



## RESEARCH ARTICLE

# Multivariate analysis of drought stress indices to assess bread wheat (*Triticum aestivum* L.) germplasm under rainfed conditions

Vishnu Kumar\*,<sup>s</sup>, Vikram Singh<sup>1</sup>, Arun Gupta, Vikas Gupta and Gyanendra Pratap Singh<sup>s</sup>

## Abstract

Water stress impacts wheat yields and poses a serious threat to stabilizing the global food supply. In the present investigation, 71 diverse bread wheat germplasm accessions were evaluated at Karnal and Hisar under normal and rainfed (four environments) conditions during *rabi*, 2022-23. The overall pooled grain yield showed a reduction of 15.6% under water stress, while days to heading, tiller count and plant height were reduced by 8.1, 15.5 and 12.5%, respectively. The genotypes, namely K8027, HI1531, PBW175, UAS375, WH1142, HI1612 and K1317 showed higher NDVI values both at heading and grain filling stages under water stress conditions. Grain yield showed positive and significant associations with 1000-grain wt. ( $r=0.62^{***}$ ) and no. of tillers/m ( $r=0.53^{***}$ ). In the principal component biplots, 14 drought stress indices were grouped into three clusters. The genotypes K9465, HD2987 and K8027 were the three top rankers for drought susceptibility index; however, they showed yield reduction of 10.4, 5.5 and 17.4% over the best check NIAW3170. Most of the drought stress indices considered only the grain yield reduction as selection criteria; however, the yield potential coupled with drought tolerance is desired to get favourable gene constellations. The drought resistance index (DRI) was highly successful in identifying high-yielding and drought-tolerant genotypes. The genotypes *viz.*, HD3171, MP1358, 20<sup>th</sup> HTWYT-48, 29<sup>th</sup> SAWYT-316, WAP91 and K1317, appeared to be high-yielding and water stress-tolerant. The traits, such as 1000 grain weight, no. of grains/spike and tiller count, can be targeted as on-farm selection criteria under water stress and to train genomic models.

**Keywords:** Wheat, water stress, drought stress indices, PCA biplot

## Introduction

Wheat is an important cereal crop, occupying 220 m ha of area worldwide, followed by maize (208 m ha) and rice (168.35 m ha). However, in terms of global production, it ranked third (799 m t) after maize (1241.55 m t) and rice (800 m t) with an average productivity of 3.65 t ha<sup>-1</sup> (FAOSTAT, 2025). Wheat is a staple food crop and caters to the daily dietary needs of 35% of the global population (Allahverdiyev et al. 2025; Sangha et al. 2025). India is the second largest wheat producer and an all-time high wheat production of 117.5 m t has been projected for the year 2024-25. India shares nearly 13.8, 31.4 and 69% of the global, Asian and South Asian wheat production. Global wheat production has witnessed an increase of 36% since 2000. Similarly, Asian, South-Asian and Indian wheat production has also witnessed an increase of 38.3, 45.6 and 45%, respectively. However, the global and Asian wheat acreages showed a marginal increase of 2.45 and 2.38% over the year 2000. In contrast, South-Asian and Indian wheat areas were increased by 11.39 and 14.26%, respectively. In contrast, the Indian population showed an upward growth of 38% during the past 25 years, which mainly depends on wheat for food

and protein needs (Mottaleb et al. 2023). Kumar et al. (2025) reported that food demand will rise by 60 to 70% of the present supply by 2050 to feed the global population. Wheat has seen remarkable growth; however, the challenges, like

---

ICAR-Indian Institute of Wheat & Barley Research, Karnal 132001, Haryana, India

<sup>1</sup>Vikram Singh, Chaudhary Charan Singh Haryana Agricultural University, Hisar 125004, Haryana, India

<sup>s</sup>Present address: Vishnu Kumar and Gyanendra Pratap Singh, ICAR-National Bureau of Plant Genetic Resources, Pusa Campus, New Delhi 110012, India

**\*Corresponding Author:** Vishnu Kumar, ICAR-Indian Institute of Wheat & Barley Research, Karnal 132001, Haryana, India, E-Mail: vishnupbg@gmail.com

**How to cite this article:** Kumar, V., Singh, V., Gupta, A., Gupta, V., Singh, G.P. 2025. Multivariate analysis of drought stress indices to assess bread wheat (*Triticum aestivum* L.) germplasm under rainfed conditions. Indian J. Genet. Plant Breed., **85**(4): 546-554.

**Source of support:** ICAR, DARE, Govt.of India

**Conflict of interest:** None.

**Received:** Aug. 2025 **Revised:** Nov. 2025 **Accepted:** Nov. 2025

---

soil health, depleting natural resources, global warming and changed pest and disease dynamics, need to be addressed sustainably (Qian et al. 2025).

The impact of climate change is conspicuous, where effects of drought stress are largely detrimental and can reduce crop yields to the tune of 50% under severe stress conditions (Bapela et al. 2022). Nearly 50% of the global wheat area, including Africa, Australia, regions of China, parts of India and Russia, is largely under rainfed conditions and prone to low wheat yields (Pequeno et al. 2021). Drought impacts crop growth and exerts cytotoxic effects on physiological processes, including photosynthetic rate, stomatal conductance, cell membrane stability, chlorophyll content, etc. (Sairam and Saxena 2000; Zhao et al. 2020; Nyaupane et al. 2024; Priya et al. 2025). The low seedling survivability under water stress is one of the major yield-limiting factors, along with various other factors (Tomar and Kumar 2004). The stake of irrigated land varies, from Africa (4%) of the total cropped area to South Asia (42%). In the Indian scenario, 45% of the net sown crop area is under rainfed conditions. Increasing wheat yields under changing climate and drought stress is challenging as drylands share 47% of the global land and 66% of global wheat area (Mohammadi 2018; Qiu et al. 2022; Mutanda et al. 2025; Arif et al. 2025). The water stress experienced at the post-anthesis, grain filling and terminal stages is the most critical in wheat (Farooq et al. 2014). Under such a situation of scanty irrigation and rainfed agriculture, the challenges are enormous to feed the burgeoning population and stabilize the agricultural food system. Besides improving irrigation facilities, the deployment of water stress resilient wheat genotypes is an economic, viable and eco-friendly solution.

In general, the Indian wheat genotypes, namely, K8027, K9465, HINDI62, HI1500, C306, K1317, NIAW34, NI5439 and PBW175, are considered drought-tolerant and used as one of the parents in the crossing programs. (Subrahmanyam et al. 2006; Kadam et al. 2012; Kumar et al. 2018; Pandey et al. 2023). These genotypes are low-yielding, susceptible to wheat rusts and repeated use in the breeding programs leads to the loss of genetic diversity. The narrow genetic base, poor specific combining ability and low extent of heterosis are other obstacles with the above genotypes in generating stress-tolerant materials (Balla et al. 2025). For instance, the most widely used parent, C306, is a *Ne1* carrier and shows hybrid mortality at the hybrid seedling stage (Vikas et al. 2013; Manoj et al. 2020). Here, the highly diverse nature of the experimental materials, real field conditions and application of 14 drought stress indices gives a comparative advantage over the previous studies. Therefore, the present study was aimed at investigation the response of water stress based on multivariate analysis of drought stress indices to identify high-yielding and drought-tolerant genotypes.

## Materials and methods

### Planting materials and experimental site

The present study was carried out to assess the effect of water stress on 71 bread wheat (*Triticum aestivum* L.) genotypes grown under four environments (normal and water stress) during *rabi*, 2022-23. The experiments were conducted at Karnal and Hisar locations under rainfed early sown (October 2022: water stress) and irrigated timely sown (November 2022) conditions. An extremely diverse wheat germplasm (71) comprised of registered unique germplasm for water stress (12), released cultivars (30), advance breeding lines (14) and exotic breeding lines selected from CIMMYT yield trials (11), along with four check varieties (C306, DBW110, DBW296 and NIAW3170), was evaluated under rainfed and irrigated conditions. The released cultivars were selected out of a database of nearly 450 released varieties being grown under different agro-ecologies of rainfed and restricted irrigation (water stress), timely sown (high yield potential), late and very late sown (early maturity: escape mechanism) and well reported with drought stress resilience. Similarly, 11 exotic promising lines were selected from the CIMMYT trials 29<sup>th</sup>SAWYT and 20<sup>th</sup>HTWYT evaluated during *rabi*, 2021-22, whereas 14 advanced breeding lines were identified from 115 advanced yield trials conducted during *rabi* 2021-22. The experimental materials, like unique registered germplasm, released varieties and exotic lines, were received from the gene bank of the ICAR-Indian Institute of Wheat & Barley Research, Karnal, Haryana, India. The parentage details of the studied research materials are given in Table 1. The Karnal is situated at 29.70°N & 76.99°E and Hisar is located at 29.14°N & 75.70°E. The soil type of Karnal and Hisar is sandy-loam and loamy-sand.

