COALESCENT THEORY

AN INTRODUCTION



JOHN WAKELEY

Coalescent Theory An Introduction

M.Elizabeth Halloran, Seymour Geisser

Coalescent Theory An Introduction:

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Gene Genealogies, Variation and Evolution Jotun Hein, Mikkel H. Schierup, Carsten Wiuf, 2005 Covering the growing crossover field of mathematics and biological science this text presents a practical and accessible introduction to coalescent theory

Introduction to Computational Biology Bernhard Haubold, Thomas Wiehe, 2006-08-09 Written with the advanced undergraduate in mind this book introduces into the field of Bioinformatics The authors explain the computational and conceptional background to the analysis of large scale sequence data Many of the corresponding analysis methods are rooted in evolutionary thinking which serves as a common thread throughout the book The focus is on methods of comparative genomics and subjects covered include alignments gene finding phylogeny and the analysis of single nucleotide polymorphisms SNPs The volume contains exercises questions answers to selected problems Brenner's Encyclopedia of Genetics Stanley Maloy, Kelly Hughes, 2013-03-03 The explosion of the field of genetics over the last decade with the new technologies that have stimulated research suggests that a new sort of reference work is needed to keep pace with such a fast moving and interdisciplinary field Brenner's Encyclopedia of Genetics Second Edition Seven Volume Set builds on the foundation of the first edition by addressing many of the key subfields of genetics that were just in their infancy when the first edition was published The currency and accessibility of this foundational content will be unrivalled making this work useful for scientists and non scientists alike Featuring relatively short entries on genetics topics written by experts in that topic Brenner's Encyclopedia of Genetics Second Edition Seven Volume Set provides an effective way to quickly learn about any aspect of genetics from Abortive Transduction to Zygotes Adding to its utility the work provides short entries that briefly define key terms and a guide to additional reading and relevant websites for further study Many of the entries include figures to explain difficult concepts Key terms in related areas such as biochemistry cell and molecular biology are also included and there are entries that describe historical figures in genetics providing insights into their careers and discoveries This 7 volume set represents a 25% expansion from the first edition with over 1600 articles encompassing this burgeoning field Thoroughly up to date with many new topics and subfields covered that were in their infancy or not inexistence at the time of

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Population Genetics Matthew B. Hamilton, 2021-01-26 Now updated for its second edition Population Genetics is the classic accessible introduction to the concepts of population genetics Combining traditional conceptual approaches with classical hypotheses and debates the book equips students to understand a wide array of empirical studies that are based on the first principles of population genetics Featuring a highly accessible introduction to coalescent theory as well as covering the major conceptual advances in population genetics of the last two decades the second edition now also includes end of chapter problem sets and revised coverage of recombination in the coalescent model metapopulation extinction and recolonization and the fixation index Handbook of Statistical Genomics David J. Balding, Ida Moltke, John Marioni, 2019-07-02 A timely update of a highly popular handbook on statistical genomics This new two volume edition of a classic text provides a thorough introduction to statistical genomics a vital resource for advanced graduate students early career researchers and new entrants to the field It introduces new and updated information on developments that have occurred since the 3rd edition Widely regarded as the reference work in the field it features new chapters focusing on statistical aspects of data generated by new sequencing technologies including sequence based functional assays It expands on previous coverage of the many processes between genotype and phenotype including gene expression and epigenetics as well as metabolomics It also examines population genetics and evolutionary models and inference with new chapters on the multi species coalescent admixture and ancient DNA as well as genetic association studies including causal analyses and variant interpretation The Handbook of Statistical Genomics focuses on explaining the main ideas analysis methods and algorithms citing key recent and historic literature for further details and references It also includes a glossary of terms acronyms and abbreviations and features extensive cross referencing between chapters tying the different areas together With heavy use of up to date examples and references to web based resources this continues to be a must have reference in a vital area of research Provides much needed timely coverage of new developments in this expanding area of study Numerous brand new chapters for example covering bacterial genomics microbiome and metagenomics Detailed coverage of application areas with chapters on plant breeding conservation and forensic genetics Extensive coverage of human genetic epidemiology including ethical aspects Edited by one of the leading experts in the field along with rising stars as his co editors Chapter authors are world renowned experts in the field and newly emerging leaders The Handbook of Statistical Genomics is an excellent introductory text for advanced graduate students and early career researchers involved in statistical genetics Statistics in Genetics M.Elizabeth Halloran, Seymour Geisser, 1999-06-04 Though the Genome Project will eventually

