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RESEARCH ARTICLE



Variation among wheat (*Triticum easativum* L.) genotypes in response to the drought stress: I – selection approaches

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ABSTRACT

The agronomic and physiological traits, drought tolerance indexes, principal component analysis and Ward's method were applied to assess the differences among 20 wheat genotypes in response to drought. Statistically significant correlation was observed for measured traits. Drought susceptibility index (DSI), stress tolerance index (STI) and stress index (SI) were most useful to identify genotypes differing in their response to drought. Utility of the indexes was confirmed by physiological markers of drought tolerance i.e. membrane injury and leaf water status. Variation of the genotypes in biomass and grain yield during drought stress was also verified by clustering methods. Finally, integration of physiological and statistical methods presented in this work, allows to both, indicate that tolerance to drought in wheat has a common genetic background, and select the most diverse genotypes. Based on the results, we recommend a tool for breeders, useful to select the genotypes resistant and sensitive to drought.

Abbreviations: DM: dry matter; DSI: drought susceptibility index; FWC: field water capacity; GY: grain yield; GMP: geometric mean productivity index; H: plant height; LI: leakage index related to membrane injury; M_{PRO}: mean productivity index; M_{HAR}: harmonic mean index; NoT: number of tillers; NoG, W-1000: number of grains and weight of 1000 grains, respectively; NoL_{MT}, NoL_{AT}, NoL_T: number of leaves on main tiller, adventitious tillers and total leaf number, respectively; PCA: principal component analysis; RTC: relative trait change; RWC, RT, WD: relative water content, relative turgidity and water deficit, respectively SI: stress index; SPAD: leaf greening; STI: stress tolerance index; TI: tolerance index; WCA: Ward's cluster analysis.

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Introduction

It has been shown that the plant responses to environmental stresses depend on the plant species, genotype, and age, level and duration of stress factors, climate and soil conditions. On the other hand, plant susceptibility to the stress is determined by the genetic factors encoding morphological, phenological, physiological, biochemical and molecular traits. The plants have also developed different active mechanisms, which inhibit or alleviate the harmful effects of stresses (Levitt 1980; Boyer 1982; Asharaf 2010; King 2011).

Among various environmental stresses, soil drought is the most common environmental phenomenon affecting a plant growth, development, and a main factor limiting a grain yield. It has been shown that 20–25% of the global wheat growing area is exposed to drought and the grain yield reduction ranges from 10% to 76% depending on a plant growth stage and a stress severity. Differences in a tolerance to drought within particular genotypes and plant species were proved in many studies, including wheat (Winter et al. 1988; Reynolds et al. 1998; Paknejad et al. 2007), maize (Martiniello and Lorenzoni 1985; Lorens et al. 1987; Grzesiak 1990; Grzesiak et al. 2012), rape seed (Richards 1978), oat (Larsson and Górny 1988; Marcińska et al. 2017), coconut (Gomez et al. 2008) and triticale (Royo et al. 2000; Grzesiak et al. 2012, 2017).

The aim of different breeding programs, including wheat, is to apply selection tests to improve quantity, quality and stability of yield under soil drought and furthermore to develop new drought-adapted genotypes. Methods of screening for drought tolerance within a large number of genotypes should be easy, rapid and inexpensive (Hanson and Nelson 1985; Palta 1990; Evans et al. 1991). In field conditions, proper testing can be performed, providing a number of methodological problems is solved in order to control the water content in the soil, i.e. controlled irrigation and/or reduction of a rainfall inflow. The relations between the plant yield obtained under drought and under optimal soil moistening were preferred among the indicators of drought tolerance in a field. Such markers, however, are not accurate enough or they are too simplified to show important relations between the crop developmental processes and soil-water-plant status. In contrast, some suitable quantitative formulas describing the relations in natural conditions were presented in studies by Fischer and Maurer (1978), Hanson and Nelson (1985), Winter et al. (1988), Stanley (1990) and in FAO reports by Doorenbos and Pruitt (1977), Doorenbos and Kassam (1986). It was found that Drought Susceptibility Index (DSI), Stress Tolerance Index (STI), and Geometric Mean Productivity index (GMP) are efficient for identification of genotypes with good performance to both, non-stress

(control) and stress (drought) conditions (Khalili et al. 2004, Golbashy et al. 2010). Furthermore, Moghaddam and Hadi-Zadeh (2002) found that stress tolerance index (STI) was also useful in selection of maize genotypes significantly differing in a response under stress and non-stress conditions. In laboratory conditions, the testing method should be characterized by a significant correlation with drought resistance observed in a field (Sullivan and Ross 1979; Bouslama and Schapauch 1984; Hanson and Nelson 1985; Blum 1988, 1996; Kpoghomou et al. 1990; Chaves et al. 2002; Grzesiak et al. 2012). The important laboratory protocols for screening of drought tolerance in crop plants include e.g. seed germination in osmotic medium (mannitol, PEG), growth or survival of young seedlings subjected to water stress and high temperature stress (Sullivan and Ross 1979; Blum and Ebercon 1981; Martiniello and Lorenzoni 1985; Clarke and McCaig 1982; Blum 2005), control of membrane injury, leaf water status, leaf temperature and leaf gas exchange (Farquhar et al. 1993; Passioura et al. 1993; Dubey 1997). Other traits that may seem promising as screening tests are various parameters of chlorophyll *a* fluorescence (Reynolds et al. 1998; Hura et al. 2007) and association between dark respiration and heat dissipation under drought (Gerik and Eastin 1985, Reynolds et al. 1998). The numerous publications also pointed out, that root system structure (morphology and anatomy) and water status in a continuum of soil-root-leaf can determine plant survival during drought stress (Yamauchi 1993, Masle 2002), but in this case plant responses can be modified by intensity of other physiological processes, such as leaf gas exchange, stress-induced membrane injury and symbiosis between the roots and mycorrhizal fungi (Kono et al. 1987, Masle 2002, Ogawa et al. 2014, Grzesiak et al. 2017).

A large set of data from screening tests requires a correct statistical analysis to formulate the conclusions concerning tolerant and sensitive genotypes. According to Finlay and Wilkinson (1963), regression analysis was a useful tool in identification of three different adaptation mechanisms to drought conditions in barley genotypes grown at various locations in Australia. The first group of genotypes produced a high yield in conditions of optimal water supply, but showed a drastic reduction in yield under drought. The second group of genotypes had a lower yield in the control conditions, however, the yield reduction during drought was not as significant as in the first group. The third group of genotypes had stable yield under both, control and drought conditions, thus plants have not use favorable water supply conditions to increase the yield. Some other selection tools based on mathematical relationship between stress and non-stress conditions have classified plants in accordance to their responses to four groups: (i) genotypes with good performance in stress and non-stress condition, (ii) genotypes with good performance only in non-stress condition or (iii) genotypes with good performance in only stress condition and (iv) genotypes with weak performance in stress and non-stress conditions (Golbashy et al. 2010, Grzesiak et al. 2012).

The objective of this study was to evaluate the correlation between selected physiological traits and various drought tolerance indexes for 20 wheat cultivars grown in drought conditions. Measurements of different physiological plant responses to drought allowed to recognize the mechanisms, which abolish or reduce the harmful effects of water deficit in the soil or plant tissues. The purpose of our study was also to demonstrate that on the basis of physiological traits

it is possible to select wheat genotypes, which exhibit different tolerance to drought stress as well as determine which tests are most useful for selection of drought-resistant genotypes.

Materials and methods

Plant materials

Experiments were carried out on 20 wheat genotypes: 14 commercial cultivars and 6 breeding forms obtained from DANKO Plant Breeders Ltd; and Smolice Plant Breeders Ltd. IHAR (Table 1).

Experimental conditions

The experiment was carried out in a horticulture tent under natural conditions from March 31st to August 16th, 2017. Plants were grown in plastic pots (21 cm high, 13 cm diameter) filled with a mixture of universal potting soil (Holas, Poland) and quartz-sand (Chemoform AG, Germany < 0.1 cm) in proportion 1:2 (v/v). Air-dried soil substrate was sieved through a 0.25 cm mesh. A compaction of soil substrate in pots was 1.3 g cm^{-3} . Field soil water capacity (FWC) for soil mixture was determined according to Kopecky methods. Air-dried soil samples (100 cm^3) were placed inside metal cylinders, with a 1 mm hole at the bottom. Cylinders with the samples had been soaked 30 min in a container filled with water. After 8 h, maximal soil water content in the samples was $0.43 \text{ (g cm}^{-3}\text{)}$ and after next 48 h it decreased to $0.21 \text{ (g cm}^{-3}\text{)}$. Following Hillel and van Bavel (1976), the values measured after 48 h were assumed to be 100% of soil field water capacity (FWC). For each pot six pre-germinated grains were planted in soil at the depth of 3–4 cm.

Meteorological data (air temperature, relative humidity) was measured continuously by sensors located in the experimental tent. The weather data collected by the data logger were used to establish daily maximum and minimum temperature, relative humidity and the index of Growth Degree-Days (GDD). The concept of GDD assumes that plant growth is related directly to the average daily temperature. The degree-days for each day are summed up, or accumulated throughout the growing season. If the mean daily temperature is equal to or less than the base temperature (for wheat, the base temperature is equal to -5°C) the degree-day value was equals 0 (Edey 1977).

$$\text{GDD} = n_1 + n_2 + n_3 + \dots n_X$$

where n is the successive days of vegetation with mean daily temperature $(t_{\min} + t_{\max})/2$.

