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Molecular Description Of Biological Membrane Components By Computer Aided Conformational Analysis

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Handbook of Nonmedical Applications of Liposomes Danilo D. Lasic, Yechezkel Barenholz, 2023-01-27 First published in 1996 liposomes have become an important model in fundamental biomembrane research including biophysical biochemical and cell biological studies of membranes and cell function They are thoroughly studied in several applications such as drug delivery systems in medical applications and as controlled release systems microencapsulating media signal carriers support matrices and solubilizers in other applications While medical applications have been extensively reviewed in recent literature there is a need for easily accessible information on applications for liposomes beyond pharmacology and medicine The Handbook of Nonmedical Applications of Liposomes fills this void This unique new handbook series presents recent developments in the use of liposomes in many scientific disciplines from studies on the origin of life protein function and vesicle shapes to applications in cosmetics diagnostics ecology bioreclamation and the food industry In these volumes

many of the top experts contribute extensive reviews of their work Biological Membranes Kenneth M. Merz, Benoit Roux, 2012-12-06 The interface between a living cell and the surrounding world plays a critical role in numerous complex biological processes Sperm egg fusion virus cell fusion exocytosis endocytosis and ion permeation are a few examples of processes involving membranes In recent years powerful tools such as X ray crystal lography electron microscopy nuclear magnetic resonance and infra red and Raman spectroscopy have been developed to characterize the structure and dy namics of biomembranes Despite this progress many of the factors responsible for the function of biomembranes are still not well understood The membrane is a very complicated supramolecular liquid crystalline structure that is largely composed of lipids forming a bilayer to which proteins and other biomolecules are anchored Often the lipid bilayer environment is pictured as a hydropho bic structureless slab providing a thermodynamic driving force to partition the amino acids of a membrane protein according to their solubility However much of the molecular complexity of the phospholipid bilayer environment is ignored in such a simplified view It is likely that the atomic details of the polar head group region and the transition from the bulk water to the hydrophobic core of the membrane are important An understanding of the factors responsible for the function of biomembranes thus requires a better characterization at the molecular level of how proteins interact with lipid molecules of how lipids affect protein structure and of how lipid molecules might regulate protein function Molecular and Cellular Mechanisms of H+ Transport Barry H. Hirst, 2013-06-29 Reviewed here is the current knowledge of proton transport mechanisms in mammals The emphasis is on gastric acid secretion and the role of the H K ATPase but molecular and cellular information on other P V and F type H ATPases in bone kidney plants and yeast as well as other cation ATPases are included for important comparisons The role of proton anion antiports symports and channels in proton transport is discussed Further attention is given to the regulation of proton transport mechanisms and cellular mechanisms to resist damage from highly acidic environments Physical Chemistry of Biological Interfaces Adam Baszkin, Willem Norde, 1999-11-22 An introduction to the most important fundamental concepts of physicochemical interface science and a description of experimental techniques and applications of surface science in relation to biological systems It explores artificial assemblies of lipids proteins and polysaccharides that perform novel functions that living systems cannot duplicat Thermodynamics of Membrane Receptors and Channels Meyer B. Jackson, 1992-11-18 Thermodynamics of Membrane Receptors and Channels synthesizes a wealth of new information regarding the biophysics of membrane proteins New insights provided by molecular genetics single channel recording and high resolution structural techniques are discussed from a conceptual perspective Basic theoretical topics are introduced developed and then extensively illustrated with recent results from the literature or data from the authors own laboratories Theoretical and experimental information is incorporated into in depth discussions of ion permeation mechanisms ion channel and receptor conformational changes aggregate activity of complexes of lipids and proteins and how coupling is achieved between different energy modes in the many transduction systems residing in

