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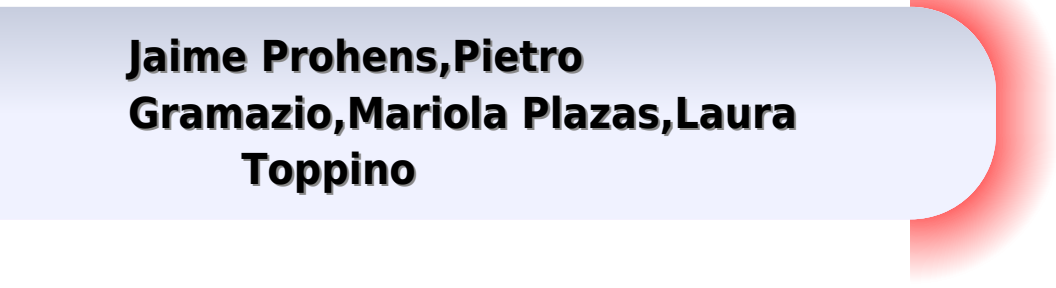
Alien Introgression in Wheat

Cytogenetics, Molecular Biology, and
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And Alien Introgression Wheat Cytogenetics Molecular

**Jaime Prohens, Pietro
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Alien Introgression in Wheat Márta Molnár-Láng, Carla Ceoloni, Jaroslav Doležal, 2015-11-20 This book provides an overview of the latest advancements in the field of alien introgression in wheat The discovery and wide application of molecular genetic techniques including molecular markers in situ hybridization and genomics has led to a surge in interspecific and intergeneric hybridization in recent decades The work begins with the taxonomy of cereals especially of those species which are potential gene sources for wheat improvement The text then goes on to cover the origin of wheat breeding in connection with alien introgressions and the problems of producing intergeneric hybrids and backcross derivatives These problems can include crossability sterility and unequal chromosome transmission The work then covers alien introgressions according to the related species used as well as new results in the field of genomics of wild wheat relatives and introgressions

Alien Introgression in Wheat Márta Molnár-Láng, Carla Ceoloni, Jaroslav Doležal, 2015 This book provides an overview of the latest advancements in the field of alien introgression in wheat The discovery and wide application of molecular genetic techniques including molecular markers in situ hybridization and genomics has led to a surge in interspecific and intergeneric hybridization in recent decades The work begins with the taxonomy of cereals especially of those species which are potential gene sources for wheat improvement The text then goes on to cover the origin of wheat breeding in connection with alien introgressions and the problems of producing intergeneric hybrids and backcross derivatives These problems can include crossability sterility and unequal chromosome transmission The work then covers alien introgressions according to the related species used as well as new results in the field of genomics of wild wheat relatives and introgressions

Molecular Advances in Wheat and Barley Manuel Martinez, 2019-08-20 Allohexaploid bread wheat and diploid barley are two of the most cultivated crops in the world This book reports novel research and reviews concerning the use of modern technologies to understand the molecular bases for wheat and barley improvement The contributions published in this book illustrate research advances in wheat and barley knowledge using modern molecular techniques These molecular approaches cover genomic transcriptomic proteomic and phenomic levels together with new tools for gene identification and the development of novel molecular markers Overall the contributions for this book lead to a further understanding of regulatory systems in order to improve wheat and barley performance

Aegilops: Promising Genesources to Improve Agronomical and Quality Traits of Wheat Peter Shewry, Marianna Rakszegi, István Molnár, Eva Darko, Vijay Kumar Tiwari, 2020-09-02 This eBook is a collection of articles from a Frontiers Research Topic Frontiers Research Topics are very popular trademarks of the Frontiers Journals Series they are collections of at least ten articles all centered on a particular subject With their unique mix of varied contributions from Original Research to Review Articles Frontiers Research Topics unify the most influential researchers the latest key findings and historical advances in a hot research area Find out more on how to host your own Frontiers Research Topic or contribute to one as an author by

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Chromosome Manipulation for Plant Breeding

Purposes Pilar Prieto, 2021-01-29 The ability to exploit the potential of wild relatives carrying beneficial traits is a major goal in breeding programs. However, it relies on the possibility of the chromosomes from the crop and wild species in interspecific crosses to recognize, associate, and undergo crossover formation during meiosis, the cellular process responsible for producing gametes with half the genetic content of their parent cells. Unfortunately, in most cases, a barrier exists preventing successful hybridization between the wild and crop chromosomes. Understanding the mechanisms controlling chromosome associations during meiosis are of great interest in plant breeding and will allow chromosome manipulation to introduce genetic variability from related species into a crop. In addition to interspecific hybrids, other materials such as natural and synthetic polyploids and introgression lines derived from allopolyploids among others are powerful tools in the framework of plant breeding. For example, an extra pair of alien chromosomes in the full genome complement of a crop species has been frequently used as a first step to access genetic variation from the secondary gene pool in breeding programs. In addition, such introgression lines are also pivotal in the study of interspecific genetic interactions, the chromosomal location of genetic markers, and in the study of chromosome structure and behavior in somatic and meiotic cells. Contained in this Special Issue are accounts of original research including new tools to identify chromosome introgressions and the development and characterization of introgression lines and interspecific hybrids carrying desirable agronomic traits for plant breeding purposes. Also included are reviews about the chromosome engineering of tropical cash crops and the effect of chromosome structure on chromosome associations and recombination during meiosis to allow chromosome manipulation in the framework of plant breeding.

Plant Genetics and Biotechnology in Biodiversity Rosa Rao, Giandomenico

Corrado, 2018-08-09 This book is a printed edition of the Special Issue Plant Genetics and Biotechnology in Biodiversity that was published in Diversity.

