MARC MANGEL

The Theoretical **Biologist's** Toolbox

Quantitative **Methods for Ecology and Evolutionary** Biology

Sarah P. Otto, Troy Day

The Theoretical Biologist's Toolbox Marc Mangel,2006-07-27 Mathematical modelling is widely used in ecology and evolutionary biology and it is a topic that many biologists find difficult to grasp In this new textbook Marc Mangel provides a no nonsense introduction to the skills needed to understand the principles of theoretical and mathematical biology Fundamental theories and applications are introduced using numerous examples from current biological research complete with illustrations to highlight key points Exercises are also included throughout the text to show how theory can be applied and to test knowledge gained so far Suitable for advanced undergraduate courses in theoretical and mathematical biology this book forms an essential resource for anyone wanting to gain an understanding of theoretical ecology and evolution

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 Bulletin ,2006 A Biologist's Guide to Mathematical Modeling in Ecology and Evolution Sarah P. Otto, Troy Day, 2007-03-12 Thirty years ago biologists could get by with a rudimentary grasp of mathematics and modeling Not so today In seeking to answer fundamental questions about how biological systems function and change over time the modern biologist is as likely to rely on sophisticated mathematical and computer based models as traditional fieldwork In this book Sarah Otto and Troy Day provide biology students with the tools necessary to both interpret models and to build their own The book starts at an elementary level of mathematical modeling assuming that the reader has had high school mathematics and first year calculus Otto and Day then gradually build in depth and complexity from classic models in ecology and evolution to more intricate class structured and probabilistic models The authors provide primers with instructive exercises to introduce readers to the more advanced subjects of linear algebra and probability theory Through examples they describe how models have been used to understand such topics as the spread of HIV chaos the age structure of a country speciation and extinction Ecologists and evolutionary biologists today need enough mathematical training to be able to assess the power and limits of biological models and to develop theories and models themselves This innovative book will be an indispensable guide to the world of mathematical models for the next generation of biologists A how to guide for developing new mathematical models in biology Provides step by step recipes for constructing and analyzing models Interesting biological applications Explores classical models in ecology and evolution Questions at the end of every chapter Primers cover important mathematical topics Exercises with answers Appendixes summarize useful rules Labs and advanced material available Quantitative Ecology and Evolutionary Biology Otso Ovaskainen, Henrik Johan de Knegt, Maria del Mar Delgado, 2016-09-01 This novel interdisciplinary text achieves an integration of empirical data and theory with the aid of mathematical models and statistical methods The emphasis throughout is on spatial ecology and evolution especially on the interplay between environmental heterogeneity and biological processes. The book provides a coherent theme by interlinking the modelling approaches used for different

subfields of spatial ecology movement ecology population ecology community ecology and genetics and evolutionary ecology each being represented by a separate chapter Each chapter starts by describing the concept of each modelling approach in its biological context goes on to present the relevant mathematical models and statistical methods and ends with a discussion of the benefits and limitations of each approach The concepts and techniques discussed throughout the book are illustrated throughout with the help of empirical examples This is an advanced text suitable for any biologist interested in the integration of empirical data and theory in spatial ecology evolution through the use of quantitative statistical methods and mathematical models. The book will also be of relevance and use as a textbook for graduate level courses in spatial ecology ecological modelling theoretical ecology and statistical ecology An Introduction to Methods and Models in Ecology, Evolution, and Conservation Biology Stanton Braude, Bobbi S. Low, 2010-01-04 An innovative introduction to ecology and evolution This unique textbook introduces undergraduate students to quantitative models and methods in ecology behavioral ecology evolutionary biology and conservation It explores the core concepts shared by these related fields using tools and practical skills such as experimental design generating phylogenies basic statistical inference and persuasive grant writing And contributors use examples from their own cutting edge research providing diverse views to engage students and broaden their understanding This is the only textbook on the subject featuring a collaborative active learning approach that emphasizes hands on learning Every chapter has exercises that enable students to work directly with the material at their own pace and in small groups Each problem includes data presented in a rich array of formats which students use to answer questions that illustrate patterns principles and methods Topics range from Hardy Weinberg equilibrium and population effective size to optimal foraging and indices of biodiversity The book also includes a comprehensive glossary In addition to the editors the contributors are James Beck Cawas Behram Engineer John Gaskin Luke Harmon Jon Hess Jason Kolbe Kenneth H Kozak Robert J Robertson Emily Silverman Beth Sparks Jackson and Anton Weisstein Provides experience with hypothesis testing experimental design and scientific reasoning Covers core quantitative models and methods in ecology behavioral ecology evolutionary biology and conservation Turns discussion sections into thinking labs Professors A supplementary Instructor's Manual is available for this book It is restricted to teachers using the text in courses For information on how to obtain a copy refer to http press princeton edu class use solutions html Handbook of Quantitative Ecology Justin Kitzes, 2022-08-16 An essential guide to quantitative research methods in ecology and conservation biology accessible for even the most math averse student or professional Quantitative research techniques have become increasingly important in ecology and conservation biology but the sheer breadth of methods that must be understood from population modeling and probabilistic thinking to modern statistics simulation and data science and a lack of computational or mathematics training have hindered quantitative literacy in these fields In this book ecologist Justin Kitzes addresses those challenges for students and practicing scientists alike Requiring only basic algebra and the ability to use a spreadsheet

