Methods in Molecular Biology™

VOLUME 137

Developmental Biology Protocols

Volume III

Edited by

Rocky S. Tuan Cecilia W. Lo



<u>Developmental Biology Protocols Volume Iii Methods In</u> <u>Molecular Biology</u>

Neil Osheroff, Mary-Ann Bjornsti

Developmental Biology Protocols Volume Iii Methods In Molecular Biology:

Developmental Biology Protocols Rocky S. Tuan, Cecilia W. Lo, 2000-01-21 This three volume set consisting of 142 chapters is intentionally broad in scope because of the nature of modern developmental biology **Developmental Biology Protocols** Rocky S. Tuan, Cecilia W. Lo, 2008-02-05 Developmental biology is one of the most exciting and fast growing fields today In part this is so because the subject matter deals with the innately fascinating biological events changes in form structure and function of the org ism The other reason for much of the excitement in developmental biology is that the field has truly become the unifying melting pot of biology and provides a framework that integrates anatomy physiology genetics biochemistry and cellular and mole lar biology as well as evolutionary biology No longer is the study of embryonic development merely embryology In fact development biology has produced portant paradigms for both basic and clinical biomedical sciences alike Although modern developmental biology has its roots in experimental emb ology and the even more classical chemical embryology the recent explosive and remarkable advances in developmental biology are critically linked to the advent of the cellular and molecular biology revolution. The impressive arsenal of experient and analytical tools derived from cell and molecular biology which promise to continue to expand together with the exponentially developing sophistication in fu tional imaging and information technologies guarantee that the study of the devel ing embryo will contribute one of the most captivating areas of biological research in the next millennium Molecular Methods in Developmental Biology Matt Guille, 2008-02-03 The process whereby a single cell the fertilized egg develops into an adult has fascinated for centuries Great progress in understanding that process h ever has been made in the last two decades when the techniques of molecular biology have become available to developmental biologists By applying these techniques the exact nature of many of the interactions responsible for forming the body pattern are now being revealed in detail Such studies are a large and it seems ever expanding part of most life science groups It is at newcomers to this field that this book is primarily aimed A number of different plants and animals serve as common model org isms for developmental studies In Molecular Methods in Developmental Bi ogy Xenopus and Zebrafish a range of the molecular methods applicable to two of these organisms are described these are the South African clawed frog Xenopus laevis and the zebrafish Brachydanio rerio The embryos of both of these species develop rapidly and externally making them particularly suited to investigations of early vertebrate development However both Xenopus and zebrafish have their own advantages and disadvantages Xenopus have large robust embryos that can be manipulated surgically with ease but their pseudotetraploidy and long generation time make them unsuitable candidates for genetics This disadvantage may soon be overcome by using the diploid Xenopus tropicalis and early experiments are already underway The transp ent embryos of zebrafish render them well suited for in situ hybridization and immunohistochemistry and good for observing mutations in genetic screens Developmental Biology Protocols Rocky S. Tuan, Cecilia W. Lo, 2000 Annotation This book combines expert reviews of the best contemporary methods for studying the disease process in ovarian cancer with detailed technical protocols for its analysis The book focuses on the major problems still facing researchers on reproducible experimental methods to illuminate those problems and on gene therapeutic approaches to its treatment The ELISA Guidebook John R. Crowther, 2008-02-04 John R Crowther provides today's premier practical guide to the understanding and application of ELISA Updating and greatly expanding his widely appreciated earlier publication ELISA Theory and Practice 1995 this important work introduces chapters on such major new topics as checkerboard titrations quality control of testing kit production and control novel monoclonal antibodies validation of assays statistical requirements for data examination and epidemiological considerations With its numerous worked examples detailed instructions and extensive illustrations The ELISA Guidebook offers a powerful synthesis of all the basic concepts and practical experimental details investigators need to understand develop and apply the new ELISA methodology successfully in day to day basic and clinical research Chromatin Protocols Peter B. Becker, 2008-02-03 More than 40 years after the discovery of the nucleosome as the fun mental unit of chromatin the multifaceted problem of how variations in ch matin structure affect the activity of the eukaryotic genome has not been solved However during the past few years research on chromatin structure and fu tion has gained considerable momentum and impressive progress has been made at the level of concept development as well as filling in crucial detail The structure of the nucleosome has been visualized at unprecedented reso tion Powerful multisubunit enzymes have been identified that alter histone DNA interactions in ways that expose regulatory sequences to factors initi ing and regulating such nuclear processes as transcription Though the imp tance of posttranslational modifications of histones notably their acetylation has long been known the finding that a number of bona fide regulators increase transcription by acetylating nucleosomes has lent new support to the old idea that the process of gene regulation is intimately related to the nature of the chromatin environment A wealth of nonhistone proteins contribute to a continuum of structures with distinct biochemical properties and varying degrees of DNA condensation Perhaps the most important conclusion from a large number of studies is a fresh appreciation of the dynamic nature of chromatin structure the built in flexibility providing the basis for regulation **Chaperonin Protocols** Christine Schneider.2008-02-05 **Neurotrophin Protocols** Robert A. Rush, 2008-02-03 The past decade has seen an extraordinary growth in research interest in neurotrophic factors and the study of the neurotrophin family has led this activity Nevertheless this area of research has often struggled as a result of techniques that were either inadequate or just emerging from other research fields and disciplines Neurotrophin Protocols has brought together many leaders in the neurotrophin field who detail their special expertise in a wide variety of techniques Though most procedures are valid across many diff ent fields of research some of those described here have been developed to address particular issues within the neurotrophic factor field The protocols cover a broad range of biochemical histological and biological techniques that are often required by the modern laboratory However all have been written with sufficient detail to allow any laboratory to achieve proficiency without

