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This book assesses the potential effects of biotechnological approaches, particularly genetic modification, on the present state of fiber crop cultivation and sustainable production. Leading international researchers discuss and explain how biotechnology can affect and solve problems in connection with fiber crops. The topics covered include biology, biotechnology, genomics and applications of fiber crops like cotton, flax, jute and bamboo. Providing complete, comprehensive and broad subject-based reviews, the book offers a valuable resource for students, teachers, and researchers including agriculturists, biotechnologists and botanists, as well as industrialists and government agencies involved in the planning of fiber crop cultivation.

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Nuclear Trafficking is a summary of the state of knowledge in nuclear trafficking, and is organized into five parts. The book begins by discussing the diffusion and signal-mediated transport through the pores. It then looks into the detailed accounts of pore structure and composition, nuclear localization signals, signal binding proteins, RNA efflux, and biochemical factors influencing nucleocytoplasmic exchange. This book will be very useful to those people new to this field of interest.

Featuring current resources used to discover new legume family genes and to understand genes and their interactions, Legume Genomics: Methods and Protocols provides techniques from expert researchers to study these plants that are so vitally important for food, feed, human nutrition, bioenergy, and industrial purposes. This detailed volume covers genome characterization and analysis, transcriptome analysis and miRNA identification/analysis, forward and reverse genetics, molecular markers, as well as transformation strategies used to investigate gene function and many other topics. Written in the highly successful Methods in Molecular Biology series format, chapters include introductions to their respective topics, lists of the necessary materials and reagents, step-by-step, readily reproducible laboratory protocols, and tips on troubleshooting and avoiding known pitfalls. Authoritative and useful, Legume Genomics: Methods and Protocols aims to serve plant molecular biologists, molecular breeders, plant physiologists and biochemists, developmental biologists, and those interested in plant-microbe interactions.

This volume is an exhaustive source of information on the control and regulation of flowering. It presents data on the factors controlling flower induction and how they may be affected by climate and chemical treatments. For each plant, specific information is provided on all aspects of flower development, including sex expression, requirements for flowering initiation and development, photoperiod, light density, vernalization, and other temperature effects and interactions. Individual species are described from the standpoint of juvenility and maturation, morphology, induction and morphogenesis to anthesis. All information is presented alphabetically for easy reference.

This text provides an overview of the emerging field of the study of complex plant genomes and the laboratory techniques that will enhance genetic improvement of cotton. It also addresses important issues and opportunities that face the cotton industry. Topics include the approaches utilized in breeding and development of cotton cultivars in Australia; the expression and regulation of lipid transfer protein genes in cotton fibre; and RFLP diversity in cotton.

It is perhaps not surprising that plants have evolved a mechanism to sense the light environment about them and to modify growth for optimal use of the available 'life-giving' light. Green plants, and ultimately all forms of life, depend on the energy of sunlight fixed during photosynthesis. Unlike animals that use behaviour to find food, sedentary plants use physiology to optimize their growth and development for light absorption. By appreciating the quality, quantity, direction and duration of light, plants can control such complex processes as germination, growth and flowering. To perceive the light environment several receptor pigments have evolved, including the red/far-red reversible phytochrome and the blue/UV-absorbing photoreceptors (Part 1). The quantification of light (Part 2) and importance of instrumentation for photomorphogenesis research are introduced in Part 3. Isolation and characterization of phytochrome is a classic example of how photobiological techniques can predict the nature of an unknown photoreceptor. Current knowledge of the phytochrome photoreceptor family is given in Part 4 and that of blue/UV receptors in Part 5. Part 6 deals with the coaction of photoreceptors. The light environment and its perception is addressed in Part 7. Molecular and genetic approaches and the photoregulation of gene

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expression compose Part 8. Part 9 contains further selected topics: photomodulation of growth phototropism, photobiology of stomatal movements, photomovement, photocontrol of flavonoid biosynthesis, photobiology of fungi and photobiology of ferns. The 28 chapters written by leading experts from Europe, Israel, Japan and the USA, provide an advanced treatise on the exciting and rapidly developing field of plant photomorphogenesis.

