

Cooperativity Theory In Biochemistry Steady State

Linear Aggregation Theory in Cell Biology Terrell L.

Hill.2012-12-06 During the past few decades we have witnessed an era of remarkable growth in the field of molecular biology. In 1950 very little was known of the chemical constitution of biological systems, the manner in which information was transmitted from one organism to another, or the extent to which the chemical basis of life is unified. The picture today is dramatically different. We have an almost bewildering variety of information detailing many different aspects of life at the molecular level. These great advances have brought with them some breathtaking insights into the molecular mechanisms used by nature for replicating, distributing, and modifying biological information. We have learned a great deal about the chemical and physical nature of the macromolecular nucleic acids and proteins, and the manner in which carbohydrates, lipids, and smaller molecules work together to provide the molecular setting of living systems. It might be said that these few decades have replaced a near vacuum of information with a very large surplus. It is in the context of this flood of information that this series of monographs on molecular biology has been organized. The idea is to bring together in one place, between the covers of one book, a concise assessment of the state of the subject in a well-defined field. This will enable the reader to get a sense of historical perspective-what is known about the field today-and a description of the frontiers of research where our knowledge is increasing steadily.

Cooperativity Theory in Biochemistry Terrell L. Hill.1985-01-01
Mechanisms of Cooperativity and Allosteric Regulation in

Proteins Max F. Perutz, Max Ferdinand Perutz. 1990-05-17
The Good, The Bad and The Ugly: Multiple Roles of Bacteria in Human Life Tatiana Venkova, Chew Chieng Yeo, Manuel Espinosa. 2018-10-17

Bacteria are among the earliest forms of life on Earth. Notwithstanding their small size and primitive origin, bacteria still have a tremendous impact on everyday human life. Over the centuries, research into bacteria have provided and enriched the fundamental biological knowledge due to their readily measured processes and effects on higher organisms. Although molecular genetics and microbiology were among the scientific fields that have mostly benefited from the discoveries made in bacteria, our current state of knowledge has gone beyond what anyone could have ever imagined. The present Research Topic aims to cover new and exciting broad aspects of the importance of bacteria to human life, both positive and negative influences. Regulation of bacterial gene expression, replication and segregation control mechanisms, cell to cell communication via quorum sensors, and the relatively recent finding of bacterial immunity via CRISPR, have led to the development of many, and very important new tools in biotechnology and the emerging field of molecular medicine. The battle against infectious diseases has also benefited from the genetic approaches that have been developed in the quest for finding new targets and novel drugs against pathogenic bacteria. At the next level, the human microbiome project has opened up new avenues in understanding the role of bacteria in human health and wellbeing. Finally, the relationship between bacterial infections and human cancers will also be covered, a subject that is still under verification through rigorous experimental approaches. Special emphasis will be given to the bacterial accessory genome, i.e the mobilome, as the primary cause of health-threatening antimicrobial resistance and the production of toxins and virulence factors. Taking into account the evolutionary importance of horizontal gene transfer and the additional

beneficial roles of certain bacterial mobile genetic elements, they help project best “the Good, the Bad and the Ugly” outline of this topic. At the time this eBook is about to be published, our Research Topic has registered nearly 55, 000 views.

Catalysis in Chemistry and Biochemistry Theory and

Experiment A. Pullman.2012-12-06 Proceedings of the Twelfth Jerusalem Symposium on Quantum Chemistry and Biochemistry held in Jerusalem, Israel, April 2-4, 1979

Biothermodynamics Michael L. Johnson,Jo M. Holt,Gary K.

Ackers.2011 The use of thermodynamics in biological research can be equated to an energy book-keeping system. While the structure and function of a molecule is important, it is equally important to know what drives the energy force. This volume presents sophisticated methods for estimating the thermodynamic parameters of specific protein-protein, protein-DNA and small molecule interactions. * Elucidates the relationships between structure and energetics and their applications to molecular design, aiding researchers in the design of medically important molecules * Provides a must-have methods volume that keeps MIE buyers and online subscribers up-to-date with the latest research * Offers step-by-step lab instructions, including necessary equipment, from a global research community
Molecular Biology .1990

