

Computational Biology

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Distributed and Sequential Algorithms for Bioinformatics



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**Jakub Kurzak,David A. Bader,Jack
Dongarra**

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Intelligent Systems Design and Applications Ajith Abraham,Aswani Kumar Cherukuri,Patricia Melin,Niketa Gandhi,2019-04-11 This book highlights recent research on Intelligent Systems and Nature Inspired Computing It presents 212 selected papers from the 18th International Conference on Intelligent Systems Design and Applications ISDA 2018 and the 10th World Congress on Nature and Biologically Inspired Computing NaBIC which was held at VIT University India ISDA NaBIC 2018 was a premier conference in the field of Computational Intelligence and brought together researchers engineers and practitioners whose work involved intelligent systems and their applications in industry and the real world Including contributions by authors from over 40 countries the book offers a valuable reference guide for all researchers students and practitioners in the fields of Computer Science and Engineering

Scientific Computing with Multicore and Accelerators

Jakub Kurzak,David A. Bader,Jack Dongarra,2010-12-07 The hybrid heterogeneous nature of future microprocessors and large high performance computing systems will result in a reliance on two major types of components multicore manycore central processing units and special purpose hardware massively parallel accelerators While these technologies have numerous benefits they also pose substantial perfo

Grid and Cooperative Computing ,2004

Operations Research in

Medicine-computing and Optimization in Medicine and Life Sciences Eva K. Lee,2006

Encyclopedia of Genetics,

Genomics, Proteomics and Bioinformatics, 8 Volume Set Michael J. Dunn,Lynn B. Jorde,Peter F. R. Little,Shankar Subramaniam,2005-11-11 Available in print and online this unique reference brings together all four fields of genetics genomics proteomics and bioinformatics to meet your dynamic research requirements It brings together the latest concepts in these vibrant areas and ensures a truly multidisciplinary approach Topics include genetic variation and evolution epigenetics the human genome expression profiling proteome families structural proteomics gene finding gene structure protein function and annotation and more The work incorporates a vast amount of topical information profiles cutting edge techniques and presents the very latest findings from an international team of over five hundred contributors With articles for both students and more experienced scientists this is a key reference source for everyone Contains more than 450 articles covering all aspects of genomics proteomics bioinformatics and related technologies Includes a glossary containing over 550 clear and concise definitions I am pleased to recommend it heartily as a essential reference tool should remain the definitive work for many years to come THE CHEMICAL EDUCATOR Jorde and co editors have done a remarkable job in coordinating this information distilling it into a package that is both easy to navigate and over flowing in discovery ELECTRIC REVIEW

The Good, The Bad and The Ugly: Multiple Roles of Bacteria in Human Life Tatiana Venkova,Chew Chieng Yeo,Manuel Espinosa,2018-10-17 Bacteria are among the earliest forms of life on Earth Notwithstanding their small size and primitive origin bacteria still have a tremendous impact on everyday human life Over the centuries research into bacteria have provided and enriched the fundamental biological knowledge due to their readily measured processes and effects on higher

organisms Although molecular genetics and microbiology were among the scientific fields that have mostly benefited from the discoveries made in bacteria our current state of knowledge has gone beyond what anyone could have ever imagined The present Research Topic aims to cover new and exciting broad aspects of the importance of bacteria to human life both positive and negative influences Regulation of bacterial gene expression replication and segregation control mechanisms cell to cell communication via quorum sensors and the relatively recent finding of bacterial immunity via CRISPR have led to the development of many and very important new tools in biotechnology and the emerging field of molecular medicine The battle against infectious diseases has also benefited from the genetic approaches that have been developed in the quest for finding new targets and novel drugs against pathogenic bacteria At the next level the human microbiome project has opened up new avenues in understanding the role of bacteria in human health and wellbeing Finally the relationship between bacterial infections and human cancers will also be covered a subject that is still under verification through rigorous experimental approaches Special emphasis will be given to the bacterial accessory genome i e the mobilome as the primary cause of health threatening antimicrobial resistance and the production of toxins and virulence factors Taking into account the evolutionary importance of horizontal gene transfer and the additional beneficial roles of certain bacterial mobile genetic elements they help project best the Good the Bad and the Ugly outline of this topic At the time this eBook is about to be published our Research Topic has registered nearly 55 000 views

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Sequential Algorithms for Bioinformatics Kayhan Erciyes,2015-10-31 This unique textbook reference presents unified coverage of bioinformatics topics relating to both biological sequences and biological networks providing an in depth analysis of cutting edge distributed algorithms as well as of relevant sequential algorithms In addition to introducing the latest algorithms in this area more than fifteen new distributed algorithms are also proposed Topics and features reviews a range of open challenges in biological sequences and networks describes in detail both sequential and parallel distributed algorithms for each problem suggests approaches for distributed algorithms as possible extensions to sequential algorithms when the distributed algorithms for the topic are scarce proposes a number of new distributed algorithms in each chapter to serve as potential starting points for further research concludes each chapter with self test exercises a summary of the key points a comparison of the algorithms described and a literature review **Computational Intelligence** Kurosh Madani, António Dourado, Agostinho Rosa, Joaquim Filipe, Janusz Kacprzyk, 2015-11-19 The present book includes a set of selected extended papers from the fifth International Joint Conference on Computational Intelligence IJCCI 2013 held in Vilamoura Algarve Portugal from 20 to 22 September 2013 The conference was composed by three co located conferences The International Conference on Evolutionary Computation Theory and Applications ECTA the International Conference on Fuzzy Computation Theory and Applications FCTA and the International Conference on Neural Computation Theory and Applications NCTA Recent progresses in scientific developments and applications in these three areas are reported in this book IJCCI received

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Proceedings of the ... Annual International Conference on Research in Computational Molecular Biology ,2004

Philosophical Transactions ,2008 Each issue of Transactions B is devoted to a specific area of the biological sciences including clinical science All papers are peer reviewed and edited to the highest standards Published on the 29th of each month Transactions B is essential reading for all biologists *Science* John Michels (Journalist),2005 A weekly record of scientific progress *ACM SIGPLAN Notices* ,2006 *Parallel Computing for Bioinformatics and Computational Biology* Albert Y. Zomaya,2006-05-24 Discover how to streamline complex bioinformatics applications with parallel computing This publication enables readers to handle more complex bioinformatics applications and larger and richer data sets As the editor clearly shows using powerful parallel computing tools can lead to significant breakthroughs in deciphering genomes understanding genetic disease designing customized drug therapies and understanding evolution A broad range of bioinformatics applications is covered with demonstrations on how each one can be parallelized to improve performance and gain faster rates of computation Current parallel computing techniques and technologies are examined including distributed computing and grid computing Readers are provided with a mixture of algorithms experiments and simulations that provide not only qualitative but also quantitative insights into the dynamic field of bioinformatics Parallel Computing for Bioinformatics and Computational Biology is a contributed work that serves as a repository of case studies collectively demonstrating how parallel computing streamlines difficult problems in bioinformatics and produces better results Each of the chapters is authored by an established expert in the field and carefully edited to ensure a consistent approach and high standard throughout the publication The work is organized into five parts Algorithms and models Sequence analysis and microarrays Phylogenetics Protein folding Platforms and enabling technologies Researchers educators and students in the field of bioinformatics will discover how high performance computing can enable them to handle more complex data sets gain deeper insights and make new discoveries

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