

Calcium Signaling Protocols

Second Edition

Edited by

David G. Lambert



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Calcium Signaling Protocols Methods In Molecular Biology Vol 312

Jessica J Manson



Calcium Signaling Protocols Methods In Molecular Biology Vol 312:

Calcium Signaling Protocols David G. Lambert, 2008-02-04 In the first edition of Calcium Signaling Protocols I began by writing The regulation of intracellular Ca is a common theme presented in many papers over the last 20 or so years and the description of the Ca sensitive indicator dye fura 2 in 1985 resulted in a massive increase in these types of studies This statement is as true in 2005 as it was in 1999 but 20 or so years is now 30 years There has been some reorganization of the volume such that there are now 22 chapters including five new ones all written by experts in their field These new chapters include use of the FlexStation and electrophysiological measurement of Ca channel activity The book is broken into six parts Part I is a general coverage of basic theory and the simplest use of fluorescent indicators Part II covers specialist measurement systems and Part III covers measurement of Ca channel activity Assessment of release of stored Ca is covered in some detail in Part IV with Parts V and VI covering specialist measurement techniques and Ca sensitive targets Putting a book like this together even as a second edition takes time and I am again indebted to the individual authors for their help and patience I am also very grateful to Professor John M Walker the series editor for his continued help and advice over the course of this project

Arabidopsis Protocols, 2nd Edition Julio Salinas, Jose J. Sanchez-Serrano, 2008-02-04 For several decades *Arabidopsis thaliana* has been the organism of choice in the laboratories of many plant geneticists physiologists developmental biologists and biochemists around the world During this time a huge amount of knowledge has been acquired on the biology of this plant species which has resulted in the development of molecular tools that account for much more efficient research The significance that *Arabidopsis* would attain in biological research may have been difficult to foresee in the 1980s when its use in the laboratory started In the meantime it has become the model plant organism much the same way as *Drosophila* *Caenorhabditis* or mouse have for animal systems Today it is difficult to envision research at the cutting edge of plant biology without the use of *Arabidopsis* Since the first edition of *Arabidopsis Protocols* appeared new developments have fostered an impressive advance in plant biology that prompted us to prepare *Arabidopsis Protocols Second Edition* Completion of the *Arabidopsis* genome sequence offered for the first time the opportunity to have in hand all of the genetic information required for studying plant function In addition the development of whole systems approaches that allow global analysis of gene expression and protein and metabolite dynamics has encouraged scientists to explore new scenarios that are extending the limits of our knowledge

Ubiquitin-Proteasome Protocols Cam Patterson, Douglas M. Cyr, 2008-02-04 A collection of cutting edge techniques for studying ubiquitin dependent protein degradation via the proteasome The topics covered range broadly from basic biochemistry to cellular assays to discovery techniques using mass spectrometric analysis These biochemical and cellular methods are necessary to explore the ubiquitin proteasome system and ubiquitin proteasome dependent functions State of the art and user friendly *Ubiquitin Proteasome Protocols* offers novice and experienced bench scientists alike a thorough compendium of readily reproducible techniques that will accelerate

discovery enhance productivity and permit manipulation of the system for varied research purposes *DNA Repair Protocols* Daryl S. Henderson, 2008-02-03 The first edition of this book published in 1999 and called DNA Repair Protocols Eukaryotic Systems brought together laboratory based methods for studying DNA damage and repair in diverse eukaryotes namely two kinds of yeast a nematode a fruit fly a toad three different plants and human and murine cells This second edition of DNA Repair Protocols covers mammalian cells only and hence its new subtitle Mammalian Systems There are two reasons for this fresh emphasis both of them pragmatic to cater to the interests of what is now a largely mammalocentric DNA repair field and to expedite editing and production of this volume Although DNA Repair Protocols Mammalian Systems is a smaller book than its predecessor it actually contains a greater variety of methods Fourteen of the book's thirty two chapters are entirely new and areas of redundancy present in the first edition have been eliminated here for example now just two chapters describe assays for nucleotide excision repair NER rather than seven All eighteen returning chapters have been revised many of them extensively In order to maintain a coherent arrangement of topics the four part partitioning seen in the first edition was dispensed with and chapters concerned with ionizing radiation damage and DNA strand breakage and repair were relocated to near the front of the book Finally an abstract now heads each chapter *Cytochrome P450 Protocols* Ian R. Phillips, 2008-02-04 For this second edition of their much praised Cytochrome P450 the editors have collected accounts of the essential core techniques that use the latest methodologies for the investigation of P450s Highlights include protocols for spectral analysis and purification of P450s enzymatic assays of P450s and flavin containing monooxygenases FMOs expression of P450s and FMOs in heterologous systems and the production and use of anti-peptide antibodies Additional chapters contain readily reproducible techniques for the transfection of hepatocytes for gene regulation studies P450 reporter gene assays in situ hybridization and analysis of genetic polymorphisms Although the emphasis is on P450s of mammalian origin many of the readily reproducible methods described are suitable for P450s from any source

