

Current Topics in Microbiology and Immunology



Michael G. Katze *Editor*

Systems Biology

 Springer

Systems Biology Current Topics In Microbiology And Immunology

Karthik Raman



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An Introduction to Computational Systems Biology Karthik Raman,2021-05-30 This book delivers a comprehensive and insightful account of applying mathematical modelling approaches to very large biological systems and networks a fundamental aspect of computational systems biology The book covers key modelling paradigms in detail while at the same time retaining a simplicity that will appeal to those from less quantitative fields Key Features A hands on approach to modelling Covers a broad spectrum of modelling from static networks to dynamic models and constraint based models Thoughtful exercises to test and enable understanding of concepts State of the art chapters on exciting new developments like community modelling and biological circuit design Emphasis on coding and software tools for systems biology Companion website featuring lecture videos figure slides codes supplementary exercises further reading and appendices <https://ramanlab.github.io/SysBioBook> An Introduction to Computational Systems Biology Systems Level Modelling of Cellular Networks is highly multi disciplinary and will appeal to biologists engineers computer scientists mathematicians and others

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Systems Biology Michael G. Katze,2013-01-01 First systems biology is an inter disciplinary approach requiring the combined talents of biologists mathematicians and computer scientists Second systems biology is holistic with the goal of obtaining a comprehensive understanding of the workings of biological systems This is achieved through the acquisition of massive amounts of data by high throughput technologies oligonucleotide microarrays mass spectrometry and next generation sequencing and the analysis of this data through sophisticated mathematical algorithms It is perhaps the use of mathematics to integrate abundant and diverse types of data and to generate models of interconnected molecular networks that best characterizes systems biology *Periodontal Medicine and Systems Biology* Brian Henderson,Michael Curtis,Robert Seymour,Nikolaos Donos,2009-07-07 This book provides a systems based approach to periodontology and offers a scientific roadmap of the interactions which can lead to periodontal disease The content is divided into five sections The first introduces the reader to the concept of systems theory and its mathematical foundation The second section provides the reader with a current view of periodontal medicine including the microbiology

molecular genetics relationship to systemic disease and current and future therapies Periodontitis is caused by members of the oral microbiota and the third section provides the reader with various views of the relationship of the microbiota to the host The fourth section moves from the bacterium to the host and its immune responses to altered host bacteria interactions The final section deals specifically with bone destruction in periodontitis and brings the reader up to date with the current view of the control network that exists between mesenchymal cells such as osteoblasts immune cells and osteoclast precursor cells that controls bone remodelling in health and disease

Omics Approaches in Biofilm Research Siddhardha Busi, Subhaswaraj Pattnaik, Ram Prasad, 2025-08-08 The increased incidence of microorganisms selective pressure to traditional antibiotics has led to the emergence of multi drug resistance MDR phenomena and has become a global health issue with a catastrophic influence on millions of lives as well as the global economy The inherent tendency of pathogenic microorganisms to infer MDR could be attributed to their ability to form recalcitrant biofilm matrices The biofilm matrix not only advocates chronic nosocomial infections but also critically provides protection against environmental stress including antibiotic therapies Biofilm mediated MDR has posed a serious challenge to human well being Henceforth it is important to understand the pathophysiology of biofilms and the concomitant development of diagnostic therapeutic modalities to counteract biofilm mediated chronic infections The lack of understanding on biofilm biology has a critical negative influence on diagnostic and therapeutic efforts Therefore it is imperative to discover the right course of action to understand biofilm mechanics The advent of Omics based approaches has provided a holistic realization to understand biofilm ecology with special reference to the pathophysiological interactions of antibiotic resistant genes protein protein interactions and response based interactions with therapeutic agents upon infection The inherent ability of several Omics based approaches has provided a comprehensive understanding of biofilm dynamics at various levels of organization such as genes mRNA proteins and their regulation Omics based tools such as metagenomics transcriptomics proteomics metabolomics etc have provided a new horizon to understand and tackle the biofilm mediated antibiotic resistance The integrated approach to consider multi Omics tools e g genomics transcriptomics proteomics lipidomics metabolomics etc has further improved our understanding of the mechanisms associated with biofilm resistome profile The applications of transcriptomics proteomics and metabolomics profiles of biofilm matrices could provide new dimensions in relation to the characteristic properties of different ARGs their relative expression profiles and their metabolic intervention in biofilm mechanics Also advanced integrated Phenomics Lipidomics and Culturomics approaches could provide novel avenues to understand the diverse range of biofilm phenotypes their macromolecular reorganization profiles and molecular tools for identification of microbial species in the complex biofilm microenvironment Based on the advancement in omics based tools *Omics Approaches in Biofilm Research Perspectives and Applications* integrates the current knowledge of biofilm microenvironment and innovative strategies to address biofilm mediated drug resistance This work provides a comprehensive platform to enhance our

knowledge diagnosis and strategies to mitigate biofilms and associated diseases

Mathematical Modeling of the Immune System in Homeostasis, Infection and Disease Gennady Bocharov, Burkhard Ludewig, Andreas Meyerhans, Vitaly Volpert, 2020-02-24 The immune system provides the host organism with defense mechanisms against invading pathogens and tumor development and it plays an active role in tissue and organ regeneration Deviations from the normal physiological functioning of the immune system can lead to the development of diseases with various pathologies including autoimmune diseases and cancer Modern research in immunology is characterized by an unprecedented level of detail that has progressed towards viewing the immune system as numerous components that function together as a whole network Currently we are facing significant difficulties in analyzing the data being generated from high throughput technologies for understanding immune system dynamics and functions a problem known as the curse of dimensionality As the mainstream research in mathematical immunology is based on low resolution models a fundamental question is how complex the mathematical models should be To respond to this challenging issue we advocate a hypothesis driven approach to formulate and apply available mathematical modelling technologies for understanding the complexity of the immune system Moreover pure empirical analyses of immune system behavior and the system's response to external perturbations can only produce a static description of the individual components of the immune system and the interactions between them Shifting our view of the immune system from a static schematic perception to a dynamic multi level system is a daunting task It requires the development of appropriate mathematical methodologies for the holistic and quantitative analysis of multi level molecular and cellular networks Their coordinated behavior is dynamically controlled via distributed feedback and feedforward mechanisms which altogether orchestrate immune system functions The molecular regulatory loops inherent to the immune system that mediate cellular behaviors e g exhaustion suppression activation and tuning can be analyzed using mathematical categories such as multi stability switches ultra sensitivity distributed system graph dynamics or hierarchical control GB is supported by the Russian Science Foundation grant 18 11 00171 AM is also supported by grants from the Spanish Ministry of Economy Industry and Competitiveness and FEDER grant no SAF2016 75505 R the Mar a de Maeztu Programme for Units of Excellence in R D MDM 2014 0370 and the Russian Science Foundation grant 18 11 00171

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