Record Nr. UNINA9910483429603321 Computational Methods in Systems Biology: International Conference, **Titolo** CMSB 2006, Trento, Italy, October 18-19, 2006, Proceedings / / edited by Corrado Priami Berlin, Heidelberg:,: Springer Berlin Heidelberg:,: Imprint: Springer, Pubbl/distr/stampa **ISBN** 3-540-46167-1 Edizione [1st ed. 2006.] 1 online resource (X, 323 p.) Descrizione fisica Collana Lecture Notes in Bioinformatics;; 4210 Disciplina 572.80285 Soggetti **Bioinformatics** Computer simulation **Bioinformatics** Computational biology Software engineering Database management Computational Biology/Bioinformatics Simulation and Modeling Computer Appl. in Life Sciences Software Engineering **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 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 Biclustering: An Application to Oligonucleotide Microarray Data Analysis.