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Springer Protocols

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Marlon J. Hinner *Editors*

Site-Specific Protein Labeling

Methods and Protocols

 Humana Press

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A. Kristina Downing



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Site-Specific Protein Labeling Arnaud Gautier, Marlon J. Hinner, 2015-01-06 This detailed volume provides in depth protocols for protein labeling techniques and applications with an additional focus on general background information on the design and generation of the organic molecules used for the labeling step Chapters provide protocols for labeling techniques and applications with an additional focus on general background information on the design and generation of the organic molecules used for the labeling step Written in the highly successful Methods in Molecular Biology series format chapters include introductions to their respective topics lists of the necessary materials and reagents step by step readily reproducible laboratory protocols and tips on troubleshooting and avoiding known pitfalls Authoritative and practical Site Specific Protein Labeling Methods and Protocols provides a comprehensive overview on the most relevant and established labeling methodologies and helps researchers to choose the most appropriate labeling method for their biological question

Proximity Labeling Murat Sunbul, Andres Jäschke, 2019-05-23 This book provides detailed protocols and untold tips and tricks regarding the most well known examples of proximity labeling methods in which the protein of interest is genetically fused to or labeled with an enzyme that can generate short lived reactive species to non specifically label molecules within a certain radius of up to twenty nanometers Beginning with peroxidase based proximity labeling methods the volume continues with BioID proximity labeling methods that describe the proximity ligation assay to detect RNA DNA interactions UV cross linking to demonstrate RNA protein interactions and how chemical and enzymatic reactivities can be improved upon DNA DNA and protein protein interactions as well as proximity induced self labeling where the radius of labeling is zero 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 readily reproducible laboratory protocols and tips on troubleshooting and avoiding known pitfalls Authoritative and practical Proximity Labeling Methods and Protocols serves as an ideal guide for researchers exploring the crucial roles that proximity driven reactions play in biological systems **Carbohydrate-Protein**

Interactions D. Wade Abbott, Wesley F. Zandberg, 2023-05-06 This second edition provides new and updated tools for studying protein carbohydrate interactions ranging from traditional biochemical methods to state of the art techniques This book focuses on four different research themes detailing methods for screening and quantifying CAZyme activity investigating the interactions between proteins carbohydrate ligands methods for the visualization of carbohydrates protein carbohydrate complexes structural and omic approaches for studying systems of CAZymes Written in the format of the highly successful Methods in Molecular Biology series each chapter includes an introduction to the topic lists necessary materials and methods includes tips on troubleshooting and known pitfalls and step by step readily reproducible protocols Authoritative and cutting edge Carbohydrate Protein Interactions Methods and Protocols Second Edition aims to be comprehensive guide for researchers in the field **SMC Proteins** Yasuto Murayama, Yasutaka Kakui, 2025-11-20 The aim

of this volume is to provide a wide range of experimental methods for analyzing molecular functions of SMC complexes such as microscopic imaging protein localization chromosome conformation analyses by deep DNA sequencing and biochemical reconstitution with purified proteins or cell lysate Written in the highly successful Methods in Molecular Biology series format the chapters include brief introductions to the material lists of necessary materials and reagents step by step readily reproducible laboratory protocols and a Notes section which highlights tips on troubleshooting and avoiding known pitfalls Authoritative and cutting edge SMC Proteins Methods and Protocols aims to be comprehensive guide for researchers in the field

Protein-Protein Interactions Haiyan Fu, 2008-02-03 As the mysteries stored in our DNA have been more completely revealed scientists have begun to face the extraordinary challenge of unraveling the intricate network of protein-protein interactions established by that DNA framework It is increasingly clear that proteins continuously interact with one another in a highly regulated fashion to determine cell fate such as proliferation differentiation or death These protein-protein interactions enable and exert stringent control over DNA replication RNA transcription protein translation macromolecular assembly and degradation and signal transduction essentially all cellular functions involve protein-protein interactions Thus protein-protein interactions are fundamental for normal physiology in all organisms Alteration of critical protein-protein interactions is thought to be involved in the development of many diseases such as neurodegenerative disorders cancers and infectious diseases Therefore examination of when and how protein-protein interactions occur and how they are controlled is essential for understanding diverse biological processes as well as for elucidating the molecular basis of diseases and identifying potential targets for therapeutic interventions Over the years many innovative biochemical biophysical genetic and computational approaches have been developed to detect and analyze protein-protein interactions This multitude of techniques is mandated by the diversity of physical and chemical properties of proteins and the sensitivity of protein-protein interactions to cellular conditions

