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Xiaolu Zhang,Kim-Kwang Raymond Choo

Normal Mode Analysis Qiang Cui,Ivet Bahar,2005-12-12 Rapid developments in experimental techniques continue to push back the limits in the resolution, size, and complexity of the chemical and biological systems that can be investigated. This challenges the theoretical community to develop innovative methods for better interpreting experimental results. Normal Mode Analysis (NMA) is one such technique. Capable of providing unique insights into the structural and dynamical properties of complex systems, it is now finding a wide range of applications in chemical and biological problems. From the fundamental physical ideas to cutting-edge applications and beyond, this book presents a broad overview of normal mode analysis and its value in state-of-the-art research. The first section introduces NMA, examines NMA algorithm development at different resolutions, and explores the application of those techniques in the study of biological systems. Later chapters cover method developments based on or inspired by NMA but going beyond the harmonic approximation inherent in standard NMA techniques. Normal mode analysis complements traditional approaches with computational efficiency and applicability to large systems that are beyond the reach of older methods. This book offers a unique opportunity to learn from the experiences of an international, interdisciplinary panel of top researchers and explore the latest developments and applications of NMA to biophysical and chemical problems.

The Potato Genome Swarup Kumar Chakrabarti,Conghua Xie,Jagesh Kumar Tiwari,2017-12-26 This book describes the historical importance of potato (Solanum tuberosum L.),potato genetic resources and stocks (including S. tuberosum group Phureja

DM1-3 516 R44, a unique doubled monoploid homozygous line) used for potato genome sequencing. It also discusses strategies and tools for high-throughput sequencing, sequence assembly, annotation, analysis, repetitive sequences and genotyping-by-sequencing approaches. Potato (Solanum tuberosum L.; 2n = 4x = 48) is the fourth most important food crop of the world after rice, wheat and maize and holds great potential to ensure both food and nutritional security. It is an autotetraploid crop with complex genetics, acute inbreeding depression and a highly heterozygous nature. Further, the book examines the recent discovery of whole genome sequencing of a few wild potato species genomes, genomics in management and genetic enhancement of Solanum species, new strategies towards durable potato late blight resistance, structural analysis of resistance genes, genomics resources for abiotic stress management, as well as somatic cell genetics and modern approaches in true-potato-seed technology. The complete genome sequence provides a better understanding of potato biology, underpinning evolutionary process, genetics, breeding and molecular efforts to improve various important traits involved in potato growth and development.

The Art of Mac Malware Patrick Wardle,2022-07-12 A comprehensive guide to the threats facing Apple computers and the foundational knowledge needed to become a proficient Mac malware analyst. Defenders must fully understand how malicious software works if they hope to stay ahead of the increasingly sophisticated threats facing Apple products today. The Art of Mac Malware: The Guide to Analyzing Malicious Software is a comprehensive handbook to cracking open these malicious programs and seeing what’s inside. Discover the secrets of nation state backdoors, destructive ransomware, and

subversive cryptocurrency miners as you uncover their infection methods, persistence strategies, and insidious capabilities. Then work with and extend foundational reverse-engineering tools to extract and decrypt embedded strings, unpack protected Mach-O malware, and even reconstruct binary code. Next, using a debugger, you'll execute the malware, instruction by instruction, to discover exactly how it operates. In the book's final section, you'll put these lessons into practice by analyzing a complex Mac malware specimen on your own. You'll learn to: Recognize common infections vectors, persistence mechanisms, and payloads leveraged by Mac malware Triage unknown samples in order to quickly classify them as benign or malicious Work with static analysis tools, including disassemblers, in order to study malicious scripts and compiled binaries Leverage dynamical analysis tools, such as monitoring tools and debuggers, to gain further insight into sophisticated threats Quickly identify and bypass anti-analysis techniques aimed at thwarting your analysis attempts A former NSA hacker and current leader in the field of macOS threat analysis, Patrick Wardle uses real-world examples pulled from his original research. The Art of Mac Malware: The Guide to Analyzing Malicious Software is the definitive resource to battling these ever more prevalent and insidious Apple-focused threats.

