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World Health Organization,World Health Organization. Global Influenza Programme

Wireless Algorithms, Systems, and Applications Kuai Xu, Haojin Zhu, 2015-07-31 This book constitutes the proceedings of the 10th International Conference on Wireless Algorithms, Systems, and Applications, WASA 2015, held in Qufu, Shandong, China, in August 2015. The 36 revised full papers presented together with 5 revised short papers and 42 invited papers were carefully reviewed and selected from 133 initial submissions. The papers present current trends, challenges, and state-of-the-art solutions related to various issues in wireless networks. Topics of interests include effective and efficient state-of-the-art algorithm design and analysis, reliable and secure system development and implementations, experimental study and testbed validation, and new application exploration in wireless networks. .

Oseltamivir John Oxford, Anna Palmer, Nigel Higson, 2004-11 Drugs in Context is an independent, peer reviewed and practice orientated review of all the significant data on a specific drug, placed in the context of the disease and clinical practice setting. Oseltamivir; Editorial: The coming years will see GPs prescribing more of the new anti-influenza drugs in primary care, both in treating the most vulnerable patients and in preventing virus spread in families and elderly residential homes. Disease Overview: The threat from influenza is greatly increased in those with other chronic underlying medical conditions, particularly cardiovascular disease, diabetes and chronic obstructive pulmonary disease. Drug Review: Reported benefits of oseltamivir treatment include faster resolution of symptoms such as cough, fever and sore throat, plus a speedier return to good health, normal sleeping patterns and a restoration of activity to pre-illness levels. Improving Practice: We can only hope that government policy may shift over the coming years to encourage the routine vaccination of more children and adults.

Pandemic Influenza Preparedness and Response World Health Organization, World Health Organization. Global Influenza Programme, 2009 This guidance is an update of WHO global influenza preparedness plan: the role of WHO and recommendations for national measures before and during pandemics, published March 2005 (WHO/CDS/CSR/GIP/2005.5).

Modeling, Design and Simulation of Systems Mohamed Sultan Mohamed Ali, Herman Wahid, Nurul Adilla Mohd Subha, Shafishuhaza Sahlan, Mohd Amri Md. Yunus, Ahmad Ridhwan Wahap, 2017-08-24 This two-volume set CCIS 751 and CCIS 752 constitutes the proceedings of the 17th Asia Simulation Conference, AsiaSim 2017, held in Malacca, Malaysia, in August/September 2017. The 124 revised full papers presented in this two-volume set were carefully reviewed and selected from 267 submissions. The papers contained in these proceedings address challenging issues in modeling and simulation in various fields such as embedded systems; symbiotic simulation; agent-based simulation; parallel and distributed simulation; high performance computing; biomedical engineering; big data; energy, society and economics; medical processes; simulation language and software; visualization; virtual reality; modeling and Simulation for IoT; machine learning; as well as the fundamentals and applications of computing.

Emerging Infectious Diseases , 2015

Case Studies in Bayesian Statistics Constantine Gatsonis, Robert E. Kass, Alicia Carriquiry, Andrew Gelman, David Higdon, Donna K. Pauler, Isabella Verdinelli, 2018-08-17 This volume contains invited case studies with the accompanying discussion as well as contributed papers selected by a refereeing process of 6th Workshop on Case Studies in Bayesian Statistics was held at the Carnegie Mellon University in October, 2001.

Drinking Water and Infectious Disease Paul Raymond Hunter, Michael Waite, Elettra Ronchi, 2002-07-30 In many countries, considerable uncertainty still exists about the contribution of drinking water to sporadic cases of disease. The Organization for Economic Cooperation and Development (OECD), in cooperation with the World Health Organization (WHO), led the Workshop on Molecular Technologies for Safe Drinking Water in 1998 to address the role of w

Algorithms in Bioinformatics Aaron Darling, Jens Stoye, 2013-08-16 This book constitutes the refereed proceedings of the 13th International

Workshop on Algorithms in Bioinformatics, WABI 2013, held in Sophia Antipolis, France, in September 2013. WABI 2013 is one of seven workshops which, along with the European Symposium on Algorithms (ESA), constitute the ALGO annual meeting and highlights research in algorithmic work for bioinformatics, computational biology and systems biology. The goal is to present recent research results, including significant work-in-progress, and to identify and explore directions of future research. The 27 full papers presented were carefully reviewed and selected from 61 submissions. The papers cover all aspects of algorithms in bioinformatics, computational biology and systems biology.

