

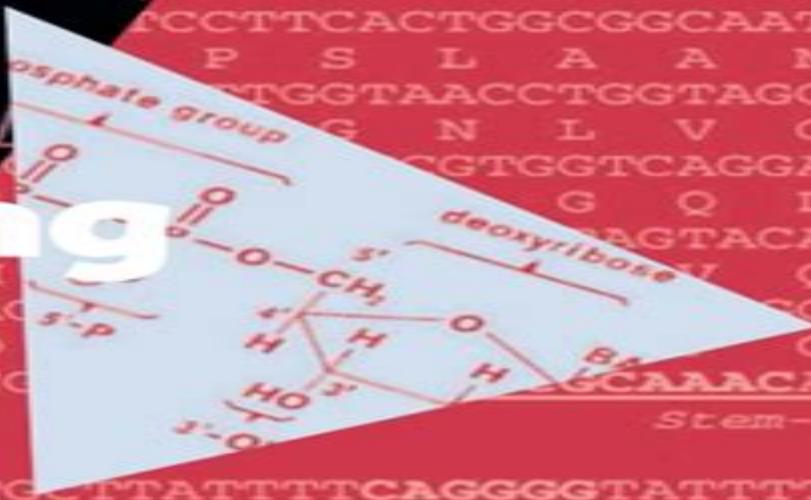
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Dominic W.S. Wong

 Springer



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The ABCs of Gene Cloning Dominic W. S. Wong, 2018-05-26 Clear and concise this easy to use book offers an introductory course on the language of gene cloning covering microbial plant and mammalian systems It presents the nuts and bolts of gene cloning in a well organized and accessible manner Part I of this book outlines the essentials of biology and genetics relevant to the concept of gene cloning Part II describes common techniques and approaches of gene cloning ranging from the basic mechanics of DNA manipulation vector systems process transformation to gene analysis Part III IV present application technologies of major impact in agriculture biomedicine and related areas The ABCs of Gene Cloning Third Edition contains updates including a tutorial chapter on gene vector construction methodologies on exome sequencing in finding disease genes revised topics on gene therapy and whole genome sequencing new developments for gene targeting and genome editing as well as the current state of next generation sequencing With more than 140 illustrations this new edition provides an invaluable text for students and anyone who have interest in gaining proficiency in reading and speaking the language of gene cloning

The ABCs of Gene Cloning Dominic Wong, 2007-12-31 Clear and concise this easy to use text offers an introductory course on the language of gene cloning covering microbial plant and animal systems The essential concepts in biology relevant to the understanding of gene cloning are presented in a well organized and accessible manner This updated version of the first edition is an invaluable book for nonscientists as well as scientists with little background knowledge in gene cloning providing a wealth of information for anyone wishing to gain proficiency in reading and speaking the language of gene cloning

ABC Transporters: Biochemical, Cellular, and Molecular Aspects Suresh V. Ambudkar, Michael M. Gottesman, 1998-07-31 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 More than 285 volumes have been published all of them still in print and much of the material is relevant even today truly an essential publication for researchers in all fields of life sciences Prokaryotic ABC Transporters Eukaryotic ABC Transporters Nonmammalian ABC Transport Systems Mammalian P Glycoproteins Multidrug Resistance Associated Protein Cystic Fibrosis Transmembrane Conductance Regulator Sulfonylurea Receptor Intracellular ABC Transporters

Gene Cloning Terence A. Brown, 1995

Synthetic Biology and Metabolic Engineering in Plants and Microbes Part B: Metabolism in Plants, 2016-07-29 Synthetic Biology and Metabolic Engineering in Plants and Microbes Part B the latest volume in the *Methods in Enzymology* series continues the legacy of this premier serial with quality chapters authored by leaders in the field This volume covers research methods synthetic biology and metabolic engineering in plants and microbes and includes sections on such topics as the usage of integrases in microbial engineering biosynthesis and engineering of tryptophan derived metabolites regulation and discovery of fungal natural products and elucidation and localization of plant pathways Continues the legacy of this premier

serial with quality chapters authored by leaders in the field of enzymology Contains two volumes covering research methods in synthetic biology and metabolic engineering in plants and microbes Includes sections on such topics as the uses of integrases in microbial engineering biosynthesis and engineering of tryptophan derived metabolites regulation and discovery of fungal natural products and elucidation and localization of plant pathways **Nucleic Acids Abstracts** ,1997

