

1. Record Nr.	UNISA996465894603316
Titolo	Combinatorial Pattern Matching [[electronic resource]] : 18th Annual Symposium, CPM 2007, London, Canada, July 9-11, 2007, Proceedings / / edited by Bin Ma, Kaizhong Zhang
Pubbl/distr/stampa	Berlin, Heidelberg : , : Springer Berlin Heidelberg : , : Imprint : Springer, , 2007
ISBN	3-540-73437-6
Edizione	[1st ed. 2007.]
Descrizione fisica	1 online resource (XII, 368 p.)
Collana	Theoretical Computer Science and General Issues, , 2512-2029 ; ; 4580
Disciplina	005.1
Soggetti	Pattern recognition systems Algorithms Natural language processing (Computer science) Data mining Bioinformatics Artificial intelligence—Data processing Automated Pattern Recognition Natural Language Processing (NLP) Data Mining and Knowledge Discovery Computational and Systems Biology Data Science
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	Invited Talks (Abstracts) -- A Combinatorial Approach to Genome-Wide Ortholog Assignment: Beyond Sequence Similarity Search -- Stringology: Some Classic and Some Modern Problems -- Algorithmic Problems in Scheduling Jobs on Variable-Speed Processors -- Session 1: Alogirthmic Techniques I -- Speeding Up HMM Decoding and Training by Exploiting Sequence Repetitions -- On Demand String Sorting over Unbounded Alphabets -- Session 2: Approximate Pattern Matching -- Finding Witnesses by Peeling -- Cache-Oblivious Index for Approximate String Matching -- Improved Approximate String Matching and Regular Expression Matching on Ziv-Lempel Compressed Texts -- Self-normalised Distance with Don't Cares -- Session 3: Data

Compression I -- Move-to-Front, Distance Coding, and Inversion
 Frequencies Revisited -- A Lempel-Ziv Text Index on Secondary
 Storage -- Dynamic Rank-Select Structures with Applications to Run-
 Length Encoded Texts -- Most Burrows-Wheeler Based Compressors
 Are Not Optimal -- Session 4: Computational Biology I -- Non-
 breaking Similarity of Genomes with Gene Repetitions -- A New and
 Faster Method of Sorting by Transpositions -- Finding Compact
 Structural Motifs -- Session 5: Computational Biology II -- Improved
 Algorithms for Inferring the Minimum Mosaic of a Set of Recombinants
 -- Computing Exact p-Value for Structured Motif -- Session 6:
 Algorithmic Techniques II -- Improved Sketching of Hamming Distance
 with Error Correcting -- Deterministic Length Reduction: Fast
 Convolution in Sparse Data and Applications -- Guided Forest Edit
 Distance: Better Structure Comparisons by Using Domain-knowledge --
 Space-Efficient Algorithms for Document Retrieval -- Session 7: Data
 Compression II -- Compressed Text Indexes with Fast Locate --
 Processing Compressed Texts: A Tractability Border -- Session 8:
 Computational Biology III -- Common Structured Patterns in Linear
 Graphs: Approximation and Combinatorics -- Identification of
 Distinguishing Motifs -- Algorithms for Computing the Longest
 Parameterized Common Subsequence -- Fixed-Parameter Tractability
 of the Maximum Agreement Supertree Problem -- Session 9: Pattern
 Analysis -- Two-Dimensional Range Minimum Queries -- Tiling
 Periodicity -- Fast and Practical Algorithms for Computing All the Runs
 in a String -- Longest Common Separable Pattern Among Permutations
 -- Session 10: Suffix Arrays and Trees -- Suffix Arrays on Words --
 Efficient Computation of Substring Equivalence Classes with Suffix
 Arrays -- A Simple Construction of Two-Dimensional Suffix Trees in
 Linear Time.

Sommario/riassunto

The papers contained in this volume were presented at the 18th Annual
 Symposium on Combinatorial Pattern Matching (CPM 2007) held at the
 University of Western Ontario, in London, Ontario, Canada from July 9
 to 11, 2007. All the papers presented at the conference are original
 research contributions on computational pattern matching and analysis,
 data compression and compressed text processing, suffix arrays and
 trees, and computational biology. They were selected from 64
 submissions. Each submission was reviewed by at least three reviewers.
 The committee decided to accept 32 papers. The programme also
 included three invited talks by Tao Jiang from the University of
 California, Riverside, USA, S. Muthukrishnan from Rutgers University,
 USA, and Frances Yao from City University of Hong Kong, Hong Kong.
 Combinatorial Pattern Matching addresses issues of searching and
 matching strings and more complicated pattern structures such as trees,
 regular expressions, graphs, point sets, and arrays. The goal is to derive
 non-trivial combinatorial properties of such structures and to exploit
 these properties in order to either achieve superior performance for the
 corresponding computational problems or pinpoint conditions under
 which searches cannot be performed efficiently.

2. Record Nr.	UNISALENTO991004284337407536
Autore	Faretra, Giuseppe
Titolo	Esercizi e complementi di analisi infinitesimale : anno accademico 1944-45 / Giuseppe Faretra
Pubbl/distr/stampa	Città di Castello ; Bari : L. Macri, 1945
Descrizione fisica	180 p. ; 26 cm
Classificazione	LC QA300-QA316 AMS 26-XX
Disciplina	515.33
Soggetti	Calculus - Testbooks
Lingua di pubblicazione	Italiano
Formato	Materiale a stampa
Livello bibliografico	Monografia
Note generali	On title page: R. Università degli studi, Bari