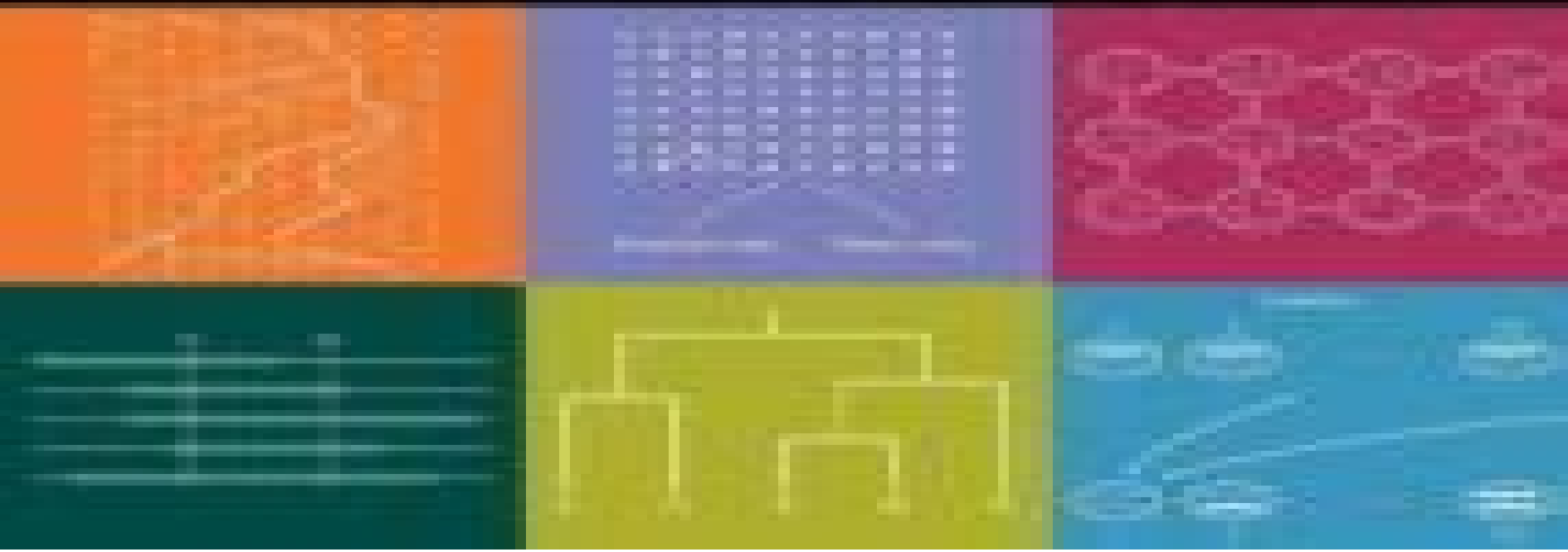


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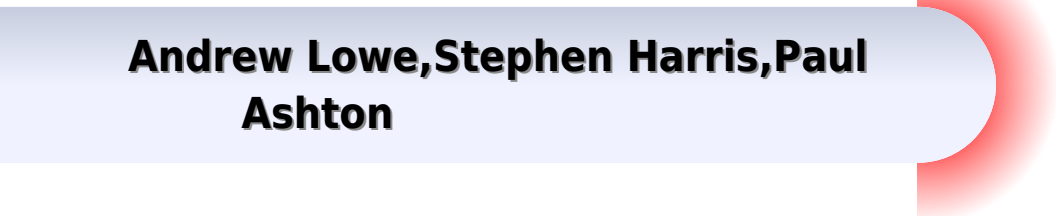
AN APPROACH TO THE EVOLUTION OF GENETIC VARIATION

JOSEPH NEALE
MICHAEL H. SCHNEIDER
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Sequence Variation Genealogies And Evolution

**Andrew Lowe, Stephen Harris, Paul
Ashton**



Sequence Variation Genealogies And Evolution:

