

# Quantitative Genetics, Genomics and Plant Breeding



Edited by  
**Manjit S. Kang**



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# Quantitative Genetics Genomics And Plant Breeding

**Manjit S Kang**



## **Quantitative Genetics Genomics And Plant Breeding:**

**Quantitative Genetics, Genomics and Plant Breeding** Manjit S. Kang, 2020 Quantitative genetics and breeding have made a major contribution to crop improvement over the years but the genomics revolution has dramatically changed the field this second edition of a popular book explains how traditional and genomic techniques have been combined to advance the field Quantitative Genetics, Genomics, and Plant Breeding Amanda Greene, 2018 Quantitative Genetics, Genomics, and Plant Breeding, 2015 Quantitative Genetics, Genomics and Plant Breeding, 2nd Edition Manjit S. Kang, 2020-04-01 This book presents state of the art authoritative chapters on contemporary issues in the broad areas of quantitative genetics genomics and plant breeding Section 1 Chapters 2 to 12 emphasizes the application of genomics and genome and epigenome editing techniques in plant breeding bioinformatics quantitative trait loci mapping and the latest approaches of examining and exploiting genotype environment interactions Section 2 Chapters 13 to 20 represents the intersection of breeding genetics and genomics This section describes the use of cutting edge molecular breeding and quantitative genetics techniques in wheat rice maize root and tuber crops and pearl millet Overall the book focuses on using genomic information to help evaluate traits that can combat biotic abiotic stresses genome wide association mapping high throughput genotyping phenotyping biofortification use of big data orphan crops and gene editing techniques The examples featured are taken from across crop science research and cover a wide geographical base Genetics, Genomics and Breeding of Maize Ramakrishna Wusirika, Martin Bohn, Jinsheng Lai, Chittaranjan Kole, 2014-08-05 Sequencing of the maize genome has opened up new opportunities in maize breeding genetics and genomics research This book highlights modern trends in development of hybrids analysis of genetic diversity molecular breeding comparative and functional genomics epigenomics and proteomics in maize The use of maize in biofuels phytoremediation and **Genetics, Genomics, and Breeding of Tomato** Barbara E. Liedl, Joanne A. Labate, John R. Stommel, Ann Slade, Chittaranjan Kole, 2013-01-17 This volume covers the advances in the study of tomato diversity and taxonomy It examines the mapping of simple and complex traits classical genetics and breeding association studies molecular breeding positional cloning and structural and comparative genomics The contributors also discuss transcriptomics proteomics metabolomics and bioin **Molecular Genetics, Genomics and Biotechnology of Crop Plants Breeding** Søren K. Rasmussen, 2020-05-27 This Special Issue on molecular genetics genomics and biotechnology in crop plant breeding seeks to encourage the use of the tools currently available It features nine research papers that address quality traits grain yield and mutations by exploring cytoplasmic male sterility the delicate control of flowering in rice the removal of anti nutritional factors the use and development of new technologies for non model species marker technology site directed mutagenesis and GMO regulation genomics selection and genome wide association studies how to cope with abiotic stress and an exploration of fruit trees adapted to harsh environments for breeding purposes A further four papers review the genetics of pre harvest spouting readiness for climate

smart crop development genomic selection in the breeding of cereal crops and the large numbers of mutants in straw lignin biosynthesis and deposition

**Cereal Genomics** Pushpendra K. Gupta, R.K. Varshney, 2006-01-19 Cereals make an important component of daily diet of a major section of human population so that their survival mainly depends on the cereal grain production which should match the burgeoning human population. Due to painstaking efforts of plant breeders and geneticists at the global level, cereal production in the past witnessed a steady growth. However, the cereal production in the past has been achieved through the use of high yielding varieties which have a heavy demand of inputs in the form of chemical fertilizers, herbicides, and insecticides/pesticides, leading to environmental degradation. In view of this, while increasing cereal production, one also needs to keep in mind that agronomic practices used for realizing high productivity do not adversely affect the environment. Improvement in cereal production in the past was also achieved through the use of alien genetic variation available in the wild relatives of these cereals, so that conservation and sustainable use of genetic resources is another important area which is currently receiving the attention of plant breeders. The work leading to increased cereal production in the past received strong support from basic research on understanding the cereal genomes which need to be manipulated to yield more from low inputs without any adverse effects as above. Through these basic studies, it also became fairly apparent that the genomes of all cereals are related and were derived from the same lineage million of years ago.

**Physiology and Biotechnology Integration for Plant Breeding** Henry T. Nguyen, Abraham Blum, 2004-01-14 Global demand for wheat, rice, corn, and other essential grains is expected to steadily rise over the next twenty years. Meeting this demand by increasing production through increased land use is not very likely, and while better crop management may make a marginal difference, most agriculture experts agree that this anticipated deficit must be met.

