Phenotypic screening of 32 West African Sorghum Genotypes for Drought Tolerance

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Abstract

This study evaluated the agro-physiological responses of 32 sorghum genotypes subjected to water stress applied at the critical stage of transition to panicle initiation. The objective was to identify sorghum adaptation mechanisms under such stress and to determine potentially tolerant genotypes. Analysis of physiological and agro-morphological traits revealed significant inter-genotypic variability. Under stress, leaf water potential decreased from -0.59 to -4.84 MPa, indicating differentiated tolerance levels. Genotypes V1, V2, V12, V16, V22, and V28 maintained good water status and exhibited the lowest rates of leaf desiccation after stress. Stress also induced reductions in stomatal conductance (-25%), photosynthesis (-13%), and transpiration (-40%), reflecting adaptive strategies in the genotypes. However, a marked decrease in grain yield (-47%) was observed, underscoring the limits of adaptive mechanisms to sustain productivity. Combined analysis of tolerance indices (SSI, STI) and agro-physiological traits identified genotype V26 as elite, combining low stress sensitivity with high yield. Other tolerant but less productive genotypes (V2, V6, V10, V11, V12, V14, V16, V18, V24, V30, V32) may serve as gene reservoirs for breeding improvement. In addition, principal component analysis distinguished three groups of genotypes according to their adaptive profiles. These findings highlight the relevance of an integrated approach combining agro-physiological traits, tolerance indices (SSI, STI), and multivariate analyses for the selection of genotypes adapted to water stress conditions.

Keywords: Sorghum, water stress, physiological adaptation, yield, tolerant genotypes.

1. Introduction

Sorghum (Sorghum bicolor [L.] Moench) is the fifth most cultivated cereal worldwide after wheat, maize, rice, and barley (FAO, 2015). It is also the most important crop in the semi-arid Sudanian-Sahelian zones of Africa and Asia (Mindaye et al., 2016). It is primarily grown for both its grain and biomass, which are used for human and animal consumption. Despite the multiple benefits of this crop, its production remains threatened by various biotic and abiotic constraints. The effects of these constraints are increasingly exacerbated by climate change, mainly through the increased frequency of extreme events such as droughts and heavy rainfall (Deng et al., 2024; Qiu et al., 2021). Furthermore, several authors (Ben Mariem et al., 2021; Pickson et al., 2023) have reported that climate change, through rising temperatures, negatively impacts agricultural production, particularly cereals. Sorghum plants employ different morphological and physiological adaptation strategies to survive intermittent or continuous drought occurring during their growth cycle. One key adaptation strategy is the reduction of water loss through transpiration when atmospheric demand increases (Raymundo et al., 2024). Several studies have previously reported mechanisms underlying this trait and its positive impact on sorghum yield (Choudhary et al., 2013; Sinclair et al., 2017; Mwamahonje et al., 2021). These investigations summarize the expression of adaptation mechanisms through stomatal closure, development of a deep root system, and reduction of leaf area under water stress. Consequently, drought tolerance in sorghum is reinforced by multiple genes, each contributing partially to this tolerance (Phuong et al., 2019). Genetic improvement of sorghum for drought tolerance thus relies on integrating numerous traits, such as the "staygreen" trait, leaf rolling, and reduced transpiration. However, these traits are negatively correlated with sorghum yield under water-stressed conditions (Mwamahonje et al., 2021). Moreover, sorghum's response to water stress depends on the species, growth stage, type, intensity, and duration of the stress (Dos Santos et al., 2022; Gano et al., 2021a). Several recent studies have shown that exposure of sorghum to water stress shortly after germination leads to significant reductions in seedling growth, leaf emergence, transpiration, and photosynthesis, often accompanied by early leaf senescence (D. Chen et al., 2015; Zhang et al., 2024). Additionally, Gano et al. (2021) found that early water stress, around 30 days after germination, causes a significant reduction in vegetative growth, decreased photosynthesis, early leaf senescence, and lower grain yield. Other authors (de Souza et al., 2021; Kamal et al., 2018; Sanjari et al., 2021; Tovignan et al., 2016) reported that post-flowering drought negatively affects sorghum, mainly reducing grain yield, biomass, and stem sugar accumulation. However, limited data exist on the impact of early drought occurring during the

transition stage toward panicle initiation and structuring. This stage, which corresponds to the differentiation of the growth point and marks the beginning of panicle development, is particularly sensitive to water stress (Abreha et al., 2022; Tovignan et al., 2016). A better understanding of drought effects at this critical stage is essential, as it could compromise the formation of reproductive organs, leading to reduced panicle weight, fewer fertile flowers, and consequently lower final yield (Ndlovu et al., 2024; Tovignan et al., 2016). Identifying sorghum varieties tolerant to water stress at this specific stage would help guide breeding programs toward targeted genetic improvement, thereby enhancing crop resilience to climatic hazards in the Sahelian and Sudanian-Sahelian regions. The objective of this study was to evaluate the effect of water deficit occurring during the transition stage toward panicle initiation and structuring on sorghum, in order to determine the main adaptation mechanisms and the useful methods and criteria for agro-physiological phenotyping under water stress conditions. It also aimed to select, using drought tolerance indices, the sorghum genotypes that remain productive under drought conditions.

2. Materials andmethods

2.1. Location of the study

The present study was conducted at the experimental site of the National University of Agriculture (UNA) located in Kétou, in the Plateau department, southeastern Benin. The site is situated at the following geographic coordinates: latitude 7°18′26″ N (7.3072) and longitude 2°36′28″ E (2.6077). The region has a tropical climate with a bimodal rainfall pattern, characterized by two main climatic variants: that of the Middle Zou and the southeastern plateaus. The climate calendar includes a long rainy season from March to July, followed by a short dry season in August. A second, shorter rainy season occurs from September to October, preceding a long dry season from November to February. The average annual rainfall in the commune is estimated at approximately 1073 mm, distributed over 65 rainy days (INSAE, 2016). Furthermore, the experimental site's soil exhibited the physicochemical characteristics listed in Table 1.

Table 1:Physicochemical characteristics of the experimental soil

Parameters	Unit	Values	Standards
MO	(%)	0.59	2-3
NT		0.05	0.1-0.15
Ca2+	Cmol/kg	1.5	2.3-3.5

Mg2+		0.5	1-1.5
K+		0.2	0.2-0.4
Na+		0.1	0.3-0.7
CEC		5.2	10-25
pH	-	5.5	6.5-7.5
Pass	mg/kg	6.428	10-15

MO: Organic matter; NT: Total nitrogen; Ca²⁺: Exchangeable calcium; Mg²⁺: Exchangeable magnesium; K*: Exchangeable potassium; Na*: Exchangeable sodium; CEC: Cation exchange capacity; pH: Hydrogen potential (soil acidity); P ass: Available phosphorus.

2.2. Plant material.

The plant material consisted of thirty-two (32) sorghum genotypes obtained from four agricultural research institutions in West Africa. Table 2 presents the characteristics of the genotypes and their origin.

Table 2: Characteristics and origin of the evaluated genotypes

Co			Country	
de			of	
s	Genotypes	Pedigree	Origin	Home Institutions
		(POPD08-611/02-SB-		
	ICSB	F5DT-12B)-7-4-3-1-10-6-		
V1	176008	6-10	Mali	ICRISAT
		(POPD08-622/02-SB-		
	ICSB	F5DT-12B)-1-3-1-3-6-7-		
V2	176003	7-3	Mali	ICRISAT
		(POPD08-622/02-SB-		
	ICSB	F5DT-12B)-1-3-1-3-6-7-		
V3	176005	7-3	Mali	ICRISAT
		(POPD08-611/02-SB-		
	ICSB	F5DT-12B)-3-1-7-2-8-7-		
V4	176006	1-7	Mali	ICRISAT
	ICSB	(POPD08-611/PR3009B)-		
V5	176002	7-3-1-4-1-3-21-3	Mali	ICRISAT
		(POPD08-611/02-SB-		
	ICSB	F5DT-12B)-11-1-5-2-5-4-		
V6	176016	14-3	Mali	ICRISAT
V7	12B	B line	Mali	ICRISAT
		(POPD08-611/02-SB-		
	ICSB	F5DT-12B)-11-5-2-9-8-4-		
V8	176031	9-7	Mali	ICRISAT
	ICSB	(POPD08-611/PR3009B)-		
V9	176001	7-3-1-1-6-3-19-8	Mali	ICRISAT

V1		D.F.	NT: :	Institute for Agricultural
0	45	R line	Nigeria	Research (IAR)
V1 1	ISS 455	R line	Mali	ICRISAT
		[GPN01 S01-267-9-3-1-4-		
		sibvr//(Sambalma(4)/GPN		
V1	ICSV	01 S01 267-9-3-1-7)]-5-2-		
2	1360964	1-1	Mali	ICRISAT
V1		SS07(MadouM)Ban-13-v-		
3		5-Balla Berthe-v	Mali	ICRISAT
V1	SAMSORG	R line		Institute for Agricultural
4	3		Nigeria	Research (IAR)
V1		R line	6	Institut d'Economie Rurale
5			6 Mali	(IER)
V1	Niobougoum	R line	6	Institut d'Economie Rurale
6	a		6 Mali	(IER)
V1	015-SB-CS-	R line	6	Institut d'Economie Rurale
7	F7-127		6 Mali	(IER)
V1			6	Institut d'Economie Rurale
- 8	Seguifa	MALISOR 92-1	6 Mali	(IER)
V1		R line	6	Institut d'Economie Rurale
9	Diamadjigui		6 7 Iali	(IER)
V2		R line	Burkina	Institut de l'Environnement et de
0	SARIASO14		Faso	Recherches Agricoles (INERA)
	019-SB-CS-	R line		
V2	AVANCE-		6	Institut d'Economie Rurale
1	22		6 Mali	(IER)
V2	Tiandougou		6	Institut d'Economie Rurale
2	Coura	04-SB-F5DT-105	6 7 Iali	(IER)
V2		R line	Burkina	Institut de l'Environnement et de
3	Sariaso 16		Faso	Recherches Agricoles (INERA)
V2		R line		
4	Lata 3		Mali	ICRISAT
		(Tiandougou Coura/015-		
V2	ICSV	CS-SB-BC1F1-15)-B-		
5	206056	SS1-7-2-2	Mali	ICRISAT
V2		[(SPV 35 x E35-1) x CS		
6	ICSV 111	3541]	Mali	ICRISAT
V2				
7	ISS 3187	R line	Mali	ICRISAT
V2		GR/(GR/SC566-14)		Institut d'Economie Rurale
8	BC36-080	BC1F3:5-BC36-080	Mali	(IER)
V2	ICSV	(Narichita/PI 639719 02		
9	206084	SD)-B-SS1-SS1-3-1-3	Mali	ICRISAT
V3	SAMSORG	l		Institute for Agricultural
0	49	R line	Nigeria	Research (IAR)
V	Soubatimi	Check	Mali	
31				ICRISAT
V3	Jakumbè	Check	Mali	ICRISAT

2 | |

2.3. Methodology

2.3.1. Experimental Design

The trial was conducted using a randomized complete block design (RCBD) with a two-factor factorial arrangement: water regime and genotype. The experiment was carried out in two distinct environments, namely a control (well-watered) environment and a waterstressed environment, each comprising three replicate blocks. The 32 sorghum genotypes were sown in November to avoid the water stress period coinciding with rainfall. Each genotype was sown on an experimental plot 3.6 m long, consisting of 10 hills spaced 0.4 m apart. The plots consisted of a single row with an inter-row spacing of 0.8 m. Genotypes were randomly assigned within each block and replicate. The two water treatments (normal irrigation and water stress) were separated by a 10 m buffer to prevent accidental irrigation of stressed plots. For the control treatment, plants were irrigated twice a week with 25 mm per irrigation until physiological maturity. In contrast, for the water-stressed treatment, irrigation was withheld for one month starting from 45 days after sowing (DAS). After this stress period, optimal irrigation was resumed until physiological maturity, following Gano et al. (2021). In total, 350 mm of water were applied to plants under water stress, compared to 550 mm for well-watered plants. Field management was limited to fertilization and weeding. Fertilization followed the recommendations of the National Agricultural Research Institute of Benin (INRAB), with 100 kg/ha of NPK applied 14 days after sowing, followed by 50 kg/ha of urea applied at 45 DAS. Weeding was carried out as needed.

