

Genom 2005 340

Ezio Portis,Alberto Acquadro,Sergio Lanteri

Genetic Diagnosis of Endocrine Disorders Roy E. Weiss, Samuel Refetoff, 2015-10-09 Genetic Diagnosis of Endocrine Disorders, Second Edition provides users with a comprehensive reference that is organized by endocrine grouping (i.e., thyroid, pancreas, parathyroid, pituitary, adrenal, and reproductive and bone), discussing the genetic and molecular basis for the diagnosis of various disorders. The book emphasizes the practical nature of diagnosing a disease, including which tests should be done for the diagnosis of diabetes mellitus in adults and children, which genes should be evaluated for subjects with congenital hypothyroidism, which genetic tests should be ordered in obese patients or for those with parathyroid carcinoma, and the rationale behind testing for multiple endocrine neoplasias. Offers a clear presentations of pharmacogenetics and the actual assays used in detecting endocrine diseases Teaches the essentials of the genetic basis of disease in each major endocrine organ system Offers expert advice from genetic counselors on how to use genetic information in counseling patients Includes new chapters on the genetics of lipid disorders and glycogen storage diseases, genetics of hypoglycemia, and whole genome/exome sequencing

Origin Jennifer Raff, 2022-02-08 AN INSTANT NEW YORK TIMES BESTSELLER! From celebrated anthropologist Jennifer Raff comes the untold story—and fascinating mystery—of how humans migrated to the Americas. ORIGIN is the story of who the first peoples in the Americas were, how and why they made the crossing, how they dispersed south, and how they lived based on a new and powerful kind of evidence: their complete genomes. ORIGIN provides an overview of these new histories throughout North and South America, and a glimpse into how the tools of genetics reveal details about human history and evolution. 20,000 years ago, people crossed a great land bridge from Siberia into Western Alaska and then dispersed southward into what is now called the Americas. Until we venture out to other worlds, this remains the last time our species has populated an entirely new place, and this event has been a subject of deep fascination and controversy. No written records—and scant archaeological evidence—exist to tell us what happened or how it took place. Many different models have been proposed to explain how the Americas were peopled and what happened in the thousands of years that followed. A study of both past and present, ORIGIN explores how genetics is currently being used to construct narratives that profoundly impact Indigenous peoples of the Americas. It serves as a primer for anyone interested in how genetics has become entangled with identity in the way that society addresses the question Who is indigenous?

Molecular Medical Microbiology Yi-Wei Tang, Musa Hindiyeh, Dongyou Liu, Andrew Sails, Paul Spearman, Jing-Ren Zhang, 2023-11-21 Molecular Medical Microbiology, Third Edition presents the latest release in what is considered to be the first book to synthesize new developments in both molecular and clinical research. The molecular age has brought about dramatic changes in medical microbiology, along with great leaps in our understanding of the mechanisms of infectious disease. This third edition is completely updated, reviewed and expanded, providing a timely and helpful update for microbiologists, students and clinicians in the era of increasing use of molecular techniques, changing epidemiology and prevalence, and increasing resistance of many pathogenic bacteria. Written by experts in the field, chapters include cutting-edge information and clinical overviews for each major bacterial group, along with the latest updates on vaccine development, molecular technology and diagnostic technology. Completely updated and revised edition of this comprehensive and accessible reference on molecular medical microbiology Includes full color presentations throughout Delves into in-depth discussions on individual pathogenic bacteria in a system-oriented approach Includes a clinical overview for each major bacterial group Presents the latest information on vaccine development, molecular technology and diagnostic technology Provides more than 100 chapters on all major groups of bacteria

Recent Trends in Carbohydrate Chemistry Amélia Pilar Rauter, Bjørn E. Christensen, László Somsák, Paul Kosma, Roberto Adamo, 2020-04-29 Recent Trends in Carbohydrate Chemistry: Synthesis and Biomedical Applications of Glycans and Glycoconjugates covers biomedically relevant bacterial cell wall carbohydrates including recent findings on biosynthetic aspects, advances in the chemical assembly of bacterial lipopolysaccharide fragments and teichoic acids, and modern NMR approaches to unravel structural details. The first part introduces and provides the relevant background for synthetic glycoconjugate vaccines. The second section focuses on synthetic carbohydrate-based vaccines of therapeutic potential that are licensed or under development. This second volume of Recent Trends in Carbohydrate Chemistry is ideal for researchers working as synthetic organic chemists, as well as those interested in glycoconjugation, protein chemists, immunologists, and microbiologists, in academia as well as in industry. Highlights important features of bacterial glycoproteins Illustrates modern chemical synthesis and structural analysis of bacterial glycans Demonstrates the importance of carbohydrate chemistry for the synthesis of lipopolysaccharides and teichoic acid Covers recent findings on glycan ligation Gives an overview of the most recent developments on carbohydrate-based vaccines

