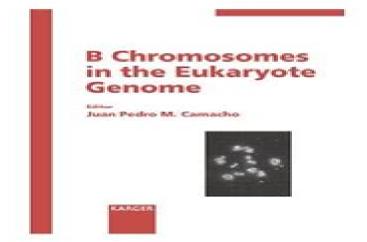
# B Chromosomes in the Eukaryote Genome



# **B** Chromosomes In The Eukaryote Genome

**Brendan G. Carr** 

#### **B Chromosomes In The Eukaryote Genome:**

*B Chromosomes in the Eukaryote Genome* ,2004 *B Chromosomes in the Eukaryote Genome* Juan Pedro M. Camacho, 2004 B chromosomes are additional passengers found in the karyotypes of about 15% of eukaryote species They are best understood as genome parasites exploiting the host genome because of their transmissional advantage and are frequently deleterious for the organism carrying them Since the last compilation of B chromosome literature more than 20 years ago many new findings have contributed to building an increasing body of knowledge of these mysterious genetic elements This single topic issue of Cytogenetic and Genome Research updates these new data in its 40 papers authored by researchers from 19 countries It provides up to date reviews on plants fish amphibians and mammals and for the first time on Sternorrhyncha hemipterans decapodan crustaceans helminth parasites and lizards It also presents new original data on many aspects of B chromosomes including their definition new occurrences frequency structure composition origin effects on the host genome transmission and population dynamics Unique in its field this publication is recommended reading for cytogeneticists interested in chromosome polymorphisms for evolutionary biologists interested in selfish genetic elements and for geneticists interested in the intragenomic conflict The Eukaryote Genome in Development and Evolution John Bernard, 2012-12-06 The mind unlearns with difficulty what has long been impressed upon it Seneca Reductionism is without question the most successful analytical approach available to the experimental scientist With the advent of techniques for cloning and sequencing DNA and the development of a variety of molecular probes for localizing macromolecules in cells and tissues the biologist now has available the most powerful reductionist tools ever invented The application of these new technologies has led to a veritable explosion of facts regarding the types and organization of nucleotide sequences present in the genomes of eukaryotes These data offer a level of precision and predictability which is unparalleled in biology Recombinant DNA techniques were initially developed to gather information about the structure and organization of the DNA sequences within a genome The power and potential of these techniques however extend far beyond simple data collection of this kind In an attempt to use the new technology as a basis for analyzing development and evolution attention was first focused on the topic of gene regulation an approach that had proven so successful in prokaryotes It is now clear that this has not been an adequate approach Lewin 1984 has quoted Brenner as stating at the beginning it was said that the answer to the understanding of development was going to come from a knowledge of the molecular mechanisms of gene control I doubt whether anyone believes this any more Evolution. Composition and Regulation of Supernumerary B Chromosomes Andreas Houben, Neil Jones, Cesar Martins, Vladimir Trifonov, 2019-04-25 Supernumerary B chromosomes Bs are dispensable genetic elements found in thousands of species of plants and animals and some fungi Since their discovery more than a century ago they have been a source of puzzlement as they only occur in some members of a population and are absent from others When they do occur they are often harmful and in the absence of

selfishness based on mechanisms of mitotic and meiotic drive there appears to be no obvious reason for their existence Cytogeneticists have long wrestled with questions about the biological existence of these enigmatic elements including their lack of any adaptive properties apparent absence of functional genes their origin sequence organization and co evolution as nuclear parasites Emerging new technologies are now enabling researchers to step up a gear to look enthusiastically beyond the previous limits of the horizon and to uncover the secrets of these silent chromosomes This book provides a comprehensive guide to theoretical advancements in the field of B chromosome research in both animal and plant systems

