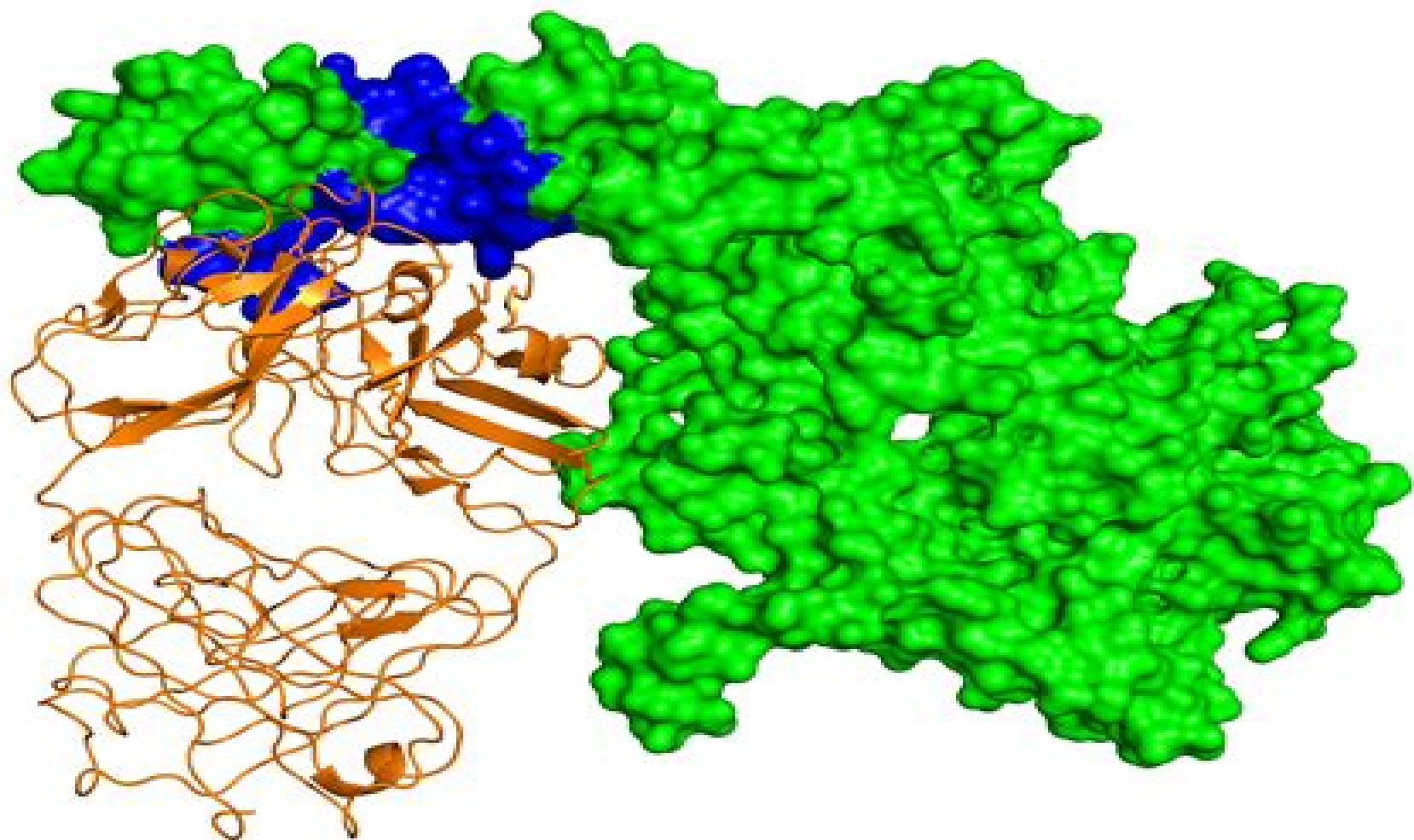




Small Molecule Protein Interaction

Michael D. Wendt



Small Molecule Protein Interaction:

