

RSC Biomolecular Sciences

Edited by Luis Moroder and Johannes Buchner

Oxidative Folding of Peptides and Proteins



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Oxidative Folding Of Peptides And Proteins Rsc Rsc Biomolecular Sciences:

Oxidative Folding of Peptides and Proteins Luis Moroder, Johannes Buchner, 2008-12-10 The formation of disulfide bonds is probably the most influential modification of peptides and proteins An elaborate set of cellular machinery exists to catalyze and guide this process In recent years significant developments have been made in both our understanding of the in vivo situation and the in vitro manipulation of disulfide bonds This is the first monograph to provide a comprehensive overview of this exciting and rapidly developing area It offers in depth insights into the mechanisms of in vivo and in vitro oxidative folding of proteins as well as mono and multiple stranded peptides Procedures applied for laboratory and industrial purposes are also discussed by top experts in the field The book describes the enzymes involved in the correct oxidative folding of cysteine containing proteins in prokaryotes and eukaryotes It then goes on to discuss the mimicking of these enzymes for successful in vitro folding of proteins including synthetic replicates and to deal with important issues concerning cysteine rich peptides The ability of natural bioactive peptides to fold correctly and in high yields to form defined structural motifs using cysteine sequence patterns is still puzzling With this in mind synthetic procedures for establishing native cysteine frameworks are discussed using selected examples such as the potential of selenocysteines The biotechnological and pharmaceutical relevance of proteins peptides their variants and synthetic replicates is continuously increasing Consequently this book is invaluable for peptide and protein chemists involved in related research and production

Oxidative Folding of Peptides and Proteins Luis Moroder, Johannes Buchner (Prof.), 2009 With contributions from experts in the field this book provides a comprehensive overview of the oxidative folding of cysteine rich peptides

Natural Product Chemistry for Drug Discovery Antony D. Buss, Mark S. Butler, 2010 This text provides a comprehensive summary of where natural product chemistry is today in drug discovery It covers emerging technologies and case studies and is a source of up to date information on the topical subject of natural products

RNA Helicases Eckhard Jankowsky, 2010-07-01 RNA helicases and RNA helicase like proteins are the largest group of enzymes in eukaryotic RNA metabolism and although they are subject to intense ongoing research there is much confusion about function and classification of these enzymes Although these enzymes are essential for virtually all processes involving RNA there is no overview detailing structure function and or biological roles of these pivotal proteins This book provides the first comprehensive and systematic overview of biology mechanism and structure of RNA helicases and RNA helicase like enzymes Research into RNA helicases takes place in many different fields from cell and developmental biology to mechanistic enzymology and structural biology and this book integrates the knowledge of these diverse fields into one valuable resource It also provides an informative overview on the entire group of enzymes Individual chapters on each subfamily of RNA helicases and RNA helicase like proteins are written by experts in the respective fields All chapters are systematically integrated and the reader is guided by a didactic introductory chapter The main strengths of the book are the combination of systematics and details that will allow the reader to gain insight into

results from diverse fields while maintaining a view of the entire field It will be a key reference for academics advanced students researchers and professionals working in or joining this field

Recent Developments in Biomolecular NMR Marius Clore, Jennifer Potts, 2012 NMR spectroscopy is widely used in biomolecular science particularly for structure determination of proteins nucleic acids and carbohydrates Much of the innovation within NMR spectroscopy has been within the field of protein NMR spectroscopy an important technique in structural biology Filling a gap in the literature this book draws together experts in the field to discuss the real advances in NMR methods that have occurred or have an impact on the biomolecular field in the last few years The coverage includes recent developments in using NMR for determination of protein structures membrane proteins the dynamics of RNA and advances in NMR in drug discovery Edited by leading biological NMR spectroscopists the book is essential reference for researchers in industry and academia interested in or joining this bioanalytical field

RNA Polymerases as Molecular Motors Henri C. Buc, Terence Strick, 2009-04-16 This book focuses on two aspects of the transcription cycle the recognition of the promoter by the enzyme and the achievement of consecutive translocation steps during elongation of the RNA product

Innovations in Biomolecular Modeling and Simulations Tamar Schlick, 2012

Biophysical Approaches Determining Ligand Binding to Biomolecular Targets Alberto Podjarny, Annick P Dejaegere, Bruno Kieffer, 2011-04-01 The binding of small ligands to biological molecules is central to most aspects of biological function The past twenty years has seen the development of an increasing armoury of biophysical methods that not only detect such binding but also provide varying degrees of information about the kinetics thermodynamics and structural aspects of the process These methods have received increasing attention with the growth in more rational approaches to drug discovery and design This book reviews the latest advances in the application of biophysics to the study of ligand binding It provides a complete overview of current techniques to identify ligands characterise their binding sites and understand their binding mechanisms Particular emphasis is given to the combined use of different techniques and their relative strengths and weaknesses Consistency in the way each technique is described makes it easy for readers to select the most suitable protocol for their research The introduction explains why some techniques are more suitable than others and emphasizes the possible synergies between them The following chapters all written by a specialist in the particular technique focus on each method individually The book finishes by describing how several complimentary techniques can be used together for maximum effectiveness This book is suitable for biomolecular scientists at graduate or post doctoral level in academia and industry Biologists and chemists will also find it a useful introduction to the techniques available