RNA Biochemistry and Biotechnology Jan Barciszewski,Brian F.C. Clark,1999-08-31 RNA Biochemistry and Biotechnology describes various aspects of nucleic acid and protein structure, mainly RNA structure and proteins, interacting with specific RNA species. Papers deal with DNA protein interactions, telomerase, aminoacyl-tRNA synthetases, elongation factor Tu, DNA repair, RNA structure, NMR technology,

RNA aptamer interaction of biological macromolecules with metal ions. Two papers deal with theoretical aspects of RNA structure production and computer modelling. Many papers describe the possibility of commercial application of RNA biotechnology. One article discusses the impact of direct democracy on basic science supporting biotechnology. Readership: Advanced graduate students, Ph.D. students and young scientists as well as specialists in the field.

Human Monoclonal Antibodies Michael Steinitz,2013-09-14 The introduction of monoclonal antibodies revolutionized immunology. The development of human monoclonal antibodies was inspired primarily by the enormous clinical benefits promised by these reagents which can be used as anti-inflammatory reagents, anti-tumor reagents and reagents for passive immunization in a variety of pathologies. Human Monoclonal Antibodies: Methods and Protocols presents technical protocols of cellular and molecular methods for the production, purification and application of human monoclonal antibodies, as well as review articles on related topics of human monoclonal and polyclonal antibodies. Written in the successful Methods in Molecular Biology series format, chapters include introductions to their respective topics, lists of the necessary materials and reagents, step-by-step, readily reproducible protocols, and notes on troubleshooting and avoiding known pitfalls. Authoritative and easily accessible, Human Monoclonal Antibodies: Methods and Protocols seeks to serve both professionals and novices with its well-honed methodologies which will prove invaluable in a clinical setting.

Bioinformatics and Human Genomics Research Diego A. Forero,2021-12-23 Advances in high-throughput biological methods have led to the publication of a large

number of genome-wide studies in human and animal models. In this context, recent tools from bioinformatics and computational biology have been fundamental for the analysis of these genomic studies. The book Bioinformatics and Human Genomics Research provides updated and comprehensive information about multiple approaches of the application of bioinformatic tools to research in human genomics. It covers strategies analysis of genome-wide association studies, genome-wide expression studies and genome-wide DNA methylation, among other topics. It provides interesting strategies for data mining in human genomics, network analysis, prediction of binding sites for miRNAs and transcription factors, among other themes. Experts from all around the world in bioinformatics and human genomics have contributed chapters in this book. Readers will find this book as quite useful for their in silico explorations, which would contribute to a better and deeper understanding of multiple biological processes and of pathophysiology of many human diseases.

Wine Tasting Ronald S. Jackson,2016-12-22 From OIV-award-winning author, Ronald S. Jackson, Wine Tasting: A Professional Handbook, Third Edition, is an essential guide for any professional or serious connoisseur seeking to understand both the theory and practice of wine tasting. From techniques for assessing wine properties and quality, including physiological, psychological, and physicochemical sensory evaluation, to the latest information on the types of wine, the author guides the reader to a clear and applicable understanding of the wine tasting process. With its inclusion of illustrative data and testing technique descriptions, the book is ideal for both those who train tasters, those involved in designing wine tastings, and the connoisseur seeking to maximize their

perception and appreciation of wine. Contains revised and updated coverage, notably on the physiology and neurology of taste and odor perception Includes expanded coverage of the statistical aspect of wine tasting (specific examples to show the process), qualitative wine tasting, wine language, the origins of wine quality, and food and wine combination Provides a flow chart of wine tasting steps and production procedures Presents practical details on wine storage and the problems that can occur both during and following bottle opening

Digital Forensic Education Xiaolu Zhang,Kim-Kwang Raymond Choo,2019-07-24 In this book, the editors explain how students enrolled in two digital forensic courses at their institution are exposed to experiential learning opportunities, where the students acquire the knowledge and skills of the subject-matter while also learning how to adapt to the ever-changing digital forensic landscape. Their findings (e.g., forensic examination of different IoT devices) are also presented in the book. Digital forensics is a topic of increasing importance as our society becomes “smarter” with more of the “things” around us been internet- and inter-connected (e.g., Internet of Things (IoT) and smart home devices); thus, the increasing likelihood that we will need to acquire data from these things in a forensically sound manner. This book is of interest to both digital forensic educators and digital forensic practitioners, as well as students seeking to learn about digital forensics.

