

1. Record Nr.	UNINA9910462946703321
Autore	Verney Kevern <1960, >
Titolo	African Americans and US popular culture / / Kevern Verney
Pubbl/distr/stampa	London ; ; New York : , : Routledge, , 2003
ISBN	1-136-47527-3 0-415-27528-8 1-315-01531-5
Descrizione fisica	1 online resource (138 p.)
Collana	Introductions to history
Disciplina	305.896/073
Soggetti	African Americans in popular culture African Americans - Race identity African Americans - Intellectual life Racism in popular culture - United States Popular culture - United States - History Electronic books. United States Race relations United States Civilization African American influences
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 (p. 115-120) and index.
Nota di contenuto	Cover; African Americans and US Popular Culture; Title Page; Copyright Page; Table of Contents; Preface; Introduction: emancipation and segregation; 1. Migration and urbanization, 1915-30; 2. The Great Depression and the Second World War, 1930-45; 3. The Civil Rights era, 1945-65; 4. Black Power, 1965-76; 5. African Americans in US society since 1976; Conclusion: from Ragtime to Rap; Further reading; Select bibliography; Index
Sommario/riassunto	This volume is an authoritative introduction to the history of African Americans in US popular culture, examining its development from the early nineteenth century to the present. Kevern Verney examines: * the role and significance of race in all major forms of popular culture, including sport, film, television, radio and music* how the entertainment industry has encouraged racism through misrepresentations and caricatured images of African Americans.

2. Record Nr.	UNINA9910451319503321
Titolo	Computational systems bioinformatics [[electronic resource]] : CSB2006 conference proceedings, Stanford CA, 14-18 August 2006 // editors, Peter Markstein, Ying Xu
Pubbl/distr/stampa	London, : Imperial College Press, c2006
ISBN	1-281-86737-3 9786611867379 1-86094-757-3
Descrizione fisica	1 online resource (416 p.)
Collana	Series on advances in bioinformatics and computational biology ; ; v. 4
Altri autori (Persone)	MarksteinPeter XuYing <1960->
Disciplina	572.80285
Soggetti	Bioinformatics Computational biology Biological systems - Computer simulation Biological systems - Simulation methods Electronic books.
Lingua di pubblicazione	Inglese
Formato	Materiale a stampa
Livello bibliografico	Monografia
Note generali	"Fifth Annual Computational Systems Bioinformatics Conference, CSB2006"--p. [v]. Sponsored by Hewlett-Packard Company and Microsoft Research--p. [v]. Previous CSB Conference (4th) held in Stanford, Calif. Aug. 8-11, 2005 (IEEE Computer Society Bioinformatics Conference). Includes bibliographical references and index.
Nota di bibliografia	
Nota di contenuto	CONTENTS ; Committees ; Referees ; Preface ; Keynote Addresses ; Exploring the Ocean's Microbes: Sequencing the Seven Seas ; Don't Know Much About Philosophy: The Confusion Over Bio- Ontologies ; Invited Talks

Biomedical Informatics Research Network (BIRN): Building a National Collaboratory for BioMedical and Brain Research
Protein Network Comparative Genomics ;
Systems Biology in Two Dimensions: Understanding and Engineering Membranes as Dynamical Systems
; Bioinformatics at Microsoft Research
Movie Crunching in Biological Dynamic Imaging
Engineering Nucleic Acid-Based Molecular Sensors for Probing and Programming Cellular Systems
; Reactome: A Knowledgebase of Biological Pathways
; Structural Bioinformatics
Effective Optimization Algorithms for Fragment-Assembly based Protein Structure Prediction
Transmembrane Helix and Topology Prediction Using Hierarchical SVM Classifiers and an Alternating Geometric Scoring Function
; Protein Fold Recognition Using the Gradient Boost Algorithm
A Graph-Based Automated NMR Backbone Resonance Sequential Assignment A Data-Driven Systematic Search Algorithm for Structure Determination of Denatured or Disordered Proteins
; Multiple Structure Alignment by Optimal RMSD Implies that the Average Structure is a Consensus
Identification of α -Helices from Low Resolution Protein Density Maps

Sommario/riassunto

This volume contains about 40 papers covering many of the latest developments in the fast-growing field of bioinformatics. The contributions span a wide range of topics, including computational genomics and genetics, protein function and computational proteomics, the transcriptome, structural bioinformatics, microarray data analysis, motif identification, biological pathways and systems, and biomedical applications. There are also abstracts from the keynote addresses and invited talks. The papers cover not only theoretical aspects of bioinformatics but also delve into the application of new