Gene Genealogies, Variation and Evolution: A primer in coalescent theory Jotun Hein, Mikkel Schierup, Carsten Wiuf, 2004-12-09 Authored by leading experts this seminal text presents a straightforward and elementary account of coalescent theory which is a central concept in the study of genetic sequence variation observed in a population Rich in examples and illustrations it is ideal for a graduate course in statistics population molecular and medical genetics bioscience and medicine and for students studying the evolution of human population and disease It is also an invaluable reference for bioscientists and statisticians in the pharmaceutical industry and academia Coalescent theory is a central concept in the study of genetic sequence variation that probabilistically describes the genealogy relating the sampled sequences In this text besides fulfilling the glaring need for such a book the authors present this theory in a straightforward and elementary manner and describe the statistical and computational methods used in modelling and analyzing genetic sequence variation Rich in examples and illustrations the book covers basic concepts complications arising from geographical structure and recombination before considering aspects of statistical inference based on these models The book ends with chapters on Gene Mapping which combines sequence variation data with phenotypic data such as disease to define areas of the genome where genes are responsible for the trait and Human Evolution a research area that is experiencing a renaissance due to the enormous amounts of data produced in molecular studies Authored by leading experts this seminal text presents a straightforward and elementary account of coalescent theory which is a central concept in the study of genetic sequence variation observed in a population It is highly suitable for a graduate course in statistics population molecular and medical genetics bioscience and medicine and students studying the evolution of human population and disease and will be an invaluable reference for bioscientists and statisticians in the pharmaceutical industry and academia an excellent and timely book that should appeal to a variety of people in genetics and applied mathematics Professor Montgomery Slatkin Berkeley the authors are outstanding experts in the field and the book is topical and timely Professor David Balding Imperial College Hein Schierup and Wiuf have written the first general book on the coalescent It is an engaging combination of clear mathematical derivation and real data examples Professor Joe Felsenstein University of Washington

Biotic Evolution and Environmental Change in Southeast Asia David Gower, Kenneth Johnson, James Richardson, Brian Rosen, Lukas Rüber, Suzanne Williams, 2012-07-19 The flora and fauna of Southeast Asia are exceptionally diverse The region includes several terrestrial biodiversity hotspots and is the principal global hotspot for marine diversity but it also faces the most intense challenges of the current global biodiversity crisis Providing reviews syntheses and results of the latest research into Southeast Asian earth and organismal history this book investigates the history present and future of the fauna and flora of this bio and geodiverse region Leading authorities in the field explore key topics including palaeogeography palaeoclimatology biogeography population genetics and conservation biology illustrating research approaches and themes

with spatially taxonomically and methodologically focused case studies The volume also presents methodological advances in population genetics and historical biogeography Exploring the fascinating environmental and biotic histories of Southeast Asia this is an ideal resource for graduate students and researchers as well as environmental NGOs **Molecular**

Approaches to Ecology and Evolution R. deSalle, B. Schierwater, 2012-12-06 Four years ago we edited a volume of 36 papers entitled Molecular Approaches to Ecology and Evolution Schierwater et al 1994 in which we attempted to put together a diverse array of papers that demonstrated the impact that the technological revolution of molecular biology has had on the field of evolutionary biology and ecology The present volume borrows from that theme but attempts to focus more sharply on the impact that molecular biology has had on our understanding of different hierarchical levels important in evolutionary and ecological studies Because DNA sequence variation is at the heart of every paper in the present volume we feel it necessary to examine how DNA has affected study at various levels of biological organization The majority of the chapters in the present volume follow themes established in the earlier volume all chapters by authors in the previous volume are either fully updated or entirely new and expand into areas that we felt were important for a more complete understanding of the impact of DNA technology on ecology and evolution The collection of papers in this volume cover a diverse array of ecological and evolutionary questions and demonstrates the breadth of coverage molecular technology has imparted on modern evolutionary biology There are also a broad range of hierarchical questions approached by the 17 papers in this volume Probability Models for DNA Sequence Evolution Rick Durrett, 2013-03-09 Our basic question is

Given a collection of DNA sequences what underlying forces are responsible for the observed patterns of variability To approach this question we introduce and analyze a number of probability models the Wright Fisher model the coalescent the infinite alleles model and the infinite sites model We study the complications that come from nonconstant population size recombination population subdivision and three forms of natural selection directional selection balancing selection and background selection These theoretical results set the stage for the investigation of various statistical tests to detect departures from neutral evolution The final chapter studies the evolution of whole genomes by chromosomal inversions reciprocal translocations and genome duplication Throughout the book the theory is developed in close connection with data from more than 60 experimental studies from the biology literature that illustrate the use of these results This book is written for mathematicians and for biologists alike We assume no previous knowledge of concepts from biology and only a basic knowledge of probability a one semester undergraduate course and some familiarity with Markov chains and Poisson processes Rick Durrett received his Ph D in operations research from Stanford University in 1976 He taught in the UCLA mathematics department before coming to Cornell in 1985 He is the author of six books and 125 research papers and is the academic father of more than 30 Ph D students His current interests are the use of probability models in genetics and ecology and decreasing the mean and variance of his golf **Cumulated Index Medicus**, 1993 **Documenting**