biomembranes Thermodynamics of Membrane Receptors and Channels will be valuable both as a learning aid and a reference for biophysicists neuroscientists cell biologists physiologists and other researchers investigating any aspects of biomembranes Modelling the Dynamics of Biological Systems Erik Mosekilde, Ole G. Mouritsen, 2012-12-06 The development of a proper description of the living world today stands as one of the most significant challenges to physics A variety of new experimental techniques in molecular biology microbiol ogy physiology and other fields of biological research constantly expand our knowledge and enable us to make increasingly more detailed functional and structural descriptions Over the past decades the amount and complexity of available information have multiplied dramatically while at the same time our basic understanding of the nature of regulation behavior morphogenesis and evolution in the living world has made only modest progress A key obstacle is clearly the proper handling of the available data This requires a stronger emphasis on mathematical modeling through which the consistency of the adopted explanations can be checked and general principles may be extracted As an even more serious problem however it appears that the proper physical concepts for the development of a theoretically oriented biology have not hitherto been available Classical mechanics and equilibrium thermody namics for instance are inappropriate and useless in some of the most essential biological contexts Fortunately there is now convincing evidence that the concepts and methods of the newly developed fields of nonlinear dynam ics and complex systems theory combined with irreversible thermodynamics and far from equilibrium statistical mechanics will enable us to move ahead with many of these problems **Permeability and Stability of Lipid Bilayers** E. Anibal Disalvo, Sidney A. Simon, 2017-12-14 This book presents a comprehensive and coherent picture of how molecules diffuse across a liquid that is on average only two molecules thick It begins by characterizing bilayers structurally using X ray diffraction and then mechanically by measuring elastic moduli and mechanisms of failure Emphasis is placed on the stability and mechanical properties of plant membranes that are subject to very large osmotic and thermal stresses Using this information the transport of molecules of increasing complexity across bilayers is analyzed The Molecular Biology of Chloroplasts and Mitochondria in **Chlamydomonas** J.-D. Rochaix, M. Goldschmidt-Clermont, Sabeeha Merchant, 2006-04-11 Provides a thorough overview of current research with the green alga Chlamydomonas on chloroplast and mitochondrial biogenesis and function with an emphasis on the assembly and structure function relationships of the constituents of the photosynthetic apparatus Contributions emphasize the multidisciplinary nature of current research in photosynthesis combining molecular genetics biochemical biophysical and physiological approaches The 36 articles address topics including nuclear genome organization RNA stability and processing splicing translation protein targeting in the chloroplast photosystems pigments glycerolipids the ATP synthase and ferrodoxin and thioredoxin Further contributions address new measurements methods for photosynthetic activity in vivo starch biosynthesis the responses of Chlamydomonas to various stress conditions nitrogen assimilation and mitochondrial genetics Annotation copyrighted by Book News Inc Portland OR Advances in

Computational Biology H.O. Villar, 1996-05-31 The second volume in a series which aims to focus on advances in computational biology This volume discusses such topics as statistical analysis of protein sequences progress in large scale sequence analysis and the architecture of loops in proteins Thermal Biophysics of Membranes Thomas Heimburg, 2008-02-08 An overview of recent experimental and theoretical developments in the field of the physics of membranes including new insights from the past decade The author uses classical thermal physics and physical chemistry to explain our current understanding of the membrane He looks at domain and raft formation and discusses it in the context of thermal fluctuations that express themselves in heat capacity and elastic constants Further topics are lipid protein interactions protein binding and the effect of sterols and anesthetics Many seemingly unrelated properties of membranes are shown to be intimately intertwined leading for instance to a coupling between membrane state domain formation and vesicular shape This also applies to non equilibrium phenomena like the propagation of density pulses during nerve activity Also included is a discussion of the application of computer simulations on membranes For both students and researchers of biophysics biochemistry physical chemistry and soft matter physics The Amphipathic Helix Richard M. Epand, 2024-12-06 The Amphipathic Helix is a comprehensive volume discussing amphipathic helices in systems as diverse as serum lipoproteins lung surfactant cytotoxic peptides ion channels mitochondrial targeting peptide hormones G proteins T cell recognition DNA binding proteins and antifreeze proteins The book also includes general introductory material that defines amphipathic helices discusses methods to identify amphipathic helical segments from the amino acid sequence of a protein illustrates how amphipathic helices can be used in the de novo design of peptide and protein structures and describes how these helices stabilize protein structures. There is also a section on techniques to determine helix orientation in a membrane environment using polarized attenuated total reflection infrared spectroscopy or solid state NMR spectroscopy Recent developments on all these topics have been discussed by leading experts in this reference for researchers and students in Biological and Biomedical Infrared Spectroscopy Andreas Barth, Parvez I. biochemistry biophysics and pharmacology Haris, 2009 Although infrared spectroscopy has been applied with success to the study of important biological and biomedical processes for many years key advances in this vibrant technique have led to its increasing use ranging from characterization of individual macromolecules DNA RNA lipids proteins to human tissues cells and their components Infrared spectroscopy thus has a significant role to play in the analysis of the vast number of genes and proteins being identified by the various genomic sequencing projects Whilst this book gives an overview of the field it highlights more recent developments such as the use of bright synchrotron radiation for recording infrared spectra the development of two dimensional infrared spectroscopy and the ability to record infrared spectra at ultra fast speeds The main focus is on the mid infrared region since the great majority of studies are carried out in this region but there is increasing use of the near infrared for biomedical applications and a chapter is devoted to this part of the spectrum Biological and Biomedical Infrared Spectroscopy is