Differential Display Methods and Protocols Peng Liang, Jonathan Meade, Arthur B. Pardee, 2008-02-04 Since the first edition of this book dedicated to differential display DD technology was published in 1997 we have witnessed an explosive interest in studying differential gene expression The gene hunting euphoria was initially powered by the invention of DD which was gradually overtaken by DNA microarray technology in recent years Then why is there still the need for second edition of this DD book First of all DD still enjoys a substantial lead over DNA microarrays in the ISI citation data see Table 1 despite the hundreds of millions of dollars spent each year on arrays This may come as a surprise to many but to us it implies that many of the DNA microarray studies went unpublished owing to their unfulfilled promises 1 Second unlike DNA microarrays DD is an open ended gene discovery method that does not depend on prior genome sequence information of the organism being studied As such DD is applicable to the study of all living organisms from bacteria fungi insects fish plants to mammals even when their genomes are not sequenced Second DD is more accessible technically and financially to most cost

conscious cottage industry academic laboratories So clearly DD still has its unique place in the modern molecular biological toolbox for gene expression analysis

NanoBiotechnology Protocols Sandra J Rosenthal, David W Wright, 2008-02-04 Hands on experts in nanomaterial synthesis and application describe in detail the key experimental techniques currently employed in novel materials synthesis dynamic cellular imaging and biological assays The author s emphasize diverse strategies to synthesize and functionalize the use of nanoparticles for biological applications Additional chapters focus on the use of biological components peptides antibodies and DNA to synthesize and organize nanoparticles to be used a building block in larger assemblies These new materials make it possible to image cellular processes for longer durations leading to high throughput cellular based screens for drug discovery drug delivery and diagnostic applications Highlights include overview chapters on quantum dots and DNA nanotechnology and cutting edge techniques in the emerging nanobiotachnology arena

Human Retrovirus Protocols Tuofu Zhu, 2008-02-04 A cutting edge collection of basic and state of the art methods optimized for investigating the molecular biology of this class of retrovirus These readily reproducible techniques range from methods for the isolation and detection of human retroviruses to cutting edge methods for exploring the interplay between the viruses and the host Here the researcher will find up to date techniques for the isolation and propagation of HIV HTLV and foamy virus from a variety of sources There are also assays for determining the cell tropism of HIV 1 the coreceptor usage of HIV 1 and human gene expression with HIV 1 infection by microarrays as well as for phenotyping HIV 1 infected monocytes and examining their fitness Highlights include the detection and quantification of HIV 1 in resting CD4 a new cloning system for making recombinent virus cDNA microarrays and the determination of genetic polymorphisms in two recently identified HIV 1 co factors that are critical for HIV 1 infection

Phosphodiesterase Methods and Protocols Claire Luginier, 2008-02-04 Research leaders in the PDE field describe new concepts and techniques for investigating the role of PDEs in orchestrating normal and pathophysiological responses Presented in step by step detail these readily reproducible methods allow the measurement of cyclic nucleotide variations in living cells as well as their visualization in a spatio temporal manner the localization and characterization of their activities in tissues and living cells and the assessment of targeted PDEs in creating specific tools and drugs