Cotton, a source of natural fiber for textile industry, has a long breeding history aiming at increasing cotton fiber yield and its quality. Newly developed cotton varieties poorly respond in low-input environments. Secondly, the impact of changing climate may threaten the cotton production in the future. To address these challenges, efforts toward the development of resilient cotton varieties have been initiated using genetic and modern genomic approaches. In this book, research updates on cotton fiber types and properties, DNA markers for selecting desirable cotton plants, and cotton fiber genomics were compiled. Also, the modern breeding trends including development of transgenic cotton and the biosafety studies and possibilities of improving cotton genome using modern genome editing tools were also compressively discussed.

Genetic approaches to understanding plant growth and development have always benefitted from screens that are simple, quantitative and rapid. Visual screens and morphometric analysis have yielded a plethora of interesting mutants and traits that have provided insight into complex regulatory pathways, and yet many genes within any given plant genome remain undefined. The premise underlying High Throughput Phenotyping in Plants: Methods and Protocols is that the higher the resolution of the phenotype analysis the more likely that new genes and complex interactions will be revealed. The methods described in this volume can be generally classified as quantitative profiling of cellular components, ranging from ions to small molecule metabolites and nuclear DNA, or image capture that ranges in resolution from chlorophyll fluorescence from leaves and time-lapse images of seedling shoots and roots to individual plants within a population at a field site. Written in the successful Methods in Molecular Biology™ series format, chapters include introductions to their respective topics, lists of the necessary materials and reagents, step-by-step, readily reproducible protocols, and notes on troubleshooting and avoiding known pitfalls. Authoritative and easily accessible, High Throughput Phenotyping in Plants: Methods and Protocols serves as an invaluable guide to plant researchers and all scientists who wish to better understand plant growth and development.

Brassica crop species and their allies (*Raphanus*, *Sinapis*, *Eruca*, etc.) are important sources of edible roots, stems, leaves, buds and inflorescences, as well as of edible or industrial oils, condiments and forage. Many well known names of plants or plant products, such as kale, cabbage, brocolli, cauliflower, Brussels sprouts, kohlrabi, Chinese cabbage, turnip, rape, rutabaga, swede, colza or rapeseed, canola, mustard, rocket, etc. are directly associated to this botanical group. The scientific interest for this botanical group has run parallel to its economical importance, and research achievements in our days would have certainly appeared unimaginable only two decades ago. As the end of the millenium approaches, entirely new fields (transformation, somatic fusion, etc.) have been added to the classical ones. Thus, nobody can doubt the opportuneness of this book, which combines and presents both the basic and applied biological aspects of the Brassica species.

Cotton fiber is the most important natural fiber used in the textile industry. The physical structure and chemical compositions of cotton fibers have been extensively studied. Newer high speed spinning instruments are being deployed around the world that demand longer, stronger and finer fibers. Consequently, genetic improvement in fiber quality has been stressed. With improvement in fiber quality has come the realization that further fiber improvement will require a better understanding of fiber development and biology. As a consequence, cotton fiber developmental biology, genetics and

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genomics have become focal points in the cotton research community. As the longest single-celled plant hair, cotton fiber has been used as an experiment model to study trichome initiation and elongation in plants. This book provides a comprehensive update on cotton fiber physics, chemistry and biology that form the three sections of the book. In the physics section, the physical structure of cotton fiber is first illustrated in great detail. Then a suite of fiber properties and their measuring methods are described. The pros and cons of each method are outlined. New methods to measure physical properties of single fiber and young developing fibers are included. In the chemistry section, the chemical compositions of cotton fibers are described in detail. This knowledge is necessary for efficient modification of cotton fibers for better and broader utilization. The advancement in cotton fiber modification using chemical and enzymatic methods opened new ways to utilize cotton fibers. In the biology section, the book first introduces the utilization of naturally occurring color cottons. Color cottons possess unique attributes such as better fire retardant ability. Advancement in understanding fiber color genetics and biochemical pathways and new utilization of color cottons are discussed. Recent technological advancements in molecular biology and genomics have enabled us to study fiber development in great depth. Many genes and quantitative trait loci related to fiber quality attributes have been identified and genetically mapped. Some of these genes and QTLs are being used in breeding. Progresses in cotton fiber improvement using breeding and biotechnology are discussed in the last chapter. This book serves as a reference for researchers, students, processors, and regulators who either conduct research in cotton fiber improvement or utilize cotton fibers.

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