### Experimental details

The rainfed experiments were sown during 25-30 October 2022 and no supplementary irrigation was applied during the entire crop period. During the rainfed crop period (up to 20<sup>th</sup> March 2023), the crop received total rainfall of 14.3 mm and 24.0 mm at Hisar and Karnal locations, respectively. The details of weather parameters recorded are presented in the Supplementary Fig. 1. Under normal irrigated conditions, the irrigated experiments were sown during 05-10 Nov. 2022, following the standard package of practices with full optimum irrigations. The experiments were conducted in the augmented block design (ABD) with 05 blocks, where 04 checks (C306, DBW110, DBW296 and NIAW3170) were repeated in each block randomly in all four environments. The paired rows of 3 m row length spaced 20 cm apart were grown at each location.

### Statistical analysis

The data were recorded for days to heading (DH), days to maturity (DM), effective tillers count/m (TPM), spike length

**Table 1.** Details of the bread wheat germplasm evaluated for water stress tolerance

S No.	Genotype	Sym.	Year	Trait/production condition	ID for registered germplasm	Sr No.	Genotype	Sym.	Year	Trait/production condition	ID for registered germplasm
<b>Trait specific germplasm</b>											
1	HN1DI62	G1	1997	Drought tolerance	IC296681	37	GW 513	G37	2021	IR-TS-CZ	
2	KRL-99	G2	2007	Tolerant to abiotic stress	IC546936	38	HD 3298	G38	2021	VLS-IR-NWPZ	
3	WCF8-HT13	G3	2005	Drought tolerance	IC443622	39	MP 1358	G39	2021	RI-TS-PZ	
4	AKAW3717	G4	2010	Drought tolerance	IC0582907	40	HI 1633	G40	2021	IR-LS-PZ	
5	DWRL1	G5	2012	Lodging resistance	IC0590878	41	HI 1634	G41	2021	IR-LS-CZ	
6	WCF 12-19	G6	2011	Drought tolerance	IC0594378	42	HI 1636	G42	2021	IR-TS-CZ	
7	WCF 12-208	G7	2011	Drought tolerance	IC0594379	43	20 <sup>th</sup> HTWYT-43	G43	2021	CIMMYT	
8	WCF 12-61	G8	2011	Drought tolerance	IC0594377	44	20 <sup>th</sup> HTWYT-2	G44	2021	CIMMYT	
9	WCF 12-7	G9	2011	Drought tolerance	IC0594376	45	20 <sup>th</sup> HTWYT-48	G45	2021	CIMMYT	
10	EC 531185	G10	2018	Drought tolerance	EC531185	46	20 <sup>th</sup> HTWYT-25	G46	2021	CIMMYT	
11	HTW63	G11	2019	Drought tolerance	IC36761	47	20 <sup>th</sup> HTWYT-41	G47	2021	CIMMYT	
12	RW5	G12	2020	Drought tolerance	IC635019	48	20 <sup>th</sup> HTWYT-13	G48	2021	CIMMYT	
<b>Advanced Breeding Lines</b>											
13	HD2781	G13	2002	RF-TS-PZ		49	20 <sup>th</sup> HTWYT-32	G49	2021	CIMMYT	
14	HD2987	G14	2011	RF/RI-TS-PZ		50	20 <sup>th</sup> HTWYT-42	G50	2021	CIMMYT	
15	HI 1500	G15	2003	RF-TS-CZ		51	29 <sup>th</sup> SAWYT-334	G51	2021	CIMMYT	
16	HI 1531	G16	2006	RF/RI-TS-CZ		52	29 <sup>th</sup> SAWYT-320	G52	2021	CIMMYT	
17	UAS 375	G17	2018	RF-TS-CZ		53	29 <sup>th</sup> SAWYT-303	G53	2021	CIMMYT	
18	HUW234	G18	1986	IR-LS-NEPZ		54	29 <sup>th</sup> SAWYT-341	G54	2021	CIMMYT	
19	HUW468	G19	1999	IR-TS-NEPZ		55	29 <sup>th</sup> SAWYT-310	G55	2021	CIMMYT	
20	HW 2004	G20	1997	RF-TS-CZ		56	29 <sup>th</sup> SAWYT-316	G56	2021	CIMMYT	
21	K8027	G21	1989	RF-TS-NEPZ		57	WAP83	G57	2021	Indigenous sel.	
22	K9465	G22	1998	RF-LS-NEPZ		58	WAP71	G58	2021	Indigenous sel.	
23	NI 5439	G23	1975	RF/RI-TS-PZ		59	WAP92	G59	2021	Indigenous sel.	
24	NIAW 34	G24	1997	IR-LS-MH		60	WAP84	G60	2021	Indigenous sel.	
						61	WAP75	G61	2021	Indigenous sel.	

Contd....

(SL), no. of grains/spike (GPS), and 1000-grain wt. (TKW) and grain yield/plot (GY). In addition, NDVI values and canopy temperature at heading and grain filling stages were also observed with the hand-held devices following standard procedures. The data of all four environments were analysed using the R package Augmented RCBD and pooled analysis of variance was employed as suggested by Federer et al. (2001). The correlation plots were generated using the R package "Metan" in version R 4.1.3. The principal component analysis (PCA) was performed in R 4.1.3 employing FactoMineR, Factoextra and ggplot2 packages. Here, 14 drought stress indices, namely, yield index (YI), yield stability index (YSI), tolerance (TOL), mean productivity (MP), Harmonic mean (HM), drought susceptibility index (DSI), geometric mean productivity (GMP), drought resistance index (DRI), drought tolerance index (DTI), relative drought index (RDI), stress susceptibility per cent index (SSPI), mean relative performance (MRP), percent yield reduction (PYR) and abiotic tolerance index (ATI) were calculated in MS-Excel. These indices were estimated as given in Farshadfar et al. (2014), Singh et al. (2018) and Lamba et al. (2023).

## Results

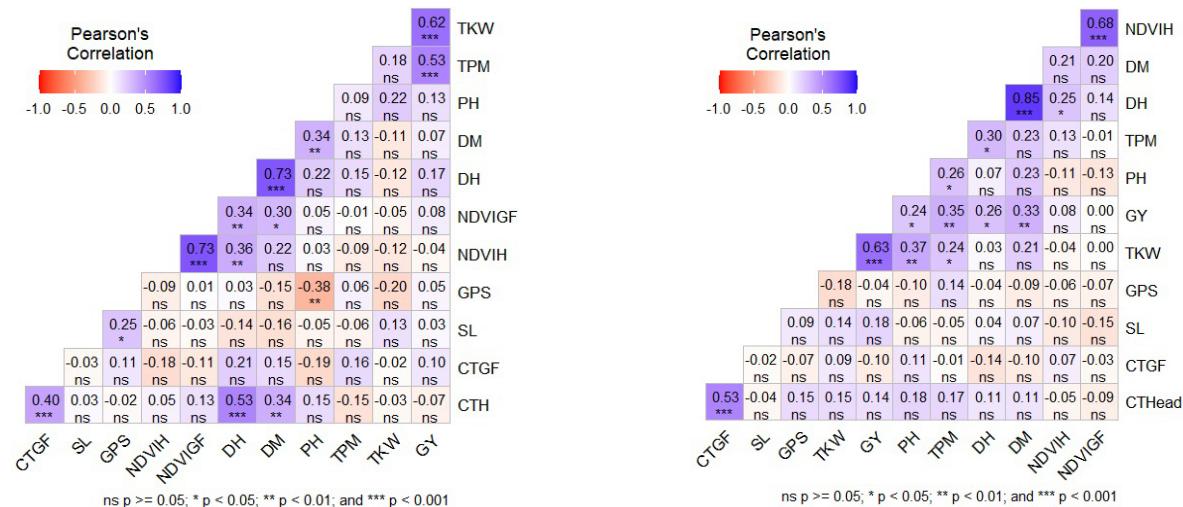
## ***Analysis of variance and mean grain yield***

The individual locations analysis of variance (ANOVA) depicted significant mean squares for treatments (eliminating blocks), test entries and check genotypes under stress and favourable environments (Table 2). The combined ANOVA revealed significant mean squares for genotypes, environments and genotype  $\times$  environment interactions ( $p < 0.001$ ), indicating substantial genetic variations and impact of rainfed conditions on the yield potentials (Supplementary Table S1).

