

Computational Biology

K. Erciyes

Distributed and Sequential Algorithms for Bioinformatics



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Distributed and 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 **Parallel**

Computing for Bioinformatics and Computational Biology Albert Y. Zomaya, 2006-04-21 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

Computational Science - ICCS 2003. Part 3. Peter Sloot, 2003-05-22 The four volume set LNCS 2657 LNCS 2658 LNCS 2659 and LNCS 2660 constitutes the refereed proceedings of the Third International Conference on Computational Science ICCS 2003 held concurrently in Melbourne Australia and in St Petersburg Russia in June 2003 The four volumes present more than 460 reviewed contributed and invited papers and span the whole range of computational science from foundational issues in computer science and algorithmic mathematics to advanced applications in virtually all application

fields making use of computational techniques These proceedings give a unique account of recent results in the field

Deep Learning and Parallel Computing Environment for Bioengineering Systems Arun Kumar

Sangaiah,2019-07-26 Deep Learning and Parallel Computing Environment for Bioengineering Systems delivers a significant forum for the technical advancement of deep learning in parallel computing environment across bio engineering diversified domains and its applications Pursuing an interdisciplinary approach it focuses on methods used to identify and acquire valid potentially useful knowledge sources Managing the gathered knowledge and applying it to multiple domains including health care social networks mining recommendation systems image processing pattern recognition and predictions using deep learning paradigms is the major strength of this book This book integrates the core ideas of deep learning and its applications in bio engineering application domains to be accessible to all scholars and academicians The proposed techniques and concepts in this book can be extended in future to accommodate changing business organizations needs as well as practitioners innovative ideas Presents novel in depth research contributions from a methodological application perspective in understanding the fusion of deep machine learning paradigms and their capabilities in solving a diverse range of problems Illustrates the state of the art and recent developments in the new theories and applications of deep learning approaches applied to parallel computing environment in bioengineering systems Provides concepts and technologies that are successfully used in the implementation of today s intelligent data centric critical systems and multi media Cloud Big data 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

Encyclopedia of Parallel Computing David Padua,2011-09-08 Containing over 300 entries in an A Z format the Encyclopedia of Parallel Computing provides easy intuitive access to relevant information for professionals and researchers seeking access to any aspect within the broad field of parallel computing Topics for this comprehensive reference were selected written and peer reviewed by an international pool of distinguished researchers in the field The Encyclopedia is broad in scope covering machine organization programming languages algorithms and applications Within each area concepts designs and specific implementations are presented The highly structured essays in this work comprise synonyms a definition and discussion of the topic bibliographies and links to related literature Extensive cross references to other entries within the Encyclopedia support efficient user friendly searchers for immediate access to useful information Key concepts presented in the Encyclopedia of Parallel Computing include laws and metrics specific numerical and non numerical algorithms asynchronous algorithms libraries of subroutines benchmark suites applications sequential consistency and cache coherency machine classes such as clusters shared memory multiprocessors special purpose machines and dataflow

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Distributed Sequential Algorithms Bioinformatics Computational Book Review: Unveiling the Magic of Language

In an electronic era where connections and knowledge reign supreme, the enchanting power of language has are more apparent than ever. Its power to stir emotions, provoke thought, and instigate transformation is actually remarkable. This extraordinary book, aptly titled "**Distributed Sequential Algorithms Bioinformatics Computational**," compiled by a highly acclaimed author, immerses readers in a captivating exploration of the significance of language and its profound effect on our existence. Throughout this critique, we will delve to the book is central themes, evaluate its unique writing style, and assess its overall influence on its readership.

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