Drought stress

Soil drought conditions were achieved by limited watering of the pots, i.e. soil water content was set at the level of 30–35% FWC. Drought treatment (D) had been applied during 5-week period from 28th to 63rd day of plant growth. After this period, soil water content was restored to the level of 65–70% FWC. The control content of soil water (C) was maintained continuously, from sowing to harvest, at the level of 65–70% FWC.

Table 1. Dry matter (DM) and grain yield (GY) of 20 wheat genotypes grown under control (C) and drought stress (D) and ANOVA analyze of variance.

No	Genotype	Dry matter (DM-g plant ⁻¹)			Grain yield (GY-g plant ⁻¹)		
		Control (C)	Drought (D)	RTC	Control (C)	Drought (D)	RTC
1	Telimena	1.52 m	0.59 abcd	0.61	0.89 de	0.01 a	0.98
2	Arabella	1.26 hijk	0.58 abcd	0.54	1.07 f	0.35 d	0.67
3	Mandaryna	1.28 ijkl	0.60 abcd	0.53	1.08 f	0.43 de	0.60
4	Sirocco	1.61 n	0.60a bcd	0.63	0.96 def	0.12 bc	0.87
5	Sharki	1.02 bc	0.58a bcd	0.43	0.76 c	0.08 ab	0.59
6	Cornetto	1.06 cd	0.57 abcd	0.46	0.71 bc	0.11 b	0.60
7	Tybalt	1.09 cde	0.52 ab	0.52	0.80 cd	0.37 d	0.60
8	Goplana	1.30 jkl	0.52 ab	0.60	0.92 de	0.19 bc	0.79
9	Harenda	1.17 fg	0.62 bcd	0.47	0.77 c	0.47 e	0.56
10	Kandela	1.20 ghij	0.59 abcd	0.51	0.91 de	0.49 e	0.58
11	Struna	1.21 ghij	0.89 ef	0.26	0.94 de	0.57ef	0.41
12	Jarlanka	1.10 cdef	0.51 ab	0.54	0.86 d	0.39 d	0.55
13	SMJ 2115	0.95 ab	0.71d	0.25	0.87 de	0.59 f	0.32
14	SMJ 3016	1.25 ghij	0.56 ab	0.55	0.51bc	0.23 c	0.55
15	SMJ 1314	1.32 jkl	0.70 cd	0.47	0.78 c	0.39 d	0.50
16	Ostka smolicka	1.19 gh	0.60 abcd	0.50	0.59 bc	0.34 d	0.49
17	Tdoz 1384	0.92 a	0.47 a	0.49	0.32 a	0.16 bc	0.49
18	Tdoz 1403	1.68 n	0.81d	0.52	0.99 def	0.03a	0.97
19	Tdzo 1508	1.21 ghij	1.05 e	0.13	0.80 cd	0.49 e	0.39
20	Tonika	1.68 n	1.02 e	0.39	0.79 cd	0.11b	0.86
Mean		1.25	0.65	0.47	0.82	0.29	0.62
CV (%)		17.7	25.2	26.7	25.2	63.7	34.5
Anova							
Variable	df	Dry matter (DM)		Grain yield (GY)			
		C	D	C	D		
Genotype (G)	19	**	***	**	**		
Treatment (T)	1	*	*	NS	*		
G × T	19	*	*	*	*		

Notes: CV – coefficient of variation; df – number of freedom degree. Different letters in column indicate significant differences according Duncan test at $p < .05$ level. Based on decrease of relative traits changes (RTC) of DM or GY, the genotypes are divided into four groups: Light grey (>0.30), Medium grey ($<0.30 > 0.50$), Dark Grey ($<0.50 > 0.70$) and Normal Grey (>0.70). *, **, *** – statistical significance at level $p < .1$, .05, .01, respectively. NS – no significant differences.

Drought tolerance indexes were calculated on the basis of DM or GY using the following equations (Fischer and Maurer 1978; Golbashy et al. 2010; Grzesiak et al. 2012).

- Drought Susceptibility Index (DSI)

$$DSI = (1 - D/C)/(1 - xD/xC)$$

- Stress Tolerance Index (STI)

$$STI = (C \times D)/(xC)$$

- Mean Productivity Index (M_{PRO})

$$M_{PRO} = (C + D)/2$$

- Harmonic Mean Index (M_{HAR})

$$M_{HAR} = 2(C \times D)/(C + D)$$

- Tolerance Index (TI)

$$TI = C - D$$

- Geometric Mean Productivity Index (GMP)

$$GMP = \sqrt{C \times D}$$

- Stress Index (SI)

$$SI = D/C$$

where C, D – dry matter of aboveground part of the plant (DM) or grain yield (GY) in control (C) and drought (D) treatment, respectively. xC, xD – average values for all examined genotypes of dry mater of aboveground part of the plant (DM) or grain yield (GY) in control (C) and drought (D) treatment, respectively.

Relative trait changes (RTC) was calculated as $(C-D)/C$

Measurements

Physiological traits: Dry matter of above-ground part of the plants was analyzed directly after drought treatment, whereas plant height (H), number of tillers (NoT), number of leaves on main tiller (NoL_{MT}), number of leaves on adventitious tillers (NoL_{AT}), total leaf number (NoL_T), grain yield (GY), number of grains (NoG), and weight of 1000 grains (W-1000) were determinate after plant harvesting. Plants were dried at 65°C for 72 h. For each genotype and treatment data were collected for 9 plants grown in 3 pots.

Leaf greening (SPAD) was measured directly after drought treatment on well-developed leaf using SPAD CL-01 meter (Hansatech, Norfolk, UK).

Leaf water status was determined as a relative water content (RWC), relative turgidity (RT), and water deficit (WD). According to the protocol published by Weatherley (1950) and Barrs (1986) leaf fragments about 5 cm long were collected between 11:00 pm and 1:00 am from most recent fully expanded leaf. Each analysis was performed in 5 biological replicates. Parameters were calculated as it follows:

$$RWC = f_w - d_w/f_w$$

$$RT = f_w - d_w/t_w - d_w$$

$$WD = 100 - RT$$

where f_w – fresh weight, d_w – dry weight and t_w – fully turgid weight measured after 24 h leaf floated in water at 4°C and ambient light.

Relative loss of intracellular electrolytes from leaf tissues (LI-leakage index) was measured with the conductivity meter OK-102/1 (Radelkis, Hungary). Leaf fragments about 7 cm long were taken between 11:00 pm and 1:00 am on most recent

fully expanded leaf and incubated in test tubes filled with 30 cm³ redistilled water. After 24 h, initial conductance was measured. Final conductance was measured after autoclaving tubes with leaves at 110° C for 15 min and cooling them to room temperature. There were 5 replicates for each treatments. LI was calculated according to the formula described by Sullivan and Ross (1979) and Blum and Ebercon (1981).

$$LI = ((D_1/D_2) - (C_1/C_2))/(1 - (C_1/C_2)) \times 100$$

where *C* and *D* refer to the conductivity of control (*C*) and drought (*D*) treatment, respectively, and subscript 1 and 2 refer to initial and final conductance, respectively.

Statistical analysis

To determine statistical significance of obtained data, ANOVA analysis of variance was performed with STATISTICA 12.0 software (Statsoft, Tulsa, OK, USA). Means were compared using Duncan's multiple range test at 0.05 level of probability. Provided *F*-values were significant, linear correlation coefficients together with the probability levels of sensitivity to drought stress indexes were calculated. Principal component analysis (PCA) was applied to assign the ranks to genotypes and to classify which of them were more susceptible or resistant to drought stress. PCA utilizes orthogonal transformation to convert a set of possibly correlated variables into a set of linearly uncorrelated variables called principal components. This transformation is defined in such a way that the first principal component has the largest possible variance. PCA is sensitive to the relative scaling of the original variables. Ward's cluster analysis (WCA) was also applied as a criterion in hierarchical cluster analysis. The nearest-neighbor chain algorithm was used to find the same clustering in time proportional to the size of the input distance matrix and space linear in the number of points being clustered.

Results

Climatic parameters

The changes of daily maximum and minimum air temperature, relative humidity (RH) and the values of growth degree days (GDD) are shown in Figure 1. During the experiment period, the average monthly air temperature values were 8.2°C, 14.0°C, 16.8°C, 18.8°C and 18.1°C, respectively, from April to August 2017. Changes in average monthly air temperature in the period from April to May were similar, and from June to August were higher, in comparison to the long-term norm for the Krakow area (www.weatheronline.pl). The value of the GDD index in the 2017 growing season reached the value of around 2400 and was close to the optimal value for early and moderate – early spring wheat cultivars (McMaster et al. 2003).