2.3.2. Collected Data

Meteorological Data

The automatic weather station at CRA-PP Pobè, located near the experimental site, was used to continuously record the climatic variables necessary for characterizing water stress. Measured variables included air temperature (Temp, in °C), relative humidity (RH, in %), solar radiation (Rad, in W·m⁻²), rainfall (Pluv, in mm), and wind speed (Vent, in m·s⁻¹).

Soil Moisture Monitoring

Soil moisture was monitored using two complementary tools: piezometers and a multifunction digital soil tester (Sonkir MS02). In each experimental block, three piezometers were installed at strategic locations to ensure representative measurements of soil moisture conditions. Each piezometer was installed vertically to a depth of 1.20 m. These devices

allowed tracking of soil moisture changes at different times during the experiment, particularly during the water stress period. Measurements were taken daily to assess deep soil moisture and evaluate the effects of water deficit on the soil moisture profile.

Additionally, the multifunction digital soil tester was used for spot measurements of surface soil moisture (topsoil layer), temperature, and pH. This device features a backlit LCD screen with digital readout, allowing simultaneous display of the three parameters.

Morphological Parameters

Morphological traits of the plants were measured weekly after the induction of water stress on five tagged plants per plot. Evaluated parameters included the number of leaves emerged (NFA), collar diameter (DAC, in cm), and plant height (HP, in cm). The number of dried leaves (NFD) was recorded at the end of the stress period. Specific leaf area (SLA, in cm²·g⁻¹) was determined from the last ligulated leaf by dividing its leaf area by its dry biomass.

Physiological parameters

Physiological parameters were measured at the end of the water stress period, always on the last ligulated leaf, using the portable photosynthesis system ADC BioScientific LCpro-SD. Measurements included: photosynthetic capacity (**A**, in µmol·m⁻²·s⁻¹), leaf temperature (Tleaf, in °C), transpiration (**E**, in mmol·m⁻²·s⁻¹), and stomatal conductance (**gs**, in mol·m⁻²·s⁻¹). Additionally, leaf water potential (\mathbf{Yf}, in MPa) was measured for each genotype and irrigation regime using a pressure chamber (model 3005F01) between 2:00 and 6:00 a.m. Values were recorded in bars and then converted to megapascals (1 bar = 0.1 MPa). The classification of Kramer & Boyer (1995) and Blum (2010) (Table 3) was used to assess levels of water stress according to the leaf water potential (\mathbf{Yf}) values of the different genotypes under each irrigation regime.

Tableau 3: Classification of water stress levels based on leaf water potential (Ψf):

Value of Ψf (MPa)	Drought stress level
−0.1 to −0.5	No stress / well-watered
-0.5 to -1.0	Mild to moderate stress
-1.0 to -1.5	Moderate to severe stress
< -1.5 to -2.5	Severe stress
< -2.5	Very severe / critical stress

Phenological Parameters

Phenological observations were conducted on each genotype to assess the impact of water stress on the duration of developmental phases. Collected data included: the date of first flowering (DAPF, in days after sowing), the date of 50% flowering (D50%F, in days after sowing), the date of 100% flowering (D100%F, in days after sowing), and the date of physiological maturity (DM, in days after sowing).

Yield Parameters

Yield components were measured after panicle drying. The following variables were recorded: - Panicle length (LP, in cm), - Panicle width (IP, in cm), - Panicle weight (PP, in g), - Grain yield per hectare (RDT, in t/ha).

Calculation of Drought Tolerance and Sensitivity Indices

a) Recovery Index after Stress (IDR):

The Recovery Index (IDR) was used to assess the physiological capacity of genotypes to recover after water stress, according to Strauss et al. (2006) and Oukarroum et al. (2007). Formula: IDR = log A + 2 log B, where A is the stressed/control ratio at the end of stress, and **B** is the same ratio two weeks after rewatering.

b) Stress Intensity Index (SI):

 $SI = 1 - (RDT_str / RDT_etm)$, with $RDT_str = \frac{44}{mean}$ yield under stress conditions and RDT_etm = mean yield under normal conditions. The closer SI is to 1, the more severe the

c) Stress Susceptibility Index (SSI):

 $SSI = [1 - (RDT_str / RDT_etm)] / SI$, where RDT_str is the yield of each genotype under water stress, RDT_etm is the yield under normal conditions, and SI is the stress intensity. Genotypes with SSI < 1 are considered tolerant, while those with $SSI \geq 1$ are considered

d) Stress Tolerance Index (STI):

 $STI = (RDT_str \times RDT_etm) / (RDT_m)^2, \text{ where } RDT_str = \text{ yield of genotype } x \text{ under water stress, } RDT_etm = \text{ yield under normal conditions, and } RDT_m = \text{ mean yield of the } x \text{ where } RDT_m = \text{ mean yield of the } x \text{ where } RDT_m = \text{ mean yield of the } x \text{ where } RDT_m = \text{ mean yield of the } x \text{ where } RDT_m = \text{ mean yield of the } x \text{ where } x \text{ where } RDT_m = \text{ mean yield of the } x \text{ where } RDT_m = \text{ mean yield of the } x \text{ where } RDT_m = \text{ mean yield of the } x \text{ where } RDT_m = \text{ mean yield of the } x \text{ where } RDT_m = \text{ mean yield of the } x \text{ where } RDT_m = \text{ mean yield of the } x \text{ where } RDT_m = \text{ mean yield of the } x \text{ where } RDT_m = \text{ mean yield of the } x \text{ where } RDT_m = \text{ mean yield of the } x \text{ where } RDT_m = \text{ mean yield of the } x \text{ where } RDT_m = \text{ mean yield of the } x \text{ where } RDT_m = \text{ mean yield of the } x \text{ where } RDT_m = \text{ mean yield of the } x \text{ where } RDT_m = \text{ mean yield of the } x \text{ where } RDT_m = \text{ mean yield of the } x \text{ where } RDT_m = \text{ mean yield of the } x \text{ where } RDT_m = \text{ mean yield of the } x \text{ where } RDT_m = \text{ mean yield of the } x \text{ where } RDT_m = \text{ mean yield of the } x \text{ where } RDT_m = \text{ mean yield of the } x \text{ where } RDT_m = \text{ mean yield of the } x \text{ where } RDT_m = \text{ mean yield of the } x \text{ where } RDT_m = \text{ mean yield of the } x \text{ where } RDT_m = \text{ mean yield of the } x \text{ where } RDT_m = \text{ mean yield of the } x \text{ where } RDT_m = \text{ mean yield of the } x \text{ where } RDT_m = \text{ mean yield of the } x \text{ where } RDT_m = \text{ mean yield of the } x \text{ where } RDT_m = \text{ mean yield of the } x \text{ where } RDT_m = \text{ mean yield of the } x \text{ where } RDT_m = \text{ mean yield of the } x \text{ where } RDT_m = \text{ mean yield of the } x \text{ where } RDT_m = \text{ mean yield of the } x \text{ where } RDT_m = \text{ mean yield of the } x \text{ where } RDT_m = \text{ mean yield of the } x \text{ where } RDT_m = \text{ mean yield of the } x \text{ where } RDT_m = \text{ mean yield of the } x \text{ where } RDT_m = \text{ mean yield o$ trial under normal conditions. Higher STI values indicate greater tolerance and productivity.

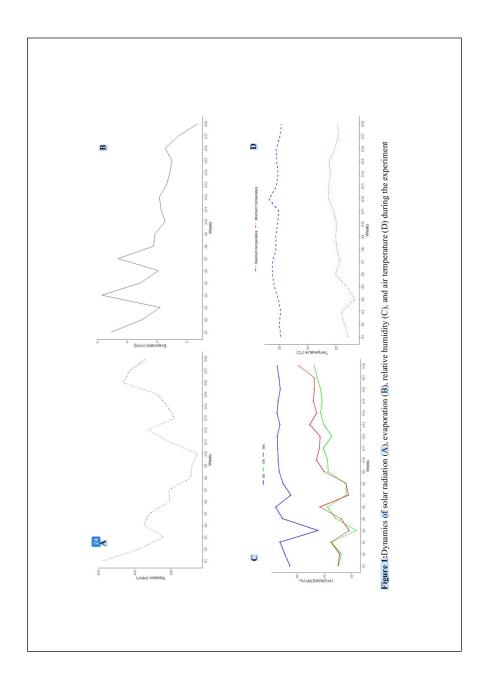
2.3.3. Statistical Analysis

An analysis of variance (ANOVA) was performed for each measured parameter to test statistical differences among genotypes within the same water regime, between different water regimes, and for their interactions. In addition, a combined ANOVA was conducted following the method recommended by McIntosh (1983) to assess genotype × water regime interactions across the data from both environments. The homogeneity of residual variances was verified using Bartlett's test (1937). When the data did not meet the assumptions of homogeneity or normality, a non-parametric analysis was carried out using the Kruskal-Wallis test to detect significant differences between treatments. Treatment means were compared at a probability threshold of 5% (p < 0.05). Furthermore, a principal component analysis (PCA) was conducted to explore multivariate relationships among the measured traits. This analysis was complemented by hierarchical clustering (HCA) to group genotypes according to their response to water stress. The R packages used for these multivariate analyses were FactoMineR and factoextra. Finally, a correlation analysis was performed to evaluate the relationships between morphological, physiological, phenological, and yield traits, in order to better understand the mechanisms of drought tolerance. All statistical analyses were carried out using R software (version 4.x).

