

AN OVERVIEW OF BREEDING FOR DROUGHT STRESS TOLERANCE IN COTTON

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Abstract: Drought is a main non-living factor that causes severe crop yield loss globally. Given the strengthening and reappearance of drought events and their impacts, it's important to deepen our understanding as a key to subsidizing mechanisms for drought training and mitigation plans. Pakistan is ranked maximum of the top 5 biggest cotton manufacturers, the seventh largest material producer international, and cotton contributes 10% to the country-wide GDP compared to the overall agriculture area GDP percentage of 18.9%. Cotton farming performs a tremendous role in presenting direct livelihood to 11 million farmers. The cotton crop, in particular, is confined to northern, imperative, and southern zones, with approximately 90 in keeping with cent of the area coming beneath 3 zones. Regardless of this, its cumulative, not apparent impact and multidimensional nature significantly impact the cotton plant's morphological, physiological, biochemical, and molecular attributes with a detrimental impact on photosynthetic capability. Dealing with water scarcity, plants evolve various complicated resistance and edition mechanisms, including physiological and biochemical responses, which range with species stage. The sophisticated adaptation mechanisms and regularity community that improve the water stress tolerance and version in plants are briefly discussed. Growth pattern and structural dynamics, reduction in transpiration loss via altering stomatal conductance and distribution, leaf rolling, root-to-shoot ratio dynamics, root duration increment, accumulation of like-minded solutes, enhancement in transpiration performance, osmotic and hormonal regulation, and behind-schedule senescence are the techniques that are followed using cotton plant life underneath water deficit. Approaches for drought stress resistance we develop transgenic cotton plants which which can tolerate drought stress to improve cotton quality with good yield.

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Introduction

With the worldwide weather alternate and sturdy water demand for the improvement of destiny sustainable agriculture, drought is turning into one of the foremost abiotic stresses that seriously outcomes in decreased increase and crop yield loss (FAO, 2018). Drought can set off a complicated array of flower responses, including decreased turgor stress and photosynthesis prices, stomatal closure, and adjusted leaf gas change (Farooq et al., 2012). Plants have developed huge physio- biochemical, cell, and molecular versions of drought strains to complete their lifecycle (Naseem et al., 2020; Yamaguchi-Shinozaki and Shinozaki, 2006). Previous reviews have documented the one-of-a-kind plant's drought-responsive genes and essential mechanisms (Balqees et al., 2020; Li et al., 2017; Mittal et al., 2014; Xu et al., 2018). However, the underlying molecular mechanisms responding to drought strain are poorly elucidated. Mining the key genes to increase drought-resistant varieties is still a cost-

effective and green technique for drought-resistant breeding. Drought can weaken meal protection and contributes to rural flight and migration, mainly in developing countries. Semiarid regions constitute approximately 15% of the globe and encompass approximately 15% of the global population (Idrees et al., 2022; Safriel et al., 2005; Zahoor et al., 2022). Agricultural production in those regions, characterized in big elements with rainfed systems, is exceedingly imposed to weather trade (Porter et al., 2014; Schwinning et al., 2004). additionally, the future conditions of improved temperature and reduced precipitation in these regions may cause important screw-ups (Dai, 2013; Huang et al., 2016; Iqra et al., 2020). In Brazil, weather exchange can also affect agricultural income, meals safety, and, consequently, the neighborhood economy, particularly within the Brazilian Semiarid region (BSR) (Marengo et al., 2019), located in the Northeast. The BSR, which focuses maximum on the poorest population, is the most susceptible to the



consequences of droughts (Marengo et al., 2017). The recurrence of extreme droughts negatively affects the growth of flowers, lowering their productiveness, and contributing to land degradation, particularly on arid and semi-arid lands (Mariano et al., 2018; Tomasella et al., 2018; Vicente-Serrano et al., 2015).

Cotton

Cotton is one of the widespread fibre and gold mine crops of India and plays a critical position in the cotton industrial and agricultural financial system, which provides employment to about eleven million farmers and circuitously about forty-50 million humans are employed in numerous ranges of processing and alternate of cotton and its prototypical (MURALI and KHAN, 2022; Rafi et al., 2022). Cotton is considered as 'White gold' being cultivated by the archaic and principal producer, patron and exporter of cotton globally (Atif et al., 2022; Dupdal and Patil, 2018; Fatima et al., 2022a; Fatima et al., 2022b; Masood et al., 2022). In India, cotton is cultivated on 12.90 million hectares, constituting forty-one per cent of the world's location. In addition, at some point in 2019-20, the cotton area reached a new report of 13.40 million ha with 361 lakh bales manufacturing (one hundred seventy kgs in step with bale).

