Mathematical Models In Biology

Dynamic Models in BiologyStochastic Models in BiologyA Primer in Mathematical Models in BiologyMathematical Models in BiologyMathematical Models in BiologyNeutral Models in BiologyMathematical Models in BiologyTheoretical Models in BiologyMathematical Models in BiologyLinear Models in BiologyMathematical Models in Biology and MedicineMathematical Models for Society and BiologyModels and Modeling in the SciencesDynamical Models in BiologyMathematical Modeling in Systems BiologySingle-Cell-Based Models in Biology and MedicineBiological Modeling and SimulationStructured Population Models in Biology and EpidemiologyModeling Biological Systems:Theoretical Models in Biology Stephen P. Ellner Narendra S. Goel Lee A. Segel Elizabeth S. Allman Elisabeth S. Allman Matthew H. Nitecki Elizabeth Spencer Allman Glenn W. Rowe Valeria Zazzu Michael R. Cullen IFIP-TC4 Working Conference on Mathematical Models in Biology and Medicine$ (1972 : Varna, Bulgarie) Edward Beltrami Stephen M. Downes Miklós Farkas Brian P. Ingalls Alexander Anderson Russell Schwartz Pierre Magal James W. Haefner Glenn W. Rowe

from controlling disease outbreaks to predicting heart attacks dynamic models are increasingly crucial for understanding biological processes many universities are starting undergraduate programs in computational biology to introduce students to this rapidly growing field in dynamic models in biology the first text on dynamic models specifically written for undergraduate students in the biological sciences ecologist stephen ellner and mathematician john guckenheimer teach students how

to understand build and use dynamic models in biology developed from a course taught by ellner and guckenheimer at cornell university the book is organized around biological applications with mathematics and computing developed through case studies at the molecular cellular and population levels the authors cover both simple analytic models the sort usually found in mathematical biology texts and the complex computational models now used by both biologists and mathematicians linked to a site with computer lab materials and exercises dynamic models in biology is a major new introduction to dynamic models for students in the biological sciences mathematics and engineering

stochastic models in biology describes the usefulness of the theory of stochastic process in studying biological phenomena the book describes analysis of biological systems and experiments though probabilistic models rather than deterministic methods the text reviews the mathematical analyses for modeling different biological systems such as the random processes continuous in time and discrete in state space the book also discusses population growth and extinction through malthus law and the work of macarthur and wilson the text then explains the dynamics of a population of interacting species the book also addresses population genetics under systematic evolutionary pressures known as deterministic equations and genetic changes in a finite population known as stochastic equations the text then turns to stochastic modeling of biological systems at the molecular level particularly the kinetics of biochemical reactions the book also presents various useful equations such as the differential equation for generating functions for birth and death processes the text can prove valuable for biochemists cellular biologists and researchers in the medical and chemical field who are tasked to perform data analysis

a textbook on mathematical modelling techniques with powerful applications to biology combining theoretical exposition with exercises and examples

this introductory textbook on mathematical biology focuses on discrete models across a variety of biological subdisciplines biological topics treated include linear and non linear models of populations markov models of molecular evolution phylogenetic tree construction genetics and infectious disease models the coverage of models of molecular evolution and phylogenetic tree construction from dna sequence data is unique among books at this level computer investigations with matlab are incorporated throughout in both exercises and more extensive projects to give readers hands on experience with the mathematical models developed matlab programs accompany the text mathematical tools such as matrix algebra eigenvector analysis and basic probability are motivated by biological models and given self contained developments so that mathematical prerequisites are minimal

neutral models are constructed to help scientists understand complex patterns of form structure or behavior that may not be observed directly in this unique volume eight distinguished scientists present a comprehensive study of the use of neutral models in testing biological theories they describe the principles of model testing

and explore how they are applied to research in molecular biology genetics ecology evolution and paleontology in addition to the editors the contributors include stephen stigler david raup paul harvey l b slobodkin stuart kauffman william wimsatt and james crow

