



mathematics of evolution & phylogeny

EDITED BY
OLIVIER GASCUEL



Mathematics Of Evolution And Phylogeny

JA Banks



Mathematics Of Evolution And Phylogeny:

Mathematics of Evolution and Phylogeny Olivier Gascuel, 2005-02-24 This book considers evolution at different scales sequences genes gene families organelles genomes and species The focus is on the mathematical and computational tools and concepts which form an essential basis of evolutionary studies indicate their limitations and give them orientation Recent years have witnessed rapid progress in the mathematics of evolution and phylogeny with models and methods becoming more realistic powerful and complex Aimed at graduates and researchers in phylogenetics mathematicians computer scientists and biologists and including chapters by leading scientists A Bergeron D Bertrand D Bryant R Desper O Elemento N El Mabrouk N Galtier O Gascuel M Hendy S Holmes K Huber A Meade J Mixtacki B Moret E Mossel V Moulton M Pagel M A Poursat D Sankoff M Steel J Stoye J Tang L S Wang T Warnow Z Yang this book of contributed chapters explains the basis and covers the recent results in this highly topical area

Mathematics of Evolution and Phylogeny Olivier Gascuel, 2023 Presenting a consideration of evolution at the level of sequences genes gene families organelles genomes and species the central focus of this text is on the mathematical and computational tools and concepts which form an essential basis of evolutionary studies

Phylogeny Mike Steel, 2016-09-29 Phylogenetics is a topical and growing area of research Phylogenies phylogenetic trees and networks allow biologists to study and graph evolutionary relationships between different species These are also used to investigate other evolutionary processes for example how languages developed or how different strains of a virus such as HIV or influenza are related to each other This self contained book addresses the underlying mathematical theory behind the reconstruction and analysis of phylogenies The theory is grounded in classical concepts from discrete mathematics and probability theory as well as techniques from other branches of mathematics algebra topology differential equations The biological relevance of the results is highlighted throughout The author supplies proofs of key classical theorems and includes results not covered in existing books emphasizes relevant mathematical results derived over the past 20 years and provides numerous exercises examples and figures

Reconstructing Evolution Olivier Gascuel, Mike Steel, 2007-06-28 Evolution is a complex process acting at multiple scales from DNA sequences and proteins to populations of species Understanding and reconstructing evolution is of major importance in numerous subfields of biology For example phylogenetics and sequence evolution is central to comparative genomics attempts to decipher genomes and molecular epidemiology Phylogenetics is also the focal point of large scale international biodiversity assessment initiatives such as the Tree of Life project which aims to build the evolutionary tree for all extant species Since the pioneering work in phylogenetics in the 1960s models have become increasingly sophisticated to account for the inherent complexity of evolution They rely heavily on mathematics and aim at modelling and analyzing biological phenomena such as horizontal gene transfer heterogeneity of mutation and speciation and extinction processes This book presents these recent models their biological relevance their mathematical basis their properties and the algorithms to infer them from data A number of

subfields from mathematics and computer science are involved combinatorics graph theory stringology probabilistic and Markov models information theory statistical inference Monte Carlo methods continuous and discrete algorithmics This book arises from the Mathematics of Evolution Phylogenetics meeting at the Mathematical Institute Henri Poincaré Paris in June 2005 and is based on the outstanding state of the art reports presented by the conference speakers Ten chapters based around five themes provide a detailed overview of key topics from the underlying concepts to the latest results some of which are at the forefront of current research

