

Computational Biology

K. Erciyes

# Distributed and Sequential Algorithms for Bioinformatics



 Springer

# Distributed Sequential Algorithms Bioinformatics Computational

**Hongru Du**



## **Distributed Sequential Algorithms Bioinformatics Computational:**

**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

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**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

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**Machine Learning Used in Biomedical Computing and Intelligence Healthcare, Volume I** Honghao Gao,Ying Li,Zijian Zhang,Wenbing Zhao,2021-06-17 *A Bioinformatics Discovery-oriented Computing Framework* Jake Yue Chen,2001

**Bioinformatics** Bertil Schmidt,2011 A compilation of recent approaches from prominent researchers this referencediscusses how to take advantage of bioinformatics applications and algorithmson a variety of modern parallel architectures

**Grid and Cooperative Computing** ,2004

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**String Processing and Information Retrieval Symposium and International Workshop on Groupware** ,1999 This text contains proceedings from the 1999 String Processing and Information Retrieval Symposium SPIRE 99 and the 1999 International Workshop on Groupware CRIWG 99 It is aimed at professors researchers practitioners students and other computing professionals

**Proceedings of the ... Annual International Conference on Research in Computational Molecular Biology** ,2004 *Proceedings of the Eighth Annual International Conference on Research in Computational Molecular Biology* Philip E Bourne,RECOMB04: The Eighth Annual International Conference on Research in Computational Molecular Biology 2004,2004

**International Conference on Mathematical Biology 2007** Kamel A.M. Atan,I.S. Krishnarajah,R. Isamidin,2008-02-05 Mathematical biology is an interdisciplinary area that focuses on the application of mathematics to biology systems Mathematical biology spans all levels of biological organization and biological function from the configuration of biological macromolecules to the entire ecosphere over the course of evolutionary time The International Conference on Mathematical Biology 2007 provides the opportunity to bring together the people projects and issues from all over the world to share experiences and examine the challenge of applying mathematics to biological problems

**BIBE 2004** IEEE Computer Society,IEEE International Symposium on Bioinformatics and Bioengineering,IEEE Neural Networks Society,2004

High Performance Computing ,2004

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

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