

ISSN 2757-5675

DOI: http://dx.doi.org/10.52520/masjaps.138 Derleme Makalesi

Abiotic Stress Factors & Diseases of Durum Wheat (Triticum turgidum ssp. durum)

Nazlı KALENDER1*, Yusuf DOĞAN1

¹Mardin Artuklu University, Vocational Higher School Of Kızıltepe, Mardin

*Corresponding author: nazlibudakalender@hotmail.com

Geliş Tarihi: 28.03.2021 Abstract Kabul Tarihi: 30.04.2021

When considered on global scale, durum wheat is considered a minor wheat crop but is an important cereal in semiarid parts of Middle East, North Africa, South Europe and North America. Durum wheat is well adapted to high temperatures and semiarid conditions but under Mediterranean conditions, water deficiency and high temperature during reproduction period limit durum production severely. Apart from abiotic factors, Fusarium head blight, stripe rust, leaf rust and stem rust are major diseases resulting with serious yield losses in durum wheat. Abiotic stress factors, especially drought and high temperature, and major diseases of durum wheat effective worldwide, are the subject of this review.

Keywords: Durum wheat, Triticum turgidum ssp. durum, drought, high temperature, diseases

INTRODUCTION

Globally, durum wheat is still considered a minor wheat crop and research efforts on durum is mostly conducted in relation with bread wheat (Beres et al., 2020). Durum is an important cereal in semiarid parts of Middle East, North Africa and South Mediterranean Europe in basin (Hammami & Sissons, 2020). Major consumption areas of durum are pasta, bulgur, couscous and bread production (De Vita & Taranto, 2019). The genetic improvement of durum wheat requires identification of stable **OTLs** (quantitative trait loci) and linked markers. This helps understand the genetic basis of important traits more and identify a method for selection during breeding. Meta-QTL analysis approach is useful to be used in the markerassisted selection for this aim (Soriano et al., 2021). Association mapping allows choose more a accurate to characterization of QTLs in genetic background (Cane et al., 2014). Highdensity genetic linkage maps of durum are particularly useful in detection of quantitative and qualitative QTLs for important agronomical traits and to identify candidate genes (Colasuonno et al., 2014). Selection of single durum wheat plants is a step at early generations during breeding but populations are highly heterogeneous. Instead, leaf weight and spike partitioning of single plants at anthesis have significant positive relation to crop yield (Pedro et al., 2012).

Effects of high temperature and drouht on durum wheat

Genomic applications of durum have potential for exploitation of genetic resources and understanding important complex traits such as tolerance to abiotic and biotic stress factors because durum wheat is generally cropped in medium-low precipitation areas in the world (Maccaferri et al., 2014). Durum adapted wheat is well to high temperatures and semiarid conditions compared to bread wheat but climate change is threatening durum wheat production (De Vita & Taranto, 2019). It is generally assumed that durum has higher tolerance to stress compared to bread wheat (Marti & Slafer, 2014). Durum has great economic importance developing countries in in Mediterranean region but decreases in its production is expected in future due to climate change (Dettori et al., 2011). Mediterranean basin, especially North Africa is highly vulnerable to climate change (Chourghal et al., 2016). Crops get affected from climate change due to inter-relationship of crop development, atmospheric CO2 growth, levels. climatic conditions, reduction in water resources and increase in temperature (Ventrella et al., 2012). Under Mediterranean conditions, water deficiency and high temperature during reproduction period limit cereal crop production severely (Liu et al., 2019). Water deficit is the major limiting factor in Mediterranean basin and durum yield is greatly reduced by drought in this zone. Grain filling rate and duration and distribution of assimilates in stems have important effects under stress conditions on yield performance of durum wheat (El Fakhri et al., 2012).



Fig. 1. New botanical varieties and forms of durum wheat (*T. durum* Desf.): Upper right to left 1) *T. durum* var. *falcaticaerulescens*, 2) *T. durum* var. *falcatiprovinciale*, 3) *T. durum* var. *falcataffine*, 4) *T. durum* var. *falcaterythromelan*. Lower right to left 1) T. durum var. *muticerythromelan*, 2) T. durum var. *muticoleucomelan*, 3) *T. durum* var. *caumelanopus*, 4) *T. durum* var. *Falcativalencia* (Lyapunova, 2017).

