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Class-tested and coherent, this textbook teaches classical and web information retrieval, including web search and the related areas of text classification and text clustering from basic concepts. It gives an up-to-date treatment of all aspects of the design and implementation of systems for gathering, indexing, and searching documents; methods for evaluating systems; and an introduction to the use of machine learning methods on text collections. All the important ideas are explained using examples and figures, making it perfect for introductory courses in information retrieval for advanced undergraduates and graduate students in computer science. Based on feedback from extensive classroom experience, the book has been carefully structured in order to make teaching more natural and effective. Slides and additional exercises (with solutions for lecturers) are also available through the book's supporting website to help course instructors prepare their lectures.

Non-communicable diseases (NCDs) are the leading cause of death worldwide, contributing to over 73% of all deaths annually. Each day NCDs cause more than 100,000 deaths, 80% of which occur in low- and middle-income countries. NCDs, however, are largely preventable, and a great deal of technical knowledge exists about how to prevent and manage them. Why, then, have we, as a global community, not been more successful at reducing this NCD burden? Does a universal problem not have a universal solution? Created by an international consortium of experts, this informative and accessible book provides practical guidelines, key learning points, and dynamic, real-world case studies to aid NCD program managers, policy officers and decision-makers in low- and middle-income countries, so that they can assess interventions for the prevention and control of NCDs. The book was commissioned by the Prince Mahidol Award Conference (PMAC), an annual international conference centred on policy of global significance related to public health. Non-Communicable Disease Prevention: Best Buys, Wasted Buys and Contestable Buys emphasises the importance of context in NCD control and prevention, arguing that the success of an intervention lies in an ability to respond to local needs and environments. The book comprises ten chapters, which collectively explore the reasons behind, and strategies for, preventing and managing the NCD burden. It spans key themes such as political economy, the transferability of economic evidence, the role of cross-sectoral policies, the importance of deliberative processes, and health technology assessment. This book is written for the benefit of the global health community, and is primarily targeted at those individuals who are involved in NCD programs. This book will also be of interest to NCD champions, policy advocates, and educators spearheading the movement for increased visibility of NCDs.

Volume 5.

Since the first attempts at structure-based drug design about four decades ago, molecular modelling techniques for drug design have developed enormously, along with the increasing computational power and structural and biological information of active compounds and potential target molecules. Nowadays, molecular modeling can be considered to be an integral component of the modern drug discovery and development toolbox. Nevertheless, there are still many methodological challenges to be overcome in the application of molecular modeling approaches to drug discovery. The eight original research and five review articles collected in this book provide a snapshot of the state-of-the-art of molecular modeling in drug design, illustrating recent advances and critically discussing important challenges. The topics covered include virtual screening and pharmacophore modelling, chemoinformatic applications of artificial intelligence and machine learning, molecular dynamics simulation and enhanced sampling to investigate contributions of molecular flexibility to drug–receptor interactions, the modeling of drug–receptor solvation, hydrogen bonding and polarization, and drug design against protein–protein interfaces and membrane protein receptors.

Since the first edition of this invaluable book in 2012, third-party funding has become more mainstream in international arbitration practice. However, since even the existence of a third-party funding agreement in a dispute is often kept secret, it can be difficult to glean the specifics of successful funding agreements. This welcome book, now updated, expertly reveals the nuances of third-party funding in international arbitration, examines the phenomenon in key jurisdictions, and provides a reliable resource for users and potential users that may wish to tap into and make use of this distinctive funding tool. Focusing on Australia, the United Kingdom, the United States, Germany, the Netherlands, Canada, and South Africa, the authors analyze and assess the legal regime based upon legislation, judicial opinions, ethics opinions, and practitioner anecdotes describing the state of third-party funding in each jurisdiction. In addition to updating summaries of the law of the various jurisdictions, the second edition includes a new chapter addressing third-party funding in investor-state arbitration. Among the issues raised and examined are the following: · payment of adverse costs; · “Before-the-Event” (BTE) and “After-the-Event” (ATE) insurance; · attorney financing; pro bono representation, contingency representation, conditional fee arrangements; · loans; · ethical doctrines affecting the third-party funding industry; · possible future bundling, securitization, and trading of legal claims; · risk that the funder may put its own interests ahead of the client’s interests; and · whether the existence of a funding agreement must or should be disclosed to the decision maker. The second edition also includes discussion of recent institutional developments as they relate to third-party funding, including the work of the ICCA-Queen Mary Task Force on Third-Party Funding and how third-party funding is being incorporated into arbitral rules and investment treaties. Ably providing a thorough understanding of what third-party funding entails and what legal parameters exist, this book will be of compelling interest to parties aiming to take advantage of the high values, speed, reduced evidentiary costs, outcome predictability, industry expertise, and high award enforceability characteristic of the third-party funding arrangements available in international arbitration.