Biochemistry of Metabolism David D. Davies.2014-05-10 The Biochemistry of Plants: A Comprehensive Treatise, Volume 11: Biochemistry of Metabolism provides information pertinent to the chemical and biochemical aspects of metabolism. This book discusses the control mechanisms of metabolism. Organized into nine chapters, this volume begins with an overview of the history of biochemistry and discusses the developments in the kinetics of regulatory enzymes. This text then examines a theory that explains how subunit interactions modulate the rate of conversion of a substrate into a product. Other chapters consider some relation between cell-wall elongation and cell-wall charge density

and explore the subcellular localization of the enzymes of glycolysis. This book discusses as well the regulation of glycolysis and the pentose phosphate pathway. The final chapter deals with the pathways of C1 metabolism that are of prime importance, as the synthesis of several cellular constituents depends directly or indirectly on folate metabolism. This book is a valuable resource for plant biochemists, neurobiochemists, molecular biologists, senior graduate students, and research workers.

Technical Book Review Index .1986

Introduction to Systems Biology Sangdun Choi.2008-05-17 This book provides an introductory text for undergraduate and graduate students who are interested in comprehensive biological systems. The authors offer a broad overview of the field using key examples and typical approaches to experimental design. The volume begins with an introduction to systems biology and then details experimental omics tools. Other sections introduce the reader to challenging computational approaches. The final sections provide ideas for theoretical and modeling optimization in systemic biological researches. The book is an indispensable resource, providing a first glimpse into the state-of-the-art in systems biology.

Thermal Physics Charles Kittel,Herbert Kroemer.1980-01-15 CONGRATULATIONS TO HERBERT KROEMER, 2000 NOBEL LAUREATE FOR PHYSICS For upper-division courses in thermodynamics or statistical mechanics, Kittel and Kroemer offers a modern approach to thermal physics that is based on the idea that all physical systems can be described in terms of their discrete quantum states, rather than drawing on 19th-century classical mechanics concepts.

Molecular Driving Forces Ken Dill,Sarina Bromberg.2010-10-21 Molecular Driving Forces, Second Edition E-book is an introductory statistical thermodynamics text that describes the principles and forces that drive chemical and biological processes. It demonstrates how the complex behaviors of

molecules can result from a few simple physical processes, and how simple models provide surprisingly accurate insights into the workings of the molecular world. Widely adopted in its First Edition, *Molecular Driving Forces* is regarded by teachers and students as an accessible textbook that illuminates underlying principles and concepts. The Second Edition includes two brand new chapters: (1) *Microscopic Dynamics* introduces single molecule experiments; and (2) *Molecular Machines* considers how nanoscale machines and engines work. The *Logic of Thermodynamics* has been expanded to its own chapter and now covers heat, work, processes, pathways, and cycles. New practical applications, examples, and end-of-chapter questions are integrated throughout the revised and updated text, exploring topics in biology, environmental and energy science, and nanotechnology. Written in a clear and reader-friendly style, the book provides an excellent introduction to the subject for novices while remaining a valuable resource for experts.

Chromatin Kensal E. van Holde. 2012-12-06 Contemporary views on the structure and function of chromatin are presented and the history of the development of these ideas as well as the nature of the nucleic acid and protein components of chromatin are reviewed. The structure of chromatin is studied at several levels, and its modes of transcription and replication are analyzed.

Chromatin provides researchers with a critical evaluation of current knowledge. It combines much information that has never before been assembled, and evaluates and interrelates it in a critical way. This has not been done before so that readers are not only provided with an overview, but with extensive references to the literature (there are about 2000 references in all).

Proteins Paulo Almeida. 2016-03-14 *Proteins: Concepts in Biochemistry* teaches the biochemical concepts underlying protein structure, evolution, stability, folding, and enzyme kinetics, and explains how interactions in macromolecular structures determine protein function. Intended for a one-

semester course in biochemistry or biophysical chemistry with a focus on proteins, this textbo

Cooperativity Theory in Biochemistry T.L. Hill.2013-03-07

During the past few decades we have witnessed an era of remarkable growth in the field of molecular biology. In 1950 very little was known of the chemical constitution of biological systems, the manner in which information was transmitted from one organism to another, or the extent to which the chemical basis of life is unified. The picture today is dramatically different. We have an almost bewildering variety of information detailing many different aspects of life at the molecular level. These great advances have brought with them some breath-taking insights into the molecular mechanisms used by nature for replicating, distributing and modifying biological information. We have learned a great deal about the chemical and physical nature of the macromolecular nucleic acids and proteins, and the manner in which carbohydrates, lipids and smaller molecules work together to provide the molecular setting of living systems. It might be said that these few decades have replaced a near vacuum of information with a very large surplus. It is in the context of this flood of information that this series of monographs on molecular biology has been organized. The idea is to bring together in one place, between the covers of one book, a concise assessment of the state of the subject in a well-defined field. This will enable the reader to get a sense of historical perspective-what is known about the field today-and a description of the frontiers of research where our knowledge is increasing steadily.