Marine Niche: Applications in Pharmaceutical Sciences Neelam M Nathani,Chandrashekar Mootapally,Indra R. Gadhvi,Bharat Maitreya,Chaitanya G. Joshi,2020-11-09 This book offers a comprehensive study of biological molecules acquired

from marine organisms, which have been exploited for drug discovery with the aim to treat human diseases. Biomolecules have potential impacts on a diverse range of fields, including medical and pharmaceutical science, industrial science, biotechnology, basic research, molecular science, environmental science and climate change, etc. To understand and effectively apply medicinally important biomolecules, multidisciplinary approaches are called for. The ocean remains a rich biological resource, and the vast untapped potential of novel molecules from marine bio-resources has caught the interest of more and more researchers. These novel biological compounds have never been found in terrestrial or other ecosystems, but only in this rich niche. Advances in sampling techniques and technologies, along with increased funding for research and nature conservation, have now encouraged scientists to look deeper in the waters. Aquaculture supports both tremendous seafood production and the bulk production of marine-derived drugs. Furthermore, molecular methods are now being extensively employed to explore the untapped marine microbial diversity. With the help of molecular and biotech tools, the ability of marine organisms to produce new biosynthetic drugs can be greatly enhanced. This book provides an extensive compilation of the latest information on marine resources and their undisputedly vital role in the treatment of diverse ailments.

Information, Communication and Computing Technology Saroj Kaushik,Daya Gupta,Latika Kharb,Deepak Chahal,2017-10-10 This book constitutes the refereed proceedings of the Second International Conference on Information, Communication and Computing Technology, ICICCT 2017, held in New Delhi, India, in May 2017. The 29 revised full papers and the 5 revised short papers presented in this volume were carefully

reviewed and selected from 219 submissions. The papers are organized in topical sections on network systems and communication security; software engineering; algorithm and high performance computing.

Apache Security Ivan Ristic,2005 The complete guide to securing your Apache web server--Cover.

Leptospira and Leptospirosis Ben Adler,2014-11-11 This volume covers all aspects of infection by pathogenic Leptospira species, the causative agents of the world's most widespread zoonosis. Topics include aspects of human and animal leptospirosis as well as detailed analyses of our current knowledge of leptospiral structure and physiology, epidemiology, pathogenesis, genomics, immunity and vaccines. Updates are presented on leptospiral systematics, identification and diagnostics, as well as practical information on culture of Leptospira. Contact information is also provided for Leptospira reference centers. All chapters were written by experts in the field, providing an invaluable reference source for scientists, veterinarians, clinicians and all others with an interest in leptospirosis.

Educational Technology United States. Congress. Senate. Committee on Appropriations. Subcommittee on Departments of Labor, Health and Human Services, Education, and Related Agencies,1996 This document presents witness testimony and supplemental materials from a Congressional hearing called to assess the effectiveness of federally funded educational technology programs, particularly Technology for Education, Star Schools, Ready to Learn Television, and Mathline. Other educational technologies were introduced at the hearing as well, so that all could be prioritized in light of new

presidential budget requests. Senators whose remarks or questions comprise part of the document include Thad Cochran, James Jeffords, Ernest Hollings, and Mark Hatfield. Other witnesses include: (1) Madeleine Kunin, Department of Education; (2) Tom Burnham, Mississippi superintendent of education; (3) Gary Vance, Satellite Educational Resources Consortium; (4) Walt Hindenlang, Hughes Electronics Galaxy Institute for Education; (5) Benito Casados, Hughes Electronics Galaxy Institute for Education; (6) Carolyn Reid-Wallace, Corporation for Public Broadcasting; (7) Joy Rouse, St. Louis County, Missouri Board of Education; (8) Beryl Jackson, PBS Mathline; (9) Joan Miller, West Sylvan Middle School, Portland, Oregon; (10) Margaret G. Kelly, International Society for Technology in Education; (11) Jeanne Hayes, Quality Education Data; (12) Anne Miller, Eastmen Kodak Co.; (13) Kathleen Fulton, Office of Technology Assessment; (14) Linda Morra, General Accounting Office; and (15) John Cradler. (Contains 30 tables.) (BEW)