This book describes the current state of international grape genomics, with a focus on the latest findings, tools and strategies employed in genome sequencing and analysis, and genetic mapping of important agronomic traits. It also discusses how these are having a direct impact on outcomes for grape breeders and the international grape research community. While *V. vinifera* is a model species, it is not always appreciated that its cultivation usually requires the use of other *Vitis* species as rootstocks. The book discusses genetic diversity within the *Vitis* genus, the available genetic resources for breeding, and the available genomic resources for other *Vitis* species. Grapes (*Vitis vinifera* spp. *vinifera*) have been a source of food and wine since their domestication from their wild progenitor (*Vitis vinifera* ssp. *sylvestris*) around 8,000 years ago, and they are now the world’s most valuable horticultural crop. In addition to being economically important, *V. vinifera* is also a model organism for the study of perennial fruit crops for two reasons: Firstly, its ability to be transformed and micropropagated via somatic embryogenesis, and secondly its relatively small genome size of 500 Mb. The economic importance of grapes made *V. vinifera* an obvious early candidate for genomic sequencing, and accordingly, two draft genomes were reported in 2007. Remarkably, these were the first genomes of any fruiting crop to be sequenced and only the fourth for flowering plants. Although riddled with gaps and potentially omitting large regions of repetitive sequences, the two genomes have provided valuable insights into grape genomes. Cited in over 2,000 articles, the genome has served as a reference in more than 3,000 genome-wide transcriptional analyses. Further, recent advances in DNA sequencing and bioinformatics are enabling the assembly of reference-grade genome references for more grape genotypes revealing the exceptional extent of structural variation in the species.

Noninvasive mechanical ventilation is an effective technique for the management of patients with acute or chronic respiratory failure. This comprehensive and up-to-date book explores all aspects of the subject. The opening sections are devoted to theory and equipment, with detailed attention to the use of full-face masks or helmets, the range of available ventilators, and patient-ventilator interactions. Clinical applications are then considered in depth in a series of chapters that address the use of noninvasive mechanical ventilation in chronic settings and in critical care, both within and outside of intensive care units. Due attention is also paid to weaning from conventional mechanical ventilation, potential complications, intraoperative applications, and staff training. The closing chapters examine uses of noninvasive mechanical ventilation in neonatal and pediatric care. This book, written by internationally recognized experts, will be an invaluable guide for both clinicians and researchers.

This book addresses the diversity of tropical microorganisms and its applications in agriculture, renewable energy production and environmental protection. It covers several tropical habitats such as rain forests, mangroves, sea and river waters and describes how microorganisms isolated from these regions can be used to control insects and plant diseases, to improve sugar cane and biofuels production among other applications. The book also aims to bring researchers' attention to the potential of tropical microorganisms for biotechnological purposes, an area that is still far from being well explored.

This detailed volume guides readers through strategic planning and user-friendly guidelines in order to select the most suitable CRISPR-Cas system and target sites with high activity and specificity. Methods covering CRISPR gRNA design, CRISPR delivery, CRISPR activity quantification (indel quantification), and examples of applying CRISPR gene editing in human pluripotent stem cells, primary cells, gene therapy, and genetic screening are included. Written for the highly successful Methods in Molecular Biology series, chapters include introductions to their respective topics, lists of the necessary materials and reagents, step-by-step, readily reproducible laboratory protocols, and tips on troubleshooting and avoiding known pitfalls. Authoritative and invaluable, CRISPR Gene Editing: Methods and Protocols will assist undergraduates, graduates, and researchers with detailed guidelines and methods for the vitally important CRISPR gene editing field. Chapter 3 is available open access under a CC BY 4.0 license via link.springer.com.

Provides a professional-level reference to the Samsung ARTIK API, as well as to other aspects of interest to developers such as the file systems, the operating system internals, various available interfaces, input/output, and the hardware itself. This is the perfect book for experienced programmers and developers who want to jump in and work with Samsung's new ARTIK product line to create Internet of Things devices and applications. It is also a perfect follow-up resource for new-to-the-field developers who are just getting past the beginning stages of learning the ARTIK. Samsung ARTIK Reference begins with a concise overview of the hardware and the various developer reference boards that are available. Attention then shifts to operating system internals, modes such as sleep and startup, and the various file systems and their parameters that are available for developers to adjust. Also included is a reference of API calls, guidance on input and output, documentation of serial, audio, graphic, and other interfaces. There is extensive reference to online resources with annotation and commentary guiding the learning process in many directions for further study. What You Will Learn Install the ARTIK toolkit and prepare to develop Manipulate the inner workings of the ARTIK operating system Look up and refer to details of the ARTIK API specification Perform input and output over the peripheral interface buses Build embeddable applications in support of IoT devices Embed the ARTIK modules into your own hardware products Who This Book Is For Samsung ARTIK Reference is for experienced developers wanting to understand and begin working with ARTIK. The book is especially of interest to those wishing to interact with ARTIK modules from within their own applications and web services.

