

RECOMBINATION VARIABILITY AND EVOLUTION

ALGORITHMS
OF ESTIMATION AND
POPULATION-GENETIC
MODELS

A. B. KOROT, I. A. PREYCEL
AND S. I. PREYCEL



CHAPMAN & HALL

Recombination Variability And Evolution Algorithms Of Estimation And Population Genetics

Daniel F McAuley



Recombination Variability And Evolution Algorithms Of Estimation And Population Genetics:

Recombination Variability and Evolution A.B. Korol, S.I. Preigel, 1994-08-31 Using an interdisciplinary approach the authors provide an adaptationist interpretation of the basic features of recombination its evolutionary significance as a key process in reproduction and its importance in genetic mapping The book synthesizes much recent information in the fields of evolutionary genetics of recombination the analysis of genetic markers and breeding applications The authors analyse recombination through a consideration of computer models large Drosophila populations and an empirical approach to current theories Practically orientated readers will be interested in the discussion of a wide spectrum of mapping methods and the new algorithms proposed for genetic mapping of quantitative loci

Genetics of Adaptation Rodney Mauricio, 2005-07-20 An enduring controversy in evolutionary biology is the genetic basis of adaptation Darwin emphasized many slight differences as the ultimate source of variation to be acted upon by natural selection In the early 1900 s this view was opposed by Mendelian geneticists who emphasized the importance of macromutations in evolution The Modern Synthesis resolved this controversy concluding that mutations in genes of very small effect were responsible for adaptive evolution A decade ago Allen Orr and Jerry Coyne reexamined the evidence for this neo Darwinian view and found that both the theoretical and empirical basis for it were weak Orr and Coyne encouraged evolutionary biologists to reexamine this neglected question what is the genetic basis of adaptive evolution In this volume a new generation of biologists have taken up this challenge Using advances in both molecular genetic and statistical techniques evolutionary geneticists have made considerable progress in this emerging field In this volume a diversity of examples from plant and animal studies provides valuable information for those interested in the genetics and evolution of complex traits

Molecular Genetics, Structures, Mechanisms, and Functions Tariq Ahmad Bhat, Abdul Rauf Shakoori, Jameel M. Al-Khayri, 2025-06-30 This new comprehensive two volume set covers all the classical and advanced aspects of molecular genetics and gene manipulation putting this information in one place for beginners experts and those venturing into the fascinating science of molecular biology Volume 2 Applications and Exploring the Nucleus continues the coverage of generic engineering dealing with the concept of genes their relationship with chromosomes and their functional manifestation to the benefit of organisms at large and for human in particular Topics include Mendel s Laws of Inheritance which explains the inheritance of traits visible through generations genome diversity and evolution genetic protein synthesis recombination and evolution of DNA transposable elements in genetics chromosomal aberrations and more The volume also addresses genetic engineering in agricultural science for increased crop yields to reduce costs for food or drug production to reduce the need for pesticides to enhance crop quality etc Volume 1 Principles of Gene Manipulation and Genomics provides an overview of the future of genetic engineering and delves into the role of biotechnology and its applications in genetic engineering It discusses the tools of recombinant technology which have brought about revolution in our understanding of various complex biological

phenomena Chapters cover mutagenesis construction and sequencing of DNA libraries along with applications of genetic engineering for improving health preventing genetic diseases enhancing food resources managing environmental bioremediation and more Topics include genetic engineering tools for restriction enzymes and vectors gene and cell division mutation detection and screening in plants population genetics sexuality in bacteria and more Several chapters focus on the tools of recombinant technology such as restriction enzymes vectors etc that have paved the way for creating organisms of choice and opened new horizons in the field of medicine agriculture and industry for human welfare Providing a wealth of knowledge Molecular Genetics Structures Mechanisms and Functions will be a valuable asset for researchers and scientists working in the field of genetics molecular genetics mutation breeding and plant breeding as well as for faculty and students

Evolutionary Theory and Processes: Modern Perspectives Solomon P. Wasser, 2012-12-06 This volume consists of papers written by evolutionary molecular and organismal biologists geneticists ecologists behavioural ecologists morphologists mathematicians theoreticians and experimentalists in honour of Professor Eviatar Eibi Nevo on the occasion of his seventieth birthday The contributors are only a small subset of Eibi's many friends collaborators and students not that one can distinguish these categories among Eibi's colleagues His widespread influence and activity both in Israel and more generally as a leading evolutionary biologist is indicated by his many co authors on books and papers and by his many students integrated in teaching and research This volume presents some of the most recent dramatic results of molecular genomic and organismal evolutionary processes It represents analyses experiments observations reviews discussions and forecasts of evolutionary theory comprising both novel methods and results reanalyzed and reviewed data sets based on comparative experimental and theoretical studies utilizing model organisms across phylogeny including bacteria fungi plants animals and humans It elucidates the revolution in molecular biology that ushered in our understanding of the evolutionary process over time and space The topics discussed include major problems of evolutionary theory concerning origins phylogeny relative importance of evolutionary forces structure and function adaptation and speciation in space and time in changing and stressful environments A major emerging generalization is the nonrandomness of genome structure highlighting the importance of natural selection as a major organizing evolutionary force not only at the phenotypic level but most importantly at the interlinked genotypic molecular level The integration between the molecular and organismal levels unifies life which is subjected to the mechanism of natural selection as a major orienting evolutionary force

