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Yohei Yamauchi, Maria João Amorim

Molecular Biology of Plant Tumors Günter Kahl, Josef S. Schell, 2014-06-28 Molecular Biology of Plant Tumors provides an opportunity to learn in detail about the latest insights into the mechanism of transformation of plant cells by *Agrobacterium tumefaciens*. The study of the molecular mechanism responsible for the crown gall phenomenon (induced by *Agrobacterium tumefaciens*) illustrates the point that the fundamental study of the cause(s) and mechanism(s) of abnormal growth might be one of the most efficient ways to understand cellular differentiation and the molecular basis of gene expression. The book is organized into three parts that contain research on abnormal plant growth, crown gall tumors, and potential vectors for genetic engineering in agriculture. The genetic structure responsible for the neoplastic transformation of plant cells in crown galls is a bacterial plasmid (called Ti for tumor-inducing). Research described in this volume demonstrates that these Ti plasmids were designed by evolution as natural gene vectors with which some bacteria can introduce active genes into plants. These transferred genes are maintained by integration in the plant genome and their expression is directly or indirectly responsible for the tumorous growth pattern.

The Biology of Seaweeds Christopher S. Lobban, Michael James Wynne, Lobban, 1981-01-01

The Evolution of Insect Life Cycles Fritz Taylor, Richard Karban, 2012-12-06 This book was developed out of a symposium at the XVII International Congress of Entomology held in Hamburg, Germany, on August 21, 1984. This symposium was organized by Drs. William Bradshaw and Hugh Dingle, who subsequently asked us to edit the proceedings. The chapters represent, for the most part, papers that were read in Hamburg but have been expanded and updated. The goal of this volume is to provide a comprehensive view of current research on insect life cycles, including field and laboratory studies, broad comparisons among species or local populations, and intensive studies of single populations, as well as theoretical research. Of necessity, given the magnitude of research now being carried out on insects, some important research programs are not included, and therein lie the makings of future volumes. This volume is divided into three parts. The first part, Geographical Patterns in Insect Life Cycles, explores various applications of a comparative method that has been valuable in investigating the potential for variability in life history parameters and the relation of these parameters to important variables in the environment.

Phospholipids Handbook Gregor Cevc, 2018-04-27 Employing a multidisciplinary approach to phospholipid research,

this work catalogues the current knowledge of this class of molecules and details the general, chemical, physical and structural properties of phospholipid monolayers and bilayers. Phospholipid applications are also covered.

Guidebook to Protein Toxins and Their Use in Cell Biology Rino Rappuoli, Cesare Montecucco, 1997-05-29 Many bacteria, animals, and plants produce toxins that can prove lethal to other organisms. Toxins are a form of biological warfare that helps their producer to survive and so confer an evolutionary advantage. They display an extraordinary range of complexity, from the formic acid provided by ants to bacterial proteins composed of thousands of amino acids. This Guidebook considers the more complex protein and peptide toxins and groups them according to their mode of action. Topics covered include: membrane-permeabilizing toxins; toxins affecting signal transduction and protein synthesis; cytoskeleton-affecting toxins; toxins affecting the immune and inflammatory response. This class of biomolecules will be of interest to a wide range of researchers in cell biology, neuroscience, and toxicology.

Plant Proteomics Ganesh K. Agrawal, Randeep Rakwal, 2008-10-03 Confidently face the challenges of proteomics research specific to plant science with the information in *Plant Proteomics*, which will introduce you to the techniques and methodologies required for the study of representative plant species. Read about proteomics studies in Arabidopsis, rice, and legumes and find information about common technologies like mass spectrometry and gel electrophoresis. Discover expression proteomics, functional proteomics, structural proteomics, bioinformatics, and systems biology, understand how to conduct proteomics studies in developing countries and underfunded laboratories, and gain access to guidelines for sample preparation.