Cotton manufacturing and Pakistan illustration

The observation uses Pakistan's cotton manufacturing as a standard case to research production elements from far nearer input issues and performance courting attitude to symbolize the non-meals crop in a developing united states perspective for the following motives (Arshad et al., 2022). First, Pakistan is ranked among the top 5 largest cotton producers, the seventh largest cloth producer worldwide. Cotton contributes 10% to the country-wide GDP compared to the general agriculture quarter GDP percentage of 18.9% (Azumah et al., 2019). This area also contributes to 42.3% of the hard work pressure with employment. It gives the raw substances for lots of fee-delivered sectors (Sohaib and Jamil, 2017), with 55% of the overseas profits contribution, a proportion majorly ruled by using cotton-based finished products (Arshad et al., 2021). Due to the fact Pakistan is a growing us of a those states reputedly prefer the export; in comparison, Pakistan has been uploading raw cotton for decades. Absolutely, Pakistan has not been exporting uncooked cotton since 2010, it is the fourth united states that end up finishing stalk (USDA cotton outlook 2019). Second, the main reason for increasing the import of uncooked cotton is that Pakistan's cotton yield per hectare has had a step-by-step reducing trend over time, and the numbers are a few of the lowest within the world; even the nations which have a miles smaller geographic area have a higher yield according to hectare than Pakistan (Arshad et al., 2022). On average, the yield of cotton in Pakistan is 730 kg/ha

with 10,671 million bales, that's 1.5–2% decrease than to the relaxation of the world, or even irrigated areas of the countries are lagging in phrases of lint per hectare from the rain-fed cotton-developing areas of the arena. As a result, cotton cultivation has emerged as less attractive than developing different plants, which ends up minimizing the vicinity of cotton vegetation (international textile information document 2017, USDA monetary survey of Pakistan (2019–2020). it is real that the area of cotton crops has been declining in Pakistan due to the fact 2004–2005; unluckily, this situation has been irritated since 2013–2014, and through the continued year, 14.2 percentage sowing area of the crop had decreased (Nadeem et al., 2014). Earlier than 2014–2015, it became 2.902 million hectares, whilst presently 2.489 million hectares of cotton are cultivated in Pakistan. Third, this low productivity of Pakistan's cotton crop in particular consequences from inefficient use of a couple of elements related to irrigation water, plant populace, disorder safety, plant nutrients, useful resource management competencies, and insufficient era, resulting in an opening among potential production and real production (agriculture statistics of Pakistan (2019). however, as a general trend, Pakistan farmers are looking to treat this low yield per hectare via traditional practices, along with increasing the use of crop inputs (intensification) or bringing new lands into manufacturing (extensification), which results in continuously deteriorating of the rural ecological environment. Such practices lessen the monetary impact of agriculture and damage the sources and environment on which agriculture depends for survival and development (Chen and Breedlove, 2020).

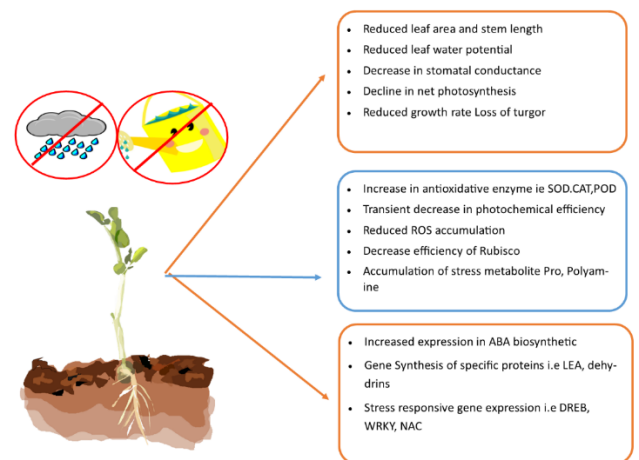


Figure 1 Effects of Drought on cotton

Effects of drought on cotton: Cotton (*Gossypium hirsutum* L.) is considered a main cash crop in the arena, with a major contribution from China, America, India, Pakistan and Brazil (Meyer, 2019).

Climate trade has threatened global agriculture because of the severity of biotic and abiotic stresses. Principal threats to cotton manufacturing in China, India, America, Pakistan, Brazil and different tropical regions include much less availability of irrigation water, exchange in frequency and depth of rainfall, drought and heat spells and salinization (Ton, 2011). Cotton has an indeterminate boom addiction and is susceptible to environmental anomalies (Rehman and Farooq, 2019). Being a crop of tropical and subtropical areas, cotton is moderately tolerant to drought strain specially for vegetative increase, but its reproductive increase is surprisingly sensitive to drought pressure (Iqbal et al., 2017; Niu et al., 2018; Wang et al., 2016). Drought is the maximum essential abiotic stress, and ~20% of the worldwide land area is dealing with moderate to excessive drought (Barichivich et al., 2019), that's projected to increase in the future. As cotton is a crop of tropical and sub-tropical weather (regions which can be more at risk of drought strain), its production is threatened by lengthy dry spells in the destiny (Meyer, 2019). To plot an approach to decrease the outcomes of drought stress on fibre production (cotton is a fibre crop), it's vital to understand the impact of drought on fibre improvement and yield. The major effects of drought are shown in **Error! Reference source not found.**