Gene Families:

Studies Of Dna, Rna, Enzymes & Proteins Graeme L Hammond, Roger S Holmes, Hwa A Lim, Zhi Hong Xu, Guo Xiong Xue, Yong Biao Xue, 2001-06-22 This archival volume is an invaluable collection of rigorously reviewed articles by experts in the fields of gene families DNA RNA and proteins to commemorate the passing of a giant of science Professor Clement L Markert 1917 1999 In 1959 Clement Markert and Freddy Moller developed the concept of the isozyme which paved the way for extensive studies of enzyme protein and gene multiplicity across all living organisms This important scientific discovery

has had a profound influence on the biological sciences for more than 40 years and has provided the basis for regular international meetings to discuss the biological and biomedical implications of enzyme multiplicity More recently this concept has been extended to a wide range of gene families of DNA RNA proteins and enzymes

Male Fertility and Infertility T. D. Glover, C. L. R. Barratt, 2009-03-19 This contemporary account of male fertility provides a much needed bridge between those seeking to understand the subject from an evolutionary and biological perspective and those with clinical responsibility for the investigation and treatment of infertility Accordingly the first half of the book deals with the evolutionary aspects of male reproduction and sperm competition sperm production and delivery in man and other animals spermatogenesis and epididymal function sperm transport in the female tract and the apparent decline in human sperm count The second part of the book puts greater emphasis on clinical problems and opens with a discussion of intracytoplasmic sperm injection its value and limitations This is followed by a review of modern developments in the genetics of male fertility and proceeds to a further chapter on the role of surgical procedures used in the treatment Semen analysis is critically reviewed and the molecular techniques now being used in preimplantation diagnosis and in the study of mitochondrial inheritance are fully described Taken together these chapters written by an international team of authors illustrate the breadth of vision needed to tackle the problem of male fertility

Peyote Beatriz Caiuby Labate, Clancy Cavnar, 2016-01-18 This book explains the role that peyote a hallucinogenic cactus plays in the religious and spiritual fulfillment of certain peoples in the United States and Mexico and examines pressing issues concerning the regulation and conservation of peyote as well as issues of indigenous and religious rights Why is mescaline an internationally controlled substance derived from peyote given exemptions for religious use by indigenous groups in Mexico and by the pan indigenous Native American Church in the United States and Canada What are the intersections of peyote use constitutional law and religious freedom And why are natural populations of peyote in decline so much so that in Mexico peyote is considered a species needing special protection This fascinating book addresses these questions and many more It also examines the delicate relationship between the needs of the plant as a species and the needs of man to consume the species for spiritual purposes The authors of this work integrate the history of peyote regulation in the United States and the special trust responsibility relationship between the American Indians and the government into their broad examination of peyote a hallucinogenic cactus containing mescaline that grows naturally in Mexico and southern Texas The book's chapters document how when it comes to peyote multiple stakeholders interests are in conflict as is often the case with issues that involve ethnic identity religion constitutional interpretation and conservation The expansion of peyote traditions also serves as a foundation for examining issues of international human rights law and protections for religious freedom within the global milieu of cultural transnationalism

[Encyclopedia of Biodiversity](#) Simon A. Levin, 2001 Includes articles on agriculture ecology forests wetlands and environment as well as organisms

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

Human Population Genetics and Genomics Alan R. Templeton, 2018-11-08 Human Population Genetics and Genomics provides researchers students with knowledge on population genetics and relevant statistical approaches to help them become more effective users of modern genetic genomic and statistical tools In depth chapters offer thorough discussions of systems of mating genetic drift gene flow and subdivided populations human population history genotype and phenotype detecting selection units and targets of natural selection adaptation to temporally and spatially variable environments selection in age structured populations and genomics and society As human genetics and genomics research often employs tools and approaches derived from population genetics this book helps users understand the basic principles of these tools In addition studies often employ statistical approaches and analysis so an understanding of basic statistical theory is also needed Comprehensively explains the use of population genetics and genomics in medical applications and research Discusses the relevance of population genetics and genomics to major social issues including race and the dangers of modern eugenics proposals Provides an overview of how population genetics and genomics helps us understand where we came from as a species and how we evolved into who we are now