Genetics, Genomics and Breeding of Potato James M. Bradeen, Chittaranjan Kole, 2016-04-19 In this volume, world leaders in potato research review historical and contemporary discoveries resulting in a range of advances. Topics include nutritional quality, yield, disease and insect resistance, processing, plant growth and development, and other aspects. The book also examines research yielding significant molecular resources that facilit

Vertebrate Genomes Jean-Nicolas Volff, 2006-01-01 The publication of the first draft of the human genome five years ago marked the start of a new era of vertebrate comparative genomics. Genome draft sequences for chimpanzee, mouse, rat, dog, chicken as well as for several fish species are now available for the reconstruction of the evolutionary history of vertebrate genomes over the last 450 million years. This volume of Genome Dynamics presents a timely update on the past, present and future of major vertebrate genome projects. In 16 chapters, renowned specialists provide informative reviews on the current status of mammalian, bird, amphibian and fish genomics. In addition, a chapter on urochordate genomes is included. The reader will also find a useful overview of resources available for each genome as well as a description of the medical and economical perspectives opened by the different projects. This book is recommended reading for biologists interested in genetics, genomics, evolution and zoology.

Sourcebook of Models for Biomedical Research P. Michael Conn, 2008-03-07 The collection of systems represented in this volume is a unique effort to reflect the diversity and utility of models used in biomedicine. That utility is based on the consideration that observations made in particular organisms will provide insight into the workings of other, more complex systems. This volume is therefore a comprehensive and extensive collection of these important medical parallels.

Principles of Addiction Medicine Richard K. Ries, Shannon C. Miller, David A. Fiellin, 2009 This respected text from the American Society of Addiction Medicine is valuable for all physicians and mental-health personnel who specialize in addiction medicine and who treat patients with addiction disorders. The chapters blend scientific principles underlying addiction with the practical essentials of clinical addiction medicine. Many of the contributors are affiliated with leading government agencies that study addiction and its science, such as the National Institute on Alcohol Abuse and Alcoholism and the National Institute on Drug Abuse. The book will appeal to a wide and interdisciplinary range of professionals, especially those with interest or duties relating to addiction-related disorders, and in particular physicians seeking certification status via either the American Board of Addiction Medicine or the American Board of Psychiatry and Neurology. A companion Website will offer the fully searchable text.

Fungal Diseases of Rice and Their Management Deepti Srivastava,Md. Shamim,Malik Mobeen Ahmad,R. S. Upadhyay,2024-01-09 Rice is a widely consumed crop around the world that has tremendous importance and is cultivated almost everywhere except Antarctica. However, various biotic and abiotic stresses have a negative effect on rice cultivation, seriously reducing its yield. This volume examines the bacterial and fungal pathogens that cause rice diseases and explores how to manage these diseases. It covers the economic and environmental impact of rice fungal diseases on global food security and proceeds to delve into diagnostic methods for rice fungal pathogen detection and discusses current strategic and applied biotechnological methods for the effective management of rice fungal diseases.

Human Herpesviruses Ann Arvin,Gabriella Campadelli-Fiume,Edward Mocarski,Patrick S. Moore,Bernard Roizman,Richard Whitley,Koichi Yamanishi,2007-08-16 This comprehensive account of the human herpesviruses provides an encyclopedic overview of their basic virology and clinical manifestations. This group of viruses includes human simplex type 1 and 2, Epstein-Barr virus, Kaposi's Sarcoma-associated herpesvirus, cytomegalovirus, HHV6A, 6B and 7, and varicella-zoster virus. The viral diseases and cancers they cause are significant and often recurrent. Their prevalence in the developed world accounts for a major burden of disease, and as a result there is a great deal of research into the pathophysiology of infection and immunobiology. Another important area covered within this volume concerns antiviral therapy and the development of vaccines. All these aspects are covered in depth, both scientifically and in terms of clinical guidelines for patient care. The text is illustrated generously throughout and is fully referenced to the latest research and developments.

The Tomato Genome Mathilde Causse,Jim Giovannoni,Mondher Bouzayen,Mohamed Zouine,2016-11-21 This book describes the strategy used for sequencing, assembling and annotating the tomato genome and presents the main characteristics of this sequence with a special focus on repeated sequences and the ancestral polyploidy events. It also includes the chloroplast and mitochondrial genomes. Tomato (*Solanum lycopersicum*) is a major crop plant as well as a model for fruit development, and the availability of the genome sequence has completely changed the paradigm of the species' genetics and genomics. The book describes the numerous genetic and genomic resources available, the identified genes and quantitative trait locus (QTL) identified, as well as the strong synteny across Solanaceae species. Lastly, it discusses the consequences of the availability of a high-quality genome sequence of the cultivated species for the research community. It is a valuable resource for students and researchers interested in the genetics and genomics of tomato and Solanaceae.