3. Results

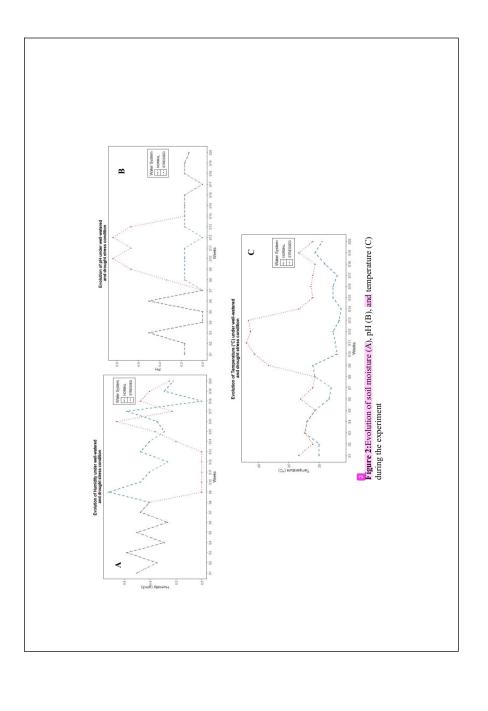
3.1. Evolution of weather conditions during the experiment.

The evolution of climatic conditions during the experiment is presented in Figure 1. Solar radiation (Figure 1A) shows a daily variation ranging from 10 W/m²/day to about 430 W/m²/day. The highest increases in radiation were recorded at the beginning of the experiment, followed by a gradual decline reaching minimum values around the 10th week, before progressively rising again towards the end. Evaporation (Figure 1B) follows a similar trend, with values ranging from 2 mm/day to about 0.5 mm/day. The highest evaporation rates were observed between the 1st and 10th weeks of the experiment, before gradually decreasing towards the last weeks. Relative humidity (Figure 1C) fluctuated between 60% and 80% at 8 a.m., between 20% and 40% at 1 p.m., and between 40% and 60% at 6 p.m. Lower values, dropping to around 30%, were recorded between the 1st and 10th weeks of the experiment, before progressively increasing towards the end. Air temperature (Figure 1D) oscillated around 35 °C for the maximum and around 15–20 °C for the minimum.



.3.2. Dynamics of soil moisture, pH, and temperature during the experiment

The evolution of the edaphic parameters measured during the experiment is presented in the figure. Soil moisture, measured through piezometers placed at a depth of 1.20 m (Figure 2A), shows variations between 0 and 0.5 cm, indicating the presence or absence of moisture at this depth. The analysis of this figure reveals that in the control plot (nonstressed), water levels ranging from 0.2 to 0.5 cm were recorded throughout the entire experimental period, highlighting a permanent presence of soil moisture. In contrast, in the stressed plot, values of 0 cm were recorded between the 9th and 13th week, indicating the absence of soil moisture. This period corresponds to the stress treatment (cessation of irrigation). Soil pH (Figure 2B) ranged between 6 and 6.3 in the control plot and between 6 and 6.8 in the stressed plot during the experimental period. Soil temperature (Figure 2C) varied between 15 °C and 34 °C in both plots, except for an increase recorded between the 7th and 14th week in the stressed plot.



3.4. Response of genotypes' leaf water potential under irrigation regimes. Table 4: Response of leaf water potential of genotypes under different irrigation regimes

Genotypes		ds		ww	ΔWS
	Ψf (MPa)	Stress level	Ψf (MPa)	Stress level	ΔWS
V1	0.93	Moderate	0.16	No stress	5
V2	1.18	Moderate	0.2	No stress	4.89
V3	1.28	Severe	0.1	No stress	11.78
V4	0.91	Moderate	0.18	No stress	4.13
V5	1.01	Severe	0.17	No stress	5.07
V6	1.09	Severe	0.13	No stress	7.17
V7	1.53	Severe	0.11	No stress	12.8
V8	0.99	Moderate	0.11	No stress	7.9
V9	1.54	Severe	0.13	No stress	10.5
V10	1.28	Severe	0.1	No stress	11.78
V11	0.89	Moderate	0.11	No stress	7
V12	0.92	Moderate	0.1	No stress	8.22
V13	1.16	Severe	0.13	No stress	7.67
V14	1.79	Severe	0.19	No stress	8.47
V15	1.08	Severe	0.21	No stress	4.11
V16	0.72	Moderate	0.13	No stress	4.42
V17	1.03	Severe	0.13	No stress	6.75
V18	1.76	Severe	0.1	No stress	16.50
V19	1.78	Severe	0.13	No stress	12.33
V20	1.41	Severe	0.14	No stress	8.77
V21	1.3	Severe	0.1	No stress	12
V22	0.98	Moderate	0.1	No stress	8.78
V23	4.84	Very Severe	0.13	No stress	35.33
V24	1.14	Severe	0.17	No stress	5.87
V25	1.4	Severe	0.12	No stress	10.4
V26	1.88	Severe	0.1	No stress	17.78
V27	3.53	Very Severe	0.12	No stress	27.9
V28	0.68	Moderate	0.1	No stress	5.78
V29	0.59	Moderate	0.13	No stress	3.42
V30	2.67	Very Severe	0.13	No stress	19
V31	0.94	Moderate	0.14	No stress	5.54
V32	1.61	Severe	0.1	No stress	15.1
Overall Main	1.4325	Severe	0.13125	No stress	
Genotype	ns	-	ns	-	-
Е	***	-		-	-

Yf: Leaf water potential; **ds**: Water-stress regime; **ww**: Well-watered regime; ΔWS : Percentage of variation due to stress; ***: significance at p = 0.001; **: significance at p = 0.01; ns: not significant, E: environment; G: genotypes.

The Table 4 presents the leaf water potential (Ψf) of the 32 evaluated sorghum genotypes. Analysis of this table indicates a highly significant difference (p < 0.001) in leaf water potential between the two water regimes. Under well-watered conditions (ww), all genotypes exhibited an average Ψf of -0.13125 MPa, indicating no water stress. In contrast, under water-stress conditions (ds), Ψf values ranged from -0.59 MPa to -4.84 MPa, reflecting varying levels of stress among genotypes. Specifically, genotypes $\sqrt{1}$, V2, V4, V8, V11, V12, V16, V22, V28, V29, and V31 experienced moderate stress (Ψf between -0.59 MPa and -0.99 MPa), whereas V3, $\sqrt{5}$, V6, V7, V9, V10, V13, V14, V15, V17, V19, V20, V21, V24, V25, V26, and V32 were severely affected (Ψf between -1.00 MPa and -1.88 MPa). These genotypes, although showing marked physiological responses to drought, remain within tolerance limits compared to V23, V27, and V30, which recorded the highest stress levels (Ψf below -2.00 MPa).

3.5. Effect of water stress on the agro-physiological traits of sorghum evaluated under well-watered and water-stress conditions

Table 5: Statistical parameters and mean performance of agro-physiological traits of sorghum evaluated under well-watered and water-stress conditions.

Parameters	ds	ww	ΔWS	Significatively
HP	144.18	182.60	- 21.04	***
NFA	13.21	13.21	0	ns
DAC	18.4	19.66	- 6.41	ns
NFD	5.8	2.78	108.63	***
SLA	145.90	121.71	19 87	***
DAPE	65.29	64.93	0.55	ns
DM	100.79	101.17	-0.3	ns
RDT	1610.68	3045.03	- 47.10	***
gs	0.06	0.08	- 25	***
A	12.68	14.73	- 13.91	***
E	2.53	4.24	- 40.33	***
Tleaf	42.34	39.69	6.67	***

ds: Stressed condition; ww: Well-watered condition; \(\Delta\WS\): Percentage variation due to stress; *** significance at \(p = 0.01\); ** significance at \(p = 0.01\); ns: not significant; NFA: Number of leaves emerged; DAC: Stem diameter (mm); HP: Plant height at maturity (cm); NFD: Number of dried leaves at the end of stress; SLA: Specific leaf area of the last ligulated leading (cm²-g-¹); A: Photosynthetic capacity (\(\mu\)mol-m-²-s-¹); Tleaf: Leaf temperature (°C); E: Transpiration (mmol-m-²-s-¹); gs: Stomatal conductance (mol-m-²-s-¹); DAPF: Days to first flowering (days after sowing); DM: Days to physiological maturity (days after sowing); RDT: Grain yield (t/ha).

Comparative analysis of the agro-physiological parameters of sorghum evaluated under well-watered (ww) and water-stress (ds) conditions reveals significant effects of water stress on several traits (Table 5, Figure 3). Plant height at maturity (HP) significantly decreased under stress, with a 21% reduction (144.18 cm vs. 182.60 cm; ***). This trend is confirmed by the height development throughout the vegetative cycle (Figure 3C), where stressed plants exhibit lower growth. In contrast, the number of leaves emerged (NFA) showed no significant difference, indicating that water stress did not affect initial leaf development (Figure 3A). Similarly, stem diameter (DAC) slightly decreased by 6% under stress, but this reduction was not statistically significant (Figure 3B).

Moreover, the number of dried leaves (NFD) strongly increased under stress, from 2.78 to 5.8 leaves, representing a 108% rise (***), reflecting pronounced foliar desiccation due to water

deficit. Specific leaf area (SLA) increased significantly by nearly 20 % (145.90 cm²·g⁻¹ vs. 121.71 cm²·g⁻¹; ***), suggesting morphological adjustments of leaves in response to stress. Phenological traits, including days to first flowering (DAPE) and physiological maturity (DM), showed no significant differences between the two regimes, indicating that water stress did not affect the plant development cycle.Regarding physiological performance, water stress caused a significant decrease in stomatal conductance (gs) by 25 % (0.06 vs. 0.08 mol·m²·s⁻¹; ***), photosynthetic capacity (A) by 13 % (12.68 vs. 14.73 μ mol·m²·s⁻¹; ***), and transpiration (E) by 40 % (2.53 vs. 4.24 μ mol·m²·s⁻¹; ***). Consequently, leaf temperature (Tleaf) increased significantly by 6 % under stress (42.34 °C vs. 39.69 °C; ***), reflecting reduced cooling via transpiration. Grain yield (RDT) was drastically reduced by 47 % under water stress (1610.68 kg/ha vs. 3045.03 kg/ha; ***).

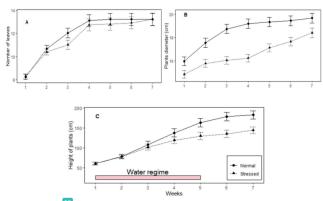


Figure 3: Changes in Number of Leaves (A), Stem Diameter (B), and Plant Height (C) under Well-Watered and Water-Stress Conditions.