Enzymes in Food and Beverage Processing Muthusamy Chandrasekaran,2015-10-23 Biotechnology, particularly eco-friendly enzyme technologies, has immense potential for the augmentation of diverse food products utilizing vast biodiversity, resolving environmental problems owing to waste disposal from food and beverage industries. In addition to introducing the basic concepts and fundamental principles of enzymes, Enzymes in Foo

Medicinal Plants and Fungi: Recent Advances in Research and Development Dinesh Chandra Agrawal,Hsin-Sheng Tsay,Lie-Fen Shyur,Yang-Chang Wu,Sheng-Yang Wang,2017-11-03 This book highlights the latest international research on different

aspects of medicinal plants and fungi. Studies over the last decade have demonstrated that bioactive compounds isolated from medicinal fungi have promising antitumor, cardiovascular, immunomodulatory, anti-allergic, anti-diabetic, and hepatoprotective properties. In the light of these studies, the book includes chapters (mostly review articles) by eminent researchers from twelve countries across the globe working in different disciplines of medicinal plants and fungi. It discusses topics such as the prevention of major neurodegenerative and neurotoxic mechanisms by Centella asiatica; the medicinal properties and therapeutic applications of several mushrooms species found in different parts of the world; and fungal endophytes as a source of bioactive metabolites including anticancer and cardioprotective agents. There are also chapters on strategies for identifying bioactive secondary metabolites of fungal origin; the use of genomic information to explore the biotechnological potential of medicinal mushrooms; and solid state fermentation of agro-industrial and forestry residues for the production of medicinal mushrooms. It is a valuable resource for the researchers, professionals and students working in the area of medicinal plants and fungi.

Mastering Docker - Third Edition Russ McKendrick,Scott Gallagher,2018-10-25 Master Docker and leverage its power in your day-to-day workflow Key Features Explore tools such as Docker Engine, Machine, Compose, and Swarm Discover how Docker can be integrated into your daily workflows Learn to leverage Docker Swarm and Kubernetes Book Description Docker has been a game-changer when it comes to how modern applications are deployed and created. It has now grown into a key driver of innovation beyond system administration, with an impact on the world of web development. But how

can you make sure you're keeping up with the innovations it's driving, or be sure you're using it to its full potential? Mastering Docker shows you how; this book not only demonstrates how to use Docker more effectively, but also helps you rethink and reimagine what's possible with it. You will cover concepts such as building, managing, and storing images, along with best practices to make you confident, before delving more into Docker security. You'll find everything related to extending and integrating Docker in new and innovative ways. Docker Compose, Docker Swarm, and Kubernetes will help you take control of your containers in an efficient manner. By the end of the book, you will have a broad, yet detailed, sense of what's possible with Docker, and how seamlessly it fits in with a range of other platforms and tools. What you will learn Become fluent with the basic components and concepts of Docker Learn the best ways to build, store, and distribute containers Understand how Docker can fit into your development workflow Secure your containers and files with Docker's security features Solve architectural problems using the first and third clustering tool Leverage Linux containers and create highly scalable applications Who this book is for If you are an I.T professional and recognize Docker's importance for innovation in everything from system administration to web development, but aren't sure how to use it to its full potential, Mastering Docker is for you.

RNA 3D Structure Analysis and Prediction Neocles Leontis,Eric Westhof,2012-06-05 With the dramatic increase in RNA 3D structure determination in recent years, we now know that RNA molecules are highly structured. Moreover, knowledge of RNA 3D structures has proven crucial for understanding in atomic detail

how they carry out their biological functions. Because of the huge number of potentially important RNA molecules in biology, many more than can be studied experimentally, we need theoretical approaches for predicting 3D structures on the basis of sequences alone. This volume provides a comprehensive overview of current progress in the field by leading practitioners employing a variety of methods to model RNA 3D structures by homology, by fragment assembly, and by de novo energy and knowledge-based approaches.