Understanding Small Molecule-protein Interactions Raeanne L. Napoleon, 2012 Abstract The binding of small molecules to a protein is among the most important phenomena in the chemistry of life the activity and functionality of many proteins depend critically on binding small molecules A deep understanding of protein small molecule interactions and the interplay between ligation and function can give valuable insight into key systems of interest The complete characterization of any small molecule protein interaction requires quantification of many interactions and the pursuit of such information is the purpose of this body of work The discovery of binding regions on proteins or hot spots is an important step in drug development To this end a highly regarded and robust fragment based protocol has been developed for the detection of hot spots Firstly we use this protocol in conjunction with other computation techniques such as homology modeling to locate the allosteric binding site of L phenylalanine in Phenylalanine Hydroxylase Secondly computational fragment mapping was employed to locate the site of allostery for Ras an important signaling protein Lastly the identification of hot spots for many unligated protein targets is presented highlighting the importance of a reliable way to predict druggability computationally The second part of this dissertation shifts focus to the development of electrostatic models of small molecules It is widely believed that classical potentials can describe neither vibrational frequency shifts in condensed phases nor the response of vibrational frequencies to an applied electric field the vibrational Stark effect In this work an improved classical molecular electrostatic model for the CO ligand was developed to faithfully model these phenomena This model is found to predict the vibrational Stark effect and Fe CO binding energy with unprecedented accuracy for such a classical model As an extension of this work a geometrically dependent water potential was developed This work has shown that comparison of results obtained from current water models against experimentally determined proton momentum distributions is an invaluable benchmark

Targeting Protein-Protein Interactions by Small Molecules Chunquan Sheng, Gunda I. Georg, 2018-06-26 This book comprehensively reviews the state of the art strategies developed for protein protein interaction PPI inhibitors and highlights the success stories in new drug discovery and development Consisting of two parts with twelve chapters it demonstrates the design strategies and case studies of small molecule PPI inhibitors The first part discusses various discovery strategies for small molecule PPI inhibitors such as high throughput screening hot spot based design computational approaches and fragment based design The second part presents recent advances in small molecule inhibitors focusing on clinical candidates and new PPI targets This book has broad appeal and is of significant interest to the pharmaceutical science and medicinal chemistry communities *Small Molecule — Protein Interactions* Herbert Waldmann, Marcus Koppitz, 2013-03-09 Based on the international workshop on Small Molecule Protein Interactions held in Berlin April 24 26 2002 researchers from industry and academic laboratories describe novel and efficient ways selecting promising new drug targets and developing small molecule inhibitors against them The structure of the book corresponds to the different aspects of the drug discovery process

All chapters are written by leading experts in the field who present and discuss the most recent state of the art tools and techniques for the development of novel drugs. The value of the book lies in surveying and summarizing the approaches taken by different companies and institutions giving the reader a balanced view on the use of the latest techniques on the one hand and experience based assistance in selecting appropriate tools for their own work on the other hand. *Small-Molecule Inhibitors of Protein-Protein Interactions* Lyubomir Vassilev, David Fry, 2011-01-18 In this volume the editors have collected the knowledgeable insights of a number of leaders in this field researchers who have achieved success in addressing the difficult problem of inhibiting protein protein interactions. These researchers describe their unique approaches and share experiences results thoughts and opinions. The content of the articles is rich and in terms of scope ranges from generalized approaches to specific case studies. There are various focal points including methodologies and the molecules themselves. Ultimately there are numerous lessons to be taken away from this collection and the editors hope that this snapshot of the current state of the art in developing protein protein inhibitors not only pays tribute to the past successes but also generates excitement about the future potential of this field. **Methods for Detection of Small Molecule-protein Interactions**

Yan Guan, 2015 Detection of molecular interactions is critical for understanding many biological processes for detecting disease biomarkers and for screening drug candidates. Fluorescence based approach can be problematic especially when applied to the detection of small molecules. Various label free techniques such as surface plasmon resonance technique are sensitive to mass making it extremely challenging to detect small molecules. In this thesis novel detection methods for molecular interactions are described. First a simple detection paradigm based on reflectance interferometry is developed. This method is simple low cost and can be easily applied for protein array detection. Second a label free charge sensitive optical detection CSOD technique is developed for detecting of both large and small molecules. The technique is based on that most molecules relevant to biomedical research and applications are charged or partially charged. An optical fiber is dipped into the well of a microplate. It detects the surface charge of the fiber which does not decrease with the size mass of the molecule making it particularly attractive for studying small molecules. Third a method for mechanically amplification detection of molecular interactions MADMI is developed. It provides quantitative analysis of small molecules interaction with membrane proteins in intact cells. The interactions are monitored by detecting a mechanical deformation in the membrane induced by the molecular interactions. With this novel method small molecules and membrane proteins interaction in the intact cells can be detected. This new paradigm provides mechanical amplification of small interaction signals allowing us to measure the binding kinetics of both large and small molecules with membrane proteins and to analyze heterogeneous nature of the binding kinetics between different cells and different regions of a single cell. Last by tracking the cell membrane edge deformation binding caused downstream event granule secretory has been measured. This method focuses on the plasma membrane change when granules fuse with the cell. The fusion of granules increases the plasma membrane area and thus the

