



Identification of Drought Tolerant Wheat Genotypes through Multi-Index Approach under Different Irrigation Regimes

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ABSTRACT

Wheat is a globally important cereal crop that serves as a major staple for billions of people. However, its productivity in rainfed areas is increasingly threatened by drought (one of the most severe abiotic stresses), which is exacerbated by climate change. This study aimed to evaluate the drought tolerance of 33 exotic wheat genotypes along with seven local checks under irrigated and rainfed conditions during the 2023–24 growing season at the Balochistan Agricultural Research and Development Center (BARDC), Quetta. The experiment was conducted using a factorial RCBD design with three replicates. Grain yield under normal (Y_{pi}) and stress (Y_{si}) conditions was used to compute 11 drought tolerance indices, including Mean Productivity (MP), Geometric Mean Productivity (GMP), Harmonic Mean (HARM), Stress Tolerance Index (STI), and others. Genotypes BARDC-26, BARDC-31, BARDC-24, BARDC-9, BARDC-14, and BARDC-19 consistently exhibited the highest values across multiple indices (MP, GMP, MRP, HARM, MSTIK1, MSTIK2, and REI), as well as superior grain yield under both conditions, classifying them as drought-tolerant and high-yielding. In contrast, BARDC-16, BARDC-15, BARDC-12, BARDC-8, and BARDC-2 exhibited low index values and were identified as drought-sensitive. Correlation analysis revealed strong positive relationships between both Y_{pi} and Y_{si} with MP, GMP, HARM, REI, and MSTIK1, while TOL was negatively correlated with Y_{si}, underscoring its inefficiency in selecting drought-tolerant genotypes. PCA showed that the first two components explained 97.7% of the total variation, with drought indices such as MP, GMP, and STI contributing most to genotype differentiation. Cluster analysis grouped the genotypes into four distinct clusters, with Cluster I (including BARDC-26, BARDC-31, BARDC-24, BARDC-9, BARDC-14, BARDC-25, and BARDC-19) comprising the most stable and drought-tolerant genotypes. These findings underscore the effectiveness of multivariate drought indices in identifying stable, high-yield genotypes for both irrigated and rainfed environments. The selected genotypes offer strong potential for future breeding programs targeting drought resilience in arid and semi-arid regions of Balochistan.

INTRODUCTION

Wheat (*Triticum aestivum* L.), often referred to as the king of cereals, holds a vital position among cereal crops globally. (Abd El-Mohsen *et al.*, 2015; Ahmed *et al.*, 2023). It plays a vital role in ensuring global food security, contributing approximately 20% of the world's calorie intake and 21% of its protein requirements (Shiferaw *et al.*, 2013; Nawaz *et al.*, 2025). As the primary staple for nearly two billion individuals, representing approximately 36% of the world's population, wheat is indispensable to human nutrition and livelihoods (Kumar *et al.*, 2024; Guo *et al.*, 2020). Global wheat production reached a record 791.02 million metric tonnes in 2023-24, with projections for 2024-25 estimating a slight increase to 793.24 million

metric tonnes (Ulukan, 2024). In Pakistan, the cultivated area under wheat increased by 6.6% from 9.0 million hectares in 2022-23 to 9.6 million ha during 2023-24. During the same period, the production area rose from 28.2 million metric tonnes to 31.4 million metric tonnes, an 11.6% increase. Wheat contributes 2.2% to Pakistan's GDP and accounts for 9.0% of the value added in the agricultural sector (Bajkani *et al.*, 2025). Despite its global and national importance, wheat production faces mounting threats from climate change. Rising temperatures and prolonged dry spells increasingly disrupt wheat growth, development, and yield. Drought, driven by climate change, is among the most significant abiotic stressors, accounting for approximately 60% of

crop failures due to climate disasters (Lao *et al.*, 2021; Shahi *et al.*, 2022). In developing regions, drought is the prominent cause of crop yield reduction, a trend that is expected to intensify with ongoing global warming (Shahrokhi *et al.*, 2020). To meet the escalating food demands of a growing global population, particularly in water-scarce regions, it is imperative to enhance wheat productivity and its resilience to drought conditions (Del Pozo *et al.*, 2016; Shahi *et al.*, 2024a). In Pakistan, drought stress has emerged as a critical challenge, especially in arid regions such as Balochistan (Naz *et al.*, 2020; Muhammad *et al.*, 2025; Khan *et al.*, 2013). Rainfed agriculture is heavily reliant on precipitation and is particularly vulnerable to different stresses at different crop stages (Mahmood *et al.*, 2019). During 2023–24, wheat cultivated under rainfed conditions covered 1.14 million hectares and yielded 1.89 million metric tons. In Balochistan alone, 30,000 hectares of rainfed wheat produced only 34,000 metric tons (Durrani *et al.*, 2025).

Drought stress can have more detrimental effects on plant growth than all other environmental stresses combined (Al-Maskri *et al.*, 2016; Shahi *et al.*, 2024b). It is a widespread issue affecting virtually all wheat-growing regions by inducing severe osmotic stress and limiting productivity (Khan *et al.*, 2011; Abhinandan *et al.*, 2018; Ahmed *et al.*, 2024). Wheat is especially sensitive to water deficit at the seedling, mid-season, and reproductive growth stages, making drought a major concern across developmental phases (Mujtaba *et al.*, 2016). Drought negatively affects plant growth parameters, notably plant height, by limiting water availability, disrupting photosynthesis, and hindering nutrient uptake (Sarto *et al.*, 2017; Naveed *et al.*, 2025). Water scarcity during the pre-anthesis stage may delay or suspend flowering, along with causing seed abortion (Ahmed *et al.*, 2020), and stress during the reproductive phase often results in more severe yield losses than the vegetative phase (Wang *et al.*, 2017; Dhakal, 2021). To mitigate drought-induced yield reductions, researchers have increasingly employed drought tolerance indices to identify genotypes capable of maintaining productivity under water stress. These indices assess the relationship between yield under stress (Y_s) and non-stress (Y_p) conditions, offering valuable insights into genotypic performance (Bennani *et al.*, 2017; Abd El-Aty *et al.*, 2024).

