

Alessandra Rogato · Valeria Zazzu
Mario Guarracino *Editors*

Dynamics of Mathematical Models in Biology

Bringing Mathematics to Life

 Springer

Mathematical Models Biology Bringing Mathematics

Peipei Pang



Mathematical Models Biology Bringing Mathematics:

Mathematical Models in Biology Valeria Zazzu, Maria Brigida Ferraro, Mario R. Guarracino, 2015-11-26 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.

Dynamics of Mathematical Models in Biology Alessandra Rogato, Valeria Zazzu, Mario Guarracino, 2016-11-03 This volume focuses on contributions from both the mathematics and life science community surrounding the concepts of time and dynamicity of nature two significant elements which are often overlooked in modeling process to avoid exponential computations. The book is divided into three distinct parts: dynamics of genomes and genetic variation, dynamics of motifs and dynamics of biological networks. Chapters included in dynamics of genomes and genetic variation analyze the molecular mechanisms and evolutionary processes that shape the structure and function of genomes and those that govern genome dynamics. The dynamics of motifs portion of the volume provides an overview of current methods for motif searching in DNA, RNA and proteins a key process to discover emergent properties of cells tissues and organisms. The part devoted to the dynamics of biological networks covers networks aptly discusses networks in complex biological functions and activities that interpret processes in cells. Moreover chapters in this section examine several mathematical models and algorithms available for integration analysis and characterization. Once life scientists began to produce experimental data at an unprecedented pace it became clear that mathematical models were necessary to interpret data to structure information with the aim to unveil biological mechanisms discover results and make predictions. The second annual Bringing Maths to Life workshop held in Naples Italy October 2015 enabled a bi-directional flow of ideas from and international group of mathematicians and biologists. The venue allowed mathematicians to introduce novel algorithms methods and software that may be useful to model aspects of life science and life scientists posed new challenges for mathematicians. [Mathematical Models in Biology](#) Leah Edelstein-Keshet, 1987-12-01 This book is an introduction for readers interested in biological applications of mathematics and modeling in biology showing how relatively simple mathematics can be applied to a variety of models.

Despite the great advances that have taken place the simple lessons described in the text are still important and informative

A Course in Mathematical Biology Gerda de Vries, Thomas Hillen, Mark Lewis, Johannes Müller, Birgitt Schöfnisch, 2006-07-01 This is the only book that teaches all aspects of modern mathematical modeling and that is specifically designed to introduce undergraduate students to problem solving in the context of biology Included is an integrated package of theoretical modeling and analysis tools computational modeling techniques and parameter estimation and model validation methods with a focus on integrating analytical and computational tools in the modeling of biological processes Divided into three parts it covers basic analytical modeling techniques introduces computational tools used in the modeling of biological problems and includes various problems from epidemiology ecology and physiology All chapters include realistic biological examples including many exercises related to biological questions In addition 25 open ended research projects are provided suitable for students An accompanying Web site contains solutions and a tutorial for the implementation of the computational modeling techniques Calculations can be done in modern computing languages such as Maple Mathematica and MATLAB

Mathematical Models for Society and Biology Edward Beltrami, 2013-06-19 *Mathematical Models for Society and Biology 2e* is a useful resource for researchers graduate students and post docs in the applied mathematics and life science fields Mathematical modeling is one of the major subfields of mathematical biology A mathematical model may be used to help explain a system to study the effects of different components and to make predictions about behavior *Mathematical Models for Society and Biology 2e* draws on current issues to engagingly relate how to use mathematics to gain insight into problems in biology and contemporary society For this new edition author Edward Beltrami uses mathematical models that are simple transparent and verifiable Also new to this edition is an introduction to mathematical notions that every quantitative scientist in the biological and social sciences should know Additionally each chapter now includes a detailed discussion on how to formulate a reasonable model to gain insight into the specific question that has been introduced Offers 40% more content 5 new chapters in addition to revisions to existing chapters Accessible for quick self study as well as a resource for courses in molecular biology biochemistry embryology and cell biology medicine ecology and evolution bio mathematics and applied math in general Features expanded appendices with an extensive list of references solutions to selected exercises in the book and further discussion of various mathematical methods introduced in the book

Introduction to Mathematical Biology Ching Shan Chou, Avner Friedman, 2016-04-27 This book is based on a one semester course that the authors have been teaching for several years and includes two sets of case studies The first includes chemostat models predator prey interaction competition among species the spread of infectious diseases and oscillations arising from bifurcations In developing these topics readers will also be introduced to the basic theory of ordinary differential equations and how to work with MATLAB without having any prior programming experience The second set of case studies were adapted from recent and current research papers to the level of the students Topics have been selected based on public

health interest This includes the risk of atherosclerosis associated with high cholesterol levels cancer and immune interactions cancer therapy and tuberculosis Readers will experience how mathematical models and their numerical simulations can provide explanations that guide biological and biomedical research Considered to be the undergraduate companion to the more advanced book *Mathematical Modeling of Biological Processes* A Friedman C Y Kao Springer 2014 this book is geared towards undergraduate students with little background in mathematics and no biological background

