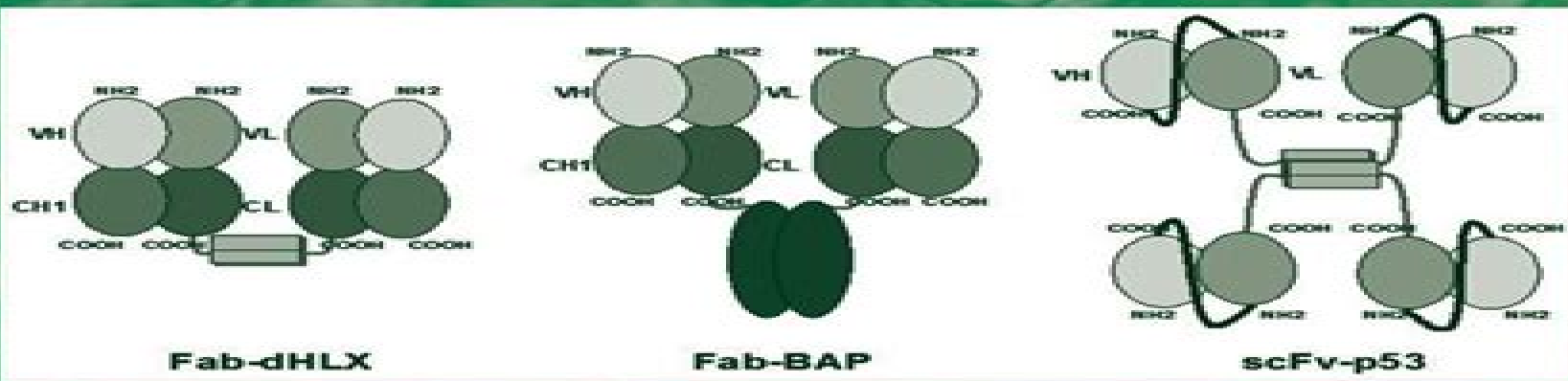


Monoclonal Antibodies

Methods and Protocols

Edited by

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Monoclonal Antibodies Methods And Protocols Methods In Molecular Biology Vol 378

Dario Leister, Johannes M. Herrmann



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Monoclonal Antibodies Maher Albitar, 2008-02-02 This book examines a collection of state of the art methods that employ monoclonal antibodies in a clinical setting The chapters offer in depth description for generating mouse and recombinant humanized antibodies and a comprehensive review of how antibodies are being used in bead based methods for measuring proteins This field will continue to expand and provide new and innovative techniques in the laboratory and as a basis that complements targeted therapy

PCR Primer Design Anton Yuryev, 2008-02-03 In the past decade molecular biology has been transformed from the art of cloning a single gene to a statistical science measuring and calculating properties of entire genomes New high throughput methods have been developed for genome sequencing and studying the cell at different systematic levels such as transcriptome proteome metabolome and other omes At the heart of most high throughput methods is the technique of polymerase chain reaction PCR PCR Primer Design focuses on primer design which is critical to both the efficiency and the accuracy of the PCR With intricate descriptions of basic approaches as well as specialized methods this volume is an exceptional reference for all those involved in studying the genome In PCR Primer Design authors describe basic approaches for PCR primer design in addition to specialized methods These state of the art methods can be used for both genome scale experiments and for small scale individual PCR amplifications This volume will be useful for organizations performing whole genome studies companies designing instruments that utilize PCR and individual scientists geneticists molecular biologists molecular geneticists and more who routinely use PCR in their research

Mass Spectrometry Data Analysis in Proteomics Rune Matthiesen, 2008-02-02 Mass Spectrometry Data Analysis in Proteomics is an in depth guide to the theory and practice of analyzing raw mass spectrometry MS data in proteomics As MS is a high throughput technique proteomic researchers must attend carefully to the associated field of data analysis and this volume outlines available bioinformatics programs algorithms and databases available for MS data analysis General guidelines for data analysis using search engines such as Mascot X tandem and VEMS are provided with specific attention to identifying poor quality data and optimizing search parameters Several different types of MS data are discussed followed by a description of optimal methods for conversion of raw data into peak lists for input to search engines Choosing the most accurate and complete databases is emphasized and a report of available sequence databases is included Methods for assembling expressed sequence tags ESTs into assembled nonredundant databases are provided along with protocols for further processing the sequences into a format suitable for MS data Mass Spectrometry Data Analysis in Proteomics describes publicly available applications whenever possible

Protein Phosphatase Protocols Greg Moorhead, 2008-02-05 Protein Phosphatase Protocols presents a broad range of protocols for the study of protein phosphatases all written by experts and innovators from phosphatase laboratories around the world This volume is a compendium of resources for the study of protein phosphatases and their potential as drug targets Experimental methodologies are taken from proteomics bioinformatics genomics biochemistry RNAi and genetics