Development V.E.A. Russo, D.J. Cove, L.G. Edgar, R. Jaenisch, F. Salamini, 2012-12-06 Studies of selected model organisms have led to a greatly improved understanding not only of the molecular events which take place during development under the control of specific sets of genes, but also of epigenetic and environmental influences on these processes. This book gives an overview of all three of these levels and their relative importance, and as such will give the reader a clear picture of one of the most exciting areas of current biological research.

Zebra Mussels Biology, Impacts, and Control Thomas F. Nalepa, Don W. Schloesser, 1992-11-24 The introduction and rapid spread of the zebra mussel in North American waters has caused great concern among industrial and recreational users of these waters. This bivalve mollusk is a biofouler that attaches to any firm substrate (e.g. rocks, piers, water intake pipes, boat hulls) and has already created significant problems for raw water users such as water treatment plants and power plants. *Zebra Mussels: Biology, Impacts and Control* provides essential information regarding the biology of the zebra mussel in North America and Europe, presents case studies of environmental and industrial impacts, and outlines control strategies. Summary articles detail its life history, origins, and morphology. The book also examines techniques used to culture and maintain this organism in the laboratory. Thirty-two color plates illustrate some of the dramatic problems created by the

explosive population growth of this species. Zebra Mussels: Biology, Impacts, and Control is an important resource for ecologists, conservationists, environmental consultants, water quality engineers, regulatory officials, power utilities, and libraries.

The Immune System F. Melchers, K. Rajewsky, 2012-12-06 The cells of the immune system generate a large variety of binding sites which differ in their binding specificities and can therefore react specifically with a large variety of ligands. These binding sites are part of receptor molecules, enabling the system to react to the universe of antigens. The classical antigen receptor is the antibody molecule, and accordingly the first session of this colloquium deals with a classical subject, namely antibody structure. Dramatic recent advances in this field make it possible to interrelate primary and three-dimensional structure both to each other and to function, i.e. the binding of antigen and possible reactions occurring in the antibody molecule upon antigen binding. The latter point is of particular interest since it may be relevant not only for effector functions of antibodies such as the binding of complement, but also for the triggering of a lymphocyte through its antibody receptor for antigen.

Molecular Genetics of Plant-Microbe Interactions Desh Pal S. Verma, N. Brisson, 2012-12-06 Increased interest in the basic biology of plants and microorganisms stems from the fact that crop productivity is directly affected by plant-microbe interactions. In spite of the fact that plants exist in the environment amongst diverse species of microorganisms, only a few ever establish a direct relationship. Emerging awareness concerning the indirect effect of microbial association on plant growth and the possibility of using one microbe against another for controlling pathogenic interactions is at the genesis of new fields of studies. The primary reason for a microbe to associate with photoautotrophic organisms (plants) is to tap its nutritional requirements, fixed carbon, as a source of energy. By hook or by crook, a microbe must survive. Some have evolved mechanisms to exploit plants to develop a niche for their biotrophic demands. When in contact with a living plant, microorganisms may live in a passive association using exudates from the plant, invade it pathogenically or coexist with it in symbiosis. The plant responds to the interloper, either reacting in a hypersensitive manner to contain the invasion of pathogens, or by inducing a set of genes that leads toward symbiosis, or by simply succumbing to the invader. Thus, prior to contact with the plant, microorganism is able to sense the presence of the host and activate accordingly a set of genes required for the forthcoming interaction, whether symbiotic or pathogenic.

The Use of Radioautography in Investigating Protein Synthesis C. P. Leblond, Katherine Brehme Warren, 1965-01-01 The Use of Radioautography in Investigating Protein Synthesis is a collection of papers that discusses the tracing of labeled proteins through photographic emulsion, particularly radioautography or autoradiography. The papers discuss proteins at the site of their synthesis, their subsequent migration through the cell, their secretions, and the formation of the basement lamella in salamanders. One paper outlines the primary aims in specimen preparation for electron microscope

autoradiographic techniques and the procedures to achieve them. Another paper reports on migrating proteins (CNP) and nonmigrating proteins of the *Amoeba proteus* nucleus in which nonmigrating proteins are sedimentable by centrifugation in situ in the nucleus. One paper investigates the extent of protein synthesis for cell renewal in thoracic duct lymphocytes (TDL) or that for export or intracellular turnover. The protein synthesized by the large cell population in TDL is associated with mitosis and cell renewal. A major portion of that synthesized by the medium cells (80 per cent) and practically all of that produced by small cells are for purposes other than cell renewal. The collection can prove beneficial to biochemists, microbiologists, bio-physicists, cytologists, histologists, pathologists, cellular researchers, and academicians involved in the study of cellular biology.