Morphological and physiological traits

The measured traits for control (*C*) and drought (*D*) treatment as well as relative trait change (RTC) are shown in Table 1 and Table 2. In control conditions, the highest values of coefficient of variation (*CV* > 20%) were observed for dry matter (DM), grain yield (GY), number of tillers (NoT), and number of grains (NoG). Furthermore, values of *CV* after *D* for traits DM, NoT, NoL_{AT}, SPAD, GY, NoG and W-1000 were higher than for *C* (Tables 1 and 2). Results presented

as the RTC show statistically significant correlation for DM, GY and NoT, while for the other traits the correlation coefficients were statistically insignificant (Table 3). Based on both, RTC for DM and RTC for GY (Table 1), distribution of the genotypes have a higher resolution than for only one parameter, due to a range of RTC values for DM was lower (0.13–0.63), while range of RTC values for GY was higher (0.32–0.98). Therefore, the genotypes could be divided into four groups. In the first group, RTC was lower than 0.30 for the genotypes: Tdoz 1508, SMJ 2115, Struna. In the second group, RTC ranged from 0.30–0.50 for genotypes: SMJ 2115, Tdoz 1508, Tonika, Struna, Sharki, Cornetto, Harenda, Ostka smolicka, Tdoz 1384. In the third group, RTC ranged from 0.50 to 0.70 for genotypes: SMJ 1314, Tdoz 1384, Ostka smolicka, Kandela, Tybalt, Tdoz 1403, Mandaryna, Arabella, Jarlanka, SMJ 3016, Harenda, Sharki, Cornetto, Goplana, Telimena, Sirocco. In the fourth group RTC was higher than 0.70 for genotypes Goplana, Tonika, Sirocco, Tdoz 1403, Telimena. Moreover, an interesting interaction between yield of the plants grown under *C* and *D* conditions was observed. Telimena, Sirocco, Goplana, Tdoz 1403, and Tonika represent genotypes accumulating the highest dry mass under *C* conditions, but at the same time, the greatest reduction of dry mass after *D* was observed. In contrast, SMJ 2115 had significantly lower mean dry mass under *C*, but at the same time the effect of the drought was minor.

Drought tolerance indexes

On the basis of DM and GY, different drought tolerance indexes were calculated to identify the drought resistant and sensitive genotypes (Table 4). Relatively high variability of the coefficient of variation was observed for presented indexes. ANOVA analysis showed also the statistically significant differences between genotypes (*G*), whereas a significant variability for the treatments (*T*) and the interactions *G* × *T* was not found in the case of all indexes. In Table 5, correlation coefficients between different drought tolerance indexes are presented. In most cases, the correlation coefficients proved that indexes provided the suitable criteria for the selection of genotypes. The correlation coefficient between drought tolerance indexes calculated on the basis of DM and GY was statistically significant (*r* = 0.460) and indicate a similar pattern of variation between indexes calculated for DM (directly after drought) and those calculated for GY (after plant harvesting). Based on DM under the control treatment (Table 1), the genotypes were divided into two groups. The group A with high yielding (*A* > from the genotypic mean value) included Tdoz 1403, Tonika, Sirocco, Telimena, SMJ 1314, Goplana, Mandaryna, Arabella, and SMJ 3016; while the group B with low yielding (*B* < from the mean genotypic value) included Struna, Tdoz 1508, Kandela, Ostka smolicka, Harenda, Jarlanka, Tybalt, Cornetto, Sharki, SMJ 2115 and Tdoz 1384. Within the group A, drought caused the highest decrease in DM for Sirocco, Telimena and Goplana (40%), and for Tdoz 1403, Arabella, Mandaryna, SMJ 3016 (up to 45%), while the lowest decrease of DM was recorded for Tonika (61%) and SMJ 1304 (53%). In the group B, the highest decrease of DM, up to about 45%, was recorded in Tybalt, Kandela and Jarlanka, and to about 55% in Ostka smolicka, Tdoz 1384, Sharki, Cornetto and Harenda. In this group the lowest decrease of DM was recorded in Tdoz 1508 (87%), SMJ 2115 and Struna (75%).

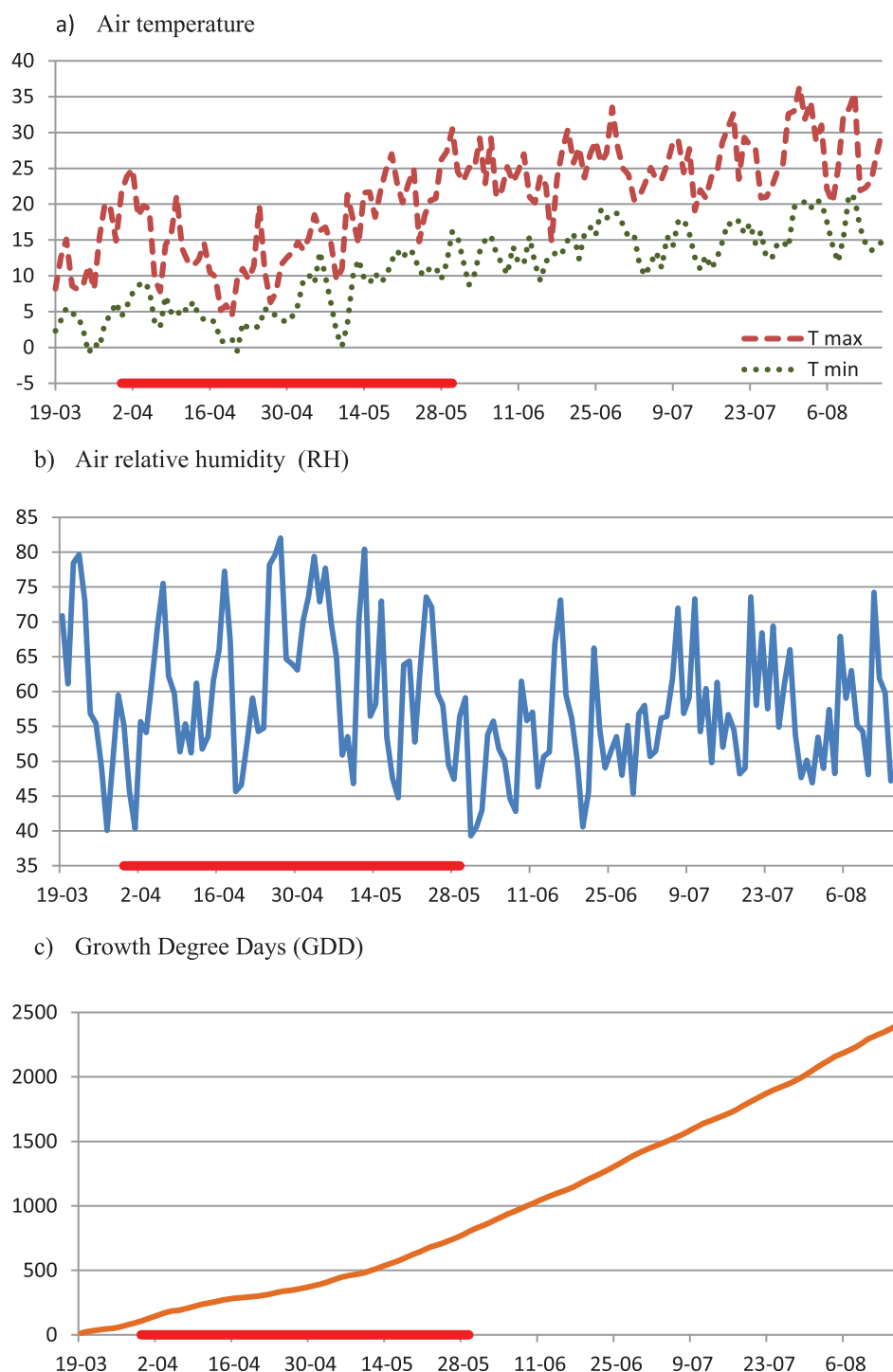


Figure 1. Changes of daily maximum and minimum air temperature (a), air relative humidity RH (b) and growth degree days GDD (c). Red line on x-axis marks the period of drought.

On the basis of the value of drought tolerance indexes (Table 4), it was also possible to ranked genotypes according to their degree of susceptibility to drought within the groups with high (A) or low (B) yielding. DSI, TI and SI indicated Sirocco, Telimena, Arabela as a drought sensitive genotypes in the group A, and Harenda, Jarlanka and Tybalt in the group B; while Tonika and SMJ 1304 as the resistant genotypes in group A, and Tdoz 1508, SMJ 2115, Struna in the group B. The division into resistant and sensitive genotypes based on the STI, M_{PRO} M_{HAR} and GMP indicated Tonika, Todz 1403, Sirocco as drought-resistant genotypes in the group A, and Todz 1508, Struna in the group B, however, only Tdoz 1384, Jarlanka, Tybalt, Sharki and Cornetto from the group B were classified as drought sensitive genotypes.

Principal component analysis (PCA) and Ward's cluster analysis (WCA)

PCA analysis was used to identify resistant and sensitive genotypes among drought-stress treated plants. The orthogonal transformation was defined in such a way that the first principal component has the largest variance. According to physiological traits in the PCA analysis (Figure 2(a,b)), the first factor (PC1) explains 47,4% of the total variance of variables, and the second factor (PC2) about 20,6%. In total, both PCs explain 68,0% of the total variance of all analyzed variables. The biplot analysis showed the correlation between stress index (SI) calculated for all of the examined traits (Figure 2(a)). The sharp angle means a positive

Table 2. Mean values, range of mean, LSD_{0.05}, and coefficient of variation (CV) for the physiological traits of wheat genotypes grown in control (C) and drought (D) conditions.

