

1. Record Nr.	UNISALENT0991003689689707536
Autore	Peterson, Douglas L.
Titolo	The English lyric from Wyatt to Donne : A history of the plain and eloquent styles
Pubbl/distr/stampa	Princeton : Princeton university press, 1967
Descrizione fisica	VI, 391 p. ; 22 cm
Disciplina	821.04
Soggetti	Poesia inglese - Lirica - Storia
Lingua di pubblicazione	Inglese
Formato	Materiale a stampa
Livello bibliografico	Monografia
2. Record Nr.	UNINA9910483429603321
Titolo	Computational Methods in Systems Biology : International Conference, CMSB 2006, Trento, Italy, October 18-19, 2006, Proceedings / / edited by Corrado Priami
Pubbl/distr/stampa	Berlin, Heidelberg : , : Springer Berlin Heidelberg : , : Imprint : Springer, , 2006
ISBN	3-540-46167-1
Edizione	[1st ed. 2006.]
Descrizione fisica	1 online resource (X, 323 p.)
Collana	Lecture Notes in Bioinformatics, , 2366-6331 ; ; 4210
Altri autori (Persone)	PriamiCorrado
Disciplina	572.80285
Soggetti	Bioinformatics Computer simulation Software engineering Database management Computational and Systems Biology Computer Modelling Software Engineering Database Management
Lingua di pubblicazione	Inglese
Formato	Materiale a stampa

Livello bibliografico**Note generali****Nota di bibliografia****Nota di contenuto****Monografia****Bibliographic Level Mode of Issuance: Monograph**

Includes bibliographical references and index.

Modal Logics for Brane Calculus -- Deciding Behavioural Properties in Brane Calculi -- Probabilistic Model Checking of Complex Biological Pathways -- Type Inference in Systems Biology -- Stronger Computational Modelling of Signalling Pathways Using Both Continuous and Discrete-State Methods -- A Formal Approach to Molecular Docking -- Feedbacks and Oscillations in the Virtual Cell VICE -- Modelling Cellular Processes Using Membrane Systems with Peripheral and Integral Proteins -- Modelling and Analysing Genetic Networks: From Boolean Networks to Petri Nets -- Regulatory Network Reconstruction Using Stochastic Logical Networks -- Identifying Submodules of Cellular Regulatory Networks -- Incorporating Time Delays into the Logical Analysis of Gene Regulatory Networks -- A Computational Model for Eukaryotic Directional Sensing -- Modeling Evolutionary Dynamics of HIV Infection -- Compositional Reachability Analysis of Genetic Networks -- Randomization and Feedback Properties of Directed Graphs Inspired by Gene Networks -- Computational Model of a Central Pattern Generator -- Rewriting Game Theory as a Foundation for State-Based Models of Gene Regulation -- Condition Transition Analysis Reveals TF Activity Related to Nutrient-Limitation-Specific Effects of Oxygen Presence in Yeast -- An In Silico Analogue of In Vitro Systems Used to Study Epithelial Cell Morphogenesis -- A Numerical Aggregation Algorithm for the Enzyme-Catalyzed Substrate Conversion -- Possibilistic Approach to Bioclustering: An Application to Oligonucleotide Microarray Data Analysis.

Sommario/riassunto

This book constitutes the refereed proceedings of the International Conference on Computational Methods in Systems Biology, CMSB 2006, held in Trento, Italy, in October 2006. The 22 fully revised papers presented together with 2 invited talks were carefully reviewed and selected from 68 submissions. The papers present a variety of techniques from computer sciences, such as language design, concurrency theory, software engineering, and formal methods.