

Full Length Research Paper

# Indoor characterization of three durum wheat genotypes exposed to drought and heat stress during early vegetative growth stages

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Selection of wheat varieties that have improved adaptation to abiotic stress is important for increasing and stabilizing yields under fluctuating environmental conditions, especially as global climate changes. A trial to estimate adaptation of wheat (*Triticum turgidum* subsp. *durum*) genotypes to abiotic stress has been performed, in a growth chamber. By counting the number of dead (yellow) plants, together with yellow and green leaves, and hence traits that easily can be also detected by automatized phenotyping platforms, were analyzed for the effects of optimal watering, progressive water deficit and different levels of heat stress. "Trinakria" variety and two Trinakria mutants ("Water-mutant" and "Hg-mutant") altered for water-related physiological traits were examined. The use of very genetically close genotypes had the aim to minimize differences in stress response due to asynchronous phenological development and to evaluate better the protocol usefulness to detect minimal phenotypic differences, such as those found between advanced breeding lines, at the final stages of a breeding program. Results showed that Trinakria had a significantly greater % of green leaves under drought stress and retained green leaf after heat stress ceased. In contrast, the two mutants had improved plant survival after moderate heat stress. In conclusion, an examination of leaf color changes under moderate water deficit and heat stress was sufficient in a differential comparison of genotypic performances.

Key words: Abiotic stress, leaf color, phenotyping, wheat.

## INTRODUCTION

Despite world-wide efforts to select high yielding varieties, a decline in wheat production has been observed from the beginning of this millennium; mainly due to a lack of varieties that resist abiotic stress (Dalal et al., 2017). Changes in weather and climate, probably

related to global warming, are shown by an increasing incidence of extreme weather phenomena, even during phenological phases in which the problem of dehydration stress was rare. At the vegetative phase, dehydration stress can modify the growth and development of plants

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Author(s) agree that this article remain permanently open access under the terms of the <u>Creative Commons Attribution</u> <u>License 4.0 International License</u> (Wittmer et al., 1982; Figueroa-Bustos et al., 2019), so affecting up to 56% of the final yield (Gallagher et al., 1976). According to the concepts advanced by Negin and Moshelion (2017), plants may differ for abiotic stress tolerance, resilience or resistance. Tolerance is the ability of the plant to continue photosynthesis, under stress conditions. Resilience denotes the capability to recover and continue growth when moisture is present after drought. Resistance is the plant capability to withstand extreme stress that generally occur at the end of the growth cycle (terminal stresses), and to complete the growth cycle even if most of the leaves (green biomass) has been lost.

In field conditions, the great variability for heat and water stress types occurring on, together with strong genotype x environment interactions and dependence of phenotype on multiple quantitative traits, make complex the selection for improved agronomic performance (Dhanda et al., 2004). For this reason, since the 1970s (Pomeroy and Fowler, 1973), pre-breeding phenotyping under controlled environmental conditions has been commonly employed for functional characterization of varieties, progenies of crosses, mutants, etc. Nowadays, high-throughput non-destructive phenotyping technologies have greatly increased the number of experimental analyses of the wilting process (Humplik et al., 2015; Watt et al., 2020). Controlled environments provide greater reproducibility of experimental conditions and allow multiple stresses to be tested. However, for both non-automated and automated systems, either used in field or indoor, the developmental stage of the plants, stress history, spatial and temporal randomization of plants and micro-environmental fluctuations affect the phenotype which is scored (Yeh et al., 2012). Using separate pots to impose stress on plant, with different morphological-physiological traits, results in application of stresses which are not comparable in timing relative to development stages, and different intensity of the stress. Thus, those plants with greater leaf area, with thinner laminae and/or increased stomatal conductance, and well developed roots, will suffer onset of a water deficit more rapid and greater stress intensity, due to a greater velocity of water loss (Lawlor, 2012). Finally, the trait type to be measured by pre-field screening should be evaluated based on the required performance of plants in field. As an example, a tolerant plant that does not change its physiological activity under early drought and hence has no heavy green leaf loss, will have only a small reduction in yield. Analogously, resilient plants that show the ability to recover their functional activity soon after the stress has ended, are suitable for cultivation in environments where stress is short and intermittent. Highly resistant plants that survive and produce seeds also if with heavy loss of leaves, could have stable yield in cultivation environments where stress generally occurs at the end of the growing cycle (Negin and Moshelion, 2017).

