

1. Record Nr.	UNISALENTO991000195439707536
Autore	INdAM meeting Mathematical models for smart materials <2001 ;
Titolo	Cortona, Italy> Mathematical models and methods for smart materials : Cortona, Italy, 25-29 June 2001 / editors Mauro Fabrizio, Barbara Lazzari, Angelo Morro
Pubbl/distr/stampa	River Edge, N. J. : World Scientific, c2002
ISBN	9812382356
Descrizione fisica	xv, 374 p. : ill. ; 24 cm
Collana	Series on advances in mathematics for applied sciences ; 62
Classificazione	AMS 74M05 LC QA401.I5
Altri autori (Persone)	Fabrizio, Mauro Morro, Angelo Lazzari, Barbara
Altri autori (Enti)	Istituto nazionale di alta matematica Francesco Severi
Disciplina	629.8312
Soggetti	Mathematical models - Mathematical models - Congresses Smart structures Functions (Mathematics)
Lingua di pubblicazione	Inglese
Formato	Materiale a stampa
Livello bibliografico	Monografia
Nota di bibliografia	Includes bibliographical references

2. Record Nr.	UNINA9910784810403321
Autore	Raychaudhuri Soumya
Titolo	Computational text analysis for functional genomics and bioinformatics [[electronic resource] /] / Soumya Raychaudhuri
Pubbl/distr/stampa	Oxford, : Oxford University Press, 2006
ISBN	0-19-191670-6 1-280-96507-X 9786610965076 0-19-151377-6 1-4294-7040-2
Descrizione fisica	1 online resource (325 p.)
Collana	Oxford scholarship online
Disciplina	572.86
Soggetti	Bioinformatics Computational biology Genomics - Data processing
Lingua di pubblicazione	Inglese
Formato	Materiale a stampa
Livello bibliografico	Monografia
Note generali	Previously issued in print: 2006.
Nota di bibliografia	Includes bibliographical references and index.
Nota di contenuto	Contents; List of Figures; List of Plates; List of Tables; 1 An introduction to text analysis in genomics; 2 Functional genomics; 3 Textual profiles of genes; 4 Using text in sequence analysis; 5 Text-based analysis of a single series of gene expression measurements; 6 Analyzing groups of genes; 7 Analyzing large gene expression data sets; 8 Using text classification for gene function annotation; 9 Finding gene names; 10 Protein interaction networks; 11 Conclusion; Index
Sommario/riassunto	This work brings together the two disparate worlds of computational text analysis and biology with careful and thorough examples on real genomics data.