The pooled mean grain yield under water stress was obtained as 610 g, which was reduced by 15.6% to the irrigated conditions. Karnal and Hisar mean rainfed grain yields were observed as 578 g and 642 g, respectively. The test genotype WAP91 ranked first (844 g), followed by HD3171 (831 g), 29<sup>th</sup> SAWYT-316 (820 g), WAP92 (815 g) and MP1358 (799 g) under stress environment. The check entry NIAW3170 performed better (754 g) than other checks, namely C306 (628 g), DBW296 (587 g) and DBW110 (569 g). The genotypes, namely 20<sup>th</sup> HTWYT-43 (G43), showed the highest yield reduction under water stress (49.5%), followed by WCF12-61 (G8: 38.03%), AKAW3717 (G4: 37.20%) and HI1612 (G32: 36.9%). Under irrigated conditions, the genotype WAP92 ranked first with 945 g pooled grain yield, followed by 29<sup>th</sup> SAWYT-310 (935 g), 29<sup>th</sup> SAWYT-303 (915 g), HI1612 (909 g) and HD3298 (888 g). Location-wise, the genotypes, namely 29<sup>th</sup> SAWYT-310, WAP92, 29<sup>th</sup> SAWYT-303, HI1612 and MP1358 were high yielding at Karnal, while at Hisar the genotypes WAP91, WAP114, WAP75, WAP66 and HI1612 performed better.

**Table 2.** Analysis of variance for individual environments under water stress and irrigated conditions

Source	DF	Mean Sum of Squares (MSS)			
		Rainfed environments		Irrigated environments	
		Karnal	Hisar	Karnal	Hisar
Block (ignoring Treatments)	4	29386.32 **	226989.84 **	75909.27 **	256183.16 **
Treatment (eliminating Blocks)	70	14025.18 **	27382.96 **	11544.36 **	24136.14 **
Block (eliminating Treatments)	4	8027.2	6181.8	1701.2	3330.45
Treatment (ignoring Blocks)	70	15245.7 **	40000.56 **	15784.82 **	38584.87 **
Test and Test vs. Check	67	14114.99 **	25480.8 **	10908.88 **	23639.6 **
Treatment: Check	3	12019.4 *	69864.58 **	25736.8 **	35225.65 **
Treatment: Test	66	15452.53 **	39072.37 **	15518.89 **	38385.58 **
Treatment: Test vs. Check	1	11274.25 ns	11668.86 *	3480.71 ns	61815.76 **
Residuals	12	3437.07	2427.5	2714.13	4637.98

**Fig. 1.** Correlations under rainfed and irrigated condition

### Yield attributes

The genotype 20<sup>th</sup> HTWYT-32 showed early days to heading (85 days), followed by WCF12-7, EC531185, 29<sup>th</sup> SAWYT-303, WCF12-19 and WCF12-61 (all  $\leq$  88 days) under congenial conditions. These genotypes exhibited physiological maturity at 135 days, except 29<sup>th</sup> SAWYT-303 (138 days). Under water stress condition, 13 genotypes, including 20<sup>th</sup> HTWYT-13, 25, 32, 41, 43 & 48, GW513, EC531185, WCF12-7, HI1634, HD2987, HUW234 and K9465 showed early spike emergence ( $\leq$  80 days). The genotype DWRL-1 was observed with the shortest plant height under rainfed (58 cm) and irrigated (82 cm) conditions. The other genotypes with short plant height ( $\leq$  95 cm) under irrigated conditions were KRL-99, WCF12-19 and HD2987. Under stress environment, 15 bread wheat genotypes (HD2987, EC531185, NIAW34, 20<sup>th</sup> HTWYT-25, HI1633, HUW468, KRL-99, etc.) were recorded with plant height  $\leq$  90 cm. The highest grains/spike were observed for DWRL-1 (80) under moisture stress, followed by 29<sup>th</sup> SAWYT-320 (73), HI1612 (72), WAP71 (71) and WAP91

(68) under water stress. The genotypes WAP91, WAP83, and HD3298 depicted higher tillers/m under stress conditions. Among check varieties, NIAW3170 had the highest TKW (45 g) and 18 genotypes, including 11 exotic CIMMYT selections, 03 indigenous selections (WAP48, 75 & 96) and 04 released varieties (HI1621, HI1628, DBW296 and K8027) showed higher TKW ( $>45$  g) under rainfed conditions.

### Physiological characters

NDVI at heading stage ranged from 0.62 (K8027) to 0.36 (20<sup>th</sup> HTWYT-2) under water stress, while it varied from 0.78 (HI1531) to 0.38 (HI1628) under irrigated conditions. The genotypes, namely K8027, HI1531, PBW175, UAS375, WH1142, HI1612, K1317, HD3271, WAP96, PBW175, PBW596 and HI1621 showed higher NDVI values both at heading and grain filling stages under water stress conditions. All the check varieties exhibited almost similar NDVI at heading (0.47) and grain filling stages (0.37-0.41) under moisture stress conditions. Under irrigated (IR) condition, the genotypes viz., HI1531

showed the highest NDVI (0.70) at heading stage, followed by UAS375 (0.68), HD3271 (0.68), WAP96 (0.68), K1317 (0.67), 20<sup>th</sup> HTWYT-25 (0.67), WAP91 (0.67), HI1612 (0.67) and HINDI62 (0.67). Whereas, under irrigated conditions, the genotypes HD3271, UAS375, K9465, HI1531, PBW596 and 29<sup>th</sup> SAWYT-341, WH1142, HUW234 and HI1612 were found with higher NDVI values (0.54). For canopy temperature at heading stage under stress environment, HUW234, 20<sup>th</sup> HTWYT-43, NIAW34, WAP91 showed cooler canopy, while the genotypes HUW234, 20<sup>th</sup> HTWYT-43, HI1628, WCF12-7 and WCF12-19 were promising at the grain filling stage. Under IR conditions, the genotypes EC531185, K1317, WCF12-7, 20<sup>th</sup> HTWYT-48, K9465, HD3171 and HUW234 showed lower CT at the heading and grain filling stages.

### Drought stress indices

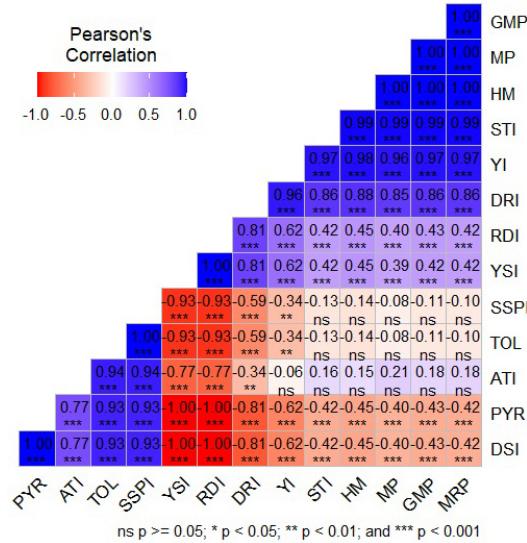
Here, 14 widely accepted drought stress indices were estimated and presented in the Supplementary Table S2. The yield index (YI) ranged from 0.39 (DWRL-1) to 1.38 (WAP91) and the stress-tolerant genotypes identified with higher YI are WAP91 (1.38), HD3171 (1.36), 29<sup>th</sup> SAWYT-316 (1.34), WAP92 (1.34) and MP1358 (1.31). YSI, MP, HM and GMP ranged from 0.50 (20<sup>th</sup> HTWYT-43) to 1.06 (K9465), 276.7 (DWRL-1) to 879.7 (WAP92), 271 (DWRL-1) to 874.9 (WAP92) and 273.8 (DWRL-1) to 877.3 (WAP92), respectively. Among the checks, the lowest DSi was depicted by NIAW3170 (0.78), whereas the genotype K9465 ranked first, followed by HD2987, K8027, 29<sup>th</sup> SAWYT-316, WAP91, HD3171, 20<sup>th</sup> HTWYT-48 and PBW596 (all<0.30). Based on SSPI, PYR and ATI, the genotype K9465 ranked first, followed by HD2987 and K8027.

