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RNA Bioinformatics Ernesto Picardi, 2016-09-24 This volume provides an overview of RNA bioinformatics methodologies, including basic strategies to predict secondary and tertiary structures, and novel algorithms based on massive RNA sequencing. Interest in RNA bioinformatics has rapidly increased thanks to the recent high-throughput sequencing technologies allowing scientists to investigate complete transcriptomes at single nucleotide resolution. Adopting advanced computational technics, scientists are now able to conduct more in-depth studies and present them to you in this book. Written in the highly successful *Methods of Molecular Biology* series format, chapters include introductions to their respective topics, lists of the necessary materials and equipment, step-by-step, readily reproducible bioinformatics protocols, and key tips to avoid known pitfalls. Authoritative and practical, *RNA Bioinformatics* seeks to aid scientists in the further study of bioinformatics and computational biology of RNA.

Advocate of Peace and Universal Brotherhood ,1900

Includes the Annual report of the American Peace Society.

The Pangenome Hervé Tettelin, Duccio

Medini, 2020-04-30 This open access book offers the first comprehensive account of the pan-genome concept and its manifold implications. The realization that the genetic repertoire of a biological species always encompasses more than the genome of each individual is one of the earliest examples of big data in biology that opened biology to the unbounded. The study of genetic variation observed within a species challenges existing views and has profound consequences for our understanding of the fundamental mechanisms underpinning bacterial biology and evolution. The underlying rationale extends well beyond the initial prokaryotic focus to all kingdoms of life and evolves into

similar concepts for metagenomes, phenomes and epigenomes. The book's respective chapters address a range of topics, from the serendipitous emergence of the pan-genome concept and its impacts on the fields of microbiology, vaccinology and antimicrobial resistance, to the study of microbial communities, bioinformatic applications and mathematical models that tie in with complex systems and economic theory. Given its scope, the book will appeal to a broad readership interested in population dynamics, evolutionary biology and genomics.

Bacterial Physiology and Metabolism Byung Hong Kim, Geoffrey Michael Gadd, 2008-02-21 Recent determination of genome sequences for a wide range of bacteria has made in-depth knowledge of prokaryotic metabolic function essential in order to give biochemical, physiological, and ecological meaning to the genomic information. Clearly describing the important metabolic processes that occur in prokaryotes under different conditions and in different environments, this advanced text provides an overview of the key cellular processes that determine bacterial roles in the environment, biotechnology, and human health. Prokaryotic structure is described as well as the means by which nutrients are transported into cells across membranes. Glucose metabolism through glycolysis and the TCA cycle are discussed, as well as other trophic variations found in prokaryotes, including the use of organic compounds, anaerobic fermentation, anaerobic respiratory processes, and photosynthesis. The regulation of metabolism through control of gene expression and control of the activity of enzymes is also covered, as well as survival mechanisms used under starvation conditions.

The Universal Chronicle Donald Davis Eddy, 1979
Quadratic Forms and Their Applications Eva Bayer-

Fluckiger,Conference on Quadratic Forms and Their Applications,David Lewis,Andrew Ranicki,Conference on Quadratic Forms and Their Applications Staff,2000 This volume outlines the proceedings of the conference on 'Quadratic Forms and Their Applications' held at University College Dublin. It includes survey articles and research papers ranging from applications in topology and geometry to the algebraic theory of quadratic forms and its history. Various aspects of the use of quadratic forms in algebra, analysis, topology, geometry, and number theory are addressed. Special features include the first published proof of the Conway-Schneeberger Fifteen Theorem on integer-valued quadratic forms and the first English-language biography of Ernst Witt, founder of the theory of quadratic forms.

Proceedings of the XSEDE16 Conference on Diversity, Big Data, and Science at Scale Kelly Gaither,2016-07-17 XSEDE16: Diversity, Big Data, and Science at Scale Jul 17, 2016-Jul 21, 2016 Miami, USA. You can view more information about this proceeding and all of ACM's other published conference proceedings from the ACM Digital Library: <http://www.acm.org/dl>.

Proceedings of the XIII International Symposium SymOrg 2012: Innovative Management and Business Performance ,2012-06-03

Metals and Related Substances in Drinking Water Prosun Bhattacharya,Ingegerd Rosborg,Arifin Sandhi,Colin Hayes,Maria Joao Benoliel,2011-11-25 Part of Metals and Related Substances in Drinking Water Set - buy all five books together to save over 30%! Metals and Related Substances in Drinking Water comprises the proceedings of COST Action 637 - METEAU, held in Kristianstad, Sweden, October 13-15, 2010. This book collates the understanding of the various factors which control metals and related

substances in drinking water with an aim to minimize environmental impacts. Metals and Related Substances in Drinking Water: Provides an overview of knowledge on metals and related substances in drinking water. Promotes good practice in controlling metals and related substances in drinking water. Helps to determining the environmental and socio-economic impacts of control measures through public participation Introduces the importance of mineral balance in drinking water especially when choosing treatment methods Shares practitioner experience. The proceedings of this international conference contain many state-of-the-art presentations by leading researchers from across the world. They are of interest to water sector practitioners, regulators, researchers and engineers.