result in the sequencing of the human genome as well as the genomes of several other organisms there will still be a need for good statistics for family studies of complex diseases. The papers in this volume are contributions by some of the leading researchers in the field to the current topics in statistical genetics. One section deals with DNA sequence matching and issues related to forensics while another deals with statistical problems of modeling phylogenies and inferential difficulties related to the complex tree structures produced as well as the method of coalescence A Primer of Molecular Population Genetics Asher D. Cutter, 2019-06-06 What are the genomic signatures of adaptations in DNA How often does natural selection dictate changes to DNA How does the ebb and flow in the abundance of individuals over time get marked onto chromosomes to record genetic history Molecular population genetics seeks to answer such questions by explaining genetic variation and molecular evolution from micro evolutionary principles It provides a way to learn about how evolution works and how it shapes species by incorporating molecular details of DNA as the heritable material It enables us to understand the logic of how mutations originate change in abundance in populations and become fixed as DNA sequence divergence between species With the revolutionary advances in genomic data acquisition understanding molecular population genetics is now a fundamental requirement for today s life scientists These concepts apply in analysis of personal genomics genome wide association studies landscape and conservation genetics forensics molecular anthropology and selection scans This book introduces in an accessible way the bare essentials of the theory and practice of molecular population genetics

Topological Data Analysis for Genomics and Evolution Raul Rabadan, Andrew J. Blumberg, 2019-12-19 An introduction to geometric and topological methods to analyze large scale biological data includes statistics and genomic applications Conservation Biology Scott P. Carroll, Charles W. Fox, 2008-09-15 The main goal of this book is to encourage and formalize the infusion of evolutionary thinking into mainstream conservation biology It reviews the evolutionary foundations of conservation issues and unifies conceptual and empirical advances in evolutionary conservation biology The book can be used either as a primary textbook or as a supplementary reading in an advanced undergraduate or graduate level course likely to be called Conservation Biology or in some cases Evolutionary Ecology The focus of chapters is on current concepts in evolution as they pertain to conservation and the empirical study of these concepts The balanced treatment avoids exhaustive reviews and overlapping duplication among the chapters Little background in genetics is assumed of the reader Human Evolutionary Genetics Mark Jobling, Edward Hollox, Toomas Kivisild, Chris Tyler-Smith, 2013-06-25 Human Evolutionary Genetics is a groundbreaking text which for the first time brings together molecular genetics and genomics to the study of the origins and movements of human populations Starting with an overview of molecular genomics for the non specialist which can be a useful review for those with a more genetic background the book shows h Encyclopedia of Evolutionary Biology, 2016-04-14 Encyclopedia of Evolutionary Biology Four Volume Set is the definitive go to reference in the field of evolutionary biology It provides a fully comprehensive review of the field in an easy to