intended for use both by research scientists already active in the use of biological infrared spectroscopy and for those coming new to the technique Graduate students will also find it useful as an introduction to the technique **Biotechnology in Animal Husbandry** R. Renaville, A. Burny, 2006-04-11 Animal biotechnology is a broad umbrella encompassing the polarities of fundamental and applied research including molecular modelling molecular and quantitative genetics gene manipulation development of diagnostics and vaccines and manipulation of tissue or digestion metabolism by growth promoters Although animal biotechnology in the broadest sense is not new what is new is the level of complexity and precision involved in scientists current ability to manipulate living organisms This new book sets out to show that the important ideas in animal biotechnology are exciting and relevant to everyday experience It represents an important update of the literature for research workers lecturers and advisers in animal science but is also a core text for advanced undergraduate courses in animal science and biotechnology It will be an essential acquisition for librarians in agriculture and veterinary science

The Molecular Dynamics of Liquid Crystals G.R. Luckhurst, C.A. Veracini, 2012-12-06 Liquid crystalline phases are now known to be formed by an ever growing range of quite diverse materials these include those of low molecular weight as well as the novel liquid crystalline polymers such phases can also be induced by the addition of a solvent to amphiphilic systems leading to lyotropic liquid crystals Irrespective of the structure of the constituent molecules these numerous liquid cl Stailine phases are characterised by their long range orientational order In addition certain phases exhibit elements of long range positional order Our understanding both experimental and theoretical at the molecular level of the static behaviour of these fascinating and important materials is now well advanced In contrast the influence of the long range order both orientational and positional on the molecular dynamics in liquid Cl Stais is less well understood In an attempt to address this situation a NATO Advanced Study Institute devoted to liquid ctystal dynamics was held at n Ciocco Barga Italy in September 1989 This brought together experimentalists and theoreticians concerned with the various dynamical processes occurring in all liquid crystals. The skills of the participants was impressively wide ranging they spanned the experimental techniques used in the study of molecular dynamics the nature of the systems investigated and the theoretical models employed to understand the results While much was learnt it was also recognised that much more needed to be done in order to advance our understanding of molecular dynamics in liquid Cl Stais **Protein-Lipid Interactions** Anthony Watts, 1993-10-08 Protein lipid interactions as a field of study is now a mature area and this volume of NewComprehensive Biochemistry has been published with two objectives in mind Firstly to look to thefuture and try to envisage how the subject may develop in the near to medium future Secondly to present contrasting or complementary views on the same system For example the acetylcholine receptor is discussed from a predominantly structural aspect by Barrantes and from the kinetic standpoint by Rankin et al The volume not only gives an update on specific aspects of the field but also shows the way in whichthe phenomenon of protein lipid interactions is now seemingly infiltrating many areas of biomembraneresearch from recombinant DNA studies protein

in Chemical Physics M.P. Allen,D.J. Tildesley,2012-12-06 Computer Simulation in Chemical Physics contains the proceedings of a NATO Advanced Study Institute held at CORISA Alghero Sardinia in September 1992 In the five years that have elapsed since the field was last summarized there have been a number of remarkable advances which have significantly expanded the scope of the methods Good examples are the Car Parrinello method which allows the study of materials with itinerant electrons the Gibbs technique for the direct simulation of liquid vapor phase equilibria the transfer of scaling concepts from simulations of spin models to more complex systems and the development of the configurational biased Monte Carlo methods for studying dense polymers The field has also been stimulated by an enormous increase in available computing power and the provision of new software All these exciting developments an more are discussed in an accessible way here making the book indispensable reading for graduate students and research scientists in both academic and industrial settings

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