In Vitro Transcription and Translation Protocols Guido Grandi, 2007-05-03 This book is a highly anticipated update of the previous edition It provides molecular biology laboratories with the most powerful techniques for exploiting in vitro transcription and translation systems It has been completely updated with new chapters and topics

Cardiac Gene Expression Jun Zhang, Gregg Rokosh, 2008-02-03 Cardiac Gene Expression Methods and Protocols presents both cutting edge and established methods for studying cardiac gene expression The protocols provide a template for solid research and cover the process through screening analysis characterization and functional confirmation of novel genes or known genes with a new function Section I Cardiac Gene Expression Profiling The Global Perspective discusses several different approaches to examining identifying and analyzing changes in transcriptome gene expression Section II Cardiac Gene Regulation Gene Specific mRNA Measurement in the Myocardium outlines more sensitive and gene targeted expression methods Section III Cardiac Gene Regulation Promoter Characterization in the Myocardium provides protocols for the study of underlying gene regulation mechanisms by focusing on the interaction of transcription factors with their cognate cis binding elements Section IV In Silico Assessment of Regulatory cis Elements and Gene Regulation and Section V Cardiac Single Network Polymorphisms emphasize new analytical approaches for deciphering the functional elements buried in the 3 billion nucleotides of the human genome and other model genomes The concluding section Gene Overexpression and Targeting in the Myocardium highlights methods that facilitate overexpression or cardiac specific targeted gene deletion

Metabolomics Wolfram Weckwerth, 2008-02-04 Metabolomics Methods and Protocols examines the state of the art in metabolomic analysis Leading researchers in the field present protocols for the application of complementary analytical methods such as gas chromatography mass spectrometry GC MS Metabolomics Methods and Protocols contains forward looking protocols which provide the essential groundwork for future efforts in elucidating the structure of the unknowns detected in metabolomic studies

Adhesion Protein Protocols Amanda S. Coutts, 2008-02-03 The second edition of Adhesion Protein Protocols combines traditional techniques with cutting edge and novel techniques that can be adapted easily to different molecules and cell types The topics discussed include novel techniques for studying cell cell adhesion neutrophil chemotaxis in vitro assays used to study leukocyte migration through monolayers of cultured endothelial cells and novel techniques to purify pseudopodia from migratory cells The protocols discussed in this volume are suitable for both novice and expert scientists who will gain further insight into the complex and incompletely understood processes involved in cellular adhesion

Gene Function Analysis Michael F. Ochs, 2007-08-23 With the advent of high throughput technologies following completion of the human genome project and similar projects the number of genes of interest has expanded and the traditional methods for gene function analysis cannot achieve the throughput necessary for large scale exploration This book brings together a number of recently developed techniques for looking at gene function including computational biochemical and biological methods and protocols

High Content Screening D. Lansing Taylor, 2008-02-04 There has always been some

tension between proponents of hypothesis driven and discovery driven research in the broad field of life sciences Academic research has been primarily focused on hypothesis driven research However the success of the human genome project a discovery driven research approach has opened the door to adding other types of discovery driven research to a continuum of research approaches In contrast drug discovery research in the pharmaceutical industry has embraced discovery driven research for many years A good example has been the discovery of active compounds from large chemical libraries through screening campaigns The success of the human genome project has also demonstrated the need for both academic researchers and industrial researchers to now understand the functions of genes and gene products The cell is the basic unit of life and it has been at the cellular level where function can be demonstrated most cost effectively and rapidly High content screening HCS was developed by Cellomics Inc in the mid 1990s to address the need for a platform that could be used in the discovery driven research and development required to understand the functions of genes and gene products at the level of the cell

Microarrays Jang B. Rampal, 2007-07-27 Microarray Technology Volumes 1 and 2 present information in designing and fabricating arrays and binding studies with biological analytes while providing the reader with a broad description of microarray technology tools and their potential applications The first volume deals with methods and protocols for the preparation of microarrays The second volume details applications and data analysis which is important in analyzing the enormous data coming out of microarray experiments Among the topics discussed in Volume 1 Synthesis Methods are matrices in the synthesis of microarrays array optimization processes array based comparative genomic hybridization 60 mer oligonucleotide probes bifunctional reagents NTMTA and NTPAC and high density arrays using digital microarray synthesis platforms Other topics include multiplex ligation dependent probe amplification MLPA hybridization conditions in situ synthesized oligo arrays peptide arrays high density replication tools HDRT protocols for the quantification of oligo hybridization glyco bead arrays and an investigation into the emerging nano technology Microarray Technology Volumes 1 and 2 provide ample information to all levels of scientists from novice to those intimately familiar with array technology