Table 6: Drought Recovery Index (IDR) of Sorghum Genotypes for Plant Height, Stem Diameter, and Number of Leaves Emerged

Genotypes	IDR_ Height	IDR_ Diameter	IDR NFA	Codes	IDR_ Height	IDR_ Diameter	IDR_ NFA
V1	-0.45	-0.08	0.03	V17	-0.38	-0.09	-0.06

-0.24	-0.06	0.02	V18	-0.35	-0.13	-0.05
-0.40	-0.10	0.00	V19	-0.46	-0.06	-0.03
-0.22	-0.18	-0.04	V20	-0.22	0.13	-0.21
-0.24	-0.09	0.04	V21	-0.17	-0.14	-0.10
-0.25	0.19	0.00	V22	-0.25	-0.08	0.06
-0.51	-0.11	0.06	V23	-0.05	-0.28	0.04
-0.29	-0.10	0.05	V24	-0.06	-0.12	-0.01
-0.57	-0.23	-0.04	V25	-0.54	-0.16	-0.04
-0.12	0.01	-0.13	V26	-0.25	-0.06	0.06
-0.45	-0.25	0.02	V27	0.04	-0.06	-0.10
-0.19	-0.20	0.00	V28	-0.81	-0.44	-0.15
-0.29	-0.12	0.05	V29	-0.33	-0.07	-0.06
-0.35	-0.04	0.07	V30	-0.10	-0.07	0.00
-0.31	-0.15	0.05	V31	-0.38	-0.12	-0.04
-0.38	-0.08	-0.12	V32	-0.36	-0.17	0.12
	-0.40 -0.22 -0.24 -0.25 -0.51 -0.29 -0.57 -0.12 -0.45 -0.29 -0.35 -0.31	-0.40	-0.40 -0.10 0.00 -0.22 -0.18 -0.04 -0.24 -0.09 0.04 -0.25 0.19 0.00 -0.51 -0.11 0.06 -0.29 -0.10 0.05 -0.57 -0.23 -0.04 -0.12 0.01 -0.13 -0.45 -0.25 0.02 -0.19 -0.20 0.00 -0.29 -0.12 0.05 -0.35 -0.04 0.07 -0.31 -0.15 0.05	-0.40 -0.10 0.00 V19 -0.22 -0.18 -0.04 V20 -0.24 -0.09 0.04 V21 -0.25 0.19 0.00 V22 -0.51 -0.11 0.06 V23 -0.29 -0.10 0.05 V24 -0.57 -0.23 -0.04 V25 -0.12 0.01 -0.13 V26 -0.45 -0.25 0.02 V27 -0.19 -0.20 0.00 V28 -0.29 -0.12 0.05 V29 -0.35 -0.04 0.07 V30 -0.31 -0.15 0.05 V31	-0.40 -0.10 0.00 V19 -0.46 -0.22 -0.18 -0.04 V20 -0.22 -0.24 -0.09 0.04 V21 -0.17 -0.25 0.19 0.00 V22 -0.25 -0.51 -0.11 0.06 V23 -0.05 -0.29 -0.10 0.05 V24 -0.06 -0.57 -0.23 -0.04 V25 -0.54 -0.12 0.01 -0.13 V26 -0.25 -0.45 -0.25 0.02 V27 0.04 -0.19 -0.20 0.00 V28 -0.81 -0.29 -0.12 0.05 V29 -0.33 -0.35 -0.04 0.07 V30 -0.10 -0.31 -0.15 0.05 V31 -0.38	-0.40 -0.10 0.00 V19 -0.46 -0.06 -0.22 -0.18 -0.04 V20 -0.22 0.13 -0.24 -0.09 0.04 V21 -0.17 -0.14 -0.25 0.19 0.00 V22 -0.25 -0.08 -0.51 -0.11 0.06 V23 -0.05 -0.28 -0.29 -0.10 0.05 V24 -0.06 -0.12 -0.57 -0.23 -0.04 V25 -0.54 -0.16 -0.12 0.01 -0.13 V26 -0.25 -0.06 -0.45 -0.25 0.02 V27 0.04 -0.06 -0.19 -0.20 0.00 V28 -0.81 -0.44 -0.29 -0.12 0.05 V29 -0.33 -0.07 -0.35 -0.04 0.07 V30 -0.10 -0.07 -0.31 -0.15 0.05 V31 -0.38 -0.12

NFA: Number of emerged leaves; DRI: Drought Recovery Index

Table 6 presents the Drought Recovery Index (DRI), calculated according to the formula of Oukarroum et al. (2007) on plant height, stem diameter, and the number of emerged leaves of the different genotypes. DRI values close to zero indicate a good recovery capacity, whereas values near –1 indicate poor recovery. Considering plant height, DRI values range from –0.81 (V28) to 0.04 (V27), with the majority of genotypes showing negative values, reflecting partial or limited recovery after stress. However, several genotypes exhibit relatively good recovery, notably V27 (0.04), V23 (–0.05), V24 (–0.06), and V10 (–0.12). In contrast, genotypes V28 (–0.81), V9 (–0.57), and V25 (–0.54) show very low recovery for plant height.Regarding stem diameter, DRI values are generally negative, except for V6 (0.19), V20 (0.13), and V10 (0.01), which exhibit positive recovery. The lowest values are observed in V28 (–0.44), V11 (–0.25), and V23 (–0.28), indicating poor recovery for this trait. Conversely, several genotypes show DRI values close to zero or slightly positive for the number of emerged leaves (NFA), indicating good recovery of leaf activity after stress for genotypes V32 (0.12), V14 (0.07), V22, V7, V26 (0.06), and V8, V13, V15 (0.05).

3.7. Variability of physiological adaptation performances of the different genotypes under well-watered and water-stress conditions.

Table 7: Physiological performance of the different genotypes evaluated under well-watered and water-stress conditions.

VI 41.5 40.58 V2 41.82 40.22 V3 43.08 39.28 V4 40.19 39.12 V5 43.67 40.02 V6 42.3 40.02 V6 42.3 40.24 V7 43.49 39.24 V7 43.49 39.24 V8 39.96 40.24 V9 42.16 39.39 V10 43.23 40.43 V11 43.23 40.43 V12 41.88 40.45 V13 41.28 40.06 V14 42.12 41.72 V15 42.25 39.97	AWS -2.20 -3.84 -8.82 -2.66 -8.35 -9.78 -9.78 -6.56	2.98 2.98 2.26 2.26	MM		ľ								
41.8 43.08 43.08 40.19 43.67 43.67 43.29 42.16 42.16 42.16 42.16 43.23 42.18 41.28 41.28	-2.20 -3.84 -8.82 -2.66 -8.35 -4.86 -9.78 0.69	2.98		ΔWS	qs	WW	VWS	qs	WW	VWS	qs	WW	ΔWS
41.82 43.08 40.19 40.19 42.33 42.36 42.16 41.06 42.16 42.16 42.16 42.12 42.13 42.13 42.13 42.13	-3.84 -8.82 -2.66 -2.66 -8.35 -4.86 -9.78 0.69 -6.56	2.26	3.57	-16.66	90.0	0.07	-16.83	14.03abcde	12.97	8.21	155.15	124.19	24.92
43.08 40.19 40.19 43.67 42.3 43.49 39.96 42.16 41.06 41.28 42.12 42.25	-8.82 -2.66 -8.35 -4.86 -9.78 0.69 -6.56	2.26	3.65	-18.39	0.07	0.07	-6.01	13.73abcde	14.76	-7.02	143.00	166.45	-14.09
40.19 43.67 42.3 43.49 39.96 43.16 41.08 41.28 41.28 42.12 42.12	-2.66 -8.35 -4.86 -9.78 0.69 -6.56	2.11	4.63	-51.20	0.05	60.0	-45.51	10.71 ^{ef}	14.73	-27.31	170.64	104.47	63.33
43.67 42.3 43.49 43.96 41.06 41.06 43.23 41.28 42.12 42.12	-8.35 -4.86 -9.78 0.69 -6.56		3.62	-41.76	0.05	0.1	-52.39	11.45bcdef	14.15	-19.09	145.66	130.47	11.64
42.3 43.49 43.49 42.16 42.16 41.28 41.28 42.12 42.12	-4.86 -9.78 0.69 -6.56	2.94	5.14	-42.82	0.07	0.1	-31.90	14.06abcde	14.03	0.25	178.64	147.44	21.16
43.49 39.96 42.16 41.06 43.23 41.28 41.28 42.12	-9.78 0.69 -6.56 -7.28	2.70	4.14	-34.84	90.0	80.0	-30.54	12.88abcdef	12.15	5.98	145.27	96.38	50.72
39.96 42.16 41.06 43.23 41.28 41.28 42.12 42.12	0.69 -6.56 -7.28	1.83	4.00	-54.34	0.04	0.07	-38.80	9.84 ^f	14	-29.69	178.10	95.96	91.59
42.16 41.06 43.23 41.88 41.28 42.12 42.25	-6.56	2.22	3.68	-39.63	0.05	60.0	-49.89	11.05cdef	13.65	-19.08	161.36	115.42	39.81
41.06 43.23 41.88 41.28 42.12 42.25	-7.28	2.65	4.67	-43.33	90.0	0.1	-36.55	12.7abcdef	16.01	-20.64	104.31	137.17	-23.96
43.23 41.88 41.28 42.12 42.25		2.65	3.92	-32.39	0.07	80.0	-12.84	14.38abc	15.66	-8.21	167.28	104.63	59.88
41.88 41.28 42.12 42.25	-5.80	2.78	4.64	-40.07	90.0	60.0	-33.80	12.59abcdef	14.41	-12.63	135.54	99.54	36.17
41.28 42.12 42.25	-3.42	2.52	3.72	-32.25	0.05	0.07	-35.21	12.23abcdef	15.82	-22.73	194.92	69:56	103.71
42.12	-2.94	2.36	3.21	-26.45	0.05	90.0	-23.16	12.25abcdef	15.23	-19.57	186.60	148.34	25.79
42.25	-0.94	2.77	4.01	-30.82	0.05	80.0	-36.69	11.88abcdcf	14.96	-20.54	149.11	146.65	1.68
43.73	-5.39	2.91	3.79	-23.43	0.07	0.07	0.85	14.47 ^{ab}	11.87	21.85	129.65	130.09	-0.33
77:51	10.18	2.56	5.03	-49.14	90.0	0.1	-32.56	13.6abcde	16.48	-17.49	144.99	140.37	3.29
V17 42.09 36.53	13.21	1.93	4.36	-55.68	90.0	0.00	-37.19	12.3abcdef	12.76	-3.58	151.05	124.93	20.91
V18 44.16 41.39	-6.28	2.67	4.46	-40.21	0.05	80.0	-34.64	10.88 ^{def}	16.7	-34.83	153.54	106.87	43.67
V19 42.91 40.79	-4.95	2.99	3.71	-19.20	90.0	0.07	-3.33	12.65abcdef	15.56	-18.74	139.12	143.75	-3.23
V20 42.71 37.84	-11.40	1.92	5.42	-64,60	0.05	0.11	-55.88	11.96abcdef	15.59	-23.33	116.74	109.18	6.93