In this context, indices such as Mean Productivity (MP), Geometric Mean Productivity (GMP), and Harmonic Mean (HARM) provide a general measure of overall performance across environments (Yadav *et al.*, 2024). The Stress Susceptibility Index (SSI), Stress Tolerance Index (TOL), and Yield Reduction Ratio (YR) specifically quantify yield decline under drought conditions, highlighting genotypic tolerance or vulnerability (Mubushar *et al.*, 2022). The Stress Tolerance Index (STI) and its modified forms (MSTIk1 and MSTIk2) are particularly effective in identifying genotypes that perform well under both stress and non-stress conditions. Additional indices such as the Yield Index (YI), Relative Drought Index (RDI), and Drought Resistance Index (DI) enable comparative evaluation across genotypes relative to the mean, facilitating the selection of drought-resilient lines (Sofi *et al.*, 2021). Therefore, the present study was undertaken to

evaluate drought-tolerant wheat genotypes using multiple drought tolerance indices, compare the effectiveness of different drought resistance indices, and recommend superior genotypes suitable for cultivation in drought-prone regions.

MATERIALS AND METHODS

Study Site Characteristics

The experiment was conducted at the research farm of the Balochistan Agricultural Research and Development Centre (BARDC), Quetta, Pakistan (30° .1952296' N, 66° .9637362' E). The site is located in a semi-arid agro-climatic zone at an elevation of 1,680 m (5,510 ft) above sea level. The region experiences an average annual precipitation of 200 to 250 mm (8 to 10 in), with the majority occurring during the winter months. Summers are typically warm, with average temperatures ranging between 25°C and 35°C but getting hotter due to climate change with a 1–2-degree increase, whereas winters are cold, with temperatures often dropping below the freezing point, ranging from -5°C and 10°C.

Table 1

Monthly Average Temperature and Rainfall Data During 2023/24 Growing Season at BARDC Quetta.

| Months | Temperature (°C) | | Precipitation (mm) |
|----------|------------------|---------------|--------------------|
| | Min (Average) | Max (Average) | |
| November | 7.3 | 22.2 | 9.38 |
| December | 2.9 | 18.7 | 3.29 |
| January | 3 | 16.8 | 23.68 |
| February | 2.8 | 15.9 | 51.2 |
| March | 5.7 | 16.9 | 66.4 |
| April | 10.3 | 22.3 | 128.6 |
| May | 17.2 | 31.4 | 0 |

The soil at the experimental site was characterized as silty loam. It exhibited a slightly alkaline pH (7.98), low levels of organic matter (0.91%), nitrogen (0.05%), and phosphorus (5.4 mg/kg), whereas potassium levels were adequate (Table 2). Overall, the soil was considered marginally fertile and suboptimal for crop growth without any agronomic interventions.

Table 2

Soil Physiochemical Properties of Experimental Site.

| Properties | Values |
|---|------------|
| Textural Class | Silty Loam |
| pH | 7.98 |
| Electrical Conductivity (dS m ⁻¹) | 0.43 |
| Organic Matter (%) | 0.69 |
| Total Nitrogen (%) | 0.05 |
| Available Phosphorus (mg kg ⁻¹) | 5.4 |
| Available Potassium (mg kg ⁻¹) | 117 |

Field Experiment and Treatments

The experiment was conducted during the Rabi season of 2023-24 using a Randomized Complete Block Design (RCBD) with three replications, arranged in a factorial layout involving two irrigation regimes (irrigated and rainfed). A total of 40 wheat genotypes were evaluated, comprising 33 test genotypes and seven check varieties.

The experimental field was initially prepared by ploughing twice using a cultivator. Sowing was performed on 1st November using a manual hand drill, with each plot consisting of two rows, 3 m in length, spaced 20 cm apart. Three irrigations were applied under the irrigated treatment, whereas only one irrigation of 15 mm (equivalent to approximately half an inch) was applied under the rainfed condition on December 17th.

Drought Tolerance Indices

Grain yield data were recorded for each genotype under both irrigated and rainfed conditions. These yield values were used to calculate 11 drought tolerance/resistance indices, which were used to evaluate the performance of genotypes under water-deficit and non-stressed environments. The indices computed are listed in Table 3.

Table 3
Stress Indices, Their Abbreviations, Formulas and References.

| S.No | Stress Indices | Formula | Reference |
|------|--|--|------------------------------|
| 1 | Stress Tolerance (TOL) | $Y_{pi} - Y_{si}$ | (Rosielle and Hamblin, 1981) |
| 2 | Mean Productivity (MP) | $(Y_{pi} + Y_{si})/2$ | (Rosielle and Hamblin, 1981) |
| 3 | Mean Relative Performance (MRP) | $(Y_{si}/Y_s) + (Y_{pi}/Y_p)$ | (Hossain et al., 1990) |
| 4 | Geometric Mean Productivity (GMP) | $\sqrt{Y_{pi} \times Y_{si}}$ | (Fernandez, 1993) |
| 5 | Relative Efficiency Index (REI) | $Y_{si}/Y_s \times Y_{pi}/Y_p$ | (Hossain et al., 1990) |
| 6 | Modified Stress Tolerance Index 1 (MSTIK1) | $((Y_{pi})^2/(Y_p)^2) \times (Y_{si} \times Y_{pi})/(Y_p)^2$ | (Farshadfar and Sutka, 2003) |
| 7 | Modified Stress Tolerance Index 2 (MSTIK2) | $((Y_{si})^2/(Y_s)^2) \times (Y_{si} \times Y_{pi})/(Y_p)^2$ | (Farshadfar and Sutka, 2003) |
| 8 | Harmonic Mean Of Yield (HARM) | $2 \times (Y_{pi} \times Y_{si})/(Y_{pi} + Y_{si})$ | (Dadbakhsh et al., 2011) |
| 9 | Relative Drought Index (RDI) | $(Y_{si}/Y_{pi})/(Y_s/Y_p)$ | (Fischer and Wood, 1979) |
| 10 | Drought Resistance Index (DI) | $Y_{si} \times (Y_{si}/Y_{pi})/(Y_s)$ | (Lan, 1998) |
| 11 | Golden Mean (GM) | $(Y_{pi} + Y_{si})/(Y_{pi} - Y_{si})$ | (Moradi et al., 2012) |

