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Bioinformatics Programming in Python Ruediger-Marcus Flaig,2008-03-31 This first introductory book designed to train novice programmers is based on a student course taught by the author and has been optimized for biology students without previous experience in programming By interspersing theory chapters with numerous small and large programming exercises the author quickly shows readers how to do their own programming and throughout uses anecdotes and real life examples from the biosciences to spice up the text This practical book thus teaches essential programming skills for life scientists who want or need to write their own bioinformatics software tools Mastering Python for Bioinformatics Ken Youens-Clark,2021-05-05 Life scientists today urgently need training in bioinformatics skills Too many bioinformatics programs are poorly written and barely maintained usually by students and researchers who ve never learned basic programming skills This practical guide shows postdoc bioinformatics professionals and students how to exploit the best parts of Python to solve problems in biology while creating documented tested reproducible software Ken Youens Clark author of Tiny Python Projects Manning demonstrates not only how to write effective Python code but also how to use tests to write and refactor scientific programs You ll learn the latest Python features and tools including linters formatters type checkers and tests to create documented and tested programs You ll also tackle 14 challenges in Rosalind a problem solving platform for learning bioinformatics and programming Create command line Python programs to document and validate parameters Write tests to verify refactor programs and confirm they re correct Address bioinformatics ideas using Python data structures and modules such as Biopython Create reproducible shortcuts and workflows using makefiles Parse essential bioinformatics file formats such as FASTA and FASTQ Find patterns of text using regular expressions Use higher order

functions in Python like filter map and reduce

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

Reproducible Bioinformatics with Python Ken Youens-Clark, 2021-08-17 Life scientists today urgently need training in bioinformatics skills Too many bioinformatics programs are poorly written and barely maintained usually by students and postdoc researchers who've never learned basic programming skills This practical guide shows how to exploit the best parts of Python for solving problems in biology while also creating documented tested reproducible software Ken Youens Clark author of Tiny Python Projects Manning demonstrates how to write effective Python code and how to use tests to write and refactor scientific programs You'll learn the latest Python features and tools such as linters formatters type checkers and tests to write documented and tested programs Create command line Python programs that document and validate parameters Write tests to verify refactor programs and confirm they're correct Address bioinformatics ideas using Python data structures strings lists and sets and modules such as Biopython Create reproducible shortcuts and workflows using makefiles Parse essential bioinformatics file formats such as FASTA FASTQ and SwissProt Find patterns of text using regular expressions Use higher order functions in Python like filter and map

Python Programming for Biology Tim J. Stevens, Wayne Boucher, 2015-02-12 Do you have a biological question that could be readily answered by computational techniques but little experience in programming Do you want to learn more about the core techniques used in computational biology and bioinformatics Written in an accessible style this guide provides a foundation for both newcomers to computer programming and those interested in learning more about computational biology The chapters guide the reader through a complete beginners course to programming in Python with an introduction to computing jargon descriptions of core bioinformatics methods with working Python examples scientific computing techniques including image analysis statistics and machine learning This book also functions as a language reference written in straightforward English covering the most common Python language elements and a glossary of computing and biological terms This title will teach undergraduates postgraduates and professionals working in the life sciences how to program with Python a powerful flexible and easy to use language

Bioinformatics with Python Cookbook Tiago Antao, 2018-11-30 Discover modern next generation sequencing libraries from Python ecosystem to analyze large amounts of biological data Key Features Perform complex bioinformatics analysis using the most important Python libraries and applications Implement next generation sequencing

metagenomics automating analysis population genetics and more Explore various statistical and machine learning techniques for bioinformatics data analysis Book Description Bioinformatics is an active research field that uses a range of simple to advanced computations to extract valuable information from biological data This book covers next generation sequencing genomics metagenomics population genetics phylogenetics and proteomics You ll learn modern programming techniques to analyze large amounts of biological data With the help of real world examples you ll convert analyze and visualize datasets using various Python tools and libraries This book will help you get a better understanding of working with a Galaxy server which is the most widely used bioinformatics web based pipeline system This updated edition also includes advanced next generation sequencing filtering techniques You ll also explore topics such as SNP discovery using statistical approaches under high performance computing frameworks such as Dask and Spark By the end of this book you ll be able to use and implement modern programming techniques and frameworks to deal with the ever increasing deluge of bioinformatics data What you will learn Learn how to process large next generation sequencing NGS datasets Work with genomic dataset using the FASTQ BAM and VCF formats Learn to perform sequence comparison and phylogenetic reconstruction Perform complex analysis with proteomics data Use Python to interact with Galaxy servers Use High performance computing techniques with Dask and Spark Visualize protein dataset interactions using Cytoscape Use PCA and Decision Trees two machine learning techniques with biological datasets Who this book is for This book is for Data data Scientistsscientists Bioinformatics bioinformatics analysts researchers and Python developers who want to address intermediate to advanced biological and bioinformatics problems using a recipe based approach Working knowledge of the Python programming language is expected

Python for the Life Sciences Alexander Lancaster, Gordon Webster, 2019-09-28 Treat yourself to a lively intuitive and easy to follow introduction to computer programming in Python The book was written specifically for biologists with little or no prior experience of writing code with the goal of giving them not only a foundation in Python programming but also the confidence and inspiration to start using Python in their own research Virtually all of the examples in the book are drawn from across a wide spectrum of life science research from simple biochemical calculations and sequence analysis to modeling the dynamic interactions of genes and proteins in cells or the drift of genes in an evolving population Best of all Python for the Life Sciences shows you how to implement all of these projects in Python one of the most popular programming languages for scientific computing If you are a life scientist interested in learning Python to jump start your research this is the book for you What You ll Learn Write Python scripts to automate your lab calculations Search for important motifs in genome sequences Use object oriented programming with Python Study mining interaction network data for patterns Review dynamic modeling of biochemical switches Who This Book Is For Life scientists with little or no programming experience including undergraduate and graduate students postdoctoral researchers in academia and industry medical professionals and teachers lecturers A comprehensive introduction to using Python for computational biology A lovely book with humor and

perspective John Novembre Associate Professor of Human Genetics University of Chicago and MacArthur Fellow Fun entertaining witty and darn useful A magical portal to the big data revolution Sandro Santagata Assistant Professor in Pathology Harvard Medical School Alex and Gordon s enthusiasm for Python is contagious Glenys Thomson Professor of Integrative Biology University of California Berkeley *Computing for Biologists* Ran Libeskind-Hadas, Eliot Bush, 2014-09-22 Computing is revolutionizing the practice of biology This book which assumes no prior computing experience provides students with the tools to write their own Python programs and to understand fundamental concepts in computational biology and bioinformatics Each major part of the book begins with a compelling biological question followed by the algorithmic ideas and programming tools necessary to explore it the origins of pathogenicity are examined using gene finding the evolutionary history of sex determination systems is studied using sequence alignment and the origin of modern humans is addressed using phylogenetic methods In addition to providing general programming skills this book explores the design of efficient algorithms simulation NP hardness and the maximum likelihood method among other key concepts and methods Easy to read and designed to equip students with the skills to write programs for solving a range of biological problems the book is accompanied by numerous programming exercises available at www.cs.hmc.edu/CFB

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