Mathematical Models in the Biosciences I Michael Frame, 2021-06-22 An award winning professor's introduction to essential concepts of calculus and mathematical modeling for students in the biosciences This is the first of a two part series exploring essential concepts of calculus in the context of biological systems Michael Frame covers essential ideas and theories of basic calculus and probability while providing examples of how they apply to subjects like chemotherapy and tumor growth chemical diffusion allometric scaling predator prey relations and nerve impulses Based on the author's calculus class at Yale University the book makes concepts of calculus more relatable for science majors and premedical students

Mathematical Modeling in Systems Biology Brian P. Ingalls, 2013-07-05 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

New Approaches to Identifying Structures Using Geometric Structure Analysis: Design and Adaptation Karabutov, Nikolay Nikolayevich, 2025-09-10 An actual problem of identification theory is considered related to the non formalized task of evaluating the model structure Novel approaches to structural identification SI propose solutions to various problems of identification theory based on the analysis of geometric frameworks GFs This formalized approach to the structural identifiability SID for nonlinear dynamical systems of various classes shows that structural identifiability follows

from SI Additionally based on the GF estimates for the Lyapunov exponents LEs of dynamical systems are shown to be recoverable detectable and identifiable When combined with synthesized methods and algorithms they can be applied to the construction of mathematical models for complex processes and systems Thus they can be used in decision making systems process forecasting control of nonlinear systems and processing of heterogeneous time series Novel Approaches to Structural Identification Using Geometric Framework Analysis proposes various solutions to the problem of identification theory It discusses the development of adaptive identification and control systems for analyzing complex processes and systems Covering topics such as parametric restrictions distributed lags and interconnected systems this book is an excellent resource for data analysis specialists mathematical software developers professionals researchers scholars academicians and more

Dynamic Models in Biology Stephen P. Ellner, John Guckenheimer, 2006-04-16 What are dynamic models Matrix models and structured population dynamics Membrane channels and action potentials Cellular Dynamics pathways of gene expression Dynamical systems Differential equations models for infectious disease Spatial patterns in biology Agent based and other computational models for complex systems Building dynamic models *Exploring Mathematical Modeling in Biology Through Case Studies and Experimental Activities* Rebecca Sanft, Anne Walter, 2020-03-30 Exploring Mathematical Modeling in Biology through Case Studies and Experimental Activities provides supporting materials for courses taken by students majoring in mathematics computer science or in the life sciences The book's cases and lab exercises focus on hypothesis testing and model development in the context of real data The supporting mathematical coding and biological background permit readers to explore a problem understand assumptions and the meaning of their results The experiential components provide hands on learning both in the lab and on the computer As a beginning text in modeling readers will learn to value the approach and apply competencies in other settings Included case studies focus on building a model to solve a particular biological problem from concept and translation into a mathematical form to validating the parameters testing the quality of the model and finally interpreting the outcome in biological terms The book also shows how particular mathematical approaches are adapted to a variety of problems at multiple biological scales Finally the labs bring the biological problems and the practical issues of collecting data to actually test the model and or adapting the mathematics to the data that can be collected Presents a single volume on mathematics and biological examples with data and wet lab experiences suitable for non experts Contains three real world biological case studies and one wet lab for application of the mathematical models Includes R code templates throughout the text which are also available through an online repository along with the necessary data files to complete all projects and labs [Plant Genetics and Biotechnology in Biodiversity](#) Rosa Rao, Giandomenico Corrado, 2018-08-09 This book is a printed edition of the Special Issue Plant Genetics and Biotechnology in Biodiversity that was published in *Diversity* *Trends in Biomathematics: Modeling Health Across Ecology, Social Interactions, and Cells* Rubem P. Mondaini, 2025-09-26 This volume compiles selected peer reviewed papers presented at the 24th International

Symposium on Mathematical and Computational Biology BIOMAT 2024 held from October 27 to November 1 2024 at the Orthodox Academy of Crete in Kolympari Crete Island Greece The book covers a wide range of topics from epidemiological modeling and optimal infection control to the application of machine learning and artificial intelligence in cell biology imaging It also explores the dynamics of disease spread protein structure modeling and mathematical models of HIV 1 COVID 19 monkeypox and measles featuring contributions from some of the most esteemed researchers in the field as well as findings from a new generation of researchers fostering cross disciplinary collaborations Carefully edited this volume will appeal to both researchers and students looking for topics for further study Previous BIOMAT volumes from 2018 to 2024 are also available from Springer

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

Mathematical Models in Biology Elizabeth Spencer Allman, John A. Rhodes, 2004 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

Biosorption Processes for Heavy Metal Removal Saini, Pinki, 2024-03-11 Persistent and non degradable heavy metals stand as pollutants with the potential for severe ecological repercussions when released into the environment Municipal and industrial wastewater face a high risk of contamination by these hazardous substances posing a formidable challenge to water treatment technologies The imperative is clear effective and affordable methods for effluent treatment and metal recovery are essential for meeting regulatory standards and unlocking the latent value of valuable metals within the waste However new methods of accomplishing this challenge are necessary for increasing the effectiveness in both cost and application