cell edge expands The expansion is localized at the vesicle release location Granule size was calculated based on measured edge expansion The membrane deformation due to the granule release is real time monitored by this method

Studies on Small Molecule-protein Interactions with a Note on the Use of Tracers in Transport Systems Erik Olof Arvidsson,1965

Protein Interactions as Targets in Drug Discovery Rossen Donev,2020-04-18 Protein Interactions as Targets in Drug Discovery Volume 121 is dedicated to the design of therapeutics both experimental and computational that target protein interactions Chapters in this new release include Trends in structure based drug design with protein targets From fragment to peptide protein interaction addressing the structural basis of binding using Supervised Molecular Dynamics SuMD Protein protein and protein ligand interactions identification of potential inhibitors through computational analysis Aromatic aromatic interactions in protein drug and protein protein interactions Role of protein protein interaction in allosteric drug design within the human methyltransferase and much more Integrates experimental and computational methods for studying protein interactions and their modulation by potential therapeutics Contains timely chapters written by well renown authorities in their field Covers information that is well supported by a number of high quality illustrations figures and tables Targets a very wide audience of specialists researchers and students

Design and Synthesis of Small-molecule Protein-protein Interaction Antagonists Xu Han,2014 Protein protein interactions play a crucial role in a wide range of biological processes Research on the design and synthesis of small molecules to modulate these proteinprotein interactions can lead to new targets and drugs to modulate their function In chapter one we discuss the design and synthesis of small molecules to probe a proteinprotein interaction in a voltage gated Ca²⁺ channel Virtual screening identified a compound BTT 3 that contained a 3,4 dihydro 3,4 pyrazole core This compound had modest biological activity when tested in a fluorescence polarization FP assay The synthetic route to BTT 3 consisted of six steps In addition analogs of BTT 3 were made for a structure activity study to establish the importance of a carboxylate moiety We also synthesized a biotinylated benzophenone photo affinity probe and linked it to BTT 3 to identify additional protein targets of the compound In Chapter two small molecule antagonists targeting uPA uPAR protein protein interaction are presented A total of 500 commercially available compounds were previously identified by virtual screening and tested by a FP assay Three classes of compounds were found with biological activity The first class of compounds contains pyrrolidone core structures represented by IPR 1110 the second class has a novel pyrrolo 3,4 c pyrazole ring system represented by xv IPR 1283 and the last series had compounds with a 1,2 disubstituted 1,2 dihydropyrrolo 3,4 b indol 3,4H one core structure represented by IPR 540 Each of these three compounds were synthesized and assessed by FP and ELISA assays A binding mode of IPR 1110 with uPA was subsequently proposed Based on this binding mode another 61 IPR 1110 derivatives were synthesized by us to illustrate the SAR activity Analogs of the other two series were also synthesized

Protein Surface Recognition Ernest Giralt,Mark Pecuh,Xavier Salvatella,2011-07-07 A new perspective on the design of molecular therapeutics is emerging This new

strategy emphasizes the rational complementation of functionality along extended patches of a protein surface with the aim of inhibiting protein protein interactions The successful development of compounds able to inhibit these interactions offers a unique chance to selectively intervene in a large number of key cellular processes related to human disease Protein Surface Recognition presents a detailed treatment of this strategy with topics including an extended survey of protein protein interactions that are key players in human disease and biology and the potential for therapeutics derived from this new perspective the fundamental physical issues that surround protein protein interactions that must be considered when designing ligands for protein surfaces examples of protein surface small molecule interactions including treatments of protein natural product interactions protein interface peptides and rational approaches to protein surface recognition from model to biological systems a survey of techniques that will be integral to the discovery of new small molecule protein surface binders from high throughput synthesis and screening techniques to in silico and in vitro methods for the discovery of novel protein ligands Protein Surface Recognition provides an intellectual tool kit for investigators in medicinal and bioorganic chemistry looking to exploit this emerging paradigm in drug discovery Protein Interactions Peter