search structure Under the collective leadership of fifteen distinguished section editors it is comprised of articles written by leading experts in the field providing a full review of the current status of each topic The articles are up to date and fully illustrated with in text references that allow readers to easily access primary literature While all entries are authoritative and valuable to those with advanced understanding of evolutionary biology they are also intended to be accessible to both advanced undergraduate and graduate students Broad topics include the history of evolutionary biology population genetics quantitative genetics speciation life history evolution evolution of sex and mating systems evolutionary biogeography evolutionary developmental biology molecular and genome evolution coevolution phylogenetic methods microbial evolution diversification of plants and fungi diversification of animals and applied evolution Presents fully comprehensive content allowing easy access to fundamental information and links to primary research Contains concise articles by leading experts in the field that ensures current coverage of each topic Provides ancillary learning tools like tables illustrations and multimedia features to assist with the comprehension process Handbook of the History and Philosophy of Mathematical Practice Bharath Sriraman, 2024-04-26 The purpose of this unique handbook is to examine the transformation of the philosophy of mathematics from its origins in the history of mathematical practice to the present It aims to synthesize what is known and what has unfolded so far as well as to explore directions in which the study of the philosophy of mathematics as evident in increasingly diverse mathematical practices is headed Each section offers insights into the origins debates methodologies and newer perspectives that characterize the discipline today Contributions are written by scholars from mathematics history and philosophy as well as other disciplines that have contributed to the richness of perspectives abundant in the study of philosophy today who describe various mathematical practices throughout different time periods and contrast them with the development of philosophy Editorial Advisory Board Andrew Aberdein Florida Institute of Technology USA Jody Azzouni Tufts University USA Ot vio Bueno University of Miami USA William Byers Concordia University Canada Carlo Cellucci Sapienza University of Rome Italy Chandler Davis University of Toronto Canada 1926 2022 Paul Ernest University of Exeter UK Michele Friend George Washington University USA Reuben Hersh University of New Mexico USA 1927 2020 Kyeong Hwa Lee Seoul National University South Korea Yuri Manin Max Planck Institute for Mathematics Germany 1937 2023 Athanase Papadopoulos University of Strasbourg France Ulf Persson Chalmers University of Technology Sweden John Stillwell University of San Francisco USA David Tall University of Warwick UK 1941 2024 This book with its exciting depth and breadth illuminates us about the history practice and the very language of our subject about the role of abstraction of proof and manners of proof about the interplay of fundamental intuitions about algebraic thought in contrast to geometric thought The richness of mathematics and the philosophy encompassing it is splendidly exhibited over the wide range of time these volumes cover from deep platonic and neoplatonic influences to the most current experimental approaches Enriched as well with vivid biographies and brilliant personal essays written by and about people who play an important role in our tradition

this extraordinary collection of essays is fittingly dedicated to the memory of Chandler Davis Reuben Hersh and Yuri Manin Barry Mazur Gerhard Gade University Professor Harvard University This encyclopedic Handbook will be a treat for all those interested in the history and philosophy of mathematics Whether one is interested in individuals from Pythagoras through Newton and Leibniz to Grothendieck fields geometry algebra number theory logic probability analysis viewpoints from Platonism to Intuitionism or methods proof experiment computer assistance the reader will find a multitude of chapters that inform and fascinate John Stillwell Emeritus Professor of Mathematics University of San Francisco Recipient of the 2005 Chauvenet Prize Dedicating a volume to the memory of three mathematicians Chandler Davis Reuben Hersh and Yuri Manin who went out of their way to show to a broader audience that mathematics is more than what they might think is an excellent initiative Gathering authors coming from many different backgrounds but who are very strict about the essays they write was successfully achieved by the editor in chief The result a great source of potential inspiration Jean Pierre Bourguignon Nicolaas Kuiper Honorary Professor at the Institut des Hautes tudes Scientifiques A Kinetic View of Statistical Physics Pavel L. Krapivsky, Sidney Redner, Eli Ben-Naim, 2010-11-18 Aimed at graduate students this book explores some of the core phenomena in non equilibrium statistical physics It focuses on the development and application of theoretical methods to help students develop their problem solving skills The book begins with microscopic transport processes diffusion collision driven phenomena and exclusion It then presents the kinetics of aggregation fragmentation and adsorption where the basic phenomenology and solution techniques are emphasized The following chapters cover kinetic spin systems both from a discrete and a continuum perspective the role of disorder in non equilibrium processes hysteresis from the non equilibrium perspective the kinetics of chemical reactions and the properties of complex networks The book contains 200 exercises to test students understanding of the subject A link to a website hosted by the authors containing supplementary material including solutions to some of the exercises can be found at www cambridge org 9780521851039 **Problem** Solving Handbook in Computational Biology and Bioinformatics Lenwood S. Heath, Naren Ramakrishnan, 2010-10-20 Bioinformatics is growing by leaps and bounds theories algorithms statistical techniques are constantly evolving Nevertheless a core body of algorithmic ideas have emerged and researchers are beginning to adopt a problem solving approach to bioinformatics wherein they use solutions to well abstracted problems as building blocks to solve larger scope problems Problem Solving Handbook for Computational Biology and Bioinformatics is an edited volume contributed by world renowned leaders in this field This comprehensive handbook with problem solving emphasis covers all relevant areas of computational biology and bioinformatics Web resources and related themes are highlighted at every opportunity in this central easy to read reference Designed for advanced level students researchers and professors in computer science and bioengineering as a reference or secondary text this handbook is also suitable for professionals working in this industry Branching Processes in Biology Marek Kimmel, David E. Axelrod, 2015-02-17 This book provides a theoretical background