### Morpho-physiological correlations

Under moisture stress conditions, days to heading (DH) showed positive and significant correlations ( $r=0.73^{**}$ ) with days to maturity (DM), while plant height (PH) showed negative correlations with grains/spike ( $r=-0.38^{**}$ ) (Figure 1). Grain yield (GY) showed positive and significant associations with 1000-grain wt. (TKW:  $r=0.62^{***}$ ) and tillers/m ( $r=0.53^{***}$ ). While under irrigated conditions, GY showed positive and significant associations with DH ( $r=0.26^*$ ), DM ( $r=0.33^{**}$ ), tiller count (0.35\*\*), PH (0.24\*) and TKW (0.63\*\*). Under water stress, NDVI at heading (NDVIH) was found to be positively associated with days to heading, NDVI at grain filling, whereas the canopy temperature at heading (CTH) exhibited a positive relationship with DH, DM and CT at grain filling (CTGF). NDVIH and CTH showed a positive association with NDVIGF and CTGF, respectively, under favourable conditions.

### Drought stress indices correlations

Yield index showed positive and significant correlations with YSI ( $r=0.62^{***}$ ), RDI ( $r=0.62^{***}$ ), DRI ( $0.96^{***}$ ), STI ( $r=0.97^{***}$ ), HM ( $r=0.98^{***}$ ), MP ( $r=0.96^{***}$ ), GMP and MRP (both  $r=0.97^{***}$ ) (Figure 2). Drought susceptibility index (DSi) exhibited a positive and significant association with PYR,



**Fig. 2.** Correlations among 14 drought stress indices

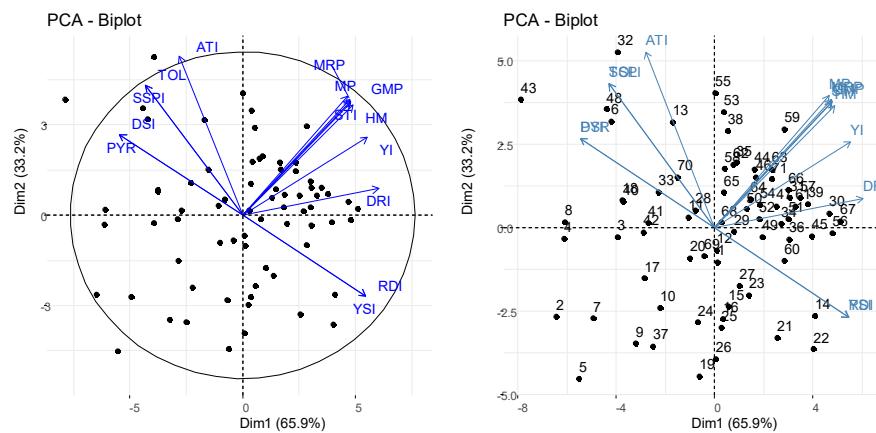
ATI, TOL and SSPI, while it depicted negative correlations with YI, YSI, RDI, DRI, STI, HM, MP, GMP and RMP. Here, the complete associations ( $r=1.00^{***}$ ) were observed between DSi and PYR, TOL and SSPI, YSI and RDI and among HM, MP, GMP and MRP.

### Principal Component analysis (PCA)

Indices-based PCA was carried out to understand the distribution of the genotypes (Fig. 3). The initial two principal components (PCs) captured 65.9% and 33.2% of the total variation. The standard deviation (SD) based PCA biplot revealed that the estimated 14 drought stress indices were grouped into three clusters. The indices, namely PYR, DSi, SSPI, TOL and ATI, were grouped into a single segment. An individual cluster was obtained for HM, GMP, MRP, MP, STI, YI and DRI. The indices RDI and YSI were clustered into a different quadrant. The genotypes G30 (HD3171), G39 (MP1358), G45 (20<sup>th</sup> HTWYT-48), G56 (29<sup>th</sup> SAWYT-316) and G67 (WAP91) were found high yielding and water stress tolerant.

### Discussion

The pooled crop yield showed a reduction of 15.6% under water stress, while days to heading, tiller count and plant height were reduced by 8.1, 15.5, and 12.5%, respectively. The wheat genotypes, WAP84, PBW596, 20<sup>th</sup> HTWYT-48, HD3171, WAP91, 29<sup>th</sup> SAWYT-316, K8027, K9465 and HD2987 were identified with the lowest grain yield reduction (<5.0%). It was observed that the drought susceptible genotypes depicted a significant decrease in tiller count, plant height and 1000 grain wt. Whereas, the tolerant genotypes were found with the moderate reduction for plant height and tiller count, but largely maintained 1000 grain wt. under water stress. Sarwar et al. (2023) reported that the tillering stage is extensively susceptible to drought stress and reduces



**Fig. 3.** PCA biplot for drought stress indices and grouping of genotypes

tiller count significantly. The commonly used parents in the drought stress breeding programs, namely HI1500, NI5439, NIAW34 and PBW175, also showed minimum grain yield reduction (<10%). Whereas, the most popular two genotypes, HINDI62 and C306, exhibited 12.8% and 15.3% yield loss under water stress. However, all of these genotypes were lower yielders ( $\geq 25\%$ ) than the best check, NIAW3170.

Garg et al. (2012) summarized that high proline, lower  $\text{H}_2\text{O}_2$  and increased antioxidant enzymatic activity are some of the key characteristics for drought tolerance in C306 and HD2382. Morphologically, root system architecture, leaf senescence, stay green habit and rapid flowering (early maturity) triggered by earliness *per se* (*Eps*) and photoperiod (*Ppd*) gene(s) are some effective mechanisms to tackle water stress (Farooq et al. 2014). Here, the early flowering by 5-10 days was observed under water stress; however, most of the tolerant genotypes compensated for grain yield by a

prolonged grain filling period. Singh et al. (2012) reported that C306 can alleviate drought effects by curtailing lipid peroxidation. The well-reported biochemical mechanisms of drought tolerance are functional protection of proteins, accumulation of osmolytes, induction of antioxidant defence and hormonal regulations (Fleury et al. 2010; Budak et al. 2013; Nezhadahmadi et al. 2013; Mwadzingeni et al. 2016; Adel and Carels 2023).

