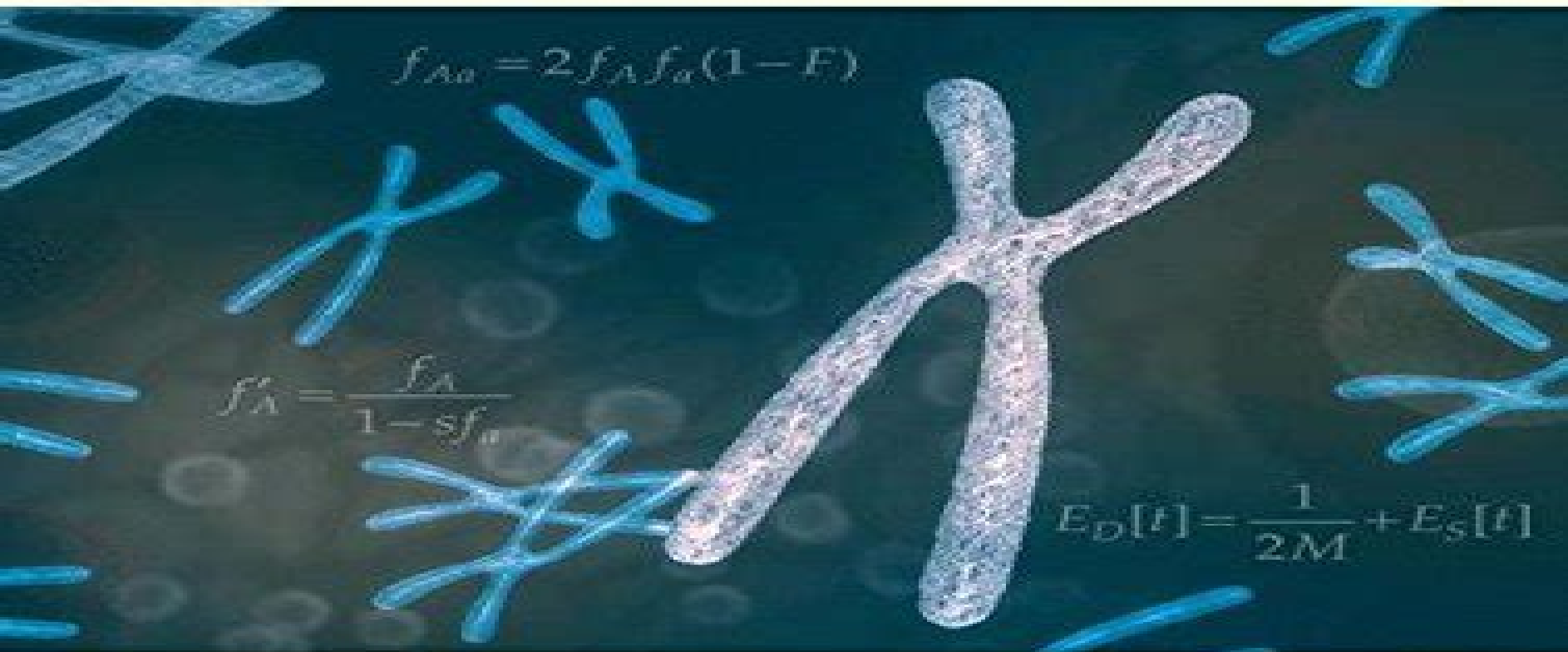


An Introduction to Population Genetics

THEORY AND APPLICATIONS



Rasmus Nielsen • Montgomery Slatkin

An Introduction To Population Genetics Theory And Applications

Om P. Rajora



An Introduction To Population Genetics Theory And Applications:

An Introduction to Population Genetics Rasmus Nielsen, Montgomery Slatkin, 2013-07-18 This book covers both classical population genetics theory developed in terms of allele and haplotype frequencies and modern population genetics theory developed in terms of coalescent theory It features applications of theory to problems that arise in the study of human and other populations and assumes little prior knowledge of mathematics

