

Methods in ENZYMOLOGY

Volume 248

Proteolytic Enzymes:
Aspartic and Metallo Peptidases

Edited by

Alan J. Barrett



ACADEMIC PRESS

Proteolytic Enzymes Vol 248 Aspartic And Metallo Peptidases

Eckhard Jankowsky



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Proteolytic Enzymes Erwin E. Sterchi, Walter Stöcker, 2013-11-11 Researchers may find themselves confronted with proteases either because they play an essential role in a particular process they are studying or because they interfere with that process In either case they may need to investigate or inhibit the proteolytic activity Others may wish to use proteolytic enzymes as laboratory tools This book has been written with these investigators in mind and includes assay methods using natural and artificial substrates genetic based assays and strategies for the inhibition purification and crystallization of proteases In selected chapters the use of proteolytic enzymes to analyze proteins segregate cells or in peptide synthesis is covered

Fragment Based Drug Design, 2011-02-28 There are numerous excellent reviews on fragment based drug discovery FBDD but there are to date no hand holding guides or protocols with which one can embark on this orthogonal approach to complement traditional high throughput screening methodologies This *Methods in Enzymology* volume offers the tools practical approaches and hit to lead examples on how to conduct FBDD screens The chapters in this volume cover methods that have proven to be successful in generating leads from fragments including chapters on how to apply computational techniques nuclear magnetic resonance surface plasma resonance thermal shift and binding assays protein crystallography and medicinal chemistry in FBDD Also elaborated by experienced researchers in FBDD are sample preparations of fragments proteins and GPCR as well as examples of how to generate leads from hits Offers the tools practical approaches and hit to lead examples on how to conduct FBDD screens The chapters in this volume cover methods that have proven to be successful in generating leads from fragments including chapters on how to apply computational techniques nuclear magnetic resonance surface plasma resonance thermal shift and binding assays protein crystallography and medicinal chemistry in FBDD

Two-Component Signaling Systems, Part C, 2010-03-01 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 This volume the third in a three volume treatment edited by the same group of editors includes a wide range of methods including those dealing with the Sln 1 kinase pathway triazole sensitivity in *C. albicans* and histidine kinases in cyanobacteria circadian clock Includes time tested core methods and new innovations applicable to any researcher studying two component signaling systems or histidine kinases Methods included are useful to both established researchers and newcomers to the field Relevant background and reference information given for procedures can be used as a guide to developing protocols in a number of disciplines

Lipidomics and Bioactive Lipids: Mass Spectrometry Based Lipid Analysis, 2007-11-26 This volume in the well established *Methods in Enzymology* series features methods for the study of lipids using mass spectrometry techniques Articles in this volume cover topics such as Qualitative Analysis and Quantitative Assessment of Changes in Neutral Glycerol Lipid Molecular Species within Cells Glycerophospholipid identification and quantitation by electrospray ionization mass spectrometry

Detection and Quantitation of Eicosanoids via High Performance Liquid Chromatography Electrospray Ionization Mass Spectrometry Structure specific quantitative methods for lipidomic analysis of sphingolipids by tandem mass spectrometry Analysis of Ubiquinones Dolichols and Dolichol Diphosphate Oligosaccharides by Liquid Chromatography Electrospray Ionization Mass Spectrometry Extraction and Analysis of Sterols in Biological Matrices by High Performance Liquid Chromatography Electrospray Ionization Mass Spectrometry The Lipid Maps Initiative in Lipidomics Basic analytical systems for lipidomics by mass spectrometry in Japan The European Lipidomics Initiative Enabling technologies Lipidomic analysis of Signaling Pathways Bioinformatics for Lipidomics Mediator Lipidomics Search Algorithms for Eicosanoids Resolvins and Protectins A guide to biochemical systems modeling of sphingolipids for the biochemist and Quantitation and Standardization of Lipid Internal Standards for Mass Spectroscopy **Computer Methods, Part C** ,2010-12-24 The combination of faster more advanced computers and more quantitatively oriented biomedical researchers has recently yielded new and more precise methods for the analysis of biomedical data These better analyses have enhanced the conclusions that can be drawn from biomedical data and they have changed the way that experiments are designed and performed This volume along with the 2 previous Computer Methods volumes for the Methods in Enzymology series aims to inform biomedical researchers about recent applications of modern data analysis and simulation methods as applied to biomedical research Presents step by step computer methods and discusses the techniques in detail to enable their implementation in solving a wide range of problems Informs biomedical researchers of the modern data analysis methods that have developed alongside computer hardware Presents methods at the nuts and bolts level to identify and resolve a problem and analyze what the results mean

Biology of Serpins James Whisstock, Phillip Bird, 2011-06-11 Serpins are a group of proteins with similar structures that were first identified as a set of proteins able to inhibit proteases The acronym serpin was originally coined because many serpins inhibit chymotrypsin like serine proteases This volume of Methods in Enzymology is split into 2 parts and comprehensively covers the subject **Constitutive Activity in Receptors and Other Proteins, Part A** ,2010-11-26 This volume of Methods in Enzymology covers the current methodology for the detection and assessment of constitutively active proteins The chapters written by expert authors who are leaders in the field provide hints and tricks not available in primary research publications It is extensively referenced with useful figures and tables throughout the volume Expert authors who are leaders in the field Extensively referenced and useful figures and tables Provides hints and tricks to facilitate reproduction of methods

Methods in Methane Metabolism Amy Claire Rosenzweig, Stephen W. Ragsdale, 2011 Produced by microbes on a large scale methane is an important alternative fuel as well as a potent greenhouse gas This volume focuses on microbial methane metabolism which is central to the global carbon cycle Both methanotrophy and methanogenesis are covered in detail Topics include isolation and classification of microorganisms metagenomics approaches biochemistry of key metabolic enzymes gene regulation and genetic systems and field measurements The state of the art