Traits	Control (C)				Drought (D)			
	Mean	Range	LSD _{0.05} V	CV (%)	Mean	Range	LSD _{0.05} V	CV (%)
Traits measured directly after drought								
DM – Biomass dry matter (g plant ⁻¹)	1.25	0.92–1.68	0.39	26.9	0.65	0.51–1.02	0.13	37.8
H – Seedlings height (cm)	36.6	28.3–51.6	3.89	16.5	23.2	17.0–31.6	1.80	17.6
NoT – Number of tillers	2.3	1.3–3.1	0.49	21.0	1.6	0.9–2.8	0.41	33.6
NoL _{MT} – No. of leaves on main tiller	5.8	4.3–6.4	0.88	8.1	6.1	4.5–7.2	1.11	10.6
NoL _{AT} – No. of leaves on adventitious tillers	7.4	4.3–10.0	2.18	19.1	4.7	2.1–9.6	2.39	42.6
NoL _T – Total leaf number	13.2	10.4–16.1	1.07	10.5	10.8	8.3–15.6	2.13	19.2
SPAD – Leaf greening	19.0	14.5–26.2	3.11	15.8	12.7	7.7–20.3	3.18	26.6
Traits measured after harvesting								
GY – Grain weight (g plant ⁻¹)	0.81	0.32–1.08	0.08	25.2	0.30	0.01–0.11	0.07	55.4
NoG – Number of grain	27.0	7.8–38.7	3.07	29.4	13.1	1.1–20.5	4.18	48.4
W-1000 – Weight of 1000 grain (g plant ⁻¹)	31.0	23.5–43.3	5.55	17.0	22.9	9.6–37.0	5.05	33.0

Table 3. Relative change of measured traits (RTC), mean values, LSD_{0.05}, coefficient of variation (CV) and correlation coefficient for traits of 20 wheat genotypes.

No.	Genotype	DM	GY	H	NoT	Leaf number			SPAD	Grain	
						NoL _{MT}	NoL _{AT}	NoL _T		NoG	W-1000
1	Telimena	0.61	0.98	0.50	0.57	0.07	0.57	0.36	0.49	0.96	0.60
2	Arabella	0.54	0.67	0.33	0.43	–0.12	0.53	0.31	0.18	0.65	0.08
3	Mandaryna	0.53	0.60	0.56	0.12	0.30	0.88	0.25	0.20	0.60	0.02
4	Sirocco	0.63	0.87	0.50	0.50	–0.12	0.46	0.22	0.33	0.86	0.08
5	Sharki	0.43	0.59	0.29	0.47	–0.24	0.88	0.19	0.46	0.57	0.06
6	Cornetto	0.46	0.60	0.37	0.67	–0.07	0.70	0.35	0.06	0.53	0.16
7	Tybalt	0.52	0.60	0.36	0.54	–0.04	0.59	0.34	0.29	0.45	0.27
8	Goplana	0.60	0.79	0.49	0.46	–0.10	0.54	0.27	0.37	0.71	0.28
9	Harenda	0.47	0.56	0.35	0.28	0.03	0.38	0.20	0.34	0.24	0.42
10	Kandela	0.51	0.58	0.32	0.41	–0.07	0.43	0.22	0.20	0.29	0.41
11	Struna	0.26	0.41	0.26	0.08	–0.03	–0.12	–0.08	0.04	0.45	–0.08
12	Jarlanka	0.54	0.55	0.31	0.48	–0.26	0.58	0.21	0.34	0.31	0.34
13	SMJ 2115	0.25	0.32	0.12	–0.46	0.02	0.05	0.03	0.30	0.34	–0.02
14	SMJ 3016	0.55	0.55	0.27	–0.13	0.03	0.51	0.30	–0.03	–0.68	0.73
15	SMJ 1314	0.47	0.50	0.35	0.23	–0.28	0.43	0.16	0.49	0.35	0.23
16	Ostka smolicka	0.50	0.49	0.41	0.19	–0.09	0.18	0.06	0.50	–0.08	0.53
17	Tdzo 1384	0.49	0.49	0.37	0.45	–0.06	0.61	0.32	0.51	0.44	0.10
18	Tdzo 1403	0.52	0.97	0.41	0.00	0.00	0.01	0.01	0.65	0.93	0.51
19	Tdzo 1508	0.13	0.39	0.25	–0.18	0.03	–0.30	–0.15	0.17	0.52	–0.27
20	Tonika	0.39	0.86	0.33	0.10	0.05	0.20	0.14	0.51	0.73	0.49
	Mean	0.47	0.62	0.36	0.26	–0.05	0.41	0.18	0.32	0.46	0.25
	L.S.D (0.05)	0.13	0.18	0.11	0.08	0.07	0.25	0.12	0.10	0.33	0.39
	% CV	26.7	30.0	28.5	113.2	–272.8	77.2	78.0	57.2	80.4	103.9
Correlation coefficient											
Trait	GY 2	H 3	NoT 4	NoL _{MT} 5	NoL _{AT} 6	NoL _T 7	SPAD 8	NoG 9	W-1000 10		
1 – DM	0.634***	0.717***	0.638***	–0.118 ^{NS}	0.690***	0.774***	0.231 ^{NS}	0.077 ^{NS}	0.591***		
2 – GY		0.657***	0.401*	0.114 ^{NS}	0.219 ^{NS}	0.358 ^{NS}	0.444**	0.602***	0.480**		
3 – H			0.538**	0.134 ^{NS}	–0.246 ^{NS}	–0.081 ^{NS}	–0.092 ^{NS}	0.026 ^{NS}	–0.033 ^{NS}		
4 – NoT				–0.395*	0.670***	0.702***	0.127 ^{NS}	0.331 ^{NS}	0.134 ^{NS}		
5 – NoL _{MT}					–0.120 ^{NS}	–0.048 ^{NS}	–0.226 ^{NS}	0.054 ^{NS}	0.025 ^{NS}		
6 – NoL _{AT}						0.869***	–0.002 ^{NS}	0.010 ^{NS}	0.173 ^{NS}		
8 – NoL _T							–0.040 ^{NS}	0.001 ^{NS}	0.373 ^{NS}		
9 – SPAD								0.427**	0.283 ^{NS}		
10 – NoG									–0.312 ^{NS}		

Notes: *, **, *** – statistical significance for correlation coefficient at level $p < .50$, .10 and .01, respectively. NS – not significant differences.

correlation between the measured parameters, the obtuse angle – negative, while the right angle means no correlation between the parameters. Positive correlation was recorded among of dry matter (DM) with all traits except NL_{MT} and W-1000 and among grain yield (GY) with all traits except NL_{AT}. Furthermore, PCA analysis allowed to group wheat genotypes according to their phenotypic similarities (Figure 2(b)). Group I (the highest PC1 and PC2) includes: Struna and Tdzo 1508; group II (the highest PC1 and the lowest PC2) Ostka smolicka, SMJ 2115, SMJ 3016 and SMJ 1314; group III (the lowest PC1 and PC2) Arabella, Sharki, Cornetto, Tybalt, Harenda, Kandela, Tdzo 1384, and Jarlanka; and group IV (the lowest PC1 and the highest PC2) consisted genotype Telimena, Sirocco, Goplana, Mandaryna, Tdzo 1403, and Tonika.

The PCA was also carried out for seven different drought tolerance indexes calculated for DM (Figure 3(a,b)). The first factor of PCA analysis explains about 72.6% of the total variance of variables, and the second factor about 26.7%. In total, both main axes explain 99.0% of the total variance of all analyzed variables. Positive correlation was observed among the drought susceptibility index (DSI) only with TI, stress tolerance index (STI) with GMP, MPRO, MHAR and SI, tolerance index (TI) only with MPRO, geometric mean productivity (GMP) with MPRO and MHAR, mean productivity index (MPRO) with MHAR and SI and harmonic mean index (MHAR) with SI (Figure 3(a)). According to PCA analysis wheat genotypes were classified into 4 groups (Figure 3(b)). Group I (the highest PC1 and PC2) consisted genotypes Cornetto, SMJ 3016, Tdzo 1314, group II (the highest PC1 and

Table 4. Drought tolerance indexes calculated for dry biomass of above-ground plant part (DM) and grain yield (GY) of 20 wheat genotypes grown in control (C) and drought (D) conditions and analyze of variance ANOVA.

No	Genotype	Based on biomass dry matter (DM) of above-ground plant part							Based on grain yield (GY)						
		DSI	STI	M _{PRO}	M _{HAR}	TI	GMP	SI	DSI	STI	M _{PRO}	M _{HAR}	TI	GMP	SI
1	Telimena	1.291	0.571	1.054	0.845	0.938	0.944	0.384	1.552	0.019	0.452	0.027	0.876	0.111	0.016
2	Arabella	1.126	0.471	0.922	0.798	0.677	0.858	0.463	1.063	0.569	0.708	0.524	0.719	0.609	0.326
3	Mandaryna	1.116	0.491	0.939	0.816	0.681	0.875	0.467	0.951	0.704	0.751	0.611	0.649	0.678	0.397
4	Sirocco	1.316	0.619	1.105	0.874	1.011	0.983	0.372	1.373	0.184	0.544	0.221	0.838	0.346	0.130
5	Sharki	0.909	0.380	0.801	0.740	0.444	0.770	0.567	0.933	0.362	0.421	0.145	0.450	0.247	0.409
6	Cornetto	0.974	0.386	0.814	0.740	0.493	0.776	0.536	0.952	0.303	0.406	0.185	0.426	0.274	0.397
7	Tybalt	1.101	0.361	0.803	0.701	0.572	0.750	0.475	0.949	0.389	0.583	0.504	0.480	0.542	0.398
8	Goplana	1.262	0.431	0.908	0.740	0.782	0.820	0.398	1.253	0.266	0.555	0.314	0.731	0.417	0.205
9	Harenda	0.981	0.470	0.900	0.816	0.550	0.857	0.532	0.884	0.400	0.622	0.586	0.432	0.604	0.440
10	Kandela	1.064	0.455	0.896	0.792	0.609	0.842	0.493	0.918	0.529	0.698	0.635	0.529	0.666	0.418
11	Struna	0.554	0.692	1.051	1.027	0.320	1.039	0.736	0.650	0.849	0.755	0.710	0.400	0.732	0.588
12	Jarlanka	1.125	0.360	0.805	0.697	0.590	0.749	0.464	0.860	0.512	0.624	0.536	0.468	0.578	0.455
13	SMJ 2115	0.524	0.432	0.829	0.812	0.237	0.820	0.751	0.508	0.781	0.727	0.701	0.279	0.714	0.678
14	SMJ 3016	1.158	0.445	0.901	0.770	0.688	0.833	0.447	0.865	0.076	0.338	0.338	0.182	0.338	0.452
15	SMJ 1314	0.978	0.593	1.010	0.916	0.614	0.962	0.534	0.784	0.468	0.585	0.521	0.387	0.552	0.503
16	Ostka smolicka	1.038	0.457	0.894	0.797	0.588	0.844	0.505	0.781	0.273	0.465	0.430	0.294	0.447	0.505
17	Tdzo 1384	1.033	0.276	0.694	0.620	0.454	0.656	0.507	0.774	0.079	0.240	0.214	0.156	0.227	0.509
18	Tdoz 1403	1.086	0.876	1.248	1.095	0.872	1.169	0.482	1.526	0.048	0.510	0.061	0.956	0.177	0.032
19	Tdoz 1508	0.273	0.812	1.128	1.123	0.157	1.125	0.869	0.616	0.601	0.646	0.608	0.313	0.626	0.610
20	Tonika	0.824	1.098	1.350	1.269	0.660	1.309	0.607	1.362	0.131	0.450	0.190	0.684	0.293	0.136
Mean		0.987	0.530	1.578	0.849	0.597	0.899	0.529	0.978	0.377	0.552	0.408	0.512	0.463	0.380
CV (%)		16.9	19.1	36.1	17.8	17.8	25.2	23.9	30.0	65.8	26.2	49.2	44.8	39.7	49.0
ANOVA															
Genotype (G)		19	***	*	**	*	*	*	***	**	***	*	*	*	*
Treatment (T)		1	**	NS	NS	*	*	*	**	*	NS	NS	*	*	*
G × T		19	**	*	NS	*	NS	NS	*	NS	NS	*	NS	*	NS

