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R E Dinnebier,S J L Billinge

<u>Introduction To Geographical Information Systems</u> Prithvish Nag And Smita Sengupta,Prithvish Nag,2008 In Indian context.

Geographical Information System Concepts And Business Opportunities Prithvish Nag And Smita Sengupta,2007 In Indian context.

The Wildlife Techniques Manual Nova J. Silvy,2012-03 A standard text in a variety of courses, the Techniques Manual, as it is commonly called, covers every aspect of modern wildlife management and provides practical information for applying the hundreds of methods described in its pages. To effectively incorporate the explosion of new information in the wildlife profession, this latest edition is logically organized into a two-volume set: Volume 1 is devoted to research techniques and Volume 2 focuses on management methodologies.

Mammalian Genomics Anatoly Ruvinsky,Jennifer A. Marshall Graves,2005 Organization of the Mammalian Genome; Linkage mapping ; Mapping genomes at the chromosome leveI ; Mapping genomes at the molecular level ; DNA sequence of the human and other mammalian genomes; Expression of the Mammalian Genomes ; The transcriptome ; The proteome ; The epigenome: epigenetic reguIation of gene expression in mammalian species ; Regulation of genome activity and genetic networks in mammals ; Inducing alterations in the mammalian genome for investigating the functions : of genes ; Evolution of the Mammalian Genome ; O A comparative analysis of mammalian genomics: prokaryote and eukaryote perspectives ; Elements and mechanisms of genome change ; DNA sequence evolution and phylogenetic footprinting ; Evolution of the mammalian karyotype ; Compara tive gene mapping, chromosome painting and the reconstruction of the ancestral mammalian karyotype ; Genome Analysis and Bioinformatics ; Bioinformatics: from computational

analysis through to integrated systems ; Genetic databases ; Gene predictions and annotations ; The Fruits of Mammalian Genomics ; Genomic research and progress in understanding inherited disorders in humans and other mammals ; Pharmacogenomics ; O Genome scanning for quantitative trait loci ; Mammalian popuIation genetics and genomics.

Models and Methods in Social Network Analysis Peter J. Carrington,John Scott,Stanley Wasserman,2005-02-07 Models and Methods in Social Network Analysis, first published in 2005, presents the most important developments in quantitative models and methods for analyzing social network data that have appeared during the 1990s. Intended as a complement to Wasserman and Faust's Social Network Analysis: Methods and Applications, it is a collection of articles by leading methodologists reviewing advances in their particular areas of network methods. Reviewed are advances in network measurement, network sampling, the analysis of centrality, positional analysis or blockmodelling, the analysis of diffusion through networks, the analysis of affiliation or 'two-mode' networks, the theory of random graphs, dependence graphs, exponential families of random graphs, the analysis of longitudinal network data, graphical techniques for exploring network data, and software for the analysis of social networks.

Information Technology Laboratory ,1998

Genetic Epidemiology Melissa A. Austin,Terri H. Beaty,W. David Dotson,2013 Genetic epidemiology plays a key role in discovering genetic factors influencing health and disease, and in understanding how genes and environmental risk factors interact. There is growing interest in this field within public health, with the goal of translating the results into promoting health and preventing disease in both families and populations. This textbook provides graduate students with a working knowledge of genetic epidemiology research methods. Following an overview of the field,

the book reviews key genetic concepts, provides an update on relevant genomic technology, including genome-wide chips and DNA sequencing, and describes methods for assessing the magnitude of genetic influences on diseases and risk factors. The book focuses on research study designs for discovering disease susceptibility genes, including family-based linkage analysis, candidate gene and genome-side association studies, assessing gene-environment interactions and epistasis, studies of Non-Mendelian inheritance, and statistical analyses of data from these studies. Specific applications of each research method are illustrated using a variety of diseases and risk factors relevant to public health, and useful web-based genetic analysis software, human reference panels, and repositories, that can greatly facilitate this work, are described.

