

Methods in
Molecular Biology 925

Springer Protocols



Nora Engel *Editor*

Genomic Imprinting

Methods and Protocols

 Humana Press

Genomic Imprinting Methods And Protocols Methods In Molecular Biology

Wolfgang Guggemos



Genomic Imprinting Methods And Protocols Methods In Molecular Biology:

Genomic Imprinting Nora Engel,2012-08-21 Genomic imprinting the process by which the non equivalence of the paternal and maternal genomes is established has been fascinating us for over three decades and has provided many emerging scientists with the chance to hit their stride in a frontier posing many unexpected questions and even more surprising answers In *Genomic Imprinting Methods and Protocols* experts in the field provide a survey of the technologies that are being applied to advance the study of imprinting This detailed volume features new technologies that are accelerating the pace of discovery of imprinted genes and characterization of their epigenetic profile bioinformatic procedures for prediction and comparative analyses of imprinted genes as well as methods in embryology and basic molecular biology that have been employed for many years some appearing in new versions for small cell numbers Written in the highly successful *Methods in Molecular Biology*TM 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 Authoritative and easy to use *Genomic Imprinting Methods and Protocols* will aid scientists in unveiling both much awaited answers and all new questions to keep this vital field busy for many exciting years to come

Methods in Molecular Biology: Genomic imprinting: methods and protocols John M. Walker,1984

Genomic Imprinting Andrew Ward,2001-10-10 Genomic imprinting is the process by which gene activity is regulated according to parent of origin Usually this means that either the maternally inherited or the paternally inherited allele of a gene is expressed while the opposite allele is repressed The phenomenon is largely restricted to mammals and flowering plants and was first recognized at the level of whole genomes Nuclear transplantation experiments carried out in mice in the late 1970s established the non equivalence of the maternal and paternal genomes in mammals and a similar conclusion was drawn from studies of interploidy crosses of flowering plants that extend back to at least the 1930s Further mouse genetic studies involving animals carrying balanced translocations reviewed in Chapter 3 indicated that imprinted genes were likely to be widely scattered and would form a minority within the mammalian genome The first imprinted genes were identified in the early 1990s over forty are now known in mammals and the list continues steadily to expand

PCR Cloning Protocols Bing-Yuan Chen,Harry W. Janes,2008-02-05 *PCR Cloning Protocols* Second Edition updates and expands Bruce White's best selling *PCR Cloning Protocols* 1997 with the newest procedures for DNA cloning and mutagenesis Here the researcher will find readily reproducible methods for all the major aspects of PCR use including PCR optimization computer programs for PCR primer design and analysis and novel variations for cloning genes of special characteristics or origin with emphasis on long distance PCR and GC rich template amplification Also included are both conventional and novel enzyme free and restriction site free procedures to clone PCR products into a range of vectors as well as state of the art protocols to facilitate DNA mutagenesis and recombination and to clone the challenging uncharacterized DNA flanking a known DNA fragment

PCR Detection of Microbial Pathogens Konrad Sachse, Joachim Frey, 2003 Hands on laboratory experts present a set of classic PCR based methods for the identification and detection of important animal and food microbial pathogens including several zoonotic agents These proven techniques can be precisely applied to a wide variety of microbes among them *Campylobacter* spp *Chlamydiae* *Toxigenic Clostridia* *Escherichia coli* STEC *Listeria monocytogenes* *Mycoplasmas* *Salmonellae* and *Yersinia enterocolitica* Additional chapters review the specificity and performance of diagnostic PCR analysis the pre PCR processing of samples the critical aspects of standardizing PCR methods and the general issues involved in using PCR technology for microbial diagnosis

Biostatistical Methods Stephen W. Looney, 2008-02-03 Leading biostatisticians and biomedical researchers describe many of the key techniques used to solve commonly occurring data analytic problems in molecular biology and demonstrate how these methods can be used in the development of new markers for exposure to a risk factor or for disease outcomes Major areas of application include microarray analysis proteomic studies image quantitation genetic susceptibility and association evaluation of new biomarkers and power analysis and sample size

Posttranslational Modification of Proteins Christoph Kannicht, 2008-02-04 Christoph Kannicht and a panel of highly experienced researchers describe readily reproducible methods for detecting and analyzing the posttranslational modifications of protein particularly with regard to protein function proteome research and the characterization of pharmaceutical proteins