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Gene Flow Wei Wei, C. Neal Stewart Jr., 2021-11-02 Gene flow is a natural process that occurs spontaneously and enables the evolution of life However with the release of genetically modified organisms concerns have focused on introduced foreign transgenes and their dispersal in nature through gene flow This book examines gene flow of transgenes such as herbicide resistance genes with the goal of understanding the factors that may affect the process of gene flow A greater biological understanding is essential to make sound management regulatory decisions when also taking into consideration the processes that happen in conventional plants Monitoring modelling and mitigation are the three most closely related elements of gene flow The book includes both scientific reviews and perspectives on gene flow and experimental case studies including studies of gene flow in soybean and poplar The authors present diverse views and research methodologies to understand transgene flow

Computational and Evolutionary Analysis of HIV Molecular Sequences Allen G. Rodrigo, Gerald H. Learn Jr., 2007-05-08 Computational and Evolutionary Analysis of HIV Molecular Sequences is for all researchers interested in HIV research even those who only have a nodding acquaintance with computational biology or those who are familiar with some but not all aspects of the field HIV research is unusual in that it brings together scientists from a wide range of disciplines clinicians pathologists immunologists epidemiologists virologists computational biologists structural biologists evolutionary biologists statisticians and mathematicians This book seeks to bridge the gap between these groups in both subject matter and terminology Focused largely on HIV genetic variation Computational and Evolutionary Analysis of HIV Molecular Sequences covers such issues as sampling and processing sequences population genetics phylogenetics and drug targets

Beyond Equilibria Hamish G. Spencer, 2025-06-17 Ecologists and evolutionary biologists especially those with theoretical and mathematical leanings have tended to dismiss historical approaches as descriptive smacking of particularities allowing few generalizations or even being untestable and even antithetical to scientific explanation By contrast the concept of biological equilibrium pervades every aspect of ecology and evolution from the popular notion of the balance of nature to sophisticated mathematical analyses of critical points in the state space of scientific models This novel book explains the problems associated with an over reliance on equilibrium

centred explanations creating a persuasive argument that an explicit consideration of history i.e looking beyond the equilibrium will often lead to a deeper more nuanced understanding of almost every eco evolutionary system The author introduces the concept of flavours of history the variety of ways in which history can play a critical role in biological processes Like the flavours of our food these historical flavours may not always be distinct they may be blended or overlapping and any number may be present However distinguishing between historical flavours is of practical use in explicating ecological and evolutionary phenomena Beyond Equilibria Historical Approaches to Ecology and Evolution is aimed at a broad audience of graduate students and researchers in ecology and evolution Although there is extensive discussion of mathematical models the material is presented in such a way that no detailed knowledge is needed The book will be particularly suitable for a graduate level discussion course

Human Population Genetics John H. Relethford, 2012-03-27 Introductory guide to human population genetics and microevolutionary theory Providing an introduction to mathematical population genetics Human Population Genetics gives basic background on the mechanisms of human microevolution This text combines mathematics biology and anthropology and is best suited for advanced undergraduate and graduate study Thorough and accessible Human Population Genetics presents concepts and methods of population genetics specific to human population study utilizing uncomplicated mathematics like high school algebra and basic concepts of probability to explain theories central to the field By describing changes in the frequency of genetic variants from one generation to the next this book hones in on the mathematical basis of evolutionary theory Human Population Genetics includes Helpful formulae for learning ease Graphs and analogies that make basic points and relate the evolutionary process to mathematical ideas Glossary terms marked in boldface within the book the first time they appear In text citations that act as reference points for further research Exemplary case studies Topics such as Hardy Weinberg equilibrium inbreeding mutation genetic drift natural selection and gene flow Human Population Genetics solidifies knowledge learned in introductory biological anthropology or biology courses and makes it applicable to genetic study NOTE errata for the first edition can be found at the author's website http://employees.oneonta.edu/relethjh/HPG_errata.pdf

