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Ibrokhim Y. Abdurakhmonov

The Epidemiology, Diagnosis and Prevention of Infectious Diseases in Livestock Satoshi

Sekiguchi, Van Giap Nguyen, Anuwat Wiratsudakul, 2022-02-24

Class 3.2 Hydrolases VII Dietmar Schomburg, Ida Schomburg, 2003-06-18 The Springer Handbook of Enzymes provides concise data on some 5,000 enzymes sufficiently well characterized – and here is the second, updated edition. Their application in analytical, synthetic and biotechnology processes as well as in food industry, and for medicinal treatments is added. Data sheets are arranged in their EC-Number sequence. The new edition reflects considerable progress in enzymology: the total material has more than doubled, and the complete 2nd edition consists of 39 volumes plus Synonym Index. Starting in 2009, all newly classified enzymes are treated in Supplement Volumes.

Networks of Networks in Biology Narsis A. Kiani, David Gomez-Cabrero, Ginestra Bianconi, 2021-04

Introduces network inspired approaches for the analysis and integration of large, heterogeneous data sets in the life sciences.

Application of Clinical Bioinformatics Xiangdong Wang, Christian Baumgartner, Denis C.

Shields, Hong-Wen Deng, Jacques S Beckmann, 2016-03-31 This book elucidates how genetic, biological and medical information can be applied to the development of personalized healthcare, medication and therapies. Focusing on aspects of the development of evidence-based approaches in bioinformatics and computational medicine, including data integration, methodologies, tools and models for clinical and translational medicine, it offers an essential introduction to clinical bioinformatics for clinical researchers and physicians, medical students and teachers, and scientists working with human disease-based omics and bioinformatics. Dr. Xiangdong Wang is a distinguished Professor of Medicine. He is Director of Shanghai Institute of Clinical Bioinformatics, Director of Fudan University Center for Clinical Bioinformatics, Deputy Director of Shanghai Respiratory Research Institute, Director of Biomedical Research Center, Fudan University Zhongshan Hospital, Shanghai, China; Dr. Christian Baumgartner is a Professor of Health Care and Biomedical Engineering at Institute of Health Care Engineering with European Notified Body of Medical Devices, Graz University of Technology, Graz, Austria; Dr. Denis Shields is a Professor of Clinical Bioinformatics at Conway Institute, Belfield, Dublin, Ireland; Dr. Hong-Wen Deng is a Professor at Department of Biostatistics and Bioinformatics, Tulane University School of Public Health and Tropical Medicine, USA; Dr. Jacques S Beckmann is a Professor and Director of Section of Clinical Bioinformatics, Swiss Institute of Bioinformatics, Switzerland.

Applied Genomics of Foodborne Pathogens Xiangyu Deng, Henk C. den Bakker, Rene S.

Hendriksen, 2017-01-23 This book provides a timely and thorough snapshot into the emerging and fast evolving area of applied genomics of foodborne pathogens. Driven by the drastic advance of whole genome shot gun sequencing (WGS) technologies, genomics applications are becoming increasingly valuable and even essential in studying, surveying and controlling foodborne microbial pathogens. The vast opportunities brought by this trend are often at odds with the lack of bioinformatics know-how among food safety and public health professionals, since such expertise is not part of a typical food microbiology curriculum and skill set. Further complicating the challenge is the large and ever evolving body of bioinformatics tools that can obfuscate newcomers to this area. Although reviews, tutorials and books are not in short supply in the fields of bioinformatics and genomics, until now there has not been a comprehensive and customized source of information designed for and accessible to microbiologists interested in applying cutting-edge genomics in food safety and public health research. This book fills this void with a well-selected collection of topics, case studies, and bioinformatics tools contributed by experts at the forefront of foodborne pathogen genomics research.

Recent Advances in Veterinary Immunology Concepts and Methodology Armin Saalmueller, Dirk

Werling, Crystal L. Loving, Philip John Griebel, 2021-03-03 We acknowledge the initiation and support of this Research Topic by the International Union of Immunological Societies (IUIS). We hereby state publicly that the IUIS has had no editorial input in articles included in this Research Topic, thus ensuring that all aspects of this Research Topic are evaluated objectively, unbiased by any specific policy or opinion of the IUIS.