Modelling, Simulation and Control of Thermal Energy Systems Kwang Y. Lee,Damian Flynn,Hui Xie,Li Sun,2020-11-03 Faced with an ever-growing resource scarcity and environmental regulations, the last 30 years have witnessed the rapid development of various renewable power sources, such as wind, tidal, and solar power generation. The variable and uncertain nature of these resources is well-known, while the utilization of power electronic converters presents new challenges for the stability of the power grid. Consequently, various control and operational strategies have been proposed and implemented by the industry and research community, with a growing requirement for flexibility and load regulation placed on conventional thermal power generation. Against this background, the modelling and control of conventional thermal engines, such as those based on diesel and gasoline, are experiencing serious obstacles when facing increasing environmental concerns. Efficient control that can fulfill the requirements of high efficiency, low pollution, and long durability is an emerging requirement. The

modelling, simulation, and control of thermal energy systems are key to providing innovative and effective solutions. Through applying detailed dynamic modelling, a thorough understanding of the thermal conversion mechanism(s) can be achieved, based on which advanced control strategies can be designed to improve the performance of the thermal energy system, both in economic and environmental terms. Simulation studies and test beds are also of great significance for these research activities prior to proceeding to field tests. This Special Issue will contribute a practical and comprehensive forum for exchanging novel research ideas or empirical practices that bridge the modelling, simulation, and control of thermal energy systems. Papers that analyze particular aspects of thermal energy systems, involving, for example, conventional power plants, innovative thermal power generation, various thermal engines, thermal energy storage, and fundamental heat transfer management, on the basis of one or more of the following topics, are invited in this Special Issue: • Power plant modelling, simulation, and control; • Thermal engines; • Thermal energy control in building energy systems; • Combined heat and power (CHP) generation; • Thermal energy storage systems; • Improving thermal comfort technologies; • Optimization of complex thermal systems; • Modelling and control of thermal networks; • Thermal management of fuel cell systems; • Thermal control of solar utilization; • Heat pump control; • Heat exchanger control.

Cancer Evolution Charles Swanton, 2017 Tumor progression is driven by mutations that confer growth advantages to different subpopulations of cancer cells. As a tumor grows, these subpopulations expand, accumulate new mutations, and are subjected to selective pressures from the environment, including anticancer interventions.

This process, termed clonal evolution, can lead to the emergence of therapy-resistant tumors and poses a major challenge for cancer eradication efforts. Written and edited by experts in the field, this collection from Cold Spring Harbor Perspectives in Medicine examines cancer progression as an evolutionary process and explores how this way of looking at cancer may lead to more effective strategies for managing and treating it. The contributors review efforts to characterize the subclonal architecture and dynamics of tumors, understand the roles of chromosomal instability, driver mutations, and mutation order, and determine how cancer cells respond to selective pressures imposed by anticancer agents, immune cells, and other components of the tumor microenvironment. They compare cancer evolution to organismal evolution and describe how ecological theories and mathematical models are being used to understand the complex dynamics between a tumor and its microenvironment during cancer progression. The authors also discuss improved methods to monitor tumor evolution (e.g., liquid biopsies) and the development of more effective strategies for managing and treating cancers (e.g., immunotherapies). This volume will therefore serve as a vital reference for all cancer biologists as well as anyone seeking to improve clinical outcomes for patients with cancer.

Molecular Basis of Stage Conversion in Apicomplexan Parasites Maria E. Francia, Mathieu Gissot, Katarzyna Kinga Modrzyńska, Mattie Christine Pawlowic, 2021-06-08

Select Thermodynamic Models for Process Simulation Jean-Charles de Hemptinne, Jean-Marie Ledanois, 2012 The selection of the most adequate thermodynamic model in a process simulation is an issue that most process engineer has to face sooner or later. This book, conceived as a practical guide, aims at providing

adequate answers by analysing the questions to be looked at. The analysis (first chapter) yields three keys that are further discussed in three different chapters. (1) A good understanding of the properties required in the process, and their method of calculation is the first key. The second chapter provides to that end in a synthetic manner the most important equations that are derived from the fundamental principles of thermodynamics. (2) An adequate description of the mixture, which is a combination of models and parameters, is the second key. The third chapter makes the link between components and models, both from a numerical (parameterisation) and physical (molecular interactions) point of view. Finally, (3) a correct view of the phase behaviour and trends in regard of the process conditions is the third key. The fourth chapter illustrates the phase behaviour and makes model recommendations for the most significant industrial systems. A decision tree is provided at the end of this chapter. In the last chapter, the key questions are reviewed for a number of typical processes. This book is intended for process engineers, who are not specialists of thermodynamics but are confronted with this kind of problems and need a reference book, as well as process engineering students who will find an original approach to thermodynamics, complementary of traditional lectures