Analysis of Phylogenetics and Evolution with R Emmanuel Paradis, 2006-11-25 As a result the inference of phylogenies often seems divorced from any connection to other methods of analysis of scientific data Felsenstein Once calculation became easy the statistician's energies could be voted to understanding his or her dataset Venables Ripley The study of the evolution of life on Earth stands as one of the most complex fields in science It involves observations from very different sources and has implications far beyond the domain of basic science It is concerned with processes occurring on very long time spans and we now know that it is also important for our daily lives as shown by the rapid evolution of many pathogens As a field ecologist for a long time I was remotely interested in phylogenetics and other approaches to evolution Most of the work I accomplished during my doctoral studies involved field studies of small mammals and estimation of demographic parameters Things changed in 1996 when my interest was attracted by the question of the effect of demographic parameters on bird diversification This was a new issue for me so I searched for relevant data analysis methods but I failed to find exactly what I needed I started to conduct my own research on this problem to propose some at least partial solutions This work made me realize that this kind of research critically depends on the available software and it was clear to me that what was offered to phylogeneticists at this time was inappropriate

Phylogenetics Charles Semple, Mike Steel, Both in the Department of Mathematics and Statistics Mike Steel, 2003 Phylogenetics is the reconstruction and analysis of phylogenetic evolutionary trees and networks based on inherited characteristics It is a flourishing area of interaction between mathematics statistics computer science and biology The main role of phylogenetic techniques lies in evolutionary biology where it is used to infer historical relationships between species However the methods are also relevant to a diverse range of fields including epidemiology ecology medicine as well as linguistics and cognitive psychology This graduate level book based on the author's lectures at The University of Canterbury New Zealand focuses on the mathematical aspects of phylogenetics It brings together the central results of the field providing proofs of the main theorem outlines their biological significance and indicates how algorithms may be derived The presentation is self contained and relies on discrete mathematics with some probability theory A set of exercises and at least one specialist topic ends each chapter This book is intended for biologists interested in the mathematical theory behind phylogenetic methods and for mathematicians statisticians and computer scientists eager to learn about this emerging area of discrete mathematics

Phylogenetics in the 24th volume in the Oxford Lecture Series in Mathematics and its Applications This series contains short

books suitable for graduate students and researchers who want a well written account of mathematics that is fundamental to current to research The series emphasises future directions of research and focuses on genuine applications of mathematics to finance engineering and the physical and biological sciences

Mathematical Approaches to Polymer Sequence Analysis and Related Problems Renato Bruni, 2010-10-17 An edited volume describing the latest developments in approaching the problem of polymer sequence analysis with special emphasis on the most relevant biopolymers peptides and DNA but not limited to them The chapters will include peptide sequence analysis DNA sequence analysis analysis of biopolymers and nonpolymers sequence alignment problems and more

Tutorials in Mathematical Biosciences IV Avner Friedman, 2007-11-21 This book offers an introduction to fast growing research areas in evolution of species population genetics ecological models and population dynamics It reviews the concept and methodologies of phylogenetic trees introduces ecological models examines a broad range of ongoing research in population dynamics and deals with gene frequencies under the action of migration and selection The book features computational schemes illustrations and mathematical theorems

Computational Phylogenetics Tandy Warnow, 2018 This book presents the foundations of phylogeny estimation and technical material enabling researchers to develop improved computational methods

Encyclopedia of Bioinformatics and Computational Biology, 2018-08-21 Encyclopedia of Bioinformatics and Computational Biology ABC of Bioinformatics Three Volume Set combines elements of computer science information technology mathematics statistics and biotechnology providing the methodology and in silico solutions to mine biological data and processes The book covers Theory Topics and Applications with a special focus on Integrative omics and Systems Biology The theoretical methodological underpinnings of BCB including phylogeny are covered as are more current areas of focus such as translational bioinformatics cheminformatics and environmental informatics Finally Applications provide guidance for commonly asked questions This major reference work spans basic and cutting edge methodologies authored by leaders in the field providing an invaluable resource for students scientists professionals in research institutes and a broad swath of researchers in biotechnology and the biomedical and pharmaceutical industries Brings together information from computer science information technology mathematics statistics and biotechnology Written and reviewed by leading experts in the field providing a unique and authoritative resource Focuses on the main theoretical and methodological concepts before expanding on specific topics and applications Includes interactive images multimedia tools and crosslinking to further resources and databases

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