Irrigation required

Generally, cotton growth and development are divided into 5 ranges with appreciation to irrigation requirements, i.e., planting to emergence, emergence to first development, first square to the first flower, the first flower to height bloom and peak bloom first to boll open (Bauer et al., 2012). At planting time, irrigation is essential for crop germination and establishment. From emergence to the first square, drought strain has little impact on yield, or even mild drought stress is useful to sell root growth (Bauer et al., 2012; Zonta et al., 2017). The subsequent two levels, i.e., first rectangular to first flower and first flower to peak bloom, are the most sensitive degrees to abiotic stresses, specifically drought and heat strain. At these stages, drought stress impacts the development of fruiting websites and motive abortion of the current fruiting structure and, as a result, causes reduced yield. (Bauer et al., 2012; Iqbal et al., 2017; Snowden et al., 2014; Zonta et al., 2017). At the ultimate stage, i.e., top bloom to first boll open, crop water requirement is reduced, and slight drought pressure has highly less effect on yield, however it impacts fiber characteristics (Bauer et al., 2012; Hussain et al., 2020; Snowden et al., 2014). Moisture pressure adversely has effects fibre duration, power and micronaire price, where the effect located extra stated on top fruiting branches than decreased ones (Wang et al., 2016). The availability of enough water is essential in any respect fiber developmental degrees for higher fibre yield and great (Rehman and Farooq, 2019; Zou et

al., 2016) cause abnormality in photosynthesis, stomatal conductance, ATP synthesis carbohydrate cycle and translocation disturb the biomass limited water supply throughout the reproductive level influences the pollen feature because of decreased activities of starch synthesis enzymes, downregulation of sucrose synthase and invertase gene, curtailed starch accumulation in pistils result in negative pollen tube boom, instigating reproductive failure (Zou et al., 2016), poor fibre yield and quality of cotton

Mechanism Of Drought Tolerance

Various accommodative mechanisms that make plants more tolerant to the unfavourable effects of drought stress have been developed through development (Batool et al., 2020). Stress avoidance, escape, and tolerance are the three important schemes that plants use when unmasking drought pressure. Consequently, flora comments to drought pressure vary from the molecular to plant level (Galindo et al., 2018). The mechanisms of the plant getaway, avoidance and tolerance towards drought pressure are discussed in the following sections.

Escape

To get away the unfavorable outcomes of drought pressure on plant productiveness, a few floras make use of mechanisms regarding speedy plant development and shortening of the lifestyles cycle, self-reproduction, and seasonal boom earlier than the start of the driest part of the year (Álvarez et al., 2018). among these mechanisms, early flowering is perhaps the pleasant possible break out adaptive mechanism in plant life (Tekle and Alemu, 2016), although this mechanism can mean a big reduction within the period of the plant growing period and the final plant productivity in a few instances.

Avoidance

beneath the avoidance strategy, plant water ability is maintained high through a reduction within the stomatal transpiration losses and the increase of water uptake from nicely established root structures (Dobra et al., 2010). but, overdevelopment of those systems has a price for the plant in terms of discounts in plant productivity and reduced average size of vegetative and reproductive elements of the plant (Wasaya et al., 2018).

Tolerance

An adaptive tolerance mechanism on the photosynthetic machinery stage includes discounts inside the plant leaf region and boundaries in the enlargement of new leaves. similarly, trichomes manufacturing on both facet of the leaves are exomorphic attributes that permit the plant to tolerate water deficits in dry environments (Zhang et al., 2019). Those structures lessen the leaf temperature via increasing the rate of mild reflection in the leaf and additionally by including any other extra layer of resistance to the water loss. therefore the fee of water loss thru leaf transpiration is decreased (Tiwari et al., 2021). However, it is widely familiar that

adjustments inside the root machine, including root size, density, period, proliferation, expansion and growth rate, represent the primary strategy for drought-tolerant plant life to manage against water deficits (Tzortzakis et al., 2020).

Alleviate the drought stress to reduce the cotton yield loss

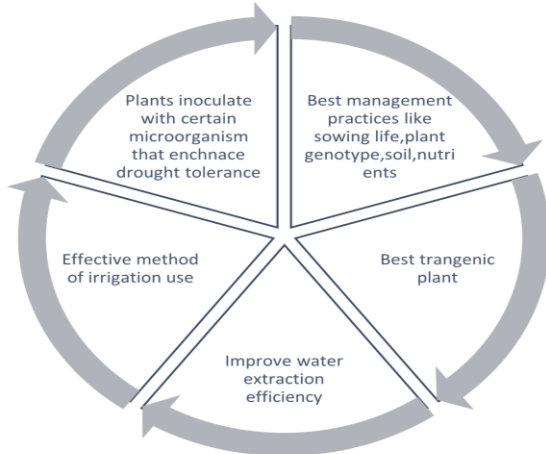


Figure 2 How to improve the effects of drought stress

Drought Stress can be minimized by following step as shown in **Error! Reference source not found.**