In the present study, 14 drought stress indices were calculated to identify drought-tolerant genotypes. Drought susceptibility index (DSI) is most widely used in the stress breeding programs (Fischer and Maurer 1978; Mall et al. 2011; Mohammed et al. 2021; Negisho et al. 2022); however, it does not provide substantial information for yield advantage coupled with stress resilience. For instance, here the genotypes K9465, HD2987 and K8027 were the three top rankers for DSI but showed yield reduction of

**Table 3.** Per cent reduction under water stress and ranking based on drought resistance index (DRI)

Geno Sym.	Genotype	DSI rank	DRI rank	Yield advantage over the best check	Percent reduction under water stress						
					GY	DH	PH	TPM	SL	GPS	TKW
G67	WAP91	5	1	11.9	2.2	7.8	13.4	6.9	8.3	6.7	3.9
G56	29 <sup>th</sup> SAWYT-316	4	2	8.7	2.0	9.3	19.4	11.7	1.9	10.2	0.4
G30	HD 3171	6	3	10.2	3.7	7.2	10.0	15.4	4.3	4.3	2.3
G45	20 <sup>th</sup> HTWYT-48	7	4	3.3	3.9	11.9	14.0	12.6	14.3	0.3	2.1
G39	MP 1358	13	5	5.9	6.6	7.0	13.9	17.0	14.9	1.2	2.4
G57	WAP83	18	7	4.9	7.8	8.5	13.0	10.1	11.1	4.2	1.3
G61	WAP75	17	9	2.7	7.8	12.2	12.2	8.3	8.2	14.4	0.8
G59	WAP92	35	10	8.0	13.8	5.9	12.9	14.8	6.3	11.9	3.3
G31	K 1317	22	11	2.3	9.0	6.2	12.7	19.7	0.0	3.5	0.4
G66	WAP96	24	12	2.9	9.6	8.3	17.3	7.4	8.0	1.9	1.4
G63	WAP115	30	18	1.0	12.7	8.6	19.0	9.8	11.8	28.9	8.1
G71	NIAW3170 (c)	29	19	-	12.0	7.2	4.2	18.6	11.1	8.1	0.4

10.4, 5.5, and 17.4% over the best check NIAW3170 under rainfed conditions. These genotypes also appeared as stress-tolerant in the estimated indices, namely YSI, TOL, RDI, SSPI, PYR and ATI (Supplementary Table 3). In general, the genotypes having minimum yield variations under water stress and irrigated conditions were identified by the above stress indices and presented as drought-tolerant. However, the yield potential of the genotypes cannot be ignored, particularly in the stress breeding programs. The high-yielding potential and drought-tolerant parents are likely to throw more desirable segregants in the filial generations with the favourable gene constellations. Mohammadi et al. (2016) reported that SSI, YSI and TOL showed variable concordance values and were inaccurate under drought conditions. Based on yield and stress resilience, the drought resistance index (DRI) was more successful in identifying tolerant genotypes under moisture stress.

The genotypes viz., WAP91 (G67), 29<sup>th</sup> SAWYT-316 (G56), HD3171 (G30), 20<sup>th</sup> HTWYT-48 (G45) and MP1358 (G39) were the top five rankers based on DRI (Table 3). All these genotypes revealed a grain yield advantage (3.3–11.9%) over the best check NIAW3170, coupled with minimum grain yield reduction under water stress (2.2–6.6%). While comparing DRI and DSI, we found that WAP92 (G59) ranked 10<sup>th</sup> for DRI and 35<sup>th</sup> in DSI, depicting a yield advantage of 8.0% over the best check; however, it showed a 13.8% yield reduction. The other promising genotypes identified based on DRI were, namely, WAP75, WAP83, WAP96 and K1317. The other stress indices, YI, MP, HM, GMP and MRP, also substantiated better yielding stress-tolerant genotypes, but were not found as effective as DRI. The PCA biplots generated for genotypes and stress indices also classified DRI with an acute angle, long vector length and better representativeness.

The exotic lines, 29<sup>th</sup> SAWYT-316 (G56) and 20<sup>th</sup> HTWYT-48 (G45), were studied for genealogy and the water resilience in G45 could be attributed to one of the parents, CROC\_1/ *Ae. squarrosa*. The genotype G56 probably possessed water stress tolerance gene(s) and genomic regions from the climate resilient genotypes, Kachu and Kiritati. Similarly, the released wheat variety HD3171 possibly gained water resilience from HD2879, while MP1358 from Kachu and K1215. The genealogy found is in agreement with the findings of Meena et al. (2025), who reported the common parent Kiritati in the parentage of the water-use-efficient bread wheat genotypes, 40<sup>th</sup> ESWYT-07 and 40<sup>th</sup> ESWYT-37. Among the employed drought tolerance indices, the drought resistance index (DRI) was the most successful in identifying stress-tolerant and high-yielding genotypes. The genotypes, namely HD3171, MP1358, 20<sup>th</sup> HTWYT-48, 29<sup>th</sup> SAWYT-316, K1317 and WAP91 appeared as drought tolerant. The yield attributes 1000 grain weight, grains/spike and tillers count can be targeted as selection under drought stress breeding programs and to train genomic models.

## Supplementary materials

The Supplementary Tables S1 to S3 and Supplementary Fig. 1 are provided, which can be accessed at [www.isgp.org](http://www.isgp.org)

## Authors' contribution

Conceptualization of research (VK, VS, AG); Designing of the experiments (VK, VS, VG, SK); Contribution of experimental materials (VK, AG); Execution of field experiments and data collection (VK, VS, VG); Analysis of data and interpretation (VK, GPS); Preparation of the manuscript (VK, GPS).

## References

Adel S. and Carels N. 2023. Plant tolerance to drought stress with emphasis on wheat. *Plants*, **12**(11): 2170. <https://doi.org/10.3390/plants12112170>.

Allahverdiyev T. I., Rzayev F. H., Gasimov E. K. and Huseynova I. M. 2025. Effect of drought stress on some biochemical parameters and ultrastructure of bread wheat genotypes. *Cereal Res. Commun.*, 1-15. <https://doi.org/10.1007/s42976-025-00681-0>

Arif M., Haroon M., Nawaz A. F., Abbas H., Xu R. and Li L. 2025. Enhancing wheat resilience: biotechnological advances in combating heat stress and environmental challenges. *Plant Mol. Biol.*, **115**(2): 41. <https://doi.org/10.1007/s11103-025-01569-7>

Balla M. Y., Kamal N. M., Tahir I. S. A., Gorafi Y. S. A., Abdalla M. G. A and Tsujimoto H. 2025. Intraspecific variation for heat stress tolerance in wild emmer-derived durum wheat populations. *Front. Plant Sci.*, **16**: 1523562. Doi: 10.3389/fpls.2025.1523562

Bapela T., Shimelis H., Tsilo T. J and Mathew I. 2022. Genetic improvement of wheat for drought tolerance: Progress, challenges and opportunities. *Plants* **11**(10): 1331. <https://doi.org/10.3390/plants11101331>

Budak H., Kantar M. and Yucebilgili K. 2013. Drought tolerance in modern and wild wheat. *Sci. World J.*, **2013**(1): 548246. <https://doi.org/10.1155/2013/548246>

FAOSTAT. 2025. Crop production data from <https://www.fao.org/faostat/en/#data/QCL>. Accessed on 11th July, 2025.

Farooq M., Hussain M. and Siddique K. H. 2014. Drought stress in wheat during flowering and grain-filling periods. *Crit. Rev. Plant Sci.*, **33**(4): 331-349. <https://doi.org/10.1080/07352689.2014.875291>

Fischer R. A. and Maurer R. 1978. Drought resistance in spring wheat cultivars. I. Grain yield responses. *Aust. J. Agril. Res.*, **29**(5): 897-912. <https://doi.org/10.1071/AR9780897>

Fleury D., Jefferies S., Kuchel H. and Langridge P. 2010. Genetic and genomic tools to improve drought tolerance in wheat. *J Exp Bot.*, **61**(12): 3211-3222. <https://doi.org/10.1093/jxb/erq152>

Garg B., Jaiswal J. P., Misra S., Tripathi B. N. and Prasad M. 2012. A comprehensive study on dehydration-induced antioxidative responses during germination of Indian bread wheat (*Triticum aestivum* L. em Thell) cultivars collected from different agro-climatic zones. *Physiol. Mol. Biol. Plants*, **18**: 217-228. <https://doi.org/10.1007/s12298-012-0117-7>

Kadam S., Singh K., Shukla S., Goel S., Vikram P., Pawar V., et al. 2012. Genomic associations for drought tolerance on the short arm of wheat chromosome 4B. *Funct. Integr. Genomics*, **12**: 447-464. <https://doi.org/10.1007/s10142-012-0276-1>

Kumar J., Gunapati S., Kianian S. F. and Singh S. P. 2018. Comparative

analysis of transcriptome in two wheat genotypes with contrasting levels of drought tolerance. *Protoplasma*, **255**: 1487-1504. <https://doi.org/10.1007/s00709-018-1237-x>

Kumar S., Vishwakarma H., Panzade K. P., Jaiswal N. and Aggarwal D. 2025. Stress-associated gene TaTT1 from wheat confers heat stress tolerance in the prokaryotic system and transgenic tobacco (*Nicotiana tabacum* L.). In *In Vitro Cellular Develop. Biol. Plant.*, 1-15. doi: 10.1007/s11627-024-10479-0

Mall A., Swain P., Das S., Singh O. and Kumar A. 2011. Effect of drought on yield and drought susceptibility index for quality characters of promising rice genotypes. *Cereal Res. Commun.*, **39(1)**: 22-31. <https://doi.org/10.1556/crc.39.2011.1.3>

Manoj N. V., Chaudhary H. K., Sharma P., Singh K. and Sood V. K. 2020. Revealing allelic expressivity and distribution of genes for hybrid necrosis in north-west Himalayan diverse wheat (*Triticum aestivum* L.) gene pools. *Cereal Res. Commun.*, **48**: 25-31. <https://doi.org/10.1007/s42976-019-00008-w>

Meena R. P., Kumar V., Venkatesh K., Sujatha H. T., Tripathi S. C., Nargund R. and Nataraj V. 2025. Multi-trait-based identification of water use efficient genotypes in bread wheat (*Triticum aestivum* L.). *Indian J. Genet. Plant Breed.*, **85(3)**: 418-425.

