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NUCLEAR MAGNETIC RESONANCE OF BIOLOGICAL MACROMOLECULES

Volume 338

Thomas L. James

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Nuclear Magnetic Resonance of Biological Macromolecules, Part A ,2001-07-12 This volume and its companion Volume 339 supplement Volumes 176 177 239 and 261 Chapters are written with a hands on perspective That is practical applications with critical evaluations of methodologies and experimental considerations needed to design execute and interpret NMR experiments pertinent to biological molecules

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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 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 Phase II Conjugation Enzymes and Transport Systems Helmut Sies, Lester Packer, 2005-11-28 This volume on conjugation enzymes and transporters serves to bring together current methods and concepts in an interesting important and rapidly developing field of cell and systems biology Phase II Conjugation Enzymes and Transport Systems focuses on the so called Phase II enzymes of drug metabolism xenobiotics which has important ramifications for endogenous metabolism and nutrition Also included are aspects on Phase III transport systems This volume of Methods in Enzymology presents current knowledge and methodology on glucuronidation sulfation

acetylation and transport systems in this field of research Together with the volumes on Quinones and Quinone Enzymes volumes 378 and 382 and on Glutathione Transferases and gamma Glutamyl Transpeptidases volume 401 the state of knowledge on proteomics and metabolomics of many pathways of waste product elimination enzyme protein induction and gene regulation and feedback control is provided This volume will help stimulate future investigations and speed the advance of knowledge in systems biology A laboratory standard for more than 40 years Over 400 volumes strong Also available on 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 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 **Functional Glycomics** Minoru Fukuda, 2006-12-08 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 Glycomics vol 416 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 **Mass Spectrometry: Modified Proteins and Glycoconjugates** A.L. Burlingame, 2005-12-13 This volume provides comprehensive treatment of tools and proper usage for the identification of proteins affinity chromatography and studies the complexity of protein machines and assemblages assignment of the most common protein posttranslational modifications phosphorylation and glycosylation and glycolipidomics Part 2 of 2 volumes about Mass Spectrometry Discusses peptide and protein cleanup and preparation requirements for mass spectrometry Explains protein enzymic and chemical digestion strategies Includes case studies of Ubiquitin and Protein Degradation Raymond Joseph Deshaies, 2005 Ubiquitin and protein assemblages and machines 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 Degration series Topics include E1 Enzymes E2 Enzymes E3 Enzymes Proteasomes and Isopeptidases **Embryonic Stem Cells**, 2006-12-08 This is the first of three planned volumes in the Methods in Enzymology series on the topic of stem cells This volume is a unique anthology of stem cell techniques written by experts from the top laboratories in the world The contributors not only have hands on experience in the field but often have developed the original approaches that they share with great attention to detail The chapters provide a brief review of each field followed by a cookbook and handy illustrations The collection of protocols includes the isolation and maintenance of stem cells from various species using conventional and novel methods such as derivation of ES cells from single blastomeres differentiation of stem cells into specific tissue types isolation and maintenance of somatic stem cells stem cell specific techniques and approaches to tissue engineering using stem cell derivatives. The reader will find that some of the topics are covered by more than one group of authors and complement each other Comprehensive step by step protocols and informative illustrations can be easily followed by even the least experienced researchers in the field and allow the setup and troubleshooting of these state of the art technologies in other laboratories Provides complete coverage spanning from derivation isolation of stem cells and including differentiation protocols characterization and maintenance of derivatives and tissue engineering Presents the latest most innovative technologies Addresses therapeutic relevance including FDA compliance and tissue engineering 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 **Advanced Bacterial Genetics: Use of Transposons and Phage for Genomic Engineering**, 2007-02-27 The critically acclaimed laboratory standard for more than fifty 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 over 400 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. This new volume presents methods related to the use of bacterial genetics for genomic engineering The book includes sections on strain collections and genetic nomenclature transposons and phage Other Experimental Protocols, 2006-12-12 This is the third of three planned volumes in the Methods in Enzymology series on the topic of stem cells This volume is a unique anthology of stem cell techniques written by experts from the top laboratories in the world The contributors not only have hands on experience in the field but often have developed the original approaches that they share with great attention to detail The chapters provide a brief review of each field followed by a cookbook and handy illustrations The collection of protocols includes the isolation and maintenance of stem cells from various species using conventional and novel methods such as derivation of ES cells from single blastomeres differentiation of stem cells into specific tissue types isolation and maintenance of somatic stem cells stem cell specific techniques and approaches to tissue engineering using stem cell derivatives. The reader will find that some of the topics are covered by more than one group of authors and complement each other Comprehensive step by step protocols and informative illustrations can be easily followed by even the least experienced researchers in the field and allow the setup and troubleshooting of these state of the art technologies in other laboratories Provides complete coverage spanning from derivation isolation of stem cells and including differentiation protocols characterization and maintenance of derivatives and tissue engineering Presents the latest most innovative technologies Addresses therapeutic relevance including FDA compliance and tissue engineering Adult Stem Cells, 2006-12-11 This is the second of three planned volumes in the Methods in Enzymology series on the

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and gene discovery Early commitment steps and generation of chimeric mice Differentiation to mesoderm derivatives Gene discovery by manipulation of mouse embryonic stem cells *Protein Phosphatases* Susanne Klumpp, Josef Krieglstein, 2003 Protein phosphatases are a group of enzymes responsible for the dephosphorylation of various proteins and enzymes in a cell This role is an extremely important one since protein phosphorylation and dephosphorylation is required for the regulation of a large number of cellular activities Classification of Protein Phosphatases Analysis Technology Cell and Molecular Imaging Technology Assay of Protein Phosphatases MS and MNR Genomics Proteomics cDNA Microarray Analysis Cellular Regulation Substrates Inhibitors Regulation Protein Interaction Biological Function Antisense Studies Transgenic and Knockout Animal Models in Vivo Therapeutic Approaches <u>Protein Engineering</u> Dan Robertson, Joseph P. Noel, 2004-09-29 This MIE volume covers methods for a multitude of topics among which are computational methods laboratory methods enzyme optimization binding proteins antibodies and screening technologies Table of Contents Methodology Applications Opzimization and Screening Applications Directed Evolution of Enzymatic Function Applications Evolution of Biosynthetic Pathways Devices Antibodies and Vaccines

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