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Nota di contenuto	CPM's 20th Anniversary: A Statistical Retrospective -- Quasi-distinct Parsing and Optimal Compression Methods -- Generalized Substring Compression -- Text Indexing, Suffix Sorting, and Data Compression: Common Problems and Techniques -- Contracted Suffix Trees: A Simple and Dynamic Text Indexing Data Structure -- Linear Time Suffix Array Construction Using D-Critical Substrings -- On the Value of Multiple Read/Write Streams for Data Compression -- Reoptimization of the Shortest Common Superstring Problem -- LCS Approximation via

Embedding into Local Non-repetitive Strings -- An Efficient Matching Algorithm for Encoded DNA Sequences and Binary Strings -- Fast Searching in Packed Strings -- New Complexity Bounds for Image Matching under Rotation and Scaling -- Online Approximate Matching with Non-local Distances -- Faster and Space-Optimal Edit Distance “1” Dictionary -- Approximate Matching for Run-Length Encoded Strings Is 3sum-Hard -- Modeling and Algorithmic Challenges in Online Social Networks -- Permuted Longest-Common-Prefix Array -- Periodic String Comparison -- Deconstructing Intractability: A Case Study for Interval Constrained Coloring -- Maximum Motif Problem in Vertex-Colored Graphs -- Fast RNA Structure Alignment for Crossing Input Structures -- Sparse RNA Folding: Time and Space Efficient Algorithms -- Multiple Alignment of Biological Networks: A Flexible Approach -- Graph Mining: Patterns, Generators and Tools -- Level-k Phylogenetic Networks Are Constructable from a Dense Triplet Set in Polynomial Time -- The Structure of Level-k Phylogenetic Networks -- Finding All Sorting Tandem Duplication Random Loss Operations -- Average-Case Analysis of Perfect Sorting by Reversals -- Statistical Properties of Factor Oracles -- Haplotype Inference Constrained by Plausible HaplotypeData -- Efficient Inference of Haplotypes from Genotypes on a Pedigree with Mutations and Missing Alleles (Extented Abstract).

Sommario/riassunto

It is our great pleasure to introduce the proceedings of the 20th anniversary edition of the Annual Symposium on Combinatorial Pattern Matching (CPM). The meeting was held in Lille, France, hosted by the Laboratoire d’Informatique Fondamentale de Lille (LIFL) affiliated with the Université de Lille 1 and the French Centre National de Recherche Scientifique (CNRS), as well as by INRIA Lille - Nord Europe. Started in 1990 as a summer school with about 30 invited participants, CPM quickly evolved into a representative annual international conference. Principally motivated by combinatorial algorithms for search problems in strings (texts, sequences), the scope of CPM extended to more complex data structures such as trees, graphs, two-dimensional arrays, or sets of points. Those studies resulted in a rich collection of algorithmic techniques and data structures, making bridges to other parts of the theory of discrete algorithms and algorithm engineering. Today, the area of combinatorial pattern matching is a well-identified active field of algorithmic research. Importantly, this development has been fertilized by a number of major application areas providing direct motivations and fruitful feedback to the CPM problematics. Those applications include data compression, computational biology, Internet search, data mining, information retrieval, coding, natural language processing, pattern recognition, music analysis, and others. On the one hand, all these areas make use of combinatorial pattern matching techniques and, on the other hand, raise new pattern matching problems. For example, the fast progress in computational molecular biology, triggered in the 1990s by the availability of mass genomic data, considerably influenced the combinatorial pattern matching field: as an illustration, about one-third of the papers presented in this volume deal with problems related to bioinformatics applications.
