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Andreas Holzinger, Martina Ziefle, Martin Hitz, Matjaz Debevc

Object-Based Image Analysis Thomas Blaschke, Stefan Lang, Geoffrey Hay, 2008-08-09 This book brings together a collection of invited interdisciplinary persp- tives on the recent topic of Object-based Image Analysis (OBIA). Its c- st tent is based on select papers from the 1 OBIA International Conference held in Salzburg in July 2006, and is enriched by several invited chapters. All submissions have passed through a blind peer-review process resulting in what we believe is a timely volume of the highest scientific, theoretical and technical standards. The concept of OBIA first gained widespread interest within the GIScience (Geographic Information Science) community circa 2000, with the advent of the first commercial software for what was then termed 'obje- oriented image analysis'. However, it is widely agreed that OBIA builds on older segmentation, edge-detection and classification concepts that have been used in remote sensing image analysis for several decades. Nevert-less, its emergence has provided a new critical bridge to spatial concepts applied in multiscale landscape analysis, Geographic Information Systems (GIS) and the synergy between image-objects and their radiometric char- teristics and analyses in Earth Observation data (EO).

Oxidative Damage to Nucleic Acids Mark D. Evans, Marcus S. Cooke, 2009-06-11 This book provides up-to-date coverage of selected topics in nucleic acid oxidation. The topics have been selected to cover everything from basic chemical mechanisms, repair of damage and the biological and pathological meaning of DNA oxidation. The chapters are authored by leading, research active, international experts in the respective topics.

Web Data Management Serge Abiteboul, Ioana Manolescu, Philippe Rigaux, Marie-Christine Rousset, Pierre Senellart, 2011-11-28 The Internet and World Wide Web have revolutionized access to information. Users now store information across multiple platforms from personal computers to smartphones and websites. As a consequence, data management concepts, methods and techniques are increasingly focused on distribution concerns. Now that information largely resides in the network, so do the tools that process this information. This book explains the foundations of XML with a focus on data distribution. It covers the many facets of distributed data management on the Web, such as description logics, that are already emerging in today's data integration applications and herald tomorrow's semantic Web. It also introduces the machinery used to manipulate the unprecedented amount of data collected on the Web. Several 'Putting into Practice' chapters describe detailed practical applications of the technologies and techniques. The book will serve as an introduction to the new, global, information systems for Web professionals and master's level courses.

Pacific Shipper, 1953

Mechanical Fatigue of Metals José A.F.O. Correia, Abílio M.P. De Jesus, António Augusto Fernandes, Rui Calçada, 2019-05-07 This volume contains the proceedings of the XIX International Colloquium on Mechanical Fatigue of Metals, held at the Faculty of Engineering of the University of Porto, Portugal, 5-7 September 2018. This International Colloquium facilitated and encouraged the exchange of knowledge and experiences among the different communities involved in both basic and applied research in the field of the fatigue of metals, looking at the problem of fatigue exploring analytical and numerical simulative approaches. Fatigue damage represents one of the most important types of damage to which structural materials are subjected in normal industrial services that can finally result in a sudden and unexpected abrupt fracture. Since metal alloys are still today the most used materials in designing the majority of components and structures able to carry the highest service loads, the study of the different aspects of metals fatigue attracts permanent attention of scientists, engineers and designers.

Cape Times Peninsula Directory,

Ontology Matching Jérôme Euzenat, Pavel Shvaiko, 2013-11-08 Ontologies tend to be found everywhere. They are viewed as the silver bullet for many applications, such as database integration, peer-to-peer systems, e-commerce, semantic web services, or social networks. However, in open or evolving systems, such as the semantic web, different parties would, in general, adopt different ontologies. Thus, merely using ontologies, like using XML, does not reduce heterogeneity: it just raises heterogeneity problems to a higher level. Euzenat and Shvaiko's book is devoted to ontology matching as a solution to the semantic heterogeneity problem faced by computer systems. Ontology matching aims at finding correspondences between semantically related entities of different ontologies. These correspondences may stand for equivalence as well as other relations, such as consequence, subsumption, or disjointness, between ontology entities. Many different matching solutions have been proposed so far from various viewpoints, e.g., databases, information systems, and artificial intelligence. The second edition of Ontology Matching has been thoroughly revised and updated to reflect the most recent advances in this quickly developing area, which resulted in more than 150 pages of new content. In particular, the book includes a new chapter dedicated to the methodology for performing ontology matching. It also covers emerging topics, such as data interlinking, ontology partitioning and pruning, context-based matching, matcher tuning, alignment debugging, and user involvement in matching, to mention a few. More than 100 state-of-the-art matching systems and frameworks were reviewed. With Ontology Matching, researchers and practitioners will find a reference book that presents currently available work in a uniform framework. In particular, the work and the techniques presented in this book can be equally applied to database schema matching, catalog integration, XML schema matching and other related problems. The objectives of the book include presenting (i) the state of the art and (ii) the latest research results in ontology matching by providing a systematic and detailed account of matching techniques and matching systems from theoretical, practical and application perspectives.

