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Molecular Biology of The Cell Bruce Alberts, 2002

PCR Applications Michael A. Innis, David H. Gelfand, John J. Sninsky, 1999-05-11 PCR is the most powerful technique currently used in molecular biology. It enables the scientist to quickly replicate DNA and RNA on the benchtop. From its discovery in the early 80's, PCR has blossomed into a method that enables everything from ready mutation of DNA/RNA to speedy analysis of tens of thousands of nucleotide sequences daily. PCR Applications examines the latest developments in this field. It is the third book in the series, building on the previous publications PCR Protocols and PCR Strategies. The manual discusses techniques that focus on gene discovery, genomics, and DNA array technology, which are contributing factors to the now-occurring bioinformatics boom. Key Features * Focuses on gene discovery, genomics, and DNA array technology * Covers quantitative PCR techniques, including the use of standards and kinetic analysis includes statistical refinement of primer design parameters * Ilustrates techniques used in microscopic tissue samples, such as single cell PCR, whole cell PCR, laser capture microdissection, and in situ PCR Entries provide information on: * Nomenclature * Expression * Sequence analysis * Structure and function * Electrophysiology * Parmacology * Information retrieval

p53 Protocols Sumitra Deb,Swati Palit Deb,2008-02-02 Since the discovery of p53 as a tumor suppressor, numerous methods have evolved to reveal the unique structural features and biochemical functions of this protein. Several unique properties of p53 posed a challenge to understa- ing its normal function in the initial phase of its research. The low levels of p53 in normal cells, its stabilization under situations of genotoxic stress, induction of growth arrest, and apoptosis with stabilization of the protein, obstructed the visibility of its normal, unmutated function. The property of p53 that can sense a promoter and transactivate or inhibit is still not well understood. It is still not known whether it is the absence of the protein that causes tumorigenesis, or if its mutants have a dominant role in inducing cancer. p53 Protocols comprises eighteen chapters for the study of the diverse properties of p53 and related proteins. The methods included are invaluable for delineating the function of other proteins that may function as tumor suppr- sors or growth suppressors. The chapters are not presented in any schematic order, for the importance and diversity of the functions of p53 make it imp- sible to organize them suitably. We have made a sincere effort to collect the methods most useful to those investigators working on tumor suppressors or growth suppressors. The purpose of p53 Protocols is not only to provide investigators with methods to analyze similar biochemical functions, but also to familiarize them with the associated problems that arose during the course of investigations.

Chemotaxis Tian Jin, Dale Hereld, 2010-11-16 Fundamental to the development and vital functions of organisms, the migration of motile cells due to the detection of shallow gradients of specific chemical signals in their environments, or chemotaxis, can be clearly seen as a major force in cell biology. In Chemotaxis: Methods and Protocols, expert researchers in the field provide state-of-the-art methods for investigating cell migration behaviors, studying molecular components involved in detecting extracellular signals and directing cell movement, visualizing spatiotemporal dynamics of the components in signaling networks of chemotaxis in real time, and constructing quantitative models that simulate chemoattractant-induced cell responses. Written in the highly successful Methods in Molecular BiologyTM series format, chapters include introductions to their respective topics, lists of the necessary materials and reagents, step-by-step, readily reproducible laboratory protocols, and notes on troubleshooting and avoiding known pitfalls. Comprehensive and cutting-edge, Chemotaxis: Methods and Protocols serves scientists with practical guidance on the diverse methodologies that are currently propelling chemotaxis research forward in order to further our understanding of this vital biological system.

Stem Cells Ariff Bongso, Eng Hin Lee, 2011 Stem cell biology has drawn tremendous interest in recent years as it promises cures for a variety of incurable diseases. This book deals with the basic and clinical aspects of stem cell research and involves work on the full spectrum of stem cells isolated today. It also covers the conversion of stem cell types into a variety of useful tissues which may be used in the future for transplantation therapy. It is thus aimed at undergraduates, postgraduates, scientists, embryologists, doctors, tissue engineers and anyone who wishes to gain some insight into stem cell biology. This book is important as it is comprehensive and covers all aspects of stem cell biology, from basic research to clinical applications. It will have 33 chapters written by renowned stem cell scientists worldwide. It will be up-to-date and all the chapters include self-explanatory figures, color photographs, graphics and tables. It will be easy to read and give the reader a complete understanding and state of the art of the exciting science and its applications.