Population Biology Simon A. Levin, American Mathematical Society, 1984-12-31 The lecture notes contained in this volume were presented at the AMS Short Course on Population Biology held August 6-7 1983 in Albany New York in conjunction with the summer meeting of the American Mathematical Society These notes will acquaint the reader with the mathematical ideas that pervade almost every level of thinking in population biology and provide an introduction to the many applications of mathematics in the field Research mathematicians college teachers of mathematics and graduate students all should find this book of interest Population biology is probably the oldest area in mathematical biology but remains a constant source of new mathematical problems and the area of biology best integrated with mathematical theory The need for mathematical approaches has never been greater as evolutionary theory is challenged by new interpretations of the paleontological record

and new discoveries at the molecular level as world resources for feeding populations become limiting as the problems of pollution increase and as both animal and plant epidemiological problems receive closer scrutiny A background of advanced calculus introduction to ordinary and partial differential equations and linear algebra will make the book accessible All of the papers included have high research value A list of the contents follows **Einführung in die Mathematik für Biologen** E. Batschelet, 2013-11-11 **Using the Biological Literature** Diane Schmidt, 2014-04-14 The biological sciences cover a broad array of literature types from younger fields like molecular biology with its reliance on recent journal articles genomic databases and protocol manuals to classic fields such as taxonomy with its scattered literature found in monographs and journals from the past three centuries Using the Biological Literature A Practical Guide Fourth Edition is an annotated guide to selected resources in the biological sciences presenting a wide ranging list of important sources This completely revised edition contains numerous new resources and descriptions of all entries including textbooks The guide emphasizes current materials in the English language and includes retrospective references for historical perspective and to provide access to the taxonomic literature It covers both print and electronic resources including monographs journals databases indexes and abstracting tools websites and associations providing users with listings of authoritative informational resources of both classical and recently published works With chapters devoted to each of the main fields in the basic biological sciences this book offers a guide to the best and most up to date resources in biology It is appropriate for anyone interested in searching the biological literature from undergraduate students to faculty researchers and librarians The guide includes a supplementary website dedicated to keeping URLs of electronic and web based resources up to date a popular feature continued from the third edition *Mathematical Properties of Population-Genetic Statistics* Noah A. Rosenberg, 2025-05-20 A pedagogical monograph showing how to use the mathematical properties of population genetic statistics to better interpret genetic data **Parallel Problem Solving from Nature - PPSN III** Yuval Davidor, Hans-Paul Schwefel, Reinhard Männer, 1994-09-21 The challenges in ecosystem science encompass a broadening and strengthening of interdisciplinary ties the transfer of knowledge of the ecosystem across scales and the inclusion of anthropogenic impacts and human behavior into ecosystem landscape and regional models The volume addresses these points within the context of studies in major ecosystem types viewed as the building blocks of central European landscapes The research is evaluated to increase the understanding of the processes in order to unite ecosystem science with resource management The comparison embraces coastal lowland forests associated wetlands and lakes agricultural land use and montane and alpine forests Techniques for upscaling focus on process modelling at stand and landscape scales and the use of remote sensing for landscape level model parameterization and testing The case studies demonstrate ways for ecosystem scientists managers and social scientists to cooperate Mathematische Modelle in der Biologie Wilfried Nöbauer, 2013-07-02 Der Einsatz mathematischer Methoden zur Formulierung und Lösung von Problemen aus dem Bereich der Biowissenschaften auch aus dem Bereich der Medizin hat

in den letzten Jahrzehnten zunehmend an Bedeutung gewonnen Während die mathematische Behandlung einiger Teilgebiete der Biowissenschaften über erste Anfänge noch nicht hinaus gekommen ist gibt es andere Teilgebiete insbesondere die Genetik die Ökologie und die Epidemiologie für die sich bereits eine umfangreiche mathematische Theorie entwickelt hat Dementsprechend gibt es auch schon eine Reihe von Monographien über Teilgebiete der Biomathematik Es gibt aber noch sehr wenige Einführungen in das Gesamtgebiet der Biomathematik das heißt also Bücher die ohne zu sehr ins Detail zu gehen einen Einblick in die bisherigen Ansätze auf diesem Gebiet geben Das vorliegende Buch ist als Beitrag zur Schließung dieser Lücke gedacht Wir wollen in ihm daher die Möglichkeiten der mathematischen Modellbildung im Bereich der Biologie an charakteristischen Beispielen demonstrieren und analysieren beschränken uns dabei aber bewusst auf relativ einfache Modelle um den mathematischen Aufwand in Grenzen zu halten Darüber hinaus werden wir im Anhang des Buches eine Zusammenstellung der verwendeten Mathematik geben soweit sie nicht zum allgemein bekannten Schullehrstoff gehört Wir hoffen dadurch das Buch auch für den Biologen ohne zu große Mühe verständlich wird

