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| Nota di contenuto | Shift-And Approach to Pattern Matching in LZW Compressed Text -- A General Practical Approach to Pattern Matching over Ziv-Lempel Compressed Text -- Pattern Matching in Text Compressed by Using Antidictionaries -- On the Structure of Syntenic Distance -- Physical Mapping with Repeated Probes: The Hypergraph Superstring Problem -- Hybridization and Genome Rearrangement -- On the Complexity of |

Positional Sequencing by Hybridization -- GESTALT: Genomic Steiner Alignments -- Bounds on the Number of String Subsequences -- Approximate Periods of Strings -- Finding Maximal Pairs with Bounded Gap -- A Dynamic Data Structure for Reverse Lexicographically Sorted Prefixes -- A New Indexing Method for Approximate String Matching -- The Compression of Subsegments of Images Described by Finite Automata -- Ziv Lempel Compression of Huge Natural Language Data Tries Using Suffix Arrays -- Matching of Spots in 2D Electrophoresis Images. Point Matching Under Non-uniform Distortions -- Applying an Edit Distance to the Matching of Tree Ring Sequences in Dendrochronology -- Fast Multi-dimensional Approximate Pattern Matching -- Finding Common RNA Secondary Structures from RNA Sequences -- Finding Common Subsequences with Arcs and Pseudoknots -- Computing Similarity between RNA Structures.