Drought and/or heat stress during growth affect the processing quality of durum. In a study of Li et al., (2013), lactic acid retention capacity and mixograph peak time increased under drought and decreased under heat stress. Heat and drought stress sharply reduced grain yield but increased yellowness (Li et al., 2013). 154 durum landraces and 18 modern cultivars from 20 Mediterranean countries were used in field experiments under rainfed conditions during three years in Spain conducted by Nazco et al., Environmental (2012). conditions effected grain protein content, grain yield and grain flour yellowness most. Landraces from the eastern countries Mediterranean recieved

highest mean quality index (protein content, gluten strength, yellow color index and thousand kernel weight) and individual quality trait variability, but grains were relatively small. Landraces of western Mediterranean countries had heavier grains and higher grain filling rates. Modern cultivars, as a group, were the most productive and showed high quality index, but they had the lowest grain protein content and phenotypic variability (Nazco et al., 2012).

Diseases of durum wheat

Fungi from durum wheat grown soil was isolated from soil and plant samples and classified by PCR amplification and sequencing in the study of Vujanovic et al., (2012). Total

46 fungal species were identified. Most of them were from Ascomycota phyla. A were from Zygomycota and few Basidiomycota phyla. Penicillium, Fusarium and Geomyces spp. were abundant throughout growth stages and plant organs. Seventeen species were found potential pathogens of durum where *Fusarium* are the most abundant. Other pathogenic taxa were Pyrenophora, Alternaria, Nigrospora, Microdochium, Bipolaris, Phaeosphaeria, Arthrinium and Cladosporium taxa (Vujanovic et al., 2012). Fusarium head blight (Fusarium graminearum) is a major diseases of durum wheat reducing yield and quality. Mycotoxins also contaminate grains (Gorczyca et al., 2018). Deoxynivalenol is the most important mycotoxin produced by Fusarium moulds (Bensassi et al., 2010). There is an unfavorable correlation with plant height and heading date for durum wheat on the effect of Fusarium head blight. Selection of multiple traits is hard for breeders due to associations of different characters (Moreno-Amores et al., 2020). Breeding for resistance to *Fusarium* in durum is prevented by the lack of resistance resources. Information on resistance

QTL for Fusarium head blight in durum is also limited (Buerstmayr et al., 2012). Improvement for *Fusarium* resistance is hard particularly due to the limited genetic variation in the durum wheat species (Buerstmayr et al., 2013). Fusarium head blight threatens durum production in many growth regions (Zhang et al., 2014). Genomic selection response is higher than phenotypic selection response to obtain Fusarium resistance (Steiner et al., 2019). Stripe rust (Puccinia striiformis f. tritici Eriks.), leaf rust (Puccinia triticina Eriks.), and stem rust (Puccinia graminis f. sp. tritici) are major diseases making serious yield losses in durum wheat (Singh et al., 2013). Identifying new genes is essential for resistance against the diseases (Xu et al., 2013). Stem rust was historically a destructive diseases of durum wheat worldwide (Haile et al., 2012). A stem rust race named TTKSK (Ug99) was identified in 1999 in Uganda. It is still virulent on many resistance genes and is rapidly spreading to other countries (Simons et al., 2011). More than 50 resistance loci for stem rust was identified in wheat but just a few are effective for Ug99 race (Letta et al., 2013).



Fig. 2. Uredinia of leaf rust on flag leaves of wheat (Photo: Mark Hughes, USDA) (Kolmer, 2013).

Resistance breeding to stripe rust is a major objective for durum wheat (Liu et al., 2017). Leaf rust, is an important diseases of common wheat and durum wheat worldwide (Chhetri et al., 2017). Resistance to leaf rust is one of a main target for durum wheat and association mapping on germplasms is used as an approach to discover and validate major genes and QTLs (Maccaferri et al., 2010).

CONCLUSIONS

Durum wheat is well adapted to high temperatures and semiarid conditions compared to bread wheat but climate change is threatening durum wheat production. Especially especially North Africa durum production is highly vulnerable. Apart from abiotic factors, Fusarium head blight, stripe rust, leaf rust and stem rust are major diseases resulting with serious yield losses in durum wheat. Innovative solutions to these diseases are required for durum wheat.

