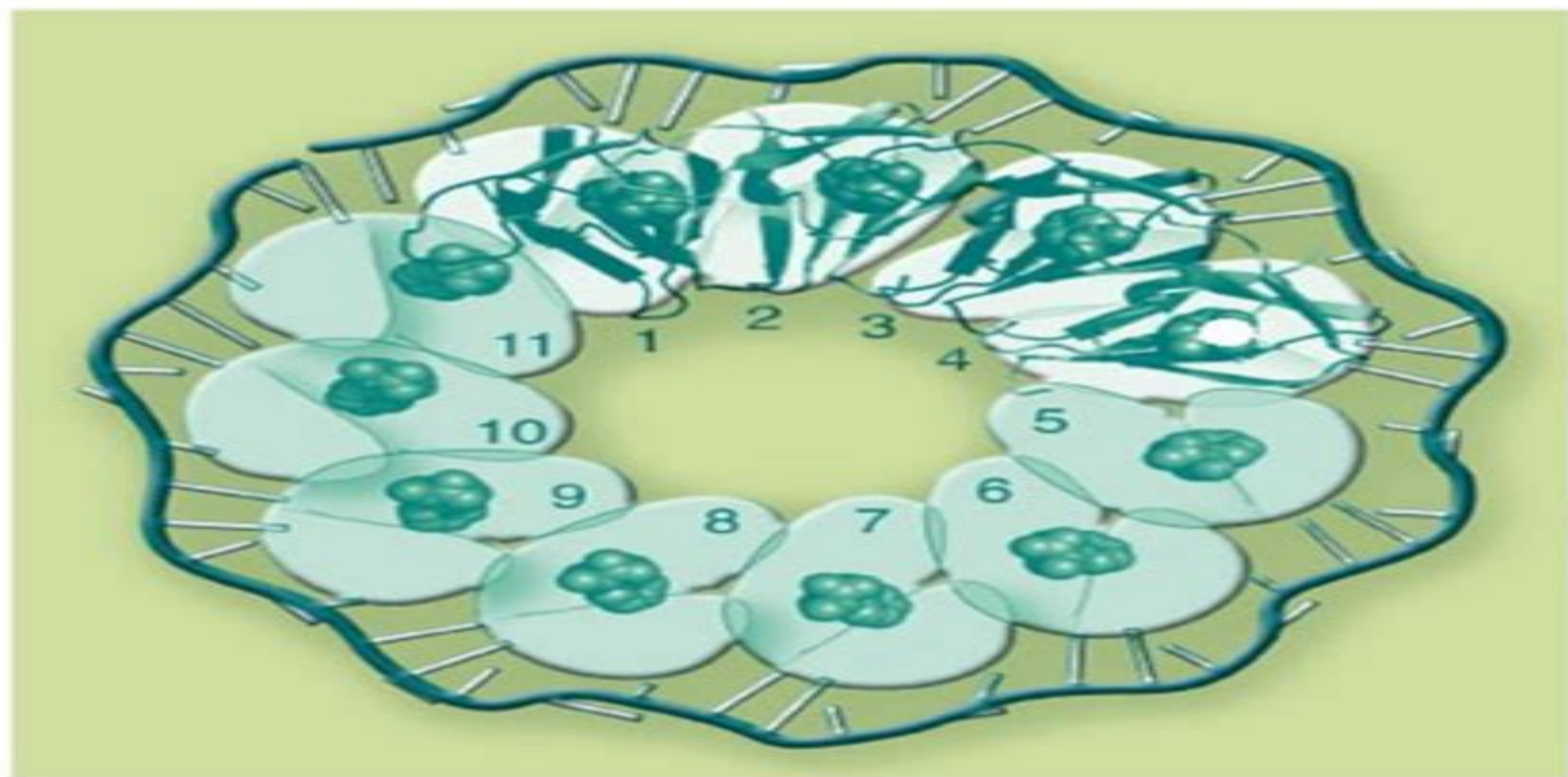


RSC Biomolecular Sciences

Edited by Phoebe A Rice and Carl C Correll

Protein–Nucleic Acid Interactions

Structural Biology



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Protein Nucleic Acid Interactions Structural Biology Rsc Biomolecular Sciences

Diane Schmidt



Protein Nucleic Acid Interactions Structural Biology Rsc Biomolecular Sciences:

