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| 1. Record Nr.           | UNISA990000555260203316                      |
| Autore                  | BACH, Emmon                                  |
| Titolo                  | An introduction to transformational grammars |
| Pubbl/distr/stampa      | New York : Holt- Rinehart- Winston, 1964     |
| Descrizione fisica      | X, 205 p. : ill ; 21 cm                      |
| Disciplina              | 401.   |
| Collocazione            | 401 BAC<br>IV.2. 383(XII W 13)               |
| Lingua di pubblicazione | Inglese                                      |
| Formato                 | Materiale a stampa                           |
| Livello bibliografico   | Monografia                                   |
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| 2. Record Nr.      | UNINA9910877186203321  |
| Titolo             | Structural bioinformatics  |
| Pubbl/distr/stampa | [Place of publication not identified], : Wiley Blackwell, 2009   |
| ISBN               | 1-280-55693-5<br>9786610556939<br>0-471-32634-8<br>0-471-72120-4   |
| Descrizione fisica | 1 online resource (674 pages)  |
| Collana            | Methods of biochemical analysis Structural bioinformatics  |
| Disciplina         | 572.8/733  |
| Soggetti           | Structural bioinformatics<br>Molecular Biology<br>Computational Biology<br>Biochemistry<br>Biology<br>Genetics<br>Biological Science Disciplines<br>Chemistry<br>Natural Science Disciplines<br>Disciplines and Occupations<br>Biology - General<br>Health & Biological Sciences |

Lingua di pubblicazione	Inglese
Formato	Materiale a stampa
Livello bibliografico	Monografia
Note generali	Bibliographic Level Mode of Issuance: Monograph
Sommario/riassunto	<p>From the Foreword: "[A] must read for all of us committed to understanding the interplay of structure and function...[T]he individual chapters outline the suite of major basic life science questions, such as the status of efforts to predict protein structure and how proteins carry out cellular functions, and also the applied life science questions such as how structural bioinformatics can improve health care through accelerating drug discovery." This book provides a basic understanding of the theories, associated algorithms, resources, and tools used in structural bioinformatics. The reader emerges with the ability to make effective use of protein, DNA, RNA, carbohydrate, and complex structures to better understand biological function. Moreover, it draws a clear connection between structural studies and the rational design of new therapies.</p>