Next-Generation Strategies for Crop Improvement Yogesh Ahlawat,Darshna Chaudhary,Pawan K. Jaiwal,2025-09-30 This edited volume serves as a comprehensive collection of methodologies paradigms and case studies explaining novel approaches for crop improvement programs It summarizes the collective knowledge of leading experts in agronomy genetics biotechnology and related fields offering a holistic perspective on the multifaceted aspects of modern crop enhancement In an era of climate change increasing population and environmental degradation the need for revolutionary advancements in agriculture has never been more urgent This book delves into the transformative potential of emerging methodologies from precision breeding and genetic engineering to computational biology and synthetic biology Through compelling case studies and insightful analyses readers discover how novel approaches are revolutionizing crop breeding cultivation and resilience The central theme of the book is the promotion of sustainable agricultural practices and the development of resilient crop varieties capable of withstanding environmental stressors pests and diseases The book fosters interdisciplinary collaboration crucial for tackling complex agricultural challenges by combining insights from diverse disciplines including bioinformatics computational biology and ecological sciences From disease resistant crops to stress tolerant varieties the advanced strategies of crop improvement offer avenues for sustainable agriculture and global food security This book is essential reading for students academics researchers scientists and policymakers interested in and working in the fields of agriculture or biotechnology Wiley Encyclopedia of Molecular Medicine, Volume 2 Wiley,2001-11-01 Gene Cloning and DNA Analysis T. A. Brown,2015-10-13 Known world wide as the standard introductory text to this important and exciting area the seventh edition of Gene Cloning and DNA Analysis addresses new and growing areas of research whilst retaining the philosophy of the previous editions Assuming the reader has little prior knowledge of the subject its importance the principles of the techniques used and their applications are all carefully laid out with over 250 clearly presented four colour illustrations In addition to a number of informative changes to the text throughout the book the chapters on DNA sequencing and genome studies have been rewritten to reflect the continuing rapid developments in this area of DNA analysis In depth description of the next generation sequencing methods and descriptions of their applications in studying genomes and transcriptomes New material on the use of ChiP seq to locate protein binding sites Extended coverage of the strategies used to assemble genome sequences Description of how the Neanderthal genome has been sequenced and what that sequence tells us about interbreeding between Neanderthals and Homo sapiens Gene Cloning and DNA Analysis remains an essential introductory text to a wide range of biological sciences students including genetics and genomics molecular biology

biochemistry immunology and applied biology It is also a perfect introductory text for any professional needing to learn the basics of the subject All libraries in universities where medical life and biological sciences are studied and taught should have copies available on their shelves

ABC Proteins I Barry Holland, Susan P. C. Cole, Karl Kuchler, Christopher F. Higgins, 2003-01-07 ABC Proteins is an in depth up to date analysis of all that is known about the subject to date It discusses and compares evolution biology and mechanism of action of all known ABC proteins including the first structural studies as well as clinical implications It will be useful to anyone trying to stay abreast of the latest findings This book is sure to become a classic and will regularly be updated

Phylogeny and Evolution of ABC Transporters Fundamental Aspects of the Mechanism of Action of ABC Transporters Prokaryote ABC Transporters Non Mammalian Transporters Multidrug Transporters ABC Transporters Physiological Roles and Human Disease Full color throughout

Genetics Abstracts, 2001 American Book Publishing Record, 2002

Recombinant DNA Methodology Jo-Anne R. Dillon, A. Nasim, Earle R. Nestmann, 1985-01-21 Basic techniques Gene cloning in *Saccharomyces cerevisiae* Gene cloning in *Pseudomonas aeruginosa* Mutant construction by in vitro mutagenesis

