Alfonsas Juška

Analysis of biological processes

Verbal description, schemes, and mathematical models

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Molecular Pathology William B. Coleman, Gregory J. Tsongalis, 2017-11-09 As the molecular basis of human disease becomes better characterized and the implications for understanding the molecular basis of disease becomes realized through improved diagnostics and treatment Molecular Pathology Second Edition stands out as the most comprehensive textbook where molecular mechanisms represent the focus It is uniquely concerned with the molecular basis of major human diseases and disease processes presented in the context of traditional pathology with implications for translational molecular medicine The Second Edition of Molecular Pathology has been thoroughly updated to reflect seven years of exponential changes in the fields of genetics molecular and cell biology which molecular pathology translates in the practice of molecular medicine The textbook is intended to serve as a multi use textbook that would be appropriate as a classroom teaching tool for biomedical graduate students medical students allied health students and others such as advanced undergraduates Further this textbook will be valuable for pathology residents and other postdoctoral fellows that desire to advance their understanding of molecular mechanisms of disease beyond what they learned in medical graduate school In addition this textbook is useful as a reference book for practicing basic scientists and physician scientists that perform disease related basic science and translational research who require a ready information resource on the molecular basis of various human diseases and disease states Explores the principles and practice of molecular pathology molecular pathogenesis molecular mechanisms of disease and how the molecular pathogenesis of disease parallels the evolution of the disease Explains the practice of molecular medicine and the translational aspects of molecular pathology Teaches from the perspective of integrative systems biology Enhanced digital version included with purchase **Biological Networks** Rudiyanto Gunawan, Neda Bagheri, 2019-01-10 This book is a printed edition of the Special Issue Biological Networks that was published in Processes Intelligent Computing Theories and Application De-Shuang Huang, Kang-Hyun Jo, Juan Carlos Figueroa-García, 2017-07-18 This three volume set LNCS 10361 LNCS 10362 and LNAI 10363 constitutes the refereed proceedings of the 13th International Conference on Intelligent Computing ICIC 2017 held in Liverpool UK in August 2017 The 221 full papers and 15 short papers of the three proceedings volumes were carefully reviewed and selected from 639 submissions This second volume of the set comprises 74 papers The papers are organized in topical sections such as Pattern Recognition Image Processing Virtual Reality and Human Computer Interaction Healthcare Informatics Theory and Methods Genetic Algorithms Blind Source Separation Intelligent Fault Diagnosis Machine Learning Knowledge Discovery and Data Mining Gene Expression Array Analysis Systems Biology Modeling Simulation and Optimization of Biological Systems Intelligent Computing in Computational Biology Computational Genomics Computational Proteomics Gene Regulation Modeling and Analysis SNPs and Haplotype Analysis Protein Protein Interaction Prediction Protein Structure and Function Prediction Next Gen Sequencing and Metagenomics Structure Prediction and Folding Biomarker Discovery Applications of Machine Learning Techniques to Computational Proteomics Genomics and Biological Sequence Analysis Biomedical Image

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Information and Communication Technologies for Agriculture—Theme III: Decision Dionysis D. Bochtis, Claus Grøn Sørensen, Spyros Fountas, Vasileios Moysiadis, Panos M. Pardalos, 2022-04-27 This volume is the third III of four under the main themes of Digitizing Agriculture and Information and Communication Technologies ICT The four volumes cover rapidly developing processes including Sensors I Data II Decision III and Actions IV Volumes are related to digital transformation within agricultural production and provision systems and in the context of Smart Farming Technology and Knowledge based Agriculture Content spans broadly from data mining and visualization to big data analytics and decision making alongside with the sustainability aspects stemming from the digital transformation of farming The four volumes comprise the outcome of the 12th EFITA Congress also incorporating chapters that originated from select presentations of the Congress The focus of this book III is on the transformation of collected information into valuable decisions and aims to shed light on how best to use digital technologies to reduce cost inputs and time toward becoming more efficient and transparent Fourteen chapters are grouped into 3 Sections The first section of is dedicated to decisions in the value chain of agricultural products The next section titled Primary Production elaborates on decision making for the improvement of processes taking place with the farm under the implementation of ICT The last section is devoted to the development of innovative decision applications that also consider the protection of the environment recognizing its importance in the preservation and considerate use of resources as well as the mitigation of adverse impacts that are related to agricultural production Planning and modeling the assessment of agricultural practices can provide farmers with valuable information prior to the execution of any task This book provides a valuable reference for them as well as for those directly involved with decision making in planning and assessment of agricultural production Specific advances covered in the volume Modelling and Simulation of ICT based agricultural systems Farm Management Information Systems FMIS Planning for unmanned aerial systems Agri robotics awareness and planning Smart livestock farming Sustainable strategic planning in agri production Food business information Computational Systems Biology Andres Kriete, Roland Eils, 2013-11-26 This comprehensively revised second systems edition of Computational Systems Biology discusses the experimental and theoretical foundations of the function of biological systems at the molecular cellular or organismal level over temporal and spatial scales as systems biology advances to provide clinical solutions to complex medical problems In particular the work focuses on the engineering of biological systems and network modeling Logical information flow aids understanding of basic building blocks of life through disease phenotypes Evolved principles gives insight into underlying organizational principles of biological organizations and systems processes governing functions such as adaptation or response patterns Coverage of technical tools and systems helps researchers to understand and resolve specific systems biology problems using advanced computation Multi scale modeling on disparate

scales aids researchers understanding of dependencies and constraints of spatio temporal relationships fundamental to biological organization and function Computational Methods in Systems Biology Alessandro Abate, Tatjana Petrov, Verena Wolf, 2020-10-01 This book constitutes the refereed proceedings of the 18th International Conference on Computational Methods in Systems Biology CMSB 2020 held in Konstanz Germany in September 2020 The 17 full papers and 5 tool papers were carefully reviewed and selected from 30 submissions In addition 3 abstracts of invited talks and 2 tutorials have been included in this volume Topics of interest include formalisms for modeling biological processes models and their biological applications frameworks for model verification validation analysis and simulation of biological systems high performance computational systems biology and parallel implementations model inference from experimental data model integration from biological databases multi scale modeling and analysis methods computational approaches for synthetic biology and case studies in systems and synthetic biology The conference was held virtually due to the COVID 19 pandemic **Systems** Biology Edda Klipp, Wolfram Liebermeister, Christoph Wierling, Axel Kowald, 2016-03-28 This advanced textbook is tailored for an introductory course in Systems Biology and is well suited for biologists as well as engineers and computer scientists It comes with student friendly reading lists and a companion website featuring a short exam prep version of the book and educational modeling programs The text is written in an easily accessible style and includes numerous worked examples and study questions in each chapter For this edition a section on medical systems biology has been included

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