Biologics, **Biosimilars**, **and Biobetters** Iqbal Ramzan,2021-02-03 A comprehensive primer and reference, this book provides pharmacists and health practitioners the relevant science and policy concepts behind biologics, biosimilars, and biobetters from a practical and clinical perspective. Explains what pharmacists need to discuss the equivalence, efficacy, safety, and risks of biosimilars with physicians, health practitioners, and patients about Guides regulators on pragmatic approaches to dealing with these drugs in the context of rapidly evolving scientific and clinical evidence Balances scientific information on complex drugs with practical information, such as a checklist for pharmacists

<u>PCR Protocols</u> 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.

Production of Recombinant Proteins Gerd Gellissen, 2006-03-06 While the choices of microbial and eukaryotic expression systems for production of recombinant proteins are many, most researchers in academic and industrial settings do not have ready access to pertinent biological and technical information since it is normally scattered throughout the scientific literature. This book closes the gap by providing information on the general biology of the host organism, a description of the expression platform, a methodological section -- with strains, genetic elements, vectors and special methods, where applicable -- as well as examples of proteins produced with the respective platform. The systems thus described are well balanced by the inclusion of three prokaryotes (two Gramnegatives and one Gram-positive), four yeasts, two filamentous fungi and two higher eukaryotic cell systems --

mammalian and plant cells. Throughout, the book provides valuable practical and theoretical information on the criteria and schemes for selecting the appropriate expression platform, the possibility and practicality of a universal expression vector, and on comparative industrial-scale fermentation, with the production of a recombinant Hepatitis B vaccine chosen as an industrial example. With a foreword by Herbert P. Schweizer, Colorado State University, USA: As a whole, this book is a valuable and overdue resource for a varied audience. It is a practical guide for academic and industrial researchers who are confronted with the design of the most suitable expression platform for their favorite protein for technical or pharmaceutical purposes. In addition, the book is also a valuable study resource for professors and students in the fields of applied biology and biotechnology.

Architectural Design Portable Handbook Andy Pressman, 2001-03-15 Integrate key information to facilitate optimal design solutions Essential for any working architect, Architectural Design Portable Handbook guides you through projects every step of the way, summarizing, synthesizing, and systematizing the core tasks of design. Developed by noted architect Andy Pressman, this handy take-along reference is certain to become an indispensable tool. You'll find value-added features such as quick tips and case studies, quotes from leading architects, checklists, and a customizable layout that encourages you to add your own helpful notes and reminders. And that's in addition to expert, time- and error-sparing information on: * Design strategies * Site analysis methods * The use of CAD and other graphics * Working with clients * User-needs evaluation * Data compilation * Concept development * Exploring design alternatives * Computing and design * Presentations * More!

Nuclear Medicine Resources Manual International Atomic Energy Agency,2021-03-22 Medical imaging is crucial in a variety of medical settings and at all levels of health care. In public health and preventive medicine as well as in both curative and palliative care, effective decisions depend on correct diagnoses. This edition addresses the most current needs and offers guidance on clinical practice, radiation safety and patient protection, human resource development and training required for the overall practice of nuclear medicine.

Liquid Life Rachel Armstrong, 2019 If we lived in a liquid world, the concept of a machine would make no sense. Liquid life is metaphor and apparatus that discusses the consequences of thinking, working, and living through liquids. It is an irreducible, paradoxical, parallel, planetary-scale material condition, unevenly distributed spatially, but temporally continuous. It is what remains when logical explanations can no longer account for the experiences that we recognize as part of being alive. Liquid Life references a third-millennial understanding of matter that seeks to restore the agency of the liquid soul for an ecological era, which has been banished by reductionist, brute materialist discourses and mechanical models of life. Offering an alternative worldview of the living realm through a new materialist and liquid study of matter, Armstrong conjures forth examples of creatures that do not obey mechanistic concepts like predictability, efficiency, and rationality. With the advent of molecular science, an increasingly persuasive ontology of liquid technologies can be identified. Through the lens of lifelike dynamic droplets, the agency for these systems exists at the interfaces between different fields of matter/energy that respond to highly local effects, with no need for a central organizing system.Liquid Life seeks an alternative partnership between humanity and the natural world. It provokes a re-invention of the languages of the living realm to open up alternative spaces for exploration, including contributor Rolf Hughes' angelology of language, which explores the transformative invocations of prose poetry, and Simone Ferracina's graphical notations that help shape our concepts of metabolism, upcycling, and designing with fluids. A conceptual and practical toolset for thinking and designing, liquid life reunites us with the irreducible soul substance of living things, which will neither be simply solved, nor go away.