Mycologia Balcanica, 2004
Population Genetics and Microevolutionary Theory Alan R. Templeton, 2006-09-29 The advances made possible by the development of molecular techniques have in recent years revolutionized quantitative genetics and its relevance for population genetics Population Genetics and Microevolutionary Theory takes a modern approach to population genetics incorporating modern molecular biology species level evolutionary biology and a thorough acknowledgment of quantitative genetics as the theoretical basis for population genetics Logically organized into three main sections on population structure and history genotype phenotype interactions and selection adaptation Extensive use of real examples to illustrate concepts Written in a clear and accessible manner and devoid of complex mathematical equations Includes the author's introduction to background material as well as a conclusion for a handy overview of the field and its modern applications Each chapter ends with a set of review questions and answers Offers helpful general references and Internet links

European Journal of Entomology, 2005 **Genetic Architecture and Evolution of Complex Traits and Diseases in Diverse Human Populations** Mashaal Sohail, Jeremy Berg, Diego Ortega-Del Vecchyo, 2022-03-31 Genetics of Populations Philip W. Hedrick, 2005 Genetics and Evolution Encyclopedia of Virology, 2021-02-24 Encyclopedia of Virology Fourth Edition Five

Volume Set builds on the solid foundation laid by the previous editions expanding its reach with new and timely topics In five volumes the work provides comprehensive coverage of the whole virosphere making this a unique resource Content explores viruses present in the environment and the pathogenic viruses of humans animals plants and microorganisms Key areas and concepts concerning virus classification structure epidemiology pathogenesis diagnosis treatment and prevention are discussed guiding the reader through chapters that are presented at an accessible level and include further readings for those needing more specific information More than ever now with the Covid19 pandemic we are seeing the huge impact viruses have on our life and society This encyclopedia is a must have resource for scientists and practitioners and a great source of information for the wider public Offers students and researchers a one stop shop for information on virology not easily available elsewhere Fills a critical gap of information in a field that has seen significant progress in recent years Authored and edited by recognized experts in the field with a range of different expertise thus ensuring a high quality standard

The Phylogenetic Handbook Marco Salemi, Anne-Mieke Vandamme, Philippe Lemey, 2009-03-26 A broad hands on guide with detailed explanations of current methodology relevant exercises and popular software tools

Catalyzing Inquiry at the Interface of Computing and Biology National Research Council, Division on Engineering and Physical Sciences, Computer Science and Telecommunications Board, Committee on Frontiers at the Interface of Computing and Biology, 2006-01-01 Advances in computer science and technology and in biology over the last several years have opened up the possibility for computing to help answer fundamental questions in biology and for biology to help with new approaches to computing Making the most of the research opportunities at the interface of computing and biology requires the active participation of people from both fields While past attempts have been made in this direction circumstances today appear to be much more favorable for progress To help take advantage of these opportunities this study was requested of the NRC by the National Science Foundation the Department of Defense the National Institutes of Health and the Department of Energy The report provides the basis for establishing cross disciplinary collaboration between biology and computing including an analysis of potential impediments and strategies for overcoming them The report also presents a wealth of examples that should encourage students in the biological sciences to look for ways to enable them to be more effective users of computing in their studies

Theories of Population Variation in Genes and Genomes Freddy Bugge Christiansen, 2014-12-17 This textbook provides an authoritative introduction to both classical and coalescent approaches to population genetics Written for graduate students and advanced undergraduates by one of the world's leading authorities in the field the book focuses on the theoretical background of population genetics while emphasizing the close interplay between theory and empiricism Traditional topics such as genetic and phenotypic variation mutation migration and linkage are covered and advanced by contemporary coalescent theory which describes the genealogy of genes in a population ultimately connecting them to a single common ancestor Effects of selection particularly genomic effects are discussed with

reference to molecular genetic variation The book is designed for students of population genetics bioinformatics evolutionary biology molecular evolution and theoretical biology as well as biologists molecular biologists breeders biomathematicians and biostatisticians Contains up to date treatment of key areas in classical and modern theoretical population genetics Provides in depth coverage of coalescent theory Discusses genomic effects of selection Gives examples from empirical population genetics Incorporates figures diagrams and boxed features throughout Includes end of chapter exercises Speaks to a wide range of students in biology bioinformatics and biostatistics Handbook on Analyzing Human Genetic Data Shili Lin, Hongyu Zhao, 2009-10-13 The discipline of statistical genetics is highly computational Be it exact computational methods simulation based or a hybrid of the two computational packages are indispensable tools and constant companions of researchers in the field This handbook is intended to provide human geneticists and other biomedical researchers with guidance on selections of appropriate computational methods and software packages for their specific genetic problems It may also be used by students and other learners as a reference in conjunction with a more theoretical and or methodologically oriented text book This book tries to strike a balance between methodological expositions and practical guidelines for software selections Wherever possible comparisons among the competing methods and software are made to highlight the relative advantages and disadvantages of the approaches so that the reader can make informed choices to best match their specific needs Human genetics has been undergoing an evolution in the past several years as new knowledge and technologies are transforming the field leading to numerous new discoveries of genes associated with complex traits such as cancer obesity and diabetes Many recent genome wide association studies employ the case control design where the study subjects consist of unrelated affected individuals and normal controls For each individual a large number of genetic markers are queried

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