Methods for improvement

Conventional breeding

Interspecific populations derived from crossings between *G. hirsutum* and *G. barbadense* explain the conventional trouble of limited genetic range in Upland cotton genetic and QTL mapping in the early days. Those two grown species are searched for exclusive trends further to supply the DNA-degree polymorphism required for genetic map creation. As previously said, *G. hirsutum* breeding has confused maximal output and extensive adaptability, whereas *G. barbadense* breeding emphasises fibre first-class. As a result, most genetic mapping (Reinisch et al., 1994; Wang et al., 2015) and molecular quantitative genetic research of fibre properties have used populations derived from interspecific hybridization related to wild and domesticated forms of *G. barbadense* crossed with Upland cotton. QTLs for several fibre nice variables mapped in cotton have been summarized using Chee and Campbell (Campbell, 2021). A few important QTLs for fibre duration, energy, and fineness have now been found, and several were verified (Cao et al., 2015). This information offers cotton growers greater options to enhance positive fibre features in upland cotton by introducing genes from *G. barbadense* with minimum disruption to the favourable allelic combinations developed over a century of selective choice. Inbred backcross populations generated from crossing Upland types with the allotetraploid species *G. barbadense*, *G. tomentose*, and *G. musteline* were evolved as a part of a collaborative attempt to

minimize Upland cotton's genetic susceptibility (Paterson et al., 2004). Inbred backcrossing became used to reduce reproductive limitations due to interspecific introgression between those species (Baohua and Peng, 2010; Tanksley and Nelson, 1996). one could study very tiny portions of introjected DNA for agronomic or fibre exceptional overall performance and examine them for QTLs by way of establishing a complete set of close to Isogenic Introgression strains from the BC2 or BC3 families. Because recombination and segregation have split the donor genome into smaller components, the hobby of specific genetic loci can be more surely described than in preceding generations. Rong, Feltus, Waghmare, Pierce, Chee, Draye, Saranga, Wright, Wilkins and may (Rong et al., 2007) said the alignment of 432 fiber QTLs recognized in 10 interspecific *G. hirsutum* by way of *G. barbadense* populations right into a consensus map, presenting further records on the genetic dissection of each of the fiber residences.

Non-Conventional Breeding

Molecular markers are very useful for molecular characterization and identity of genetic variants and have been used in marker-assisted choice (MAS) and genome fingerprinting (Kalia et al., 2011; Munir et al., 2022). molecular markers are vital in genomics studies that they may or might not link with the phenotypic expression of a man or woman in an organism (Agarwal et al., 2008). In cotton genomes, the maximum crucial molecular markers are polymerase chain response (PCR)-primarily based markers because of their high effectiveness and utilization, which include inter simple collection repeats (ISSRs) (Ahmad et al., 2021; Pradeep Reddy et al., 2002), amplified fragment period polymorphism (AFLP) (Abdalla et al., 2001; Álvarez and Wendel, 2006), simple sequence repeats (SSRs) (Zhou et al., 2014) and random amplified polymorphic DNA (RAPD) some of the genomic assets, there are approximately 16 162 SSRs and 312 mapped cotton RFLP sequences to be had publicly. The RFLP, SSR, AFLP, AFLP and RAPD markers were applied in distinct mapping populations to increase linkage maps. It has been reported that the identity of DNA markers is related to over 29 important developments, such as fibre first-class and yield, leaf and flower morphology, trichomes density and distribution, and ailment resistance (Farooq et al., 2021; Shaheen et al., 2012; Tahir et al., 2020).

Conclusion

Drought stress affects the production of cotton vegetative, reproductive, and fiber quality and causes low yield that's why we adopted the best method to alleviate the drought stress and developed a good variety of cotton by best management practices like sowing life, plant genotype, soil, nutrients, the best transgenic plant, improve water extraction efficiency, effective method of irrigation use, plants

inoculated with a certain microorganism that enhance drought tolerance.

Conflict of interest

The authors declared absence of conflict of interest.

