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Mathematical Epidemiology of Infectious Diseases

Model Building, Analysis
and Interpretation

O. Diekmann, J. A. P. Heesterbeek

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Mathematical Epidemiology Of Infectious Diseases

Model Building Analysis And Interpretation

Guillaume Favre

Mathematical Epidemiology Of Infectious Diseases Model Building Analysis And Interpretation:

Mathematical Epidemiology of Infectious Diseases O. Diekmann, J. A. P. Heesterbeek, 2000-04-07 Mathematical Epidemiology of Infectious Diseases Model Building Analysis and Interpretation O Diekmann University of Utrecht The Netherlands J A P Heesterbeek Centre for Biometry Wageningen The Netherlands The mathematical modelling of epidemics in populations is a vast and important area of study It is about translating biological assumptions into mathematics about mathematical analysis aided by interpretation and about obtaining insight into epidemic phenomena when translating mathematical results back into population biology Model assumptions are formulated in terms of usually stochastic behaviour of individuals and then the resulting phenomena at the population level are unravelled Conceptual clarity is attained assumptions are stated clearly hidden working hypotheses are attained and mechanistic links between different observables are exposed Features Model construction analysis and interpretation receive detailed attention Uniquely covers both deterministic and stochastic viewpoints Examples of applications given throughout Extensive coverage of the latest research into the mathematical modelling of epidemics of infectious diseases Provides a solid foundation of modelling skills The reader will learn to translate model analyse and interpret with the help of the numerous exercises In literally working through this text the reader acquires modelling skills that are also valuable outside of epidemiology certainly within population dynamics but even beyond that In addition the reader receives training in mathematical argumentation The text is aimed at applied mathematicians with an interest in population biology and epidemiology at theoretical biologists and epidemiologists Previous exposure to epidemic concepts is not required as all background information is given The book is primarily aimed at self study and ideally suited for small discussion groups or for use as a course text

Mathematical Epidemiology of Infectious Diseases O. Diekmann, J. A. P. Heesterbeek, 2000-04-07 Mathematical Epidemiology of Infectious Diseases Model Building Analysis and Interpretation O Diekmann University of Utrecht The Netherlands J A P Heesterbeek Centre for Biometry Wageningen The Netherlands The mathematical modelling of epidemics in populations is a vast and important area of study It is about translating biological assumptions into mathematics about mathematical analysis aided by interpretation and about obtaining insight into epidemic phenomena when translating mathematical results back into population biology Model assumptions are formulated in terms of usually stochastic behaviour of individuals and then the resulting phenomena at the population level are unravelled Conceptual clarity is attained assumptions are stated clearly hidden working hypotheses are attained and mechanistic links between different observables are exposed Features Model construction analysis and interpretation receive detailed attention Uniquely covers both deterministic and stochastic viewpoints Examples of applications given throughout Extensive coverage of the latest research into the mathematical modelling of epidemics of infectious diseases Provides a solid foundation of modelling skills The reader will learn to translate model analyse and interpret with the help of the numerous exercises In literally working through this text the reader acquires modelling skills

that are also valuable outside of epidemiology certainly within population dynamics but even beyond that In addition the reader receives training in mathematical argumentation The text is aimed at applied mathematicians with an interest in population biology and epidemiology at theoretical biologists and epidemiologists Previous exposure to epidemic concepts is not required as all background information is given The book is primarily aimed at self study and ideally suited for small discussion groups or for use as a course text

Mathematical Epidemiology of Infectious Diseases O. Diekmann,2000

Provides systematic coverage of the mathematical theory of modelling epidemics in populations with a clear and coherent discussion of the issues concepts and phenomena Mathematical modelling of epidemics is a vast and important area of study and this book helps the reader to translate model analyse and interpret with numerous applications examples and exercises to aid understanding Publisher description

An Introduction to Mathematical Epidemiology Maia Martcheva,2015-10-20

The book is a comprehensive self contained introduction to the mathematical modeling and analysis of infectious diseases It includes model building fitting to data local and global analysis techniques Various types of deterministic dynamical models are considered ordinary differential equation models delay differential equation models difference equation models age structured PDE models and diffusion models It includes various techniques for the computation of the basic reproduction number as well as approaches to the epidemiological interpretation of the reproduction number MATLAB code is included to facilitate the data fitting and the simulation with age structured models

[A Primer on Population Dynamics Modeling](#)