Genetically Incorporated Non-Canonical Amino Acids Yu-Hsuan Tsai, Simon J. Elsäßer, 2023-06-05 This detailed volume explores non-canonical amino acids ncAAs through their site-specific incorporation by genetic code expansion GCE The collection provides a broad resource of methods for implementing GCE in E. coli mammalian cells and animals highlighting specific applications ranging from fluorescence labeling to photocontrol and the study of protein post-translational modification 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 and tips on troubleshooting and avoiding known pitfalls Authoritative and practical Genetically Incorporated Non-Canonical Amino Acids Methods and Protocols serves as an ideal source of methodologies that can be adapted and extended migrated to different model systems and combined in new ways to help explore a wide range of biological questions and to augment industrial and pharmaceutical protein engineering

Protein NMR Techniques A. Kristina Downing, 2008-02-03 When I was asked to edit the second edition of Protein NMR Techniques my first thought was

that the time was ripe for a new edition The past several years have seen a surge in the development of novel methods that are truly revolutionizing our ability to characterize biological macromolecules in terms of speed accuracy and size limitations I was particularly excited at the prospect of making these techniques accessible to all NMR labs and for the opportunity to ask the experts to divulge their hints and tips and to write practically about the methods I commissioned 19 chapters with wide scope for Protein NMR Techniques and the volume has been organized with numerous themes in mind Chapters 1 and 2 deal with recombinant protein expression using two organisms E coli and P pastoris that can produce high yields of isotopically labeled protein at a reasonable cost Staying with the idea of isotopic labeling Chapter 3 describes methods for perdeuteration and site specific protonation and is the first of several chapters in the book that is relevant to studies of higher molecular weight systems A different but equally powerful method that uses molecular biology to edit the spectrum of a large molecule using segmental labeling is presented in Chapter 4 Having successfully produced a high molecular weight target for study the next logical step is data acquisition Hence the final chapter on this theme Chapter 5 describes TROSY methods for structural studies

Tau Protein Caroline Smet-Nocca,2024-03-21 This volume explores the latest advancements and techniques to study Tau protein that include basic and advanced methods and protocols from in vitro assays to in vivo models that address the molecular and functional aspects of tau physiopathology and many of its related technical issues The chapters in this book are organized into five parts Part One describes conformational and functional studies of native tau protein using wet and non wet lab protocols Part Two looks at in vitro methods to monitor or control the formation of Tau oligomers and fibrils and the fibrillization process Part Three provides protocols for the characterization and in vitro introduction of post translational modifications in Tau protein for further functional studies Part Four describes analytical tools for the detection of Tau proteins under various forms factors associated with Tau pathology and MAPT gene studies Finally Part Five explores cellular and in vivo models for the investigations of Tau physiopathology Written in the highly successful Methods in Molecular Biology series format chapters include introductions to their respective topics lists of the necessary materials and reagents step by step readily reproducible laboratory protocols and tips on troubleshooting and avoiding known pitfalls Cutting edge and comprehensive Tau Proteins Methods and Protocols Second Edition is a valuable tool for any researcher interested in learning more about this important and developing field related to Tau protein as a relevant and attractive target for neurodegeneration therapies

Methods in Protein Biochemistry Harald Tschesche,2011-12-23 This book presents a survey of recent developments in protein biochemistry Top researchers in the field of protein biochemistry describe modern methods to address the challenges of protein purification by three phase partitioning and their folding and degradation by the functions of chaperones The significance of peptide purity for fibril formation is addressed as well as the use of target oriented peptide arrays in palliative approaches in mucoviscidose The design and application of protein epitope mimetics just as the structural resolving of the misfolding of various mutant

proteins in serpinopathies enlarge our tools in resolving pathophysiological imbalances **Glycovirology** Yasuo Suzuki, 2022-09-29 This volume details applications in molecular biological techniques and focuses on applications to determine the involvement of glycans in virus interactions Chapters guide readers through glycan analysis glycan distribution analysis glycan and lectin microarray preparation of recombinant viral domain protein reverse genetics and receptor binding virus host interactions and receptor binding and sialidase assays and inhibitors Written in the successful Methods in Molecular Biology series format chapters include introductions to their respective topics lists of the necessary materials and reagents step by step readily reproducible protocols and notes on troubleshooting and avoiding known pitfalls Authoritative and cutting edge Glycovirology Methods and Protocols aims to be an essential resource for researchers who wish to learn more about glycovirology

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