Mohammadi R. 2016. Efficiency of yield-based drought tolerance indices to identify tolerant genotypes in durum wheat. *Euphytica*, **211(1)**: 71-89. <https://doi.org/10.1007/s10681-016-1727-x>

Mohammadi R. 2018. Breeding for increased drought tolerance in wheat: a review. *Crop Pasture Sci.*, **69(3)**: 223-241. <https://doi.org/10.1071/CP17387>

Mohammed S., Huggins T., Mason E., Beecher F., Chick C., Sengodan P., et al. 2021. Mapping the genetic loci regulating leaf epicuticular wax, canopy temperature, and drought susceptibility index in *Triticum aestivum*. *Crop Sci.*, **61(4)**: 2294-2305. <https://doi.org/10.1002/csc.20458>

Mottaleb K. A., Kruseman G., Frija A., Sonder K. and Lopez-Ridaura S. 2023. Projecting wheat demand in China and India for 2030 and 2050: Implications for food security. *Front. Nutr.*, **9**: 1077443. doi: 10.3389/fnut.2022.1077443

Mutanda M., Shimelis H., Chaplot V., Shamuyarira K. W. and Figlan S. 2025. Agronomic performance and water use efficiency of newly developed wheat populations under drought-stressed and non-stressed conditions. *Discov. Appl. Sci.*, **7(3)**: 176. <https://doi.org/10.1007/s42452-025-06605-1>

Mwadzingeni L., Shimelis H., Dube E., Laing M. D. and Tsilo T. J. 2016. Breeding wheat for drought tolerance: Progress and technologies. *J. Integr. Agric.*, **15(5)**: 935-943. [https://doi.org/10.1016/S2095-3119\(15\)61102-9](https://doi.org/10.1016/S2095-3119(15)61102-9)

Negisho K., Shibu S., Matros A., Pillen K., Ordon F. and Wehner G. 2022. Association mapping of drought tolerance indices in Ethiopian durum wheat (*Triticum turgidum* ssp. *durum*). *Front Plant Sci.*, **13**: 838088. <https://doi.org/10.3389/fpls.2022.838088>

Nezhadahmadi A., Prodhan Z. H. and Faruq G. 2013. Drought tolerance in wheat. *Sci. World J.*, **2013(1)**: 610721. <https://doi.org/10.1155/2013/610721>

Nyaupane S., Poudel M. R., Panthi B., Dhakal A., Paudel H. and Bhandari R. 2024. Drought stress effect, tolerance, and management in wheat—a review. *Cogent Food Agric.*, **10(1)**: 2296094. <https://doi.org/10.1080/23311932.2023.2296094>

Pandey A., Masthigowda M. H., Kumar R., Mishra S., Khobra R., Pandey G. C., et al. 2023. Explicating drought tolerance of wheat (*Triticum aestivum* L.) through stress tolerance matrix. *Plant Physiol. Rep.*, **28(1)**: 63-77. <https://doi.org/10.1007/s40502-022-00707-3>

Pequeno D. N., Hernandez-Ochoa I. M., Reynolds M., Sonder K., MoleroMilanA., Robertson R. D. et al. 2021. Climate impact and adaptation to heat and drought stress of regional and global wheat production. *Environ. Res. Lett.*, **16(5)**: 054070. <https://doi.org/10.1088/1748-9326/abd970>

Priya M., Farooq M. and Siddique K. H. 2025. Enhancing Tolerance to Combined Heat and Drought Stress in Cool-Season Grain Legumes: Mechanisms, Genetic Insights and Future Directions. *Plant Cell Environ.*, <https://doi.org/10.1111/pce.15382>

Qian D., Wang M., Niu Y., Yang Y. and Xiang Y. 2025. Sexual reproduction in plants under high temperature and drought stress. *Cell Rep.*, **44(3)**. <https://doi.org/10.1016/j.celrep.2025.115390>

Qiu W., Ma X., Cao H., Huang T., She X., Huang M. et al. 2022. Improving wheat yield by optimizing seeding and fertilizer rates based on precipitation in the summer fallow season in drylands of the Loess Plateau. *Agric. Water Manag.*, **264**: 107489. <https://doi.org/10.1016/j.agwat.2022.107489>

Sairam R. K. and Saxena D. C. 2000. Oxidative stress and antioxidants in wheat genotypes: possible mechanism of water stress tolerance. *J. Agron. Crop Sci.*, **184(1)**: 55-61. <https://doi.org/10.1046/j.1439-037x.2000.00358.x>

Sangha J. S., Wang W., Knox R., Ruan Y., Cuthbert R. D., Isidro-Sanchez J., et al. 2025. Phenotypic plasticity of bread wheat contributes to yield reliability under heat and drought stress. *Plos One*, **20(3)**: e0312122. <https://doi.org/10.1371/journal.pone.0312122>

Sarwar M., Saleem M. F., Ullah N., Khan M. J., Maqsood H., Ahmad H., et al. 2023. Silver nanoparticles protect tillering in drought-stressed wheat by improving leaf water relations and physiological functioning. *Funct. Plant Biol.*, **50(11)**: 901-914. <https://doi.org/10.1071/FP23036>

Singh S., Gupta A. K. and Kaur N. 2012. Differential responses of anti-oxidative defence system to long-term field drought in wheat (*Triticum aestivum* L.) genotypes differing in drought tolerance. *J. Agron. Crop Sci.*, **198(3)**: 185-195. <https://doi.org/10.1111/j.1439-037X.2011.00497.x>

Subrahmanyam D., Subash N., Haris A. and Sikka A. K. 2006. Influence of water stress on leaf photosynthetic characteristics in wheat cultivars differing in their susceptibility to drought. *Photosynthetica*, **44**: 125-129. <https://doi.org/10.1007/s11099-005-0167-y>

Tomar S. M. S. and Kumar G. T. 2004. Seedling survivability as a selection criterion for drought tolerance in wheat. *Plant Breed.*, **123(4)**: 392-394. <https://doi.org/10.1111/j.1439-0523.2004.00993.x>

Vikas V. K., Tomar S. M. S., Sivasamy M., Kumar J., Jayaprakash P., Kumar A., et al. 2013. Hybrid necrosis in wheat: evolutionary significance or potential barrier for gene flow?. *Euphytica*, **194**: 261-275. <https://doi.org/10.1007/s10681-013-0952-9>

Zhao W., Liu L., Shen Q., Yang J., Han X., Tian F. and Wu J. 2020. Effects of water stress on photosynthesis, yield, and water use efficiency in winter wheat. *Water*, **12(8)**: 2127. <https://doi.org/10.3390/w12082127>

**Supplementary Table S1.** Combined analysis of variance for stress and irrigated conditions

Source	df	SS	MSS	F Value	Pr
Environment (Env.)	1	905018	905018	273.90	P<.001
Genotype (Geno)	70	5384351	76919	12.35	P<.001
Rep (Env.)	2	324333	162167	49.07	P<.001
Geno × Env.	70	435857	6227	1.88	P<.001
Pooled Error	140	462583	3304.17	-	-