Protein-Nucleic Acid Interactions Phoebe A. Rice, Carl C. Correll, 2008-05-22 The structural biology of protein nucleic acid interactions is in some ways a mature field and in others in its infancy High resolution structures of protein DNA complexes have been studied since the mid 1980s and a vast array of such structures has now been determined but surprising and novel structures still appear quite frequently High resolution structures of protein RNA complexes were relatively rare until the last decade Propelled by advances in technology as well as the realization of RNA's importance to biology the number of example structures has ballooned in recent years New insights are now being gained from comparative studies only recently made possible due to the size of the database as well as from careful biochemical and biophysical studies As a result of the explosion of research in this area it is no longer possible to write a comprehensive review Instead current review articles tend to focus on particular subtopics of interest This makes it difficult for newcomers to the field to attain a solid understanding of the basics One goal of this book is therefore to provide in depth discussions of the fundamental principles of protein nucleic acid interactions as well as to illustrate those fundamentals with up to date and fascinating examples for those who already possess some familiarity with the field The book also aims to bridge the gap between the DNA and the RNA views of nucleic acid protein recognition which are often treated as separate fields However this is a false dichotomy because protein DNA and protein RNA interactions share many general principles This book therefore includes relevant examples from both sides and frames discussions of the fundamentals in terms that are relevant to both The monograph approaches the study of protein nucleic acid interactions in two distinctive ways First DNA protein and RNA protein interactions are presented together Second the first half of the book develops the principles of protein nucleic acid recognition whereas the second half applies these to more specialized topics Both halves are illustrated with important real life examples The first half of the book develops fundamental principles necessary to understand function An introductory chapter by the editors reviews the basics of nucleic acid structure Jen Jacobsen and Jacobsen discuss how solvent interactions play an important role in recognition illustrated with extensive thermodynamic data on restriction enzymes Marmorstein and Hong introduce the zoology of the DNA binding domains found in transcription factors and describe the combinatorial recognition strategies used by many multiprotein eukaryotic complexes Two chapters discuss indirect readout of DNA sequence in detail Berman and Lawson explain the basic principles and illustrate them with in depth studies of CAP while in their chapter on DNA bending and compaction Johnson Stella and Heiss highlight the intrinsic connections between DNA bending and indirect readout Horvath lays out the fundamentals of protein recognition of single stranded DNA and single stranded RNA and describes how they apply in a detailed analysis of telomere end binding proteins Nucleic acids adopt more complex structures Lilley describes the conformational properties of helical junctions and how proteins recognize and cleave them Because RNA readily folds due to the stabilizing role of its 2 hydroxyl groups Li discusses how proteins

recognize different RNA folds which include duplex RNA With the fundamentals laid out discussion turns to more specialized examples taken from important aspects of nucleic acid metabolism Schroeder discusses how proteins chaperone RNA by rearranging its structure into a functional form Berger and Dong discuss how topoisomerases alter the topology of DNA and relieve the superhelical tension introduced by other processes such as replication and transcription Dyda and Hickman show how DNA transposases mediate genetic mobility and Van Duyne discusses how site specific recombinases cut and paste DNA Horton presents a comprehensive review of the structural families and chemical mechanisms of DNA nucleases whereas Li in her discussion of RNA protein recognition also covers RNA nucleases Lastly Ferr D Amar shows how proteins recognize and modify RNA transcripts at specific sites The book also emphasises the impact of structural biology on understanding how proteins interact with nucleic acids and it is intended for advanced students and established scientists wishing to broaden their horizons

Protein-Nucleic Acid Interactions Phoebe A. Rice, Carl C. Correll, 2008

Nucleic Acid-metal Ion Interactions Nicholas V. Hud, 2009 Provides a perspective on nucleic acid metal ion interactions with an emphasis on experimental biophysical studies which will prove indispensable to biophysicists and molecular biologists

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

Computational and Structural Approaches to Drug Discovery Robert M. Stroud, Janet Finer-Moore, 2008

