

Screening of salt-tolerance of maize varieties based on the value of the membership function and under GGE biplot analysis (#90694)

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Screening of salt-tolerance of maize varieties based on the value of the membership function and under GGE biplot analysis

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This study aimed to screen for highly salt-tolerant maize varieties using four NaCl treatments at concentrations of 0, 60, 120, and 180 mMol/L. Agronomic traits, physiological and biochemical indices related to salt tolerance were measured, and salt tolerance was evaluated using principal component analysis, membership function method, and GGE biplot analysis. A total of 41 local maize varieties were assessed based on their **D values**. Results show: (1) The results revealed significant differences in plant height, stem thickness, germ length, etc. among the different NaCl treatments. As the salt concentration increases, the lengths of embryo shoot and root, leaf area, etc. decreased, while electrical conductivity and salt damage index increased. (2) Correlations were found among most of the indicators, and principal component analysis identified eight main components. (3) In the GGE biplot analysis, Youqi 909, Xuanhe 8, and Qunze 888 exhibited superior salt tolerance and adaptability in salt stress environments. The optimal stress concentration was determined to be 120 mMol/L NaCl solution. (4) The top five varieties for salt tolerance and stability were Youqi 909, Xuanhe 8, Qunze 888, and Wugu 568, followed by New Jade 66. This study further validated the reliability of GGE biplot analysis in germplasm selection, expanded the genetic resources of salt-tolerant maize in Xinjiang, and provided theoretical references and germplasm utilization for introducing maize into saline-alkali areas in southern Xinjiang. These research findings contributed to improving our understanding of maize salt tolerance and promoting its cultivation in harsh environments.

Screening of Salt - tolerance of Maize Varieties based on the Value of the Membership Function and under GGE biplot analysis

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Abstract

This study aimed to screen for highly salt-tolerant maize varieties using four NaCl treatments at concentrations of 0, 60, 120, and 180 mMol/L. Agronomic traits, physiological and biochemical indices related to salt tolerance were measured, and salt tolerance was evaluated using principal component analysis, membership function method, and GGE biplot analysis. A total of 41 local maize varieties were assessed based on their D values. Results show: (1) The results revealed significant differences in plant height, stem thickness, germ length, etc. among the different NaCl treatments. As the salt concentration increases, the lengths of embryo shoot and root, leaf area, etc. decreased, while electrical conductivity and salt damage index increased. (2) Correlations were found among most of the indicators, and principal component analysis identified eight main components. (3) In the GGE biplot analysis, Youqi 909, Xuanhe 8, and Qunze 888 exhibited superior salt tolerance and adaptability in salt stress environments. The optimal stress concentration was determined to be 120 mMol/L NaCl solution. (4) The top five varieties for salt tolerance and stability were Youqi 909, Xuanhe 8, Qunze 888, and Wugu 568, followed by New Jade 66. This study further validated the reliability of GGE biplot analysis in germplasm selection, expanded the genetic resources of salt-tolerant maize in Xinjiang, and provided theoretical references and germplasm utilization for introducing maize into saline-alkali areas in southern Xinjiang. These research findings contributed to improving our understanding of maize salt tolerance and promoting its cultivation in harsh environments.

Introduction

The sustainability of global agriculture has been severely compromised by soil salinisation, resulting in significant reductions in crop yield and quality worldwide [1,2]. The area of saline land in the world is as much as 954 million hectares, and the area of saline land in China reaches 37 million hectares, accounting for 4.9% of arable land [3-4]. In the process of growth and development, salt stress causes ionic toxicity, osmotic stress, oxidation stress, etc., resulting in cell membrane damage, root drying, metabolic disorders, and reduced plant photosynthesis, which seriously affects plant growth. [5]. Maize (*Zea mays* L.), a widely cultivated crop globally, plays a central role in the economic development of China. It is a moderately saline crop with low salt tolerance and is severely affected by salt stress at the seedling stage. It is severely affected by salt stress at the seedling stage and is a moderately salt-sensitive crop with poor salt tolerance. It is of great significance to study the salt tolerance of maize varieties in China and its comprehensive evaluation methods and identification indexes for the breeding, production and application of salt-tolerant germplasm resources.