Basic Cell Culture Protocols Cheryl D. Helgason, Cindy L. Miller, 2016-08-23 At some point in their careers, virtually every scientist and technician, as well as many medical professionals, regardless of their area of specialization have a need to utilize cell culture systems. Updating and significantly expanding upon the previous editions, Basic Cell Culture Protocols, Fourth Edition provides the novice cell culturist with sufficient information to perform the basic techniques, to ensure the health and identity of their cell lines, and to be able to isolate and culture specialized primary cell types. The intent of this extensive volume is to generate a valuable resource containing clear methodologies pertinent to current areas of investigation, rather than attempting to educate cell culturists on specific cell types or organ systems. Written in the highly successful Methods in Molecular BiologyTM, 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. Comprehensive and up-to-date, Basic Cell Culture Protocols, Fourth Edition compiles the essential techniques needed to approach this vital laboratory activity with full success.

Gene Quantification Francois Ferre, 2012-12-06 Geneticists and molecular biologists have been interested in quantifying genes and their products for many years and for various reasons (Bishop, 1974). Early molecular methods were based on molecular hybridization, and were devised shortly after Marmur and Doty (1961) first showed that denaturation of the double helix could be reversed - that the process of molecular reassociation was exquisitely sequence dependent. Gillespie and Spiegelman (1965) developed a way of using the method to titrate the number of copies of a probe within a target sequence in which the target sequence was fixed to a membrane support prior to hybridization with the probe - typically a RNA. Thus, this was a precursor to many of the methods still in use, and indeed under development, today. Early examples of the application of these methods included the measurement of the copy numbers in gene families such as the ribosomal genes and the immunoglo bulin family. Amplification of genes in tumors and in response to drug treatment was discovered by this method. In the same period, methods were invented for estimating gene num bers based on the kinetics of the reassociation process - the so-called Cot analysis. This method, which exploits the dependence of the rate of reassociation on the concentration of the two strands, revealed the presence of repeated sequences in the DNA of higher eukaryotes (Britten and Kohne, 1968). An adaptation to RNA, Rot analysis (Melli and Bishop, 1969), was used to measure the abundance of RNAs in a mixed population.

<u>MicroRNA and Cancer</u> Wei Wu,2010-10-21 The tiny microRNAs (miRNAs) can have huge impacts on the regulation of a variety of genes and play crucial roles in the fundamental cellular processes. Recent miRNA studies change the landscape of cancer genetics by scrutinizing the alterations of genome-wide miRNA expressions in most common cancers and their regulatory functions during the development of cancer. The connections between miRNAs and cancer are widespread enough to warrant more comprehensive investigations in the systems biology perspective. In MicroRNA and Cancer: Methods and Protocols, internationally renowned experts provide the latest miRNA knowledge, the various techniques and methodologies currently available for cancer research application. Ranging from the fundamental concepts to practical applications, this book presents: • Overview of microRNA biogenesis,

computational prediction of new miRNAs in the cancer genome, and miRNA-based therapeutic approaches for cancer treatment • Detailed experimental protocols in miRNA detection with novel and high-throughput technology, miRNA library cloning, miRNA epigenetic regulation, and miRNA pathway study • Stepwise computational and bioinformatic procedures for miRNA complex networks in cancer genomes with a variety of softwares and programs • Cross-cited notes on troubleshooting and avoiding known pitfalls Authoritative and cutting-edge, MicroRNA and Cancer: Methods and Protocols serves researchers with the basic principles of experimental and computational methods for microRNA study in cancer research and provides a firm grounding for those who wish to further develop their own applications and tailor them to their own specific research needs.

NMR-Based Metabolomics G. A. Nagana Gowda, Daniel Raftery, 2020-08-29 This book provides broad coverage of nuclear magnetic resonance (NMR) spectroscopy-based methods and applications for the analysis of metabolites in a wide range of biological samples, from biofluids, cells, animal models, human, to plants and foods. The applications range from mechanistic understanding, biomarker discovery, environmental studies, and drug discovery to nutrition, while NMR methods include global, targeted, and isotope tracer-based techniques. Written for the highly successful Methods in Molecular Biology series, 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 practical, NMR-Based Metabolomics: Methods and Protocols serves as a wealth of information for beginners as well as advanced practitioners and also as stepping stones for further advances in the field of metabolomics.