Statistical analysis

The collected data were subjected to multivariate statistical analyses to comprehensively evaluate genotypic responses under irrigated and rainfed conditions. Analysis of variance (ANOVA) was performed using Statistics 8.1 to determine the significance of the differences among genotypes and treatments. Pearson’s correlation analysis was conducted to assess the relationships between drought tolerance indices and yield components. Principal component analysis (PCA) was employed to reduce data dimensionality and identify the most influential indices contributing to drought tolerance. Hierarchical cluster analysis, using Ward’s method, was performed to classify genotypes into distinct groups based on their performance

under stress and non-stress conditions. All statistical analyses were performed using the Jamovi software.

RESULTS

In the current study, genotypes BARDC-26, BARDC-24, BARDC-9, BARDC-31, BARDC-25, and BARDC-19 showed superior performance with consistently high MP, GMP, HARM, and STI values (Table 5). For instance, BARDC-26 recorded the highest GMP (5316.2), HARM (5229.7), MP (5404.2), and STI (2.55) values. Similarly, BARDC-24 and BARDC-9 also showed strong performance across these indices, with GMP values of 5119.9 and 5104.8, respectively, and STI values of 2.45 and 2.44.

Table 4
Mean Values of Yield in Non-Stressed (Y_{pi}), Yield in Stressed (Y_{si}), Tolerance Index (TOL), Mean Productivity (MP), Mean Relative Performance (MRP), Geometric Mean Productivity (GMP), Relative Efficiency Index (REI), Modified Stress Tolerance.

| Genotype | Y _{si} | Y _{pi} | TOL | MP | MRP | GMP | REI | MSTIK1 | MSTIK2 | HARM | RDI | DI | GM |
|----------|-----------------|-----------------|--------|--------|------|--------|------|--------|--------|--------|------|------|-------|
| BARDC-1 | 3612.5 | 5087.5 | 1475.0 | 4350.0 | 2.05 | 4287.0 | 1.05 | 0.89 | 0.79 | 4225.0 | 0.94 | 0.71 | 5.90 |
| BARDC-2 | 2675.0 | 4616.7 | 1941.7 | 3645.9 | 1.70 | 3514.2 | 0.71 | 0.49 | 0.29 | 3387.3 | 0.77 | 0.43 | 3.76 |
| BARDC-3 | 4116.7 | 4591.7 | 475.0 | 4354.2 | 2.09 | 4347.7 | 1.08 | 0.74 | 1.05 | 4341.2 | 1.19 | 1.02 | 18.33 |
| BARDC-4 | 3391.7 | 4833.3 | 1441.6 | 4112.5 | 1.94 | 4048.8 | 0.94 | 0.71 | 0.62 | 3986.2 | 0.93 | 0.66 | 5.71 |
| BARDC-5 | 3541.7 | 3766.7 | 225.0 | 3654.2 | 1.76 | 3652.5 | 0.76 | 0.35 | 0.55 | 3650.7 | 1.25 | 0.92 | 32.48 |
| BARDC-6 | 4083.3 | 5208.3 | 1125.0 | 4645.8 | 2.21 | 4611.6 | 1.22 | 1.07 | 1.16 | 4577.7 | 1.04 | 0.88 | 8.26 |
| BARDC-7 | 3133.3 | 5016.7 | 1883.4 | 4075.0 | 1.91 | 3964.7 | 0.90 | 0.74 | 0.51 | 3857.4 | 0.83 | 0.54 | 4.33 |
| BARDC-8 | 2991.7 | 3900.0 | 908.3 | 3445.9 | 1.63 | 3415.8 | 0.67 | 0.33 | 0.34 | 3386.0 | 1.02 | 0.63 | 7.59 |
| BARDC-9 | 4358.3 | 5979.2 | 1620.9 | 5168.8 | 2.44 | 5104.8 | 1.49 | 1.73 | 1.62 | 5041.7 | 0.97 | 0.88 | 6.38 |
| BARDC-10 | 2933.3 | 4500.0 | 1566.7 | 3716.7 | 1.74 | 3633.2 | 0.76 | 0.50 | 0.37 | 3551.5 | 0.86 | 0.53 | 4.74 |
| BARDC-11 | 3966.7 | 4816.7 | 850.0 | 4391.7 | 2.09 | 4371.1 | 1.09 | 0.83 | 0.99 | 4350.6 | 1.09 | 0.90 | 10.33 |
| BARDC-12 | 3175.0 | 3658.3 | 483.3 | 3416.7 | 1.64 | 3408.1 | 0.67 | 0.29 | 0.38 | 3399.6 | 1.15 | 0.76 | 14.14 |
| BARDC-13 | 2900.0 | 4991.7 | 2091.7 | 3945.9 | 1.84 | 3804.7 | 0.83 | 0.67 | 0.40 | 3668.6 | 0.77 | 0.46 | 3.77 |
| BARDC-14 | 4183.3 | 6141.7 | 1958.4 | 5162.5 | 2.43 | 5068.8 | 1.47 | 1.80 | 1.47 | 4976.8 | 0.90 | 0.79 | 5.27 |
| BARDC-15 | 3183.3 | 3541.7 | 358.4 | 3362.5 | 1.61 | 3357.7 | 0.65 | 0.26 | 0.37 | 3352.9 | 1.19 | 0.79 | 18.76 |
| BARDC-16 | 2441.7 | 4662.5 | 2220.8 | 3552.1 | 1.64 | 3374.1 | 0.65 | 0.46 | 0.22 | 3205.0 | 0.69 | 0.35 | 3.20 |

