

Sidney P. Colowick and Nathan O. Kaplan

Methods in ENZYMOLLOGY

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Cumulative Subject Index

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WJ Hussar



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Ubiquitin and Protein Degradation Raymond Joseph Deshaies, 2005 Ubiquitin and Protein Degradation Part B will cover chemical biology ubiquitin derivatives and ubiquitin like proteins deubiquitinating enzymes proteomics as well as techniques to monitor protein degradation The chapters are highly methodological and focus on application of techniques Second part of the Ubiquitin and Protein Degradation series Topics include E1 Enzymes E2 Enzymes E3 Enzymes Proteasomes and Isopeptidases Sphingolipid Metabolism and Cell Signaling Alfred Harrison Merrill, Yusuf A. Hannun, 2000 This volume contains information on analyzing sphingolipids sphingolipid transport and trafficking and sphingolipid protein interactions and cellular targets Its companion Volume 311 presents methods used in studying enzymes of sphingolipid biosynthesis and turnover including inhibitors of some of these enzymes genetic approaches and organic and enzymatic syntheses of sphingolipids and analogs The critically acclaimed laboratory standard for more than forty years 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 Now with more than 300 volumes all of them still in print the series contains much material still relevant today truly an essential publication for researchers in all fields of life sciences *Two-Component Signaling Systems, Part B*, 2007-07-03 Multicellular organisms must be able to adapt to cellular events to accommodate prevailing conditions Sensory response circuits operate by making use of a phosphorylation control mechanism known as the two component system Sections in Two Component Signaling Systems Part B include Structural Approaches Reconstitution of Heterogeneous Systems Intracellular Methods and Assays Genome Wide Analyses of Two Component Systems Presents detailed protocols Includes troubleshooting tips **GTPases Regulating Membrane Dynamics** W. E. Balch, Channing J. Der, Alan Hall, 2005-12-13 Provides a comprehensive set of articles describing the use and application of state of the art methodologies to identify and characterize these GTPases and their expanding list of regulators and effectors This work also includes methodologies focused on biochemical molecular and advanced imaging techniques *Glycobiology* Minoru Fukuda, 2006-11-14 In the past decade there has been an explosion of progress in understanding the roles of carbohydrates in biological systems This explosive progress was made with the efforts in determining the roles of carbohydrates in immunology neurobiology and many other disciplines examining each unique system and employing new technology This volume represents the first of three in the Methods in Enzymology series including Glycomics vol 416 and Functional Glycomics vol 417 dedicated to disseminating information on methods in determining the biological roles of carbohydrates These books are designed to provide an introduction of new methods to a large variety of readers who would like to participate in and contribute to the advancement of glycobiology The methods covered include structural analysis of carbohydrates biological and chemical synthesis of carbohydrates expression and determination of ligands for carbohydrate binding proteins gene expression profiling including micro array and generation of gene knockout mice and their phenotype

analyses **Energetics of Biological Macromolecules, Part C** ,2000-08-09 Volume 323 of Methods in Enzymology is dedicated to the energetics of biological macromolecules Understanding the molecular mechanisms underlying a biological process requires detailed knowledge of the structural relationships within the system and an equally detailed understanding of the energetic driving forces that control the structural interactions This volume presents modern thermodynamic techniques currently being utilized to study the energetic driving forces in biological systems It will be a useful reference source and textbook for scientists and students whose goal is to understand the energetic relationships between macromolecular structures and biological functions This volume supplements Volumes 259 and Volume 295 of Methods in Enzymology Key Features Probing Stability of Helical Transmembrane Proteins Energetics of Vinca Alkaloid Interactions with Tubulin Deriving Complex Ligand Binding Formulas Mathematical Modeling of Cooperative Interactions in Hemoglobin Analysis of Interactions of Regulatory Protein TyrR with DNA Parsing Free Energy of Drug DNA Interactions Use of Fluorescence as Thermodynamics Tool *Glycomics* Minoru Fukuda,2006-11-13 In the past decade there has been an explosion of progress in understanding the roles of carbohydrates in biological systems This explosive progress was made with the efforts in determining the roles of carbohydrates in immunology neurobiology and many other disciplines examining each unique system and employing new technology This volume represents the second of three in the Methods in Enzymology series including Glycobiology vol 415 and Functional Glycomics vol 417 dedicated to disseminating information on methods in determining the biological roles of carbohydrates These books are designed to provide an introduction of new methods to a large variety of readers who would like to participate in and contribute to the advancement of glycobiology The methods covered include structural analysis of carbohydrates biological and chemical synthesis of carbohydrates expression and determination of ligands for carbohydrate binding proteins gene expression profiling including micro array and generation of gene knockout mice and their phenotype analyses **Amyloid, Prions, and Other Protein Aggregates, Part C** ,2006-10-06 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 *Regulators and Effectors of Small GTPases: Ras Family* ,2006-06-07 The Ras superfamily 150 human members encompasses Ras GTPases involved in cell proliferation Rho GTPases involved in regulating the cytoskeleton Rab GTPases involved in membrane targeting fusion and a group of GTPases including Sar1 Arf Arl and dynamin involved in vesicle budding fission These GTPases act as molecular switches and their activities are controlled by a large number of