Genetics of Populations Philip Hedrick,2011-08-24 The Fourth Edition of Genetics of Populations is the most current, comprehensive, and accessible introduction to the field for advanced undergraduate and graduate students, and researchers in genetics, evolution, conservation, and related fields. In the past several years, interest in the application of population genetics principles to new molecular data has increased greatly, and Dr. Hedrick's new edition exemplifies his commitment to keeping pace with this dynamic area of study. Reorganized to allow students to focus more sharply on key material, the Fourth Edition integrates coverage of theoretical issues with a clear presentation of experimental population genetics and empirical data. Drawing examples from both recent and classic studies, and using a variety of organisms to illustrate the vast developments of population genetics, this text provides students and researchers with the most comprehensive resource in the field.

Genetics of Populations Philip W. Hedrick,2009-12-29 The Fourth Edition of Genetics of Populations is the most current, comprehensive, and accessible introduction to the field for advanced undergraduate and graduate students, and researchers in genetics, evolution, conservation, and related fields. In the past several years, interest in the application of population genetics principles to new molecular data has increased greatly, and Dr. Hedrick's new edition exemplifies his commitment to keeping pace with this dynamic area of study. Reorganized to allow students to focus more sharply on key material, the Fourth Edition integrates coverage of theoretical issues with a clear presentation of experimental population genetics and empirical data. Drawing examples from both recent and classic studies, and using a variety of organisms to illustrate the vast developments of population genetics, this text provides students and researchers with the most comprehensive resource in the field.

Animal Genomics for Animal Health Marie-Hélène Pinard,2008-01-01 It includes a review of the state of the art in animal genomics and its applications to animal health. The contributions describe the new tools available, such as HapMaps for chicken and cattle, and show how the understanding of gene structure and function can be successfully applied to delineate the molecular mechanisms of disease and determine complex phenotypes associated with health traits. A critical evaluation of future needs and future applications of animal genomics is also presented. The integration of animal genomics in animal health research is likely to revolutionize the way scientists approach the challenges of discovering highly effective drugs and vaccines for animal diseases.-BOOK JACKET.

Broadening the Genetic Base of Grain Cereals Mohar Singh,Sandeep Kumar,2016-09-13 This book offers comprehensive coverage of important grain cereals including their origin and distribution, crop gene pool, level of diversity, production constraints, traits of importance for genetic base widening, crop improvement methodologies, genome mapping, genomics for breeding, and future strategies. The chapters, contributed by eminent crop researchers from around the world, provide rare insights into the crop-specific constraints and prospects drawing from their substantial experience. As such, the book offers an essential source of information for grain cereals scientists, teachers, students, policy planners and developmental experts alike. Grain cereals, which comprise rice, wheat, maize, barley, oats, sorghum and millets, are members of the grass family. These crops are vital to human nutrition, thanks to their roles as staple food crops in different parts of the globe. Some of them are rich sources of carbohydrates, which provide energy, while others are important sources of minerals, vitamins and proteins, in addition to their medicinal properties. In most cereals, the existing variability among elite germplasm has been exploited to attain a desirable level of productivity. However, to make further breakthroughs in enhancing yield and improving stability in future crop cultivars, new sources of genes/alleles need to be identified in wild/weedy species and incorporated into the cultivated varieties. Though there have been many publications on various aspects of grain cereal improvement in the recent past, to date this essential information has remained scattered among different periodicals.

Present Knowledge in Food Safety Michael E. Knowles,Lucia Anelich,Alan Boobis,Bert Popping,2022-10-08 Present Knowledge in Food Safety: A Risk-Based Approach Through the Food Chain presents approaches for exposure-led risk assessment and the management of changes in the chemical, pathogenic microbiological and physical (radioactivity) contamination of 'food' at all key stages of production, from farm to consumption. This single volume resource introduces scientific advances at all stages of the production to improve reliability, predictability and relevance of food safety assessments for the protection of public health. This book is aimed at a diverse audience, including graduate and post-graduate students in food science, toxicology, microbiology, medicine, public health, and related fields. The book's reach also includes government agencies, industrial scientists, and policymakers involved in food risk analysis. Includes new technologies such as nanotechnology, genetic modification, and cloning Provides information on advances in pathogen risk assessment through novel and real-time molecular biological techniques, biomarkers, resistance measurement, and cell-to-cell communication in the gut Covers the role of the microbiome and the use of surrogates (especially for viruses)

