

1. Record Nr.	UNINA9910483986203321
Titolo	Cotton precision breeding / / Mehboob Ur Rahman, Yusuf Zafar and Tianzhen Zhang (editors)
Pubbl/distr/stampa	Cham, Switzerland : , : Springer, , [2021] ©2021
ISBN	3-030-64504-5
Edizione	[1st ed. 2021.]
Descrizione fisica	1 online resource (XII, 459 p. 43 illus., 37 illus. in color.)
Disciplina	633.51
Soggetti	Cotton - Genetics Cotton - Breeding Cotó Conreu del cotoner Llibres electrònics
Lingua di pubblicazione	Inglese
Formato	Materiale a stampa
Livello bibliografico	Monografia
Nota di bibliografia	Includes bibliographical references and index.
Nota di contenuto	1. Historical perspective of cotton improvement -- 2. Challenges to sustainable cotton production worldwide -- 3. New DNA markers assays -- 4. DNA markers for resistance to diseases -- 5. DNA markers for tolerance to abiotic stresses -- 6. DNA markers for cotton fiber traits -- 7. Evolutionary dynamics of Cotton Genome -- 8. Sequencing <i>Gossypium</i> genomes: Opportunities & Challenges -- 9. Chloroplast genome -- 10 Functional genomic studies in cotton -- 11. Comparative genomics studies of cotton -- 12. First generation transgenic cotton crops -- 13. Prospects of 2nd and 3rd generation transgenic crops -- 14. GM cotton for stress environments -- 15. GM technology and Fiber traits -- 16. Critical evaluation of GM cotton -- 17. Role of private sector in cotton sustainability -- 18. CRISPR-based genome editing tools for improving resistance to insect pests and diseases in cotton -- 19. Using CRISPR-based genome editing tools for improving tolerance to abiotic stresses in cotton -- 20. Application of CRISPR-based genome editing tools for improving fiber traits in cotton -- 21. Cotton crop beyond 2030.
Sommario/riassunto	Cotton, the most important natural fiber crop, has been improved by

conventional breeding—largely through planned hybridization of different cotton genotypes, since the discovery of Mendelian genetics. All these efforts resulted in the development of resilient high yielding cotton varieties. However, the progress through conventional breeding procedures is slow because of long lag periods for developing a variety, little control over the new genetic combinations, unwanted traits and lack of foolproof performance testing system. Genomic assays discovered over the last two decades have made it possible to understand the “language” of the genome by associating the genes with specific traits. Together with the more recently established gene-editing tools like CRISPR-Cas9, the cotton genome can be tailored much more precisely than ever before. In this regard, genetic information has been harnessed, through (i) sequencing of the progenitor and cultivated cotton species, (ii) ongoing mega pan-genome sequencing projects, (iii) genetic and physical mapping, and (iv) introgression of genes from alien sources, that resulted in the development of resilient cotton cultivars. These technologies have been deployed or are attempting to overcome the challenges of water shortage, excessive heat in most cotton growing regions, infectious diseases and infestation of insect pests, as well as rising production cost, for sustainable cotton production beyond 2030. In this book, new knowledge generated by the cotton research community and its application for developing resilient cotton are comprehensively summarized. This book contributed by well-known cotton researchers is a timely collection of the challenges and successes of precision cotton breeding in a changing environment.

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2. Record Nr.	UNINA9910697471203321
Titolo	Davy Crockett makes history with a little feller at the NTS [[electronic resource]]
Pubbl/distr/stampa	[Las Vegas, Nev.] : , : National Nuclear Security Administration, , [2007]
Descrizione fisica	2 unnumbered pages : digital, PDF file
Soggetti	Nuclear weapons - Nevada - Testing - History
Lingua di pubblicazione	Inglese
Formato	Materiale a stampa
Livello bibliografico	Monografia
Note generali	Title from title screen (viewed on Aug. 25, 2008). "April 2007." "DOE/NV -- 1200."