References

- Abdalla, A., Reddy, O., El-Zik, K., and Pepper, A. (2001). Genetic diversity and relationships of diploid and tetraploid cottons revealed using AFLP. *Theoretical and Applied Genetics* **102**, 222-229.
- Agarwal, M., Shrivastava, N., and Padh, H. (2008). Advances in molecular marker techniques and their applications in plant sciences. *Plant cell reports* **27**, 617-631.
- Ahmad, M., Ali, Q., Hafeez, M. M., and Malik, A. (2021). Improvement for biotic and abiotic stress tolerance in crop plants. *Biological and Clinical Sciences Research Journal* **2021**. <https://doi.org/10.54112/bcsrj.v2021i1.50>
- Álvarez, I., and Wendel, J. F. (2006). Cryptic interspecific introgression and genetic differentiation within *Gossypium aridum* (Malvaceae) and its relatives. *Evolution* **60**, 505-517.
- Álvarez, S., Rodríguez, P., Broetto, F., and Sánchez-Blanco, M. J. (2018). Long term responses and adaptive strategies of *Pistacia lentiscus* under moderate and severe deficit irrigation and salinity: Osmotic and elastic adjustment, growth, ion uptake and photosynthetic activity. *Agricultural Water Management* **202**, 253-262.
- Arshad, M. U., Yuanfeng, Z., Yufei, G., Xinya, G., Hanif, S., Ying, G., and Jun, T. (2021). The effect of climate change on cotton productivity-an empirical investigation in Pakistan. *Pakistan Journal of Agricultural Sciences* **58**, 1455-1462.
- Arshad, M. U., Zhao, Y., Hanif, O., and Fatima, F. (2022). Evolution of Overall Cotton Production and Its Determinants: Implications for Developing Countries Using Pakistan Case. *Sustainability* **14**, 840.
- Atif, M., Ahmad, F., Manzoor, M. T., Gilani, K., Ali, Q., Sarwar, M., Anjum, S., Alam, M. W., and Hussain, A. (2022). Application of bioinformatics tools to check mutation and evolution potential of chickpea chlorotic dwarf virus (CPCDV) infecting cotton and host plants. *Biological and Clinical Sciences Research Journal* **2022**. <https://doi.org/10.54112/bcsrj.v2022i1.116>
- Azumah, S. B., Donkoh, S. A., and Awuni, J. A. (2019). Correcting for sample selection in stochastic frontier analysis: insights from rice farmers in Northern Ghana. *Agricultural and food economics* **7**, 1-15.
- Balqees, N., Ali, Q., and Malik, A. (2020). Genetic evaluation for seedling traits of maize and wheat under biogas wastewater, sewage water and drought stress conditions. *Biological and Clinical Sciences Research Journal* **2020**. <https://doi.org/10.54112/bcsrj.v2020i1.38>
- Baohua, W., and Peng, W. C. (2010). Application of advanced backcross QTL analysis in crop improvement. *Journal of plant Breeding and crop Science* **2**, 221-232.
- Barichivich, J., Osborn, T., Harris, I., van der Schrier, G., and Jones, P. (2019). Drought: Monitoring global drought using the self-calibrating Palmer Drought Severity Index. *Bulletin of the American Meteorological Society* **100**, S39-S40.
- Batool, T., Ali, S., Seleiman, M. F., Naveed, N. H., Ali, A., Ahmed, K., Abid, M., Rizwan, M., Shahid, M. R., and Alotaibi, M. (2020). Plant growth promoting rhizobacteria alleviates drought stress in potato in response to suppressive oxidative stress and antioxidant enzymes activities. *Scientific Reports* **10**, 1-19.
- Bauer, P., Faircloth, W., Rowland, D., Ritchie, G., Perry, C., and Barnes, E. (2012). Water-sensitivity of cotton growth stages. *Cotton irrigation management for humid regions. Cary: Cotton Incorporated* **1**, 17-20.
- Campbell, B. T. (2021). Examining the relationship between agronomic performance and fiber quality in ten cotton breeding populations. *Crop Science* **61**, 989-1001.
- Cao, Z., Zhu, X., Chen, H., and Zhang, T. (2015). Fine mapping of clustered quantitative trait loci for fiber quality on chromosome 7 using a *Gossypium barbadense* introgressed line. *Molecular Breeding* **35**, 1-13.
- Chen, G., and Breedlove, J. (2020). The effect of innovation-driven policy on innovation efficiency: Based on the listed sports firms on Chinese new Third Board. *International Journal of Sports Marketing and Sponsorship*.
- Dai, A. (2013). Increasing drought under global warming in observations and models. *Nature climate change* **3**, 52-58.
- Dobra, J., Motyka, V., Dobrev, P., Malbeck, J., Prasil, I. T., Haisel, D., Gaudinova, A., Havlova, M., Gubis, J., and Vankova, R. (2010). Comparison of hormonal responses to heat, drought and combined stress in tobacco plants with elevated proline content. *Journal of plant physiology* **167**, 1360-1370.
- Dupdal, R., and Patil, B. (2018). Production performance and supply response of cotton in Karnataka: A case study of Dharwad district. *International Journal of Pure and Applied Biosciences* **6**, 1184-1189.
- FAO, F. (2018). The future of food and agriculture: alternative pathways to 2050. *Food and Agriculture Organization of the United Nations Rome*.