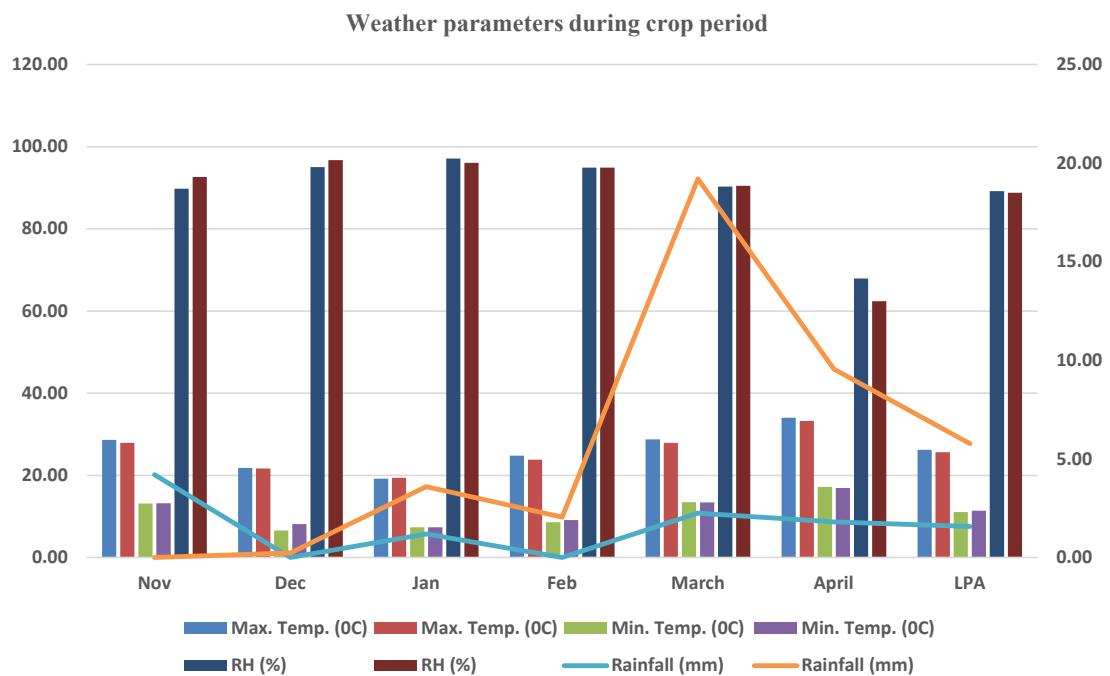
**Supplementary Table 2.** Estimated drought stress indices for bread wheat genotypes

Genotype	Sym.	YI	YSI	TOL	MP	HM	DSI	GMP	DRI	STI	RDI	SSPI	MRP	PYR	ATI
HINDI 62	G1	0.96	0.87	86.35	630.20	627.24	0.82	628.72	511.75	0.76	1.03	5.97	1.89	12.82	480.44
KRL-99	G2	0.43	0.66	137.35	332.70	318.52	2.19	325.53	173.68	0.20	0.78	9.50	0.99	34.22	395.68
WCF8-HT13	G3	0.71	0.71	174.35	523.20	508.67	1.83	515.89	311.48	0.51	0.85	12.06	1.56	28.56	795.97
AKAW3717	G4	0.57	0.63	204.35	447.20	423.86	2.38	435.37	216.69	0.36	0.74	14.13	1.33	37.20	787.33
DWRL1	G5	0.39	0.75	79.35	276.70	271.01	1.61	273.84	177.58	0.14	0.89	5.49	0.83	25.08	192.29
WCF 12-19	G6	0.84	0.65	280.35	653.20	623.12	2.27	637.98	331.74	0.78	0.77	19.39	1.94	35.34	1582.82
WCF 12-208	G7	0.53	0.73	120.35	382.20	372.73	1.74	377.43	234.42	0.27	0.86	8.32	1.14	27.21	401.98
WCF 12-61	G8	0.59	0.62	220.35	469.20	443.33	2.44	456.08	222.48	0.40	0.74	15.24	1.39	38.03	889.36
WCF 12-7	G9	0.60	0.82	81.35	407.70	403.64	1.16	405.67	300.43	0.31	0.97	5.63	1.22	18.14	292.04
EC 531185	G10	0.73	0.83	92.85	490.45	486.06	1.11	488.25	367.23	0.46	0.98	6.42	1.47	17.29	401.18
HTW63	G11	0.94	0.80	140.85	644.45	636.75	1.26	640.59	460.93	0.79	0.95	9.74	1.93	19.70	798.47
RW5	G12	0.98	0.86	95.35	643.70	640.17	0.88	641.93	513.83	0.79	1.02	6.59	1.93	13.79	541.67
HD2781	G13	1.02	0.73	229.85	735.95	718.00	1.73	726.92	453.27	1.01	0.87	15.90	2.19	27.01	1478.61
HD2987	G14	1.17	1.03	-21.65	701.70	701.53	-0.20	701.62	734.85	0.94	1.22	-1.50	2.12	-3.13	-134.42
HI 1500	G15	0.93	0.92	48.35	588.70	587.71	0.51	588.20	519.99	0.66	1.09	3.34	1.77	7.89	251.68
HI 1531	G16	0.89	0.92	44.35	565.70	564.83	0.48	565.27	502.52	0.61	1.10	3.07	1.70	7.54	221.85
UAS 375	G17	0.73	0.78	124.35	506.70	499.07	1.40	502.87	347.36	0.48	0.93	8.60	1.52	21.86	553.38
HUW 234	G18	0.78	0.70	202.35	575.70	557.92	1.92	566.74	332.67	0.61	0.83	13.99	1.71	29.89	1014.87
HUW 468	G19	0.72	0.95	25.35	454.20	453.85	0.35	454.02	417.55	0.39	1.12	1.75	1.37	5.43	101.85
HW 2004	G20	0.89	0.83	109.35	596.20	591.19	1.08	593.69	450.55	0.67	0.99	7.56	1.79	16.80	574.51
K 8027	G21	1.02	1.01	-5.65	619.70	619.69	-0.06	619.69	628.23	0.73	1.20	-0.39	1.87	-0.92	-30.98
K 9465	G22	1.11	1.06	-39.15	655.45	654.87	-0.39	655.16	716.59	0.82	1.26	-2.71	1.99	-6.16	-226.99
NI 5439	G23	1.00	0.94	41.85	632.45	631.76	0.41	632.10	572.36	0.76	1.11	2.89	1.91	6.41	234.10
NIAW 34	G24	0.81	0.89	58.35	524.70	523.08	0.68	523.89	443.32	0.53	1.06	4.04	1.58	10.53	270.52
PBW 175	G25	0.87	0.93	40.35	551.70	550.96	0.45	551.33	494.02	0.58	1.10	2.79	1.66	7.06	196.87
PBW 596	G26	0.80	0.95	23.85	502.45	502.17	0.30	502.31	467.78	0.48	1.13	1.65	1.52	4.64	106.02
PBW 660	G27	0.99	0.92	54.35	631.20	630.03	0.53	630.61	554.16	0.76	1.09	3.76	1.90	8.26	303.31
WH 1080	G28	0.97	0.81	142.35	662.20	654.55	1.24	658.36	476.31	0.83	0.96	9.84	1.98	19.41	829.36
WH 1142	G29	1.05	0.87	95.35	690.70	687.41	0.83	689.05	559.99	0.91	1.03	6.59	2.08	12.91	581.43
HD 3171	G30	1.36	0.96	32.35	847.20	846.89	0.24	847.05	799.89	1.37	1.14	2.24	2.56	3.75	242.49
K 1317	G31	1.26	0.91	76.35	809.20	807.40	0.58	808.30	701.55	1.25	1.08	5.28	2.44	9.01	546.14
HI 1612	G32	0.94	0.63	336.35	741.20	703.04	2.37	721.87	361.08	1.00	0.75	23.26	2.20	36.99	2148.68
HD 3271	G33	0.89	0.75	184.35	634.20	620.80	1.63	627.47	404.46	0.75	0.89	12.75	1.89	25.38	1023.66
HI 1621	G34	1.20	0.92	65.35	766.70	765.31	0.52	766.00	674.02	1.12	1.09	4.52	2.31	8.18	442.99
HI 1628	G35	1.15	0.83	145.35	775.70	768.89	1.10	772.29	582.58	1.14	0.98	10.05	2.33	17.13	993.38
HD 3293	G36	1.21	0.94	47.35	759.70	758.96	0.39	759.33	691.54	1.10	1.11	3.27	2.29	6.04	318.18

