1. Record Nr. UNINA9910462647303321

Autore	Faulkner Neil
Titolo	A Marxist history of the world [[electronic resource]] : from Neanderthals to neoliberals / / Neil Faulkner
Pubbl/distr/stampa	London, : PlutoPress, 2013
ISBN	1-84964-864-6 1-84964-863-8
Descrizione fisica	1 online resource (352 p.)
Collana	Counterfire
Disciplina	909
Soggetti	World history Marxian historiography 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.
Nota di contenuto	Hunters and farmers, c. 2.5 million-3000 BC -- The first class societies, c. 3000-1000 BC -- Ancient empires, c. 1000-30 BC -- The end of antiquity, c 30 BC-AD 650 -- The medieval world, c. AD 650-1500 -- European feudalism, c. AD 650-1500 -- THe first wave of bourgeois revolutions, 1517-1775 -- The second wave of bourgeois revolutions, 1775-1815 -- The rise of industrial capitalism, c. 1750-1850 -- The age of blood and iron, 1848-1896 -- Imperialism and war, 1873-1918 -- The revolutionary wave, 1917-1928 -- The Great Depression and the rise of fascism, 1929-1939 -- World War and Cold War, 1939-1967 -- The new world disorder, 1968-present -- Conslusion: Making the future -- Timeline.
Sommario/riassunto	This magisterial analysis of human history combines the insights of earlier generations of Marxist historians with radical new ideas about the historical process. Reading history against the grain, Neil Faulkner reveals that what happened in the past was not predetermined. Choices were frequent and numerous. Different outcomes - liberation or barbarism - were often possible. Rejecting the top-down approach of conventional history, Faulkner contends that it is the mass action of ordinary people that drives great events. At the beginning of the 21st century - with economic disaster, war, climate catastrophe and deep

	class divisions - humans face perhaps the greatest crisis in the long history of our species. The lesson of A Marxist History of the World is that, since we created our past, we can also create a better future.

2. Record Nr. UNINA9910483890403321

Titolo	Recent advances in biological network analysis : comparative network analysis and network module detection / / Byung-Jun Yoon, Xiaoning Qian, editors
Pubbl/distr/stampa	Cham, Switzerland : , : Springer, , [2021] Â©2021
ISBN	3-030-57173-4
Edizione	[1st ed. 2021.]
Descrizione fisica	1 online resource (XII, 217 p. 42 illus., 29 illus. in color.)
Disciplina	570.285
Soggetti	Bioinformatics
Lingua di pubblicazione	Inglese
Formato	Materiale a stampa
Livello bibliografico	Monografia
Nota di bibliografia	Includes bibliographical references.
Nota di contenuto	Chapter 1: Global Alignment of PPI Networks -- Chapter 2: Integrated Network-Based Computational Analysis for Drug Development -- Chapter 3: Effective Random Walk Models for Comparative Network Analysis -- Chapter 4: Computational Methods for Protein-Protein Interaction Network Alignment -- Chapter 5: Network Propagation for the Analysis of Multi_Omics Data -- Chapter 6: Motifs in Biological Networks -- Chapter 7: Bio Fabric Visualization of Network Alignments -- Chapter 8: Module Identification of Biological Networks via Graph Partition -- Chapter 9: Network Module Detection to Decipher the Heterogeneity of Cancer Mutations.
Sommario/riassunto	This book reviews recent advances in the emerging field of computational network biology with special emphasis on comparative network analysis and network module detection. The chapters in this volume are contributed by leading international researchers in computational network biology and offer in-depth insight on the latest techniques in network alignment, network clustering, and network module detection. Chapters discuss the advantages of the respective

techniques and present the current challenges and open problems in the field. Recent Advances in Biological Network Analysis: Comparative Network Analysis and Network Module Detection will serve as a great resource for graduate students, academics, and researchers who are currently working in areas relevant to computational network biology or wish to learn more about the field. Data scientists whose work involves the analysis of graphs, networks, and other types of data with topological structure or relations can also benefit from the book's insights. .