  Food Engineering - Volume IV Gustavo V. Barbosa-Cánovas,2009-08-10 Food Engineering is a component of Encyclopedia of Food and Agricultural Sciences, Engineering and Technology Resources in the global Encyclopedia of Life Support Systems (EOLSS), which is an integrated compendium of twenty one Encyclopedias. Food Engineering became an academic discipline in the 1950s. Today it is a professional and scientific multidisciplinary field related to food manufacturing and the practical applications of food science. These volumes cover five main topics: Engineering Properties of Foods; Thermodynamics in Food Engineering; Food Rheology and Texture; Food Process Engineering; Food Plant Design, which are then expanded into multiple subtopics, each as a chapter. These four volumes are aimed at the following five major target audiences: University and College students Educators, Professional practitioners, Research personnel and Policy analysts, managers, and decision makers and NGOs

  Cyndi's List Cyndi Howells,2001 A two volume set which provides researchers with more than 70,000 links to every conceivable genealogical resource on the Internet.

Software Engineering for Modern Web Applications: Methodologies and Technologies Brandon, Daniel M.,2008-06-30 This book presents current, effective software engineering methods for the design and development of modern Web-based applications--Provided by publisher.

Supply Market Intelligence Robert Handfield,2006-01-13 Supply Market Intelligence: A Managerial Handbook for Building Sourcing Strategies charts the course to success for executives who are seeking to lead their organizations to supply-chain maturity. This volume is intended for supply-chain executives who struggle with the challenges of an uncertain supply market environment and whose supply chain function is constantly buffeted by unexpected changes in customer requirements and commodity prices. The author offers a more effective path, describing an integrated approach to supply chain management based upon solid market intelligence. This path is derived from the authors' multiple engagements with organizations such as Suncor Energy, Bank of America, Shell Oil, Honda of America, General Motors, Boston Scientific, Visteon, Federal Express, Sonoco, Duke Energy, Caterpillar, John Deere, GlaxoSmithKline, and many others. He emphasizes that the elements of a successful plan do not arise from a simple vision, instead they require much hard work and a focused approach backed by solid leadership. This book begins by defining supply market intelligence and discussing opportunities, the establishment of a project team, and conducting an internal business intelligence assessment. The book then examines the development of business and market intelligence, supplier evaluations, and sourcing strategies. It also explores how to execute a sourcing strategy, manage a strategic supplier relationship, and redesign an organization for effective supply-chain intelligence and strategic sourcing. This volume offers a benchmarking maturity model tool that covers all facets of end-to-end supply-chain management.

Powder Diffraction R E Dinnebier,S J L Billinge,2015-11-09 Powder diffraction is a widely used scientific technique in the characterization of materials with broad application in materials science, chemistry, physics, geology, pharmacology and archaeology. Powder Diffraction: Theory and Practice provides an advanced introductory text about modern methods and applications of powder diffraction in research and industry. The authors begin with a brief overview of the basic theory of diffraction from crystals and powders. Data collection strategies are described including x-ray, neutron and electron diffraction setups using modern day apparatus including synchrotron sources. Data corrections, essential for quantitative analysis are covered before the authors conclude with a discussion of the analysis methods themselves. The information is presented in a way that facilitates understanding the information content of the data, as well as best practices for collecting and analyzing data for quantitative analysis. This long awaited book condenses the knowledge of renowned experts in the field into a single, authoritative, overview of the application of powder diffraction in modern materials research. The book contains essential theory and introductory material for students and researchers wishing to learn how to apply the frontier methods of powder diffraction

Handbook of Neuroscience for the Behavioral Sciences, Volume 1 Gary G. Berntson,John T. Cacioppo,2009-10-12 As technology has made imaging of the brain noninvasive and inexpensive, nearly every psychologist in every subfield is using pictures of the brain to show biological connections to feelings and behavior. Handbook of Neuroscience for the Behavioral Sciences, Volume I provides psychologists and other behavioral scientists with a solid foundation in the increasingly critical field of neuroscience. Current and accessible, this volume provides the information they need to understand the new biological bases, research tools, and implications of

brain and gene research as it relates to psychology.