		su	121.71	150.11 1.56	133.11 27.19	98.63 30.47	103.91 25.25	135.86 -8.03	96.21 28.18	120.93 -13.92		137.30 2.31	102.38	133.24 2 102.38 137.30	115.17 133.24 102.38 137.30
1	* *	su	124.15	5 152.46	4 169.29	2 128.68	1 130.15	124.95	1 123.32	104.10	_	1 140 47			
				1 -19.76	9 -6.94	5 -14.22	1 -21.51	-18.34	5 -15.01	4 -5.21	5 -9.04	1 6.60		5 -2187	
		us	14.73	15.91	14.49	14.66	15.71	15.8	13.66	14.94	16.05	12.91	14.85	14.75	
* *	* *	* * *	12.68	12.76abcdef	13.49abcde	12.57abcdef	12.33abcdef	12.9abcdef	11.61 abcdef	14.16abcd	14.6^{ab}	13.76abcde	11.61abcdef	11.43 ^{bcdef}	14.91
				-47.20	-12.87	-50.70	-27.31	-51.93	-3.20	-24.58	-48.16	21.15	-40.43	-41.82	-14.30
		su	0.08	0.11	0.07	0.1	80.0	0.12	90.0	80.0	0.11	0.05	0.07	0.09	80.0
I	* *	su	90.0	90.0	90.0	0.05	0.05	0.06	0.06	90.0	0.05	90.0	0.04	0.05	0.07
				-53.33	-33.15	-53.72	-33.39	-56.00	-26.66	-31.05	-53.13	16.15	-50.56	-48.40	-38.70
		su	4.24	5.37	3.78	4.71	3.92	6.23	3.31	3.94	5.45	2.59	4.34	4.31	4.32
	* * *	su	2.53	2.51	2.53	2.18	2.61	2.74	2.43	2.72	2.56	3.01	2.14	2.22	2.65
				-7.82	-9.10	-5.70	-6.07	-5.82	-7.68	-4.83	-6.40	-0.18	-9.16	-8.27	10.07
1	* * *	su	39.69	39.29	38.33	39.19	39.95	40.98	39.12	39.57	40.51	41.19	40.06	38.68	38.42
		su	42.34	42.63	42.17	41.56	42.53	43.51	42.38	41.58	43.28	41.26	44.1	42.16	42.72
GxE	H	Genotype	Overall Mean	V32	V31	V30	V29	V28	V27	V26	V25	V24	V23	V22	V21

des stressed regime; ww: well-wateryd regime; AWS: percentage change due to stress; *** significance at p = 0.001; ** significance at p = 0.01; ns: not significantly light area of the last fully expanded leaf (cm² g²¹); Ar. photosynthetic capacity (µmol·m² s²¹); Tleaf: lgf temperature (°C); E: transpiration (mmol·m² ·s²); gs. stomatal conductance (mol·m² ·s²¹); E: environment; G: genotypes. Means sharing the same letter are not significantly different.

Table 7 presents the physiological performances of the different genotypes evaluated for drought tolerance. Analysis of this table reveals a highly significant difference between the different water regimes for the physiological parameters studied (p < 0.001). Under well-watered conditions, genotypes V24, V15, and V1 stand out for their high performance in terms of photosynthesis (A), stomatal conductance (gs), and transpiration (E), whereas V3, V7, and V20 exhibit lower values. Under water-stress conditions, most genotypes showed a marked reduction in photosynthesis, stomatal conductance, and transpiration. However, V1, V15, and V24 displayed an increase in photosynthetic capacity under stress, in contrast to V3, V7, V18, and V20, which showed greater reductions (up to -35%). Regarding specific leaf area (SLA), some genotypes exhibited a marked increase in response to stress: V12 (+103.7%), V7 (+91.6%), and V6 (+50.7%). Conversely, V2, V26, and V9 experienced a decrease in leaf area.

3.8. Variability of morphological performance of the different genotypes under well-watered and water-stress conditions Table 8. Morphological performance of the different genotypes evaluated under well-watered and water-stress conditions

VI 123.2 θing T/S ya θing AWS d6 VI 123.2 θing 175.93 θing 175.93 fing 18.3 θing 18.3 θing V3 125.53 θing 171.2 fing 26.7 γ 18.6 θing 17.7 γ V4 105.03 k 127.37 ding -17.5 fing 17.7 γ 17.7 γ V5 117.13 b 144.3 θing -17.5 fing 20.9 θing V6 127.49 θing 155.33 θing -17.9 ging 20.9 θing V7 113.27 βi 144.3 θing -22.6 fing 20.6 θing V9 153.87 θing -22.6 fing 20.6 θing V10 177.7 ding 20.5 γing -20.3 fing V11 144.03 θing 20.5 γing -20.3 fing V12 177.8 bing 20.5 γing -20.3 fing V13 158.23 hing 19.5 γing -20.3 fing V14 12.4 dighu 12.0 ging -20.7 fing V16 151.4 bing 20.5 bing -20.7 fing V16 151.4 bing 20.	18.13 18.03 18.03 18.05 18.05 18.05 17.47 17.02 17.02 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.05 17.0	AWS -4.23 -0.97 -5.92 -	ds 14.67abcde 12.33hijklm 13fghij	14.33 ^{bod} 12ghij	ΔWS 2.33	5.67abcd	ww 7 33cdef	∆ WS 142.86
123.2@ibik 175.93del@ibi 29.97 122.8@ibk 148.4@id -17.25 125.53@ibik 148.4@id -17.54 105.03# 127.37 ^{id} -17.54 117.13 ^{id} 144.3Gi ^{id} -18.86 113.27 ^{id} 145.92 ^{id} -22.65 113.8G ^{id} 145.92 ^{id} -22.65 144.03 ^{id} 195.73 ^{id} -22.65 177.72 ^{id} 195.73 ^{id} -22.65 177.72 ^{id} 195.73 ^{id} -20.73 177.8 ^{id} 195.73 ^{id} -20.37 132.4 ^{id} 172.13 ^{id} -23.08 171.8 ^{id} 200.2 ^{id} -21.62 173.4 ^{id} 172.13 ^{id} -20.37 158.82 ^{id} 172.13 ^{id} -21.62 179.27 ^{id} 20.29 ^{id} -21.62 179.27 ^{id} 20.29 ^{id} -21.62 179.27 ^{id} 195.29 ^{id} -21.63 179.27 ^{id} 195.29 ^{id} -23.44 127.33 ^{id} 194.5 ^{id} -20.97		-4.23 -0.97 -5.92 -	14.67abcde 12.33hijklm 13fghij	14.33 ^{bcd} 12ghij	2.33	5.67abcd	2 22cdef	142.86
122.8\text{big}		-5.92	12.33hijklm 13fghij	12ghij	0	6. bcd	2.33	
125.53\(\text{stability} \) 171.43\(\text{stability} \) 175.43\(\text{stability} \) 175.37\(\text{stability} \) 175.44\(\text{stability} \) 177.37\(\text{stability} \) 177.49\(\text{stability} \) 177.49\(\text{stability} \) 155.33\(\text{stability} \) 179.33\(\text{stability} \) 179.33\(\text{stability} \) 179.37\(\text{stability} \) 179.37\(\text{stability} \) 179.37\(\text{stability} \) 177.72\(\text{stability} \) 195.73\(\text{stability} \) 195.73\(\text{stability} \) 177.72\(\text{stability} \) 195.73\(\text{stability} \) 177.73\(\text{stability} \) 194.03\(\text{stability} \) 172.13\(\text{stability} \) 172.46\(\text{stability} \) 172.13\(\text{stability} \) 172.13\(\text{stability} \) 172.13\(\text{stability} \) 172.13\(\text{stability} \) 172.13\(\text{stability} \) 179.27\(\text{stability} \) 179.		-5.92	13fghij		2.78	na	2.33cdef	157.14
105.03\(\text{ik}\) 175.37\(\text{id}\) 175.43\(\text{ik}\) 177.13\(\text{ik}\) 144.36\(\text{ik}\) 177.49\(\text{ik}\) 177.49\(\text{ik}\) 193.33\(\text{ik}\) 193.97\(\text{ik}\) 193.97\(\text{ik}\) 193.97\(\text{ik}\) 145.92\(\text{id}\) 122.65\(\text{ik}\) 153.87\(\text{ik}\) 145.92\(\text{id}\) 177.72\(\text{ik}\) 195.73\(\text{ik}\) 195.73\(\text{ik}\) 177.72\(\text{ik}\) 195.73\(\text{ik}\) 177.72\(\text{ik}\) 195.73\(\text{ik}\) 177.72\(\text{ik}\) 195.73\(\text{ik}\) 177.73\(\text{ik}\) 177.73\(\text{ik}\) 177.73\(\text{ik}\) 177.73\(\text{ik}\) 177.73\(\text{ik}\) 179.27\(\text{ik}\) 179.27\(\text{ik}\)		11.68		13ctgh	0.00	7.33^{a}	2 ^{def}	266.67
117.1349k			13 33cfghij	13 67cdef	24 C-	2cde	2 33cdef	114 29
127.49@wik 155.33%wikd -17.93 113.27% 169.97%wik -33.36 112.86% 145.92%wik -22.65 153.87%wike 228.11 % -32.55 177.72%wike 195.73%wike -9.20 144.03%wike 200.49%wike -28.16 177.88%wike 194.39%wike 194.39%wike 194.39%wike 158.82%wike 194.39%wike 219.23% -23.44 179.27%wike 222.93%wike 179.27%wike 222.93%wike 179.27%wike 222.93%wike 125.44%wike 222.93%wike 222.93%wike 125.34%wike 127.33%wike 127.33%wike		-3.43	15.33abc	14.67abc	4.55	5.33cde	2 ^{ef}	166.67
113.27th 169.97ghij -33.36 112.86k 145.92ki -22.65 153.87dbcde 228.11db -32.55 177.72abc 195.73bcde/g -9.20 144.034clgh 200.49abcde -28.16 177.8abcd 205.72abcd -13.57 158.82abcde 199.43bcde 203.77 158.82dbcde 199.43bcde 20.37 151.47bcde 219.2abc -26.79 179.27bcd 216.23.44 179.27bcd 226.29.3abcde 179.27bcd 226.29.3abcde 226.79 179.27bcd 226.29.3abcde 226.79 179.27bcd 226.29.3abcde 226.79 179.27bcd 226.29.3abcde 226.79 179.27bcd 226.79 179.27bcd 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79 226.79	Н	18.30	14cdefgh	14cdc	0.00	7.33ª	3.67abc	100.00
112.86 k 145.92 id -22.65 153.87 bbcde 228.11 ab -32.55 177.72 abc 195.73 bcde -9.20 144.03 deligh 200.49 abcd -13.57 158.82 abcd -13.57 158.82 abcd 199.43 bcde -20.37 158.82 abcd 199.43 bcde -20.37 158.82 abcd -20.37 158.82 abcd -20.37 158.82 abcd -20.37 158.82 abcd -20.37 171.8a bcd 219.2abc -21.62 171.8a bcd 20.9abcd -26.79 179.27 abc 20.9abcd -2.1.62 179.27 abc 20.9abcd -2.1.63 179.27 abc 20.9abcd -2.1.63 179.27 abc 20.9abcd -2.5.74 179.27 abc 20.9abcd -2.3.44 147.07 adc 192.1 adc ge -2.3.44 127.33 qbuild 184.45 ac ge -30.97 -2.62		-6.24	14.33bcdef	13.67 ^{cdef}	4.88	6.67 ^{ab}	3 abcdef	122.22
153.87#cvde 228.11#h -32.55 177.72#bc 195.73#cvde@ -9.20 144.034*cdf@ 200.49#cvde -28.16 177.88#cvd 205.72#bcvd -13.57 158.82#bcvd 172.13*cdf@ 219.24#cv -20.37 151.47*cdf@ 219.24#cv -21.62 151.47*de 206.94#cvd -26.79 179.27#bc 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79 206.79		-6.64	15.33abc	14.67abc	4.55	6.67^{ab}	2d ^{ef}	233.33
177.72abc 195.73bcdefg -9.20 144.03defgh 200.49abcde -28.16 177.8abcd 205.72abcd -13.57 158.82abcde 199.43bcde -23.03 132.44fbd 219.2abc -21.62 151.47bcde 206.9abcd -26.79 179.27bcd 206.9abcd -26.79 179.27bcd 222.93ab -19.59 147.07bcde 192.1bcdefg -23.344 127.33fbuild 184.45defgh -30.97	20.6ab 23.83abc	13.55	15abc	15abc	0.00	2cde	2.33cdef	114.29
144,03de@h 200,49abcde -28,16 177,8abcd 205,72abcd -13,57 158,82abcde 190,43bcdef -20,37 132,4e@hi 172,13e@hid -23,08 171,8abcd 219,2abc -21,62 171,8abcd 219,2abc -21,62 179,27abc 220,23bc -19,59 147,07def 192,1bcde@h -23,44 127,33dehid 184,45de@h -30,97	16.61hi 16.18fgh	5.69	9.33mn	10.33 ^{ijk}	89.6-	2 _{de}	1.67 ^f	200.00
177.8abcd 205.72abcd -13.57 158.82abcde 199.43bcdef -20.37 132.4c@my 172.13c@my -21.62 171.8abcd 219.2abc -21.62 151.47bcde 206.9bcd -26.79 179.27abc 206.9bcd -26.79 147.07adc 192.1bcdef -23.44 147.07adc 184.45dcfef -30.97	16hi 19.02 ^{defg}	15.89	12.67ghijkl	12.33ghij	2.70	5de	2.67bcdef	87.50
158.82abrede 199.43brdef -20.37	17ghi 19.5defg	12.82	12.33 ^{ijklm}	12.33 fghi	0.00	7.33ª	3.33abcde	120.00
132,4°@hii 172,13°@hii -23,08	48ab 21.98abcde	-6.82	15.67ab	14.67abc	6.82	5.33bcde	2.67bcdef	100.00
171.8abod 219.2abc -21.62	18.2bcdefgh 18.67defgh	-2.50	12.67ghijk	12ghij	5.56	6.33apc	4ab	58.33
151.47bcde 206.9bcd -26.79 179.27abc 222.93ab -19.59 147.07cdef 192.1bcdefs -23.44 127.33/ghilk 184.45defgh -30.97	19.67abcd 21.8abcde	-9.79	16.33 ^a	15.67a	4.26	5.33bcde	5a	29.9
179.27abc 222.93ab -19.59 147.07cdef 192.1bcdefs -23.44 127.33'fbihk 184.45'defsh -30.97	20.27 ^{ab} 21.2 ^{bcde}	-4.40	13.33 ^{defghi}	14.67 ^{abc}	-9.09	5.33cde	3apcde	77.78
147.07cdef 192.1bcdefg -23.44 127.33fghijk 184.45defgh -30.97	20.4ab 21.2bcdc	-3.77	15.33abc	15.67a	-2.13	6.33abc	3abcdef	111.111
127.33fghijk 184.45defgh -30.97	17.87bcdcfgh 19.27dcfg	-7.27	11.33 ^{jklmn}	11.67ghijk	-2.86	5.67bcde	2 ^{def}	183.33
	20.09abc 20.57bcdef	-2.32	14.33bcdef	14.67abc	-2.27	5 _{de}	2.33cdef	114.29
						1	7	
138 ^{ergn1} 164.67 ^{nyk} -16.19		12.23	9.67 ^{imn}	11.33hijk	14.71	6.67 ^{ab}	3.33abcd	100.00
V21 117 ^{ijk} 147ijkl -20.41 21.27 ^{abcdef}	7abcdef 23abcd	-7.54	14 ^{cdefg}	14.33 ^{bcd}	-2.33	4.67bcdc	3.33abcde	40.00

