

1. Record Nr.	UNISA996465914403316
Titolo	Algorithms in Bioinformatics [[electronic resource]] : First International Workshop, WABI 2001, Aarhus, Denmark, August 28-31, 2001, Proceedings / / edited by Olivier Gascuel, Bernard M.E. Moret
Pubbl/distr/stampa	Berlin, Heidelberg : , : Springer Berlin Heidelberg : , : Imprint : Springer, , 2001
ISBN	3-540-44696-6
Edizione	[1st ed. 2001.]
Descrizione fisica	1 online resource (X, 314 p.)
Collana	Lecture Notes in Computer Science, , 0302-9743 ; ; 2149
Disciplina	570/.285
Soggetti	Computer programming Life sciences Data structures (Computer science) Algorithms Computers Programming Techniques Life Sciences, general Data Structures and Information Theory Algorithm Analysis and Problem Complexity Data Structures Computation by Abstract Devices
Lingua di pubblicazione	Inglese
Formato	Materiale a stampa
Livello bibliografico	Monografia
Note generali	Bibliographic Level Mode of Issuance: Monograph
Nota di bibliografia	Includes bibliographical references and index.
Nota di contenuto	An Improved Model for Statistical Alignment -- Improving Profile-Profile Alignments via Log Average Scoring -- False Positives in Genomic Map Assembly and Sequence Validation -- Boosting EM for Radiation Hybrid and Genetic Mapping -- Placing Probes along the Genome Using Pairwise Distance Data -- Comparing a Hidden Markov Model and a Stochastic Context-Free Grammar -- Assessing the Statistical Significance of Overrepresented Oligonucleotides -- Pattern Matching and Pattern Discovery Algorithms for Protein Topologies -- Computing Linking Numbers of a Filtration -- Side Chain-Positioning as an Integer Programming Problem -- A Chemical-Distance-Based Test for Positive Darwinian Selection -- Finding a Maximum Compatible

Tree for a Bounded Number of Trees with Bounded Degree Is Solvable
in Polynomial Time -- Experiments in Computing Sequences of
Reversals -- Exact-IEBP: A New Technique for Estimating Evolutionary
Distances between Whole Genomes -- Finding an Optimal Inversion
Median: Experimental Results -- Analytic Solutions for Three-Taxon
MLMC Trees with Variable Rates Across Sites -- The Performance of
Phylogenetic Methods on Trees of Bounded Diameter -- (1+?)-
Approximation of Sorting by Reversals and Transpositions -- On the
Practical Solution of the Reversal Median Problem -- Algorithms for
Finding Gene Clusters -- Determination of Binding Amino Acids Based
on Random Peptide Array Screening Data -- A Simple Hyper-Geometric
Approach for Discovering Putative Transcription Factor Binding Sites --
Comparing Assemblies Using Fragments and Mate-Pairs.

2. Record Nr.	UNISOBSOBE00033661
Autore	Baudelaire, Charles
Titolo	2 / Baudelaire
Pubbl/distr/stampa	Paris, : Gallimard, 1976
ISBN	2070108538
Descrizione fisica	1691 p. ; 18 cm
Collana	Bibliothèque de la Pléiade ; 7

Lingua di pubblicazione	Francese
Formato	Materiale a stampa
Livello bibliografico	Monografia