REFERENCES

- Bensassi, F., Zaied, C., Abid, S., Hajlaoui, M.R., Bacha, H. 2010. Occurrence of deoxynivalenol in durum wheat in Tunisia. Food Control, 21(3): 281-285.
- Beres, B.L., Rahmani, E., Clarke, J.M., Grassini, P., Pozniak, C. J., Geddes, C.M., Ransom, J.K. 2020. A Systematic Review of Durum Wheat: Enhancing Production Systems by Exploring Genotype, Environment, and Management (G× E× M) Synergies. Frontiers in Plant Science, 11.
- Buerstmayr, M., Alimari, A., Steiner, B., Buerstmayr, H. 2013. Genetic mapping of QTL for resistance to Fusarium head blight spread (type 2 resistance) in a Triticum dicoccoides× Triticum durum backcross-derived population. Theoretical and applied genetics, 126(11): 2825-2834.
- Buerstmayr, M., Huber, K., Heckmann, J., Steiner, Nelson, B., J.C., Buerstmayr, H. 2012. Mapping of QTL for Fusarium head blight resistance and morphological and developmental traits in three backcross populations derived from dicoccum× Triticum Triticum durum. Theoretical and Applied Genetics, 125(8): 1751-1765.
- Cane, M. A., Maccaferri, M., Nazemi, G., Salvi, S., Francia, R., Colalongo, C., Tuberosa, R. 2014. Association mapping for root architectural traits in durum wheat seedlings as related to agronomic performance. Molecular Breeding, 34(4): 1629-1645.
- Chhetri, M., Bariana, H., Wong, D., Sohail, Y., Hayden, M., Bansal, U. 2017. Development of robust molecular markers for marker-assisted selection of leaf rust resistance gene Lr23 in common and durum wheat

breeding programs. Molecular Breeding, 37(3): 21.

- Chourghal, N., Lhomme, J. P., Huard, F., Aidaoui, A. 2016. Climate change in Algeria and its impact on durum wheat. Regional environmental change, 16(6): 1623-1634.
- Colasuonno, P., Gadaleta, A., Giancaspro, A., Nigro, D., Giove, S., Incerti, O., Blanco, A. 2014. Development of a high-density SNP-based linkage map and detection of yellow pigment content QTLs in durum wheat. Molecular Breeding, 34(4): 1563-1578.
- De Vita, P., Taranto, F. 2019. Durum wheat (Triticum turgidum ssp. durum) breeding to meet the challenge of climate change. In Advances in plant breeding strategies: cereals (pp. 471-524). Springer, Cham.
- Dettori, M., Cesaraccio, C., Motroni, A., Spano, D., Duce, P. 2011. Using CERES-Wheat to simulate durum wheat production and phenology in Southern Sardinia, Italy. Field crops research, 120(1): 179-188.
- El Fakhri, M., Mahboub, S., Benchekroun, M., Nsarellah, N. 2012. Grain filling and stem accumulation effects on durum wheat (Triticum Durum Desf.) yield under drought. Nature & Technology, (7), 67.
- Gorczyca, A., Oleksy, A., Gala-Czekaj, D., Urbaniak, M., Laskowska, M., Waśkiewicz, A., Stępień, Ł. 2018.
 Fusarium head blight incidence and mycotoxin accumulation in three durum wheat cultivars in relation to sowing date and density. The Science of Nature, 105(1): 1-11.
- Haile, J. K., Nachit, M. M., Hammer, K., Badebo, A., Röder, M.S. 2012. QTL mapping of resistance to race Ug99 of Puccinia graminis f. sp. tritici in durum wheat (Triticum durum Desf.). Molecular Breeding, 30(3): 1479-1493.
- Hammami, R., Sissons, M. 2020. Durum wheat products, couscous. Wheat Quality For Improving Processing And Human Health, 347.

