1.
Record Nr.	UNICASTO00323030
Autore	Evans, Robert J. W.
Titolo	Rodolfo 2. d'Absburgo : l'enigma di un imperatore / Robert J. W. Evans
Pubbl/distr/stampa	Bologna, : Il mulino, 1984
Titolo uniforme	Rudolf 2. and his world
ISBN	8815005625
Descrizione fisica	445 p. : c. geogr. ; 22 cm
Collana	Biblioteca storica
Disciplina	943.034
Soggetti	Rodolfo <imperatore ; 2>
Lingua di pubblicazione	Italiano
Formato	Materiale a stampa
Livello bibliografico	Monografia
Note generali	Trad. di Alfonso Prandi.

2.	Record Nr.	UNINA9910970663603321
	Titolo	Concise encyclopaedia of bioinformatics and computational biology / / edited by John M. Hancock (Department of Physiology, Development & Neuroscience University of Cambridge Cambridge, UK), Marketa J. Zvelebil (Breakthrough Breast Cancer Research Institute of Cancer Research London, UK)
	Pubbl/distr/stampa	Chichester, West Sussex, UK : , : John Wiley & Sons Ltd., , 2014
	ISBN	9780470978726 0470978724 9781118598160 1118598164
	Edizione	[Second edition]
	Descrizione fisica	1 online resource (832 pages)
	Disciplina	572.330285
	Soggetti	Bioinformatics -- Dictionaries Computational biology -- Dictionaries Systems biology -- Dictionaries Bioinformatics Computational biology Biology Health & Biological Sciences Biology - General Bioinformàtica Biologia computacional
	Lingua di pubblicazione	Inglese
	Formato	Materiale a stampa
	Livello bibliografico	Monografia
	Note generali	Description based upon print version of record. The concise encyclopaedia of bioinformatics and computational biology is a follow-up edition to the Dictionary of bioinformatics and computational biology
	Nota di contenuto	Concise Encyclopaedia of Bioinformatics and Computational Biology; Title Page; Copyright; Contents; List of Contributors; Preface; A; Ab Initio; Ab Initio Gene Prediction, see Gene Prediction, ab initio.; ABNR, see Energy Minimization.; Accuracy (of Protein Structure Prediction);

David Jones; Accuracy Measures, see Error Measures.; Adjacent Group; Aidan Budd and Alexandros Stamatakis; Admixture Mapping (Mapping by Admixture Linkage Disequilibrium); Andrew Collins, Mark McCarthy and Steven Wiltshire; Adopted-basis Newton-Raphson Minimization (ABNR), see Energy Minimization.

Affine Gap Penalty, see Gap Penalty.Affinity Propagation-based Clustering; Pedro Larranaga and Concha Bielza; Affymetrix GeneChipTM Oligonucleotide Microarray; Stuart Brown and Dov Greenbaum; Affymetrix Probe Level Analysis; Stuart Brown; After Sphere, see After State.; After State (After Sphere); Thomas D. Schneider; AIC, see Akaike Information Criterion.; Akaike Information Criterion; Pedro Larranaga and Concha Bielza; Algorithm; Matthew He; Alignment (Domain Alignment, Repeats Alignment); Jaap Heringa; Alignment Score; Laszlo Patthy; Allele-Sharing Methods (Non-parametric Linkage Analysis)

Mark McCarthy, Steven Wiltshire and Andrew CollinsAllelic Association; Mark McCarthy, Steven Wiltshire and Andrew Collins; Allen Brain Atlas; Dan Bolser; Allopatric Evolution (Allopatric Speciation); A.R. Hoelzel; Allopatric Speciation, see Allopatric Evolution.; AlogP; Bissan Al-Lazikani; Alpha carbon, see C (C-Alpha).; Alpha Helix; Roman Laskowski and Tjaart de Beer; Alternative Splicing; Enrique Blanco and Josep F. Abril; Alternative Splicing Gene Prediction, see Gene Prediction, alternative splicing.; Amide Bond (Peptide Bond); Roman Laskowski and Tjaart de Beer; Amino Acid (Residue)

Roman Laskowski, Jeremy Baum and Tjaart de BeerAmino Acid Abbreviations, see IUPAC-IUB Codes.; Amino Acid Composition; Jeremy Baum; Amino Acid Exchange Matrix (Dayhoff Matrix, Log Odds Score, PAM (Matrix), BLOSUM Matrix); Jaap Heringa; AMINO Acid Substitution Matrix, see Amino Acid Exchange Matrix.; Amino-terminus, see N-terminus.; Amphipathic; Roman Laskowski and Tjaart de Beer; Analog (Analogue); Dov Greenbaum; Ancestral Lineage, see Offspring Lineage.; Ancestral State Reconstruction; Sudhir Kumar and Alan Filipski; Anchor Points; Roland Dunbrack

Annotation Refinement Pipelines, see Gene Prediction.Annotation Transfer (Guilt by Association Annotation); Dov Greenbaum; APBIONET (Asia-Pacific Bioinformatics Network); Pedro Fernandes; Apomorphy; A.R. Hoelzel; APOLLO, see Gene Annotation, visualization tools.; Arc, see Branch (of a Phylogenetic Tree).; Are We There Yet?, see AWTY.; Aromatic; Roman Laskowski and Tjaart de Beer; Array, see Data Structure.; Artificial Neural Networks, see Neural Networks.; ASBCB (The African Society for Bioinformatics and Computational Biology); Pedro Fernandes

Association Analysis (Linkage Disequilibrium Analysis)

Sommario/riassunto

Concise Encyclopaedia of Bioinformatics and Computational Biology, 2nd Edition is a fully revised and updated version of this acclaimed resource. The book provides definitions and often explanations of over 1000 words, phrases and concepts relating to this fast-moving and exciting field, offering a convenient, one-stop summary of the core knowledge in the area. This second edition is an invaluable resource for students, researchers and academics.