| | | | | | | | | | | | | | |
|-------------------|--------|--------|--------|--------|------|--------|------|------|------|--------|------|------|-------|
| BARDC-17 | 2658.3 | 5408.3 | 2750.0 | 4033.3 | 1.86 | 3791.7 | 0.82 | 0.78 | 0.33 | 3564.5 | 0.65 | 0.36 | 2.93 |
| BARDC-18 | 3525.0 | 5541.7 | 2016.7 | 4533.4 | 2.12 | 4419.8 | 1.12 | 1.12 | 0.80 | 4309.1 | 0.84 | 0.62 | 4.50 |
| BARDC-19 | 4141.7 | 5866.7 | 1725.0 | 5004.2 | 2.36 | 4929.3 | 1.39 | 1.56 | 1.37 | 4855.5 | 0.94 | 0.81 | 5.80 |
| BARDC-20 | 3925.0 | 4166.7 | 241.7 | 4045.9 | 1.95 | 4044.0 | 0.94 | 0.53 | 0.83 | 4042.2 | 1.25 | 1.02 | 33.48 |
| BARDC-21 | 3366.7 | 4000.0 | 633.3 | 3683.4 | 1.76 | 3669.7 | 0.77 | 0.40 | 0.50 | 3656.1 | 1.12 | 0.78 | 11.63 |
| BARDC-22 | 3891.7 | 4225.0 | 333.3 | 4058.4 | 1.95 | 4054.9 | 0.94 | 0.55 | 0.82 | 4051.5 | 1.22 | 0.99 | 24.35 |
| BARDC-23 | 3791.7 | 4116.7 | 325.0 | 3954.2 | 1.90 | 3950.9 | 0.89 | 0.49 | 0.74 | 3947.5 | 1.22 | 0.96 | 24.33 |
| BARDC-24 | 4375.0 | 5991.7 | 1616.7 | 5183.4 | 2.45 | 5119.9 | 1.50 | 1.75 | 1.64 | 5057.3 | 0.97 | 0.88 | 6.41 |
| BARDC-25 | 4275.0 | 5591.7 | 1316.7 | 4933.4 | 2.34 | 4889.2 | 1.37 | 1.39 | 1.43 | 4845.5 | 1.01 | 0.90 | 7.49 |
| BARDC-26 | 4433.3 | 6375.0 | 1941.7 | 5404.2 | 2.55 | 5316.2 | 1.62 | 2.14 | 1.82 | 5229.7 | 0.92 | 0.85 | 5.57 |
| BARDC-27 | 4200.0 | 5191.7 | 991.7 | 4695.9 | 2.24 | 4669.6 | 1.25 | 1.09 | 1.26 | 4643.5 | 1.07 | 0.94 | 9.47 |
| BARDC-28 | 3791.7 | 4400.0 | 608.3 | 4095.9 | 1.96 | 4084.5 | 0.96 | 0.60 | 0.79 | 4073.3 | 1.14 | 0.90 | 13.47 |
| BARDC-29 | 2941.7 | 4825.0 | 1883.3 | 3883.4 | 1.81 | 3767.5 | 0.81 | 0.62 | 0.40 | 3655.0 | 0.81 | 0.49 | 4.12 |
| BARDC-30 | 2925.0 | 4491.7 | 1566.7 | 3708.4 | 1.74 | 3624.7 | 0.75 | 0.49 | 0.37 | 3542.9 | 0.86 | 0.53 | 4.73 |
| BARDC-31 | 4633.3 | 5808.3 | 1175.0 | 5220.8 | 2.48 | 5187.6 | 1.54 | 1.69 | 1.89 | 5154.7 | 1.06 | 1.02 | 8.89 |
| BARDC-32 | 3575.0 | 4216.7 | 641.7 | 3895.9 | 1.86 | 3882.6 | 0.86 | 0.50 | 0.63 | 3869.4 | 1.12 | 0.84 | 12.14 |
| BARDC-33 | 3300.0 | 4558.3 | 1258.3 | 3929.2 | 1.86 | 3878.5 | 0.86 | 0.58 | 0.54 | 3828.4 | 0.96 | 0.66 | 6.25 |
| AZRI-96 | 3058.3 | 4383.3 | 1325.0 | 3720.8 | 1.75 | 3661.3 | 0.77 | 0.48 | 0.41 | 3602.8 | 0.93 | 0.59 | 5.62 |
| Local White | 3916.7 | 4566.7 | 650.0 | 4241.7 | 2.03 | 4229.2 | 1.02 | 0.69 | 0.90 | 4216.8 | 1.14 | 0.93 | 13.05 |
| Pishin local | 4125.0 | 5133.3 | 1008.3 | 4629.2 | 2.20 | 4601.6 | 1.21 | 1.04 | 1.18 | 4574.2 | 1.07 | 0.91 | 9.18 |
| Shalkot-14 | 3987.5 | 4591.7 | 604.2 | 4289.6 | 2.05 | 4278.9 | 1.05 | 0.72 | 0.95 | 4268.3 | 1.15 | 0.95 | 14.20 |
| Tijaban-10 | 3875.0 | 4516.7 | 641.7 | 4195.9 | 2.01 | 4183.6 | 1.00 | 0.66 | 0.86 | 4171.3 | 1.14 | 0.92 | 13.08 |
| Wadan | 3920.8 | 4475.0 | 554.2 | 4197.9 | 2.01 | 4188.7 | 1.00 | 0.65 | 0.88 | 4179.6 | 1.16 | 0.95 | 15.15 |
| Zarghoon-21 | 3800.0 | 4800.0 | 1000.0 | 4300.0 | 2.04 | 4270.8 | 1.04 | 0.78 | 0.86 | 4241.9 | 1.05 | 0.83 | 8.60 |
| LSD (0.05) | 1659.7 | 1653.4 | 2262.1 | 1210.3 | 0.59 | 1238.3 | 0.60 | 0.98 | 1.37 | 1288.4 | 0.61 | 0.80 | 43.97 |