Multiple Sequence Alignment Methods David J Russell,2016-08-23 From basic performing of sequence alignment through a proficiency at understanding how most industry-standard alignment algorithms achieve their results, Multiple Sequence Alignment Methods describes numerous algorithms and their nuances in chapters written by the experts who developed these algorithms. The various multiple sequence alignment algorithms presented in this handbook give a flavor of the broad range of choices available for multiple sequence alignment generation, and their diversity is a clear reflection of the complexity of the multiple sequence alignment problem and the amount of information that can be obtained from multiple sequence alignments. Each of these chapters not only describes the algorithm it covers but also presents instructions and tips on using their implementation, as is fitting with its inclusion in the highly successful Methods in Molecular Biology series. Authoritative and practical, Multiple Sequence Alignment Methods provides a readily available resource which will allow practitioners to experiment with different algorithms and find the particular algorithm that is of most use in their application.

2012 4th International Conference on Cyber Conflict (CYCON 2012) ,2012 Legal

implications of territorial sovereignty in cyberspace / Wollff Heintschel von Heinegg -- When Not my problem isn't enough: political neutrality and national responsibility in cyber conflict / Jason Healey -- Neutrality in cyberspace / Wolff Heintschel von Heinegg -- Impact of cyberspace on human rights and democracy / Vittorio Fanchiotti, Jean Paul Pierini -- Russia's public stance on cyber/information warfare / Keir Giles -- French cyberdefense policy / Patrice Tromparent -- A treaty for governing cyber-weapons: potential benefits and practical limitations / Louise Arimatsu -- Internet as a critical infrastructure: a framework for the measurement of maturity and awareness in the cyber sphere / Assaf Y. Keren, Keren Elazari -- The significance of attribution to cyberspace coercion: a political perspective / Forrest Hare -- The militarisation of cyberspace: why less may be better / Myriam Dunn Cavelty -- Beyond domains, beyond commons: the context and theory of conflict in cyberspace / Jeffrey L. Caton -- Applying traditional military principles to cyber warfare / Samuel Liles, J. Eric Dietz, Marcus Rogers, Dean Larson -- The principle of maneuver in cyber operations / Scott D. Applegate -- Countering the offensive advantage in cyberspace: an integrated defensive strategy / David T. Fahrenkrug -- An analysis for a just cyber warfare / Mariarosaria Taddeo -- Socially engineered commoners as cyber warriors: Estonian future or present? / Birgy Lorenz, Kaido Kikkas -- The notion of combatancy in cyber warfare / Sean Watts -- Direct participation in cyber hostilities: terms of reference for like-minded states? / Jody Prescott -- Attack trends in present computer networks / Robert Koch, Björn Stelte, Mario Golling -- Attack as a term of art in international law: the cyber operations context / Michael N. Schmitt -- Ius ad bellum in cyberspace: some thoughts on the Schmitt-criteria for use of force / Katharina Ziolkowski -

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Deep Learning in Biology and Medicine Davide Bacciu,Paulo J. G. Lisboa,Alfredo Vellido,2021 Biology, medicine and biochemistry have become data-centric fields for which Deep Learning methods are delivering groundbreaking results. Addressing high impact challenges, Deep Learning in Biology and Medicine provides an accessible and organic collection of Deep Learning essays on bioinformatics and medicine. It caters for a wide readership, ranging from machine learning practitioners and data scientists seeking methodological knowledge to address biomedical applications, to life science specialists in search of a gentle reference for advanced data analytics.With contributions from internationally renowned experts, the book covers foundational methodologies in a wide spectrum of life sciences applications, including electronic health record processing, diagnostic imaging, text processing, as well as omics-data processing. This survey of

consolidated problems is complemented by a selection of advanced applications, including cheminformatics and biomedical interaction network analysis. A modern and mindful approach to the use of data-driven methodologies in the life sciences also requires careful consideration of the associated societal, ethical, legal and transparency challenges, which are covered in the concluding chapters of this book.

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