

1. Record Nr.	UNINA9910299787303321
Autore	Deuflhard Peter
Titolo	A Guide to Numerical Modelling in Systems Biology / / by Peter Deuflhard, Susanna Röblitz
Pubbl/distr/stampa	Cham : , : Springer International Publishing : , : Imprint : Springer, , 2015
ISBN	3-319-20059-3
Edizione	[1st ed. 2015.]
Descrizione fisica	1 online resource (X, 168 p. 42 illus., 33 illus. in color.)
Collana	Texts in Computational Science and Engineering, , 1611-0994 ; ; 12
Disciplina	515.352
Soggetti	Differential equations Applied mathematics Engineering mathematics Biomathematics Computer simulation Physics Ordinary Differential Equations Applications of Mathematics Mathematical and Computational Biology Simulation and Modeling Numerical and Computational Physics, Simulation
Lingua di pubblicazione	Inglese
Formato	Materiale a stampa
Livello bibliografico	Monografia
Note generali	Bibliographic Level Mode of Issuance: Monograph
Nota di contenuto	Preface -- Outline -- ODE Models for Systems Biological Networks -- Numerical Simulation of ODE Models -- Parameter Identification in ODE Models -- Appendix -- Software -- Index.
Sommario/riassunto	This book is intended for students of computational systems biology with only a limited background in mathematics. Typical books on systems biology merely mention algorithmic approaches, but without offering a deeper understanding. On the other hand, mathematical books are typically unreadable for computational biologists. The authors of the present book have worked hard to fill this gap. The result is not a book on systems biology, but on computational methods in systems biology. This book originated from courses taught by the

authors at Freie Universität Berlin. The guiding idea of the courses was to convey those mathematical insights that are indispensable for systems biology, teaching the necessary mathematical prerequisites by means of many illustrative examples and without any theorems. The three chapters cover the mathematical modelling of biochemical and physiological processes, numerical simulation of the dynamics of biological networks, and identification of model parameters by means of comparisons with real data. Throughout the text, the strengths and weaknesses of numerical algorithms with respect to various systems biological issues are discussed. Web addresses for downloading the corresponding software are also included. .
