Record Nr. UNINA9910437615903321 Autore Masoudi-Nejad Ali **Titolo** Next generation sequencing and sequence assembly: methodologies and algorithms / / Ali Masoudi-Nejad, Zahra Narimani, Nazanin Hosseinkhan New York, : Springer, 2013 Pubbl/distr/stampa **ISBN** 1-4614-7726-3 Edizione [1st ed. 2013.] Descrizione fisica 1 online resource (92 p.) Collana SpringerBriefs in systems biology Altri autori (Persone) NarimaniZahra HosseinkhanNazanin Disciplina 572.863 Soggetti Nucleotide sequence Human gene mapping Lingua di pubblicazione Inglese **Formato** Materiale a stampa Livello bibliografico Monografia Note generali Description based upon print version of record. Includes bibliographical references and index. Nota di bibliografia Nota di contenuto Preface -- Next generation sequencing methodologies -- Emergence of next generation sequencing -- Assembly of sequencing data -- De novo assembly algorithms -- Index. The goal of this book is to introduce the biological and technical Sommario/riassunto aspects of next generation sequencing methods, as well as algorithms to assemble these sequences into whole genomes. The book is organized into two parts; part 1 introduces NGS methods and part 2 reviews assembly algorithms and gives a good insight to these methods for readers new to the field. Gathering information, about sequencing and assembly methods together, helps both biologists and computer

scientists to get a clear idea about the field. Chapters will include information about new sequencing technologies such as Chlp-seq,

Chlp-chip, and De Novo sequence assembly.