- Farooq, M., Hussain, M., Wahid, A., and Siddique, K. (2012). Drought stress in plants: an overview. *Plant responses to drought stress*, 1-33.
- Farooq, M. U., Bashir, M. F., Khan, M. U. S., Iqbal, B., and Ali, Q. (2021). Role of CRISPR to improve abiotic stress tolerance in crop plants. *Biological and Clinical Sciences Research Journal* **2021**. <https://doi.org/10.54112/bcsrj.v2021i1.69>
- Fatima, A., Saeed, A., Khalid, M. N., Imam, M. M. F., Rafique, M. A., Sharif, M. S., Iqbal, N., Tipu, A. L. K., and Amjad, I. (2022a). Genetic studies of F2 population for fiber and yield related attributes in *Gossypium hirsutum*. *Biological and Clinical Sciences Research Journal* **2022**. <https://doi.org/10.54112/bcsrj.v2022i1.134>
- Fatima, A., Saeed, A., Ullah, M. I., Shah, S. A. H., Ijaz, M., Anwar, M. R., Khaliq, A., Chohan, S. M., Khalid, M. N., Khan, A., and Amjad, I. (2022b). Estimation of gene action for the selection of superior parents and their cross combinations for yield and fiber associated attributes in american cotton (*Gossypium hirsutum* L.). *Biological and Clinical Sciences Research Journal* **2022**. <https://doi.org/10.54112/bcsrj.v2022i1.151>
- Galindo, A., Collado-González, J., Griñán, I., Corell, M., Centeno, A., Martín-Palomo, M., Girón, I., Rodríguez, P., Cruz, Z., and Memmi, H. (2018). Deficit irrigation and emerging fruit crops as a strategy to save water in Mediterranean semiarid agrosystems. *Agricultural water management* **202**, 311-324.
- Huang, J., Ji, M., Xie, Y., Wang, S., He, Y., and Ran, J. (2016). Global semi-arid climate change over last 60 years. *Climate Dynamics* **46**, 1131-1150.
- Hussain, S., Ahmad, A., Wajid, A., Khaliq, T., Hussain, N., Mubeen, M., Farid, H. U., Imran, M., Hammad, H. M., and Awais, M. (2020). Irrigation scheduling for cotton cultivation. In "Cotton production and uses", pp. 59-80. Springer.
- Idrees, H., Shabbir, I., Khurshid, H., Khurshid, A., Tahira, R. I., Fatima, F., Younas, A., and Abbas, H. G. (2022). Seed priming of wheat through salicylic acid to induce salt stress tolerance. *Biological and Clinical Sciences Research Journal* **2022**. <https://doi.org/10.54112/bcsrj.v2022i1.95>
- Iqbal, M., Ul-Allah, S., Naeem, M., Ijaz, M., Sattar, A., and Sher, A. (2017). Response of cotton genotypes to water and heat stress: from field to genes. *Euphytica* **213**, 1-11.
- Iqra, L., Rashid, M. S., Ali, Q., Latif, I., and Malik, A. (2020). Evaluation of genetic variability for salt tolerance in wheat. *Biological and Clinical Sciences Research Journal* **2020**. <https://doi.org/10.54112/bcsrj.v2020i1.16>
- Kalia, R. K., Rai, M. K., Kalia, S., Singh, R., and Dhawan, A. (2011). Microsatellite markers: an overview of the recent progress in plants. *Euphytica* **177**, 309-334.
- Li, F., Li, M., Wang, P., Cox Jr, K. L., Duan, L., Dever, J. K., Shan, L., Li, Z., and He, P. (2017). Regulation of cotton (*Gossypium hirsutum*) drought responses by mitogen-activated protein (MAP) kinase cascade-mediated phosphorylation of Gh WRKY 59. *New Phytologist* **215**, 1462-1475.
- Marengo, J. A., Cunha, A. P., Soares, W. R., Torres, R. R., Alves, L. M., Barros Brito, S. S. d., Cuartas, L. A., Leal, K., Ribeiro Neto, G., and Alvalá, R. (2019). Increase risk of drought in the semiarid lands of Northeast Brazil due to regional warming above 4 C. In "Climate change risks in Brazil", pp. 181-200. Springer.
- Marengo, J. A., Torres, R. R., and Alves, L. M. (2017). Drought in Northeast Brazil—past, present, and future. *Theoretical and Applied Climatology* **129**, 1189-1200.
- Mariano, D. A., dos Santos, C. A., Wardlow, B. D., Anderson, M. C., Schiltmeyer, A. V., Tadesse, T., and Svoboda, M. D. (2018). Use of remote sensing indicators to assess effects of drought and human-induced land degradation on ecosystem health in Northeastern Brazil. *Remote Sensing of Environment* **213**, 129-143.
- Masood, S. A., Khaliq, A., Rauf, H. A., Mahmood, K., Ahmed, I., Hussain, N., Kanwal, S., Faheem, U., and Muhammad, T. (2022). Heat and drought forbearing, upland cotton (*Gossypium hirsutum* L.) variety; RH-668 for cultivation in semi-arid region. *Biological and Clinical Sciences Research Journal* **2022**. <https://doi.org/10.54112/bcsrj.v2022i1.121>
- Meyer, L. A. (2019). "The World and US Cotton Outlook for 2019/20."
- Mittal, A., Gampala, S. S., Ritchie, G. L., Payton, P., Burke, J. J., and Rock, C. D. (2014). Related to ABA-Insensitive3 (ABI 3)/Viviparous1 and At ABI 5 transcription factor coexpression in cotton enhances drought stress adaptation. *Plant biotechnology journal* **12**, 578-589.
- Munir, M. A., Bashir, H., Zaghum, M. J., Aziz, S., Akhtar, S., Ahmad, N. H., Kanwal, S., Kiran, S., Tipu, A. L. K., Liaqat, S., Ahmad, M. I., Latif, A., Latif, A., Nadeem, M., and Shaukat, S. (2022). Evaluation of cotton mutants for water deficit condition. *Biological and Clinical Sciences Research Journal* **2022**. <https://doi.org/10.54112/bcsrj.v2022i1.107>
- MURALI, N., and KHAN, M. (2022). Determinants of Production Performance of Cotton in

