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Biological Data Analysis A Practical Approach

**Daniel P. Berrar, Werner
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Biological Data Analysis John C. Fry, 1993 Many biologists remain unfamiliar with statistical analysis and modelling yet need to apply these techniques increasingly in their research This volume describes how to analyze biological data with commonly available software packages without making errors which can invalidate results Practical guidance is provided for planning the correct strategy for a variety of different statistical approaches and modelling problems and interpreting the results Many examples of computer commands and output are given to illustrate the different analytical approaches *Biological Data Analysis A Practical Approach* has been designed specifically to allow researchers with only a minimal knowledge of statistics to understand a variety of statistical methods and apply them directly The provision of data sets from several biological disciplines will make this book useful to all types of biologists

RNA-seq Data Analysis Eija Korpelainen, Jarno Tuimala, Panu Somervuo, Mikael Huss, Garry Wong, 2014-09-19 The State of the Art in Transcriptome Analysis RNA sequencing RNA seq data offers unprecedented information about the transcriptome but harnessing this information with bioinformatics tools is typically a bottleneck RNA seq Data Analysis A Practical Approach enables researchers to examine differential expression at gene exon and transcript level

A Practical Approach to Microarray Data Analysis Daniel P. Berrar, Werner Dubitzky, Martin Granzow, 2007-05-08 In the past several years DNA microarray technology has attracted tremendous interest in both the scientific community and in industry With its ability to simultaneously measure the activity and interactions of thousands of genes this modern technology promises unprecedented new insights into mechanisms of living systems Currently the primary applications of microarrays include gene discovery disease diagnosis and prognosis drug discovery pharmacogenomics and toxicological research toxicogenomics Typical scientific tasks addressed by microarray experiments include the identification of coexpressed genes discovery of sample or gene groups with similar expression patterns identification of genes whose expression patterns are highly differentiating with respect to a set of discerned biological entities e g tumor types and the study of gene activity patterns under various stress conditions e g chemical treatment More recently the discovery modeling and simulation of regulatory gene networks and the mapping of expression data to metabolic pathways and chromosome locations have been added to the list of scientific tasks that are being tackled by microarray technology Each scientific task corresponds to one or more so called data analysis tasks Different types of scientific questions require different sets of data analytical techniques Broadly speaking there are two classes of elementary data analysis tasks predictive modeling and pattern detection Predictive modeling tasks are concerned with learning a classification or estimation function whereas pattern detection methods screen the available data for interesting previously unknown regularities or relationships

Plant Cell Biology William V Dashek, 2010-03-09 While there are a few plant cell biology books that are currently available these are expensive methods oriented monographs The present volume is a textbook for upper undergraduate and beginning graduate students This textbook stresses concepts and is

inquiry oriented To this end there is extensive use of original research literature As we live in an era of literature explosion one must be selective These judgements will naturally vary with each investigator Input was sought from colleagues in deciding the literature to include In addition to provision of select research literature this volume presents citations and summaries of certain laboratory methods In this connection the textbook stresses quantitative data to enhance the student s analytical abilities Thus the volume contains computer spread sheets and references to statistical packages e g Harvard Graphics and Statistica

Protein Structure Prediction : A Practical Approach Michael J. E. Sternberg,1996-11-28 The three dimensional structure of proteins is a key factor in their biological activity There is an increasing need to be able to predict the structure of a protein once its amino acid sequence is known this book presents practical methods of achieving that ambitious aim using the latest computer modelling algorithms The prediction of the three dimensional structure of a protein from its sequence is a problem faced by an ever increasing number of biological scientists as they strive to utilize genetic information The increasing sizes of the sequence and structural databases the improvements in computing power and the deeper understanding of the principles of protein structure have led to major developments in the field in the last few years This book presents practical computer based methods using the latest computer modelling algorithms

Describing Species Judith E. Winston,1999 A basic practical manual for the process of describing new species this desperately needed desk reference and guide to nomenclatural procedure and taxonomic writing serves as a Strunk White of species description covering both botanical and zoological codes of nomenclature

RNA-Protein Interactions : A Practical Approach Christopher W.J. Smith,1998-07-09 RNA protein interactions play a fundamental role in gene expression and protein synthesis Recent research into the role of RNA in cells has elucidated many more vital interactions with proteins This book provides an up to date and comprehensive guide to a wide range of laboratory procedures to investigate the interactions between RNA and proteins RNA protein interactions play a vital role in gene transcription and protein expression Interactions such as the synthesis of mRNA by RNA polymerases to the essential modification of RNA by the proteins of the spliceosome complex and the highly catalytic action of the ribosome in protein synthesis are established as being fundamental to the function of RNA Recent research into for example the role of RNA as a catalyst has elucidated many more interactions with proteins that are vital to cell function RNA Protein Interactions A Practical Approach provides a clear and comprehensive guide to the experimental procedures used in studying RNA protein interactions The approaches covered range from those initially used to detect a novel RNA protein interaction various biochemical and genetic approaches to purifying and cloning RNA binding proteins through to methods for an in depth analysis of the structural basis of the interaction The volume includes a number of procedures that have not previously been covered in this type of manual These include the production of site specifically modified RNAs by enzymatic and chemical methods and in vivo screening for novel RNA protein interactions in yeast and E coli This is the first volume to gather in one place this wide array of approaches for

studying RNA protein interactions As is customary for the Practical Approach series the writing is characterized by a clear explanatory style with many detailed protocols This informative book will be a valuable aid to laboratory workers in biochemistry and molecular biology graduate students postdoctoral and senior scientists whose research encompasses this field *Mathematical Models of Plant-Herbivore Interactions* Zhilan Feng,Donald DeAngelis,2017-09-07 Mathematical Models of Plant Herbivore Interactions addresses mathematical models in the study of practical questions in ecology particularly factors that affect herbivory including plant defense herbivore natural enemies and adaptive herbivory as well as the effects of these on plant community dynamics The result of extensive research on the use of mathematical modeling to investigate the effects of plant defenses on plant herbivore dynamics this book describes a toxin determined functional response model TDFRM that helps explains field observations of these interactions This book is intended for graduate students and researchers interested in mathematical biology and ecology *Chromatin* Ralf Blossey,2017-08-04 An invaluable resource for computational biologists and researchers from other fields seeking an introduction to the topic Chromatin Structure Dynamics Regulation offers comprehensive coverage of this dynamic interdisciplinary field from the basics to the latest research Computational methods from statistical physics and bioinformatics are detailed whenever possible without lengthy recourse to specialized techniques *Python for Bioinformatics* Sebastian Bassi,2017-08-07 In today s data driven biology programming knowledge is essential in turning ideas into testable hypothesis Based on the author s extensive experience Python for Bioinformatics Second Edition helps biologists get to grips with the basics of software development Requiring no prior knowledge of programming related concepts the book focuses on the easy to use yet powerful Python computer language This new edition is updated throughout to Python 3 and is designed not just to help scientists master the basics but to do more in less time and in a reproducible way New developments added in this edition include NoSQL databases the Anaconda Python distribution graphical libraries like Bokeh and the use of Github for collaborative development

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