

Population Genetics Concise Guide 2nd

John H. Gillespie

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Population Genetics John H. Gillespie, 2004-08-06 This concise introduction offers students and researchers an overview of the discipline that connects genetics and evolution Addressing the theories behind population genetics and relevant empirical evidence John Gillespie discusses genetic drift natural selection nonrandom mating quantitative genetics and the evolutionary advantage of sex First published to wide acclaim in 1998 this brilliant primer has been updated to include new sections on molecular evolution genetic drift genetic load the stationary distribution and two locus dynamics This book is indispensable for students working in a laboratory setting or studying free ranging populations Genetics with R Áki Jarl Láruson, Floyd A. Reed, 2021 This book focuses on applying the hugely popular R software specifically to the field offering an accessible step by step guide to tackling the challenges of achieving effective data interpretation and Understanding Population Genetics Torbjörn Säll, Bengt O. Bengtsson, 2017-07-14 An inspiring introduction to a vital scientific field The reader is taken through ten mathematical derivations that lead to important results explaining in a hands on manner the key concepts and methods of theoretical population genetics. The derivations are carefully worked out and easy to follow Particular attention is given to the underlying assumptions and the mathematics used The results are discussed and broadened out with relevant current implications All topics feature questions with helpful answers The book is intended for the reader who already knows some population genetics but requires a more comprehensive understanding It is particularly suited to those who analyse genetic data and wish to better grasp what their results actually mean It will also be helpful for those who wish to understand how population genetics contributes to the explanation of evolution Or as the writers claim If one wants to understand life in all its improbable and amazing richness one must start by understanding population genetics Strickberger's Evolution Brian Hall, Benedikt Hallgrímsson, 2008 Thoroughly updated and reorganized Strickberger's Evolution Fourth Edition presents biology students with a basic introduction to prevailing knowledge and ideas about evolution discussing how why and where the world and its organisms changed throughout history Keeping consistent with Strickberger's engaging writing style the authors carefully unfold a broad range of philosophical and historical topics that frame the theories of today including cosmological and geological evolution and its impact on life the origins of life on earth the development of molecular pathways from genetic systems to organismic morphology and function the evolutionary history of organisms from microbes to animals and the numerous molecular and populational concepts that explain the earth s dynamic evolution Population Genetics John H. Gillespie, 2004-08-06 This concise introduction offers students and researchers an overview of the discipline that connects genetics and evolution Addressing the theories behind population genetics and relevant empirical evidence John Gillespie discusses genetic drift natural selection nonrandom mating quantitative genetics and the evolutionary advantage of sex First published to wide acclaim in 1998 this brilliant primer has been updated to include new sections on molecular evolution genetic drift genetic load the stationary

distribution and two locus dynamics This book is indispensable for students working in a laboratory setting or studying free ranging populations 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 Mathematical Properties of Population-Genetic Statistics Noah A. Reviews of the ISI Vol 23 2 2003 Rosenberg, 2025-05-20 A powerful new approach to interpreting population genetic data in evolution and ecology Population genetics uses statistical analysis to catalog genetic variation among populations and species Summary statistics computed from allele frequencies mathematical functions that measure features of genetic similarity and diversity are key to this global effort Yet despite their widespread use in evolutionary biology ecology and conservation biology their mathematical properties have largely been overlooked This book shows how to use the mathematical bounds on summary statistics to make better interpretations of population genetic data Noah Rosenberg discusses how the behavior of these statistics depends not only on the biology of the populations they seek to describe but also on the mathematical properties of the functions used to compute them properties that produce constraints on the values of the statistics and influence their interpretation Focusing on the concept of homozygosity a quadratic function of allele frequencies in a population he demonstrates how to account for mathematical constraints when measuring genetic similarity and diversity Rosenberg illustrates the results using examples

from empirical data and shares strategies that readers can use to apply this mathematical perspective to different kinds of summary statistics including those for measuring biodiversity in ecological communities Drawing inspiration from Charles Darwin who marveled at evolution s endless forms most beautiful and most wonderful this book presents a groundbreaking approach to the study of genetic variation It is sure to stimulate new research in population biology and deeper thinking about the meaning and interpretation of essential measurements of the world's genomes Phylogenomic Data **Acquisition** W. Bryan Jennings, 2016-12-12 Phylogenomics is a rapidly growing field of study concerned with using genome wide data usually in the form of DNA sequence loci to infer the evolution of genes genomes and the Tree of Life Accordingly this discipline connects many areas in biology including molecular and genomic evolution systems biology molecular systematics phylogeography conservation genetics DNA barcoding and others With the advent of Next Generation Sequencing in addition to advances in computer hardware and software over the past decade researchers can now generate unparalleled phylogenomic datasets that are helping to illuminate many areas in the life sciences This book is an introduction to the principles and practices of gathering these data Phylogenomic Data Acquisition Principles and Practice is intended for a broad cross section of biologists and anyone else interested in learning how to obtain phylogenomic data using the latest Encyclopedia of Biodiversity, 2013-02-05 The 7 volume Encyclopedia of Biodiversity Second Edition maintains methods the reputation of the highly regarded original presenting the most current information available in this globally crucial area of research and study It brings together the dimensions of biodiversity and examines both the services it provides and the measures to protect it Major themes of the work include the evolution of biodiversity systems for classifying and defining biodiversity ecological patterns and theories of biodiversity and an assessment of contemporary patterns and trends in biodiversity The science of biodiversity has become the science of our future It is an interdisciplinary field spanning areas of both physical and life sciences Our awareness of the loss of biodiversity has brought a long overdue appreciation of the magnitude of this loss and a determination to develop the tools to protect our future Second edition includes over 100 new articles and 226 updated articles covering this multidisciplinary field from evolution to habits to economics in 7 volumes The editors of this edition are all well respected instantly recognizable academics operating at the top of their respective fields in biodiversity research readers can be assured that they are reading material that has been meticulously checked and reviewed by experts Approximately 1 800 figures and 350 tables complement the text and more than 3 000 glossary entries explain key terms Complexity and Dynamics ,2017

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