Schuck,2007-03-20 When I was invited to edit this volume I wanted to take the opportunity to assemble reviews of different biophysical methodologies for protein interactions at a level sufficiently detailed to understand how complex systems can be studied There are several excellent introductory texts for biophysical methodologies many with hands on descriptions or embedded in general introductions to physical chemistry The goal of the present volume was to present state of the art reviews that do not necessarily enable the reader to carry out these techniques but to gain a deep understanding of the biophysical observables to stimulate creative thought on how the techniques may be applied to study a particular biological system and to foster collaboration and multidisciplinary work Reversible protein interactions involve noncovalent chemical bonds producing protein complexes with free energies not far from the order of magnitude of the thermal energy kT As a consequence they can be highly dynamic and may be controlled for example by protein expression levels and changes in the intracellular or microenvironment Reversible protein complexes may have sufficient stability to be purified for study but frequently their short lifetime essentially limits their existence to solutions of mixtures of the binding partners in which they remain populated through dissociation and reassociation processes To understand the function of such protein complexes it is important to study their structure and dynamics Biomolecules and Their Interactions Mr. Rohit Manglik,2024-04-06

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Cancer Therapeutics Kurt Deshayes Jeremy Murray Domagoj Vucic The Development of Small Molecule IAP Antagonists for the Treatment of Cancer John F Kadow David R Langley Nicholas A Meanwell Michael A Walker Kap Sun Yeung Richard Pracitto Protein Protein Interaction Targets to Inhibit HIV 1 Infection Nicholas A Meanwell David R Langley Inhibitors of Protein Protein Interactions in Paramyxovirus Fusion a Focus on Respiratory Syncytial Virus Andrew B Mahon Stephen E Miller Stephen T Joy Paramjit S Arora Rational Design Strategies for Developing Synthetic Inhibitors of Helical Protein Interfaces Michael D Wendt The Discovery of Navitoclax a Bcl 2 Family Inhibitor **Targeting Protein-Protein**

Interactions for Drug Discovery Jian Zhang, 2025-09-24 Up to date reference surveying the latest advances in the structural understanding of protein protein interactions and developments in drug discovery and therapeutics Targeting Protein Protein Interactions for Drug Discovery provides a systematic and comprehensive overview of protein protein interactions PPIs reviewing foundational concepts advanced methodologies and emerging therapeutic strategies reflecting the multidisciplinary nature of PPI research This book discusses computational methods for predicting PPI structures with a special emphasis on protein docking and deep learning based approaches diverse chemical scaffolds for PPI modulation including foldamers as inhibitors of aberrant PPIs and sulfonyl AApeptides as novel modulators and the development and application of stapled peptides as modulators of intracellular PPIs offering enhanced stability binding affinity and cellular permeability Readers will also find information on cyclic peptides focusing on their unique conformational stabilization and therapeutic potential across a range of diseases small molecule inhibitors targeting BCL family proteins revealing their potential in cancer therapy molecular glues as activators for PPIs categorized into degraders stabilizers and inhibitors based on their biological effects and the targeting of the APC Asef interaction for drug discovery in colorectal cancer therapy offering a case study of specificity and clinical relevance Targeting Protein Protein Interactions for Drug Discovery explores sample topics including Challenges and strategies of drug discovery targeting PPIs including high throughput screening and structure based drug design Fluorescence resonance energy transfer FRET technology a powerful tool for real time analysis of molecular interactions in live cells Utility of mass spectrometry MS for large scale mapping of PPI networks with high sensitivity and resolution Proximity ligation assays PLA for detecting PPIs in situ emphasizing spatial precision and adaptability for multiplexed detection Application of surface plasmon resonance SPR for characterizing PPI specificity affinity and kinetics Exploring both classical and novel approaches to PPI characterization and modulation Targeting Protein Protein Interactions for Drug Discovery offers a comprehensive reference for researchers aiming to unlock the therapeutic potential of PPIs along with educators and students engaged in the study of cellular mechanisms drug discovery and biotechnology