Conversely, genotypes BARDC-16, BARDC-17, and BARDC-2 exhibited the lowest values across these indices. For example, BARDC-16 showed the lowest GMP (3374.1), HARM (3205.0), and STI (1.64), along with the highest yield reduction (TOL = 2220.8). STI, MSTik1, MSTik2, and REI effectively differentiated the high-yielding genotypes. Genotypes BARDC-26, BARDC-31, BARDC-9, and BARDC-24 recorded higher values for MSTik1 (2.14, 1.69, 1.73, 1.75, respectively) and REI (>1.45), confirming their drought resilience and yield stability. Interestingly, check varieties such as Zarghoon-21, Wadan, and Shalkot-14 also demonstrated good performance under both conditions, suggesting their continued relevance in breeding programs and varietal recommendations for the region. For instance, Zarghoon-21 showed a GMP of 4270.8 and HARM of 4241.9, indicating good adaptability to semi-arid environments. In contrast, genotypes such as BARDC-17, despite showing high yield potential under nonstress conditions (Ypi = 5408.3), recorded poor values for drought indices (e.g., MSTik2 = 0.33, REI = 0.82), highlighting their susceptibility to water deficit. Traits such as GMP and STI are especially valuable because they combine productivity with stability, aiding in the identification of genotypes with broad adaptability. This aligns with the breeding goal of developing drought-resilient cultivars that not only survive but also thrive in water-limited environments. These findings are particularly relevant for semi-arid and drought-prone regions, such as Balochistan, where rainfed agriculture predominates. The identified high-performing genotypes can serve as potential donors in wheat improvement

programs aimed at enhancing drought resilience.

Table 5

Top-Performing Wheat Genotypes (Top 15%) Identified for Each Drought Tolerance Index under Irrigated (Ypi) and Stress (Ysi) Conditions.

| Index | Top 15% Genotypes |
|--------------|-------------------------------------|
| Ysi | BARDC-31, 26, 24, 9, 25, 27 |
| Ypi | BARDC-26, 14, 24, 9, 19, 31 |
| TOL (Bottom) | BARDC-5, 20, 23, 22, 15, 3 |
| MP | BARDC-26, 31, 24, 9, 14, 19 |
| MRP | BARDC-26, 31, 24, 9, 14, 19 |
| GMP | BARDC-26, 31, 24, 9, 14, 19 |
| REI | BARDC-26, 31, 24, 9, 14, 19 |
| MSTIK1 | BARDC-26, 14, 24, 9, 31, 19 |
| MSTIK2 | BARDC-31, 26, 24, 9, 14, 25 |
| HARM | BARDC-26, 31, 24, 9, 14, 19 |
| RDI | BARDC-20, 5, 22, 23, 15, 3 |
| DI | BARDC-20, 31, 3, 22, 23, Shalkot-14 |
| GM | BARDC-20, 5, 22, 23, 15, 3 |

Correlation Analysis

Correlation analysis among grain yield under stress (Ysi), yield under non-stress (Ypi), and the even drought tolerance indices provides valuable insight into the interrelationship of traits and their effectiveness in genotype evaluation under contrasting environmental conditions. Grain yield under Ysi exhibited strong and highly significant positive correlations with several drought indices MSTIK2 ($r = 0.929$), HARM ($r = 0.917$), GMP ($r = 0.879$), MRP ($r = 0.872$), DI ($r = 0.870$), REI ($r = 0.868$), and MP ($r = 0.829$). These strong associations highlight the relevance of these indices in reliably

predicting genotype performance under water-limited conditions in the future. Specifically, indices such as HARM, GMP, and DI are recognized for their capacity to stabilize performance under stress by balancing yield level and stress tolerance. In contrast, grain yield under non-stress conditions (Ypi) also exhibited significant positive correlations with indices such as MSTIK1 (r = 0.940), MP (r = 0.894), REI (r = 0.855), MRP (r = 0.855), GMP (r = 0.847), HARM (r = 0.796), and MSTIK2 (r = 0.744) (Figure 1). These correlations emphasize the consistency of these indices in selecting genotypes with superior yield potential under optimal (irrigated) conditions. Although a moderate positive correlation was observed between Ysi and Ypi (r = 0.493), indicating some shared influences, their responses differed under both conditions. This indicates that while some genotypes may perform well under both conditions, many exhibit divergent responses due to genotype × environment interactions. Interestingly,

TOL was negatively correlated with Ysi (r = -0.33) and positively correlated with Ypi (r = 0.656), suggesting that TOL tends to favor high-yielding genotypes under non-stress conditions but penalizes those with stability under drought conditions. Indices such as the RDI and GM exhibited opposing correlation trends: both were positively correlated with Ysi but negatively with Ypi. This pattern further underscores the stress-specific utility of these genes and highlights the trade-offs between drought adaptability and optimal yield expression. Conversely, indices such as MP, GMP, MRP, HARM, and REI maintained strong and positive correlations with both Ysi and Ypi, underscoring their robustness in different environments. Their dual sensitivity to performance in both moisture regimes makes them valuable tools for selecting genotypes with broad adaptability, which is a critical breeding objective in regions prone to fluctuating water availabilities.