Topics in Applied Statistics Mingxiu Hu, Yi Liu, Jianchang Lin, 2013-09-14 This volume presents 27 selected papers in topics that range from statistical applications in business and finance to applications in clinical trials and biomarker analysis. All papers feature original, peer-reviewed content. The editors intentionally selected papers that cover many topics so that the volume will serve the whole statistical community and a variety of research interests. The papers represent select contributions to the 21st ICSA Applied Statistics Symposium. The International Chinese Statistical Association (ICSA) Symposium took place between the 23rd and 26th of June, 2012 in Boston, Massachusetts. It was co-sponsored by the International Society for Biopharmaceutical Statistics (ISBS) and American Statistical Association (ASA). This is the inaugural proceedings volume to share research from the ICSA Applied Statistics Symposium.

Spatial Audio Woon Seng Gan, Jung-Woo Choi, 2018-03-23 This book is a printed edition of the Special Issue Spatial Audio that was published in Applied Sciences

Recent Developments in Smart Healthcare Wenbing Zhao, Xiong Luo, Tie Qiu, 2018-03-23 This book is a printed edition of the Special Issue Smart Healthcare that was published in Applied Sciences

Theory and Applications of Time Series Analysis and Forecasting Olga Valenzuela, Fernando Rojas, Luis Javier Herrera, Héctor Pomares, Ignacio Rojas, 2023-04-04 This book presents a selection of peer-reviewed contributions on the latest developments in time series analysis and forecasting, presented at the 7th International Conference on Time Series and Forecasting, ITISE 2021, held in Gran Canaria, Spain, July 19-21, 2021. It is divided into four parts. The first part addresses general modern methods and theoretical aspects of time series analysis and forecasting, while the remaining three parts focus on forecasting methods in econometrics, time series forecasting and prediction, and numerous other real-world applications. Covering a broad range of topics, the book will give readers a modern perspective on the subject. The ITISE conference series provides a forum for scientists, engineers, educators and students to discuss the latest advances and implementations in the foundations, theory, models and applications of time series analysis and forecasting. It focuses on interdisciplinary research encompassing computer science, mathematics, statistics and econometrics.

Advances in Statistical Bioinformatics Kim-Anh Do, Steven Qin, Marina Vannucci, 2013-06-10 Chapter 1 An introduction to next-generation biological platforms Virginia Mohlere, Wenting Wang, and Ganiraju Manyam The University of Texas. MD Anderson Cancer Center 1.1 Introduction When Sanger and Coulson first described a reliable, efficient method for DNA sequencing in 1975 (Sanger and Coulson, 1975), they made possible the full sequencing of both genes and entire genomes. Although the method was resource-intensive, many institutions invested in the necessary equipment, and Sanger sequencing remained the standard for the next 30 years. Refinement of the process increased read lengths from around 25 to 2 Mohlere, Wang, and Manyam almost 750 base pairs (Schadt et al., 2010, fig. 1). While this greatly increased efficiency and reliability, the Sanger method still required not only large equipment but significant human investment, as the process requires the work of several people. This prompted researchers and companies such as Applied Biosystems to seek improved sequencing techniques and instruments. Starting in the late 2000s, new instruments came on the market that, although they actually decreased read length, lessened run time and could be operated more easily with fewer human

resources (Schadt et al., 2010). Despite discoveries that have illuminated new therapeutic targets, clarified the role of specific mutations in clinical response, and yielded new methods for diagnosis and predicting prognosis (Chin et al., 2011), the initial promise of genomic data has largely remained so far unfulfilled. The difficulties are numerous--

Statistical Methods in Spatial Epidemiology Andrew B. Lawson, 2013-07-08 Spatial epidemiology is the description and analysis of the geographical distribution of disease. It is more important now than ever, with modern threats such as bio-terrorism making such analysis even more complex. This second edition of Statistical Methods in Spatial Epidemiology is updated and expanded to offer a complete coverage of the analysis and application of spatial statistical methods. The book is divided into two main sections: Part I introduces basic definitions and terminology, along with map construction and some basic models. This is expanded upon in Part II by applying this knowledge to the fundamental problems within spatial epidemiology, such as disease mapping, ecological analysis, disease clustering, bio-terrorism, space-time analysis, surveillance and infectious disease modelling. Provides a comprehensive overview of the main statistical methods used in spatial epidemiology. Updated to include a new emphasis on bio-terrorism and disease surveillance. Emphasizes the importance of space-time modelling and outlines the practical application of the method. Discusses the wide range of software available for analyzing spatial data, including WinBUGS, SaTScan and R, and features an accompanying website hosting related software. Contains numerous data sets, each representing a different approach to the analysis, and provides an insight into various modelling techniques. This text is primarily aimed at medical statisticians, researchers and practitioners from public health and epidemiology. It is also suitable for postgraduate students of statistics and epidemiology, as well professionals working in government agencies.