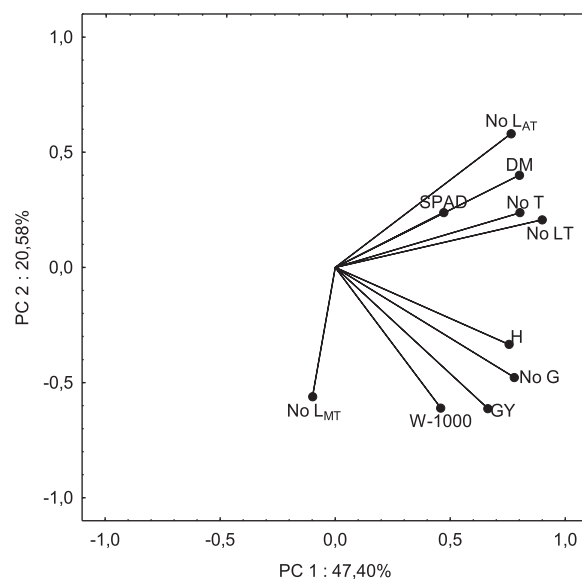
Notes: *, **, *** – statistical significance at level $p < .1$, $.05$, $.01$, respectively. NS – no significant differences.

Table 5. Linear correlation coefficient between different drought tolerance indexes of wheat genotypes (df = 18).

No	Indexes	STI	M_{PRO}	M_{HAR}	TI	GMP	SI
Based on dry matter (DM)							
1	DSI	-0.342 ^{NS}	0.025 ^{NS}	-0.481**	0.874***	-0.335 ^{NS}	-0.999***
2	STI		0.917***	0.985***	0.118***	0.995***	0.341 ^{NS}
3	M_{PRO}			0.854***	0.491**	0.929***	-0.026 ^{NS}
4	M_{HAR}				-0.030 ^{NS}	0.986***	0.480**
5	TI					0.136 ^{NS}	-0.875***
6	GMP						0.335 ^{NS}
Based on grain yield (GY)							
1	DSI	-0.673***	-0.250 ^{NS}	-0.794***	0.886***	-0.688***	-0.999***
2	STI		0.872***	0.969***	-0.272 ^{NS}	0.980***	0.673***
3	M_{PRO}			0.766***	0.218 ^{NS}	0.848***	0.250 ^{NS}
4	M_{HAR}				-0.446**	0.986***	0.794***
5	TI					-0.301 ^{NS}	-0.886***
6	GMP						0.688***

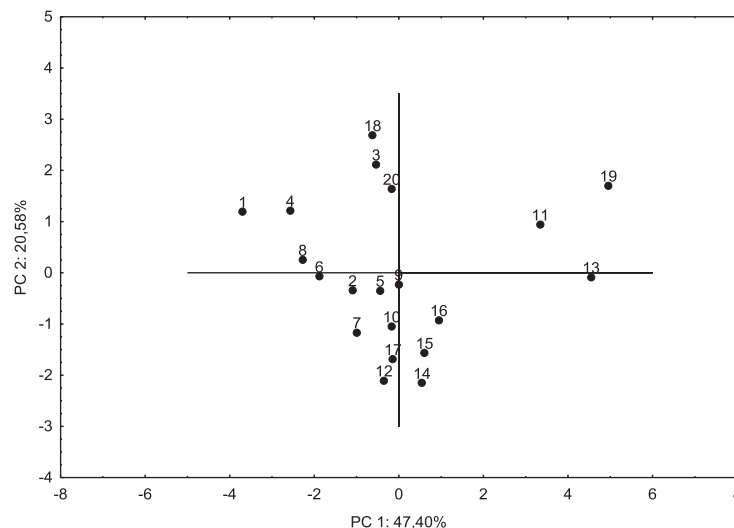
Notes: *, **, *** – statistical significance for correlation coefficient at level $p < .50$, .10 and .01, respectively. NS – not significant differences.

a. PCA for physiological traits.



DM – Dry matter
GY – Grain weight
NoG – Number of grain
W-1000 – Weight of 1000
H – Seedlings height
NoT – Number of tillers
NoLMT – Number of leaves on main tiller
NoLAT – Number of leaves on adventitious tillers
NoLT – Total leaf number
SPAD – Leaf greening

b. PCA for listed wheat genotypes



No	Genotype
1	Telimena
2	Arabella
3	Mandaryna
4	Sirocco
5	Sharki
6	Cornetto
7	Tybalt
8	Goplana
9	Harenda
10	Kandela
11	Struna
12	Jarlanka
13	SMJ 2115
14	SMJ 3016
15	SMJ 1314
16	Ostka smolicka
17	Tdoz 1384
18	Tdoz 1403
19	Tdoz 1508
20	Tonika

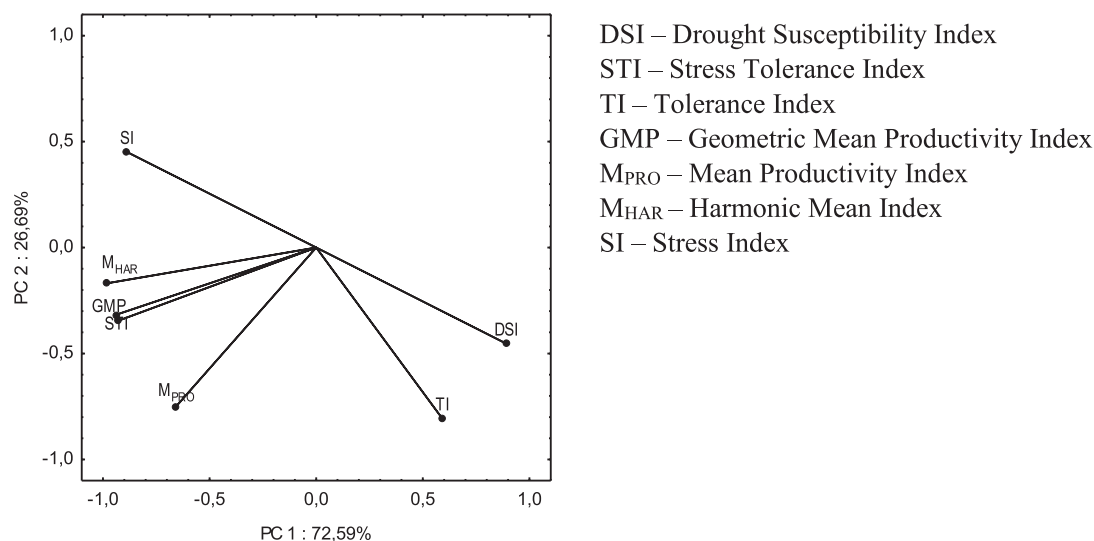
Figure 2. Principal component analysis (PCA) for physiological traits (a) and genotypes (b) based on stress indexes calculated for physiological traits.

the lowest PC2) consisted Telimena, Sirocco, Goplana, Tdoz 1403, Tonika; group III (the lowest PC1 and PC2) consisted Arabella, Mandaryna, Kandela, Struna, Jarlanka; and group IV (the lowest PC1 and the highest PC2) consisted SMJ

2115, Sharki, Tybalt, Harenda, SMJ 1314, Ostka smolicka and Tdoz 1508.