Admixture Dynamics, Natural Selection and Diseases in Admixed Populations Wenfei Jin,2015-10-20 In this thesis, Dr. Jin presents the distribution of ancestral chromosomal segments in the admixed genome, which could provide the information needed to explore population admixture dynamics. The author derives accurate population histories of African Americans and Mexicans using genome-wide single nucleotide polymorphisms (SNPs) data. Mapping the genetic background facilitates the study of natural selection in the admixed population, and the author identifies the signals of selection in African Americans since their African ancestors left for America. He further demonstrates that many of the selection signals were associated with African American-specific high-risk diseases such as prostate cancer and hypertension, suggesting an important role these disease-related genes might have played in adapting to their new environment. Lastly, the author reveals the complexity of natural selection in shaping human susceptibility to disease. The thesis significantly advances our understanding of the recent population admixture, adaptation to local environment and its health implications.

Versatile Video Coding Humberto Ochoa Dominguez,K.R. Rao,2022-09-01 Video is the main driver of bandwidth use, accounting for over 80 per cent of consumer Internet traffic. Video compression is a critical component of many of the available multimedia applications, it is necessary for storage or transmission of digital video over today's band-limited networks. The majority of this video is coded using international standards developed in collaboration with ITU-T Study Group and MPEG. The MPEG family of video coding standards begun on the early 1990s with MPEG-1, developed for video and audio storage on CD-ROMs, with support for progressive video. MPEG-2 was standardized in 1995 for applications of video on DVD, standard and high definition television,

with support for interlaced and progressive video. MPEG-4 part 2, also known as MPEG-2 video, was standardized in 1999 for applications of low- bit rate multimedia on mobile platforms and the Internet, with the support of object-based or content based coding by modeling the scene as background and foreground. Since MPEG-1, the main video coding standards were based on the so-called macroblocks. However, research groups continued the work beyond the traditional video coding architectures and found that macroblocks could limit the performance of the compression when using high-resolution video. Therefore, in 2013 the high efficiency video coding (HEVC) also known and H.265, was released, with a structure similar to H.264/AVC but using coding units with more flexible partitions than the traditional macroblocks. HEVC has greater flexibility in prediction modes and transform block sizes, also it has a more sophisticated interpolation and de blocking filters. In 2006 the VC-1 was released. VC-1 is a video codec implemented by Microsoft and the Microsoft Windows Media Video (VMW) 9 and standardized by the Society of Motion Picture and Television Engineers (SMPTE). In 2017 the Joint Video Experts Team (JVET) released a call for proposals for a new video coding standard initially called Beyond the HEVC, Future Video Coding (FVC) or known as Versatile Video Coding (VVC). VVC is being built on top of HEVC for application on Standard Dynamic Range (SDR), High Dynamic Range (HDR) and 360° Video. The VVC is planned to be finalized by 2020. This book presents the new VVC, and updates on the HEVC. The book discusses the advances in lossless coding and covers the topic of screen content coding. Technical topics discussed include: Beyond the High Efficiency Video CodingHigh Efficiency Video Coding encoderScreen contentLossless and visually lossless coding algorithmsFast coding algorithmsVisual quality assessmentOther screen content coding algorithmsOverview of JPEG Series

  DNA Fingerprinting in Plants Kurt Weising,Hilde Nybom,Markus Pfenninger,Kirsten

Wolff,Günter Kahl,2005-02-28 Given the explosive development of new molecular marker techniques over the last decade, newcomers and experts alike in the field of DNA fingerprinting will find an easy-to-follow guide to the multitude of techniques available in DNA Fingerprinting in Plants: Principles, Methods, and Applications, Second Edition. Along with step-by-step annotated p

Human Evolutionary Genetics, Second Edition Mark Jobling,Edward Hollox,Matthew Hurles,Toomas Kivisild,Chris Tyler-Smith,2013-06-23 Now in full-color, the Second Edition of Human Evolutionary Genetics has been completely revised to cover the rapid advances in the field since publication of the highly regarded First Edition. Written for upper-level undergraduate and graduate students, it is the only textbook to integrate genetic, archaeological, and linguistic perspectives on human evolution, and to offer a genomic perspective, reflecting the shift from studies of specific regions of the genome towards comprehensive genomewide analyses of human genetic diversity. Human Evolutionary Genetics is suitable for courses in Genetics, Evolution, and Anthropology. Those readers with a background in anthropology will find that the streamlined genetic analysis material contained in the Second Edition is more accessible. The new edition also integrates new technologies (including next-generation sequencing and genome-wide SNP typing) and new data analysis methods, including recent data on ancient genomes and their impact on our understanding of human evolution. The book also examines the subject of personal genomics and its implications.