		0.00581			1 000				0 99875		0 99477	CA E
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	* *	* *		* * *	* * * *		* *	* * *		* *	* *	Genotypes
1.09	2.78	5.8	0	13.21	13.21	-0.06	19.66	18.4	-0.21	182.60	144	Overall Mean
53.85	4.33 ^{ab}	6.67 ^{ab}	13.51	12.33ghij	14abcd	12.90	20.28bcdef	17.67ghi	-24.28	321.53ª	243.47abcde	V32
70.00	3.33abcd	5.67bcde	-2.17	15.33 ^{ab}	15abc	-7.08	18.37efgh	17.07 ^{fghi}	-26.57	195.53bcdef	143.58 ^{defgh}	V31
122.22	3apcde	6.67 ^{ab}	0.00	10 ^{ijk}	10klmn	-3.72	17.93 ^{cfgh}	17.27 ^{defghi} 17.93 ^{efgh}	-7.20	118.4	109.93^{jk}	V30
57.14	2.33cdef	3.67°	-4.26	15.67 ^a	15abc	-3.26	20.62 ^{bcdef}	19.95abcdefg 20.62bcdef	-20.19	183.6 ^{defgh}	146.53 ^{cdefg}	V29
37.50	2.67bcdef	3.67 ^{de}	-9.76	13.67 ^{cdef}	- 27.41 12.33 ^{hijklm}	27.41	25.33ª	18.39bcdefgh	-45.22	145.33 ^{ijkl}	79.61 ^k	V28
109.09	3.67abc	7.67 ^a	-3.45	9.67 ^{jk}	9.33 ^{mn}	-2.28	14.6 ^h	14.27i	2.40	185.8 ^{defgh}	190.27^{a}	V27
142.86	2.33cdef	5.67bcdc	5.56	12ghij	12.67ghij	-5.58	19.91 cdef	18.8abcdefg	-16.96	185.69cdefgh	154.2abcde	V26
150.00	2 ^{ef}	2 _{de}	-2.33	14.33 ^{bcd}	14cdefgh	-8.57	19.07defg	17.43defghi 19.07defg	-30.24	197.47 ^{bcde}	137.75efghi	V25
80.00	3.33abede	e _{apcd}	0.00	14 ^{cde}	14cdefg	-7.75	18.28 ^{cfgh}	16.87fghi	-6.08	204.57 ^{bcdefg}	192.13 ^{ab}	V24
320.00	1.67 ^f	7a	4.35	7.67k	 8	16.82	14.67 ^h	12.2i	4.60	154.13 ^{ijkl}	147.05 ^{cdefg}	V23
150.00	2 _{def}	5pcde	5.00	13.33detg	<u>+</u>	-4.01	20.23 beder	19.42abcuei	-16.20	145.47cde1g 173.58de1gnij	145.47 ^{cdetg}	V22

ds. water-stress condition; ww: well-watered condition; ΔWS: percentage change due to stress; *** significance at p = 0.001; ** significance at p = 0.01; ns. not significant. NFA: number of leaves emerged at maturity; DAC: stem diameter (mm); HP: plant height at maturity (cm); NFD: number of dired leaves at the end of stress; E: Environment; G: Genotype. Means sharing the same letter are not significantly different.

Table 8 presents the morphological performance of the different genotypes evaluated under well-watered (ww) and water-stress (ds) conditions. The results show highly significant differences between the water regimes for the main morphological parameters measured: plant height, stem diameter, number of leaves emerged, and number of dried leaves (p < 0.001). Moreover, genotypes displayed statistically significant differences for all parameters within the same water regime (p < 0.001).Under well-watered conditions, genotypes V32, V24, V17, and V12 exhibited the highest plant heights: 321.53 cm, 204.57 cm, 222.93 cm, and 205.72 cm, respectively. In contrast, V28, V30, and V4 showed the lowest heights, representing small-sized genotypes (less than 1.5 m). Regarding stem diameter, V28 displayed the highest value (25.33 mm) under well-watered conditions, while several genotypes, such as V23, V10, and V27, showed lower diameters. In terms of the number of leaves emerged, V15, V5, and V13 stood out with the highest leaf counts, indicating a good foliar development capacity, whereas V10, V20, and V23 had the lowest values. Water stress caused a reduction in plant height, stem diameter, and number of leaves emerged in most genotypes. However, some genotypes showed atypical responses: for example, V27 exhibited a slight height increase (+2.40%), and V20 showed a stem diameter increase (+12.23%), suggesting possible morphological adaptation strategies. Regarding the number of dried leaves, a generalized increase was observed in nearly all genotypes under water stress. The largest increases were recorded in V23 (+320%), V3 (+266.67%), V8 (+233.33%), and V10 (+200%), indicating high sensitivity to water stress. Conversely, V15 demonstrated marked tolerance, with only +6.67% of dried leaves.

3.9. Variability of Agronomic Performance of the Different Genotypes under Well-Watered and Water-Stress Conditions.

Table 9: Agronomic Performance of the Different Genotypes Evaluated under Well-Watered and Water-Stress Conditions.

Genotyp

Representations

Representations

Genotyp	٠	LP			ď			PP			RDT	
es	gp	WW	ΔWS	qs	WW	VMS	qs	WW	VMS	qs	MM	VMS
				6.17bcdef			36.11 ^{cdefg}			1576.39 ^{bcd}	2694.44 ^{ef}	
VI	33.89abc	38.67abcde	-12.36	ghij	8.89apcq	-30.63	hijk	63cdefg	-42.68	efghij	ghijk	-41.49
	31.44 ^{abcd}									1131.94ghij		
V2	efgh	33.17cdefghi	-5.19	6.78apcd	6.78bcd	0.00	29.33ghijk	31.94 ⁱ	-8.17	я	1281.25 ^m	-11.65
							36.33 bcdef	60.89 ^{def}		1354.17 ^{def}	2486.11 ^{ef}	
V3	30apcdefgh	35.56abcdefg	-15.63	7.17abcd	8.06^{apcd}	-11.03	ghijk	dg.	-40.33	ghijk	ghij	-45.53
	30.72abcd										2097.22 ^{ij}	
V4	cfgh	34.89abcdefg	-11.94	5.44 ^{fghij}	7.33abcd	-25.76	28.06hijk	44.44 ^{ghi}	-36.88	1031.25 ^{ijk}	klm	-50.83
	27.44abcd			5.61 efghi				63.78bcd			2743.06°	
V5	efgh	33.67cdefghi	-18.48	-	7.67abcd	-26.81	26.33 ^{ijk}	efg	-58.71	888.89jk	defghi	-67.59
				6.11cdefg						1229.17 ^{fghi}	1986.11 ^{ij}	
9/	34.89 ^{ab}	40.89^{ab}	-14.67	hij	6.94 pcd	-12.00	30.89ghijk	49ghi	-36.96	¥	klm	-38.11
				6.5abcdef				57.11 ^{efg}			2631.94fg	
Λ7	32abcdefg	36.56abcdef	-12.46	dg	8.22abcd	-20.95	30.33ghijk	Ņ	-46.89	1055.56hijk	hijkl	-59.89
	33.33abcd			6.11cdefg				96.78 ^{bcd}			2923.61 ^d	
N8	е	41.89^{a}	-20.42	hij	8.11abcd	-24.66	28hijk	efg	-58.07	1062.5 ^{ijk}	efghi	-63.66
	31.33abcd							62.33bcd			3055.56	
6/	efgh	41.44^{a}	-24.40	4.78 ^j	6.22^{cd}	-23.21	21.89 ^{jk}	efg	-64.88	937.5 ^{jk}	defghi	-69.32
	25.78abcd			6.28abcd							1590.28°	
V10	efgh	24.44 ^{kl}	5.45	efghi	6.78 ^{bcd}	-7.38	60.44 ^a	40.56ghi	49.04	2694.44 ^{abc}	defghi	-12.37
	23.67cdef			5.61 ^{defg}			41.44abcde	65.44 ^{bcd}		1888.89abc	3013.89°	
V11	gh	27.33hijkl	-13.41	hij	7.39abcd	-24.06	fghi	efg	-36.67	defghi	defghi	-37.33
	33.06abcd						46.22abcde	51.78fgh			1569.44 ^{jk}	
V12	e	34.33 bcdefgh	-3.72	7.33abc	7.78abcd	-5.71	fgh		-10.73	1812.5ghijk	III	-23.89
	24.22bcde						51.89abcde	87.78abc		2201.39abc	3854.17ª	
V13	lgh	32.44 ^{defghij}	-25.34	7.11 ^{ab}	10.22^{a}	-30.43	J	р	-40.89	р	bodefg	-42.88
V14	22.78cdef	24.44 ^{kl}	-6.82	5.22ghij	5.78 ^d	-9.62	31.72 ^{cfghij}	38.78hi	-18.19	1447.92 ^{cdef}	1520.83 ^{kl}	-4.79

