Record Nr. UNINA9910451155703321 BIOMAT 2005 [[electronic resource]]: proceedings of the International Titolo Symposium on Mathematical and Computational Biology, Rio de Janeiro. Brazil, 3-8 December 2005 / / edited by Rubem P. Mondaini, Rui Dilao Singapore;; Hackensack, NJ,: World Scientific, c2006 Pubbl/distr/stampa **ISBN** 1-281-92491-1 9786611924911 981-277-368-1 Descrizione fisica 1 online resource (408 p.) Altri autori (Persone) MondainiR (Rubem) DilaoRui 570.1/5118 Disciplina Soggetti Biology - Mathematical models **Biomathematics** Electronic books. Lingua di pubblicazione Inglese **Formato** Materiale a stampa Livello bibliografico Monografia "The BIOMAT 2005 International Symposium on Mathematical and Note generali Computational Biology, together with the Fifth Brazilian Symposium on Mathematical and Computational Biology, was held in the city of Petropolis, state of Rio de Janeiro, Brazil ..."--Pref. Nota di bibliografia Includes bibliographical references and index. Nota di contenuto : Editorial Board of the BIOMAT Contents : Preface Consortium ; Biological Modeling ; Modelling aspects of vascular cancer development : 1. Introduction ; 2. Brief biological background : 3. Cellular automaton model ; 4. Effects of hypoxia on cell cycle dynamics 5. The role of acidity 6. Discussion : Modelling cooperative phenomena in interacting References cell systems with cellular automata; 1. Introduction: roots of cellular automata ; 2. Cellular automaton definition : 3. Cellular automaton models of cell interaction 4. An example: a cellular automaton model of a vascular tumor growth ; A mathematical 5. Discussion : References

analysis of cylindrical shaped aneurysms ; 1. Introduction : 2. The Mathematical Model ; 3. The Dynamical Properties of the Model ; 4. Discussion ; References On the origin of metazoans 1. Introduction ; 2. Pattern Formation; 3. Coupling Pattern to Cell Shape ; 5. On Urbilateria ; 4. On Urcnidaria Appendix A ; Appendix B ; References; A software tool to model genetic regulatory networks: Applications to segmental patterning in Drosophila 1. Introduction 2. A model for the regulation of gene expression : 3. The GeNetSim ; 4. Segmental patterning in Drosophila package ; References ; The mitochondrial Eve in an exponentially growing population and a critique to the out of Africa model for human evolution

; 1. Introduction

2. Percolation and mitochondrial DNA basics

## Sommario/riassunto

This volume contains the contributions of the keynote speakers to the BIOMAT 2005 symposium, as well as a collection of selected papers by pioneering researchers. It provides a comprehensive review of the mathematical modeling of cancer development, Alzheimer's disease, malaria, and aneurysm development. Various models for the immune system and epidemiological issues are analyzed and reviewed. The book also explores protein structure prediction by optimization and combinatorial techniques (Steiner trees). The coverage includes bioinformatics issues, regulation of gene expression, evolution, d