An Introduction to Biotechnology W T Godbey, 2014-12-08 An Introduction to Biotechnology is a biotechnology textbook aimed at undergraduates. It covers the basics of cell biology, biochemistry and molecular biology, and introduces laboratory techniques specific to the technologies addressed in the book; it addresses specific biotechnologies at both the theoretical and application levels. Biotechnology is a field that encompasses both basic science and engineering. There are currently few, if any, biotechnology textbooks that adequately address both areas. Engineering books are equation-heavy and are written in a manner that is very difficult for the nonengineer to understand. Numerous other attempts to present biotechnology are written in a flowery manner with little substance. The author holds one of the first PhDs granted in both biosciences and bioengineering. He is more than an author enamoured with the wow-factor associated with biotechnology; he is a practicing researcher in gene therapy, cell/tissue engineering, and other areas and has been involved with emerging technologies for over a decade. Having made the assertion that there is no acceptable text for teaching a course to introduce biotechnology to both scientists and engineers, the author committed himself to resolving the issue by writing his own. The book is of interest to a wide audience because it includes the necessary background for understanding how a technology works. Engineering principles are addressed, but in such a way that an instructor can skip the sections without hurting course content The author has been involved with many biotechnologies through his own direct research experiences. The text is more than a compendium of information - it is an integrated work written by an author who has experienced first-hand the nuances associated with many of the major biotechnologies of general interest today.

Pesticide Chemistry Hideo Ohkawa, Hisashi Miyagawa, Phillip W. Lee, 2007-06-27 Resulting from the premier forum for pesticide development and use, this volume provides comprehensive coverage and even captures emerging technologies within the industry. All facets of pesticides are addressed here, including agriculture, agrochemicals, and environmental health aspects, as well as such global issues as food quality and safety.

The Opposite of Hate Sally Kohn, 2018-04-10 "A stunning debut by a truly gifted writer—an eye-opening read for both liberals and conservatives—and it could not come at a better time."—Adam Grant, New York Times bestselling author of Option B, with Sheryl Sandberg What is the opposite of hate? As a progressive commentator on Fox News and now CNN, Sally Kohn has made a career out of bridging intractable political differences and learning how to talk respectfully with people whose views she disagrees with passionately. Her viral TED Talk on the need to practice emotional-rather than political-correctness sparked a new way of considering how often we amplify our differences and diminish our connections. But these days even famously "nice" Kohn finds herself wanting to breathe fire at her enemies. It was time, she decided, to look into the epidemic of hate all around us and learn how we can stop it. In The Opposite of Hate, Kohn talks to leading scientists and researchers and investigates the evolutionary and cultural roots of hate and how incivility can be a gateway to much worse. She travels to Rwanda, the Middle East, and across the United States, introducing us to former terrorists and white supremacists, and even some of her own Twitter trolls, drawing surprising lessons from dramatic and inspiring stories of those who left hate behind. As Kohn confronts her own shameful moments, whether it was back when she bullied a classmate or today when she harbors deep partisan resentment, she discovers, "The opposite of hate is the beautiful and powerful reality of how we are all fundamentally linked and equal as human beings. The opposite of hate is connection." Sally Kohn's engaging, fascinating, and often funny book will open your eyes and your heart.

A Networked Self Zizi Papacharissi,2010-09-10 A Networked Self examines self presentation and social connection in the digital age. This collection brings together new work on online social networks by leading scholars from a variety of disciplines. The focus of the volume rests on the construction of the self, and what happens to self-identity when it is presented through networks of social connections in new media environments. The volume is structured around the core themes of identity, community, and culture — the central themes of social network sites. Contributors address theory, research, and practical implications of many aspects of online social networks including self-presentation, behavioral norms, patterns and routines, social impact, privacy, class/gender/race divides, taste cultures online, uses of social networking sites within organizations, activism, civic engagement and political impact.

The Polymerase Chain Reaction Kary B. Mullis, Francois Ferre, Richard A. Gibbs, 2012-02-02 James D. Watson When, in late March of 1953, Francis Crick and I came to write the first Nature paper describing the double helical structure of the DNA molecule, Francis had wanted to include a lengthy discussion of the genetic implications of a molecule whose struc ture we had divined from a minimum of experimental data and on theoretical argu ments based on physical principles. But I felt that this might be tempting fate, given that we had not yet seen the detailed evidence from King's College. Nevertheless, we reached a compromise and decided to include a sentence that pointed to the biological significance of the molecule's key feature-the complementary pairing of the bases. It has not escaped our notice, Francis wrote, that the specific pairing that we have postulated immediately suggests a possible copying mechanism for the genetic material. By May, when we were writing the second Nature paper, I was more confident that the proposed structure was at the very least substantially correct, so that this second paper

contains a discussion of molecular self-duplication using templates or molds. We pointed out that, as a consequence of base pairing, a DNA molecule has two chains that are complementary to each other. Each chain could then act . . . as a template for the formation on itself of a new companion chain, so that eventually we shall have two pairs of chains, where we only had one before and, moreover, ...

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