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Liposomes are cellular structures made up of lipid molecules. Important as a cellular model in the study of basic biology liposomes are also used in clinical applications such as drug delivery and virus studies. *Liposomes in Immunology *Liposomes in Diagnostics *Liposomes in Gene Delivery and Gene Therapy

Liposomes are cellular structures made up of lipid molecules, which are water insoluble organic molecules and the basis of biological membranes. Important as a cellular model in the study of basic biology, liposomes are also used in clinical applications such as drug delivery and virus studies. Liposomes Part F is a continuation of previous MIE Liposome volumes A through E. * One of the most highly respected publications in the field of biochemistry since 1955 * Frequently consulted and praised by researchers and reviewers alike * Truly an essential publication for anyone in any field of the life sciences

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The ability of polypeptides to form alternatively folded, polymeric structures such as amyloids and related aggregates is being increasingly recognized as a major new frontier in protein research. This new volume of Methods in Enzymology along with Part B (volume 412) on Amyloid, Prions and other Protein Aggregates continue in the tradition of the first volume (309) in containing detailed protocols and methodological insights, provided by leaders in the field, into the latest methods for investigating the structures, mechanisms of formation, and biological activities of this important class of protein assemblies. Presents detailed protocols Includes troubleshooting tips Provides coverage on structural biology, computational methods, and biology

The critically acclaimed laboratory standard, Methods in Enzymology, is one of the most highly respected publications in the field of biochemistry. Since 1955, each volume has been eagerly awaited, frequently consulted, and praised by researchers and reviewers alike. The series contains much material still relevant today - truly an essential publication for researchers in all fields of life sciences. Nuclear Magnetic Resonance of Biological Macromolecules, Part C is written with a "hands-on" perspective. That is, practical applications with critical evaluations of methodologies and experimental considerations needed to design, execute, and interpret NMR experiments pertinent to biological molecules. * One of the most highly respected publications in the field of biochemistry since 1955 * Frequently consulted, and praised by researchers and reviewers alike * Truly an essential publication for anyone in any field of the life sciences

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

In the past several years, there has been an explosion in the ability of biologists, molecular biologists and biochemists to collect vast amounts of data on their systems. Biothermodynamics, Part C presents sophisticated methods for estimating the thermodynamic parameters of specific protein-protein, protein-DNA and small molecule interactions. The use of thermodynamics in biological research is used as 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. These methods look to answer: What are the sources of energy that drive the function? Which of the pathways are of biological significance? As the base of macromolecular structures continues to expand through powerful techniques of molecular biology, such as X-ray crystal data and spectroscopy methods, the importance of tested and reliable methods for answering these questions will continue to expand as well. 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

The combination of faster, more advanced computers and more quantitatively oriented biomedical researchers has recently yielded new and more precise methods for the analysis of biomedical data. These better analyses have enhanced the conclusions that can be drawn from biomedical data, and they have changed the way that experiments are designed and performed. This volume, along with the 2 previous Computer Methods volumes for the Methods in Enzymology serial, aims to inform biomedical researchers about recent applications of modern data analysis and simulation methods as applied to biomedical research. Presents step-by-step computer methods and discusses the techniques in detail to enable their implementation in solving a wide range of problems Informs biomedical researchers of the modern data analysis methods that have developed alongside computer hardware Presents methods at the "nuts and bolts" level to identify and resolve a problem and analyze what the results mean

This volume, along with Part A and Part B, is dedicated to a description of the instruments, samples, protocols, and analyses that belong to cryo-EM. It emphasizes the relatedness of the ideas, instrumentation, and methods underlying all cryo-EM approaches, which allow practitioners to easily move between them. Within each section, the articles are ordered according to the most common symmetry of the sample to which their methods are applied. * Includes time-tested core methods and new innovations applicable to any researcher * Methods included are useful to both established researchers and newcomers to the field * Relevant background and reference information given for procedures can be used as a guide

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