regulatory molecules that affect either GTP loading guanine nucleotide exchange factors or GEFs or GTP hydrolysis GTPase activating proteins or GAPs In their active state they interact with a continually increasing functionally complex array of downstream effectors Since the last Methods in Enzymology volume on this topic in 2000 the study of Ras Family GTPases has witnessed a plethora of new directions and trends With regards to the founding member of the Ras superfamily the study of Ras in oncogenesis has seen the development and application of more advanced model cell culture and animal systems The discovery of mutationally activated B Raf in human cancers has injected renewed interest in this classical effector pathway of Ras Includes a database for Ras family proteins and their effectors and regulators Complimentary to volume 406 coverage of the Rho family Over 150 international contributors *Molecular Evolution, Producing the Biochemical Data, Part B* ,2005-06-15 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 Molecular Evolution Producing the Biochemical Data part B is a continuation of methods published in Part A 1993 volume 224 The work is a very methodological look at markers templates genomes datasets and analyses used in studies of biological diversity 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 **Two-Component Signaling Systems, Part A** ,2011-09-21 Multicellular organisms must be able to adapt to cellular events to accommodate prevailing conditions Sensory response circuits operate by making use of a phosphorylation control mechanism known as the two component system Sections include Computational Analyses of Sequences and Sequence Alignments Biochemical and Genetic Assays of Individual Components of Signaling Systems Physiological Assays and Readouts Presents detailed protocols Includes troubleshooting tips *Regulators and Effectors of Small GTPases: Rho Family* ,2006-02-21 The Ras superfamily 150 human members encompasses Ras GTPases involved in cell proliferation Rho GTPases involved in regulating the cytoskeleton Rab GTPases involved in membrane targeting fusion and a group of GTPases including Sar1 Arf Arl and dynamin involved in vesicle budding fission These GTPases act as molecular switches and their activities are controlled by a large number of regulatory molecules that affect either GTP loading guanine nucleotide exchange factors or GEFs or GTP hydrolysis GTPase activating proteins or GAPs In their active state they interact with a continually increasing functionally complex array of downstream effectors Since the last Methods in Enzymology volume on this topic in 2000 Rho GTPases have continued to receive a huge amount of attention The human genome sequence has revealed the full extent of the Rho GEF and Rho GAP families over 80 members for each and the challenge of identifying the molecular interactions and cellular pathways influenced by each of these regulators is a daunting prospect This new volume Regulators and Effectors of Small GTPases Rho Family describes some of the methods

currently being used to examine Rho family GTPase regulation at the biochemical and cellular level Describes the methods currently being used to examine Rho family GTPase regulation at the biochemical and cellular levels Includes new imaging techniques that revolutionize the ability to visualize GTPase activities Over 150 international contributors *GTPases Regulating Membrane Targeting and Fusion* W. E. Balch, Channing J. Der, Alan Hall, 2005-12-13 Rab GTPases now comprise a family of 63 members They are emerging as the key hub element controlling the membrane architecture of eukaryotic cells They are intimately involved in vesicle targeting and fusion in both the endocytic and exocytic pathways and direct the assembly and disassembly of protein complexes that include regulators GEFs and GAPs effectors tethers motors and fusion components SNAREs that control membrane targeting and fusion During the last 3 years the field has virtually exploded with the identification and characterization of many new Rab proteins and their effectors Our understanding of how Rab GTPases control membrane function remains at its infancy This volume of *Methods in Enzymology GTPases Regulating Membrane Targeting and Fusion* provides a wealth of new concepts approaches and tools to study Rab proteins in the test tube and in living cells that will be of strong benefit to both established laboratories and new investigators in the field to elucidate Rab GTPase function in cellular development differentiation and proliferation Comprehensive overview of Rab GTPase phylogeny and systems biology Identification and characterization of Rab GEFs GAPs and effectors General methodologies to study Rab GTPase function in vitro and in vivo using biochemical molecular and microscopy approaches *Liposomes, Part E*, 2005-02-28 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 Part E* is a continuation of previous *Methods in Enzymology Liposome* volumes A B C and D 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 **Regulators of G Protein Signaling, Part B**, 2004-11-16 *Regulators of G Protein Signaling Part B* continues the in depth treatment of the topic and covers the RSG protein superfamily including RZ R4 R7 R12 RhoGEF and GRK as well as other heterotrimeric G protein signaling regulators Table of Contents RZ Subfamily R4 Subfamily R7 Subfamily R12 Subfamily RhoGEF Subfamily GRK Subfamily Other RGS proteins Activators Inhibitors Other Modulators *Antisense Technology, Part B: Applications*, 1999-11-10 Antisense technology is the ability to manipulate gene expression within mammalian cells providing powerful experimental approaches for the study of gene function and gene regulation For example methods which inhibit gene expression permit studies probing the normal function of a specific product within a cell Such methodology can be used in many disciplines such as pharmacology oncology genetics cell biology developmental biology molecular biology biochemistry and neurosciences This volume will be a truly important tool in biomedically oriented research The critically acclaimed laboratory standard for more than forty years *Methods in Enzymology* is one of the most highly respected publications in the field of biochemistry Since

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