Sphingolipid Metabolism and Cell Signaling Alfred Harrison Merrill, Yusuf A. Hannun, 2000 This volume contains information on analyzing sphingolipids, sphingolipid transport and trafficking, and sphingolipid-protein interactions and cellular targets. Its companion Volume 311 presents methods used in studying enzymes of sphingolipid biosynthesis and turnover, including inhibitors of some of these enzymes, genetic approaches, and organic and enzymatic syntheses of sphingolipids and analogs. The critically acclaimed laboratory standard for more than forty years, *Methods in Enzymology* is one of the most highly respected publications in the field of biochemistry. Since 1955, each volume has been eagerly awaited, frequently consulted, and praised by researchers and reviewers alike. Now with more than 300 volumes (all of them still in print), the series contains much material still relevant today--truly an essential publication for researchers in all fields of life sciences.

Sphingolipid Metabolism and Cell Signaling, Part B, 2000-10-25 This volume contains information on analyzing sphingolipids, sphingolipid transport and trafficking, and sphingolipid-protein interactions and cellular targets. Its companion Volume 311 presents methods used in studying enzymes of sphingolipid biosynthesis and turnover, including inhibitors of some of these enzymes, genetic approaches, and organic and enzymatic syntheses of sphingolipids and analogs. The critically acclaimed laboratory standard for more than forty years, *Methods in Enzymology* is one of the most highly respected publications in the field of biochemistry. Since 1955, each volume has been eagerly awaited, frequently consulted, and praised by researchers and reviewers alike. Now with more than 300 volumes (all of them still in print), the series contains much material still relevant today--truly an essential publication for researchers in all fields of life sciences.

Influenza Virus Yohei Yamauchi, Maria João Amorim, 2025-01-31 This fully updated volume presents protocols crucial for contemporary influenza virus research, placing a special emphasis on the latest advancements in imaging technologies, including super-resolution microscopy, cryo-electron tomography (cryo-ET), freeze replica electron microscopy, and high-speed atomic force microscopy (AFM). In addition to covering traditional methods like virus purification, this edition introduces innovative approaches for visualizing molecular interactions between viral proteins and host proteins, methods for

developing physiological cell culture models and animal models, as well as concepts concerning the prediction of viral evolution and the utilization of artificial intelligence in infection biology research. 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 and readily reproducible laboratory protocols, and tips on troubleshooting and avoiding known pitfalls. Authoritative and up-to-date, Influenza Virus: Methods and Protocols, Second Edition serves as an ideal guide for researchers looking to deepen our understanding of the intricate interplay between viruses and host cells.

Evolution and the Mechanisms of Decision Making Peter Hammerstein, Jeffrey R. Stevens, 2024-03-19 A multidisciplinary examination of cognitive mechanisms, shaped over evolutionary time through natural selection, that govern decision making. How do we make decisions? Conventional decision theory tells us only which behavioral choices we ought to make if we follow certain axioms. In real life, however, our choices are governed by cognitive mechanisms shaped over evolutionary time through the process of natural selection. Evolution has created strong biases in how and when we process information, and it is these evolved cognitive building blocks—from signal detection and memory to individual and social learning—that provide the foundation for our choices. An evolutionary perspective thus sheds necessary light on the nature of how we and other animals make decisions. This volume—with contributors from a broad range of disciplines, including evolutionary biology, psychology, economics, anthropology, neuroscience, and computer science—offers a multidisciplinary examination of what evolution can tell us about our and other animals' mechanisms of decision making. Human children, for example, differ from chimpanzees in their tendency to over-imitate others and copy obviously useless actions; this divergence from our primate relatives sets up imitation as one of the important mechanisms underlying human decision making. The volume also considers why and when decision mechanisms are robust, why they vary across individuals and situations, and how social life affects our decisions.