Population Dynamics and the Tribolium Model: Genetics and Demography Robert F. Costantino, Robert A. Desharnais, 2012-12-06 The study of populations is becoming increasingly focused on dynamics We believe there are two reasons for this trend The first is the impact of nonlinear dynamics with its exciting ideas and colorful language bifurcations domains of attraction chaos fractals strange attractors Complexity which is so very much a part of biology now seems to be also a part of mathematics A second trend is the accessibility of the new concepts The barriers to communication between theorist and experimentalist seem less impenetrable The active participation of the experimentalist means that the theory will obtain substance Our role is the application of the theory of dynamics to the analysis of biological populations We began our work early in 1979 by writing an ordinary differential equation for the rate of change in adult numbers which was based on an equilibrium model proposed a decade earlier During the next few months we filled our notebooks with straightforward deductions from the model and its associated biological implications Slowly some of the biological observations were explained and papers followed on a variety of topics genetic and demographic stability stationary probability distributions for population size population growth as a birth-death process natural selection and density dependent population growth genetic disequilibrium and the stationary stochastic dynamics of adult numbers

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

An Introduction to Population

Genetics Theory J.F. Crow, 2017-01-01 This text book originally published in 1970 presents the field of population genetics starting with elementary concepts and leading the reader well into the field It is concerned mainly with population genetics in a strict sense and deals primarily with natural populations and less fully with the rather similar problems that arise in breeding live stock and cultivated plants The emphasis is on the behavior of genes and population attributes under natural selection where the most important measure is Darwinian fitness This text is intended for graduate students and advanced undergraduates in genetics and population biology This book steers a middle course between completely verbal biological arguments and the rigor of the mathematician The first two thirds of the book do not require advanced mathematical background An ordinary knowledge of calculus will suffice The latter parts of the book which deal with population stochastically use more advanced methods

Conceptual Breakthroughs in Evolutionary Genetics John C. Avise, 2014-01-18 *Conceptual Breakthroughs in Evolutionary Genetics* is a pithy lively book occupying a special niche the conceptual history of evolutionary genetics not inhabited by any other available treatment Written by a world leading authority in evolutionary genetics this work encapsulates and ranks 70 of the most significant paradigm shifts in evolutionary biology and genetics during the century and a half since Darwin and Mendel The science of evolutionary genetics is central to all of biology but many students and other practitioners have little knowledge of its historical roots and conceptual developments This book fills that knowledge gap in a thought provoking and readable format This fascinating chronological journey along the many conceptual pathways to our modern understanding of evolutionary and genetic principles is a wonderful springboard for discussions in undergraduate or graduate seminars in evolutionary biology and genetics But more than that anyone interested in the history and philosophy of science will find much of value between its covers Provides a relative ranking of 70 seminal breakthroughs and paradigm shifts in the field of evolutionary biology and genetics Modular format permits ready access to each described subject Historical overview of a field whose concepts are central to all of biology and relevant to a broad audience of biologists science historians and philosophers of science Extensively cross referenced with a guide to landmark papers and books for each topic

Sterile Insect Technique Victor A. Dyck, Jorge Hendrichs, A.S.