Figure 1

Correlation Between Grain Yield (Ypi and Ysi) and Different Stress Indices of Wheat Genotypes.

| | Ysi | Ypi | TOL | MP | MRP | GMP | REI | MSTIK1 | MSTIK2 | HARM | RDI | DI | GM |
|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|-------|-------|----|
| Ysi | 1 | | | | | | | | | | | | |
| Ypi | 0.493 | 1 | | | | | | | | | | | |
| TOL | -0.332 | 0.656 | 1 | | | | | | | | | | |
| MP | 0.830 | 0.895 | 0.251 | 1 | | | | | | | | | |
| MRP | 0.873 | 0.855 | 0.170 | 0.997 | 1 | | | | | | | | |
| GMP | 0.879 | 0.848 | 0.157 | 0.995 | 0.999 | 1 | | | | | | | |
| REI | 0.868 | 0.856 | 0.175 | 0.995 | 0.998 | 0.998 | 1 | | | | | | |
| MSTIK1 | 0.701 | 0.941 | 0.412 | 0.963 | 0.945 | 0.943 | 0.957 | 1 | | | | | |
| MSTIK2 | 0.930 | 0.745 | 0.001 | 0.955 | 0.972 | 0.974 | 0.979 | 0.902 | 1 | | | | |
| HARM | 0.918 | 0.796 | 0.067 | 0.982 | 0.994 | 0.996 | 0.992 | 0.915 | 0.984 | 1 | | | |
| RDI | 0.529 | -0.471 | -0.970 | -0.031 | 0.051 | 0.063 | 0.045 | -0.197 | 0.214 | 0.151 | 1 | | |
| DI | 0.871 | 0.009 | -0.746 | 0.453 | 0.524 | 0.533 | 0.515 | 0.275 | 0.650 | 0.604 | 0.873 | 1 | |
| GM | 0.258 | -0.556 | -0.827 | -0.224 | -0.158 | -0.157 | -0.171 | -0.350 | -0.023 | -0.091 | 0.837 | 0.642 | 1 |

Principal Component Analysis

PCA was conducted to explore the multivariate relationships among the drought tolerance indices and to identify the most influential components contributing to the variability in genotype performance. The PCA output is summarized in Table 6, which includes both the component summary and the eigenvalues. The analysis revealed that the first two principal components (PC1 and PC2) had eigenvalues greater than 1 and together explained a substantial 97.7% of the total variation among the traits studied. Specifically, PC1 accounted for 66.6% of the total variance, whereas PC2 contributed an additional 31.1% (Figure 2). This cumulative contribution indicates that most of the variation in drought tolerance among genotypes can be effectively captured in a two-dimensional space. The eigenvalue distribution further supported this result, showing a steep decline beyond PC2, with the third component (PC3) contributing only 1.82% of the variance, and subsequent components contributing less than 1% of the variance each. This clear inflection confirms that only PC1 and PC2 are meaningful for genotype differentiation and trait clustering, whereas the

remaining components represent noise or redundancy. From a biological standpoint, the dominance of PC1 suggests that it likely represents a general adaptability axis, incorporating indices associated with overall yield performance and stability across environments, such as MP, GMP, HARM, and Stress STI. These indices are known to strongly correlate with both yield under irrigated and rainfed conditions and have been widely used for the selection of genotypes with broad adaptability. In contrast, PC2 may reflect the differential stress response or specific drought adaptation, capturing variability associated with indices such as TOL, DI, and GM. These indices tend to highlight genotype behavior under stress more explicitly, thus providing complementary insights into the broader patterns observed in PC1. The high explanatory power of these two principal components validated the internal consistency of the drought tolerance indices and confirmed their utility in genotype selection. This also implies that visualizing the genotypes and indices using a biplot based on PC1 and PC2 offers a clear and interpretable clustering of genotypes into tolerant, susceptible, and stable groups. Overall, PCA was an

effective statistical tool for data reduction, trait correlation analysis, and selection strategy development in this study.

Figure 2

PCA Biplot Showing the Distribution of Wheat Genotypes and Drought Tolerance Indices Based on the First Two Principal Components.

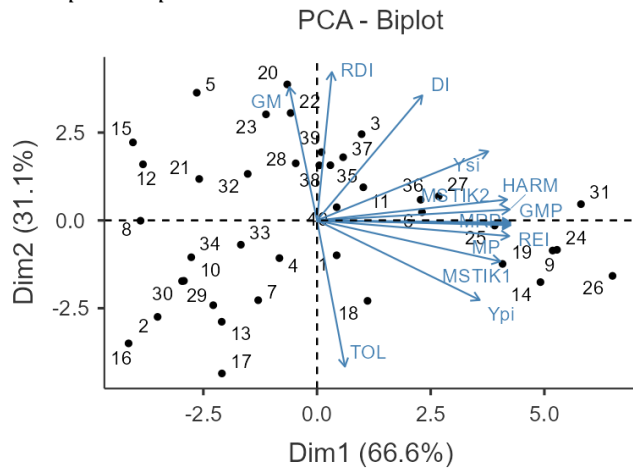


Table 6

Principal Component Analysis Summary Showing Eigenvalues and Variance Explained by the First Two Components for Drought Tolerance Indices.

