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Algorithms in Bioinformatics

Second International Workshop, WASE 2002 Rome, Italy, September, 2002 Proceedings



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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** Vincent Moulton, Mona Singh, 2010 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 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 Algorithms in Bioinformatics Steven L. Salzberg, Tandy Warnow, 2009-09-19 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 imp tantproblemsingenomics molecular biology and evolution The conference phasizes research that describes computationally e cient 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 C puter Science EATCS and the International Society for Computational Bi ogy 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 eld of submissions 34 papers were chosen for presentation at the conference and publication in the proceedings The nal programcovered a wide range of topics including gene interaction n works molecular phylogeny RNA and protein structure and genome evolution Algorithms in Bioinformatics Dan Brown, Burkhard Morgenstern, 2014-08-15 This book constitutes the refereed proceedings of the 14th International Workshop on Algorithms in Bioinformatics WABI 2014 held in Wroclaw Poland in September 2014 WABI 2014 was one of seven conferences that were organized as part of ALGO 2014 WABI is an annual conference series on all aspects of algorithms and data structure in molecular biology genomics and phylogeny data analysis The 26 full papers presented

together with a short abstract were carefully reviewed and selected from 61 submissions. The selected papers cover a wide range of topics from sequence and genome analysis through phylogeny reconstruction and networks to mass spectrometry data analysis. Algorithms in Bioinformatics Gary Benson, Roderic D. M. Page, 2003. Algorithms in Bioinformatics. Vincent Moulton, Mona Singh, 2010-08-25. This book constitutes the refereed proceedings of the 10th International Workshop on Algorithms in Bioinformatics WABI 2010 held in Liverpool UK in September 2010. The 30 revised full papers presented were carefully reviewed and selected from 83 submissions. The papers are organized in topical sections on biomolecular structure RNA protein and molecular comparison comparative genomics haplotype and genotype analysis high throughput data analysis next generation sequencing and flow cytometry networks phylogenetics and sequences strings and motifs

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Hannenhalli, 2007-08-22 The refereed proceedings from the 7th International Workshop on Algorithms in Bioinformatics are provided in this volume Papers address current issues in algorithms in bioinformatics ranging from mathematical tools to experimental studies of approximation algorithms to significant computational analyses Biological problems examined include genetic mapping sequence alignment and analysis phylogeny comparative genomics and protein structure

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