this book surveys theoretical models in three broad areas of biology the origin of life the immune system and memory in the brain introducing mathematical and mainly computational models that have been used to construct simulations most current books on theoretical biology fall into one of two categories a books that specialize in one area of biology and treat theoretical models in considerable depth and b books that concentrate on purely mathematical models with computers used only to find numerical solutions to differential equations for example although some mathmatical models are considered in this book the main emphasis is on stochastic computer modles of biological systems such techniques have a much greater potential for producting detailed realistic models of individual systems and are likely to be the preferred modelling methods of the future by considering three different areas in biology the book shows how several of these modelling techniques have been successfully applied in diverse areas put simply this book is important becuase it shows how the power of modern computers is allowing researchers in theoretical biology to break free of the constraints on modelling that were imposed by the traditional differential equation approach anyone who is interested in the theoretical models of complicated living systems should have this in his or her library g b ermentrout bulletin of mathematical biology

this book presents an exciting collection of contributions based on the workshop bringing maths to life held october 27 29 2014 in naples italy the state of the art research in biology and the statistical and analytical challenges facing huge masses of data collection are treated in this work specific topics explored in depth surround the sessions and special invited sessions of the workshop and include genetic variability via differential expression molecular dynamics and modeling complex biological systems viewed from quantitative models and microscopy images processing to name several in depth discussions of the mathematical analysis required to extract insights from complex bodies of biological datasets to aid development in the field novel algorithms methods and software tools for genetic variability molecular dynamics and complex biological systems are presented in this book researchers and graduate students in biology life science and mathematics statistics will find the content useful as it addresses existing challenges in identifying the gaps between mathematical modeling and biological research the shared solutions will aid and promote further collaboration between life sciences and mathematics

mathematical modeling for society and biology engagingly relates mathematics to compelling real life problems in biology and contemporary society it shows how mathematical tools can be used to gain insight into these modern common problems to provide effective real solutions beltrami s creative non threatening approach draws on a wealth of interesting examples pertaining to current social and biological issues central ideas appear again in different contexts throughout the book showing the general unity of the modeling process the models are strikingly novel and based on issues of real concern most have never appeared in book form

through the relevance of these models mathematics becomes not just figures and numbers but a means to a more refined understanding of the world

biologists climate scientists and economists all rely on models to move their work forward in this book stephen m downes explores the use of models in these and other fields to introduce readers to the various philosophical issues that arise in scientific modeling readers learn that paying attention to models plays a crucial role in appraising scientific work this book first presents a wide range of models from a number of different scientific disciplines after assembling some illustrative examples downes demonstrates how models shed light on many perennial issues in philosophy of science and in philosophy in general reviewing the range of views on how models represent their targets introduces readers to the key issues in debates on representation not only in science but in the arts as well also standard epistemological questions are cast in new and interesting ways when readers confront the question what makes for a good or bad model all examples from the sciences and positions in the philosophy of science are presented in an accessible manner the book is suitable for undergraduates with minimal experience in philosophy and an introductory undergraduate experience in science key features the book serves as a highly accessible philosophical introduction to models and modeling in the sciences presenting all philosophical and scientific issues in a nontechnical manner students and other readers learn to practice philosophy of science by starting with clear examples taken directly from the sciences while not comprehensive this book introduces the reader to a wide range of views on key issues in the philosophy of science

dynamic models in biology offers an introduction to modern mathematical biology this book provides a short introduction to modern mathematical methods in modeling dynamical phenomena and treats the broad topics of population dynamics epidemiology evolution immunology morphogenesis and pattern formation primarily employing differential equations the author presents accessible descriptions of difficult mathematical models recent mathematical results are included but the author s presentation gives intuitive meaning to all the main formulae besides mathematicians who want to get acquainted with this relatively new field of applications this book is useful for physicians biologists agricultural engineers and environmentalists key topics include chaotic dynamics of populations the spread of sexually transmitted diseases problems of the origin of life models of immunology formation of animal hide patterns the intuitive meaning of mathematical formulae explained with many figures applying new mathematical results in modeling biological phenomena miklos farkas is a professor at budapest university of technology where he has researched and instructed mathematics for over thirty years he has taught at universities in the former soviet union canada australia venezuela nigeria india and columbia prof farkas received the 1999 bolyai award of the hungarian academy of science and the 2001 albert szentgyorgyi award of the hungarian ministry of education a down to earth introduction to the growing field of modern mathematical biology also includes appendices which provide background material that goes beyond advanced calculus and linear algebra