need of reference to other texts Neurotrophin Protocols is divided into four sections dealing with p tein RNA recombinant and in vivo techniques Protein techniques have in general been less successfully employed than those dealing with RNA or DNA However procedures that achieve localization and quantification of the neurotrophins are now being used more extensively Their inclusion here should assist further studies at the protein level Transgenic cell lines and animals are commonplace in the scientific research literature but their inc sion in several chapters in this book provide some novel uses that are not readily available elsewhere Calpain Methods and Protocols John S. Elce, 2008-02-05 The purpose of Calpain Methods and Protocols is guite straightf ward it is to present the actual experimental methods used in many different laboratories for the study of calpain It will provide the vital experimental detail and the discussion of possible pitfalls for which the standard journals no longer provide space This will make it as easy as possible for investi tors interested in calpain to adopt established methods without repeating old mistakes and to adapt and apply these methods in novel approaches to the many outstanding calpain questions. These questions range from purely biochemical problems of protein structure and enzyme regulation at the molecular level through large areas of cell biology to applied and clinical aspects of calpain function in human dease Within this panoply of topics a wide range of investigators will find many fascinating and as yet unanswered questions about calpain Calpain Methods and Protocols will provide instant access to many essential te niques while saving them the time and effort involved in developing a new method In addition to guestions relating to the normal physiological roles of the calpains there is considerable evidence that inappropriate calpain activity may have pathological effects in many tissues for example following ischemia This provides a major stimulus for the development of specific calpain inhi tors for therapeutic purposes and for the development of methods to evaluate such inhibitors DNA Topoisomerase Protocols Neil Osheroff, Mary-Ann Bjornsti, 2008-02-05 Beginning with the Escherichia coli protein or bacterial DNA topoisomerase I an ever increasing number of enzymes have been identified that catalyze changes in the linkage of DNA strands DNA topoisomerases are ubiquitous in nature and have been shown to play critical roles in most p cesses involving DNA including DNA replication transcription and rec bination These enzymes further constitute the cellular targets of a number of clinically important antibacterial and anticancer agents Thus further studies of DNA topology and DNA topoisomerases are critical to advance our und standing of the basic biological processes required for cell cycle progression cell division genomic stability and development In addition these studies will continue to provide critical insights into the cytotoxic action of drugs that target DNA topoisomerases Such mechanistic studies have already played an important role in the development and clinical application of antimicrobial and chemotherapeutic agents The two volumes of DNA Topoisomerase Protocols are designed to help new and established researchers investigate all aspects of DNA topology and the function of these enzymes The chapters are written by prominent investigators in the field and provide detailed background information and st by step experimental protocols The topics covered in Part I DNA Topology and Enzymes range from detailed methods to analyze

various aspects of DNA structure from linking number knotting unknotting site specific recombi tion and decatenation to the overexpression and purification of bacterial and eukaryotic DNA topoisomerases from a variety of cell systems and tissues

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