Smart Intelligent Computing and Applications Suresh Chandra Satapathy, Vikrant Bhateja, Swagatam Das, 2018-10-01 The proceedings covers advanced and multi-disciplinary research on design of smart computing and informatics. The theme of the book broadly focuses on various innovation paradigms in system knowledge, intelligence and sustainability that may be applied to provide realistic solution to varied problems in society, environment and industries. The volume publishes quality work pertaining to the scope of the conference which is extended towards deployment of emerging computational and knowledge transfer approaches, optimizing solutions in varied disciplines of science, technology and healthcare.

Human Factors in Computing and Informatics Andreas Holzinger, Martina Ziefle, Martin Hitz, Matjaz Debevc, 2013-06-26 This book constitutes the refereed proceedings of the First International Conference on Human Factors in Computing and Informatics, South CHI 2013, held in Maribor, Slovenia, in July 2013. South CHI is the successor of the USAB Conference series and promotes all aspects of human-computer interaction. The 38 revised full papers presented together with 12 short papers, 4 posters and 3 doctoral thesis papers were carefully reviewed and selected from 169 submissions. The papers are organized in the following topical sections: measurement and usability evaluation; usability evaluation - medical environments; accessibility methodologies; game-based methodologies; Web-based systems and attribution research; virtual environments; design culture for ageing well: designing for situated elderliness; input devices; adaptive systems and intelligent agents; and assessing the state of HCI research and practice in South-Eastern Europe.

 $Braby \hbox{\rm '}s\ Cape\ Province\ Directory\ ,} 1988$

Ricin Toxins Nilgun E. Tumer,2020-12-14 The plant toxin ricin is one of the most potent and lethal substances known. Due to its wide availability and the ease of production, ricin has been exploited as an agent of bioterrorism and biological warfare and is classified as a level B threat. Tons of ricin are produced annually worldwide as a by-product of castor oil, leading to exposure. Currently, there is no proven safe treatment for ricin intoxication. Ricin consists of an active A subunit (RTA) covalently linked to a cell binding B subunit (RTB). RTA depurinates a universally conserved adenine in the sarcin-ricin loop (SRL) of the large rRNA and inhibits protein synthesis. RTA-antibody complexes have been explored as immunotoxins against cancer cells. The articles published in this book address the history of ricin research, how ricin enters cells and traffics to the ribosome, how it inactivates ribosomes, how it induces inflammatory signaling pathways and programmed cell death, the design of therapies against ricin intoxication, and innovative assays for ricin detection.

Cape Town Visitors' Guide and Maps ,1976

Electronic Voting Robert Krimmer, Melanie Volkamer, Véronique Cortier, Rajeev Goré, Manik Hapsara, Uwe Serdült, David Duenas-Cid, 2018-09-06 This book constitutes the refereed proceedings of the Third International Joint Conference on Electronic Voting, E-Vote-ID 2018, held in Bregenz, Austria, in October 2018. The 13 full papers presented in this volume were carefully reviewed and selected from 45 submissions. The papers deal with topics connected with electronic voting including experiences and revisions of the real uses of E-voting systems and corresponding processes in

elections.

Unraveling DNA Maxim D. Frank-Kamenetskii,1993-07-19 ... brilliant ... Yet anybody can understand it: it reads like a detective story.' John Maddox, Editor of Nature ' ... he skillfully imbues us with his joy and fascination with the living world, and the role of DNA in it.' Bruce M. Alberts, President of the National Academy of Sciences Unraveling DNA provides both laymen and scientist readers with a concise highly readable understanding of the structure, properties, and functions of the DNA molecule. The reader will find answers to all major questions about the biological, biotechnological, medical, physical, chemical, and mathematical aspects of DNA. In addition, the book includes an historical retrospective of past DNA research and forecasts future trends in the field. Written by an internationally acclaimed professor of biophysics as well as one of the world's leading authorities in DNA research, Unraveling DNA is designed to help professionals not specializing in molecular biology to understand the recent advances in this rapidly expanding field. The book is also especially useful to advanced high school students, junior college students, and university students interested in modern biology, medicine, physics, chemistry, and mathematics.

<u>History of Computer Art</u> Thomas Dreher,2020 The development of the use of computers and software in art from the Fifties to the present is explained. As general aspects of the history of computer art an interface model and three dominant modes to use computational processes (generative, modular, hypertextual) are presented. The History of Computer Art features examples of early developments in media like cybernetic sculptures, computer graphics and animation (including music videos and demos), video and computer games, reactive installations, virtual reality, evolutionary art and net art. The functions of relevant art works are explained more detailed than usual in such histories.