The Globe Artichoke Genome Ezio Portis,Alberto Acquadro,Sergio Lanteri,2019-06-08 This book presents the latest information on the genetics and genomics of the globe artichoke. It

focuses on the latest findings, tools and strategies employed in genome sequencing, physical map development and QTL analyses, as well as genomic resources. The re-sequencing of four globe artichoke genotypes, representative of the core varietal types in cultivation, as well as the genotype of cultivated cardoon, has recently been completed. Here, the five genomes are reconstructed at the chromosome scale and annotated. Moreover, functional SNP analyses highlight numerous genetic variants, which represent key tools for dissecting the path from sequence variation to phenotype, as well as for designing effective diagnostic markers. The wealth of information provided here offers a valuable asset for scientists, plant breeders and students alike.

Genomic Designing for Abiotic Stress Resistant Fruit Crops Chittaranjan Kole,2022-09-20 This book presents deliberations on molecular and genomic mechanisms underlying the interactions of crop plants to the abiotic stresses caused by heat, cold, drought, flooding, submergence, salinity, acidity, etc., important to develop resistant crop varieties. Knowledge on the advanced genetic and genomic crop improvement strategies including molecular breeding, transgenics, genomic-assisted breeding, and the recently emerging genome editing for developing resistant varieties in fruit crops is imperative for addressing FHNEE (food, health, nutrition, energy, and environment) security. Whole genome sequencing in many of these crops followed by genotyping-by-sequencing has provided precise information regarding the genes conferring resistance useful for gene discovery, allele mining, and shuttle breeding which in turn opened up the scope for 'designing' crop genomes with resistance to abiotic stresses. The seven chapters each dedicated to a fruit crop and a fruit crop group in this volume elucidate different types of abiotic stresses and their effects on and interaction with the crops; enumerate the available genetic diversity with regard to abiotic stress resistance among available cultivars; illuminate the potential gene pools for utilization in interspecific gene transfer; present brief on classical genetics of stress resistance and traditional breeding for transferring them to their cultivated counterparts; depict the success stories of genetic engineering for developing abiotic stress-resistant crop varieties; discuss on molecular mapping of genes and QTLs underlying stress resistance and their marker-assisted introgression into elite varieties; enunciate different genomics-aided techniques including genomic selection, allele mining, gene discovery, and gene pyramiding for developing adaptive crop varieties with higher quantity and quality of yields, and also elaborate some case studies on genome editing focusing on specific genes for generating abiotic stress-resistant crops.

The Phytopathogen Abhijeet Ghatak,Mohammad Ansar,2017-03-31 So often new phytopathogens emerge and appear primarily in acute form and then take a chronic form; such populations, however, in general have a limited appearance because of the lack of suitable environmental conditions. The emergence of new pathogens needs to be explored in the light of their evolutionary adaptation. This new volume focuses on the study of quantitative aspects of host-phytopathogen linkages that result in the emergence of aggressive phytopathogens. The book examines the evolution and adaptation of phytopathogens from several cropping systems.

Evolution by Gene Duplication Susumu Ohno,2013-12-11 It is said that necessity is the mother of invention. To be sure, wheels and pulleys were invented out of necessity by the tenacious minds of upright citi zens. Looking at the history of mankind, however, one has to add that leisure is the mother of cultural improvement. Man's creative genius flourished only when his mind, freed from the worry of daily toils, was permitted to entertain apparently useless thoughts. In the same manner, one might say with regard to evolution that natural selection mere(y tnodifted, while redundanry created. Natural selection has been extremely effective in policing alleHe mutations which arise in already existing gene loci. Because of natural selection, organisms have been able to adapt to changing environments, and by adaptive radiation many new species were created from a common ancestral form. Y et, being an effective policeman, natural selection is extremely conservative by nature. Had evolution been entirely dependent upon natural selection, from a bacterium only numerous forms of bacteria would have emerged. The creation of metazoans, vertebrates and finally mammals from unicellular organisms would have been quite impos sible, for such big leaps in evolution required the creation of new gene loci with previously nonexistent functions. Only the cistron which became redun dant was able to escape from the relentless pressure of natural selection, and by escaping, it accumulated formerly forbidden mutations to emerge as a new gene locus.

Delve into the emotional tapestry woven by in **Genom 2005 340** . This ebook, available for download in a PDF format (PDF Size: *), is more than just words on a page; itis a journey of connection and profound emotion. Immerse yourself in narratives that tug at your heartstrings. Download now to experience the pulse of each page and let your emotions run wild.

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