By counting the number of dead (yellow) plants, together with yellow and green leaves, and hence traits that easily can also be detected by automatized phenotyping platforms, a physiological characterization of wheat, during progressive dehydration or after heat stress treatment were analyzed. Water deficit was applied by stopping irrigation, while in a separate experiment, four levels of heat stress were applied by increasing temperature up to 46°C. To minimize differences for stress response due to non-synchronous phenological development and to evaluate the capability of our experimental conditions to detect minimal phenotypic differences (like those that can be found between advanced breeding lines, at the final stages of a breeding program), 3 very genetically closed genotypes were used. They were "Trinakria" variety and 2 mutant lines of Trinakria. The first, called "Water-mutant", has a high affinity for water fraction that is bound to the macromolecule surfaces (Rascio et al., 1999) and the second, named "Hg-mutant" is partially insensitive to HgCl<sub>2</sub>, an aquaporin inhibitor. Both traits of the 2 mutants may have protective roles against dehydration stresses. Bound water is essential for structural integrity of biomolecules (Vertucci and Leopold, 1987). Also, it may exert a passive control of osmotically active volume of the cell (Rascio et al., 2005). Aquaporins are membrane intrinsic proteins that facilitate water transport; their upregulation or down-regulation under stress conditions is thought to be important for tolerance to drought stress (Sade et al., 2009).

## MATERIALS AND METHODS

Thirty plastic pots (Figure 3) were each filled with 4 L of a mixture of soil and sand (50:50 v/v) with a maximum water-holding capacity of 0.32 g H<sub>2</sub>O/g dry weight. The soil mixed to the sand was a clayloam soil (Typic Chromoxerert), with the following physical and chemical characteristics: 36.9% clay, 50.5% silt, 12.5% sand, 15 mg/kg organic matter and pH 8. The pots were put in a 5 × 3.5 m<sup>2</sup> growth chamber, at 20°C/16°C, for a 10 h/14 h light/dark period. Plants were grown under 250-W high-pressure sodium lamps (Philips) and 400-W high pressure metal halide lamps (Philips). Radiation at the pot surface was 1000 µmol photons m<sup>-2</sup> s<sup>-1</sup> (400-700 nm) PAR. Fertilizer was applied before sowing: 18 g m<sup>-2</sup> ammonium sulfate and mineral superphosphate. Cultivation of one plant per pot may create differences among them for stress history and dehydration velocity of soil, due to different growth of plants. For this reason, all pots were subdivided in three parts. Each of them represented one replicate and contained 24 seeds: 8 of "Trinakria", 8 of "Water-mutant" and 8 of "Hg-mutant". Seeds are part of the genotypic working collections, stored at CREA-CI of Foggia (Italy). Distribution of genotypes into each pot section and of pots within the growth chamber was random. After the emergence, to avoid damages to the roots, the plants were not thinned to the same number of plants per genotype, so the final number of plants examined for each per genotype and treatment (Table 1) was different. Before drought and after heat stress, the pots were always kept well-watered, with water loss restored every 2 days to about 80% of maximum soil capacity. When most of the plants had four fully-expanded leaves and hence were at the phase 13-14 of the Zadoks' scale (Zadoks et al., 1974) 25 uniform pots, with 3-5

Table 1	. Tempera	ture cycles	during dr	ought stress,	maximur	n temperat	ure and o	duration o	of the four	heat stres	s treatments	and, nun	nber
of pots	(replicates)	and total i	number of	f plants used	ber each	genotype i	n each tr	eatment.					

Tracimont		Temperature	Time	Number of	No. of plants				
Treatmen	it i	(°C)	(min)	pots	Trinakria	Water-mutant	Hg-mutant		
Control		20		4	13	21	20		
Drought stress		20/16 (day/night)		4	19	24	26		
	Weak	44	60	5	24	24	26		
Heat	Moderate	44	165	4	17	20	20		
stress	Strong	46	60	4	15	16	20		
	Very strong	46	180	4	17	17	20		



Figure 1. % changes for % of dead plants (for plant number before stress) of well watered (control) and droughted plants at increasing days from cessation of watering.

well growth plants per genotype and without signs of disease, were selected. Four pots were kept well watered at  $20^{\circ}C/16^{\circ}C$ , for a 10 h/14 h light/dark period till the end of the experiment and used as controls.