Structural Virology Mavis Agbandje-McKenna, Robert McKenna, 2010-11-11 Over the last ten years much effort has been devoted to improving the biophysical techniques used in the study of viruses This has resulted in the visualization of these large macromolecular assemblages at atomic level thus providing the platform for functional interpretation and therapeutic design Structural Virology covers a wide range of topics and is split into three sections The

first discusses the vast biophysical methodologies used in structural virology including sample production and purification confocal microscopy mass spectrometry negative stain and cryo electron microscopy X ray crystallography and nuclear magnetic resonance spectroscopy The second discusses the role of virus capsid protein structures in determining the functional roles required for receptor recognition cellular entry capsid assembly genome packaging and mechanisms of host immune system evasion The last section discusses therapeutic strategies based on virus protein structures including the design of antiviral drugs and the development of viral capsids as vehicles for foreign gene delivery Each topic covered will begin with a review of the current literature followed by a more detailed discussion of experimental procedures a step in the viral life cycle or strategies for therapeutic development With contributions from experts in the field of structural biology and virology this exceptional monograph will appeal to biomedical scientists involved in basic and or applied research on viruses It also provides up to date reference material for students entering the field of structural virology as well as scientists already familiar with the area

DNA Conjugates and Sensors Keith R Fox, Tom Brown, 2012-11-30 Applications of nucleic acids have developed recently to provide solutions for biosensors diagnostic tools and as platforms for the assembly of complex structures These developments have been possible as their base sequence can be used to assemble precise structures following simple and predictable rules Self assembled DNA can then be amplified using polymerase chain reaction PCR and this ultimately enables the preparation of synthetic nucleic acids Their use as molecular tools or DNA conjugates has recently been enhanced by the addition of other groups including enzymes fluorophores and small molecules Written by leaders in the field this volume describes the preparation and application of these DNA conjugates Several have been used as sensors aptamers riboswitches and nanostructures based on the ability of nucleic acids to adopt specific structures in the presence of ligands whilst others link reporter groups such as proteins or fluorophores to RNA or DNA for detection single molecule studies and increasing the sensitivity of PCR The book is relevant to researchers in areas related to analytical chemistry chemical biology medicinal chemistry molecular pharmacology and structural and molecular biology

Molecular Simulations and Biomembranes Mark S. P. Sansom, Philip Charles Biggin, 2010 The need for information in the understanding of membrane systems has been caused by three things an increase in computer power methodological developments and the recent expansion in the number of researchers working on it worldwide However there has been no up to date book that covers the application of simulation methods to membrane systems directly and this book fills an important void in the market It provides a much needed update on the current methods and applications as well as highlighting recent advances in the way computer simulation can be applied to the field of membranes and membrane proteins The objectives are to show how simulation methods can provide an important contribution to the understanding of these systems The scope of the book is such that it covers simulation of membranes and membrane proteins but also covers the more recent methodological developments such as coarse grained molecular dynamics and multiscale approaches in systems biology Applications

embrace a range of biological processes including ion channel and transport proteins The book is wide ranging with broad coverage and a strong coupling to experimental results wherever possible including colour illustrations to highlight particular aspects of molecular structure With an internationally respected list of authors its publication is timely and it will prove indispensable to a large scientific readership *Quantum Tunnelling in Enzyme-catalysed Reactions* Rudolf Konrad Allemann, Nigel S. Scrutton, 2009 In recent years there has been an explosion in knowledge and research associated with the field of enzyme catalysis and H tunneling Rich in its breath and depth this introduction to modern theories and methods of study is suitable for experienced researchers those new to the subject Edited by two leading experts and bringing together the foremost practitioners in the field this up to date account of a rapidly developing field sits at the interface between biology chemistry and physics It covers computational kinetic and structural analysis of tunnelling and the synergy in combining these methods with a major focus on H tunneling reactions in enzyme systems The book starts with a brief overview of proton and electron transfer history by Nobel Laureate Rudolph A Marcus The reader is then guided through chapters covering almost every aspect of reactions in enzyme catalysis ranging from descriptions of the relevant quantum theory and quantum classical theoretical methodology to the description of experimental results The theoretical interpretation of these large systems includes both quantum mechanical and statistical mechanical computations as well as simple more approximate models Most of the chapters focus on enzymatic catalysis of hydride proton and H transfer an example of the latter being proton coupled electron transfer There is also a chapter on electron transfer in proteins This is timely since the theoretical framework developed fifty years ago for treating electron transfers has now been adapted to H transfers and electron transfers in proteins Accessible in style this book is suitable for a wide audience but will be particularly useful to advanced level undergraduates postgraduates and early postdoctoral workers

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