GW 513	G37	0.64	0.85	70.35	428.20	425.31	0.97	426.75	333.36	0.35	1.01	4.87	1.29	15.18	265.68
HD 3298	G38	1.17	0.80	177.35	799.70	789.87	1.28	794.77	569.08	1.21	0.95	12.26	2.39	19.96	1247.36
MP 1358	G39	1.31	0.93	56.85	826.95	825.97	0.43	826.46	745.45	1.31	1.11	3.93	2.49	6.65	415.79
HI 1633	G40	0.78	0.70	200.85	573.20	555.61	1.91	564.33	331.81	0.61	0.83	13.89	1.71	29.82	1003.07
HI 1634	G41	0.82	0.75	165.85	581.70	569.88	1.60	575.76	374.31	0.63	0.89	11.47	1.74	24.95	845.04
HI 1636	G42	0.79	0.75	160.85	564.20	552.74	1.60	558.44	363.06	0.60	0.89	11.12	1.68	24.95	794.91
20 <sup>th</sup> HTWYT-43	G43	0.62	0.50	369.85	561.70	500.82	3.18	530.39	190.13	0.54	0.60	25.58	1.65	49.54	1735.96
20 <sup>th</sup> HTWYT-2	G44	1.20	0.85	124.85	794.20	789.29	0.93	791.74	625.12	1.20	1.01	8.63	2.38	14.57	874.77
20 <sup>th</sup> HTWYT-48	G45	1.28	0.96	31.85	794.70	794.38	0.25	794.54	748.18	1.21	1.14	2.20	2.40	3.93	223.95
20 <sup>th</sup> HTWYT-25	G46	1.19	0.86	118.85	786.20	781.71	0.90	783.95	624.63	1.18	1.02	8.22	2.36	14.05	824.54
20 <sup>th</sup> HTWYT-41	G47	1.21	0.90	80.85	781.20	779.11	0.63	780.15	667.88	1.16	1.07	5.59	2.35	9.84	558.19
20 <sup>th</sup> HTWYT-13	G48	0.84	0.64	294.85	662.20	629.38	2.33	645.58	327.30	0.80	0.75	20.39	1.96	36.42	1684.51
20 <sup>th</sup> HTWYT-32	G49	1.13	0.91	70.85	724.20	722.47	0.60	723.33	624.53	1.00	1.08	4.90	2.18	9.33	453.52
20 <sup>th</sup> HTWYT-42	G50	1.12	0.87	102.85	735.20	731.60	0.84	733.40	594.37	1.03	1.03	7.11	2.21	13.07	667.52
29 <sup>th</sup> SAWYT-334	G51	1.23	0.92	61.85	782.70	781.48	0.49	782.09	694.63	1.17	1.10	4.28	2.36	7.60	428.07
29 <sup>th</sup> SAWYT-320	G52	1.14	0.89	85.85	740.70	738.21	0.70	739.46	621.33	1.05	1.06	5.94	2.23	10.96	561.79
29 <sup>th</sup> SAWYT-303	G53	1.18	0.79	195.10	817.20	805.56	1.37	811.36	566.16	1.26	0.93	13.49	2.44	21.33	1400.85
29 <sup>th</sup> SAWYT-341	G54	1.17	0.88	95.10	759.20	756.22	0.76	757.71	627.76	1.10	1.05	6.58	2.28	11.79	637.68
29 <sup>th</sup> SAWYT-310	G55	1.17	0.77	218.10	825.70	811.30	1.50	818.47	549.44	1.28	0.91	15.08	2.47	23.33	1579.71
29 <sup>th</sup> SAWYT-316	G56	1.34	0.98	17.10	828.20	828.11	0.13	828.16	802.90	1.31	1.16	1.18	2.50	2.04	125.32
WAP83	G57	1.30	0.92	67.10	824.20	822.83	0.50	823.52	728.80	1.30	1.09	4.64	2.48	7.82	489.01
WAP71	G58	1.11	0.82	150.10	752.70	745.22	1.16	748.95	554.77	1.07	0.97	10.38	2.26	18.13	994.84
WAP92	G59	1.34	0.86	130.10	879.70	874.89	0.88	877.29	702.47	1.47	1.02	9.00	2.64	13.77	1010.05
WAP84	G60	1.16	0.95	37.10	726.20	725.73	0.32	725.96	672.40	1.01	1.13	2.57	2.19	4.98	238.35
WAP75	G61	1.27	0.92	65.10	807.20	805.89	0.50	806.54	714.60	1.24	1.09	4.50	2.43	7.75	464.65
WAP114	G62	1.14	0.83	147.10	768.20	761.16	1.12	764.67	573.26	1.12	0.98	10.17	2.30	17.48	995.43
WAP115	G63	1.25	0.87	111.10	817.20	813.42	0.82	815.31	664.69	1.27	1.04	7.68	2.46	12.73	801.60
WAP48	G64	1.15	0.87	107.10	753.20	749.39	0.85	751.29	606.77	1.08	1.03	7.41	2.26	13.28	712.07
WAP66	G65	1.08	0.83	133.10	723.20	717.08	1.08	720.13	545.98	0.99	0.99	9.20	2.17	16.85	848.23
WAP96	G66	1.27	0.90	82.10	816.70	814.64	0.61	815.67	701.41	1.27	1.07	5.68	2.46	9.57	592.62
WAP91	G67	1.38	0.98	19.10	853.20	853.09	0.14	853.15	824.97	1.39	1.16	1.32	2.58	2.21	144.20
Checks															
C306	G68	1.03	0.85	113.10	684.05	679.38	0.98	681.71	531.67	0.89	1.01	7.82	2.05	15.27	682.31
DBW110	G69	0.93	0.85	101.00	619.30	615.18	0.97	617.24	483.03	0.73	1.01	6.98	1.86	15.08	551.69
DBW296	G70	0.96	0.76	181.40	677.90	665.76	1.51	671.80	448.61	0.86	0.91	12.54	2.03	23.60	1078.46
NIAW3170	G71	1.24	0.88	103.90	805.55	802.20	0.78	803.87	662.29	1.24	1.04	7.19	2.42	12.12	739.14

Supplementary Table 3: Drought tolerant genotypes identified based on different stress indices

YI	HD3171, MP1358, 29 <sup>th</sup> SAWYT-316, WAP92, WAP91
YSI	K9465, HD2987, K8027, 29 <sup>th</sup> SAWYT-316, WAP91
TOL	K9465, HD2987, K8027, 29 <sup>th</sup> SAWYT-316, WAP91
MP	WAP92, WAP91, 29 <sup>th</sup> SAWYT-316, MP1358, HD3171
HM	WAP92, WAP91, 29 <sup>th</sup> SAWYT-316, MP1358, HD3171
DSI	WAP91, 29 <sup>th</sup> SAWYT-316, K9465, K8027, HD2987
GMP	WAP92, WAP91, 29 <sup>th</sup> SAWYT-316, MP1358, HD3171
DRI	WAP91, 29 <sup>th</sup> SAWYT-316, HD3171, 20 <sup>th</sup> HTWYT-48, MP1358
STI	WAP92, WAP96, 29 <sup>th</sup> SAWYT-316, HD3171, MP1358
RDI	K9465, HD2987, K8027, 29 <sup>th</sup> SAWYT-316, WAP91
SSPI	K9465, HD2987, K8027, 29 <sup>th</sup> SAWYT-316, WAP91
MRP	WAP92, WAP91, 29 <sup>th</sup> SAWYT-316, MP1358, HD3171
PYR	K9465, HD2987, K8027, 29 <sup>th</sup> SAWYT-316, WAP91
ATI	K9465, HD2987, K8027, HUW468, PBW596
PCAbiplot based	WAP91, 29 <sup>th</sup> SAWYT-316, HD3171, 20 <sup>th</sup> HTWYT-48, MP1358

**Supplementary Fig. 1.** Weather parameters recorded during crop period