*Water and Agricultural Sustainability Strategies* Manjit S. Kang, 2010-03-25 According to the United Nations, 77 million people are expected to face water shortage by 2025 if people continue to use water at the current rate. More water than available would be needed to grow the world's food during the next decade. As a result of scarcity of water, global annual food production losses could reach 350 million tons by 2025. Di

**Encyclopedia of Plant and Crop Science (Print)** Robert M. Goodman, 2004-02-27

Encyclopedia of Plant and Crop Science is the first ever single source reference work to inclusively cover classic and modern studies in plant biology in conjunction with research applications and innovations in crop science and agriculture. From the fundamentals of plant growth and reproduction to developments in agronomy and agricultural science, the encyclopedia's authoritative content nurtures communication between these academically distinct yet intrinsically related fields, offering a spread of clear, descriptive, and concise entries to optimally serve scientists, agriculturalists, policy makers, students, and the general public. ALSO AVAILABLE ONLINE This Taylor Francis encyclopedia is also available through online subscription, offering a variety of extra benefits for both researchers, students, and librarians, including Citation tracking and alerts, Active reference linking, Saved searches, and marked lists. HTML and PDF format options. For more information, visit Taylor and

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Marcelo J. Carena,2009-04-21 Agriculture depends on improved cultivars and cultivars are developed through proper plant breeding Unfortunately applied plant breeding programs that are focused on cereal commodity crops are under serious erosion because of lack of funding This loss of public support affects breeding continuity objectivity and perhaps equally important the training of future plant breeders and the utilization and improvement of plant genetic resources currently available Breeding programs should focus not only on short term research goals but also on long term genetic improvement of germplasm The research products of breeding programs are important not only for food security but also for commodity oriented public and private programs especially in the fringes of crop production Breeding strategies used for long term selection are often neglected but the reality is that long term research is needed for the success of short term products An excellent example is that genetically broad based public germplasm has significantly been utilized and recycled by industry producing billions of dollars for industry and farmers before intellectual property rights were available Successful examples of breeding continuity have served the sustainable cereal crop production that we currently have The fact that farmers rely on public and private breeding institutions for solving long term challenges should influence policy makers to reverse this trend of reduced funding Joint cooperation between industry and public institutions would be a good example to follow The objective of this volume is to increase the utilization of useful genetic resources and increase awareness of the relative value and impact of plant breeding and biotechnology That should lead to a more sustainable crop production and ultimately food security Applied plant breeding will continue to be the foundation to which molecular markers are applied Focusing useful molecular techniques on the right traits will build a strong linkage between genomics and plant breeding and lead to new and better cultivars Therefore more than ever there is a need for better communication and cooperation among scientists in the plant breeding and biotechnology areas We have an opportunity to greatly enhance agricultural production by applying the results of this research to meet the growing demands for food security and environmental conservation Ensuring strong applied plant breeding programs with successful application of molecular markers will be essential in ensuring such sustainable use of plant genetic resources Applications of Bioinformatics in Rice Research Manoj Kumar Gupta,Lambodar Behera,2021-09-24 This book summarizes the advanced computational methods for mapping high density linkages and quantitative trait loci in the rice genome It also discusses the tools for analyzing metabolomics identifying complex polyploidy genomes and decoding the extrachromosomal genome in rice Further the book highlights the application of CRISPR Cas technology and methods for understanding the evolutionary development and the de novo evolution of genes in rice Lastly it discusses the role of artificial intelligence and machine learning in rice research and computational tools to analyze plant pathogen co evolution in rice crops Search for New Genes Benjamin Peary Pal,2007 Contributed papers presented at the

Symposium festschrift for Benjamin Peary Pal 1906 1989 Indian agricultural scientist Genomics of Cereal Crops Shabir Hussain Wani, Anuj Kumar, 2022-06-13 This volume details different genomic methods and resources to explore cereal genomics Chapters guide readers through crop genomes Next Generation Sequencing NGS technologies protocol for CRISPR editing transgenic wheat NGS approach virus induced gene silencing VIGS genomic tools computational prediction of ncRNAs miRNAs ceRNAs in cereal crops genotyping by sequencing GBS Bayesian method single cell sequencing genome wide association study GWAS QTL interval mapping whole genome bisulfite sequencing genome imprinting and methods for study the receptor metabolite interaction Authoritative and cutting edge Genomics of Cereal Crops aims to be a useful and practical guide to new researchers and experts looking to expand their knowledge **Transcription Factors for Abiotic Stress Tolerance in Plants** Shabir Hussain Wani, 2020-08-05 Transcription Factors for Abiotic Stress Tolerance in Plants highlights advances in the understanding of the regulatory network that impacts plant health and production providing important insights for improving plant resistance Plant production worldwide is suffering serious losses due to widespread abiotic stresses increasing as a result of global climate change Frequently more than one abiotic stress can occur at once for example extreme temperature and osmotic stress which increases the complexity of these environmental stresses Modern genetic engineering technologies are one of the promising tools for development of plants with efficient yields and resilience to abiotic stresses Hence deciphering the molecular mechanisms and identifying the abiotic stress associated genes that control plant response to abiotic stresses is a vital requirement in developing plants with increased abiotic stress resilience Addressing the various complexities of transcriptional regulation this book includes chapters on cross talk and central regulation regulatory networks the role of DOF WRKY and NAC transcription factors zinc finger proteins CRISPR CAS9 based genome editing C Repeat CRT binding factors CBFs Dehydration responsive element binding factors DREBs and factors impacting salt cold and phosphorous stress levels as well as transcriptional modulation of genes involved in nanomaterial plant interactions Transcription Factors for Abiotic Stress Tolerance in Plants provides a useful reference by unravelling the transcriptional regulatory networks in plants Researchers and advanced students will find this book a valuable reference for understanding this vital area Discusses abiotic stress tolerance and adaptive mechanisms based on the findings generated by unlocking the transcriptional regulatory network in plants Presents various kinds of regulatory gene networks identified for drought salinity cold and heat stress in plants Highlights urgent climate change issues in plants and their mitigation using modern biotechnological tools including genome editing **Fruits and Nuts** Chittaranjan Kole, 2007-03-06 Fruits and Nuts form the largest group among crop plants Several constraints such as long life cycle have caused comparatively slow research progress in the past The chapters on 20 fruit and nut crops authored by 56 renowned scientists from 12 countries include for the first time comprehensive reviews on a variety of fruits and nuts The huge amount of information hitherto dispersed in journals is now available in a clearly structured reference work **Advances in**