Algebraic Statistics for Computational Biology L. Pachter, B. Sturmfels, 2005-08-22 This book, first published in 2005, offers an introduction to the application of algebraic statistics to computational biology.

Algebraic and Geometric Methods in Statistics Paolo Gibilisco, Eva Riccomagno, Maria Piera Rogantin, Henry P. Wynn, 2010 An up-to-date account of algebraic statistics and information geometry, which also explores the emerging connections between these two disciplines.

Manual for the Laboratory Diagnosis and Virological Surveillance of Influenza World Health Organization, WHO Global Influenza Surveillance Network, 2011 WHO has developed this manual in order to strengthen the laboratory diagnosis and virological surveillance of influenza infection by providing standard methods for the collection, detection, isolation and characterization of viruses.--Publisher's description.

Viruses and Man: A History of Interactions Milton W. Taylor, 2014-07-21 Milton Taylor, Indiana University, offers an easy-to-read and fascinating text describing the impact of viruses on human society. The book starts with an analysis of the profound effect that viral epidemics had on world history resulting in demographic upheavals by destroying total populations. It also provides a brief history of virology and immunology. Furthermore, the use of viruses for the treatment of cancer (viral oncolysis or virotherapy) and bacterial diseases (phage therapy) and as vectors in gene therapy is discussed in detail. Several chapters focus on viral diseases such as smallpox, influenza, polio, hepatitis and their control, as well as on HIV and AIDS and on some emerging viruses with an interesting story attached to their discovery or vaccine development. The book closes with a chapter on biological weapons. It will serve as an invaluable source of information for beginners in the field of virology as well as for experienced virologists, other academics, students, and readers without prior knowledge of virology or molecular biology.

Handbook of Approximate Bayesian Computation Scott A. Sisson, Yanan Fan, Mark Beaumont, 2018-09-03 As the world becomes increasingly complex, so do the statistical models required to analyse the challenging problems ahead. For the very first time in a single volume, the Handbook of Approximate Bayesian Computation (ABC) presents an extensive overview of the theory, practice and application of ABC methods. These simple, but powerful statistical techniques, take Bayesian statistics beyond the need to specify overly simplified models, to the setting where the model is defined

only as a process that generates data. This process can be arbitrarily complex, to the point where standard Bayesian techniques based on working with tractable likelihood functions would not be viable. ABC methods finesse the problem of model complexity within the Bayesian framework by exploiting modern computational power, thereby permitting approximate Bayesian analyses of models that would otherwise be impossible to implement. The Handbook of ABC provides illuminating insight into the world of Bayesian modelling for intractable models for both experts and newcomers alike. It is an essential reference book for anyone interested in learning about and implementing ABC techniques to analyse complex models in the modern world.

Infectious Disease Informatics Hsinchun Chen, Daniel Zeng, Ping Yan, 2010-03-12 Computer-based infectious disease surveillance systems are capable of real-time or near real-time detection of serious illnesses and potential bioterrorism agent exposures and represent a major step forward in disease surveillance. Infectious Disease Informatics: Syndromic Surveillance for Public Health and Bio-Defense is an in-depth monograph that analyzes and evaluates the outbreak modeling and detection capabilities of existing surveillance systems under a unified framework, and presents the first book-length coverage of the subject from an informatics-driven perspective. Individual chapters consider the state of the art, including the facilitation of data collection, sharing and transmission; a focus on various outbreak detection methods; data visualization and information dissemination issues; and system assessment and other policy issues. Eight chapters then report on several real-world case studies, summarizing and comparing eight syndromic surveillance systems, including those that have been adopted by many public health agencies (e.g., RODS and BioSense). The book concludes with a discussion of critical issues and challenges, with a look to future directions. This book is an excellent source of current information for researchers in public health and IT. Government public health officials and private-sector practitioners in both public health and IT will find the most up-to-date information available, and students from a variety of disciplines, including public health, biostatistics, information systems, computer science, and public administration and policy will get a comprehensive look at the concepts, techniques, and practices of syndromic surveillance.

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