

Gary Benson
Roderic Page (Eds.)

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Algorithms in Bioinformatics

Third International Workshop, WABI 2003
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**Teresa M. Przytycka, Marie-France
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Algorithms in Bioinformatics Gary Benson, Roderic D. M. Page, 2003 Algorithms in Bioinformatics Gary Benson, Roderic Page, 2003-09-09 This book constitutes the refereed proceedings of the Third International Workshop on Algorithms in Bioinformatics WABI 2003 held in Budapest Hungary in September 2003. The 36 revised full papers presented were carefully reviewed and selected from 78 submissions. The papers are organized in topical sections on comparative genomics, database searching, gene finding and expression, genome mapping, pattern and motif discovery, phylogenetic analysis, polymorphism, protein structure, sequence alignment and string algorithms. **Algorithms in Bioinformatics** Roderic Guigo, Dan Gusfield, 2002-09-02 We are pleased to present the proceedings of the Second Workshop on Algorithms in Bioinformatics WABI 2002 which took place on September 17-21 2002 in Rome Italy. The WABI workshop was part of a three conference meeting which in addition to WABI included the ESA and APPROX 2002. The three conferences are jointly called ALGO 2002 and were hosted by the Faculty of Engineering University of Rome La Sapienza. See <http://www.dis.uniroma1.it/algo02> for more details. The Workshop on Algorithms in Bioinformatics covers research in all areas of algorithmic work in bioinformatics and computational biology. The emphasis is on discrete algorithms that address important problems in molecular biology, genomics and genetics that are founded on sound models that are computationally efficient and that have been implemented and tested in simulations and on real datasets. The goal is to present recent research results including significant work in progress and to identify and explore directions of future research. Original research papers including significant work in progress or state of the art surveys were solicited on all aspects of algorithms in bioinformatics including but not limited to exact and approximate algorithms for genomics, genetics, sequence analysis, gene and signal recognition, alignment, molecular evolution, phylogenetics, structure determination or prediction, gene expression and gene networks, proteomics, functional genomics and drug design. **Algorithms in Bioinformatics** Steven L. Salzberg, Tandy Warnow, 2009-08-28 These proceedings contain papers from the 2009 Workshop on Algorithms in Bioinformatics WABI held at the University of Pennsylvania in Philadelphia Pennsylvania during September 12-13 2009. WABI 2009 was the ninth annual conference in this series which focuses on novel algorithms that address important problems in genomics, molecular biology and evolution. The conference emphasizes research that describes computationally efficient algorithms and data structures that have been implemented and tested in simulations and on real data. WABI is sponsored by the European Association for Theoretical Computer Science EATCS and the International Society for Computational Biology ISCB. WABI 2009 was supported by the Penn Genome Frontiers Institute and the Penn Center for Bioinformatics at the University of Pennsylvania. For the 2009 conference 90 full papers were submitted for review by the Program Committee and from this strong field of submissions 34 papers were chosen for presentation at the conference and publication in the proceedings. The final program covered a wide range of topics.

including gene interaction networks molecular phylogeny RNA and protein structure and genome evolution Algorithms in Bioinformatics Mihai Pop,Hélène Touzet,2015-08-27 This book constitutes the refereed proceedings of the 15th International Workshop on Algorithms in Bioinformatics WABI 2015 held in Atlanta GA USA in September 2015 The 23 full papers presented were carefully reviewed and selected from 56 submissions The selected papers cover a wide range of topics from networks to phylogenetic studies sequence and genome analysis comparative genomics and RNA structure *Algorithms in Bioinformatics* Philipp Bücher,Bernard M.E. Moret,2006-08-30 Here are the refereed proceedings of the 6th International Workshop on Algorithms in Bioinformatics WABI 2006 held in the course of the ALGO 2006 conference meetings The book presents 36 revised full papers addressing all current issues of algorithms in bioinformatics from mathematical tools to experimental studies of approximation algorithms and reports on significant computational analyses For the first time coverage extends to machine learning approaches along with combinatorial optimization **Algorithms in Bioinformatics** Teresa M Przytycka,Marie-France Sagot,2011-08-22 **Algorithms in Bioinformatics** Teresa M. Przytycka,Marie-France Sagot,2011-08-31 This book constitutes the refereed proceedings of the 11th International Workshop on Algorithms in Bioinformatics WABI 2011 held in Saarbrücken Germany in September 2011 The 30 papers presented were carefully reviewed and selected from 77 submissions They cover aspects of algorithms in bioinformatics computational biology and systems biology *Algorithms in Bioinformatics* Aaron Darling,Jens Stoye,2013-08-16 This book constitutes the refereed proceedings of the 13th International Workshop on Algorithms in Bioinformatics WABI 2013 held in Sophia Antipolis France in September 2013 WABI 2013 is one of seven workshops which along with the European Symposium on Algorithms ESA constitute the ALGO annual meeting and highlights research in algorithmic work for bioinformatics computational biology and systems biology The goal is to present recent research results including significant work in progress and to identify and explore directions of future research The 27 full papers presented were carefully reviewed and selected from 61 submissions The papers cover all aspects of algorithms in bioinformatics computational biology and systems biology Algorithms in Bioinformatics Ben Raphael,Jijun Tang,2012-08-29 This book constitutes the refereed proceedings of the 12th International Workshop on Algorithms in Bioinformatics WABI 2012 held in Ljubljana Slovenia in September 2012 WABI 2012 is one of six workshops which along with the European Symposium on Algorithms ESA constitute the ALGO annual meeting and focuses on algorithmic advances in bioinformatics computational biology and systems biology with a particular emphasis on discrete algorithms and machine learning methods that address important problems in molecular biology The 35 full papers presented were carefully reviewed and selected from 92 submissions The papers include algorithms for a variety of biological problems including phylogeny DNA and RNA sequencing and analysis protein structure and others

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