- 1 Facing the Wall in Computationally Based Approaches to Drug Discovery Janet S Finer Moore and Jeff Blaney and Robert M Stroud
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Davis and Simon J Teague and Gerard J Kleywegt 5 Dealing with Bound Waters in a Site Do they Leave or Stay Donald Hamelberg and J Andrew McCammon 6 Knowledge Based Methods in Structure Based Design Marcel L Verdonk and Wijnand T M Mooij 7 Combating Drug Resistance Identifying Resilient Molecular Targets and Robust Drugs Celia A Schiffer 8 Docking Algorithms and Scoring Functions State of the Art and Current Limitations Gregory L Warren and Catherine E Peishoff and Martha S Head 9 Application of Docking Methods to Structure Based Drug Design Demetri T Moustakas 10 Strength in Flexibility Modeling Side Chain Conformational Change in Docking and Screening Leslie A Kuhn 11 Avoiding the Rigid Receptor Side Chain Rotamers Amy C Anderson 12 Computational Prediction of Aqueous Solubility Oral Bioavailability P450 Activity and hERG Channel Blockade David E Clark 13 Shadows on Screens Brian K Shoichet and Brian Y Feng and Kristin E D Coan 14 Iterative Docking Strategies for Virtual Ligand Screening Albert E Beuscher IV and Arthur J Olson 15 Challenges and Progresses in Calculations of Binding Free Energies What Does it Take to Quantify Electrostatic Contributions to Protein Ligand Interactions Mitsunori Kato and Sonja Braun Sand and Arie Warshel 16 Discovery and Extrapolation of Fragment Structures towards Drug Design Alessio Ciulli and Tom L Blundell and Chris Abell 17 A Link Means a Lot Disulfide Tethering in Structure Based Drug Design Jeanne A Hardy 18 The Impact of Protein Kinase Structures on Drug Discovery Chao Zhang and Sung Hou Kim

Innovations in Biomolecular Modeling and Simulations Tamar Schlick, 2012 The chemical and biological sciences face unprecedented opportunities in the 21st century A confluence of factors from parallel universes advances in experimental techniques in biomolecular structure determination progress in theoretical modeling and simulation for large biological systems and breakthroughs in computer technology has opened new avenues of opportunity as never before Now experimental data can be interpreted and further analysed by modeling and predictions from any approach can be tested and advanced through companion methodologies and technologies This two volume set describes innovations in biomolecular modeling and simulation in both the algorithmic and application fronts With contributions from experts in the field the books describe progress and innovation in areas including simulation algorithms for dynamics and enhanced configurational sampling force field development implicit solvation models coarse grained models quantum mechanical simulations protein folding DNA polymerase mechanisms nucleic acid complexes and simulations RNA structure analysis and design and other important topics in structural biology modeling The books are aimed at graduate students and experts in structural biology and chemistry and the emphasis is on reporting innovative new approaches rather than providing comprehensive reviews on each subject

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

Integration of structural biology data in lead drug discovery and optimization

Marco Nardini,Pietro Roversi,Gianluca Molla,2023-03-03

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

Using the Biological Literature

Diane Schmidt,2014-04-14 The biological sciences cover a broad array of literature types from younger fields like molecular biology with its reliance on recent journal articles genomic databases and protocol manuals to classic fields such as taxonomy with its scattered literature found in monographs and journals from the past three centuries Using the Biological Literature A Practical Guide Fourth Edition is an annotated guide to selected resources in the biological sciences presenting a wide ranging list of important sources This completely revised edition contains numerous new resources and descriptions of all entries including textbooks The guide emphasizes current materials in the English language and includes retrospective references for historical perspective and to provide access to the taxonomic literature It covers both print and electronic resources including monographs journals databases indexes and abstracting tools websites and associations providing users with listings of authoritative informational

resources of both classical and recently published works With chapters devoted to each of the main fields in the basic biological sciences this book offers a guide to the best and most up to date resources in biology It is appropriate for anyone interested in searching the biological literature from undergraduate students to faculty researchers and librarians The guide includes a supplementary website dedicated to keeping URLs of electronic and web based resources up to date a popular feature continued from the third edition

Reviewing **Protein Nucleic Acid Interactions Structural Biology Rsc Biomolecular Sciences**: Unlocking the Spellbinding Force of Linguistics

In a fast-paced world fueled by information and interconnectivity, the spellbinding force of linguistics has acquired newfound prominence. Its capacity to evoke emotions, stimulate contemplation, and stimulate metamorphosis is truly astonishing. Within the pages of "**Protein Nucleic Acid Interactions Structural Biology Rsc Biomolecular Sciences**," an enthralling opus penned by a highly acclaimed wordsmith, readers embark on an immersive expedition to unravel the intricate significance of language and its indelible imprint on our lives. Throughout this assessment, we shall delve to the book is central motifs, appraise its distinctive narrative style, and gauge its overarching influence on the minds of its readers.

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some of the most popular spin tricks include an alley oop half cab and hard way other snowboarding tricks the above list is just a glimpse into the many styles and types of tricks that you can perform on a snowboard other popular tricks that you ll see on the slopes include

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1 ollies once you gain perfection in performing the ollies you will become a pro from a novice in doing all the aerial snowboarding tips and tricks in the beginning you will use it for simple riding on the ground but later it will help you to get extra air

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