| Component | Eigenvalue | % of Variance | Cumulative % |
|-----------|------------|---------------|--------------|
| 1 | 8.66131 | 66.62545 | 66.6 |
| 2 | 4.03952 | 31.07324 | 97.7 |
| 3 | 0.23721 | 1.82471 | 99.5 |
| 4 | 0.04662 | 0.35859 | 99.9 |
| 5 | 0.01162 | 0.08940 | 100.0 |
| 6 | 0.00346 | 0.02664 | 100.0 |
| 7 | 2.34e-4 | 0.00180 | 100.0 |
| 8 | 2.00e-5 | 1.53e-4 | 100.0 |
| 9 | 1.86e-6 | 1.43e-5 | 100.0 |
| 10 | 1.23e-9 | 9.49e-9 | 100.0 |
| 11 | 1.39e-15 | 1.07e-14 | 100.0 |
| 12 | -1.29e-16 | 9.95e-16 | 100.0 |
| 13 | -1.70e-16 | 1.31e-15 | 100.0 |

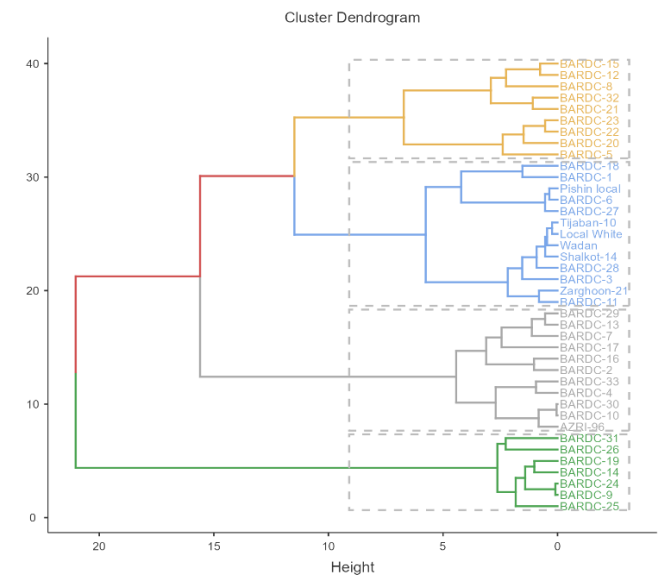
Hierarchical Cluster Analysis

Hierarchical cluster analysis was performed using Ward’s method based on grain yield under irrigation (Ypi), stress (Ysi), and associated drought tolerance indices. The resulting dendrogram (Figure 3) grouped the 40 wheat genotypes into four distinct clusters comprising seven, 11, 13, and nine genotypes, respectively. This classification reflects the genetic diversity and differential responses of the genotypes to drought stress. Cluster I included BARDC-31, BARDC-26, BARDC-19, BARDC-14, BARDC-24, BARDC-9, and BARDC-25. These genotypes showed consistently high grain yields under both stress and non-stress conditions, as well as superior values for key indices, such as GMP, MP, HARM, and GM, indicating their broad adaptability and strong drought tolerance. Moderate TOL values suggest manageable yield reduction under stress, whereas high GM values confirm yield stability. These genotypes align with Group A classification, which includes genotypes that are both drought tolerant and high yielding across environments. Hence, Cluster I genotypes are considered the most promising for cultivation in both rainfed and irrigated areas. Cluster II comprised genotypes

BARDC-29, BARDC-13, BARDC-7, BARDC-17, BARDC-16, BARDC-2, BARDC-33, BARDC-4, BARDC-30, BARDC-10, and AZRI-96. This cluster had the lowest Ysi and GMP values, along with the highest TOL and reduced REI, reflecting a poor yield performance under drought conditions.

Figure 3

Hierarchical Clustering Dendrogram of Wheat Genotypes by Cluster Analysis Based on Drought Tolerance Indices and Yield Performance in Rainfed and Irrigated Environments.



These genotypes are the most susceptible to drought, exhibiting high yield losses and limited stress resilience. Cluster III contained genotypes BARDC-18, BARDC-1, Pishin Local, BARDC-6, BARDC-27, Tijaban-10, Local White, Wadan, Shalkot-14, BARDC-28, BARDC-3, Zarghoon-21, and BARDC-11. These genotypes demonstrated moderate Ysi, MP, and GMP values, with an average DI of 0.88, indicating semi-tolerance. Although not as drought-resilient as Cluster I, these genotypes maintained reasonable productivity and could be useful in moderately stressed environments. Cluster IV grouped BARDC-15, BARDC-12, BARDC-8, BARDC-32, BARDC-21, BARDC-23, BARDC-22, BARDC-20, and BARDC-5. This cluster exhibited moderate drought indices, with Ysi, GMP, MP, and DI values slightly lower than those of Cluster III. These genotypes also fell into the semi-tolerant category, although with relatively lower stress tolerance and stability. The clustering pattern reflected clear intragroup homogeneity and intergroup variability, emphasizing that genotypes within each cluster shared similar drought response mechanisms. The dendrogram (Figure 3) clearly illustrates the genetic relationships and performance-based grouping of genotypes. This analysis confirmed the effectiveness of clustering based on drought indices for categorizing genotypes into distinct tolerance groups. Importantly, Cluster I genotypes are strongly recommended for cultivation in the drought-prone and semi-arid environments of Quetta and Balochistan because of their superior performance and yield stability across environments. The observed variability among clusters also highlights the potential for utilizing this genetic diversity in future breeding programs aimed at

improving drought resilience in wheat crops.