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

Tumor Suppressor Genes Wafik S. El-Deiry, 2008-02-03 It has become clear that tumors result from excessive cell proliferation and a corresponding reduction in cell death caused by the successive accumulation of mutations in key regulatory target genes over time During the 1980s a number of oncogenes were characterized whereas from the 1990s to the present the emphasis has shifted to tumor suppressor genes TSGs It has become clear that oncogenes and TSGs function in the same pathways providing positive and negative growth regulatory activities The signaling pathways controlled by these genes involve virtually every process in cell biology including nuclear events cell cycle cell death cytoskeletal cell membrane angiogenesis and cell adhesion effects Mutations in tumor suppressor genes have been identified in familial cancer syndromes and the same genes in many cases have been found to be mutationally inactivated in sporadically occurring cancers In their normal state TSGs control cancer development and progression as well as contribute to the sensitivity of cancers to a variety of therapeutics Understanding the classes of TSGs the biochemical pathways they function in and how they are regulated

provides an essential lesson in cancer biology We cannot hope to advance our current knowledge and to develop new and more effective therapies without understanding the relevant pathways and how they influence the present approaches to therapy Moreover it is important to be able to access not only the powerful tools now available to discover these genes but also their links to cell biology and growth control Cardiac Cell and Gene Transfer Joseph M. Metzger,2008-02-03 Heart disease is the leading cause of death in developed countries Recent experimental advances featuring cellular molecular and genetic tools and technologies offer the potential for new therapeutic strategies directed toward remediation of inherited and acquired heart diseases Whether these recent basic science advances will ultimately translate to clinical efficacy for patients with heart disease is unknown and is important to ascertain Cardiac Cell and Gene Transfer Principles Protocols and Applications is designed to provide the reader with up to date coverage of a myriad of specific methodologies and protocols for gene and cell transfer to the myocardium Each chapter features a Notes section that provides useful how to problem solving insights that are often left unstated in standard published protocols Cardiac Cell and Gene Transfer Principles Protocols and Applications addresses principles and applications of cell and gene transfer to the heart including protocols for vector production and purification Detailed step by step methods and applications for first second generation adenoviral vectors adeno associated vectors gutted adenoviral vectors and lentiviral vectors are included Additionally detailed methods for cardiac cell grafting and transplantation are provided and these chapters highlight the prospects of cell based therapies for cardiac repair The book also covers specific in vivo techniques for cardiac gene transfer and specifies subsequent cellular and organ level physiological assessment techniques and protocols Accordingly this book is designed for basic science and clinical researchers in the academic pharmaceutical and biotechnology sectors of the cardiovascular community Capillary Electrophoresis of Carbohydrates Pierre Thibault,Susumu Honda,2008-02-03 A collection of cutting edge techniques for using capillary electrophoresis CE to analyze complex carbohydrates These readily reproducible protocols provide methods for sample preparation analysis of mono and oligosaccharides glycoproteins and glycoconjugates A useful appendix describes the structures of the most commonly encountered carbohydrate residues and oligosaccharides from mammalian and bacterial origins Each protocol contains detailed information on reagents apparatus notes comments and tips on procedures

Biochemicals and Reagents for Life Science Research Sigma Chemical Company,1999 Genome Research ,2007

Plant Epigenetics and Epigenomics Charles Spillane,Peter C. McKeown,2014 Many fundamental discoveries concerning epigenetics and the elucidation of mechanisms of epigenetic regulation have developed from studies performed in plants In Plant Epigenetics and Epigenomics Methods and Protocols leading scientists in the epigenetics field describe comprehensive techniques that have been developed to understand the plant epigenetic landscape These include recently developed methods and techniques for analysis of epigenetically regulated traits such as flowering time transposon activation genomic imprinting and genome dosage effects 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 key tips on troubleshooting and avoidance of known pitfalls Authoritative and practical Plant Epigenetics and Epigenomics Methods and Protocols seeks to aid scientists in the further study of plant epigenetic phenomena using advanced contemporary methods **The British National Bibliography** Arthur James Wells,2003

American Book Publishing Record ,2006 *Nucleic Acids Abstracts* ,1996 *Current Protocols in Immunology* John E. Coligan,1991 *Current Protocols in Immunology* is a three volume looseleaf manual that provides comprehensive coverage of immunological methods from classic to the most cutting edge including antibody detection and preparation assays for functional activities of mouse and human cells involved in immune responses assays for cytokines and their receptors isolation and analysis of proteins and peptides biochemistry of cell activation molecular immunology and animal models of autoimmune and inflammatory diseases Carefully edited step by step protocols replete with material lists expert commentaries and safety and troubleshooting tips ensure that you can duplicate the experimental results in your own laboratory Bimonthly updates which are filed into the looseleaf keep the set current with the latest developments in immunology methods The initial purchase includes one year of updates and then subscribers may renew their annual subscriptions *Current Protocols* publishes a family of laboratory manuals for bioscientists including Molecular Biology Human Genetics Protein Science Cytometry Cell Biology Neuroscience Pharmacology and Toxicology *Forthcoming Books* Rose Arny,2002-02 **Genetics Abstracts** ,2001

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