Simulation in Textile Technology D Veit,2012-06-11 The use of mathematical modelling and computer simulation can vastly improve the quality, efficiency and economic success of textile technology. Simulation in textile technology provides a comprehensive review of the key principles, applications and benefits of modelling for textile production. After an introduction to modelling and simulation, Simulation in textile technology goes on to review the principles and applications of the

main types of model. The book first discusses neural networks and their applications before going on to explore evolutionary methods and fuzzy logic. It then considers computational fluid dynamics and finite element modelling. The modelling of fibrous structures and yarns are considered in the following chapters, along with wound packages, woven, braided and knitted structures. The book concludes by reviewing the simulation of textile processes and machinery. With its distinguished editor and team of expert contributors, Simulation in textile technology is a valuable reference tool for all those involved in both developing models of textile processes and those applying them to improve process efficiency and product quality. Provides a comprehensive review of the key principles, applications and benefits of modelling for textile production Discusses neural networks and their applications before going on to explore evolutionary methods and fuzzy logic Considers the modelling of fibrous structures and yarns, along with wound packages, woven, braided and knitted structures

   Dynamic Term Structure Modeling Sanjay K. Nawalkha,Gloria M. Soto,Natalia A. Beliaeva,2007-05-23 Praise for Dynamic Term Structure Modeling This book offers the most comprehensive coverage of term-structure models I have seen so far, encompassing equilibrium and no-arbitrage models in a new framework, along with the major solution techniques using trees, PDE methods, Fourier methods, and approximations. It is an essential reference for academics and practitioners alike. --Sanjiv Ranjan Das Professor of Finance, Santa Clara University, California, coeditor, Journal of Derivatives Bravo! This is an exhaustive analysis of the yield curve dynamics. It is clear, pedagogically impressive, well presented, and to the point. --Nassim Nicholas Taleb author, Dynamic Hedging and The Black Swan Nawalkha, Beliaeva, and Soto have put together a comprehensive, up-to-date textbook on modern dynamic term structure modeling. It is both

accessible and rigorous and should be of tremendous interest to anyone who wants to learn about state-of-the-art fixed income modeling. It provides many numerical examples that will be valuable to readers interested in the practical implementations of these models. --Pierre Collin-Dufresne Associate Professor of Finance, UC Berkeley The book provides a comprehensive description of the continuous time interest rate models. It serves an important part of the trilogy, useful for financial engineers to grasp the theoretical underpinnings and the practical implementation. --Thomas S. Y. Ho, PHD President, Thomas Ho Company, Ltd, coauthor, The Oxford Guide to Financial Modeling

Forest Conservation Genetics Andrew Young,David Boshier,Timothy Boyle,2000-07-24 Forest management must be sustainable not only in ecological, economic and social, but also genetic terms. Many forest managers are advocating and developing management strategies that give priority to conserving genetic diversity within production systems, or that recognise the importance of genetic considerations in achieving sustainable management. Forest Conservation Genetics draws together much previously uncollected information relevant to managing and conserving forests. The content emphasises the importance of conserving genetic diversity in achieving sustainable management. Each chapter is written by a leading expert and has been peer reviewed. Readers without a background in genetics will find the logical sequence of topics allows easy understanding of the principles involved and how those principles may impact on day-to-day forest planning and management decisions. The book is primarily aimed at undergraduate students of biology, ecology, forestry, and graduate students of forest genetics, resource management policy and/or conservation biology. It will prove useful for those teaching courses in these fields and as such help to increase the awareness of genetic factors in conservation and sustainable management, in both temperate and tropical regions.

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