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dissipation theorems age structured models and statistical field theory. This text is primarily aimed at graduate students and researchers working in mathematical biology statistical and biological physicists and applied mathematicians interested in stochastic modeling Applied probabilists should also find it of interest It provides significant background material in applied mathematics and statistical physics and introduces concepts in stochastic and nonequilibrium processes via motivating biological applications The book is highly illustrated and contains a large number of examples and exercises that further develop the models and ideas in the body of the text It is based on a course that the author has taught at the University of Utah for many years Quantitative Biology Brian Munsky, William S. Hlavacek, Lev S. Tsimring, 2018-08-21 An introduction to the quantitative modeling of biological processes presenting modeling approaches methodology practical algorithms software tools and examples of current research The quantitative modeling of biological processes promises to expand biological research from a science of observation and discovery to one of rigorous prediction and quantitative analysis The rapidly growing field of quantitative biology seeks to use biology s emerging technological and computational capabilities to model biological processes This textbook offers an introduction to the theory methods and tools of quantitative biology The book first introduces the foundations of biological modeling focusing on some of the most widely used formalisms It then presents essential methodology for model guided analyses of biological data covering such methods as network reconstruction uncertainty quantification and experimental design practical algorithms and software packages for modeling biological systems and specific examples of current quantitative biology research and related specialized methods Most chapters offer problems progressing from simple to complex that test the reader's mastery of such key techniques as deterministic and stochastic simulations and data analysis Many chapters include snippets of code that can be used to recreate analyses and generate figures related to the text Examples are presented in the three popular computing languages Matlab R and Python A variety of online resources supplement the text The editors are long time organizers of the Annual g bio Summer School which was founded in 2007 Through the school the editors have helped to train more than 400 visiting students in Los Alamos NM Santa Fe NM San Diego CA Albuguerque NM and Fort Collins CO This book is inspired by the school's curricula and most of the contributors have participated in the school as students lecturers or both Contributors John H Abel Roberto Bertolusso Daniela Besozzi Michael L Blinov Clive G Bowsher Fiona A Chandra Paolo Cazzaniga Bryan C Daniels Bernie J Daigle Jr Maciej Dobrzynski Jonathan P Doye Brian Drawert Sean Fancer Gareth W Fearnley Dirk Fey Zachary Fox Ramon Grima Andreas Hellander Stefan Hellander David Hofmann Damian Hernandez William S Hlavacek Jianjun Huang Tomasz Jetka Dongya Jia Mohit Kumar Jolly Boris N Kholodenko Markek Kimmel Micha Komorowski Ganhui Lan Heeseob Lee Herbert Levine Leslie M Loew Jason G Lomnitz Ard A Louis Grant Lythe Carmen Molina Par s Ion I Moraru Andrew Mugler Brian Munsky Joe Natale Ilya Nemenman Karol Niena towski Marco S Nobile Maria Nowicka Sarah Olson Alan S Perelson Linda R Petzold Sreenivasan Ponnambalam Arya Pourzanjani Ruy M Ribeiro William Raymond William

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