The WCA was performed based on SI calculated for physiological traits (Figure 4(a)) and seven drought tolerance

a) PCA calculated for drought tolerance indexes



b) PCA calculated for listed wheat genotypes

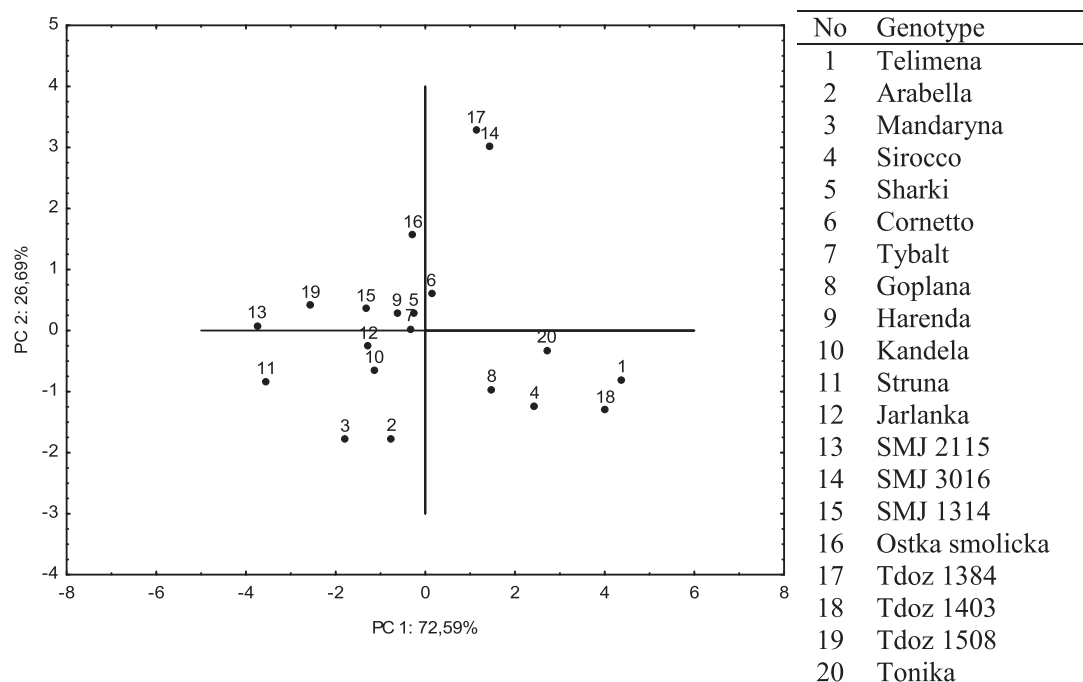
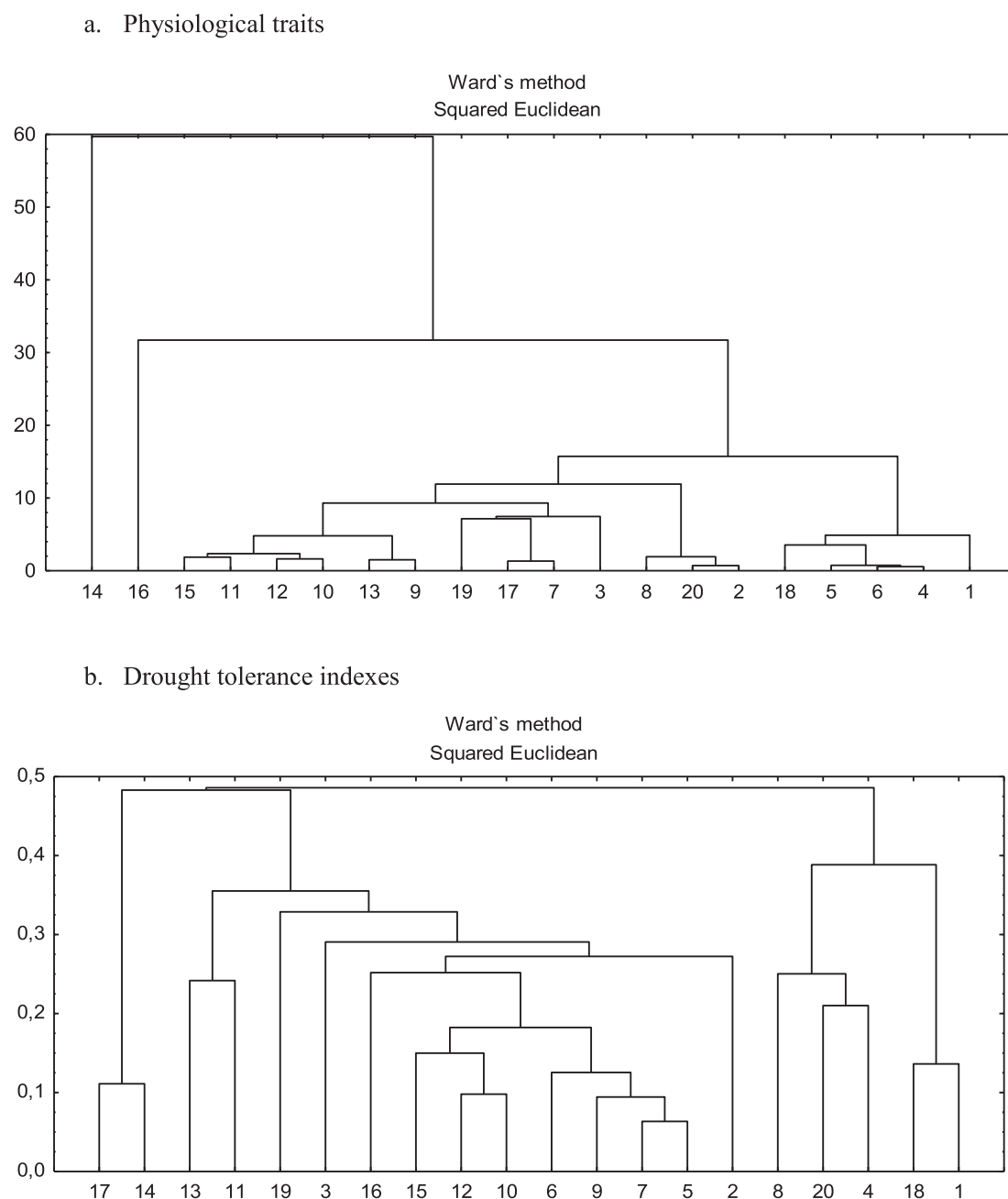


Figure 3. Principal component analysis (PCA) of drought tolerance indexes for (a) dry biomass (DM) and (b) genotypes.

indexes (Figure 4(b)). The wheat genotypes were clustered and presented in a dendrogram, and the distance between two clusters indicates the similarity of the genotypes. The genotypes classified as drought-sensitive i.e. Telimena, Sirocco, Sharki, Tdoz 1403 were located close to each other on the dendrogram. Similar trend was observed for drought-resistant genotypes, i.e. Struna, SMJ 2115 and Tdoz 1508 (Figure 4(a)). However, the WCA for seven drought tolerance indexes indicates the possibility of dividing genotypes into two groups (Figure 4(b)). The first group includes the genotypes classified as drought-sensitive i.e. Telimena, Sirocco, Goplana, Tdoz 1403 and Tonika. They had the highest DSI among genotypes. The second group, including Struna, SMJ 2115 and Tdoz 1508 genotypes, located close to each other on dendrogram, were characterized by lower DSI and small difference in DM (Table 4).

Leakage index (LI) and leaf water status (RWC, RT, WD)

For selected drought sensitive (Telimena, Sirocco, Goplana) and resistant (Struna, SMJ 2115, Tdoz 1508) genotypes we carried out measurements of membrane injury and 3 different parameters of leaf water status (Table 6, Figure 5). In group of drought sensitive genotypes, values of initial leakage were higher about 4 times after drought stress as compared to control conditions, whereas in a group of drought-resistant genotypes the values increased about 3 times. Changes in final leakage between drought sensitive and resistant genotypes were slight. As a consequence, values of membrane injury (LI) in the group of drought sensitive genotypes were about 2 times higher than for the group of drought-resistant genotypes. For the same genotypes,



1 – Telimena, 2 – Arabella, 3 – Mandaryna, 4 – Sirocco, 5 – Sharki, 6 – Cornetto, 7 – Tybalt, 8 – Goplana, 9 – Harenda, 10 – Kandela, 11 – Struna, 12 – Jarlanka, 13 – SMJ 2115, 14 – SMJ 3016, 15 – SMJ 1314, 16 – Ostka smolicka, 17 – Tdoz 1384, 18 – Tdoz 1403, 19 – Tdoz 1508, 20 – Tonika

Figure 4. Dendrogram of 20 wheat genotypes resulting from cluster analysis (Ward's methods) of stress index (SI) calculated for physiological traits (a) and drought tolerance indexes (b).

differences between sensitive and resistant ones in RWC were slight. Under control condition, RWC of sensitive genotypes was 72% and drought-resistant genotypes about 77%. Drought decreased RWC about 10% in the group sensitive genotypes and about 12% in the group of drought-resistant genotypes. Similarly, RT in seedlings grown in control conditions was about 79% and 82%, in the group drought sensitive and resistant genotypes, respectively. In the group of drought sensitive genotypes, decrease of RT in drought stressed seedlings was about 23%, whereas in the group of drought-resistant genotypes about 16%. These changes in RT significantly influenced leaf water

deficit (WD). In the group of drought sensitive genotypes, WD was higher than in drought-resistant genotypes, for both control and drought treatment. Statistical significant correlation was found between LI and each following parameter DM, GY, DSI and also between RWC and RT, WD.

Discussion

Global climate change has a negative impact on world ecosystems and the global economy (Hanson and Nelson 1985; McKersie and Leshem 1994; Ashraf 2010). Since 2000, there has been observed a decrease in yield of crops, including

Table 6. Membrane injury (LI), leaf water status (RWC, RT, WD) and linear correlation coefficient for 6 wheat genotypes. Mean values ($n = 5$) \pm SE ($df = 4$).