Transmembrane Signaling Protocols Hydar Ali, Haribabu Bodduluri, 2008-02-05 The previous edition of Transmembrane Signaling Protocols was published in 1998 Since then the human genome has been completely sequenced and new methods have been developed for the use of microarrays and proteomics to analyze global changes in gene expression and protein profiles These advances have increased our ability to understand transmembrane signaling processes in much greater detail They have also simultaneously enhanced our ability to determine the role of a large number of newly identified molecules in signaling events In addition novel video microscopy methods have been developed to image transmembrane signaling events in live cells in real time In view of these major advances it is time to update the previous edition Because of the success of that volume we have chosen to keep the essential

character of the book intact Introductory chapters from experts have been included to provide overall perspective and an overview of recent advances in signal transduction pathways The individual chapters now include comprehensive detailed methods studies in genetically tractable systems fluorescence microscopy in live single cells ex vivo analysis of primary cells from transgenic mice as well as genomic and proteomic approaches to the analysis of transmembrane signaling events We would like to express our deep gratitude to the coauthors of this publication We hope that Transmembrane Signaling Protocols Second Edition will serve as a valuable resource for future progress in the study of signal transduction pathways

Human Embryonic Stem Cell Protocols KURSAD TURKSEN, 2008-02-04 A comprehensive collection of diverse techniques for the molecular and cellular manipulation of human embryonic stem hES cells These readily reproducible methods have been optimized for the derivation characterization and differentiation of hES cells with special attention given to regenerative medicine applications A companion CD provides color versions of all illustrations in the book The protocols follow the successful Methods in Molecular Biology™ series format each offering step by step laboratory instructions an introduction outlining the principles behind the technique lists of the necessary equipment and reagents and tips on troubleshooting and avoiding known pitfalls

Receptor Binding Techniques Anthony P. Davenport, 2008-02-02 A comprehensive collection of readily reproducible methods for studying receptors in silico in vitro and in vivo These cutting edge techniques cover mining from curated databases identifying novel receptors by high throughput screening molecular methods to identify mRNA encoding receptors radioligand binding assays and their analysis quantitative autoradiography and imaging receptors by positron emission tomography PET Highlights include phenotypic characterization of receptors in knockout mice imaging receptors using green fluorescent protein and fluorescent resonance energy transfer and quantitative analysis of receptor mRNA by TaqMan PCR These book equips the researcher with techniques for exploring the unprecedented number of new receptor systems now emerging and the so called orphan receptors whose activating ligand has not been identified

Epidermal Growth Factor Tarun B. Patel, Paul J. Bertics, 2008-02-05 A comprehensive collection of optimized methods for dissecting the mechanisms that control epidermal growth factors EGF and their regulators in both normal and pathological states These readily reproducible techniques range from the study of purified EGF receptor to complex signaling and processing networks in intact cells including a chapter on the clinical and pharmacological considerations of their use in cancer therapy The protocols follow the successful Methods in Molecular Biology™ series format each offering step by step laboratory instructions an introduction outlining the principles behind the technique lists of the necessary equipment and reagents and tips on troubleshooting and avoiding known pitfalls

Cell Imaging Techniques Douglas J. Taatjes, Brooke T. Mossman, 2008-02-04 A diverse collection of state of the art methods for the microscopic imaging of cells and molecules The authors cover a wide spectrum of complementary techniques including such methods as fluorescence microscopy electron microscopy atomic force microscopy and laser scanning cytometry Additional readily reproducible protocols on confocal

scanning laser microscopy quantitative computer assisted image analysis laser capture microdissection microarray image scanning near field scanning optical microscopy and reflection contrast microscopy round out this eclectic collection of cutting edge imaging techniques now available The authors also discuss preparative methods for particles and cells by transmission electron microscopy

Xenopus Protocols X. Johné Liu, 2008-02-02 A collection of standard and cutting edge techniques for using *Xenopus* oocytes and oocytes egg extracts to reconstitute biological and cellular processes These readily reproducible methods take advantage of the oocyte s impressive protein abundance its striking protein translation capacity and its breathtaking possibilities for the assembly of infectious viral particles by single cell injection of multiple RNAs The authors focus on the versatility of frog oocytes and egg extracts in cell biology and signal transduction and cover all the major uses of oocytes extracts as experimental models