Agrobacterium: From Biology to Biotechnology

Tzvi Tzfira, Vitaly Citovsky, 2007-12-25 Agrobacterium is a plant pathogen which causes the “crown-gall” disease, a neoplastic growth that results from the transfer of a well-defined DNA segment (“transferred DNA”, or “T-DNA”) from the bacterial Ti (tumor-inducing) plasmid to the host cell, its integration into the host genome, and the expression of oncogenes contained on the T-DNA. The molecular machinery, needed for T-DNA generation and transport into

the host cell and encoded by a series of chromosomal (chv) and Ti-plasmid virulence (vir) genes, has been the subject of numerous studies over the past several decades. Today, *Agrobacterium* is the tool of choice for plant genetic engineering with an ever expanding host range that includes many commercially important crops, flowers, and tree species. Furthermore, its recent application for the genetic transformation of non-plant species, from yeast to cultivated mushrooms and even to human cells, promises this bacterium a unique place in the future of biotechnological applications. The book is a comprehensive volume describing *Agrobacterium*'s biology, interactions with host species, and uses for genetic engineering.

Gene Prediction Martin Kollmar, 2019-05-19 This volume introduces software used for gene prediction with focus on eukaryotic genomes. The chapters in this book describe software and web server usage as applied in common use-cases, and explain ways to simplify re-annotation of long available genome assemblies. Written in the highly successful *Methods in Molecular Biology* series format, chapters include introductions to their respective topics, lists of the necessary computational requirements, step-by-step, readily reproducible computational protocols, and tips on troubleshooting and avoiding known pitfalls. Cutting-edge and thorough, *Gene Prediction: Methods and Protocols* is a valuable resource for researchers and research groups working on the assembly and annotation of single species or small groups of species. Chapter 3 is available open access under a CC BY 4.0 license via link.springer.com.

A System of medicine, by many writers v. 9, 1905/11 Sir Thomas Clifford Allbutt, 1911

A System of Medicine Thomas Clifford Allbutt, Sir Humphry Davy Rolleston, 1911

Neuro-Urology Research Anne M.J.

Verstegen, 2023-05-08 Neuro-Urology Research: A Comprehensive Overview describes the current status of the neuro-urology field including the latest discoveries, explains in detail some of the neuroscience tools that can be used when studying the neural control of the lower urinary tract, and discusses potential future directions for research. The first section, Neuroscience in Urology Research, reviews neuro-regulators and the important anatomical brain sites. It also provides an overview of voluntary versus reflex micturition control and describes how bladder physiology readout is a useful tool in research. The second section, Fundamental and Translational Neuro-Urology Research, discusses the translational potential of basic research for patients and the impact of neuro-urology research on clinical practice. Chapters in this section provide more insight into pathologies endemic to specific patient populations and areas for treatment development opportunities. The third section, Neurobiological Tools Applied to Neuro-Urology Research, supplements the 'A Quick Guide to the Neuroscience Toolbox' and introduces research techniques such as RNA sequencing and calcium imaging of neural activity. The fourth and final section, aptly titled Research Directions and Research Opportunities, covers research directions that remain underexplored or have high therapeutic potential. Research in neuro-urology presents a route to understanding how bladder function is controlled and how urinary continence is maintained. This book provides a platform for researchers to initiate collaborations on the many underexplored research topics and to consolidate knowledge, ultimately lending itself to a coordinated and multi-center research approach that will benefit the field. Presents a comprehensive overview of research studies in the field of neuro-urology Discusses

approaches to conducting neuro-urology research, including models of lower urinary tract dysfunction and methods of tracking bladder physiology Proposes areas of research opportunity that, if explored, could yield effective treatment options for those affected by lower urinary tract disorders Includes an introductory guide to useful research tools and the Neuroscience Toolbox, as well as chapters specifically about RNA sequencing techniques and imaging bladder-associated neural circuits

Wheat Rusts RA McIntosh,CR Wellings,RF

Park,1995-01-01 Although stem rust has been controlled by means of resistant cultivars, leaf and stripe rust continue as problems for many growing areas of the world. Wheat Rusts: An Atlas of Resistance Genes has been prepared by specialists from one of the leading international laboratories, and illustrates with colour photographs typical resistance phenotypes associated with most known genes for resistance to the three rust diseases of wheat. Relevant details for each gene include chromosome location, aspects of genetics and pathogen variation, the effects of environment on expression, origin, availability in genetic and breeding stocks, and use in agriculture. This atlas includes an introduction to host:pathogen genetics, methodologies for wheat rust research and breeding for resistance.

Low-Oxygen Stress in Plants Joost T. van

Dongen,Francesco Licausi,2014-01-17 During the last ten years, knowledge about the multitude of adaptive responses of plants to low oxygen stress has grown immensely. The oxygen sensor mechanism has been discovered, the knowledge about the interaction network of gene expression is expanding and metabolic adaptations have been described in detail. Furthermore, morphological changes were investigated and the regulative mechanisms

triggered by plant hormones or reactive oxygen species have been revealed. This book provides a broad overview of all these aspects of low oxygen stress in plants. It integrates knowledge from different disciplines such as molecular biology, biochemistry, ecophysiology and agricultural / horticultural sciences to comprehensively describe how plants cope with low oxygen stress and discuss its ecological and agronomical consequences. This book is written for plant scientists, biochemists and scientists in agriculture and ecophysiology.

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