Interspecific Hybridization in Plant Biology

Andrew H. Paterson, Dayun Tao, Ruslan Kalendar, 2022-10-13

Wheat Evolution and Domestication Moshe Feldman, Avraham A. Levy, 2023-10-03 This open access book covers a century of research on wheat genetics and evolution starting with the discovery in 1918 of the accurate number of chromosomes in wheat. We re-evaluate classical studies that are pillars of the current knowledge considering recent genomic data in the wheat group comprising 31 species from the genera *Amblyopyrum*, *Aegilops*, *Triticum*, and other more distant relatives. For these species, we describe morphology, ecogeographical distribution, phylogeny, as well as cytogenetic and genomic features. For crops, we also address evolution under human selection, namely pre-domestication cultivation and domestication. We re-examine the genetic and archeological evidence of where, when, and how domestication occurred. We discuss unique aspects of genome evolution and maintenance under polyploidization in natural and synthetic allopolyploids of the wheat group. Finally, we propose some thoughts on the future prospects of wheat improvement. As such, it can be of great interest to wheat researchers and breeders as well as to plant scientists and students interested in plant

genetics evolution domestication and polyploidy *Introgression Breeding in Cultivated Plants* Jaime Prohens, Pietro Gramazio, Mariola Plazas, Laura Toppino, 2021-11-24 *QTL Mapping in Crop Improvement* Shabir Hussain Wani, Dechun Wang, Gyanendra Pratap Singh, 2022-11-19

QTL Mapping in Crop Improvement Present Progress and Future Perspectives presents advancements in QTL breeding for biotic and abiotic stresses and nutritional improvement in a range of crop plants. The book presents a roadmap for future breeding for resilience to various stresses and improvement in nutritional quality. Crops such as rice, wheat, maize, soybeans, common bean, and pigeon pea are the major staple crops consumed globally; hence fulfilling the nutritional requirements of global populations, particularly in the underdeveloped world, is extremely important. Sections cover the challenges facing maximized production of these crops, including diseases, insect damage, drought, heat, salinity, and mineral toxicity. Covering globally important crops including maize, wheat, rice, barley, soybean, common bean, and pigeon pea, this book will be an important reference for those working in agriculture and crop improvement. Uses the latest molecular markers to identify QTLs/genes responsible for biotic and abiotic stress tolerance in plants. Includes multiple core crops for efficient comparison and translational learning. Provides a ready reference for improving quality traits through the use of the latest technologies.

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Table of Contents And Alien Introgression Wheat Cytogenetics Molecular

1. Understanding the eBook And Alien Introgression Wheat Cytogenetics Molecular
 - The Rise of Digital Reading And Alien Introgression Wheat Cytogenetics Molecular
 - Advantages of eBooks Over Traditional Books
2. Identifying And Alien Introgression Wheat Cytogenetics Molecular
 - Exploring Different Genres
 - Considering Fiction vs. Non-Fiction
 - Determining Your Reading Goals
3. Choosing the Right eBook Platform
 - Popular eBook Platforms
 - Features to Look for in an And Alien Introgression Wheat Cytogenetics Molecular
 - User-Friendly Interface
4. Exploring eBook Recommendations from And Alien Introgression Wheat Cytogenetics Molecular

- Personalized Recommendations
- And Alien Introgression Wheat Cytogenetics Molecular User Reviews and Ratings
- And Alien Introgression Wheat Cytogenetics Molecular and Bestseller Lists
- 5. Accessing And Alien Introgression Wheat Cytogenetics Molecular Free and Paid eBooks
 - And Alien Introgression Wheat Cytogenetics Molecular Public Domain eBooks
 - And Alien Introgression Wheat Cytogenetics Molecular eBook Subscription Services
 - And Alien Introgression Wheat Cytogenetics Molecular Budget-Friendly Options
- 6. Navigating And Alien Introgression Wheat Cytogenetics Molecular eBook Formats
 - ePub, PDF, MOBI, and More
 - And Alien Introgression Wheat Cytogenetics Molecular Compatibility with Devices
 - And Alien Introgression Wheat Cytogenetics Molecular Enhanced eBook Features
- 7. Enhancing Your Reading Experience
 - Adjustable Fonts and Text Sizes of And Alien Introgression Wheat Cytogenetics Molecular
 - Highlighting and Note-Taking And Alien Introgression Wheat Cytogenetics Molecular
 - Interactive Elements And Alien Introgression Wheat Cytogenetics Molecular
- 8. Staying Engaged with And Alien Introgression Wheat Cytogenetics Molecular
 - Joining Online Reading Communities
 - Participating in Virtual Book Clubs
 - Following Authors and Publishers And Alien Introgression Wheat Cytogenetics Molecular
- 9. Balancing eBooks and Physical Books And Alien Introgression Wheat Cytogenetics Molecular
 - Benefits of a Digital Library
 - Creating a Diverse Reading Collection And Alien Introgression Wheat Cytogenetics Molecular
- 10. Overcoming Reading Challenges
 - Dealing with Digital Eye Strain
 - Minimizing Distractions
 - Managing Screen Time
- 11. Cultivating a Reading Routine And Alien Introgression Wheat Cytogenetics Molecular
 - Setting Reading Goals And Alien Introgression Wheat Cytogenetics Molecular
 - Carving Out Dedicated Reading Time
- 12. Sourcing Reliable Information of And Alien Introgression Wheat Cytogenetics Molecular

- Fact-Checking eBook Content of And Alien Introgression Wheat Cytogenetics Molecular
- Distinguishing Credible Sources
- 13. Promoting Lifelong Learning
 - Utilizing eBooks for Skill Development
 - Exploring Educational eBooks
- 14. Embracing eBook Trends
 - Integration of Multimedia Elements
 - Interactive and Gamified eBooks

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