Advances in Genetics ,1990-05-23 Advances in Genetics

Plant Responses to Environmental Stresses Lerner, 2018-04-27 Emphasizing the unpredictable nature of plant behaviour under stress and in relation to complex interactions of biological pathways, this work covers the versatility of plants in adapting to environmental change. It analyzes environmentally triggered adaptations in developmental programmes of plants that lead to permanent, heritable DNA modifications.

Insect Life Cycles Francis Gilbert, 2012-12-06 No International Congress of Entomology would now be complete without a symposium on insect life-cycles. The latest Congress, held at Vancouver, BC (Canada), in July 1988, was no exception, with a symposium on the genetics, evolution, and coordination of insect life cycles organized by Bill Bradshaw and Valerie Brown. The present volume arose from papers contributed by most of the speakers at the symposium, together with papers from other invited authors. In editing the book, I have been assisted greatly by the other authors, particularly Bill Bradshaw, Val

Brown and Fritz Taylor. All contributors agreed to referee two other chapters, a system that worked efficiently and effectively: I thank all authors for performing this task in the face of other demands on their time. I would also like to thank Philip Corbet, John Greenslade, Bryan Clarke, and Gillian Thompson of Springer for their help. Nottingham Francis Gilbert

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Introduction William E. Bradshaw 3 1 Understanding the Evolution of Insect Life-Cycles: The Role of Genetic Analysis.

The Prokaryotes M.P. Starr, H. Stolp, H.G. Trüper, A. Balows, H.G. Schlegel, 2013-11-11 The purpose of this brief Foreword is to make you, the reader, hungry for the scientific feast that follows. These two volumes on the prokaryotes offer a truly unique scientific menu—a comprehensive assembly of articles, exhibiting the biochemical depth and remarkable physiological and morphological diversity of prokaryote life. The size of the volumes might initially discourage the unprepared mind from being attracted to the study of prokaryote life, for this landmark assemblage thoroughly documents the wealth of present knowledge. But in confronting the reader with the state of the art, the Handbook also defines where new work needs to be done on well-studied bacteria as well as on unusual or poorly studied organisms. There are basically two ways of doing research with microbes. A classical approach is first to define the phenomenon to be studied and then to select the organism accordingly. Another way is to choose a specific organism and go where it leads. The pursuit of an unusual microbe brings out the latent hunter in all of us. The intellectual challenges of the chase frequently test our ingenuity to the limit. Sometimes the quarry repeatedly escapes, but the final capture is indeed a wonderful experience. For many of us, these simple rewards are sufficiently gratifying so that we have chosen to spend our scientific lives studying these unusual creatures.

Integration and Excision of DNA Molecules P.H. Hofschneider, P. Starlinger, 2012-12-06 The topic of this year's ~osbach Colloquium was DNA integration. We have tried to bring together experts from different fields of research who are studying natural processes by which DNA molecules from different sources are linked. It has been known for a long time that such linkage occurs between the chromosomes of bacteriophages and plasmids on the one hand and the chromosome of the bacterial host on the other. This process has been especially well studied in bacteriophage λ . Since it is controlled in a complicated way, we began with a lecture by M. Ptashne on these regulatory processes. H. Nash described the integration of bacteriophage λ into the bacterial chromosome. To put this site-specific process into perspective, G. Mosig lectured on genetic recombination in prokaryotes in general and K. Murray described the use of bacteriophage λ as an artificial vector for genetic engineering. A different kind of bacteriophage integration is shown by bacteriophage Mu, which is much less specific in its choice of an integration site than λ . The properties of this phage were described by P. van de Putte.

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