

Methods in Molecular Biology™

VOLUME 206

Peptide Research Protocols

Endothelin

Edited by

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Peptide Research Protocols Endothelin Methods In Molecular Biology

Subhash C. Basu, Manju Basu



Peptide Research Protocols Endothelin Methods In Molecular Biology:

Peptide Research Protocols Janet J. Maguire, Anthony P. Davenport, 2008-02-04 A panel of multidisciplinary experts describes in detail readily reproducible methods to investigate all aspects of the endothelin system from its synthesis and metabolism to its function in health and disease These methods use state of the art molecular techniques to quantify the expression of mRNA for both endothelin receptors and the endothelin converting enzymes They show how peptides precursors receptors and synthetic enzymes can be localized and quantified in plasma culture supernatants tissue homogenate and tissue sections using antibodies Several in vivo protocols illustrate the role of the endothelin peptides in healthy human individuals and describe animal models that can be used to predict the therapeutic potential of cardiovascular drugs that manipulate endothelin synthesis or function

Methods in Molecular Biology: Peptide research protocols: endothelin John M. Walker, 1984

Peptide Synthesis and Applications John Howl, 2008-02-02 Hands on experts describe in step by step detail the key methodologies of contemporary peptide synthesis and illustrate their numerous applications The techniques presented include protocols for chemical ligation the synthesis of cyclic and phosphotyrosine containing peptides lipoamino acid and sugar conjugated peptides and peptide purification and analyses Additional chapters detail methodologies and instrumentation for high throughput peptide synthesis many different applications of peptides as novel research tools and biological probes and the design and application of fluorescent substrate based peptides that can be used to determine the selectivity and activity of peptidases A practical guide to the identification of proteins using mass spectrometric analyses of peptide mixtures is also included

Liposome Methods and Protocols Subhash C. Basu, Manju Basu, 2008-02-04 In vitro utilization of liposomes is now recognized as a powerful tool in many bioscience investigations and their associated clinical studies e g liposomes in drug targeting liposomes in gene transport across plasma and nuclear membranes liposomes in enzyme therapy in patients with genetic disorders However before these areas can be effectively explored many basic areas in liposome research require elucidation including a attachment of liposomes to cell surfaces b permeation of liposomes through the plasma membranes and c stability of liposomes in cell or nuclear matrices None of these areas have been exhaustively explored and liposome researchers have ample opportunities to contribute to our knowledge The aim of *Liposome Methods and Protocols* is to bring together a wide range of detailed laboratory protocols covering different aspects of liposome biology in order to assist researchers in those rapidly advancing medical fields mentioned earlier With this goal in mind in each protocol chapter we have detailed the materials to be used followed by a step by step protocol The Notes section of each protocol is also certain to prove particularly useful since the authors include troubleshooting tips straight from their benchtops valuable information that is seldom given in restricted methods sections of standard research journals For this reason we feel that the book will prove especially useful for all researchers in the liposome field

Superantigen Protocols Teresa Krakauer, 2008-02-05 Leading researchers in the biological chemical and physical investigation of

superantigens describe in step by step detail their best experimental techniques to assess the physical characteristics and biological effects of superantigens Their protocols range from those for investigating the interactions of superantigens with cellular receptors to those for the analysis of their immunological and biological effects including methods for using BIOcore to determine binding kinetics and establishing various lymphocyte cell culture systems There are also accounts of such methods as the RNase protection assay cytokine ELISA FACS analysis and cytokine production at the single cell level

PCR Protocols John M. S. Bartlett, David Stirling, 2008-02-03 In this new edition the editors have thoroughly updated and dramatically expanded the number of protocols to take advantage of the newest technologies used in all branches of research and clinical medicine today These proven methods include real time PCR SNP analysis nested PCR direct PCR and long range PCR Among the highlights are chapters on genome profiling by SAGE differential display and chip technologies the amplification of whole genome DNA by random degenerate oligonucleotide PCR and the refinement of PCR methods for the analysis of fragmented DNA from fixed tissues Each fully tested protocol is described in step by step detail by an established expert in the field and includes a background introduction outlining the principle behind the technique equipment and reagent lists tips on trouble shooting and avoiding known pitfalls and where needed a discussion of the interpretation and use of results