Precision Medicine in Oncology Bulent Aydogan, James A. Radosevich, 2020-11-02 A FRESH EXAMINATION OF PRECISION MEDICINE'S INCREASINGLY PROMINENT ROLE IN THE FIELD OF ONCOLOGY Precision medicine takes into account each patient's specific characteristics and requirements to arrive at treatment plans that are optimized towards the best possible outcome. As the field of oncology continues to advance, this tailored approach is becoming more and more prevalent, channelling data on genomics, proteomics, metabolomics and other areas into new and innovative methods of practice. Precision Medicine in Oncology draws together the essential research driving the field forward, providing oncology clinicians and trainees alike with an illuminating overview of the technology and thinking behind the breakthroughs currently being made. Topics covered include: Biologically-guided radiation therapy Informatics for precision medicine Molecular imaging Biomarkers for treatment assessment Big data Nanoplatforms Casting a spotlight on this emerging knowledge base and its impact upon the management of tumors, Precision Medicine in Oncology opens up new possibilities and ways of working – not only for oncologists, but also for molecular biologists, radiologists, medical geneticists, and others.

Computational Pharmaceutics Defang Ouyang, Sean C. Smith, 2015-07-20 Molecular modeling techniques have been widely used in drug discovery fields for rational drug design and compound screening. Now these techniques are used to model or mimic the behavior of molecules, and help us study formulation at the molecular level. Computational pharmaceutics enables us to understand the mechanism of drug delivery, and to develop new drug delivery systems. The book discusses the modeling of different drug delivery systems, including cyclodextrins, solid dispersions, polymorphism prediction, dendrimer-based delivery systems, surfactant-based micelle, polymeric drug delivery systems, liposome, protein/peptide formulations, non-viral gene delivery systems, drug-protein binding, silica nanoparticles, carbon nanotube-based drug delivery systems, diamond nanoparticles and layered double hydroxides (LDHs) drug delivery systems. Although there are a number of existing books about rational drug design with molecular modeling techniques, these techniques still look mysterious and daunting for pharmaceutical scientists. This book fills the gap between pharmaceutics and molecular modeling, and presents a systematic and overall introduction to computational pharmaceutics. It covers all introductory, advanced and specialist levels. It provides a totally different perspective to pharmaceutical scientists, and will greatly facilitate the development of pharmaceutics. It also helps computational chemists to look for the important questions in the drug delivery field. This book is included in the Advances in Pharmaceutical Technology book series.

The Semantic Web: ESWC 2019 Satellite Events Pascal Hitzler, Sabrina Kirrane, Olaf Hartig, Victor de Boer, Maria-Esther Vidal, Maria Maleshkova, Stefan Schlobach, Karl Hammar, Nelia Lasierra, Steffen Stadtmüller, Katja Hose, Ruben Verborgh, 2019-10-11 This book constitutes the thoroughly refereed post-conference proceedings of the Satellite Events of the 16th Extended Semantic Web Conference, ESWC 2019, held in Portorož, Slovenia, in June 2019. The volume contains 38 poster and demonstration papers, 2 workshop papers, 5 PhD symposium papers, and 3 industry track papers, selected out of a total of 68 submissions. They deal with all areas of semantic web research, semantic technologies on the Web and Linked Data.

Single Stranded DNA Binding Proteins Marcos Túlio Oliveira, 2021 This volume provides a comprehensive set of protocols that can be used by any research lab to investigate diverse functional and structural properties of Single Stranded DNA Binding Proteins (SSBs) from eubacterial, archaeal, eukaryotic, mitochondrial and viral systems. Written in the highly 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 laboratory protocols, and tips on troubleshooting and avoiding known pitfalls. Authoritative and cutting-edge, Single Stranded DNA Binding Proteins aims to be a useful practical guide to researchers to help further their study in this field.

Targeted Therapies for Lung Cancer Ravi Salgia,2019-06-26 This book contextualizes translational research and provides an up to date progress report on therapies that are currently being targeted in lung cancer. It is now well established that there is tremendous heterogeneity among cancer cells both at the inter- and intra-tumoral level. Further, a growing body of work highlights the importance of targeted therapies and personalized medicine in treating cancer patients. In contrast to conventional therapies that are typically administered to the average patient regardless of the patient's genotype, targeted therapies are tailored to patients with specific traits. Nonetheless, such genetic changes can be disease-specific and/or target specific; thus, the book addresses these issues manifested in the somatically acquired genetic changes of the targeted gene. Each chapter is written by a leading medical oncologist who specializes in thoracic oncology and is devoted to a particular target in a specific indication. Contributors provide an in-depth review of the literature covering the mechanisms underlying signaling, potential cross talk between the target and downstream signaling, and potential emergence of drug resistance.

Kirsten&ajax&p2 Book Review: Unveiling the Power of Words

In some sort of driven by information and connectivity, the power of words has be evident than ever. They have the capacity to inspire, provoke, and ignite change. Such could be the essence of the book **Kirsten&ajax&p2**, a literary masterpiece that delves deep into the significance of words and their affect our lives. Compiled by a renowned author, this captivating work takes readers on a transformative journey, unraveling the secrets and potential behind every word. In this review, we will explore the book is key themes, examine its writing style, and analyze its overall effect on readers.

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