	-47.71	-46.65		-51.79		-33.40		-62.29		-6.21		-63.38		-62.35		-45.85		-38.76		-62.01		-37.33	-9.55		-84.40		-48.25		-11 18
ш	4555.56 ^a bc	3736.11 ^a bcdefgh	4451.39ª	pcq	3284.72 ^b	cdefghi	4861.11ª	p.	2201.39g	hijklm	2465.28 ^{ij}	klm		5625a	2423.61 ^{ij}	klm	2131.94 ^{ij}	klm		4625abcd	4166.67 ^a	pode	1236.11 ^m	3916.67ª	pcdet	4951.39 ^a	p q	2361.11 ^{hi}	jklm
ghij	2381.94ªb	1993.06abc defghi	2145.83abc	def	2187.5abcde	fgh	1833.33abc	defghi	2347.22abc	de		902.78 ^{jk}	2118.06abc	de	1312.5 ^{cfghij}	×	1305.56 ^{efg}	hijk	1756.94abc	defghi		2611.11 ^a	1118.06hijk		611.11 ^k		2562.5abcde	2097.22ab	odof
	-38.97	-27.08		-43.63		-41.29		-53.94		-2.87		-33.89		-60.90		-36.93		-36.49		-57.70		-32.66	-19.48		-78.41		-45.04		17.01
	93.22 ^{abc}	78.78 ^{abc} def	94.22abc	p	82.89abc	de		93.11 ^{ab}	54.11 efg	æ		53.44ghi		109.11^{a}		51.44ghi	56.33 ^{cfg}	æ	91.67abc	р	82.33abc	qe	34.22 ⁱ		88.78apc		100.89^{a}	53.56 ^{fgh}	_
×	56.89abc	57.44 ^{ab}		53.11abcde	48.67abcde	fghi	42.89abcde	fghi		52.56abcde	35.33 efghij	×	42.67abcde	fghi	32.44 ^{fghij}	×	35.78defgh	ij	38.78abcde	fghij		55.44abcd	27.56 ^{hijk}		19.17 ^k	55.44abcde	fg	44.44abcde	fehi
	-13.92	-14.10		-30.00		-15.79		-16.45		-4.88		-20.16		-21.33		-13.93		-11.59		-25.00		-12.31	-10.89		-44.85		-19.23		000
	8.78abcd	8.67abc		10^{ap}		7.39abcd		8.44 _{abcd}		6.83 pcd		6.89bcd		8.33abcd		6.78 ^{bcd}		7.67abcd		8.22abcd		7.22abcd	5.61 ^d		9.17abc		8.67abcd		p370 7
	7.56ª	7.44ab		7apc	6.22bcdef	ghij		7.0 ^{6abc}	6.5abcdef	56)		5.5efghij	6.56abcd	ef	5.83 ^{defg}	hj	6.78abcd	et	6.17bcdef	ghij	6.33pcdef	ghij	5ij		5.06 ^{hij}		7abcde		cedefehii
	-18.61	-12.67		-14.41		-11.49		-13.74		5.47		-15.61		-22.90		-2.53		-7.78		-19.27		-2.07	92.0-		-28.82		-23.75		0.51
	40abc	33.33cdefghi		39.33abcd		26.11^{ijkl}		29.11ghijkl		25.39^{ikl}		26.33^{ijkl}		33cdefghi		26.33^{ijkl}		38.56abcde		30.56^{fghijk}		24.11^{kl}	21.94		31.61 ^{etghij}		26.67^{ijkl}		21 80
gh	32.56abcd ef	29.11 ^{jabc} defgh		33.67abcd	23.11 ^{cdef}	gh	25.11abcd	efgh	26.78cabc	defgh	22.22iefg	ч	25.44 ^{abcd}	efgh	25.67 ^{abcd}	cfgh		35.56^{a}	24.67 ^{abcd}	cfgh	23.61 ^{cdef}	gh	21.78fgh		22.5 ^{idetgh}		20.33 ^h		21 7efgh
	V15	V16		V17		V18		V19		V20		V21		V22		V23		V24		V25		V26	V27		V28		V29		1720

6.39mca 8.44abcd -24.34 fight 6.67abcd 7.78abcd -14.29 39.47 66.36 **** **** **** **** **** 0.000322	8.44abcd -24.34 41.89abcool 17.78abcd -14.29 hijk 7.72 -0.19 39.47 *** 0.000322 ****	27.33hijkl -23.58 6.29mou 41.89mou 41.89mou 41.89mou 41.83mou 6.67mbod 7.78abed 1.44.29 iijk 23.44odelis 31.67 -0.13 6.29 7.72 -0.19 39.47 *** *** *** ***	-23.58 e'@h 8.44abcd -24.34 41.89abcae e@h 6.67abcd 7.78abcd -14.29 hijk -0.13 6.29 7.72 -0.19 39.47 ****
6.67 ^{abcd} 7.78a ^{bcd} -14.29 6.29 7.72 -0.19 *** ***	6.67 ^{abcd} 7.78a ^{bcd} -14.29 6.29 7.72 -0.19 *** ***	32.22de@bij _4.83	32.22de@bij _4.83
6.67mbcd 7.78abcd	-23.58	27.33 ^{hijkl} -23.5	27.33 ^{hijkl} -23.5
6.29 7.72		32.22 ^{de@hij} -4.8	32.22 ^{de@hij} -4.8
*** ***		31.67 -0.11	31.67 -0.11
6.29 ***	-23.58 0.39moz	27.33hiki -23.5	27.33hiki -23.5
	-0.13 6.67mbol	32.22defbiji -4.8	32.22defbiji -4.8
	-0.13 6.29 ****	31.67 -0.1.	31.67 -0.1.
	-23.58	27.33 ^{bijid} -23.5	27.33 ^{bijid} -23.5
	-4.83	32.22 ^{de@hij} 4.8	32.22 ^{de@hij} 4.8
	-0.13	31.67 -0.11	31.67 -0.11

ds. Water-stress regime; ww: Mell-watered regime; AWS: Percentage change due to stress; ****: significance at p = 0.001; ***: significance at p = 0.01; ns. not significant; LP: paniele length (em); -IP: paniele width (em); PP: paniele weight (g); RDT: grain yield (t/ha); E: Environment; G: Genotype. Means sharing the same letter are not significantly different.

Table 9 presents the agronomic performance results of the different sorghum genotypes evaluated under both well-watered and water-stress conditions. Statistical tests revealed highly significant differences (p < 0.001) between the two water regimes for all traits studied. Additionally, a significant interaction was observed between genotypes and water regimes for most parameters, reflecting a wide diversity of responses to stress. Under well-watered conditions, several genotypes exhibited excellent performance, such as V22 (yield: 5625 kg/ha), V25 (4625 kg/ha), V15 (4555.56 kg/ha), V17 (4451.39 kg/ha), and V31 (4305.56 kg/ha). In contrast, some genotypes like V27, V30, and V1 showed yields below 2500 kg/ha.Under drought condition, agronomic performance dropped markedly. On average, grain yield decreased from 3004.23 to 1643.45 kg/ha, representing a loss of over 45%. Similarly, panicle weight declined by nearly half (-40% on average). However, genotypic responses were not uniform. Some genotypes, such as V10, V20, and V30, showed minimal variation due to drought, maintaining yields close to the control even under stress. Their panicle weights, well above average, further confirm their resilience under stressful conditions. Conversely, other genotypes, including V9, V8, V5, and V21, experienced substantial yield reductions, in some cases exceeding 60%.

3.10. Drought Stress Tolerance Analysis Based on STI and SSI Indices in the Evaluated Sorghum Genotypes

Table10: Drought Stress Tolerance and Susceptibility Indices of the Studied Genotypes

Codes	SSI	STI	Codes	SSI	STI
V1	0.881	0.458	V17	1.100	1.030
V2	0.247	0.156	V18	0.709	0.775
V3	0.967	0.363	V19	1.322	0.961
V4	1.079	0.233	V20	0.132	0.557
V5	1.435	0.263	V21	1.346	0.240
V6	0.809	0.263	V22	1.324	1.285
V7	1.272	0.300	V23	0.973	0.343
V8	1.351	0.335	V24	0.823	0.300
V9	1.472	0.309	V25	1.316	0.876
V10	0.263	0.715	V26	0.793	1.173
V11	0.792	0.614	V27	0.203	0.149
V12	0.507	0.202	V28	1.792	0.258
V13	0.910	0.915	V29	1.024	1.368
V14	0.102	0.237	V30	0.237	0.534
V15	1.013	1.170	V31	1.092	0.971
V16	0.990	0.803	V32	0.743	0.135

STI (Stress Tolerance Index).SSI (Stress Susceptibility Index)

Table 10 presents drought tolerance indices, specifically the Stress Susceptibility Index (SSI) and the Stress Tolerance Index (STI) for the 32 evaluated genotypes. The joint analysis of these two indices allowed the discrimination of genotypes based on their performance under water-stress conditions. The SSI measures a genotype's sensitivity to stress: a value below 1 indicates good tolerance, whereas a value above 1 indicates high sensitivity. The STI evaluates the ability of a genotype to maintain high productivity under stress: a value above 1 reflects good performance under both normal and stressed conditions. Cross-interpretation of these two indices allows the classification of genotypes as follows: only genotype V26 (SSI = 0.793; STI = 1.173) combines both drought tolerance (SSI < 1) and excellent agronomic performance (STI > 1). This genotype stands out as the most promising in the study, capable of maintaining high yield under

drought, and is thus a prime candidate for breeding programs in arid environments. In contrast, genotypes V2, V6, V10, V11, V12, V14, V16, V18, V24, V30, and V32, although drought-tolerant (SSI < 1), exhibit low production potential (STI \leq 1). This limits their direct value for production, but they can still serve as valuable genetic resources for improving crop resilience. Additionally, the group comprising genotypes V22, V29, and V15 shows drought sensitivity (SSI > 1) but demonstrates good productive capacity (STI > 1). These genotypes, while sensitive to drought, possess high potential yield, making them suitable for environments with low water constraints or for crosses aiming to combine yield and tolerance. Finally, genotypes V5, V7, V8, V9, V21, V23, V25, and V28 are the most negatively affected by drought, showing both high SSI and low STI values. These genotypes are unsuitable for environments with high water stress.