Computational genomics and structural bioinformatics in personalized medicines, volume II George

Priya Doss C, Thirumal Kumar D, Balu Kamaraj, 2023-11-06

Cardiovascular Diseases Related to Diabetes and Obesity Volume II Lu Cai, Ying Xin, Jamie Lynn

Young, Rajesh Mohanraj, 2022-11-14

Phylogenetics Ibrokhim Y. Abdurakhmonov, 2017-09-06 Phylogenetics aims to study the

evolutionary relatedness of living organisms in our planet. Its application is extended to the key areas such as evolution, classification and taxonomy of living organisms; ecology, diversity, and conservation biology of agrobiocenosis; monitoring of pathogen spread, outbreaks and source of transmissions, forensic analyses, etc. Historically, phylogenetics studies were prevalently based on morphological features of species that helped to classify the 'Tree of Life' on Earth. Modern phylogenetics studies, however, rely more heavily on DNA sequences. In this Phylogenetics book, we aimed to present

readers the latest developments in phylogenetics studies that highlight multi-kingdom systems, reticulated evolution and conservation biology of living organisms as well as 'omics'-based phylogenetics advances.

Metabolomics: From Fundamentals to Clinical Applications Alessandra Sussulini, 2017-01-28 This book provides a comprehensive view of metabolomics, from the basic concepts, through sample preparation and analytical methodologies, to data interpretation and applications in medicine. It is the first volume to cover metabolomics clinical applications while also emphasizing analytical and statistical features. Moreover, future trends and perspectives in clinical metabolomics are also presented. For researches already experienced in metabolomics, the book will be useful as an updated definitive reference. For beginners in the field and graduate students, the book will provide detailed information about concepts and experimental aspects in metabolomics, as well as examples and perspectives of applications of this strategy to clinical questions.

Proteomics and its Applications in Cancer Suman S. Thakur, Harsha Gowda, D. R. Mani, Bhaswati Chatterjee, 2022-01-11

The Neurobiology and Genetics of Gilles de la Tourette Syndrome: New Avenues Through Large-Scale Collaborative Projects Peristera Paschou, Kirsten R. Müller-Vahl, 2018-03-13 Gilles de la Tourette Syndrome (TS) is a common, albeit severely under-diagnosed, neuropsychiatric disorder that is caused by a complex genetic basis, interacting with environmental factors. High comorbidity rates with other neurodevelopmental disorders such as attention deficit/hyperactivity disorder and obsessive compulsive disorder raise the intriguing hypothesis of a shared etiological background. Abnormalities of corticostriatal-thalamic-cortical circuits (CSTC) and dysfunction of both dopamine and serotonin neurotransmitter systems are assumed to be associated with TS. Recently, multiple lines of evidence also point towards an important role of additional neurotransmitters such as histamine and glutamate. For a very long time, efforts to elucidate the etiology and pathophysiology of TS have been fragmented and hampered by low statistical power. Finally, after more than two decades of active research aiming to identify the etiology and pathophysiology of TS, we are on the verge of a new era, promising exciting and rapid discoveries in the field. Investigators from around the world, representing multiple disciplines and scientific approaches, are joining their efforts in large-scale initiatives supported both by European Union and US National funding agencies, such as the European-funded EMTICS, TACTICS, and TSGeneSEE consortia, the Marie Curie Initial Training Network TS-EUROTRAIN and the European Society for the Study of TS joining forces with the NIH-funded TSAICG, GGRI, and Tic

Genetics consortia. Importantly, all these initiatives are supported by TS patient support and advocacy groups. Multiple resources are being consolidated and coming together to serve the study of TS, including large well-characterized patient cohorts, and specialized epidemiological databases, such as the unique resource of the Netherlands Twin Register. This research topic showcases current large-scale collaborative efforts aiming to elucidate the genetic and neurobiological background of TS, through diverse approaches; from genomewide association studies aiming to identify common variants associated to the disorder to neuroimaging studies and animal models. Furthermore, current approaches on the clinical assessment and management of the disorder are presented. Propelled by the gradual availability of large scale TS cohorts, novel methodologies, and importantly, sheer enthusiasm by multiple researchers working together across different countries, the new era of the neurobiology of TS holds the promise to identify novel targets for improved therapies.

Impact of System Biology and Molecular Medicine on the Management of Complex Immune Mediated Respiratory Diseases Blanca Cárđaba, Girolamo Pelaia, 2021-10-21

Beta-Cell Fate: From Gene Circuits to Disease Mechanisms Simona Chera, Kenichiro Furuyama, Hanne Scholz, Shane T. Grey, Luiza Ghila, 2022-03-16

RNA Interference Ibrokhim Y. Abdurakhmonov, 2016-04-06 RNA interference (RNAi), a hallmark of all biological sciences of twenty-first century, is an evolutionarily conserved and double-stranded RNA-dependent eukaryotic cell defense process. Opportunity to utilize an organisms own gene and to systematically induce and trigger RNAi for any desired sequence made RNAi an efficient approach for functional genomics, providing a solution for conventional longstanding obstacles in life sciences. RNAi research and application have significantly advanced during past two decades. This book RNA interference provides an updated knowledge and progress on RNAi in various organisms, explaining basic principles, types, and property of inducers, structural modifications, delivery systems/methodologies, and various successful bench-to-field or clinic applications and disease therapies with some aspects of limitations, alternative tools, safety, and risk assessment.