Host-Parasite Relationships and the Yersinia Model Akira

Wake,Herbert R. Morgan.2012-12-06 During the past few decades we have witnessed an era of remarkable growth in the field of molecular biology. In 1950 very little was known of the chemical constitution of biological systems, the manner in which information was transmitted from one organism to another, or the extent to which the chemical basis of life is unified. The picture

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Protein Compartmentalization Arnold W. Strauss, Irving Boime, Günther Kreil. 2012-12-06 During the past few decades we have witnessed an era of remarkable growth in the field of molecular biology. In 1950 very little was known of the chemical constitution of biological systems, the manner in which information was transmitted from one organism to another, or the extent to which the chemical basis of life is unified. The picture today is dramatically different. We have an almost bewildering variety of information detailing many different aspects of life at the molecular level. These great advances have brought with them some breath taking insights into the molecular mechanisms used by nature for replicating, distributing, and modifying biological information. We have learned a great deal about the chemical and physical nature of the macromolecular nucleic acids and proteins, and the manner in which carbohydrates, lipids, and

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Reviews in Computational Chemistry Kenny B.

Lipkowitz, Donald B. Boyd. 2009-09-22 THIS BOOK HAS SIX TUTORIALS AND REVIEWS WRITTEN BY INVITED EXPERTS. FIVE CHAPTERS TEACH TOPICS IN QUANTUM MECHANICS AND MOLECULAR SIMULATIONS. THE SIXTH CHAPTER EXPLAINS HOW PROGRAMS FOR CHEMICAL STRUCTURE DRAWING WORK. AN EDITORIAL DISCUSSES SOME OF THE MOST WELL-KNOWN PERSONAGES IN COMPUTATIONAL CHEMISTRY. FROM REVIEWS OF THE SERIES Anyone who is doing or intends to do computational research on molecular structure and design should seriously consider purchasing this book for his or her personal library.-JOURNAL OF

COMPUTATIONAL CHEMISTRY. These reviews are becoming regarded as the standard reference among both specialists and novices in the expanding field of computational chemistry. -

JOURNAL OF MOLECULAR GRAPHICS AND MODELLING. [This book is] written for newcomers learning about molecular modeling techniques as well as for seasoned professionals who need to acquire expertise in areas outside their own.-JOURNAL OF CHEMICAL INFORMATION AND COMPUTER SCIENCE.

Cytochromes c Geoffrey R. Moore, Graham W.

Pettigrew. 2012-12-06 Cytochromes c are haemoproteins which carry out electron transfer in a wide variety of biological systems, necessitating different kinds of cytochrome c to fulfill different biological roles. The evolutionary relationship between cytochromes c and their host organisms are described, as well as

their structural, spectroscopic and redox properties, including both electron-transfer rates and redox potentials. The treatment is aimed at the non-specialist so that both the techniques described and their application to cytochromes c can be understood. All classes of cytochrome c are dealt with to provide a comprehensive account of the field. No other text provides such a broad survey. Similar to the earlier volume *Cytochromes c: Biological Aspects* which deals with the classification, biosynthesis and biological role of cytochromes c, the present book is aimed at research workers and advanced students.

Cytochromes c Graham W. Pettigrew, Geoffrey R.

Moore.2012-12-06 Cytochrome c fulfills a central role in biological electron transport. This book draws together information from diverse disciplines in order to provide a common base for further research. The comprehensive treatment of this subject does not neglect to show the diversity of biological respirations and photosyntheses. But it also defines their unifying principles. This overview presents the evolutionary relatedness in bioenergetic systems. Such systems are discussed at the experimental level with emphasis on the interpretation of results and the methodological approaches used. No other text provides a broad survey of this central area of biology. Researchers on cytochrome c are presented with information on the impact and importance of other disciplines on their area of investigation. Advanced students gain a balanced account of biological electron transport and will be encouraged to seek new directions of research.

