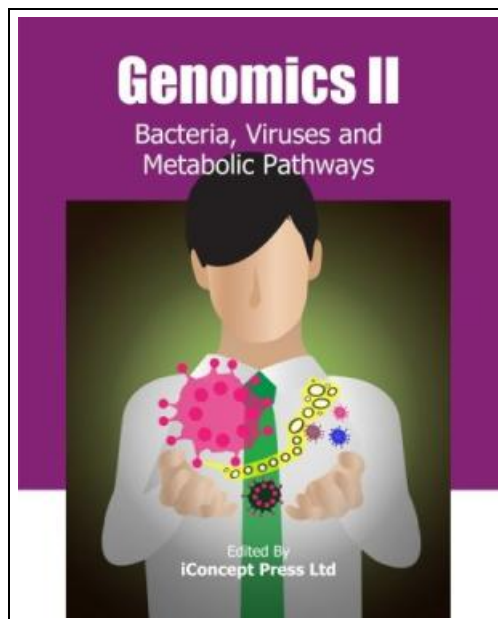


Genomics II: Bacteria, Viruses and Metabolic Pathways



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Reviews

Extensive information! Its this sort of great read through. It is amongst the most incredible book i have go through. I realized this publication from my i and dad suggested this book to understand.

(Prof. Devon Bernhard PhD)

GENOMICS II: BACTERIA, VIRUSES AND METABOLIC PATHWAYS



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Createspace, United States, 2013. Paperback. Book Condition: New. 254 x 203 mm. Language: English . Brand New Book ***** Print on Demand *****.Genomics is the study of the genomes of organisms. The field includes intensive efforts to determine the entire DNA sequence of organisms and fine-scale genetic mapping efforts. It is a discipline in genetics that applies recombinant DNA, DNA sequencing methods, and bioinformatics to sequence, assemble, and analyze the function and structure of genomes. Genomics II - Bacteria, Viruses and Metabolic Pathways is the second volume of our Genomics series. There are totally three volumes in this series. Chapter 1 describes an analysis and statistical scoring approach for cellular assay data based on single-cell information. In Chapter 2, the concept of metabolic pathways analysis is introduced. The mathematic principle of extreme pathway and elementary flux mode are compared. Chapter 3 is dedicated to the Pathway- and Network-based analysis of the high-throughput genomic data. The author introduced Reactome FI Cytoscape plugin that can construct a network based on the list of genes of interest, cluster the constructed network, and annotate network modules based on pathways and Gene Ontology terms. Chapter 4 provides a review of microarray and RNA-seq techniques for high-throughput gene expression measurements, discusses the strategies and issues of high-level analysis on gene expression data, and introduces a new algorithm for analyzing microarray data. Chapter 5 summarizes our current understanding of the intracellular defenses by APOBEC family against invading nucleic acids including endogenous retroelements that make up more than 40 of the mammalian genome. Chapter 6 discusses immunoinformatics software that can be employed to study the evolution of antigenic epitopes. Chapter 7 discusses the integration of retroviral genome into host DNA, which is a critical step in the life cycle of a retrovirus. The authors developed an assay using...



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