of branching processes and discusses their biological applications Branching processes are a well developed and powerful set of tools in the field of applied probability. The range of applications considered includes molecular biology cellular biology human evolution and medicine The branching processes discussed include Galton Watson Markov Bellman Harris Multitype and General Processes As an aid to understanding specific examples two introductory chapters and two glossaries are included that provide background material in mathematics and in biology The book will be of interest to scientists who work in quantitative modeling of biological systems particularly probabilists mathematical biologists biostatisticians cell biologists molecular biologists and bioinformaticians The authors are a mathematician and cell biologist who have collaborated for more than a decade in the field of branching processes in biology for this new edition. This second expanded edition adds new material published during the last decade with nearly 200 new references More material has been added on infinitely dimensional multitype processes including the infinitely dimensional linear fractional case Hypergeometric function treatment of the special case of the Griffiths Pakes infinite allele branching process has also been added There are additional applications of recent molecular processes and connections with systems biology are explored and a new chapter on genealogies of branching processes and their applications Reviews of First Edition This is a significant book on applications of branching processes in biology and it is highly recommended for those readers who are interested in the application and development of stochastic models particularly those with interests in cellular and molecular biology Siam Review Vol 45 2 2003 This book will be very interesting and useful for mathematicians statisticians and biologists as well and especially for researchers developing mathematical methods in biology medicine and other natural sciences Short Book Reviews of the ISI Vol 23 2 2003 Variation in the Human Genome Derek J. Chadwick, Gail Cardew, 2008-04-30 The mapping of human genes is proceeding rapidly Genes associated with specific inherited diseases are being identified often providing insight into the molecular cause of the disease At the moment however little consideration is being given to the variation present in different human populations Variation in the Human Genome discusses methods of analysing population genetic data and how contemporary genetic heterogeneity arises during the evolution and migration of human populations Specific disorders such as cystic fibrosis beta thalassaemia fragile X phenylketonuria and tumour development susceptibility are used to illustrate this genetic variability and mechanisms of gene mutation and evolution **Dynamical Systems and Differential Equations** Shouchuan Hu,Xin Lu,Weiging Xie,2005 XI Symposium on Probability and Stochastic Processes Ramsés H. Mena, Juan Carlos Pardo, Víctor Rivero, Gerónimo Uribe Bravo, 2015-07-17 This volume features a collection of contributed articles and lecture notes from the XI Symposium on Probability and Stochastic Processes held at CIMAT Mexico in September 2013 Since the symposium was part of the activities organized in Mexico to celebrate the International Year of Statistics the program included topics from the interface between statistics and stochastic processes

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