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Nota di bibliografia	Includes bibliographical references and index.
Nota di contenuto	Preface; Contents; Chapter 1 Exploring RNA Intermediate Conformations with the Massively Parallel Genetic Algorithm; Chapter 2 Introduction to Self-Assembling DNA Nanostructures for Computation and Nanofabrication; Chapter 3 Mapping Sequence to Rice FPC; Chapter 4 Graph Theoretic Sequence Clustering Algorithms and Their Applications to Genome Comparison; Chapter 5 The Protein Information Resource for Functional Genomics and Proteomics; Chapter 6 High-Grade Ore for Data Mining in 3D Structures; Chapter 7 Protein Classification: A Geometric Hashing Approach Chapter 8 Interrelated Clustering: An Approach for Gene Expression Data Analysis Chapter 9 Creating Metabolic Network Models using Text Mining and Expert Knowledge; Chapter 10 Phyloinformatics and Tree Networks; Index
Sommario/riassunto	This book contains articles written by experts on a wide range of topics that are associated with the analysis and management of biological information at the molecular level. It contains chapters on RNA and

protein structure analysis, DNA computing, sequence mapping, genome comparison, gene expression data mining, metabolic network modeling, and phyloinformatics.