Hiromi Seno,2022-11-16 This textbook provides an introduction to the mathematical models of population dynamics in mathematical biology The focus of this book is on the biological meaning translation of mathematical structures in mathematical models rather than simply explaining mathematical details and literacies to analyze a model In some recent usages of the mathematical model simply with computer numerical calculations the model includes some inappropriate mathematical structure concerning the reasonability of modeling for the biological problem under investigation For students and researchers who study or use mathematical models it is important and helpful to understand what mathematical setup could be regarded as reasonable for the model with respect to the relation between the biological factors involved in the assumptions and the mathematical structure of the model Topics covered in this book are modeling with geometric progression density effect in population dynamics deriving continuous time models from discrete time models basic modeling for birth death stochastic processes continuous time models modeling interspecific reaction for the continuous time population dynamics model competition and prey predator dynamics modeling for population dynamics with a heterogeneous structure of population qualitative analysis on the discrete time dynamical system necessary knowledge about fundamental mathematical theories to understand the dynamical nature of continuous time models The book includes popular topics in ecology and mathematical biology as well as classic theoretical topics By understanding the biological meaning of modeling for simple models readers will be able to derive a specific mathematical model for a biological problem by reasonable

modeling The contents of this book is made accessible for readers without strong Mathematical background **Handbook of Models for Human Aging** P. Michael Conn,2011-04-28 The Handbook of Models for Human Aging is designed as the only comprehensive work available that covers the diversity of aging models currently available For each animal model it presents key aspects of biology nutrition factors affecting life span methods of age determination use in research and disadvantages advantes of use Chapters on comparative models take a broad sweep of age related diseases from Alzheimer s to joint disease cataracts cancer and obesity In addition there is an historical overview and discussion of model availability key methods and ethical issues Utilizes a multidisciplinary approach Shows tricks and approaches not available in primary publications First volume of its kind to combine both methods of study for human aging and animal models Over 200 illustrations *Stochastic Processes: Modeling and Simulation* D N Shanbhag,Calyampudi Radhakrishna Rao,2003-02-24

This sequel to volume 19 of Handbook on Statistics on Stochastic Processes Modelling and Simulation is concerned mainly with the theme of reviewing and in some cases unifying with new ideas the different lines of research and developments in stochastic processes of applied flavour This volume consists of 23 chapters addressing various topics in stochastic processes These include among others those on manufacturing systems random graphs reliability epidemic modelling self similar processes empirical processes time series models extreme value therapy applications of Markov chains modelling with Monte Carlo techniques and stochastic processes in subjects such as engineering telecommunications biology astronomy and chemistry particular with modelling simulation techniques and numerical methods concerned with stochastic processes The scope of the project involving this volume as well as volume 19 is already clarified in the preface of volume 19 The present volume completes the aim of the project and should serve as an aid to students teachers researchers and practitioners interested in applied stochastic processes

Inference for Diffusion Processes Christiane Fuchs,2013-01-18 Diffusion processes are a promising instrument for realistically modelling the time continuous evolution of phenomena not only in the natural sciences but also in finance and economics Their mathematical theory however is challenging and hence diffusion modelling is often carried out incorrectly and the according statistical inference is considered almost exclusively by theoreticians This book explains both topics in an illustrative way which also addresses practitioners It provides a complete overview of the current state of research and presents important novel insights The theory is demonstrated using real data applications

Optimal Control of Age-structured Populations in Economy, Demography, and the Environment Raouf Boucekkine,Natali Hritonenko,Yuri Yatsenko,2013-05-13 This book covers a wide range of topics within mathematical modelling and the optimization of economic demographic technological and environmental phenomena Each chapter is written by experts in their field and represents new advances in modelling theory and practice These essays are exemplary of the fruitful interaction between theory and practice when exploring global and local changes The unifying theme of the book is the use of mathematical models and optimization methods to describe age structured populations in economy demography

technological change and the environment Emphasis is placed on deterministic dynamic models that take age or size structures delay effects and non standard decision variables into account In addition the contributions deal with the age structure of assets resources and populations under study Interdisciplinary modelling has enormous potential for discovering new insights in global and regional development Optimal Control of Age structured Populations in Economy Demography and the Environment is a rich and excellent source of information on state of the art modelling expertise and references The book provides the necessary mathematical background for readers from different areas such as applied sciences management sciences and operations research which helps guide the development of practical models As well as this the book also surveys the current practice in applied modelling and looks at new research areas for a general mathematical audience This book will be of interest primarily to researchers postgraduate students as well as a wider scientific community including those focussing on the subjects of applied mathematics environmental sciences economics demography management and operations research

Methods and Models in Mathematical Biology Johannes Müller,Christina Kuttler,2015-08-13 This book developed from classes in mathematical biology taught by the authors over several years at the Technische Universität München The main themes are modeling principles mathematical principles for the analysis of these models and model based analysis of data The key topics of modern biomathematics are covered ecology epidemiology biochemistry regulatory networks neuronal networks and population genetics A variety of mathematical methods are introduced ranging from ordinary and partial differential equations to stochastic graph theory and branching processes A special emphasis is placed on the interplay between stochastic and deterministic models

