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VOLUME 192

PCR Cloning Protocols

Second Edition

Edited by

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Harry W. Janes**



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Pcr Cloning Protocols Methods In Molecular Biology Vol 192

John M. S. Bartlett, David Stirling



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Translational and Experimental Clinical Research Daniel P. Schuster, William J. Powers (MD.), 2005 This volume is a comprehensive textbook for investigators entering the rapidly growing field of translational and experimental clinical research The book offers detailed guidelines for designing and conducting a study and analyzing and reporting results and discusses key ethical and regulatory issues Chapters address specific types of studies such as clinical experiments in small numbers of patients pharmacokinetics and pharmacodynamics and gene therapy and pharmacogenomic studies A major section describes modern techniques of translational clinical research including gene expression identifying mutations and polymorphisms cloning transcriptional profiling proteomics cell and tissue imaging tissue banking evaluating substrate metabolism and in vivo imaging

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

Functional Genomics Michael J. Brownstein, Arkady Khodursky, 2008-02-03 This collection of robust readily reproducible methods for microarray based studies includes expert guidance in the optimal data analysis and informatics On the methods side are proven techniques for monitoring subcellular RNA localization en masse for mapping chromosomes at the resolution of a single gene and for surveying the steady state genome wide distribution of DNA binding proteins in vivo For those workers dealing with massive data sets the book discusses the methodological aspects of data analysis and informatics in the design of microarray experiments the choice of test statistic and the assessment of observational significance data reduction and clustering

Cancer Cytogenetics John Swansbury, 2008-02-05 A collection of key cytogenetic and FISH techniques used by modern clinical laboratories in the genetic analysis of human malignancies The book's practical advice and methods are suitable for use at every level of expertise including fully established laboratories but with a sympathetic bias towards anyone considering setting up a new cytogenetics service Here the reader will find not only elementary tutorials on the fundamentals of human karyotypes and chromosome analysis but also detailed discussions on how laboratories may optimally upgrade their repertoire of capabilities to include such newer complementary techniques as CGH FISH and M FISH

Subject Guide to Books in Print, 1996

Neurogenetics Nicholas T. Potter, 2008-02-02 The rapid identification and characterization of genes of neurological relevance holds great potential for offering insight into the diagnosis management and understanding of the pathophysiologic mechanisms of neurological diseases This volume in the Methods in Molecular Biology™ series was conceived to highlight many of the contemporary methodological approaches utilized for the characterization of neurologically relevant gene mutations and their protein products Although an emphasis has been placed upon descriptions of methodologies with a defined clinical utility it is hoped that Neurogenetics Methods and Protocols will appeal not only to clinical laboratory diagnosticians but also to clinicians and to biomedical researchers with an interest in advances in disease diagnosis and the functional consequences of neurologically relevant gene mutations To meet this challenge more than 60 authors graciously accepted my invitation to contribute to the 32 chapters of this book Through their collective commitment and diligence what has emerged is a comprehensive and timely treatise that covers many methodological aspects of mutation detection and screening including discussions on quantitative PCR trinucleotide repeat detection sequence based mutation detection molecular detection of imprinted genes fluorescence in situ hybridization FISH in vitro protein expression systems and studies of protein expression and function I would like to take this opportunity to formally thank my colleagues for their effort and dedication to this work

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 Mutation Detection Protocols Bimal D. M. Theophilus, Ralph Rapley, 2008-02-02 Bimal D Theophilus and Ralph Rapley provide biological and clinical investigators with a comprehensive collection of new recent and updated PCR based screening methods suitable for detecting the presence of both known and novel mutations The methods cover point mutations e g ASO PCR SSCP DGGE chemical cleavage deletions multiplex PCR FISH blotting non sense mutations PTT and more The new and exciting techniques of DNA array analysis along with such

recently developed experimental methods as conformation sensitive gel electrophoresis are also included Each chapter explains the basic theory behind the technique and provides valuable notes essential for its successful execution

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

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