Peptides of Poisonous Amanita Mushrooms Theodor

Wieland.2012-12-06 During the past few decades we have witnessed an era of remarkable growth in the field of molecular biology. In 1950 very little was known of the chemical constitution of biological systems, the manner in which information was transmitted from one organism to another, or the extent to which the chemical basis of life is unified. The picture today is

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Molecular Biology of DNA Methylation Roger L.P. Adams, Roy H. Burdon. 2012-12-06 During the past few decades we have witnessed an era of remarkable growth in the field of molecular biology. In 1950 very little was known of the chemical constitution of biological systems, the manner in which information was transmitted from one organism to another, or the extent to which the chemical basis of life is unified. The picture today is dramatically different. We have an almost bewildering variety of information detailing many different aspects of life at the molecular level. These great advances have brought with them some breath-taking insights into the molecular mechanisms used by nature for replicating, distributing and modifying biological information. We have learned a great deal about the chemical and physical nature of the macromolecular nucleic acids and proteins, and the manner in which carbohydrates, lipids and

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Fibronectins Richard O. Hynes. 2012-12-06 Fibronectins comprise a class of high molecular weight glycoproteins present both in extracellular matrices and in soluble form in body fluids. Although they have been studied for about forty years, their real significance emerged only during the past decade. Intensive research has focused on their role in platelet function, cell migration, the cytoskeleton, reticuloendothelial function, and on alterations in fibronectin distribution during development and disease. Fibronectins have emerged as glycoproteins with a very interesting set of properties generally involving adhesion of cells to cells or to extracellular material. In more recent years, the complete sequences of several fibronectin molecules and their genes were determined, the relation between structure and function was understood and much has been learned about cell surface receptors for fibronectins and other adhesive ligands. Having been at the forefront of all these exciting developments, the author has synthesized the entire field and with all the latest information at hand for the first time given it a clear perspective.

Hemoglobin M. Weissbluth. 2012-12-06 Approximately one third of the mass of a mammalian red blood cell is hemoglobin. Its major function is to bind oxygen at the partial pressure prevailing in the lungs and to release it to the tissues where the partial pressure is lower. The process whereby hemoglobin performs this essential physiological role is characterized by a cooperative interaction among its constituent subunits. A great deal of research effort has been devoted to this interaction, going back at

least as far as the first decade of this century. Moreover, cooperativity in hemo globin is probably not unique; it may well be one instance of a general class of interactions that occur in biological molecules. Certain enzymes with a variety of regulatory and catalytic functions, for example, contain several sites which interact in a highly specific manner such that the affinity of a given site for the substrate is markedly influenced by the state of binding at the other sites. But whereas we know very little of the structure of most enzymes of this type, hemo globin is one of a very small number of biological molecules whose immensely intricate machinery has been revealed to us. We owe this insight to the group under the leadership of M. F. PERUTZ in Cambridge, England, whose research over a period of several decades culminated in a detailed description of the three dimensional structure.

Structure, Function, and Genetics of Ribosomes Boyd

Hardesty, Gisela Kramer. 2012-12-06 During the past few decades we have witnessed an era of remarkable growth in the field of molecular biology. In 1950 very little was known of the chemical constitution of biological systems, the manner in which information was transmitted from one organism to another, or the extent to which the chemical basis of life is unified. The picture today is dramatically different. We have an almost bewildering variety of information detailing many different aspects of life at the molecular level. These great advances have brought with them some breath-taking insights into the molecular mechanisms used by nature for replicating, distributing, and modifying biological information. We have learned a great deal about the chemical and physical nature of the macromolecular nucleic acids and proteins, and the manner in which carbohydrates, lipids, and smaller molecules work together to provide the molecular setting of living systems. It might be said that these few decades have replaced a near vacuum of information with a very large surplus. It is in the context of this flood of information that this series of

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Progress in Molecular and Subcellular Biology .2012-12-06

Cooperativity and Regulation in Biochemical Processes

Arieh Y. Ben-Naim.2013-03-09 This is the first book that attempts to study the origin of cooperativity in binding systems from the molecular point of view. The molecular approach provides a deeper insight into the mechanism of cooperativity and regulation, than the traditional phenomenological approach. This book uses the tools of statistical mechanics to present the molecular theory of cooperativity. Cooperativity is used in a variety of processes-such as loading and unloading of oxygen at relatively small pressure differences; maintaining an almost constant concentration of various compounds in living cells; and switching on and off the reading of genetic information. This book may be used as a textbook by graduate students in Chemistry, Biochemistry and Biophysics, and will also be of interest to researchers in theoretical biochemistry.