The Characteristics of Pediatric Soft Tissue Sarcomas: Recent Advances in Management and Treatment Jilong Yang, Shunbin Xiong, 2023-08-31 Sarcomas are forms of cancer that comprise tumors of the connective and structural tissue of the body and mainly affect children and young adults. Pediatric soft tissue sarcomas (STS) are a heterogeneous group of malignant tumors constituting about 7% of all cancer cases. Rhabdomyosarcomas (RMS) constitute about half of all soft tissue sarcomas in children, the rest being constituted by non- rhabdomyosarcoma soft tissue sarcomas

(NRSTS). The estimated 5-year survival rate is around 65.6% in children and young adults, Rhabdomyosarcoma being as low as 59.2% in some cases.

Mitochondrial DNA Herve Seligmann,Ganesh Warthi,2018-10-31 The very short genomes of mitochondria summarize the complexity of molecular biology and its interactions with cellular and whole organism biology. Studies of mitogenomes contribute to the understanding of molecular biology and evolution, and to health management. Despite or even due to their small sizes, mitogenomes continue to surprise us. Studies of mitogenomes reveal the details of molecular organization and its evolution under constraints for miniaturization.

Promoting Sustainable Local and Community Economic Development Roland V. Anglin,2017-09-25 Growing local economies, empowering communities, revitalizing downtowns, developing entrepreneurship, building leadership, and enhancing nonprofits – you can achieve all these benefits and more with a comprehensive and strategic revitalization plan. Chronicling the struggle of local revitalization as organizers move from trial and error to effective revitalization strategies, *Promoting Sustainable Local and Community Economic Development* documents the current transformation in community revitalization from market-based incentives to mixed strategies of public sector learning, partnerships, and community capacity. Knowledge about the field and what works is growing, but not always publicized and readily accessible. This reference surveys the breadth of innovative place and people development practices, presenting lessons and examples at a general and textured level, putting information about innovative ways to change, influence, and improve the economic development process within easy reach. Roland Anglin brings his unique vantage point to the topic; his experience as a practitioner and applied academic allowed him to see how community economic development practices grow over time in size, scale, and impact. He highlights the difference between what is now termed community economic development (CED) and traditional local economic development practice, specifically the priority placed on community involvement in economic development partnerships between the private sector and government. The book includes case studies that demonstrate what has and has not worked in revitalization efforts, as well as how active public and private sector partnerships have been the most effective in revitalization efforts. A Resource Guide is included at the end of the book for readers who may want a more expansive understanding of community economic development.

DNA Methyltransferases – Role and Function Albert Jeltsch,Renata Z. Jurkowska,2016-11-08 DNA methyltransferases are important enzymes in a broad range of organisms. Dysfunction of DNA

methyltransferases in humans leads to many severe diseases, including cancer. This book focuses on the biochemical properties of these enzymes, describing their structures and mechanisms in bacteria, humans and other species, including plants, and also explains the biological processes of reading of DNA methylation and DNA demethylation. It covers many emerging aspects of the biological roles of DNA methylation functioning as an essential epigenetic mark and describes the role of DNA methylation in diseases. Moreover, the book explains modern technologies, like targeted rewriting of DNA methylation by designed DNA methyltransferases, as well as technological applications of DNA methyltransferases in DNA labelling. Finally, the book summarizes recent methods for the analysis of DNA methylation in human DNA. Overall, this book represents a comprehensive state-of-the-art- work and is a must-have for advanced researchers in the field of DNA methylation and epigenetics.

Immune system modeling and analysis Ramit Mehr, Miles Davenport, 2015-04-22 The rapid development of new methods for immunological data collection – from multicolor flow cytometry, through single-cell imaging, to deep sequencing – presents us now, for the first time, with the ability to analyze and compare large amounts of immunological data in health, aging and disease. The exponential growth of these datasets, however, challenges the theoretical immunology community to develop methods for data organization and analysis. Furthermore, the need to test hypotheses regarding immune function, and generate predictions regarding the outcomes of medical interventions, necessitates the development of mathematical and computational models covering processes on multiple scales, from the genetic and molecular to the cellular and system scales. The last few decades have seen the development of methods for presentation and analysis of clonal repertoires (those of T and B lymphocytes) and phenotypic (surface-marker based) repertoires of all lymphocyte types, and for modeling the intricate network of molecular and cellular interactions within the immune systems. This e-Book, which has first appeared as a ‘Frontiers in Immunology’ research topic, provides a comprehensive, online, open access snapshot of the current state of the art on immune system modeling and analysis.

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