- Different Zones of India. *The Mysore of Agricultural Sciences* **56**, 231-235.
- Nadeem, A. H., Nazim, M., Hashim, M., and Javed, M. K. (2014). Factors which affect the sustainable production of cotton in Pakistan: a detailed case study from Bahawalpur district. In "Proceedings of the Seventh International Conference on Management Science and Engineering Management", pp. 745-753. Springer.
- Naseem, S., Ali, Q., and Malik, A. (2020). Evaluation of maize seedling traits under salt stress. *Biological and Clinical Sciences Research Journal* **2020**. <https://doi.org/10.54112/bcsrj.v2020i1.25>
- Niu, J., Zhang, S., Liu, S., Ma, H., Chen, J., Shen, Q., Ge, C., Zhang, X., Pang, C., and Zhao, X. (2018). The compensation effects of physiology and yield in cotton after drought stress. *Journal of Plant Physiology* **224**, 30-48.
- Paterson, A. H., Boman, R. K., Brown, S. M., Chee, P. W., Gannaway, J. R., Gingle, A. R., May, O. L., and Smith, C. W. (2004). Reducing the genetic vulnerability of cotton. *Crop Sci* **44**, 1900-1901.
- Porter, J. R., Xie, L., Challinor, A. J., Cochrane, K., Howden, S. M., Iqbal, M. M., Lobell, D. B., and Travasso, M. I. (2014). Food security and food production systems.
- Pradeep Reddy, M., Sarla, N., and Siddiq, E. (2002). Inter simple sequence repeat (ISSR) polymorphism and its application in plant breeding. *euphytica* **128**, 9-17.
- Rafi, R., Robina, K., Zahoor, M. J., and Abbas, H. G. (2022). Evaluation of maize and sorghum genotypes under drought, drainage and biogas waste water applications. *Biological and Clinical Sciences Research Journal* **2022**. <https://doi.org/10.54112/bcsrj.v2022i1.94>
- Rehman, A., and Farooq, M. (2019). Morphology, Physiology and Ecology of cotton. *Cotton Production*, 23-46.
- Reinisch, A. J., Dong, J.-M., Brubaker, C. L., Stelly, D. M., Wendel, J. F., and Paterson, A. H. (1994). A detailed RFLP map of cotton, *Gossypium hirsutum* x *Gossypium barbadense*: chromosome organization and evolution in a disomic polyploid genome. *Genetics* **138**, 829-847.
- Rong, J., Feltus, F. A., Waghmare, V. N., Pierce, G. J., Chee, P. W., Draye, X., Saranga, Y., Wright, R. J., Wilkins, T. A., and May, O. L. (2007). Meta-analysis of polyploid cotton QTL shows unequal contributions of subgenomes to a complex network of genes and gene clusters implicated in lint fiber development. *Genetics* **176**, 2577-2588.
- Safriel, U., Adeel, Z., Niemeijer, D., Puigdefabregas, J., White, R., Lal, R., Winslow, M., Ziedler, J., Prince, S., and Archer, E. (2005). Dryland systems. In "Ecosystems and Human Well-being: Current State and Trends.: Findings of the Condition and Trends Working Group", pp. 623-662. Island Press.
- Schwinning, S., Sala, O. E., Loik, M. E., and Ehleringer, J. R. (2004). Thresholds, memory, and seasonality: understanding pulse dynamics in arid/semi-arid ecosystems. Vol. 141, pp. 191-193. Springer.
- Shaheen, T., Tabbasam, N., Iqbal, M. A., Ashraf, M., Zafar, Y., and Paterson, A. H. (2012). Cotton genetic resources. A review. *Agronomy for sustainable development* **32**, 419-432.
- Snowden, M. C., Ritchie, G. L., Simao, F. R., and Bordovsky, J. P. (2014). Timing of episodic drought can be critical in cotton. *Agronomy Journal* **106**, 452-458.
- Sohaib, M., and Jamil, F. (2017). An insight of meat industry in Pakistan with special reference to halal meat: a comprehensive review. *Korean journal for food science of animal resources* **37**, 329.
- Tahir, T., Ali, Q., Rashid, M. S., and Malik, A. (2020). The journey of crispr-cas9 from bacterial defense mechanism to a gene editing tool in both animals and plants. *Biological and Clinical Sciences Research Journal* **2020**. <https://doi.org/10.54112/bcsrj.v2020i1.17>
- Tanksley, S., and Nelson, J. (1996). Advanced backcross QTL analysis: a method for the simultaneous discovery and transfer of valuable QTLs from unadapted germplasm into elite breeding lines. *Theoretical and Applied Genetics* **92**, 191-203.
- Tekle, A. T., and Alemu, M. A. (2016). Drought tolerance mechanisms in field crops. *World Journal of Biology and Medical Sciences* **3**, 15-39.
- Tiwari, P., Srivastava, D., Chauhan, A. S., Indoliya, Y., Singh, P. K., Tiwari, S., Fatima, T., Mishra, S. K., Dwivedi, S., and Agarwal, L. (2021). Root system architecture, physiological analysis and dynamic transcriptomics unravel the drought-responsive traits in rice genotypes. *Ecotoxicology and Environmental Safety* **207**, 111252.
- Tomasella, J., Vieira, R. M. S. P., Barbosa, A. A., Rodriguez, D. A., de Oliveira Santana, M., and Sestini, M. F. (2018). Desertification trends in the Northeast of Brazil over the period 2000–2016. *International Journal of Applied Earth Observation and Geoinformation* **73**, 197-206.
- Ton, P. (2011). "Cotton and climate change: impacts and options to mitigate and adapt. ITC, 2011. Technical paper, Doc. No." MAR-11-200. E Technical Report, Geneva, xii.

