

1. Record Nr.	UNISANNIOBVVEE022893		
Autore	Maximus : Tyrius		
Titolo	ÂMaximi Tyrii philosophi Platonici Âsermones siue disputationes 41. Ex Cosmi Paccii archiepiscopi Florentini interpretatione, ab Henrico Stephano quamplurimis in locis emendata		
Pubbl/distr/stampa	\Ginevra! : ex officina Henrici Stephani Parisiensis typographi, 1557		
Descrizione fisica	\16!, 320 p. ; 8Â°		
Collocazione	BUZ.A. 0136		
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Lingua di pubblicazione	Latino		
Formato	Materiale a stampa		
Livello bibliografico	Monografia		
Note generali	Nel tit. 41 espresso XLI Marca (X 89) sul front Cors. ; gr. ; rom Segn.: A-Xâ,.		

2. Record Nr.	UNISA996466010803316
Titolo	Comparative Genomics [[electronic resource]] : RECOMB 2004 International Workshop, RCG 2004, Bertinoro, Italy, October 16-19, 2004, Revised Selected Papers // edited by Jens Lagergren
Pubbl/distr/stampa	Berlin, Heidelberg : , : Springer Berlin Heidelberg : , : Imprint : Springer, , 2005
Edizione	[1st ed. 2005.]
Descrizione fisica	1 online resource (VIII, 134 p.)
Collana	Lecture Notes in Bioinformatics ; ; 3388
Disciplina	572
Soggetti	Biochemistry Algorithms Computer science—Mathematics Data structures (Computer science) Database management Bioinformatics Biochemistry, general Algorithm Analysis and Problem Complexity Discrete Mathematics in Computer Science Data Structures Database Management
Lingua di pubblicazione	Inglese
Formato	Materiale a stampa
Livello bibliografico	Monografia
Note generali	Bibliographic Level Mode of Issuance: Monograph
Nota di bibliografia	Includes bibliographical references and index.
Nota di contenuto	Conservation of Combinatorial Structures in Evolution Scenarios -- Toward a Phylogenetically Aware Algorithm for Fast DNA Similarity Search -- Multiple Genome Alignment by Clustering Pairwise Matches -- On the Structure of Reconciliations -- The Statistical Significance of Max-Gap Clusters -- Identifying Evolutionarily Conserved Segments Among Multiple Divergent and Rearranged Genomes -- Genome Rearrangement in Mitochondria and Its Computational Biology -- The Distribution of Inversion Lengths in Bacteria -- Estimators of Translocations and Inversions in Comparative Maps -- Databases for Comparative Analysis of Human-Mouse Orthologous Alternative Splicing.