National Library of Medicine Current Catalog National Library of Medicine (U.S.).1988

Current Catalog National Library of Medicine (U.S.). First multi-year cumulation covers six years: 1965-70.

Computational Modeling of Genetic and Biochemical

Networks James M. Bower,Hamid Bolouri.2001 How new modeling techniques can be used to explore functionally relevant molecular and cellular relationships.

Quantitative Aspects of Allosteric Mechanisms A.

Levitzki.2013-11-11 The aim of this monograph is to summarize the essential features which characterize the behavior of regulatory systems. Firstly we discuss the laws which govern ligand binding in thermodynamic terms. The basic cooperative and allosteric phenomena are described in thermodynamic terms

without assuming any particular model. Then the molecular models developed by Monod, Wyman and Changeux and by Koshland, Nemethy and Filmer are presented in detail. Special emphasis has been given to the analysis of the Hill coefficient and its meaning both in thermodynamic terms and in terms of the two allosteric models: the concerted model of Monod, Wyman and Changeux and the sequential model of Koshland, Nemethy and Filmer. Special types of cooperativities are discussed in some detail namely, cooperativity stemming from ligand coupled protein association or dissociation, negative cooperativity and half-of-the-sites reactivity. A slightly extended space was devoted to the discussion of negative cooperativity and half of-the-sites reactivity, since the existence of these phenomena and their possible biological importance is less of a common knowledge than positive cooperativity. This monograph does not attempt to be a review of specific examples analyzed according to one model or another. Rather, an attempt is made to provide the reader with the quantitative tools to analyze any specific regulatory system. Last but not least, I would like to thank Prof. F. W. Dahlquist from the Institute of Molecular Biology, the University of Oregon (Eugene) and Prof. D. E. Koshland, Jr.

Environmental Modelling and Prediction Gongbing

Peng, Lance M. Leslie, Yaping Shao. 2001-11-06 In this book the authors consider the natural environment as an integrated system. The physical, chemical and biological processes that govern the behaviour of the environmental system can thus be understood through mathematical modelling, and their evolution can be studied by means of numerical simulation. The book contains a summary of various efficient approaches in atmospheric prediction, such as numerical weather prediction and statistical forecast of climate change, as well as other successful methods in land surface modelling. The authors explore new theories and methods in environment prediction such as systems analysis and information theory. Attention is given to

new achievements in remote sensing tele-metering and geographic information systems.

Statistical Thermodynamics for Chemists and Biochemists Arieh Y. Ben-Naim.2013-11-11 This book was planned and written with one central goal in mind: to demonstrate that statistical thermodynamics can be used successfully by a broad group of scientists, ranging from chemists through biochemists to biologists, who are not and do not intend to become specialists in statistical thermodynamics. The book is addressed mainly to graduate students and research scientists interested in designing experiments the results of which may be interpreted at the molecular level, or in interpreting such experimental results. It is not addressed to those who intend to practice statistical thermodynamics per se. With this goal in mind, I have expended a great deal of effort to make the book clear, readable, and, I hope, enjoyable. This does not necessarily mean that the book as a whole is easy to read. The first four chapters are very detailed. The last four become progressively more difficult to read, for several reasons. First, presuming that the reader has already acquired familiarity with the methods and arguments presented in the first part, I felt that similar arguments could be skipped later on, leaving the details to be filled in by the reader. Second, the systems themselves become progressively more complicated as we proceed toward the last chapter.

Proceedings of the National Academy of Sciences of the United States of America .2001

Holistic Darwinism Peter Corning.2010-08-15 In recent years, evolutionary theorists have come to recognize that the reductionist, individualist, gene-centered approach to evolution cannot sufficiently account for the emergence of complex biological systems over time. Peter A. Corning has been at the forefront of a new generation of complexity theorists who have been working to reshape the foundations of evolutionary theory. Well known for his Synergism Hypothesis—a theory of complexity

in evolution that assigns a key causal role to various forms of functional synergy—Corning puts this theory into a much broader framework in Holistic Darwinism, addressing many of the issues and concepts associated with the evolution of complex systems. Corning's paradigm embraces and integrates many related theoretical developments of recent years, from multilevel selection theory to niche construction theory, gene-culture coevolution theory, and theories of self-organization. Offering new approaches to thermodynamics, information theory, and economic analysis, Corning suggests how all of these domains can be brought firmly within what he characterizes as a post-neo-Darwinian evolutionary synthesis.