Glycoviropology Protocols Richard J. Sugrue, 2008-01-18 Glycoviropology Protocols reviews the increasing importance of glycosylation to the field of viropology as well as virus replication The chapters provide an overview of glycosylation in relation to virus infection and the generic techniques that are used to analyze and characterize glycoproteins

Mitochondria Dario Leister, Johannes M. Herrmann, 2007-06-12 Mitochondrial Genomics and Proteomics Protocols offers a broad collection of methods for studying the molecular biology function and features of mitochondria In the past decade mitochondrial research has elucidated the important influence of mitochondrial processes on integral cell processes such as apoptosis and cellular aging This practical guide presents a wide spectrum of mitochondrial methods each written by specialists with solid experience and intended for implementation by novice and expert researchers alike Part I introduces major experimental model systems and discusses their specific advantages and limitations for functional analysis of mitochondria The concise

overview of general properties of mitochondrial systems is supplemented by detailed protocols for cultivation of model organisms Parts II VI comprise a robust collection of protocols for studying different molecular aspects of mitochondrial functions including genetics and microbiology biochemistry physiology dynamics and morphology and functional genomics Emphasis is placed on new and emerging topics in mitochondrial study such as the examination of apoptotic effects fusion and fission of mitochondria and proteome and transcriptome analysis

Immunoinformatics Darren R. Flower, 2008-02-07 Immunoinformatics Predicting Immunogenicity In Silico is a primer for researchers interested in this emerging and exciting technology and provides examples in the major areas within the field of immunoinformatics This volume both engages the reader and provides a sound foundation for the use of immunoinformatics techniques in immunology and vaccinology The volume is conveniently divided into four sections The first section Databases details various immunoinformatic databases including IMGT HLA IPD and SYPEITHI In the second section Defining HLA Supertypes authors discuss supertypes of GRID CPCA and hierarchical clustering methods Hla Ad supertypes MHC supertypes and Class I Hla Alleles The third section Predicting Peptide MCH Binding includes discussions of MCH binders T Cell epitopes Class I and II Mouse Major Histocompatibility and HLA peptide binding Within the fourth section Predicting Other Properties of Immune Systems investigators outline TAP binding B cell epitopes MHC similarities and predicting virulence factors of immunological interest Immunoinformatics Predicting Immunogenicity In Silico merges skill sets of the lab based and the computer based science professional into one easy to use insightful volume

Capillary Electrophoresis Philippe Schmitt-Kopplin, 2008-02-25 This book presents a selection of current capillary electrophoresis methods used to separate representative types of molecules and particles and in combination with different detection techniques It includes practical details which are hard to find elsewhere The volume is intended for beginners in the field and provides an overview of the technique and a starting point for the exploration of the defined literature on different application topics

Innate Immunity Jonathan Ewbank, Eric Vivier, 2008 Immunologists today are interested in all of the diverse cell types involved in host defense and have a deeper appreciation of the importance of innate immune mechanisms as a first line of protection against pathogens This volume thus discusses the isolation and functional characterization of cells involved in innate immunity in mouse and man including mast cells and eosinophils Other focuses include natural killer cells methods in statistics in vivo imaging genome engineering and mutagenesis and culture that are adapted to the study of innate immunity in these hosts These are complemented with a series of chapters dealing with alternative models plants worms mosquitoes flies and fish Together these approaches and models are being used to dissect the complex interplay between hosts and pathogens and contribute to developing strategies to help fight infection With chapters written by experts on the cutting edge of this technology Innate Immunity is an essential reference for immunologists histologists geneticists and molecular biologists

Stem Cell Assays Mohan C. Vemuri, 2007-08-10 The scope for improving health care using stem cell therapies is thrilling but has considerable technical

challenges and methodological constraints that need to be addressed Keeping with the tradition of Humana Press to bring these developments to the forefront in a timely manner this book presents scientific advances in stem cell methods for a wider use by novice and expert scientists through the series of Methods in Molecular Biology *Macromolecular Crystallography Protocols, Volume 2* Sylvie Doublié, 2008-02-05 In the decade since publication of the first edition of *Crystallographic Methods and Protocols* the field has seen several major developments that have both accelerated the pace of structure determination and made crystallography accessible to a broader range of investigators Volume I Preparation and Crystallization of Macromolecules is dedicated to the crystallization and ways to increase the odds of obtaining crystals in macromolecules while Volume 2 Structure Determination covers both computational methods for characterizing crystals and solving structures *Post-Transcriptional Gene Regulation* Jeffrey Wilusz, 2008 Step by step instructions that ensure successful results **Avidin-Biotin Interactions** Robert J. McMahon, 2008-02-26 This book is specifically about the application of the extremely powerful interaction between the protein avidin or its homologues and the vitamin biotin and some of its homologues With excellent descriptions of laboratory protocols written by expert researchers this volume is equally perfect for the student or the professional laboratory scientist

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