- Kolmer, J. 2013. Leaf rust of wheat: pathogen biology, variation and host resistance. Forests, 4(1): 70-84.
- Letta, T., Maccaferri, M., Badebo, A., Ammar, K., Ricci, A., Crossa, J., Tuberosa, R. 2013. Searching for novel sources of field resistance to Ug99 and Ethiopian stem rust races in durum wheat via association mapping. Theoretical and Applied Genetics, 126(5): 1237-1256.
- Li, Y. F., Wu, Y., Hernandez-Espinosa, N., Peña, R. J. 2013. Heat and drought stress on durum wheat: Responses of genotypes, yield, and quality parameters. Journal of Cereal Science, 57(3): 398-404.
- Liu, H., Able, A.J., Able, J.A. 2019. Genotypic performance of Australian durum under single and combined water-deficit and heat stress during reproduction. Scientific reports, 9(1): 1-17.
- Liu, W., Maccaferri, M., Bulli, P., Rynearson, S., Tuberosa, R., Chen, X., Pumphrey, M. 2017. Genomewide association mapping for seedling and field resistance to Puccinia striiformis f. sp. tritici in elite durum wheat. Theoretical and applied genetics, 130(4): 649-667.
- Lyapunova, O.A. 2017. Intraspecific classification of durum wheat: New botanical varieties and forms. Russian Journal of Genetics: Applied Research, 7(7): 757-762.
- Maccaferri, M., Cane, M.A., Sanguineti, M. C., Salvi, S., Colalongo, M.C., Massi, A., Tuberosa, R. 2014. A consensus framework map of durum wheat (Triticum durum Desf.) suitable for linkage disequilibrium analysis and genome-wide association mapping. BMC genomics, 15(1): 1-21.
- Maccaferri, M., Sanguineti, M.C., Mantovani, P., Demontis, A., Massi, A., Ammar, K., Tuberosa, R. 2010. Association mapping of leaf rust response in durum wheat. Molecular Breeding, 26(2): 189-228.
- Marti, J., Slafer, G.A. 2014. Bread and durum wheat yields under a wide

range of environmental conditions. Field Crops Research, 156: 258-271.

- Moreno-Amores, J., Michel, S., Miedaner, T., Longin, C. F. H., Buerstmayr, H. 2020. Genomic predictions for Fusarium head blight resistance in a diverse durum wheat panel: An effective incorporation of plant height and heading date as covariates. Euphytica, 216(2): 1-19.
- Nazco, R., Villegas, D., Ammar, K., Pena, R. J., Moragues, M., Royo, C. 2012. Can Mediterranean durum wheat landraces contribute to improved grain quality attributes in modern cultivars?. Euphytica, 185(1): 1-17.
- Pedro, A., Savin, R., Slafer, G.A. 2012. Crop productivity as related to singleplant traits at key phenological stages in durum wheat. Field Crops Research, 138: 42-51.
- Simons, K., Abate, Z., Chao, S., Zhang, W., Rouse, M., Jin, Y., Dubcovsky, J. 2011. Genetic mapping of stem rust resistance gene Sr13 in tetraploid wheat (Triticum turgidum ssp. durum L.). Theoretical and Applied Genetics, 122(3): 649-658.
- Singh, A., Pandey, M. P., Singh, A. K., Knox, R. E., Ammar, K., Clarke, J. M., Fetch, T.G. 2013. Identification and mapping of leaf, stem and stripe rust resistance quantitative trait loci and their interactions in durum wheat. Molecular breeding, 31(2): 405-418.
- Soriano, J. M., Colasuonno, P., Marcotuli, I., Gadaleta, A. 2021. Meta-QTL analysis and identification of candidate genes for quality, abiotic and biotic stress in durum wheat. Scientific reports, 11(1): 1-15.
- Steiner, B., Michel, S., Maccaferri, M., Lemmens, M., Tuberosa, R., Buerstmayr, H. 2019. Exploring and exploiting the genetic variation of Fusarium head blight resistance for genomic-assisted breeding in the elite durum wheat gene pool. Theoretical and Applied Genetics, 132(4): 969-988.

- Ventrella, D., Charfeddine, M., Moriondo, M., Rinaldi, M., Bindi, M. 2012.
 Agronomic adaptation strategies under climate change for winter durum wheat and tomato in southern Italy: irrigation and nitrogen fertilization. Regional Environmental Change, 12(3): 407-419.
- Vujanovic, V., Mavragani, D., Hamel, C. 2012. Fungal communities associated with durum wheat production system: a characterization by growth stage, plant organ and preceding crop. Crop Protection, 37: 26-34.
- Xu, L. S., Wang, M. N., Cheng, P., Kang, Z. S., Hulbert, S. H., Chen, X. M. 2013.

Molecular mapping of Yr53, a new gene for stripe rust resistance in durum wheat accession PI 480148 and its transfer to common wheat. Theoretical and Applied Genetics, 126(2): 523-533.

Zhang, Q., Axtman, J. E., Faris, J. D., Chao, S., Zhang, Z., Friesen, T. L., Xu, S.
S. 2014. Identification and molecular mapping of quantitative trait loci for Fusarium head blight resistance in emmer and durum wheat using a single nucleotide polymorphism-based linkage map. Molecular breeding, 34(4): 1677-1687.