

# Genetic Structure and Selection in Subdivided Populations

FRANÇOIS ROUSSET

MONOGRAPHS IN POPULATION BIOLOGY • 40

# Genetic Structure And Selection In Subdivided Populations

**J.F. Crow**



## **Genetic Structure And Selection In Subdivided Populations:**

**Genetic Structure and Selection in Subdivided Populations** François Rousset, 2004 Various approaches have been developed to evaluate the consequences of spatial structure on evolution in subdivided populations This book is both a review and new synthesis of several of these approaches based on the theory of spatial genetic structure François Rousset examines Sewall Wright's methods of analysis based on F statistics effective size and diffusion approximation coalescent arguments William Hamilton's inclusive fitness theory and approaches rooted in game theory and adaptive dynamics Setting these in a framework that reveals their common features he demonstrates how efficient tools developed within one approach can be applied to the others Rousset not only revisits classical models but also presents new analyses of more recent topics such as effective size in metapopulations The book most of which does not require fluency in advanced mathematics includes a self contained exposition of less easily accessible results It is intended for advanced graduate students and researchers in evolutionary ecology and population genetics and will also interest applied mathematicians working in probability theory as well as statisticians

**Genetic Structure and Selection in Subdivided Population** François Rousset, 2004 The Genetic Structure of Populations A. Jacquard, 2012-12-06 It is part of the ideology of science that it is an international enterprise carried out by a community that knows no barriers of nation or culture But the reality is somewhat different Despite the best intentions of scientists to form a single community unseparated by differences of national and political viewpoint they are in fact separated by language Scientific literature in German is not generally assimilated by French workers nor that appearing in French by those whose native language is English The problem appears to have become more severe since the last war because the ascendancy of the United States as the preeminent economic power led in a time of big and expensive science to a pre dominance of American scientific production and a growing tendency at least among English speakers to regard English as the international language of science International congresses and journals of world circulation have come more and more to take English as their standard or official language As a result students and scientific workers in the English speaking world have become more linguistically parochial than ever before and have been cut off from a considerable scientific literature Population genetics has been no exception to the rule The elegant and extremely innovative theoretical work of Malecot for example is only now being properly assimilated by population biologists outside France It was therefore with some sense of frustration that I read Prof *Ecology, Genetics and Evolution of Metapopulations* Ilkka A. Hanski, Oscar E. Gaggiotti, 2004-02-26 Spatial dynamics landscape population *Selection, Management, and Utilization of Biosphere Reserves* Jerry F. Franklin, Stanley L. Krugman, 1979 This publication is directed to the analysis of the selection management and utilization of Biosphere Reserves as viewed by scientists from the United States and the Union of Soviet Socialist Republics Soviet papers focus on types of research and monitoring programs that should be developed on Biosphere Reserves with emphasis on their use in pollutant monitoring The nature of the U S Biosphere

Reserves and their current and potential value for research education and biological preservation is considered in the U S papers An additional paper provides an international perspective on U S and Soviet collaboration in the field of Biosphere Reserves Characteristics of Soviet Biosphere Reserves are described in an appendix Modelling for Field Biologists and Other Interesting People Hanna Kokko,2007-06-28 A friendly 2007 guide to mathematical modelling for evolutionary and behavioural ecologists **Handbook of Statistical Genetics** David J. Balding,Martin Bishop,Chris Cannings,2008-06-10 The Handbook for Statistical Genetics is widely regarded as the reference work in the field However the field has developed considerably over the past three years In particular the modeling of genetic networks has advanced considerably via the evolution of microarray analysis As a consequence the 3rd edition of the handbook contains a much expanded section on Network Modeling including 5 new chapters covering metabolic networks graphical modeling and inference and simulation of pedigrees and genealogies Other chapters new to the 3rd edition include Human Population Genetics Genome wide Association Studies Family based Association Studies Pharmacogenetics Epigenetics Ethic and Insurance As with the second Edition the Handbook includes a glossary of terms acronyms and abbreviations and features extensive cross referencing between the chapters tying the different areas together With heavy use of up to date examples real life case studies and references to web based resources this continues to be must have reference in a vital area of research Edited by the leading international authorities in the field David Balding Department of Epidemiology Public Health Imperial College An advisor for our Probability Statistics series Professor Balding is also a previous Wiley author having written Weight of Evidence for Forensic DNA Profiles as well as having edited the two previous editions of HSG With over 20 years teaching experience he s also had dozens of articles published in numerous international journals Martin Bishop Head of the Bioinformatics Division at the HGMP Resource Centre As well as the first two editions of HSG Dr Bishop has edited a number of introductory books on the application of informatics to molecular biology and genetics He is the Associate Editor of the journal Bioinformatics and Managing Editor of Briefings in Bioinformatics Chris Cannings Division of Genomic Medicine University of Sheffield With over 40 years teaching in the area Professor Cannings has published over 100 papers and is on the editorial board of many related journals Co editor of the two previous editions of HSG he also authored a book on this topic **Introduction to Conservation Genetics** Richard Frankham,Jonathan D. Ballou,David Anthony Briscoe,2010 This impressive author team brings the wealth of advances in conservation genetics into the new edition of this introductory text including new chapters on population genomics and genetic issues in introduced and invasive species They continue the strong learning features for students main points in the margin chapter summaries vital support with the mathematics and further reading and now guide the reader to software and databases Many new references reflect the expansion of this field With examples from mammals birds reptiles fish amphibians plants and invertebrates this is an ideal introduction to conservation genetics for a broad audience The text tackles the quantitative aspects of conservation genetics and has a host of pedagogy to support students