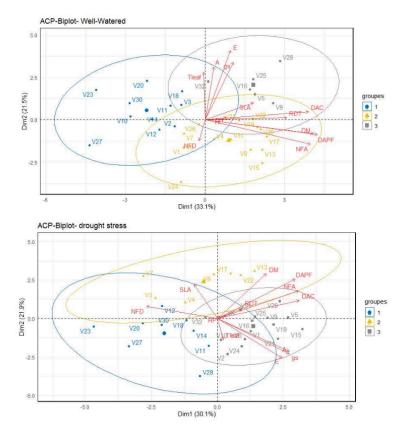
3.12. Principal Component Analysis of Agro-Physiological Traits Revealing Sorghum Genotype Grouping Under Well-Watered and drought Conditions.

Figure 4 presents the results of Principal Component Analysis (PCA) performed on the agro-physiological parameters measured in the 32 sorghum genotypes, evaluated under both normal conditions (PCA–WELL-WATERED) and water-stress conditions (PCA–WATER-STRESS).

Under well-watered conditions, the first two principal components (PCs) explained 54.6% of the total observed variability and allowed the classification of genotypes into three distinct groups. Group 1 (blue cluster) comprised genotypes V2, V3, V10, V11, V12, V14, V18, V20, V23, and V30. These genotypes were characterized by high leaf temperature but a relatively low number of dried leaves. Group 2 (green cluster), including genotypes V1, V4, V7, V8, V13, V15, V17, V19, V21, V22, V24, V26, and V29, was associated with fewer dried leaves and better agronomic performance (grain yield and stem diameter). In contrast, Group 3 (red cluster) gathered the best-performing genotypes in terms of physiological traits (V5, V9, V16, V25, V28, and V32).

Under drought conditions, the first two axes (Dim1 and Dim2) explained 52% of the total variance, also allowing the identification of three groups. Group 1 (blue cluster) consisted of genotypes V11, V12, V14, V18, V20, V23, V27, V28, and V30, distinguished by a high number of dried leaves and high specific leaf area. Group 2 (green cluster) included genotypes V3, V4,

V7, V8, V13, V17, and V22, characterized by long vegetative cycles (days to first flowering and physiological maturity). Conversely, Group 3 (red cluster) represented the most productive genotypes in terms of grain yield, panicle width, plant height, stem diameter, and number of leaves (V1, V5, V9, V10, V15, V16, V19, V21, V24, V26, and V29). Regarding correlations among variables, under well-watered conditions, physiological parameters (stomatal conductance, photosynthetic capacity, transpiration, and specific leaf area) were positively associated with grain yield, whereas the number of dried leaves was negatively correlated with yield. By contrast, under water-stress conditions, grain yield was negatively correlated with leaf temperature.



NFA: Number of leaves emerged; DAC: Stem base diameter (mm); HP: Plant height at maturity (cm); NFD: Number of dried leaves at the end of stress; SLA: Specific leaf area of the lead ligulated leaf (cm²·g⁻¹); A: Photosynthetic capacity (μmol·m⁻²·s⁻¹); Tleaf: Leaf paperature (°C); E: Transpiration (mmol·m⁻²·s⁻¹); gs: Stomatal conductance (mol·m⁻²·s⁻¹); DAPF: Days to first flowering (days after sowing); DM: Days to physiological maturity (days after sowing); RDT: Grain yield (t/ha).

Figure 4:Principal Component Analysis of the Agro-Physiological Parameters of 32 Genotypes Evaluated under Normal Conditions (PCA–Well-Watered) and Water-Stress Conditions (PCA–drought-stress)

4. Discussion

The increasing climatic constraints in the Sahelian and Sudanian-Sahelian zones, characterized by early droughts and shortened rainy seasons, deeply affect sorghum cultivation. This climatic instability highlights the need to adapt breeding systems, particularly by targeting sensitive stages such as panicle initiation, a key moment for yield formation (Abreha et al., 2022; Tovignan et al., 2016). The study showed that genotypes differ markedly in their ability to cope with water stress at such a critical stage. These differences are not only reflected in yield losses but also in fine physiological responses that may serve as a basis for breeding. For instance, some tested genotypes were able to maintain a relatively high leaf water status despite moderate to severe stress conditions. This maintenance suggests the activation of mechanisms such as early stomatal closure, reduced transpiration, or deeper rooting. These mechanisms are well described in the literature as classical strategies to limit water losses (Blum, 2010; Lehrer et al., 2025). In the present study, these responses translated into better leaf integrity, with fewer dried leaves in certain genotypes (e.g., V1, V2, or V28). These observations confirm that the ability to maintain leaf water potential is a relevant marker for varietal screening, as also demonstrated by Chen et al. (2020).

In parallel, the analysis of gas exchange highlighted differentiated behaviors among genotypes. Some, despite water constraints, maintained relatively stable photosynthesis and transpiration rates. This could be explained by partial rather than complete stomatal closure, allowing a compromise between water conservation and carbon assimilation. This phenomenon is consistent with the findings of Lopez et al. (2017), who showed that moderate reduction in transpiration without excessive impairment of photosynthesis can enhance tolerance. For example, genotypes V1 and V24 exhibited good photosynthetic stability, suggesting fine stomatal regulation and potentially a leaf architecture favorable to water-use efficiency. Conversely, other genotypes (such as V18 or V20) showed a sharp decline in photosynthesis and transpiration, indicating greater sensitivity. From a morphological standpoint, the observed reduction in plant height under stress represents a well-known adaptive response aimed at limiting transpiring surface area. This reduction, around 21% in the present study, is consistent with the findings of Somfalvi-Tóth et al. (2024), who reported that severe water stress induces a halt in cell elongation. However, stem diameter was relatively unaffected, which could indicate a redistribution of resources toward maintaining the basic structural integrity of the plant. This suggests that some genotypes adopt

"prioritization" strategies, reducing certain functions (vertical growth) to preserve others (structural stability), as noted by Tovignan et al. (2023). In terms of leaf growth, several genotypes (V32, V14, V22, V7, V26, and V8, V13, V15) showed an ability to continue or resume leaf production immediately after the stress period. This behavior, reflected by a slightly positive leaf recovery index in some genotypes, corresponds to a form of resilience already identified by Gano et al. (2021) as a key selection criterion. They showed that, despite constraints, certain varieties can restart their vegetative apparatus and potentially sustain the recovery of photosynthesis. However, it should be noted that these adaptive responses were not sufficient to offset yield losses. The present study reported an average grain yield reduction of nearly 47%, consistent with declines ranging from 36 to 87% reported by de Souza et al. (2021) and Sanjari et al. (2021) under similar conditions. This significant loss indicates that, although tolerance mechanisms contribute to survival and vegetative recovery, they do not necessarily safeguard reproductive function. This finding is particularly relevant for late-maturing varieties, whose delayed development exposes them to the premature end of the rainy season (Guan et al., 2015; Sultan et al., 2013). Late genotypes, although physiologically more robust, no longer benefit from the climatic window required for their full cycle. These results underline the need for a new integrated breeding strategy. Relying solely on classical physiological criteria (such as transpiration or stomatal conductance) is no longer sufficient; it is imperative to incorporate specific reproductive traits such as flowering stability, effective panicle number, and grain filling. Sabadin et al. (2012b) and Kapanigowda et al. (2013b) also emphasize the importance of these traits in ensuring stable yields under drought conditions. Therefore, combining morphophysiological and reproductive data, coupled with genetic and molecular analysis, represents a promising pathway toward the identification of new, more resilient sorghum genotypes.

5. Conclusion

The study revealed marked inter-genotypic variability in sorghum responses to water stress, both in agro-morphological and physiological traits. Genotype V26, in particular, stood out for its ability to maintain high yield under water deficit conditions, combining a low stress susceptibility index (SSI < 1) with a high stress tolerance index (STI > 1). This profile positions it as a priority candidate for breeding programs targeting areas with severe water constraints. Principal component analysis also identified distinct groups of genotypes according to their

adaptive profiles, confirming the relevance of an integrated approach that considers the diversity of adaptation strategies: avoidance, tolerance, or a combination of both. These findings emphasize the importance of integrating agro-physiological traits, tolerance indices (SSI, STI), and multivariate analyses for the selection of genotypes adapted to water stress conditions.

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Data Availability Statement: All the data used to support the findings of the study are available within the research article.

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Conflicts of Interest: The authors declare no conflicts of interest

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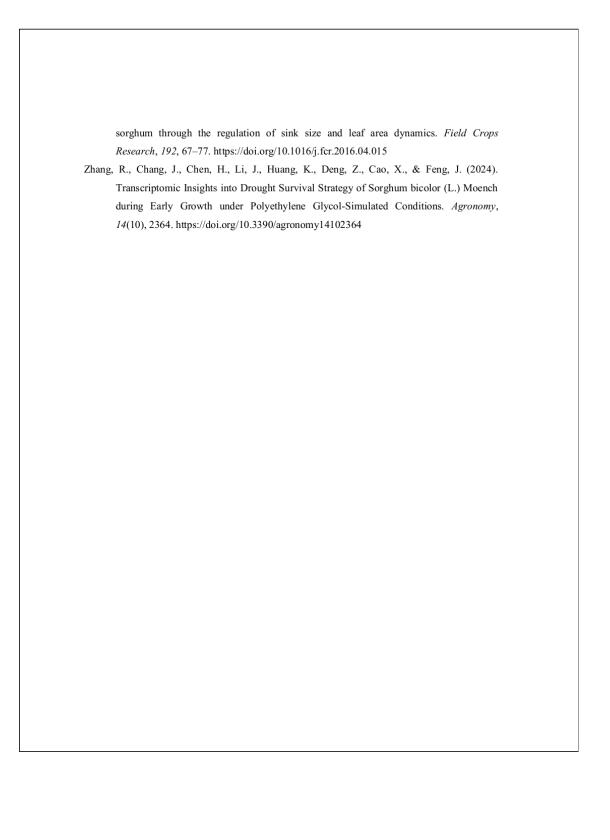
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