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| 1. Record Nr. | UNISA996465967203316 |
| Titolo | Transactions on Computational Systems Biology III [[electronic resource]] / edited by Corrado Priami, Emanuela Merelli, Pedro Pablo Gonzalez, Andrea Omicini |
| Pubbl/distr/stampa | Berlin, Heidelberg : , : Springer Berlin Heidelberg : , : Imprint : Springer, , 2005 |
| ISBN | 3-540-31446-6 |
| Edizione | [1st ed. 2005.] |
| Descrizione fisica | 1 online resource (VII, 169 p.) |
| Collana | Transactions on Computational Systems Biology, , 1861-2075 ; ; 3737 |
| Disciplina | 004.0151 |
| Soggetti | Computers Bioinformatics Proteomics Database management Computation by Abstract Devices 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 | Computer-Aided DNA Base Calling from Forward and Reverse Electropherograms -- A Multi-agent System for Protein Secondary Structure Prediction -- Modeling Kohn Interaction Maps with Beta-Binders: An Example -- Multidisciplinary Investigation into Adult Stem Cell Behavior -- Statistical Model Selection Methods Applied to Biological Networks -- Using Secondary Structure Information to Perform Multiple Alignment -- Frequency Concepts and Pattern Detection for the Analysis of Motifs in Networks -- An Agent-Oriented Conceptual Framework for Systems Biology -- Genetic Linkage Analysis Algorithms and Their Implementation -- Abstract Machines of Systems Biology. |