X-linked Adrenoleukodystrophy Doriane Trompier, Stéphane Savary, 2013-03-01 X linked adrenoleukodystrophy X ALD is the most common leukodystrophy and the most frequent peroxisomal disorder with an estimated incidence of 1 17 000 This complex neurodegenerative disorder is characterized by a huge clinical variability both in the age of onset and in symptoms The two main forms are the childhood cerebral ALD ccALD characterized by inflammatory demyelination of the central nervous system and the adult form called adrenomyeloneuropathy AMN characterized by a non inflammatory slowly progressive demyelination affecting spinal cord and peripheral nerves Adrenal insufficiency is usually associated with the nervous symptoms X ALD being the main cause of Addison s disease The main biochemical defect is the accumulation of very long chain fatty acids VLCFA fatty acids with a chain length of more than 22 carbon atoms particularly in the cholesterol ester fraction of the adrenal gland and brain white matter X ALD is associated with mutations in the ABCD1 gene which encodes for a peroxisomal ATPBinding Cassette ABC transporter predicted to allow VLCFA CoA to enter into the peroxisome for their oxidation In spite of animal models the physiopathogenesis of the disease remains poorly understood However several therapeutic strategies have been investigated Among them allogeneic bone marrow transplantation has proven effective and more recently the first gene therapy trial ended with a resounding success

Thông báo sách mới, 2008 Biotechnology, 2003 Functional and Clinical Implications of Genetic Variation in ABCC2 and Related ABC Genes Tan Duy Nguyen, 2005 The ABC superfamily of drug efflux transporters is an important class of proteins involved in drug absorption disposition and elimination We hypothesize that genetic variations in ABCC2 MRP2 and related ABC genes may contribute to interindividual variability in drug response Using an ethnically diverse set of human DNA sequences the extent of genetic variation in the coding and non coding regions of ABCC2 was determined Specifically exons exon intron boundaries and the 5 promoter region were screened for single

nucleotide polymorphisms SNPs A total of 68 variant sites were identified including 13 in the 5' promoter region, 23 in the exon-intron boundaries and 22 in the exons. Nucleotide diversity π in ABCC2 was assessed for non-synonymous and synonymous sites and π_{NS}/π_{Syn} ratios show selective pressure against amino acid changes. Variants identified in the 5' promoter region were assayed for promoter activity and several were found to have decreased activity compared to a reference sequence. The most notable of these is the haplotype containing the 1549G A, 1019A G and 24C T variants. This haplotype was found to be in tight linkage disequilibrium with the synonymous variant 3972C T which had been associated with higher irinotecan exposure. Allele-specific expression analysis of liver samples heterozygous for the 3972C T variant showed a 10% bias towards the C allele in the mRNA transcript. The allelic imbalance increased to 14% when considering liver samples containing the 1549G A, 1019A G, 24C T promoter haplotype. In a clinical study carried out to determine the genetic contribution to digoxin pharmacokinetics, a one mg dose of digoxin which undergoes minimal metabolism and is a substrate for transporters was administered to monozygotic (MZ) and dizygotic (DZ) twin pairs. Oral clearance, AUC_{0-3h} and apparent nonrenal clearance were found to have a high genetic component to their variability. Lastly, a broad genetic analysis was performed on 10 members of the ABC superfamily of drug efflux transporters. These transporters were found to be under selective pressure against amino acid changes, especially in the transmembrane domains where substrate binding occurs. Results from these studies may help in understanding their cellular function and how they may contribute to interindividual variability.

Chemical Abstracts, 2002 **Nature Encyclopedia of the Human Genome** David Neil Cooper, 2003

Nature Encyclopedia of the Human Genome is the only reference resource devoted entirely to the scientific basis and genetics and genomics research and the complex commercial, philosophical and ethical questions that arise from it. It presents a comprehensive and rigorously detailed overview of current genome science and its groundbreaking applications, examining the many topics that surround the field from the differing perspectives of history, philosophy, ethics, law, medicine, public health and industry. Core areas covered include structural genomics, functional genomics, chromosome structure and function, evolution and comparative genomics, genome mapping and sequencing, genes and disease, behavioural and psychiatric genetics, mathematical and population genetics, proteomics, bioinformatics, ethical, legal and social issues and biographies or key figures.

Proceedings of the National Academy of Sciences of the United States of America National Academy of Sciences (U.S.), 1995

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