GTPase Protocols Ed Manser, Thomas Leung, 2008-02-03 In the last 10 years researchers have firmly established key roles for R related GTPases in almost every aspect of cell biology In the 1980s the pro oncogene Ras itself was the focus of interest though in the 1990s this shifted to the increasing variety of Ras related proteins In this new decade much yet needs to be done to establish the role for all the small GTPases now uncovered by the human genome project In particular these GTPases need to be understood in the appropriate biochemical and cellular contexts In the process of trying to uncover the role of these versatile proteins a variety of novel techniques and methodologies has been developed These now enable investigators to move easily within a diversity of fields ranging from structural studies to real time in vivo analysis of a GTPase In recognition of the need for access to key background methodologies GTPase Protocols The Ras Superfamily is devoted to techniques that are presently widely used and that will continue to be the standard for researchers worldwide Each chapter is aimed at supplying detailed methodologies to allow reproduction in any laboratory while also providing the general principles on which the methods are based Some of the techniques grouped in the first section apply broadly to small GTPases whereas others in Part II are more applicable within each GTPase subfamily

The Pharmacology of Functional, Biochemical, and Recombinant Receptor Systems T. Kenakin, J. A. Angus, 2012-12-06 This the 148th volume of the Handbook of Experimental Pharmacology series focuses on the very core of pharmacology namely receptor theory It is fitting that the originator of receptor pharmacology A J CLARK authored the fourth volume of this series 63 years ago In that volume CLARK further developed his version of receptor theory first described four years earlier in his classic book *The Mode of Action of Drugs* An examination of the topics covered in volume 4 reveals a striking similarity to the topics covered in this present

volume pharmacologists today are still as interested in unlocking the secrets of dose response relationships to reveal the biological and chemical basis of drug action as they were over half a century ago Sections in that 1937 volume such as Curves relating exposure to drugs with biological effects and Implications of monomolecular theory show Clark's keen insight into the essential questions that required answers to move pharmacology forward With the advent of molecular biological cloning of human receptors has come a transformation of receptor pharmacology Thus the expression of human receptors into surrogate host cells helped unlock secrets of receptor mechanisms and stimulus transduction pathways To a large extent this eliminates the leap of faith required to apply receptor activity of drugs tested on animal receptor systems to the human therapeutic arena However a new leap of faith concerning the veracity of the effects found in recombinant systems with respect to natural ones is now required

Neuronal Cell Death Arezu Jahani-Asl, 2022-07-01 This volume covers comprehensive methods on ways to assess structural and ultrastructural changes in the mitochondria cytoskeleton and microglia using state of the art microscopy techniques including super resolution imaging electron microscopy and ultra high field MRI The chapters in this book cover topics such as analysis of neurodegeneration in the post mortem characterization of preclinical animal models in vivo modeling in cell death in different model systems and brain organoids single cell clonal analysis using Mosaic Analysis with Double Markers in genetic mouse models and genome and proteomic methods for analysis of mRNA dynamics and quantitation of targeted peptides 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 thorough Neuronal Cell Death Methods and Protocols is a valuable resource for any scientist and researcher interested in learning more about this developing field

MHC Protocols Stephen H. Powis, Robert W. Vaughan, 2008-02-05 The aim of MHC Protocols is to document protocols that can be used for the analysis of genetic variation within the human major histocompatibility complex MHC HLA region The human MHC encompasses approximately 4 million base pairs on the short arm of chromosome 6 at cytogenetic location 6p21.3 The region is divided into three subregions The telomeric class I region contains the genes that encode the HLA class I molecules HLA A B and C The centromeric class II region contains the genes encoding the HLA class II molecules HLA DR DQ and DP In between is the class III region originally identified because it contains genes encoding components of the complement pathway The entire human MHC has recently been sequenced 1 and each subregion is now known to contain many other genes a number of which have immunological functions The study of polymorphism within the MHC is well established because the region contains the highly polymorphic HLA genes HLA polymorphism has been used extensively in solid organ and bone marrow transplantation to match donors and recipients As a result large numbers of HLA alleles have been identified a process that has been further driven by recent interest in HLA gene diversity in ethnic populations The extreme genetic variation in HLA genes is believed to have been driven by the

evolutionary response to infectious agents but relatively few studies have analyzed associations between HLA genetic variation and infectious disease which has been difficult to demonstrate

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