		Electrolyte leakage			Leaf water status						
Genotype	Treatments	Initial ($\mu\text{S g}_{\text{fw}}^{-1}$)	Final ($\mu\text{S g}_{\text{fw}}^{-1}$)	LI (%)	RWC (%)	RT (%)	WD (%)				
Drought sensitive genotypes											
Telimena	Control	12.2 \pm 1.0	190.2 \pm 8.5	23.0 \pm 3.2	73.2 \pm 2.9	85.9 \pm 2.1	14.1 \pm 2.1				
	Drought	37.3 \pm 2.6	133.3 \pm 6.7		69.7 \pm 1.3	64.1 \pm 3.3	35.9 \pm 3.3				
	% control	304.5	70.1		95.3	74.6	255.1				
Sirocco	Control	12.4 \pm 1.4	167.8 \pm 2.1	21.6 \pm 1.6	72.0 \pm 3.2	72.9 \pm 2.6	27.1 \pm 2.2				
	Drought	55.1 \pm 2.3	201.6 \pm 11.9		58.8 \pm 2.9	62.6 \pm 3.1	41.2 \pm 1.8				
	% control	446.1	120.2		81.7	85.9	152.0				
Goplana	Control	13.0 \pm 1.4	208.6 \pm 2.5	21.3 \pm 2.7	70.7 \pm 4.5	77.0 \pm 2.9	23.0 \pm 2.9				
	Drought	49.8 \pm 3.0	189.8 \pm 11.6		65.0 \pm 5.0	55.6 \pm 2.1	44.4 \pm 2.1				
	% control	382.9	91.0		91.9	72.2	193.3				
Mean	Control	12.5	188.9	22.2	72.0	78.6	21.4				
	Drought	47.4	174.9		64.5	60.8	40.5				
	% control	378.2	92.6		89.6	77.3	189.3				
Drought-resistant genotypes											
Struna	Control	15.3 \pm 1.8	141.5 \pm 5.2	11.0 \pm 1.5	77.6 \pm 2.4	80.8 \pm 1.7	19.2 \pm 1.7				
	Drought	39.1 \pm 0.5	189.2 \pm 4.5		69.0 \pm 1.0	67.9 \pm 3.2	32.1 \pm 3.2				
	% control	255.3	133.7		88.9	84.1	166.9				
SMJ 2115	Control	16.3 \pm 0.7	186.2 \pm 6.6	12.9 \pm 2.1	75.2 \pm 3.3	80.3 \pm 2.3	19.7 \pm 2.3				
	Drought	42.1 \pm 3.1	205.3 \pm 11.9		67.0 \pm 4.4	64.7 \pm 1.4	35.3 \pm 1.4				
	% control	258.2	110.3		89.1	80.5	179.2				
Tdoz 1508	Control	12.2 \pm 0.1	177.6 \pm 5.7	12.3 \pm 2.9	77.5 \pm 2.4	85.7 \pm 1.7	14.3 \pm 1.7				
	Drought	45.8 \pm 5.3	249.7 \pm 22.0		66.0 \pm 3.4	75.4 \pm 2.9	24.6 \pm 2.9				
	% control	374.1	140.6		85.2	88.0	171.9				
Mean	Control	14.6	168.4	12.2	76.8	82.3	17.7				
	Drought	42.3	214.7		67.3	69.3	30.7				
	% control	289.5	127.5		87.7	84.3	172.9				
Correlation coefficient r (df = 4)											
Indexes/Traits	RWC	RT	WD	DM	GY	DSI	STI	M _{PRO}	M _{HAR}	TI	GMG
LI	0.276 ^{NS}	−0.605 ^{NS}	0.503 ^{NS}	−0.957 ^{***}	−0.969 ^{***}	0.806 [*]	−0.265 ^{NS}	−0.086 ^{NS}	−0.273 ^{NS}	0.707 ^{NS}	−0.182 ^{NS}
RWC		−0.849 ^{**}	0.872 ^{**}	−0.247 ^{NS}	−0.272 ^{NS}	0.504 ^{NS}	0.268 ^{NS}	0.361 ^{NS}	0.239 ^{NS}	0.585 ^{NS}	0.301 ^{NS}
RT			−0.719 ^{NS}	0.598 ^{NS}	0.522 ^{NS}	−0.702 ^{NS}	−0.184 ^{NS}	−0.328 ^{NS}	−0.183 ^{NS}	−0.718 ^{NS}	−0.256 ^{NS}
WD				−0.387 ^{NS}	−0.520 ^{NS}	0.576 ^{NS}	−0.106 ^{NS}	0.023 ^{NS}	−0.148 ^{NS}	0.591 ^{NS}	−0.064 ^{NS}

Notes: *, **, *** – statistical significant correlation coefficient at level $p < .5$, $.1$ and $.01$, respectively, NS – not significant differences.

rice, wheat, and maize. Following climate changes, scientists and breeders are continually improving genotypes of crop plants in terms of plant tolerance for drought by both, traditional selection methods and genetic engineering (Hanson and Nelson 1985; Blum 1996; Ehlers and Goss 2003; Jackson and Ram 2003; Fleury et al. 2010). The current research concerning plant tolerance to drought focuses mainly on traits and mechanisms that can be modified. Despite trends in science, improved agronomy practices, and developed methodology, the progress in the research related to the drought stress is slower than expected, and scientific assumptions fail in the natural environment. The tolerance of plants to drought stress is a complex feature, and finding suitably dedicated phenotypes by their exposure to stress in natural conditions, or their treatment to simultaneously or sequentially occurring stress in laboratory conditions is very difficult. The next and perhaps the most important reason of the lack of success is the fact that the genotype itself is not the only factor determining the survival in suboptimal conditions (Palta 1990; Mittler 2006; Wang et al. 2003).

In many research, mathematic formulas and statistical methods (drought tolerance indexes, correlation between biomass or grain yield and physiological traits, biplot analysis) were used to identify the genotypes responding to drought stress condition (Drikvand et al. 2012; Marcińska et al. 2017). The available formulas related to the drought stress were published in reports: Fischer and Maurer (1978); Golbashy et al. (2010); Doorenbos and Pruitt (1977), Doorenbos and Kassam (1986) Grzesiak et al. (2012). Similarly, in many studies on drought tolerance in cereals, rankings of resistant/sensitive genotypes were created for wheat

(Ahmadizadeh et al. 2002; Talebi et al. 2009; Geravandi et al. 2011), maize (Golbashy et al. 2010, Khayatnezhad et al. 2011; Grzesiak et al. 2012), triticale (Grzesiak et al. 2012), and oat (Akcura and Ceri 2011; Hisir et al. 2012; Rabiei et al. 2012). On the other hand, the effectiveness of different physiological tests has not been definitely confirmed during the evaluation of drought tolerance degree (Hanson and Nelson 1985; Bandurska and Stroński 2003).

The wheat genotypes used in our experiments show a relatively wide range of drought tolerance. In this study, differences and interactions between the physiological traits and drought tolerance indexes were found in spring wheat genotypes grown under non-stress vs drought stress conditions (Tables 1 and 2). We have shown the existence of genetic variation of a drought sensitivity degree and the possibility of selection of favorable genotypes. Using our tests and analyses, it seems to be possible to distinguish groups of wheat genotypes with a low, moderate and high susceptibility to drought. We found the genotypes that produced stably high yield of biomass (DM) and grain (GY) in both, drought and control conditions, semi-stable genotypes producing lower DM and GY, and highly sensitive genotypes with the lowest DM and GY. The stable genotypes had lower DSI, when compared to semi-stable or sensitive ones. Similarly to other authors' results, our data confirm the observations that DSI, STI and SI are useful in order to select suitable genotypes under control and drought conditions. Furthermore, index of geometric mean productivity (GMP) enable selection of the genotypes with high yield in both treatments. Statistical analysis PCA and WCA confirmed strong correlation between DSI, the number and dry weight of grain, and production of different