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Protein Engineering for Therapeutics, Part A ,2012-02-23 This volume of Methods in Enzymology looks at Protein Engineering for Therapeutics The chapters provide an invaluable resource for academics researchers and students alike With an international board of authors this volume is split into sections that cover subjects such as Antibodies Protein conjugates Peptides Enzymes and Scaffolds Chapters provide an invaluable resource for academics researchers and students alike Iinternational board of authors This volume is split into sections that cover subjects such as Antibodies Protein conjugates Peptides Enzymes and Scaffolds

Synthetic Biology, Part B Chris Voigt,2011-07-15 Synthetic biology encompasses a variety of different approaches methodologies and disciplines and many different definitions exist This Volume of Methods in Enzymology has been split into 2 Parts and covers topics such as Measuring and Engineering Central Dogma Processes Mathematical and Computational Methods and Next Generation DNA Assembly and Manipulation Encompasses a variety of different approaches methodologies and disciplines Split into 2 parts and covers topics such as measuring and engineering central dogma processes mathematical and computational methods and next generation DNA assembly and manipulation

Imaging and Spectroscopic Analysis of Living Cells, Optical and Spectroscopic Techniques P. Michael Conn,2012 This volume of Methods in Enzymology is the first of 3 parts looking at current methodology for the imaging and spectroscopic analysis of live cells The chapters provide hints and tricks not available in primary research publications It is an invaluable resource for academics researchers and students alike Expert authors who are leaders in the field Extensively referenced and useful figures and tables Provides hints and tricks to facilitate reproduction of methods

Methods in Systems Biology ,2011-09-19 Systems biology is a term used to describe a number of trends in bioscience research and a movement that draws on those trends This volume in the Methods in Enzymology series comprehensively covers the methods in systems biology With an international board of authors this volume is split into sections that cover subjects such as machines for systems biology protein production and quantification for systems biology and enzymatic assays in systems biology research This volume in the Methods in Enzymology series comprehensively covers the methods in systems biology With an international board of authors this volume is split into sections that cover subjects such as machines for systems biology protein production and quantification for systems biology and enzymatic assays in systems biology research

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Guide to Techniques in Mouse Development, Part A, 2010-08-04 Guide to Techniques in Mouse Development Part A comprehensively covers new technologies and methodologies that have appeared for the study of mouse development Update of volume 225 of Methods in Enzymology Guide to Techniques in Mouse Development edited by P M Wassarman and M L DePamphilis and published in 1993 Covers new technologies and methodologies including new techniques for the cryopreservation of gametes and embryos production of transgenic and null knockout animals use of ES cells generation of conditional inducible mutant animals use of gene trap mutagenesis analysis of allele specific expression use of new reporter constructs humanizing of transgenic animals transcript profiling of mouse development imaging of mouse development rederivation of animals and use of mouse genomics **Functional Glycomics**, 2010-10-25 In this 3 volume collection focusing on glycomics readers will appreciate how such discoveries were made and how such methods can be applied for readers own research efforts Each chapter has been designed so that enough scientific background will be given in each chapter for further development of methods by readers themselves Useful for all levels of scientists starting from the last years of colleges graduate students postdoctoral fellows to professors and to all levels of scientists in research institutes including industry **Cryo-EM Part A: Sample Preparation and Data Collection**, 2010-09-30 Cryo EM Part A Sample Preparation and Data Collection is dedicated to a description of the instruments samples protocols and analyses that belong to cryo EM It emphasizes the relatedness of the ideas instrumentation and methods underlying all cryo EM approaches which allow practitioners to easily move between them Within each section the articles are ordered according to the most common symmetry of the sample to which their methods are applied Includes time tested core methods and new innovations applicable to any researcher Methods included are useful to both established researchers and newcomers to the field Relevant background and reference information given for procedures can be used as a guide **Methods in Methane Metabolism, Part B** Amy Rosenzweig, Stephen W. Ragsdale, 2011-03-17 Produced by microbes on a large scale methane is an important alternative fuel as well as a potent greenhouse gas This volume focuses on microbial methane metabolism which is central to the global carbon cycle Both methanotrophy and methanogenesis are covered in detail Topics include isolation and classification of microorganisms metagenomics approaches biochemistry of key metabolic enzymes gene regulation and genetic systems and field measurements The state of the art techniques described here will both guide researchers in specific pursuits and educate the wider scientific community about this exciting and rapidly developing field Topics include isolation and classification of microorganisms metagenomics approaches biochemistry of key metabolic enzymes gene regulation and genetic systems and field measurements The state of the art techniques described here will

both guide researchers in specific pursuits and educate the wider scientific community about this exciting and rapidly developing field *RNA Helicases* Eckhard Jankowsky, 2012-06-22 This work integrates the current knowledge about RNA helicases from diverse fields ranging from cell and developmental biology to mechanistic enzymology and structural biology into one useful resource Protein Engineering for Therapeutics, Part B , 2012-01-25 This volume of Methods in Enzymology looks at Protein Engineering for Therapeutics The chapters provide an invaluable resource for academics researchers and students alike With an international board of authors this volume is split into sections that cover subjects such as Peptides and Scaffolds Chapters provide an invaluable resource for academics researchers and students alike International board of authors This volume is split into sections that cover subjects such as peptides and scaffolds *Biothermodynamics* Michael L. Johnson, Jo M. Holt, Gary K. Ackers, 2011

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