learning the numerical side of the subject Combined with being up to date its user friendly writing style and first class illustration programme forms a robust teaching package      **Fast Variables in Stochastic Population Dynamics** George William Albert Constable, 2015-07-25 In this thesis two variants of the fast variable elimination method are developed They are intuitive simple to implement and give results which are in very good agreement with those found from numerical simulations The relative simplicity of the techniques makes them ideal for applying to problems featuring demographic stochasticity for experts and non experts alike Within the context of mathematical modelling fast variable elimination is one of the central tools with which one can simplify a multivariate problem When used in the context of deterministic systems the theory is quite standard but when stochastic effects are present it becomes less straightforward to apply While the introductory and background chapters form an excellent primer to the theory of stochastic population dynamics the techniques developed can be applied to systems exhibiting a separation of timescales in a variety of fields including population genetics ecology and epidemiology      Plotting landscape perspectives of clearcut units Roger H. Twito, 1978

**Landscape Genetics** Niko Balkenhol, Samuel Cushman, Andrew Storfer, Lisette Waits, 2015-11-09 LANDSCAPE GENETICS CONCEPTS METHODS APPLICATIONS LANDSCAPE GENETICS CONCEPTS METHODS APPLICATIONS Edited by Niko Balkenhol Samuel A Cushman Andrew T Storfer Lisette P Waits Landscape genetics is an exciting and rapidly growing field melding methods and theory from landscape ecology and population genetics to address some of the most challenging and urgent ecological and evolutionary topics of our time Landscape genetic approaches now enable researchers to study in detail how environmental complexity in space and time affect gene flow genetic drift and local adaptation However learning about the concepts and methods underlying the field remains challenging due to the highly interdisciplinary nature of the field which relies on topics that have traditionally been treated separately in classes and textbooks In this edited volume some of the leading experts in landscape genetics provide the first comprehensive introduction to underlying concepts commonly used methods and current and future applications of landscape genetics Consistent with the interdisciplinary nature of the field the book includes textbook like chapters that synthesize fundamental concepts and methods underlying landscape genetics Part 1 chapters on advanced topics that deserve a more in depth treatment Part 2 and chapters illustrating the use of concepts and methods in empirical applications Part 3 Aimed at beginning landscape geneticists and experienced researchers alike this book will be helpful for all scientists and practitioners interested in learning teaching and applying landscape genetics      **Introduction to Theoretical Population Genetics**

Thomas Nagylaki, 2013-03-12 This book covers those areas of theoretical population genetics that can be investigated rigorously by elementary mathematical methods I have tried to formulate the various models fairly generally and to state the biological assumptions quite explicitly I hope the choice and treatment of topics will enable the reader to understand and evaluate detailed analyses of many specific models and applications in the literature Models in population genetics are highly

idealized often even over idealized and their connection with observation is frequently remote Further more it is not practicable to measure the parameters and variables in these models with high accuracy These regrettable circumstances amply justify the use of appropriate lucid and rigorous approximations in the analysis of our models and such approximations are often illuminating even when exact solutions are available However our empirical and theoretical limitations justify neither opaque incomplete formulations nor unconvincing inadequate analyses for these may produce uninterpretable misleading or erroneous results Intuition is a principal source of ideas for the construction and investigation of models but it can replace neither clear formulation nor careful analysis Fisher 1930 1958 pp x 23 24 38 not only espoused similar ideas but he recognized also that our concepts of intuition and rigor must evolve in time The book is neither a review of the literature nor a compendium of results The material is almost entirely self contained The first eight chapters are a thoroughly revised and greatly extended version of my published lecture notes Nagylaki 1977a      **General Technical Report PNW. ,1978**