- Tzortzakis, N., Chrysargyris, A., and Aziz, A. (2020). Adaptive response of a native mediterranean grapevine cultivar upon short-term exposure to drought and heat stress in the context of climate change. *Agronomy* **10**, 249.
- Vicente-Serrano, S., Cabello, D., Tomás-Burguera, M., Martín-Hernández, N., Beguería, S., Azorin-Molina, C., and Kenawy, A. E. (2015). Drought variability and land degradation in semiarid regions: Assessment using remote sensing data and drought indices (1982–2011), *Remote Sens.*, *7*, 4391–4423.
- Wang, R., Ji, S., Zhang, P., Meng, Y., Wang, Y., Chen, B., and Zhou, Z. (2016). Drought effects on cotton yield and fiber quality on different fruiting branches. *Crop Science* **56**, 1265-1276.
- Wang, S., Chen, J., Zhang, W., Hu, Y., Chang, L., Fang, L., Wang, Q., Lv, F., Wu, H., and Si, Z. (2015). Sequence-based ultra-dense genetic and physical maps reveal structural variations of allopolyploid cotton genomes. *Genome biology* **16**, 1-18.
- Wasaya, A., Zhang, X., Fang, Q., and Yan, Z. (2018). Root phenotyping for drought tolerance: a review. *Agronomy* **8**, 241.
- Xu, Y., Zheng, X., Song, Y., Zhu, L., Yu, Z., Gan, L., Zhou, S., Liu, H., Wen, F., and Zhu, C. (2018). NtLTP4, a lipid transfer protein that enhances salt and drought stresses tolerance in *Nicotiana tabacum*. *Scientific reports* **8**, 1-14.
- Yamaguchi-Shinozaki, K., and Shinozaki, K. (2006). Transcriptional regulatory networks in cellular responses and tolerance to dehydration and cold stresses. *Annu. Rev. Plant Biol.* **57**, 781-803.
- Zahoor, M. J., Robina, K., Rafi, R., and Abbas, H. G. (2022). Effects of drought and biogas waste water applications on maize seedling growth. *Biological and Clinical Sciences Research Journal* **2022**.
<https://doi.org/10.54112/bcsrj.v2022i1.93>
- Zhang, F., Wang, P., Zou, Y.-N., Wu, Q.-S., and Kuča, K. (2019). Effects of mycorrhizal fungi on root-hair growth and hormone levels of taproot and lateral roots in trifoliate orange under drought stress. *Archives of Agronomy and Soil Science* **65**, 1316-1330.
- Zhou, S., Sun, X., Yin, S., Kong, X., Zhou, S., Xu, Y., Luo, Y., and Wang, W. (2014). The role of the F-box gene TaFBA1 from wheat (*Triticum aestivum* L.) in drought tolerance. *Plant Physiology and Biochemistry* **84**, 213-223.
- Zonta, J. H., Brandao, Z. N., Rodrigues, J. I. D. S., and Sofiatti, V. (2017). Cotton response to water deficits at different growth stages. *Revista Caatinga* **30**, 980-990.
- Zou, C., Wang, Q., Lu, C., Yang, W., Zhang, Y., Cheng, H., Feng, X., Prosper, M. A., and Song, G. (2016). Transcriptome analysis reveals long noncoding RNAs involved in fiber development in cotton (*Gossypium arboreum*). *Science China Life Sciences* **59**, 164-171.



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