the highly incomplete present state of the field emphasis is placed on the underlying fundamentals and acquisitions of a broad and comprehensive grasp of the field as a whole Bioinformatics is interpreted as the application of information science to biology in which it plays a fundamental and all pervasive role This interpretation enables a remarkably unified view of the entire field of biology to be taken and hence offers an excellent entry point into the life sciences for those for whom biology is unfamiliar

An Introduction to Bioinformatics Algorithms Neil C. Jones, Pavel Pevzner, 2004

Introduction to Computational Biology Michael S. Waterman, 2018-05-02 Biology is in the midst of a era yielding many significant discoveries and promising many more Unique to this era is the exponential growth in the size of information packed databases Inspired by a pressing need to analyze that data Introduction to Computational Biology explores a new area of expertise that emerged from this fertile field the combination of biological and information sciences This introduction describes the mathematical structure of biological data especially from sequences and chromosomes After a brief survey of molecular biology it studies restriction maps of DNA rough landmark maps of the underlying sequences and

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