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Autore	Carroll Lewis <1832-1898, >
Titolo	Alice's adventures in Wonderland / / by Lewis Carroll ; illustrations by John Tenniel
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ISBN	1-4677-6830-8
Descrizione fisica	1 online resource (128 pages)
Collana	First Avenue classics
Disciplina	823.8
Lingua di pubblicazione	Inglese
Formato	Materiale a stampa
Livello bibliografico	Monografia
Nota di contenuto	Cover -- Title Page -- Copyright Page -- Table Of Contents -- Chapter I. Down The Rabbit-Hole -- Chapter II. The Pool Of Tears -- Chapter III. A Caucus-Race And A Long Tale -- Chapter IV. The Rabbit Sends In A Little Bill -- Chapter V. Advice From A Caterpillar -- Chapter VI. Pig And Pepper -- Chapter VII. A Mad Tea-Party -- Chapter VIII. The Queen's Croquet-Ground -- Chapter IX. The Mock Turtle's Story -- Chapter X. The Lobster Quadrille -- Chapter XI. Who Stole The Tarts? -- Chapter XII. Alice's Evidence -- Back Cover.

2. Record Nr.	UNINA9910484907103321
Titolo	Systems Biology and Regulatory Genomics : Joint Annual RECOMB 2005 Satellite Workshops on Systems Biology and on Regulatory Genomics, San Diego, CA, USA, December 2-4, 2005, Revised Selected Papers / / edited by Eleazar Eskin, Trey Ideker, Ben Raphael, Christopher Workman
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Edizione	[1st ed. 2006.]
Descrizione fisica	1 online resource (X, 261 p.)
Collana	Lecture Notes in Bioinformatics, , 2366-6331 ; ; 4023
Disciplina	572.865
Soggetti	Biochemistry Algorithms Computer science - Mathematics Discrete mathematics Artificial intelligence - Data processing Database management Bioinformatics Discrete Mathematics in Computer Science Data Science Database Management
Lingua di pubblicazione	Inglese
Formato	Materiale a stampa
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Note generali	Includes index.
Nota di bibliografia	Includes bibliographical references and index.
Nota di contenuto	An Interactive Map of Regulatory Networks of Pseudomonas aeruginosa Genome -- The Pathalyzer: A Tool for Analysis of Signal Transduction Pathways -- Decomposition of Overlapping Protein Complexes: A Graph Theoretical Method for Analyzing Static and Dynamic Protein Associations -- Comparison of Protein-Protein Interaction Confidence Assignment Schemes -- Characterization of the Effects of TF Binding Site Variations on Gene Expression Towards Predicting the Functional Outcomes of Regulatory SNPs -- Comparative Systems Biology of the

Sporulation Initiation Network in Prokaryotes -- Improvement of Computing Times in Boolean Networks Using Chi-square Tests -- Build a Dictionary, Learn a Grammar, Decipher Stegoscripts, and Discover Genomic Regulatory Elements -- Causal Inference of Regulator-Target Pairs by Gene Mapping of Expression Phenotypes -- Examination of the tRNA Adaptation Index as a Predictor of Protein Expression Levels -- Improved Duplication Models for Proteome Network Evolution -- Application of Expectation Maximization Clustering to Transcription Factor Binding Positions for cDNA Microarray Analysis -- Combinatorial Genetic Regulatory Network Analysis Tools for High Throughput Transcriptomic Data -- Topological Robustness of the Protein-Protein Interaction Networks -- A Bayesian Approach for Integrating Transcription Regulation and Gene Expression: Application to *Saccharomyces Cerevisiae* Cell Cycle Data -- Probabilistic in Silico Prediction of Protein-Peptide Interactions -- Improved Pattern-Driven Algorithms for Motif Finding in DNA Sequences -- Annotation of Promoter Regions in Microbial Genomes Based on DNA Structural and Sequence Properties -- An Interaction-Dependent Model for Transcription Factor Binding -- Computational Characterization and Identification of Core Promoters of MicroRNA Genes in *C. elegans*, *H. sapiens* and *A. thaliana* -- A Comprehensive Kinetic Model of the Exocytotic Process: Evaluation of the Reaction Mechanism.

Sommario/riassunto

This book constitutes the thoroughly refereed post-proceedings of two joint RECOMB 2005 satellite events - the First Annual Workshop on Systems Biology, RSB 2005, and the Second Annual Workshop on Regulatory Genomics, RRG 2005, held in San Diego, CA, USA in December 2005. The 23 revised full papers presented were carefully reviewed and selected for inclusion in the book. The papers address a broad variety of topics in systems biology and regulatory genomics including inference of gene regulatory and protein signaling networks, model prediction of drug mechanism, pathway mapping and evolution in protein interaction networks, multi-scale methods which bridge abstract and detailed models, systematic design of genome-scale experiments, modeling and recognition of regulatory elements, identification and modeling of cis-regulatory regions, modeling the structure and function of the regulatory region, and comparative genomics of regulation.