DISCUSSION

The present study demonstrates the effectiveness of multi-index technique in evaluating and distinguishing different wheat genotypes for drought tolerance under irrigated and rainfed conditions in the semi-arid environment of Quetta, Balochistan. By screening and characterizing 40 diverse wheat genotypes through 11 stress tolerance indices, we successfully categorized the genotypes into drought tolerant and drought susceptible groups, providing valuable insight for targeted wheat improvement (Singh *et al.*, 2018). The genotypes such as BARDC-26, BARDC-31, BARDC-24, BARDC-9, BARDC-14, and BARDC-19 exhibited a superior performance across indices such as MP, GMP, HARM, STI, MSTIK1, MSTIK2, and REI. Their consistent yield stability under both irrigated and rainfed conditions reflects a vigorous ability to maintain productivity despite water limitation, which is a primary breeding objective for wheat in drought-prone regions. This finding aligns with previous studies emphasizing the reliability of GMP, HARM, and STI in the accurate identification of high-yielding, stress-resilient cultivars (Yadav *et al.*, 2024; Sofi *et al.*, 2021). The correlation analysis revealed a close association between grain yield and key drought indices. Positive and significant correlations between Ysi, Ypi, and indices like MP, GMP, HARM, and REI indicate their suitability for differentiating genotypes with both high productivity and stability (Poudel *et al.*, 2021). The negative relationship between TOL and Ysi observed in this study emphasize that TOL alone may be less effective in selecting drought tolerant lines, because it might favor genotypes with high potential yield but limited stability (Bennani *et al.*, 2017). Moreover, the harmonic and geometric means served as reliable predictors for genotypic performance across variable environments, supporting their wide adoption in breeding programs (Alhag *et al.*, 2022; Chowdhury *et al.*, 2021). Furthermore, the principal component and cluster analyses further strengthen these findings. For example, PCA revealed that MP, GMP, and STI contributed most significantly to genotype differentiation, collectively explaining nearly all the variance among the tested genotypes (Sedghiyeh *et al.*, 2025; Shivramakrishnan *et al.*, 2016). Cluster analysis quickly described drought-tolerant and stable genotypes by grouping them together in Cluster I, thus offers an efficient strategy for breeding program advancement and future crossings (Thanana *et al.*, 2019; Bazzaz *et al.*, 2019). The performance of check varieties such as Zarghoon-21 and Wadan as well as high performing lines is of significance. The high and stable yields of their combined populations highlighted the presence of useful allelic variability in local gene pools that could be exploited in forward breeding (Spinoni *et al.*, 2015; Wen *et al.*, 2023). This is in line with worldwide interests in expanding the genetic base for stress tolerance in wheat by focused introgression and selection using molecular markers (Ahmed *et al.*, 2024). Physiological and agronomic, and probably genotypic-level superior drought tolerant cultivars, involve multiple adaptive traits, including root architectural improvement, osmotic adjustment or phenology optimization (Bapela *et al.*, 2022;

Alhag *et al.*, 2022). The retained yield potential of some genotypes under water deficit stresses suggests that they have the capacity for efficient water use and acclimation to stresses, in line with the mechanisms described by Al-Maskri *et al.* (2016) and Abhinandan *et al.* (2018). These traits will be major concerns in maintaining the productiveness of wheat under the prospect of the increase of the intensity of abiotic stress that may occur as a result of climate change (Rijal *et al.*, 2024). The evaluated genotypes stand out as exceptional parents for advancing new wheat varieties and for foundational breeding. By implementing multi-index statistical strategies, demonstrated here, we can swiftly and confidently identify lines that not only yield well under drought but are also broadly adaptable. Such methodologies are particularly suited to the pressing food-security issues in arid and semi-arid areas like Balochistan, where the primary agronomic target is stable production despite limited water availability (Bashir *et al.*, 2021; Blum, 2005). In addition, combining multi-index selection with cluster-based strategies creates a robust framework for enhancing drought tolerance in wheat. The standout genotypes singled out here serve not only as valuable breeding stock exhibiting immediate resilience, but they also strengthen long-term initiatives aimed at stabilizing wheat yields under rising climatic unpredictability (Devi *et al.*, 2024).

CONCLUSION

Our study evaluated 40 wheat genotypes under irrigated and rainfed conditions using drought tolerance indices, correlation analysis, PCA, and hierarchical clustering. By integrating these methods, we established a rapid and reliable framework for identifying genotypes with exceptional yield stability and drought resilience. Six genotypes (BARDC-26, BARDC-24, BARDC-9, BARDC-31, BARDC-25, and BARDC-19) consistently outperformed others, exhibiting high MP, GMP, HARM, and STI values, which underscore their adaptability across both stressed and non-stressed environments. Notably, BARDC-26 emerged as a top candidate, achieving the highest GMP, HARM, and STI scores, making it particularly valuable for drought-prone regions. Correlation analysis revealed key relationships: Ysi showed strong associations with MSTIK2, HARM, GMP, MRP, DI, REI, and MP, while Ypi correlated closely with MSTIK1, MP, GMP, MRP, and HARM. These findings emphasize the utility of multi-trait selection for ensuring stability in diverse environments. Among the indices, MP, GMP, HARM, and REI proved most reliable, displaying robust linkages with both Ysi and Ypi (a testament to their effectiveness in dual-environment screening). PCA further reinforced these insights, with the first two principal components explaining 97.7% of total variance (PC1: 66.6%; PC2: 31.1%). Yield-based indices MP, GMP, HARM, and STI were the primary drivers of this variation, solidifying their role in differentiating genotype performance under contrasting water regimes. Hierarchical clustering (Ward's method) classified the genotypes into four distinct groups. Cluster I (BARDC-31, BARDC-26, BARDC-19, BARDC-14, BARDC-24, BARDC-9, and BARDC-25) contained the most drought-tolerant and stable performers, while Clusters III and IV represented intermediate tolerance. In contrast, Cluster II grouped the

most susceptible genotypes, characterized by low Ysi, GMP, and high yield reduction under stress. In summary, our multi-analytical approach combining indices, correlation, PCA, and clustering provides a robust strategy for dissecting genotype behavior under drought. The elite

genotypes in Cluster I, particularly BARDC-26, hold significant promise for wheat breeding in water-limited regions like Balochistan. We recommend advancing these candidates to multi-location trials for potential release in semi-arid zones.

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