1. Record Nr. UNISANNIORMG0099405

Autore	Nazzaro, Anna Carla
Titolo	Oggettività giuridica dei beni produttivi : contributo allo studio dei rapporti di affitto / Anna Carla Nazzaro
Pubbl/distr/stampa	Napoli\etc.!, : Edizioni scientifiche italiane, 2002
ISBN	8849505329
Descrizione fisica	202 p. ; 24 cm.
Collana	Quaderni della Rassegna di diritto civile
Disciplina	346.04346
Soggetti	Locazione - Legislazione
Collocazione	D (C) 22 33701D (C) 22 338
Lingua di pubblicazione	Italiano
Formato	Materiale a stampa
Livello bibliografico	Monografia

2. Record Nr. UNINA9910304142603321

Autore	Will Thorsten
Titolo	Predicting Transcription Factor Complexes : A Novel Approach to Data Integration in Systems Biology / / by Thorsten Will
Pubbl/distr/stampa	Wiesbaden : , : Springer Fachmedien Wiesbaden : , : Imprint : Springer Spektrum, , 2015
ISBN	3-658-08269-0
Edizione	[1st ed. 2015.]
Descrizione fisica	1 online resource (155 p.)
Collana	BestMasters, , 2625-3577
Disciplina	570 570.285 570285
Soggetti	Bioinformatics Computational biology Biomathematics Computer Appl. in Life Sciences Mathematical and Computational Biology
Lingua di pubblicazione	Inglese
Formato	Materiale a stampa
Livello bibliografico	Monografia
Note generali	Description based upon print version of record.
Nota di bibliografia	Includes bibliographical references.
Nota di contenuto	Protein Complex Prediction -- Protein-Protein Interaction Networks -- Domain-Domain Interaction Networks -- Combinatorial Algorithms -- Algorithm Engineering.
Sommario/riassunto	In his master thesis Thorsten Will proposes the substantial information content of protein complexes involving transcription factors in the context of gene regulatory networks, designs the first computational approaches to predict such complexes as well as their regulatory function and verifies the practicability using data of the well-studied yeast S.cereviseae. The novel insights offer extensive capabilities towards a better understanding of the combinatorial control driving transcriptional regulation. Contents Protein Complex Prediction Protein-Protein Interaction Networks Domain-Domain Interaction Networks Combinatorial Algorithms Algorithm Engineering Target Groups Computational biologists and biologists working with gene regulatory networks Computer scientists interested in biological issues The Author Currently, the author is pursuing his Ph.D. at the