Handbook of Quantitative Ecology is designed to provide a practical intuitive and integrated introduction to widely used quantitative methods Kitzes builds each chapter around a specific ecological problem and arrives step by step at a general principle through the process of solving that problem Grouped into five broad categories difference equations probability matrix models likelihood statistics and other numerical methods the book introduces basic concepts starting with exponential and logistic growth and helps readers to understand the field s more advanced subjects such as bootstrapping stochastic optimization and cellular automata Complete with online solutions to all numerical problems Kitzes s Handbook of Quantitative Ecology is an ideal coursebook for both undergraduate and graduate students of ecology as well as a useful and necessary resource for mathematically out of practice scientists Statistical Methods in Molecular Evolution Rasmus Nielsen, 2005-04-21 In the field of molecular evolution inferences about past evolutionary events are made using molecular data from currently living species With the availability of genomic data from multiple related species molecular evolution has become one of the most active and fastest growing fields of study in genomics and bioinformatics Most studies in molecular evolution rely heavily on statistical procedures based on stochastic process modelling and advanced computational methods including high dimensional numerical optimization and Markov Chain Monte Carlo This book provides an overview of the statistical theory and methods used in studies of molecular evolution It includes an introductory section suitable for readers that are new to the field a section discussing practical methods for data analysis and more specialized sections discussing specific models and addressing statistical issues relating to estimation and model choice The chapters are written by the leaders of field and they will take the reader from basic introductory material to the state of the art statistical methods This book is suitable for statisticians seeking to learn more about applications in molecular evolution and molecular evolutionary biologists with an interest in learning more about the theory behind the statistical methods applied in the field The chapters of the book assume no advanced mathematical skills beyond basic calculus although familiarity with basic probability theory will help the reader Most relevant statistical concepts are introduced in the book in the context of their application in molecular evolution and the book should be accessible for most biology graduate students with an interest in quantitative methods and theory Rasmus Nielsen received his Ph D form the University of California at Berkeley in 1998 and after a postdoc at Harvard University he assumed a faculty position in Statistical Genomics at Cornell University He is currently an Ole R mer Fellow at the University of Copenhagen and holds a Sloan Research Fellowship His is an associate editor of the Journal of Molecular Evolution and has published more than fifty original papers in peer reviewed journals on the topic of this book From the reviews Overall this is a very useful book in an area of increasing importance Journal of the Royal Statistical Society I find Statistical Methods in Molecular Evolution very interesting and useful It delves into problems that were considered very difficult just several years ago the book is likely to stimulate the interest of statisticians that are unaware of this exciting field of applications It is my hope that it will also help the wet lab molecular evolutionist to better understand

mathematical and statistical methods Marek Kimmel for the Journal of the American Statistical Association September 2006 Who should read this book We suggest that anyone who deals with molecular data who does not and anyone who asks evolutionary questions who should not ought to consult the relevant chapters in this book Dan Graur and Dror Berel for Biometrics September 2006 Coalescence theory facilitates the merger of population genetics theory with phylogenetic approaches but still there are mostly two camps phylogeneticists and population geneticists Only a few people are moving freely between them Rasmus Nielsen is certainly one of these researchers and his work so far has merged many population genetic and phylogenetic aspects of biological research under the umbrella of molecular evolution Although Nielsen did not contribute a chapter to his book his work permeates all its chapters This book gives an overview of his interests and current achievements in molecular evolution In short this book should be on your bookshelf Peter Beerli for Evolution 60 2 2006

The Comparative Method in Evolutionary Biology Paul H. Harvey, Mark D. Pagel, 1991 From Darwin onward it has been second nature for evolutionary biologists to think comparatively because comparisons establish the generality of evolutionary phenomena Do large genomes slow down development What lifestyles select for large brains Are extinction rates related to body size These are all questions for the comparative method and this book is about how such questions can be answered The first chapter elaborates on suitable questions for the comparative approach and shows how it complements other approaches to problem solving in evolution The second chapter identifies the biological causes of similarity among closely related species for almost any observed character The third chapter discusses methods for reconstructing phylogenetic trees and ancestral character states The fourth chapter sets out to develop statistical tests that will determine whether different characters that exist in discrete states show evidence for correlated evolution Chapter 5 turns to comparative analyses of continuously varying characters Chapter 6 looks at allometry to exemplify the themes and methods discussed earlier while the last chapter looks to future development of the comparative approach in both molecular and organismic biology The Quantitative Method in Biology Julius MacLeod, 1919

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