Plant Genome Diversity Volume 2 Johann Greilhuber, Jaroslav Dolezel, Jonathan Wendel, 2012-11-13 This second of two volumes on Plant Genome Diversity provides in 20 chapters insights into the structural evolution of plant genomes with all its variations Starting with an outline of plant phylogeny and its reconstruction the second part of the volume describes the architecture and dynamics of the plant cell nucleus the third examines the evolution and diversity of the karyotype in various lineages including angiosperms gymnosperms and monilophytes. The fourth part presents the mechanisms of polyploidization and its biological consequences and significance for land plant evolution. The fifth part deals with genome size evolution and its biological significance Together with Volume I this comprehensive book on the plant genome is intended for students and professionals in all fields of plant science offering as it does a convenient entry into a burgeoning literature in a fast moving Protocols for Cytogenetic Mapping of Arthropod Genomes Igor V. Sharakhov, 2014-10-28 Arthropods are important field to worldwide agriculture food safety human health and energy production Besides their practical significance various species represent excellent model systems for biological investigations of evolution development physiology reproduction and social interaction For these reasons arthropod genomics is receiving increasing attention from researchers around the globe Protocols for Cytogenetic Mapping of Arthropod Genomes is a collection of up to date detailed protocols for physical chromosome mapping and their applications for studying genome organization and evolution in insects and ticks This book brings together the expertise of cytogeneticists working on diverse groups of arthropods Each chapter demonstrates approaches to tissue dissection chromosome preparation fluorescence in situ hybridization and imaging The book is a timely and complementary effort to the i5K initiative which will obtain whole genome sequences for 5 000 insect and related arthropod species This comprehensive resource provides cytogeneticists with the necessary background and protocols to understand and develop chromosome based genome assemblies from such whole genome sequence data **Genome** M. Timothy Rabanus-Wallace, Nils Stein, 2021-10-25 This book celebrates the dawn of the rye genomics era with concise comprehensive and accessible reviews on the current state of rye genomic research written by experts in the field for students researchers and growers To most rye is the key ingredient in a flavoursome bread or their favourite American whisky To a farmer rye is the remarkable grain that tolerates the harshest winters and the most unforgiving soils befitting its legacy as the life giving seed that fed the ancient civilisations of northern Eurasia Since the mid 1900s scientists have

employed genetic approaches to better understand and utilize rye but only since the technological advances of the mid 2010s has the possibility of addressing questions using rye genome assemblies become a reality Alongside the secret of its unique survival abilities rye genomics has accelerated research on a host of intriguing topics such as the complex history of rye s domestication by humans the nature of genes that switch fertility on and off the function and origin of accessory chromosomes and the evolution of selfish DNA The Evolution of the Genome T. Ryan Gregory, 2011-05-04 The Evolution of the Genome provides a much needed overview of genomic study through clear detailed expert authored discussions of the key areas in genome biology This includes the evolution of genome size genomic parasites gene and ancient genome duplications polypoidy comparative genomics and the implications of these genome level phenomena for evolutionary theory In addition to reviewing the current state of knowledge of these fields in an accessible way the various chapters also provide historical and conceptual background information highlight the ways in which the critical questions are actually being studied indicate some important areas for future research and build bridges across traditional professional and taxonomic boundaries The Evolution of the Genome will serve as a critical resource for graduate students postdoctoral fellows and established scientists alike who are interested in the issue of genome evolution in the broadest sense Provides detailed clearly written chapters authored by leading researchers in their respective fields Presents a much needed overview of the historical and theoretical context of the various areas of genomic study Creates important links between topics in order to promote integration across subdisciplines including descriptions of how each subject is actually studied Provides information specifically designed to be accessible to established researchers postdoctoral fellows and graduate students alike **Plant** Cytogenetics Hank Bass, James A. Birchler, 2011-12-02 This reference book provides information on plant cytogenetics for students instructors and researchers Topics covered by international experts include classical cytogenetics of plant genomes plant chromosome structure functional molecular cytology and genome dynamics In addition chapters are included on several methods in plant cytogenetics informatics and even laboratory exercises for aspiring or practiced instructors. The book provides a unique combination of historical and modern subject matter revealing the central role of plant cytogenetics in plant genetics and genomics as currently practiced This breadth of coverage together with the inclusion of methods and instruction is intended to convey a deep and useful appreciation for plant cytogenetics. We hope it will inform and inspire students researchers and teachers to continue to employ plant cytogenetics to address fundamental questions about the cytology of plant chromosomes and genomes for years to come Hank W Bass is a Professor in the Department of Biological Science at Florida State University James A Birchler is a Professor in the Division of Biological Sciences at the University of Genomes 3 Terence A. Brown, 2007 The VitalBook e book version of Genomes 3 is only available in the US and Missouri Canada at the present time To purchase or rent please visit http store vitalsource com show 9780815341383 Covering molecular genetics from the basics through to genome expression and molecular phylogenetics Genomes 3is the latest

edition of this pioneering textbook Updated to incorporate the recent major advances Genomes 3 is an invaluable companion for any undergraduate throughout their studies in molecular genetics Genomes 3 builds on the achievements of the previous two editions by putting genomes rather than genes at the centre of molecular genetics teaching Recognizing that molecular biology research was being driven more by genome sequencing and functional analysis than by research into genes this approach has gathered momentum in recent years

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