Robinson, 2021-01-05 The sterile insect technique SIT is an environment friendly method of pest control that integrates well into area wide integrated pest management AW IPM programmes This book takes a generic thematic comprehensive and

global approach in describing the principles and practice of the SIT The strengths and weaknesses and successes and failures of the SIT are evaluated openly and fairly from a scientific perspective The SIT is applicable to some major pests of plant animal and human health importance and criteria are provided to guide in the selection of pests appropriate for the SIT In the second edition all aspects of the SIT have been updated and the content considerably expanded A great variety of subjects is covered from the history of the SIT to improved prospects for its future application The major chapters discuss the principles and technical components of applying sterile insects The four main strategic options in using the SIT suppression containment prevention and eradication with examples of each option are described in detail Other chapters deal with supportive technologies economic environmental and management considerations and the socio economic impact of AW IPM programmes that integrate the SIT In addition this second edition includes six new chapters covering the latest developments in the technology managing pathogens in insect mass rearing using symbionts and modern molecular technologies in support of the SIT applying post factory nutritional hormonal and semiochemical treatments applying the SIT to eradicate outbreaks of invasive pests and using the SIT against mosquito vectors of disease This book will be useful reading for students in animal human and plant health courses The in depth reviews of all aspects of the SIT and its integration into AW IPM programmes complete with extensive lists of scientific references will be of great value to researchers teachers animal human and plant health practitioners and policy makers

Stochastic Models with Applications to Genetics, Cancers, AIDS and Other Biomedical Systems (second Edition) W. Y. Tan, 2015-10-28 This book presents a systematic treatment of Markov chains diffusion processes and state space models as well as alternative approaches to Markov chains through stochastic difference equations and stochastic differential equations It illustrates how these processes and approaches are applied to many problems in genetics carcinogenesis AIDS epidemiology and other biomedical systems One feature of the book is that it describes the basic MCMC Markov chain and Monte Carlo procedures and illustrates how to use the Gibbs sampling method and the multilevel Gibbs sampling method to solve many problems in genetics carcinogenesis AIDS and other biomedical systems As another feature the book develops many state space models for many genetic problems carcinogenesis AIDS epidemiology and HIV pathogenesis It shows in detail how to use the multilevel Gibbs sampling method to estimate or predict simultaneously the state variables and the unknown parameters in cancer chemotherapy carcinogenesis AIDS epidemiology and HIV pathogenesis As a matter of fact this book is the first to develop several state space models for many genetic problems carcinogenesis and other biomedical problems To emphasize special applications to medical problems in this new edition the book has added a new chapter to illustrate how to develop biologically supported stochastic models and state space models of carcinogenesis in human beings Specific examples include hidden Markov models and state space models for human colon cancer human liver cancer and some human pediatric cancers such as retinoblastoma and hepatoblastoma The book also gives examples to illustrate how to develop procedures to assess cancer risk of environmental agents through

initiation promotion protocols *Population Genetics* Matthew B. Hamilton, 2021-02-09 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 *Population Genomics* Om P. Rajora, 2019-01-07
Population genomics has revolutionized various disciplines of biology including population evolutionary ecological and conservation genetics plant and animal breeding human health medicine and pharmacology by allowing to address novel and long standing questions with unprecedented power and accuracy It employs large scale or genome wide genetic information and bioinformatics to address various fundamental and applied aspects in biology and related disciplines and provides a comprehensive genome wide perspective and new insights that were not possible before These advances have become possible due to the development of new and low cost sequencing and genotyping technologies and novel statistical approaches and software bioinformatics tools and models Population genomics is tremendously advancing our understanding the roles of evolutionary processes such as mutation genetic drift gene flow and natural selection in shaping up genetic variation at individual loci and across the genome and populations improving the assessment of population genetic parameters or processes such as adaptive evolution effective population size gene flow admixture inbreeding and outbreeding depression demography and biogeography resolving evolutionary histories and phylogenetic relationships of extant ancient and extinct species understanding the genomic basis of fitness adaptation speciation complex ecological and economically important traits and disease and insect resistance facilitating forensics genetic medicine and pharmacology delineating conservation genetic units and understanding the genetic effects of resource management practices and assisting conservation and sustainable management of genetic resources This Population Genomics book discusses the concepts approaches applications and promises of population genomics in addressing most of the above fundamental and applied crucial aspects in a variety of organisms from microorganisms to humans The book provides insights into a range of emerging population genomics topics including population epigenomics landscape genomics seascape genomics paleogenomics ecological and evolutionary genomics biogeography demography speciation admixture colonization and invasion genomic selection and plant and animal domestication This book fills a vacuum in the field and is expected to become a primary reference in Population Genomics world wide

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