Protein - Protein Interaction Meike Werther, Harald Seitz, 2008-09-29 This book covers trends in modern biotechnology It treats all aspects of this interdisciplinary technology where knowledge methods and expertise are required from chemistry biochemistry microbiology genetics chemical engineering and computer science Protein-Protein Interactions Shahid

Mukhtar,2023-07-14 This detailed volume provides a comprehensive collection of classic and cutting edge methods and techniques in mapping protein protein interactions The chapters include a variety of in vitro and in vivo experimental methods covering cell biology biochemistry and biophysics In addition the book also explores in silico methods including sequence structure and phylogenetic profile based approaches as well as gene expression and machine learning methods Written for the highly successful Methods in Molecular Biology series chapters include introductions to their respective topics lists of the necessary materials and reagents step by step and readily reproducible laboratory protocols as well as tips on troubleshooting and avoiding known pitfalls Authoritative and practical Protein Protein Interactions Methods and Protocols serves as an ideal guide for researchers working in protein science and beyond *Using Protein-protein Interactions to Influence Small Molecule Activity* Patrick Dale Braun,2003 *Comprehensive Medicinal Chemistry III*,2017-06-03 Comprehensive Medicinal Chemistry III Eight Volume Set provides a contemporary and forward looking critical analysis and summary of recent developments emerging trends and recently identified new areas where medicinal chemistry is having an impact The discipline of medicinal chemistry continues to evolve as it adapts to new opportunities and strives to solve new challenges These include drug targeting biomolecular therapeutics development of chemical biology tools data collection and analysis in silico models as predictors for biological properties identification and validation of new targets approaches to quantify target engagement new methods for synthesis of drug candidates such as green chemistry development of novel scaffolds for drug discovery and the role of regulatory agencies in drug discovery Reviews the strategies technologies principles and applications of modern medicinal chemistry Provides a global and current perspective of today s drug discovery process and discusses the major therapeutic classes and targets Includes a unique collection of case studies and personal assays reviewing the discovery and development of key drugs **Biochemistry** Deniz Ekinci,2012-03-02 Over the recent years biochemistry has become responsible for explaining living processes such that many scientists in the life sciences from agronomy to medicine are engaged in biochemical research This book contains an overview focusing on the research area of proteins enzymes cellular mechanisms and chemical compounds used in relevant approaches The book deals with basic issues and some of the recent developments in biochemistry Particular emphasis is devoted to both theoretical and experimental aspect of modern biochemistry The primary target audience for the book includes students researchers biologists chemists chemical engineers and professionals who are interested in biochemistry molecular biology and associated areas The book is written by international scientists with expertise in protein biochemistry enzymology molecular biology and genetics many of which are active in biochemical and biomedical research We hope that the book will enhance the knowledge of scientists in the complexities of some biochemical approaches it will stimulate both professionals and students to dedicate part of their future research in understanding relevant mechanisms and applications of biochemistry HIV-1 Integrase Nouri Neamati,2011-08-10 This book comprehensively covers the mechanisms of action

and inhibitor design for HIV 1 integrase It serves as a resource for scientists facing challenging drug design issues and researchers in antiviral drug discovery Despite numerous review articles and isolated book chapters dealing with HIV 1 integrase there has not been a single source for those working to devise anti AIDS drugs against this promising target But this book fills that gap and offers a valuable introduction to the field for the interdisciplinary scientists who will need to work together to design drugs that target HIV 1 integrase **The Future of Pharmaceuticals** Sarfaraz K. Niazi, 2022-02-28

Before now biological systems could only be expressed in terms of linear relationships however as knowledge grows and new techniques of analysis on biological systems is made available we are realizing the non linearity of these systems The concepts and techniques of nonlinear analysis allow for more realistic and accurate models in science The Future of Pharmaceuticals A Nonlinear Analysis provides an opportunity to understand the non linearity of biological systems and its application in various areas of science primarily pharmaceutical sciences This book will benefit professionals in pharmaceutical industries academia and policy who are interested in an entirely new approach to how we will treat disease in the future Key Features Addresses a new approach of nonlinear analysis Applies a theory of projection to chalk out the future instead of basing on linear evolution Provides an opportunity to better understand the non linearity in biological systems and its applications in various areas of science primarily pharmaceutical sciences Helps change the thought process for those looking for answers to their questions which they do not find in the linear relationship approach Encourages a broader perspective for the creative process of drug development

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