Peptide-Lipid Interactions Sidney A. Simon, Thomas J.

McIntosh. 2002-03-26 This volume contains a comprehensive overview of peptide-lipid interactions by leading researchers. The first part covers theoretical concepts, experimental considerations, and thermodynamics. The second part presents new results obtained through site-directed EPR, electron microscopy, NMR, isothermal calorimetry, and fluorescence quenching. The final part covers problems of biological interest, including signal transduction, membrane transport, fusion, and adhesion. Key Features * world-renowned experts * state-of-the-art experimental methods * monolayers, bilayers, biological membranes * theoretical aspects and computer simulations * rafts * synaptic transmission * membrane fusion * signal transduction

Biothermodynamics .2011-02-16 The use of thermodynamics in biological research can be equated to an energy book-keeping system. While the structure and function of a molecule is important, it is equally important to know what drives the energy force. This volume presents sophisticated methods for estimating the thermodynamic parameters of specific protein-protein, protein-DNA and small molecule interactions. * Elucidates the relationships between structure and energetics and their applications to molecular design, aiding researchers in the design

of medically important molecules * Provides a must-have methods volume that keeps MIE buyers and online subscribers up-to-date with the latest research * Offers step-by-step lab instructions, including necessary equipment, from a global research community

Drug-Acceptor Interactions Niels Bindslev.2017-02-10 Drug-Acceptor Interactions: Modeling theoretical tools to test and evaluate experimental equilibrium effects suggests novel theoretical tools to test and evaluate drug interactions seen with combinatorial drug therapy. The book provides an in-depth, yet controversial, exploration of existing tools for analysis of dose-response studies at equilibrium or steady state. The book is recommended reading for post-graduate students and researchers engaged in the study of systems biology, networks, and the pharmacodynamics of natural or industrial drugs, as well as for medical clinicians interested in drug application and combinatorial drug therapy. Even people without mathematical skills will be able to follow the pros and cons of reaction schemes and their related distribution equations. Chapter 9 is a hands-on guide for software to plot, fit and analyze one's own data.

Reversible Ligand Binding Andrea Bellelli,Jannette Carey.2018-01-09 Presents the physical background of ligand binding and instructs on how experiments should be designed and analyzed Reversible Ligand Binding: Theory and Experiment discusses the physical background of protein-ligand interactions—providing a comprehensive view of the various biochemical considerations that govern reversible, as well as irreversible, ligand binding. Special consideration is devoted to enzymology, a field usually treated separately from ligand binding, but actually governed by identical thermodynamic relationships. Attention is given to the design of the experiment, which aids in showing clear evidence of biochemical features that may otherwise escape notice. Classical experiments are reviewed in order to further highlight the importance of the design of the

experiment. Overall, the book supplies students with the understanding that is necessary for interpreting ligand binding experiments, formulating plausible reaction schemes, and analyzing the data according to the chosen model(s). Topics covered include: theory of ligand binding to monomeric proteins; practical considerations and commonly encountered problems; oligomeric proteins with multiple binding sites; ligand binding kinetics; hemoglobin and its ligands; single-substrate enzymes and their inhibitors; two-substrate enzymes and their inhibitors; and rapid kinetic methods for studying enzyme reactions. Bridges theory of ligand binding and allostery with experiments Applies historical and physical insight to provide a clear understanding of ligand binding Written by a renowned author with long-standing research and teaching expertise in the area of ligand binding and allostery Based on FEBS Advanced Course lectures on the topic Reversible Ligand Binding: Theory and Experiment is an ideal text reference for students and scientists involved in biophysical chemistry, physical biochemistry, biophysics, molecular biology, protein engineering, drug design, pharmacology, physiology, biotechnology, and bioengineering.

A Primer in Mathematical Models in Biology Lee A. Segel, Leah Edelstein-Keshet. 2013-01-01 This textbook introduces differential equations, biological applications, and simulations and emphasizes molecular events (biochemistry and enzyme kinetics), excitable systems (neural signals), and small protein and genetic circuits. A Primer on Mathematical Models in Biology will appeal to readers because it grew out of a course that the popular and highly respected applied mathematician Lee Segel taught at the Weizmann Institute and it represents his unique perspective; combines clear and useful mathematical methods with applications that illustrate the power of such tools; and includes many exercises in reasoning, modeling, and simulations.

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