an introduction to the mathematical concepts and techniques needed for the construction and analysis of models in molecular systems biology systems techniques are integral to current research in molecular cell biology and system level investigations are often accompanied by mathematical models these models serve as working hypotheses they help us to understand and predict the behavior of complex systems this book offers an introduction to mathematical concepts and techniques needed for the construction and interpretation of models in molecular systems biology it is accessible to upper level undergraduate or graduate students in life science or engineering who have some familiarity with calculus and will be a useful reference for researchers at all levels the first four chapters cover the basics of mathematical modeling in molecular systems biology the last four chapters address specific biological domains treating modeling of metabolic networks of signal transduction pathways of gene regulatory networks and of electrophysiology and neuronal action potentials chapters 3 8 end with optional sections that address more specialized modeling topics exercises solvable with pen and paper calculations appear throughout the text to encourage interaction with the mathematical techniques more involved end of chapter problem sets require computational software appendixes provide a review of basic concepts of molecular biology additional mathematical background material and tutorials for two computational software packages xppaut and matlab that can be used for model simulation and analysis

many different single cell based models have been developed and applied to biological and medical problems computational approaches used are monte carlo simulations energy minimisation techniques volume conservation laws solutions of the equations of motion for each individual cell or for each point on the cell membrane they differ in the level of detail that defines the cell structure and subsequently in the number of individual cells that the model can incorporate this volume presents a collection of mathematical and computational single cell based models and their application the main sections cover four general model groupings hybrid cellular automata cellular potts lattice free cells and viscoelastic cells each section is introduced by a discussion of the applicability of the particular modelling approach and its advantages and disadvantages which will make the book suitable for students starting research in mathematical biology as well as scientists modelling multicellular processes

a practice oriented survey of techniques for computational modeling and simulation suitable for a broad range of biological problems there are many excellent computational biology resources now available for learning about methods that have been developed to address specific biological systems but comparatively little attention has been paid to training aspiring computational biologists to handle new and unanticipated problems this text is intended to fill that gap by teaching students how to reason about developing formal mathematical models of biological systems that are amenable to computational analysis it collects in one place a selection of broadly useful models algorithms and theoretical analysis tools normally found scattered among many other disciplines it thereby gives the aspiring student a bag of tricks that will serve him or her well in modeling problems drawn from numerous subfields of biology these techniques are taught from the

perspective of what the practitioner needs to know to use them effectively supplemented with references for further reading on more advanced use of each method covered the text which grew out of a class taught at carnegie mellon university covers models for optimization simulation and sampling and parameter tuning these topics provide a general framework for learning how to formulate mathematical models of biological systems what techniques are available to work with these models and how to fit the models to particular systems their application is illustrated by many examples drawn from a variety of biological disciplines and several extended case studies that show how the methods described have been applied to real problems in biology

in this new century mankind faces ever more challenging environmental and publichealthproblems suchaspollution invasionbyexoticspecies theem gence of new diseases or the emergence of diseases into new regions west nile virus sars anthrax etc andtheresurgenceofexistingdiseases in uenza malaria tb hiv aids etc mathematical models have been successfully used to study many biological epidemiological and medical problems and nonlinear and complex dynamics have been observed in all of those contexts mathematical studies have helped us not only to better understand these problems but also to nd solutions in some cases such as the prediction and control of sars outbreaks understanding hiv infection and the investi tion of antibiotic resistant infections in hospitals structuredpopulationmodelsdistinguishindividualsfromoneanother cording to characteristics such as age size location status and movement to determine the birth growth and death rates interaction with each other and with environment infectivity etc the goal of structured population models is to understand how these characteristics a ect the dynamics of these models and thus the outcomes and consequences of the biological and epidemiolo cal processes there is a very large and growing body of literature on these topics this book deals with the recent and important advances in the study of structured population models in biology and epidemiology there are six chapters in this book written by leading researchers in these areas

this book surveys theoretical models in three broad areas of biology the origin of life the immune system and memory in the brain introducing mathematical and mainly computational methods that have been used to construct simulations most current books on theoretical biology fall into one of two categories a books that specialize in one area of biology and treat theoretical models in considerable depth and b books that concentrate on purely mathematical models with computers used only to find numerical solutions to differential equations for example although some mathematical models are considered in this book the main emphasis is on stochastic computer models of biological systems such techniques have a much greater potential for producing detailed realistic models of individual systems and are likely to be the preferred modelling methods of the future by considering three different areas in biology the book shows how several of these modelling techniques have been successfully applied in diverse areas put simply this book is important because it shows how the power of modern computers is allowing researchers in theoretical biology to break free of the constraints modelling that were imposed by the traditional differential equation approach

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