**Agronomy** Donald L. Sparks, 2012-04-24 *Advances in Agronomy* continues to be recognized as a leading reference and a first rate source for the latest research in agronomy As always the subjects covered are varied and exemplary of the myriad of subject matter dealt with by this long running serial Maintains the highest impact factor among serial publications in agriculture Presents timely reviews on important agronomy issues Enjoys a long standing reputation for excellence in the field

**Handbook of Statistical Genetics** David J. Balding, Martin Bishop, Chris Cannings, 2008-06-10 The Handbook for Statistical Genetics is widely regarded as the reference work in the field However the field has developed considerably over the past three years In particular the modeling of genetic networks has advanced considerably via the evolution of microarray analysis As a consequence the 3rd edition of the handbook contains a much expanded section on Network Modeling including 5 new chapters covering metabolic networks graphical modeling and inference and simulation of pedigrees and genealogies Other chapters new to the 3rd edition include Human Population Genetics Genome wide Association Studies Family based Association Studies Pharmacogenetics Epigenetics Ethic and Insurance As with the second Edition the Handbook includes a glossary of terms acronyms and abbreviations and features extensive cross referencing between the chapters tying the different areas together With heavy use of up to date examples real life case studies and references to web based resources this continues to be must have reference in a vital area of research Edited by the leading international authorities in the field David Balding Department of Epidemiology Public Health Imperial College An advisor for our Probability Statistics series Professor Balding is also a previous Wiley author having written *Weight of Evidence for Forensic DNA Profiles* as well as having edited the two previous editions of HSG With over 20 years teaching experience he s also had dozens of articles published in numerous international journals Martin Bishop Head of the Bioinformatics Division at the HGMP Resource Centre As well as the first two editions of HSG Dr Bishop has edited a number of introductory books on the application of informatics to molecular biology and genetics He is the Associate Editor of the journal *Bioinformatics* and Managing Editor of *Briefings in Bioinformatics* Chris Cannings Division of Genomic Medicine University of Sheffield With over 40 years teaching in the area Professor Cannings has published over 100 papers and is on the editorial board of many related journals Co editor of the two previous editions of HSG he also authored a book on this topic *Molecular Breeding for Sustainable Crop Improvement*

Vijay Rani Rajpal, S. Rama Rao, S.N. Raina, 2016-02-02 The world population is estimated to reach to more than 10 billion by the year 2050 These projections pose a challenging situation for the agricultural scientists to increase crops productivity to meet the growing food demands The unavailability and or inaccessibility to appropriate gene pools with desired traits required to carry out genetic improvement of various crop species make this task formidable for the plant breeders Incidentally most of the desired genes reside in the wild genetic relatives of the crop species Therefore exploration and characterization of wild genetic resources of important crop species is vital for the efficient utilization of these gene pools for sustainable genetic improvements to assure food security Further understanding the myriad

complexities of genic and genomic interactions among species more particularly of wild relatives of crop species and or phylogenetically distant germplasm can provide the necessary inputs to increase the effectiveness of genetic improvement through traditional and or genetic engineering methods This book provides comprehensive and latest insights on the evolutionary genesis of diversity access and its utilization in the evolution of various crop species A comprehensive account of various crops origin exploitation of the primary secondary and tertiary gene pools through breeding biosystematical cytogenetical and molecular phylogenetical relationships and genetic enhancement through biotechnological interventions among others have been provided as the necessary underpinnings to consolidate information on the effective and sustainable utilization of the related genetic resources The book stresses upon the importance of wild germplasm exploration characterization and exploitation in the assimilation of important crop species The book is especially intended for students and scientists working on the genetic improvement of crop species Plant Breeders Geneticists Taxonomists Molecular Biologists and Plant Biotechnologists working on crop species are going to find this book very useful



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