

1. Record Nr.	UNINA9910463010203321
Titolo	The analysis of regulatory DNA : current developments, knowledge and applications uncovering gene regulation / / Kenneth Wayne Berendzen, Joachim Kilian, Dierk Wanke, editors
Pubbl/distr/stampa	Sharjah : , : Bentham Science Publishers, , 2013
ISBN	1-60805-492-6
Descrizione fisica	1 online resource (225 p.)
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Soggetti	DNA - Analysis DNA Electronic books.
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 and index.
Nota di contenuto	Cover; Title; EUL; Content; Foreword; Preface; List of Contributors; Chapter 01; Chapter 02; Chapter 03; Chapter 04; Chapter 05; Chapter 06; Chapter 07; Chapter 08; Chapter 09; Chapter 10; Index
Sommario/riassunto	This e-book provides a resource for summarizing current knowledge eukaryotic transcription and explores cis-elements and methods for their analysis, prediction and discovery. The book also presents an overview of exploring gene regulatory networks, chromatin, and miRNAs. Information about state-of-the-art techniques for the determination of TF - cis-element interactions in vivo and in silico give cutting edge insights on how genomic-scale research is being approached