#### Stress treatments

To perform heat stress experiments, a total of 21 well-watered pots were used. Four stress types, differing for temperature (44-46°C) and heat stress duration for a minimum of 60 to a maximum of 180 min were applied and for each of them 4 or 5 pots at a time, were used (Table 1). They were transferred in a thermostatic cabinet with radiation measured at pot surface equal to 60  $\mu$ mol m<sup>-2</sup> s<sup>-1</sup> (400-700 PAR). Seven days after exposure to heat stress treatment, the number of died plants, green and yellow leaves per plant of each genotype in each pot (replicate) were counted.

For drought stress exposure, irrigation was interrupted on four pots. A total of 42 plants per genotype were used to dehydrate in the growth at 20°C/16°C, for a 10 h/14 h light/dark period. Twenty days after, when some symptoms of wilting were visible, the number of dead plants (totally yellow), the number of yellow and green leaves of surviving plants of each genotype in each pot (replicate), were counted. Leaves were classified as yellow if less than 60% of lamina was green. Then, the same measurements were repeated every 2-3 days, for 20 days.

#### Statistics

All results were analyzed using GraphPad Prism software, version 3.0. Differences among genotypes for percentage of died plants and green leaves were processed by one way analysis of variance (ANOVA), with variable number of replicates (pots) shown in Table 1. Means were compared by the multiple comparison test. Tukey regression analysis was performed to define any associations between the variables.

#### RESULTS

### Water deficit stress

The % of dead plants per genotype, concerning the total plants counted before exposure to the drought stress imposition, as shown in Figure 1. Controls had the same percentage of living plants all over the experiment. On average, 50% of plants died about 22 days after water withdrawal (DAWW), while 75% of plants died between 24 and 34 DAWW (Water-mutant and Hg-mutant, respectively). However, there were no significant

Table 2. ANOVA analysis of the number of days required to kill 50% or 75% of plants, after the watering stop.

Verieble	_	LD50%		LD75%				
	Trinakria	Water-mutant	Hg-mutant	Trinakria	Water-mutant	Hg-mutant		
Mean	29.84	31.79	34.36	35.11	36.48	39.8		
Std. Error	3.605	2.738	4.302	2.89	2.026	4.508		
Coefficient of variation	24.17%	17.23%	25.04%	16.46%	11.11%	22.65%		
F values (between genotypes)		0.3955 <sup>ns</sup>		0.5323 <sup>ns</sup>				
Р		0.6845		0.6046				

Means of values extrapolated from linear regression of the percentage of dead plants vs day, of each replicate and genotype replicate, from 27 to 32 days from cessation of watering.



**Figure 2.** Genotypic comparisons for water stress tolerance estimated as changes with time % of green leaves (for total leaves of plants) of droughted plants and well watered plants (control). Means  $\pm$  SE (n = 4).

differences between genotypes determined at increasing time intervals after water was withdrawn, on the basis of ANOVA (data not shown). Standard Error of the mean values of droughted plants increased with days of water withdrawal, because pot to pot differences in number of survived plants of each genotype were greater when drought stress intensity was stronger.

Using data collected from 13 to 32 days after water withdrawal, for each replicate of each genotype, the regression line was constructed and the number of days to have 50 and 75% of dead plants was interpolated. The estimated number of days required to kill 50% or 75% of plants did not differ significantly between genotypes (Table 2), based on ANOVA. At 22 DAWW (Figure 2) and hence under moderate drought stress, Trinakria cultivar had a significantly greater % of green leaves than the Hgmutant (F=4.32; P=0.048), but with increased water deficit it showed a similar, decreasing trend to other genotypes and hence no significant differences were observed.

#### **Heat stress**

Figure 3A shows the plant's appearance before the heat treatment (44°C for 2 h, 45'), immediately after (Figure 3B) and 7 days after (Figure 3C). The number of dead plants was counted seven days after exposure to the stress. Based on Tukey's test (Figure 4), Trinakria mortality increased significantly compared to controls with moderate heat stress. For the other 2 genotypes, % mortality was significantly higher than controls after plant exposure to strong and very strong heat stress. None of the genotypes resisted to intense heat, because all plants died after strong or very strong heat stress (Figure 4). The % of green leaves, 7 days after stress relief is shown in Figure 5. Based on Tukeys' test, Trinakria had a significant reduction of green leaf number under very strong heat stress intensity, as compared to controls. In contrast, after exposure to weak and moderate heat stress, the % of green leaves of both Water-mutant and Hg-mutant was lower than control.



