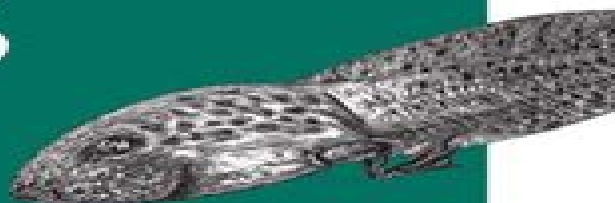


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Beginning Perl for Bioinformatics James Tisdall, 2001-10-22 With its highly developed capacity to detect patterns in data Perl has become one of the most popular languages for biological data analysis But if you re a biologist with little or no programming experience starting out in Perl can be a challenge Many biologists have a difficult time learning how to apply the language to bioinformatics The most popular Perl programming books are often too theoretical and too focused on computer science for a non programming biologist who needs to solve very specific problems Beginning Perl for Bioinformatics is designed to get you quickly over the Perl language barrier by approaching programming as an important new laboratory skill revealing Perl programs and techniques that are immediately useful in the lab Each chapter focuses on solving a particular bioinformatics problem or class of problems starting with the simplest and increasing in complexity as the book progresses Each chapter includes programming exercises and teaches bioinformatics by showing and modifying programs that deal with various kinds of practical biological problems By the end of the book you ll have a solid understanding of Perl basics a collection of programs for such tasks as parsing BLAST and GenBank and the skills to take on more advanced bioinformatics programming Some of the later chapters focus in greater detail on specific bioinformatics topics This book is suitable for use as a classroom textbook for self study and as a reference The book covers Programming basics and working with DNA sequences and strings Debugging your code Simulating gene mutations using random number generators Regular expressions and finding motifs in data Arrays hashes and relational databases Regular expressions and restriction maps Using Perl to parse PDB records annotations in GenBank and BLAST output

Beginning Perl for Bioinformatics James Tisdall, 2001 This book shows biologists with little or no programming experience how to use Perl the ideal language for biological data analysis Each chapter focuses on solving a particular problem or class of problems so you ll finish the book with the skills to tackle more advanced bioinformatics programming

Bioinformatics Thomas Dandekar, Meik Kunz, 2023-03-02 This book offers a gripping introduction to the fastest growing field of biology with easy to follow examples and a well prepared appendix for the reader to cook up and experience everything right away The book gets the reader started with the basics such as how to easily find sequence information and then analyze it In further chapters the authors go into the various analysis options from RNA DNA and proteins to entire metabolic pathways Exciting examples from biology are chosen in each chapter to illustrate the analysis Each chapter concludes with an exercise section that immediately puts what has been learned to use The subject of this book is a must for any biology student whether undergraduate or graduate as bioinformatics is now unearthing amazing insights into the molecular basis of all living things Computer science students and other students from related sciences will get a good introduction to bioinformatics as biology and current topics e g AI are systematically introduced step by step alongside the software Discover the key to life together with the authors and learn to understand the language of life This book is a translation of the original German 2nd edition

Bioinformatik by Thomas Dandekar and Meik Kunz published by Springer Verlag GmbH Germany part of Springer Nature in 2021 The translation was done with the help of artificial intelligence machine translation by the service DeepL.com A subsequent human revision was done primarily in terms of content so that the book will read stylistically differently from a conventional translation Springer Nature works continuously to further the development of tools for the production of books and on the related technologies to support the authors

Mastering Perl for Bioinformatics James Tisdall, 2003-09-25

Historically programming hasn't been considered a critical skill for biologists But now with access to vast amounts of biological data contained in public databases programming skills are increasingly in strong demand in biology research and development Perl with its highly developed capacities in string handling text processing networking and rapid prototyping has emerged as the programming language of choice for biological data analysis Mastering Perl for Bioinformatics covers the core Perl language and many of its module extensions presenting them in the context of biological data and problems of pressing interest to the biological community This book along with Beginning Perl for Bioinformatics forms a basic course in Perl programming This second volume finishes the basic Perl tutorial material references complex data structures object oriented programming use of modules all presented in a biological context and presents some advanced topics of considerable interest in bioinformatics The range of topics covered in Mastering Perl for Bioinformatics prepares the reader for enduring and emerging developments in critical areas of bioinformatics programming such as Gene finding String alignment Methods of data storage and retrieval SML and databases Modeling of networks graphs and Petri nets Graphics Tk Parallelization Interfacing with other programming languages Statistics PDL Protein structure determination Biological models of computation DNA Computers Biologists and computer scientists who have conquered the basics of Perl and are ready to move even further in their mastery of this versatile language will appreciate the author's well balanced approach to applying Perl's analytical abilities to the field of bioinformatics Full of practical examples and real world biological problem solving this book is a must for any reader wanting to move beyond beginner level Perl in bioinformatics

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