Nuclear Reprogramming Steve Pells, 2008-02-04 A wide ranging collection of readily reproducible methods for performing nuclear reprogramming by nuclear transfer in several different species by fusion through both chemical treatment and electrically shocking cells and by in vivo treatment of cells with cell extracts Several methods of monitoring nuclear reprogramming are also presented including the use of transgenic markers activation of telomerase as an ES specific marker light and electron microscopic observation of structural changes in the nucleus and verification of surface marker expression and the differentiation potential of stem cells Biochemical methods are provided for the examination of chromatin protein modifications nucleosomal footprinting transcription factor binding and the study of DNA methylation changes both at the specific locus level and at the level of the whole nucleus

PRINS and In Situ PCR Protocols Franck Pellestor, 2008-02-03 The in situ hybridization and PCR technologies are now well established molecular techniques for studying chromosomal aneuploidy and rearran ments gene localization and expression and genomic organization Over the last decade we have seen increasing applications in these fields By combining the high sensitivity of the PCR reaction and the cytological localization of target sequences both PRINS and in situ PCR techniques have provided highly powerful complements to FISH for in situ cellular and molecular investigations Both these approaches have several advantages in terms of sensitivity and specificity owing to the use of primers and to the fast kinetics of annealing and elongation reactions in situ In the first edition of PRINS and In Situ PCR Protocols edited by John R Gosden experts in the field presented in detail a variety of applications of PRINS and in situ PCR techniques in a wide range of clinical conditions Since the publication of this successful reference book there have been s nificant improvements in in situ detection techniques This completely revised and updated second edition presents a compreh sive selection of new procedures developed in the field of PRINS and in situ PCR technologies The book has two sections Part I Basic Methodology contains chapters that provide useful protocols for many variations of PRINS and in situ PCR including a new fast multicolor PRINS method and protocols for PRINS detection of unique sequences in situ

Yeast Protocols Wei Xiao, 2008-02-03 In this second edition of a widely used classic laboratory manual leading experts utilize the tremendous progress and technological

advances that have occurred to create a completely new collection of not only the major basic techniques but also advanced protocols for yeast research and for using yeast as a host to study genes from other organisms. The authors provide detailed methods for the isolation of subcellular components including organelles and macromolecules for the basic cellular and molecular analysis specific for yeast cells and for the creation of conditional mutant phenotypes that lend themselves to powerful genome manipulation. Additional protocols offer advanced approaches to study genetic interactions, DNA and chromatin metabolism, gene expression as well as the foreign genes and gene products in yeast cells.

Microfluidic Techniques Shelley D. Minteer, 2008-02-04 Hands on researchers review the principles behind successful miniaturization and describe the key techniques for miniaturizing large scale biochemical and bioanalytical methods for microchip analysis. The authors cover not only the most popular methods for the fabrication of microchips: photolithography, laser ablation and soft lithography but also microfluidic techniques for such bioanalytical assays and bioprocesses as DNA analysis, PCR, immunoassays and cell reactors. Highlights include PCR on a microchip, microscale cell culturing and the study of cellular processes on a microchip. The protocols offer step by step laboratory instructions, an introduction outlining the principles behind the technique, lists of the necessary equipment and reagents and tips on troubleshooting and avoiding known pitfalls.

In Situ Hybridization Protocols Ian A. Darby, Tim D. Hewitson, 2008-02-04 The technique of in situ hybridization in its various forms has been used routinely in many laboratories for a number of years. In the post genome era, gene arrays and proteomics have allowed us to identify hitherto unknown unrecognized pathways and mechanisms. However, rather than diminish the importance of in situ hybridization, the now widespread use of screening technologies has increased the need to temporally and spatially localize the distribution of mRNA expression. Our intention in *In Situ Hybridization Protocols* is to provide ample information for novices planning to set up the in situ hybridization technique and use it in their laboratory for the first time as well as giving updates of recent developments for those laboratories where in situ hybridization techniques are already in use. Despite its widespread significance, in situ hybridization has retained a reputation as one of the more difficult and capricious molecular biological techniques. This may in part be because of the hybrid nature of the technique which often requires a mixture of molecular biological and histological skills. The two techniques are usually taught and acquired in different streams of biological science. The step by step and detailed protocols provided in *In Situ Hybridization Protocols* by researchers active in the field should make it possible for both the molecular biologist with little experience of histology and the histologist with little experience of molecular biology to use the techniques successfully in their laboratories.

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