Domestication Melinda A. Zeder, 2006-06-20 A genetic revolution has transformed the study of the domestication of plants and animals Documenting Domestication presents the best research and resolves issues that had been intractable in the past Richard I Ford University of Michigan

Encyclopedia of Ecology Brian D. Fath, 2018-08-23 Encyclopedia of Ecology Second Edition Four Volume Set continues the acclaimed work of the previous edition published in 2008 It covers all scales of biological organization from organisms to populations to communities and ecosystems Laboratory field simulation modelling and theoretical approaches are presented to show how living systems sustain structure and function in space and time New areas of focus include micro and macro scales molecular and genetic ecology and global ecology e g climate change earth transformations ecosystem services and the food water energy nexus are included In addition new international experts in ecology contribute on a variety of topics Offers the most broad ranging and comprehensive resource available in the field of ecology Provides foundational content and suggests further reading Incorporates the expertise of over 500 outstanding investigators in the field of ecology including top young scientists with both research and teaching experience Includes multimedia resources such as an Interactive Map Viewer and links to a CSDMS Community Surface Dynamics Modeling System an open source platform for modelers to share and link models dealing with earth system processes

Molecular Evolution of the Major Histocompatibility Complex Jan Klein, Dagmar Klein, 2013-06-29 From molecules to populations and back In biology the most vigorous organisms often ensue from a union of two disparate pure lines In science too laws of hybrid vigor seem to operate at the interface between two disciplines an interface that often proves to be fertile ground for germinating concepts and new outlooks The fringes of research into the major histocompatibility complex Mhc have provided such an interface several times in the past and the encounters have invigorated fields such as transplantation biology cellular immunology and immunogenetics In the last few years a new interface has been emerging between Mhc and evolutionary genetics and particularly the branch of evolutionary genetics dealing with molecular evolution Mhc research relies upon molecular evolutionary genetics with its grand superstructure of mathematical formulations to come to grips with the events leading to and maintaining the Mhc polymorphism Without the armament of rigorous statistical procedures developed by evolutionary geneticists the intricate relationships among Mhc genes cannot be resolved It will undoubtedly be a molecular geneticist who is the final arbiter in the dispute concerning the nature of the selection pressure molding the Mhc genes And it is doubtful whether the true function of Mhc can ever be comprehended without the vantage point afforded by the elucidation of its evolutionary history

Variation and Evolution in Plants and Microorganisms National Academy of Sciences, 2000-10-11 The present book is intended as a progress report on the synthetic approach to evolution as it applies to the plant kingdom With this simple statement G Ledyard Stebbins formulated the objectives of Variation and Evolution in Plants published in 1950 setting forth for plants what became known as the synthetic theory of evolution or the modern synthesis The pervading conceit of the book was the molding of Darwin s evolution by natural

selection within the framework of rapidly advancing genetic knowledge At the time Variation and Evolution in Plants significantly extended the scope of the science of plants Plants with their unique genetic physiological and evolutionary features had all but been left completely out of the synthesis until that point Fifty years later the National Academy of Sciences convened a colloquium to update the advances made by Stebbins This collection of 17 papers marks the 50th anniversary of the publication of Stebbins classic Organized into five sections the book covers early evolution and the origin of cells virus and bacterial models protocist models population variation and trends and patterns in plant evolution

Ecological Genetics Andrew Lowe, Stephen Harris, Paul Ashton, 2009-04-01 Ecological Genetics addresses the fundamental problems of which of the many molecular markers should be used and how the resulting data should be analysed in clear accessible language suitable for upper level undergraduates through to research level professionals A very accessible straightforward text to deal with this difficult topic applying modern molecular techniques to ecological processes Written by active researchers and teachers within the field There will be an accompanying web site managed by the authors comprising of worked examples test data sets and hyperlinks to relevant web pages

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