Modeling Disease Spread and Control Tariq Halasa,Salome Dürr,2018-01-18 Mathematical models are useful tools to understand the epidemiology and agent host interaction of diseases They are developed and applied since over a century but with increasing computer capacity they become increasingly prominent as part of evidence based decision making Mathematical models are frequently used to construct preparedness and contingency plans for highly contagious diseases such as foot and mouth disease This allows proposing effective strategies to control the spread of the disease in case of an incursion and avails useful tools to support decision making during an outbreak They are also used to monitor prevent and control endemic diseases within populations or farms In addition mathematical models improve our understanding of the contact structure between farms pointing out risky elements in the contact network for disease introduction or further spread within the population This Research Topic presents valuable studies presenting different aspects and implementations of mathematical modeling for disease spread and control in the veterinary field The areas covered include model construction network analysis tools for decision makers and costeffective control of endemic diseases

Dynamical Systems and Their Applications in Biology Shigui Ruan,Gail Susan Kohl Wolkowicz,Jianhong Wu,Fields Institute for Research in Mathematical Sciences,2003-01-01 This volume is based on the proceedings of the International Workshop on Dynamical Systems and their Applications in Biology held at the Canadian

Coast Guard College on Cape Breton Island Nova Scotia Canada It presents a broad picture of the current research surrounding applications of dynamical systems in biology particularly in population biology The book contains 19 papers and includes articles on the qualitative and or numerical analysis of models involving ordinary partial functional and stochastic differential equations Applications include epidemiology population dynamics and physiology The material is suitable for graduate students and research mathematicians interested in ordinary differential equations and their applications in biology Also available by Ruan Wolkowicz and Wu is Differential Equations with Applications to Biology Volume 21 in the AMS series

Fields Institute Communications **Systems Medicine** ,2020-08-24 Technological advances in generated molecular and cell biological data are transforming biomedical research Sequencing multi omics and imaging technologies are likely to have deep impact on the future of medical practice In parallel to technological developments methodologies to gather integrate visualize and analyze heterogeneous and large scale data sets are needed to develop new approaches for diagnosis prognosis and therapy Systems Medicine Integrative Qualitative and Computational Approaches is an innovative interdisciplinary and integrative approach that extends the concept of systems biology and the unprecedented insights that computational methods and mathematical modeling offer of the interactions and network behavior of complex biological systems to novel clinically relevant applications for the design of more successful prognostic diagnostic and therapeutic approaches This 3 volume work features 132 entries from renowned experts in the fields and covers the tools methods algorithms and data analysis workflows used for integrating and analyzing multi dimensional data routinely generated in clinical settings with the aim of providing medical practitioners with robust clinical decision support systems Importantly the work delves into the applications of systems medicine in areas such as tumor systems biology metabolic and cardiovascular diseases as well as immunology and infectious diseases amongst others This is a fundamental resource for biomedical students and researchers as well as medical practitioners who need to need to adopt advances in computational tools and methods into the clinical practice Encyclopedic coverage one stop resource for access to information written by world leading scholars in the field of Systems Biology and Systems Medicine with easy cross referencing of related articles to promote understanding and further research Authoritative the whole work is authored and edited by recognized experts in the field with a range of different expertise ensuring a high quality standard Digitally innovative Hyperlinked references and further readings cross references and diagrams images will allow readers to easily navigate a wealth of information Journal of the Royal Society Interface

,2008 **Russian Journal of Numerical Analysis and Mathematical Modelling** ,2004 **Bulletin (new Series) of the American Mathematical Society** ,2005 Scientiae Mathematicae Japonicae ,2008 **Proceedings of the 2004 Pierce's Disease Research Symposium** ,2004 **Discrete and Continuous Dynamical Systems** ,2006 **The Mathematical Theory of Selection, Recombination, and Mutation** R. Bürger,2000-11-02 It is close to being a masterpiece could well be the classic presentation of the area Warren J Ewens University of Pennsylvania USA Population

genetics is concerned with the study of the genetic ecological and evolutionary factors that influence and change the genetic composition of populations. The emphasis here is on models that have a direct bearing on evolutionary quantitative genetics. Applications concerning the maintenance of genetic variation in quantitative traits and their dynamics under selection are treated in detail. Provides a unified, self-contained and in-depth study of the theory of multilocus systems. Introduces the basic population genetic models. Explores the dynamical and equilibrium properties of the distribution of quantitative traits under selection. Summarizes important results from more demanding sections in a comprehensible way. Employs a clear and logical presentation style. Following an introduction to elementary population genetics and discussion of the general theory of selection at two or more loci, the author considers a number of mutation selection models and derives the dynamical equations for polygenic traits under general selective regimes. The final chapters are concerned with the maintenance of quantitative genetic variation, the response to directional selection, the evolutionary role of deleterious mutations and other topics. Graduate students and researchers in population genetics, evolutionary theory and biomathematics will benefit from the in-depth coverage. This text will make an excellent reference volume for the fields of quantitative genetics, population and theoretical biology.

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