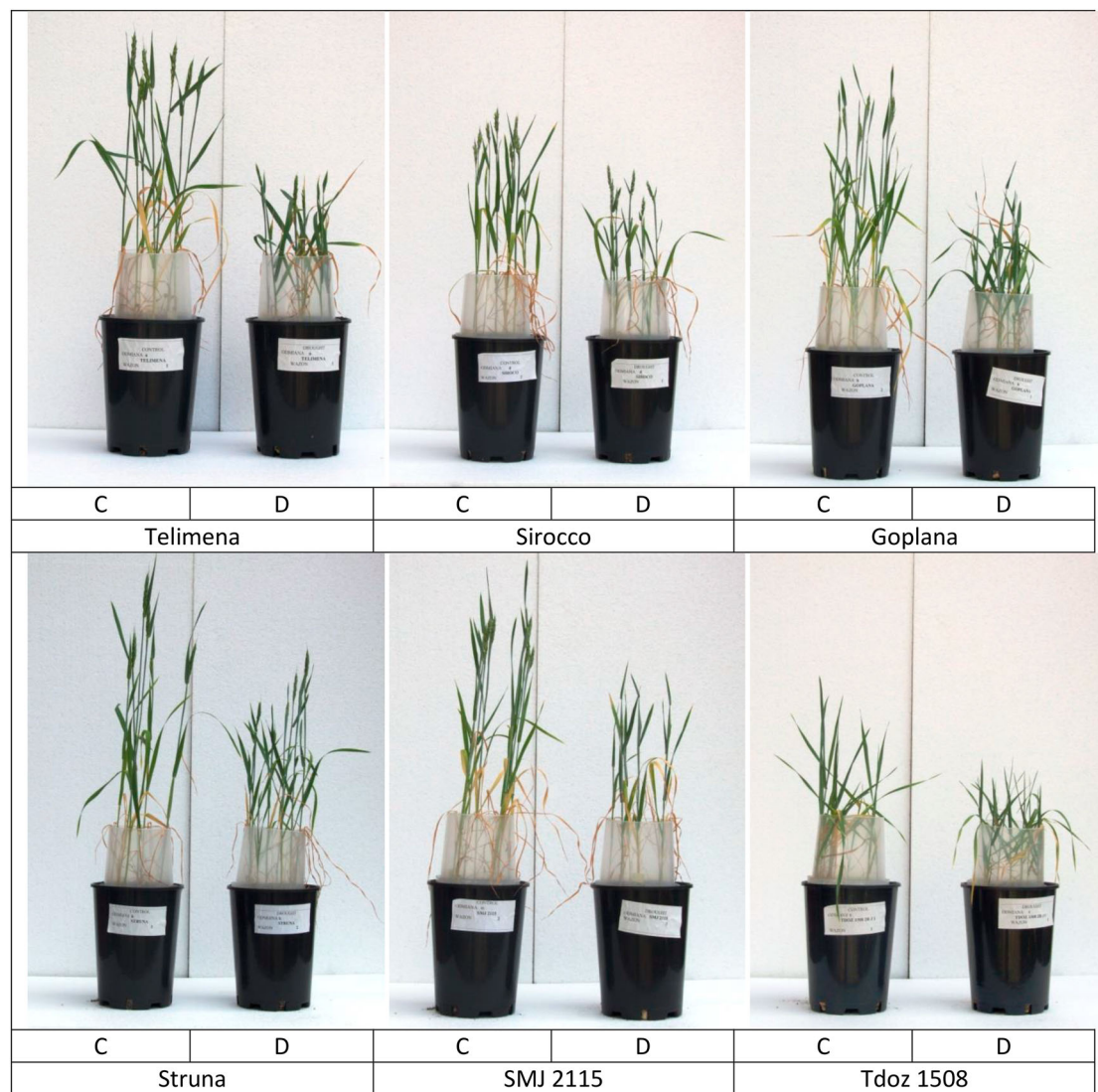


Figure 5. The photographs of 11-week-old wheat plants grown in control (C) or drought (D) conditions. Telimena, Sirocco and Goplana were selected as drought-sensitive genotypes and Struna, SMJ 2115 and Tdoz 1508 as drought-resistant genotypes.

yield components. Last but not least, the analysis allowed to select the groups of genotypes differing in resistance/sensitivity to drought stress. These findings were consistent with earlier studies (Golabadi et al. 2006; Mohammadi et al. 2010) conducted on wheat and mung bean (Zabet et al. 2003), as well as triticale and maize genotypes (Grzesiak et al. 2012).

Moreover, different plant traits can be used for a genotype screening, and to determine genotypic tolerance to stress. Biomass, grain yield, plant height, number of tillers, area and number of leaves, and leaf greening were used for this purpose in our experiments, and they were proved as a highly correlated. For example, the decrease of plant height and leaf area was related to reduction of yield.

Dehydration of plants tissues impairs various physiological processes, especially the changes in leaf water potential and photosynthesis. Decrease in leaf water content initially induces a stomatal closure, resulting in a decrease in the supply of CO_2 to the mesophyll cells and subsequently in a decrease in the rate of photosynthesis (Williams et al. 1999; Lawlor and Cornic 2002). Other mechanisms induced under prolonged or severe drought can include changes in chlorophyll synthesis, functional and structural changes in chloroplasts, disturbance in accumulation and distribution of assimilation products (Berkowitz et al. 1983; Cornic and

Masacci 1996; Muller and Whitsitt 1996; Medrano et al. 2002; Flexas and Medrano 2002). However, there is a controversy, whether drought limits photosynthesis due to leaf water status, stomatal closure, metabolic impairment, or injuries to photosynthetic apparatus. Studies of different author, show also that the biomass accumulation and changes in yield traits were associated with an increase in efficiency of PS II (Lichtenthaler and Babani 2004; Lichtenthaler et al. 2005; Hura et al. 2007; Liang et al. 2010, Rapacz et al. 2010). According to Souza et al. (2004), photosystem PSII is relatively resistant to water deficit, and the photochemical reaction are inhibited only in the prolonged drought. Some authors indicate that the stabilization of the PSII complex depends on an increase in concentration of osmotically active substances. The osmotic adjustment of cells was shown to maintain photosynthesis during drought (Shangguan and Shao 1999). Drought causes reduction in the amount of chlorophyll and the disruption of thylakoid membranes (Wright et al. 2009; Zhang et al. 2011). In response to drought, plants exhibit uncontrolled generation of reactive oxygen species (ROS) in cells and disturbances in the electron transport in the respiratory chain and in the light phase of photosynthesis (Starck 2005). ROS can also influence membrane lipid photoinhibition. In this work, the results of the

number of tillers (NoT) and leaves (NoL_{MT}, NoL_{AT}, NoL_T) indicate that within the wheat genotypes there are different strategies to counteract the harmful effects of drought. In optimal conditions, the plants maintain the most favorable relations between the size of the assimilation surface, transpiration and chlorophyll content in order to keep photosynthesis at an optimal level. In drought conditions, plants reduce the number and area of the leaves by inhibiting the growth of the leaf surface and limiting their number, e.g. by leaf drying and falling off. The results presented in Table 3 indicate a drastic decrease in GY in the drought-sensitive genotypes (Telimena, Sirocco, Goplana), which was accompanied by a relative reduction of about 50% tillers and about 30% leaves. In contrast, in drought-resistant genotypes (Struna, SMJ 2115, Tdoz 1508) yield reduction was significantly lower than in sensitive genotypes, and was accompanied by a relatively higher or similar number of tillers and the number of evolved leaves. In our earlier work, we also found the importance of the ratio of aboveground parts to the roots (Grzesiak et al. 2012). In drought-resistant genotypes of maize and triticale subjected to the drought, these relation was similar to the level of control plants, whereas the susceptible genotypes had higher ratio due to reduced mass and number of roots (Grzesiak et al. 2012).

Conductometric measurement of membrane injury (LI) was applied as a screening test for the estimation of tolerance to various stresses (Blum and Ebercon 1981; Palta 1990). The ability to maintain the structure and function of cytoplasmic membranes under water deficit is one of the most important physiological traits. Differences between sensitive and resistant genotypes might result from the fact that drought-resistant genotypes possess more efficient mechanisms protecting membrane function and structure by stronger binding of molecules to the membrane lipid-protein complex (Smirnoff and Colombe 1988; Szechyńska-Hebda et al. 2016). Drought causes a loosening of lamellar membranes in chloroplasts, certain amount of grana, and increase in the level of coarse grain matrix (Haupt-Harting and Fock 2002; Lawlor and Cornic 2002; Tang et al. 2002). Both, our results and other authors' results indicate that measurements of electrolyte leakage from plant tissues are an effective method of assessing the degree of sensitivity to drought stress. In these studies we showed high statistical significant correlations between the degree of sensitivity of various wheat genotypes with the membrane damage (LI). For the genotypes sensitive to drought (Telimena, Sirocco, Goplana) LI values were about two times higher than in the genotypes resistant to drought (Struna, SMJ 2115, Tdoz 1508). This confirms that stress-resistant genotypes can develop more effective mechanisms to protect the structure and function of membranes than stress-sensitive genotypes.

PCA and WCA were a better approach than a linear correlation in the identification of genotypes resistant and sensitive to stress both, in our research, and studies of other authors (Talebi et al. 2009; Zhang et al. 2011; Parihar et al. 2012; Zaheri and Bahraminejad 2012; Nouraein et al. 2013). According to Iezzoni and Pritts (1991) PCA analysis simplify and reduces a large data set to a small number of unrelated groups of variables, which can be scored along independent, linear axes. Variables strongly associated in the same group may share some underlying biological relationship. The response of the groups of variables to imposed conditions can be quantified using a combination of PCA analysis and

univariate statistics. Therefore, PCA analysis has been used successfully to develop novel hypotheses, and to understand the response of complex traits to imposed treatments. Our results shown that the clustering of the genotypes into groups differing in drought tolerance on the basis of PCA was on the whole similar to the results obtained from WCA. The PCA and WCA allowed to separate the groups of similarity and thus indicated that tolerance to drought in spring wheat has a common genetic background.

Conclusion

Our results confirm the existence of a wide range of genotypic variability in response to drought in cereal plants. In wheat, similarly to other crop plants, the physiological origins for this variability have not as yet been fully recognized and explained. It is a widely shared opinion that the breeding of drought-resistant crop plants will not be an easy task. Breeders are interested in improving drought resistance, while maintaining high quantity and quality of yield. Our studies contribute to the understanding of responses of different spring wheat genotypes to drought on basis of the determination of different drought tolerance indexes and physiological traits. Thus, it can be assumed that the tests performed in this work may serve as useful tools in estimating the degree of tolerance to drought stress in wheat. Physiological laboratory methods i.e. membrane injury and leaf water status were simple tools, useful in evaluation of plant responses to drought stress. Together with PCA and WCA this traits can be used as a suitable method for studying the complex resistance mechanisms and to increase yield efficiency in breeding programs. However, the variability of the groups of genotypes determined by both, PCA and WCA should be verified by physiological and molecular experiments in natural growth conditions.

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