1.	Record Nr.	UNISA996198272003316
	Titolo	Pattern Recognition in Bioinformatics [[electronic resource]] : 9th IAPR International Conference, PRIB 2014, Stockholm, Sweden, August 21-23, 2014. Proceedings / / edited by Matteo Comin, Lukas Käll, Elena Marchiori, Alioune Ngom, Jagath Chandana Rajapakse
	Pubbl/distr/stampa	Cham : , : Springer International Publishing : , : Imprint : Springer, , 2014
	ISBN	3-319-09192-1
	Edizione	[1st ed. 2014.]
	Descrizione fisica	1 online resource (XII, 135 p. 29 illus.)
	Collana	Lecture Notes in Bioinformatics ; ; 8626
	Disciplina	570.285
	Soggetti	Bioinformatics
		Health informatics
		Pattern recognition
		Data mining
		Algorithms
		Artificial intelligence
		Computational Biology/Bioinformatics
		Health Informatics
		Pattern Recognition
		Data Mining and Knowledge Discovery
		Algorithm Analysis and Problem Complexity
		Artificial Intelligence
	Lingua di pubblicazione	Inglese
	Formato	Materiale a stampa
	Livello bibliografico	Monografia
	Note generali	Includes index.
	Nota di contenuto	FULL PAPERS -- Acquiring Decision Rules for Predicting Ames-Negative Hepatocarcinogens Using Chemical-Chemical Interactions -- Using Topology Information for Protein-Protein Interaction Prediction -- Biases of drug{target interaction network data -- Logol: Expressive Pattern Matching in sequences Application to Ribosomal Frameshift Modeling -- Evolutionary Algorithm based on New Crossover for the Biclustering of Gene Expression Data -- SFFS-SW: A feature selection algorithm exploring the small-world properties of GNs -- CytomicsDB:

A Metadata-based storage and retrieval approach for High-Throughput Screening Experiments -- CUDAGRN: Parallel Speedup of Inferring Large Gene Regulatory -- Networks from Expression Data Using Random Forest -- SHORT ABSTRACTS -- Analysis of miRNA expression profiles in breast cancer using biclustering -- Gram-positive and Gram-negative Subcellular Localization Using Rotation Forest and Physicochemical-based Features -- Data Driven Feature Selection for RNA-Seq Differential Expression Analysis -- Intramuscular fat percentage estimation through ultrasound images -- An integrated approach of gene expression and DNA-methylation profiles of WNT signaling genes uncovers novel prognostic markers in Acute Myeloid Leukemia -- Improving performance of the eXtasy model by hierarchical sampling -- Popovic et al.Ensemble Neural Networks Scoring Functions for Accurate Binding Affinity -- Prediction of Protein-Ligand Complexes -- Integration of Gene Expression and DNA-methylation Profiles Improves Molecular Subtype Classification in Acute Myeloid Leukemia -- The Relative Vertex-to-Vertex Clustering Value-A New Criterion for the Fast Detection of Functional Modules in Protein Interaction Networks.

Sommario/riassunto	This book constitutes the refereed proceedings of the 8th IAPR International Conference on Pattern Recognition in Bioinformatics, PRIB 2014, held in Stockholm, Sweden in August 2014. The 9 revised full papers and 9 revised short papers presented were carefully reviewed and selected from 29 submissions. The focus of the conference was on the latest Research in Pattern Recognition and Computational Intelligence-Based Techniques Applied to Problems in Bioinformatics and Computational Biology.

2.	Record Nr.	UNINA9910783415403321
	Titolo	WebSphere commerce V5.4 catalog design and content management [[electronic resource] /] / [John Ganci ... et al.]
	Pubbl/distr/stampa	[S.l.], : IBM, International Technical Support Organization, 2003
	Edizione	[1st ed.]
	Descrizione fisica	xviii, 508 p. : ill
	Collana	IBM redbooks
	Altri autori (Persone)	GanciJohn
	Disciplina	658.8/72/02854678
	Soggetti	Electronic commerce - Computer programs
		Web servers
		Business enterprises - Computer programs
	Lingua di pubblicazione	Inglese
	Formato	Materiale a stampa
	Livello bibliografico	Monografia
	Note generali	"WebSphere software"--Cover.
		"January 2003."
		"SG24-6855-00."
	Nota di bibliografia	Includes bibliographical references and index.