**Games, Groups, and the Global Good** Simon A. Levin, 2009-06-09 How do groups form how do institutions come into being and when do moral norms and practices emerge This volume explores how game theoretic approaches can be extended to consider broader questions that cross scales of organization from individuals to cooperatives to societies Game theory strategic formulation of central problems in the analysis of social interactions is used to develop multi level theories that examine the interplay between individuals and the collectives they form The concept of cooperation is examined at a higher level than that usually addressed by game theory especially focusing on the formation of groups and the role of social norms in maintaining their integrity with positive and negative implications The authors suggest that conventional analyses need to be broadened to explain how heuristics like concepts of fairness arise and become formalized into the ethical principles embraced by a society      *Social Evolution and Inclusive Fitness Theory* James A.R. Marshall, 2019-01-08 Social behavior has long puzzled evolutionary biologists since the classical theory of natural selection maintains that individuals should not sacrifice their own fitness to affect that of others Social Evolution and Inclusive Fitness Theory argues that a theory first presented in 1963 by William D Hamilton inclusive fitness theory provides the most fundamental and general explanation for the evolution and maintenance of social behavior in the natural world James Marshall guides readers through the vast and confusing literature on the evolution of social behavior introducing and explaining the competing theories that claim to provide answers to questions such as why animals evolve to behave altruistically Using simple statistical language and techniques that practicing biologists will be familiar with he provides a comprehensive yet easily understandable treatment of key concepts and their repeated misinterpretations Particular attention is paid to how more realistic features of behavior such as nonadditivity and conditionality can complicate analysis Marshall highlights the general problem of identifying the underlying causes of evolutionary change and proposes fruitful approaches to doing so in the study of social evolution Social Evolution and Inclusive Fitness Theory describes how inclusive fitness theory addresses both simple and complex social

scenarios the controversies surrounding the theory and how experimental work supports the theory as the most powerful explanation for social behavior and its evolution      **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 cul t i vat ed plans 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      **Eco-Evolutionary Dynamics** Andrew P. Hendry, 2016-11-22 In recent years

scientists have realized that evolution can occur on timescales much shorter than the long lapse of ages emphasized by Darwin in fact evolutionary change is occurring all around us all the time This book provides an authoritative and accessible introduction to eco evolutionary dynamics a cutting edge new field that seeks to unify evolution and ecology into a common conceptual framework focusing on rapid and dynamic environmental and evolutionary change Andrew Hendry covers key aspects of evolution ecology and their interactions Topics range from natural selection adaptive divergence ecological speciation and gene flow to population and community dynamics ecosystem function plasticity and genomics Hendry evaluates conceptual and methodological approaches and draws on empirical data from natural populations including those in human disturbed environments to tackle a number of classic and emerging research questions He also discusses exciting new directions for future research at the intersection of ecology and evolution An invaluable guide for students and researchers alike Eco evolutionary Dynamics reveals how evolution and ecology interact strongly on short timescales to shape the world we see around us      Genetic Variability in Conservation and Selection Programs in the Post-Genomics Era

Maria Saura, Francesca Bertolini, Christos Palaiokostas, 2020-12-22 This eBook is a collection of articles from a Frontiers Research Topic Frontiers Research Topics are very popular trademarks of the Frontiers Journals Series they are collections of at least ten articles all centered on a particular subject With their unique mix of varied contributions from Original Research to Review Articles Frontiers Research Topics unify the most influential researchers the latest key findings and historical advances in a hot research area Find out more on how to host your own Frontiers Research Topic or contribute to one as an author by contacting the Frontiers Editorial Office frontiersin.org about contact      Selection Graham Bell, 2008

This book adopts a direct experimental approach to evolutionary questions drawing predominantly from research on microbial systems The focus is on processes and mechanisms and incorporates insights from recent advances in whole

genome sequencing bioinformatics environmental genomics and developmental genetics      *Population Genetics in Forestry*  
Hans-Rolf Gregorius, 2013-03-13 When we consider the main object of forestry the tree it immediately becomes clear why experimental population geneticists have been so hesitant in making this object a primary concern of their research Trees are very long living organisms with generation intervals frequently exceeding those of their investigators by multiples They virtually exclude therefore application of the classical methods of population genetics since these are based on observing genetic structures over generations This situation where the limits set to observation are so severe particularly requires close cooperation between theory and experiment It also requires careful consideration of results obtained for organisms other than trees in order to gain additional insights by comparing the results for trees with those for other organisms Yet the greatest challenge to population and ecological genetics probably originates from the fact that forests are very likely to be the most complex ecosystems of all even in some cases where they are subject to intense management This complexity which equally comprises biotic and abiotic factors varying both in time and space makes extremely high demands on the adaptational capacity and thus flexibility of the carriers of such an ecosystem Longevity combined with immobility during the vegetative phase however appears to contradict the obvious necessity of adaptational flexibility in forest tree populations when compared with short lived and or mobile organisms



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