Biosorption Processes for Heavy Metal Removal comprehensively explores the imperative to remove heavy metals from waste streams It provides an insightful overview of biosorbents and biosorption technology focusing on their underlying biosorption features The compilation within this book comprises of a series of review articles delving into the current understanding of biosorption mechanisms and biochemistry the efficacy of bacterial fungal and algal biomass and practical considerations for biosorbent preparation and engineering The physicochemical evaluations of biosorbents process optimization and factors influencing biosorption efficiency are also covered Furthermore the book explores biosorption applications for removing nutrients organic pollutants and metals in wastewater treatment across diverse contexts Geared towards administrators policymakers consultants industry professionals academicians scientists researchers and graduate and post graduate students in environmental sciences and related fields this book serves as their comprehensive reference

Biomath: Unveiling the Secrets of Life Through the Lens of Mathematics Pasquale De Marco, 2025-04-11 In the realm of science where disciplines intertwine and boundaries blur there lies a captivating field that bridges the gap between mathematics and biology biomathematics This book invites readers to embark on an intellectual odyssey exploring the profound connections between the elegance of mathematics and the intricacies of life From the microscopic world of cells to the vast ecosystems that shape our planet biomathematics offers a powerful lens through which we can understand the fundamental principles governing life's processes Mathematical models simulations and algorithms provide valuable tools for biologists to analyze predict and manipulate biological systems leading to advancements in medicine biotechnology and ecology This comprehensive book delves into the diverse applications of biomathematics showcasing its impact on various fields Readers will discover how mathematical models have revolutionized drug discovery enabling the development of targeted therapies with greater precision and fewer side effects They will also explore the role of biomathematics in genetic engineering where mathematical techniques are used to modify organisms and create new biological products Furthermore the book delves into the intricate world of ecology where mathematical models help scientists understand the complex interactions between species and their environment These models aid in predicting population dynamics evaluating conservation strategies and addressing global challenges such as climate change and biodiversity loss Biomathematics is not

just a collection of techniques it is a mindset a way of thinking that seeks to quantify and understand the complexities of life This book is an invitation to adopt this mindset to see the world through the eyes of a biomathematician With clear explanations engaging examples and thought provoking questions the book guides readers through the fascinating world of biomathematics inspiring them to explore the hidden connections between mathematics and life Whether you are a biologist seeking to harness the power of mathematics or a mathematician eager to explore the complexities of life this book will provide a rich source of knowledge and inspiration Embark on this captivating journey today and discover the profound beauty and elegance that lies at the intersection of mathematics and biology If you like this book write a review on google books

Proceedings of the Fourth International Congress on Mathematical Education M. Zweng, Green, Kilpatrick, Pollack, Suydam, 2012-12-06 Henry O Pollak Chairman of the International Program Committee Bell Laboratories Murray Hill New Jersey USA The Fourth International Congress on Mathematics Education was held in Berkeley California USA August 10 16 1980 Previous Congresses were held in Lyons in 1969 Exeter in 1972 and Karlsruhe in 1976 Attendance at Berkeley was about 1800 full and 500 associate members from about 90 countries at least half of these come from outside of North America About 450 persons participated in the program either as speakers or as presiders approximately 40 percent of these came from the U S or Canada There were four plenary addresses they were delivered by Hans Freudenthal on major problems of mathematics education Hermina Sinclair on the relationship between the learning of language and of mathematics Seymour Papert on the computer as carrier of mathematical culture and Hua Loo Keng on popularising and applying mathematical methods Gerge Polya was the honorary president of the Congress illness prevented his planned attendance but he sent a brief presentation entitled Mathematics Improves the Mind There was a full program of speakers panelists debates miniconferences and meetings of working and study groups In addition 18 major projects from around the world were invited to make presentations and various groups representing special areas of concern had the opportunity to meet and to plan their future activities

A Primer in Mathematical Models in Biology Lee A. Segel, Leah Edelstein-Keshet, 2013-01-01 This textbook introduces differential equations biological applications and simulations and emphasizes molecular events biochemistry and enzyme kinetics excitable systems neural signals and small protein and genetic circuits A Primer on Mathematical Models in Biology will appeal to readers because it grew out of a course that the popular and highly respected applied mathematician Lee Segel taught at the Weizmann Institute and it represents his unique perspective combines clear and useful mathematical methods with applications that illustrate the power of such tools and includes many exercises in reasoning modeling and simulations

[Using Mathematics to Understand Biological Complexity](#) Rebecca Segal, Blerta Shtylla, Suzanne Sindi, 2020-12-29 This volume tackles a variety of biological and medical questions using mathematical models to understand complex system dynamics Working in collaborative teams of six each with a senior research mentor researchers developed new mathematical models to address questions in a range of application areas Topics

include retinal degeneration biopolymer dynamics the topological structure of DNA ensemble analysis multidrug resistant organisms tumor growth modeling and geospatial modeling of malaria The work is the result of newly formed collaborative groups begun during the Collaborative Workshop for Women in Mathematical Biology hosted by the Institute of Pure and Applied Mathematics at UCLA in June 2019 Previous workshops in this series have occurred at IMA NIMBioS and MBI

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