**Figure 3.** Representative pots showing the three genotypes before the exposure exureto heat stress treatments (A), immediately after plant exposure to 44 °C for 2h, 45' (B) and 7 days after (C).



**Figure 4.** % of dead plants days after exposing or not (control) 3 wheat genotypes to different heat stress treatments. Means (n=4-5) and SE are shown. Within genotypes, bars sharing different letters are significantly different (P<0,05) according to Tukey's HSD test.

### DISCUSSION

In temperate climates, water deficit or high-temperature stresses that occur at vegetative stages are often intermittent and of low intensity, but they greatly affect crop yield. To save yield, late drought stress needs resistant plants that survive and produce seeds even if they completely lose the leaves. Tolerant and/or resilient varieties that under abiotic stress are photosynthetically active and hence do not lose their green biomass, could be better useful under early stress.

Results showed that in the system employed, water deprivation caused the death of about 50% of plants 22 days after withdrawing water. Days to kill 75% of plants, ranged from 24 days for Water-mutant, to 32 days for Hgmutant, but this trait and days to kill 50% of plants did not differ significantly between genotypes. The magnitude of genetic components of variance is, generally, lower under stress conditions than under control conditions (Dhanda et al., 2004). Moreover, the drastic treatment necessary



**Figure 5.** % of green leaves with respect to the starting number of leaves per plants, calculated at seven days after exposing or not (control) 3 wheat genotypes to different heat stress treatments. Means (n=4-5) and SE are shown. Bars with different letters are significantly different (P<0.05) according to Tukey's HSD test.

to kill plants increased the well-known variability of plant growth existing within controlled-environment chambers (Measures et al., 1973; Massonnet et al., 2010; Porter et al., 2015). Other authors (Sallam et al., 2018), by using higher average day-night temperature, a single genotype per pot and smaller pots had an average time to 50% wheat wilting, about 13 days shorter than that here. They also observed significant genotypic differences, within a ril population, derived from crosses of more genetically distant parents, as compared to wild type and mutant lines probably due to greater genotypic differences within the *ril* population; lower duration of the cultivation phase necessary to kill 50%; inability to impose the same speed of dehydration on separate pots if they contain genotypes with different morpho-physiological characteristics, already at the beginning of exposure to the stress.

In contrast to what observed for drought response, protective mechanisms that allow plant acquisition or loss of thermotolerance exist and they are under both genetic and epigenetic control (Larkindale et al., 2005; Liu et al., 2015). In this work, too genotypic differences were observed for heat stress effects on plant mortality, because Trinakria cv. performance was significantly worse than that of the mutants. Starting from moderate stress, its mortality increased compared to controls, while for the other 2 genotypes % mortality increased significantly only after the exposure to strong or very strong heat stress. Furthermore, some methodological factors could be the basis for the minor differences in the genotypic response to water stress compared to heat stress. Water withdrawal lasted twenty days. In this relatively long period, changes in the micro-environmental conditions occurred within the growth chamber which differentially modified the stress history of each pot and hence plant growth. The consequences were that, under drought conditions, large plant to plant differences were observed within replicates of the same genotype. On the contrary, thermal stress exposure lasted a few hours, after which all the surviving plants could express their recovery potentiality because optimal conditions were ensured to all plants.

Trinakria cv. Appeared to preserve better the photosynthesizing apparatus because, one week after the withdrawn of watering under weak or moderate stress, it had the same % of green leaves. In contrast, the 2 mutants had 50% fewer green leaves. The general lack of differences in genotype performance under strong high temperature stress, suggests that physiological mechanisms that differentiate the genotypes are unable to affect their performance at temperatures higher than 45°C.

### Conclusions

This study was performed to evaluate the genotypic plant

performance under abiotic stress by counting dead (vellow) plant, together with vellow and green leaves per plants of Trinakria cv., Water-mutant and Hg-mutant. Significant differences among genotypes for the examined traits have been observed. This result suggests that the used method was effective in showing differential plant performances plants under abiotic stress conditions, but further experiments in field are necessary to test the agronomic performance of the same genotypes after exposure to early stress. At the same time, to apply this approach in breeding programs by using automatized systems, pots will have to be designed for simultaneous sowing and screening of many genotypes, which will provide equal conditions of competitiveness of root systems and speed of soil dehydration. Because mutations have functional consequences in terms of abiotic stress response, these mutants are potential sources of traits to be used in traditional breeding programs.

## **CONFLICT OF INTERESTS**

The authors have not declared any conflict of interests.

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