Previous research has shown that salt damage significantly impacts maize yield stability. It affects phenotypic traits, physiological and biochemical indices of maize, especially during reproductive stages like germination and seedling growth. Salt stress causes a significant reduction in seed germination and also leads to a reduction in plant growth rate and yield, while at the same time reducing the plant's water uptake capacity [6-8]. Increased salt concentration inhibited germination and early seedling growth in several rice varieties, as shown by Song J et al [9]. Salt-tolerant varieties tend to have higher peroxidase activity under salt stress conditions [10]. The conductance and malondialdehyde (MDA) levels of leaf in Maize increased significantly after salt stress, resulting in peroxidative damage to the cell plasma membrane, and plant proline (PRO) levels increased significantly after salt stress to improve plant growth [11-12]. The salt tolerance indicators selected for the different fertility periods and treatments were

also distinct. In previous studies, germination rate, germination potential, radicle length and radicle length have been used as indicators to reflect the salt tolerance of seeds at the germination stage from various perspectives [13-14]. Seedling condition, plant height and rate of change in dry weight have also been used as grading indices to identify salt tolerance in maize seedlings by previous authors [15].

Many domestic and foreign scholars have proposed identification methods based on salinity resistance, including the membership function method as one of the important methods to identify the salinity resistance of maize, and the higher value of the membership function indicates more salinity resistance [16]. Based on the membership function values, Yu Ying et al. used an affiliation function method to classify maize inbred lines into four categories of salinity sensitivity [16]. Zhang Haiyan and colleagues, based on the salt tolerance coefficient of germination and seedling stage identification index as the evaluation basis, divided the weights of each index according to the coefficient of variation of the value of the salt tolerance coefficient of the index membership function, and classified the salt tolerance of test maize into four levels by using the weighted membership function method [17]. GGE biplot analyses have been used to classify environments, to assess the ranking of genotypes, and to determine the discriminative and representative properties of environments [18]. It can be analysed in terms of crop yield and stability [19-20], identification of the ideal environment for crop growth [21], and screening and evaluation of germplasm resources [22-23]. The "GGE biplot Chart" graphically illustrates the relationships among varieties, identification indicators or assessment methods using auxiliary lines. It provides accurate and intuitive insights into the performance of different varieties in various environments, as well as the appropriate indicators for identification or assessment methods [24].

Although there have been numerous studies on the salt tolerance mechanisms of maize, most of the previous research focused on individual stages without considering other reproductive phases. Few studies have utilized GGE biplot analysis to investigate maize germination and seedling stages under salt stress. In this study, we simulated several agronomic traits and physiological indicators related to salt tolerance in maize using NaCl solutions of 0, 60, 120, and 180 mMol/L under different salt stress conditions, building upon previous work. By incorporating fuzzy membership function and GGE biplot analysis, we comprehensively evaluated the salt tolerance of 41 maize varieties. We assessed the overall salt tolerance of these varieties, identified appropriate concentrations and key evaluation indicators, and screened out more salt-tolerant maize varieties, providing guidance for the selection and promotion of maize varieties in Xinjiang's saline-alkali areas.

Materials & Methods

Plant materials

For this trial, 41 maize varieties popularized in production were selected as test material, and the specific material information was listed in Table 1.

Experimental Design

Germination test for seeds indoors

Germination experiments were conducted in an artificial climate chamber with a day/night temperature of 25°C/15°C (day/night) and 75% humidity at the The study took place at the Key Laboratory of Genetic Improvement and Efficient Production for Specialty Crops in Arid Southern Xinjiang of Xinjiang Corps in April-May 2022. The experiment was set up with a total of four concentration gradients of 0 (CK), 60, 120 and 180 mmol/L NaCl solutions and each treatment was replicated three times. The Petri dish was first sterilised with 70% alcohol, two layers of filter paper were placed in the dish and then the seeds were placed evenly in the dish. Incubation was carried out with four concentrations of NaCl solution, 0, 60, 120 and 180

mmol/L, and the solution in each Petri dish soaked more than 1/3 of the seeds. Each day, the Petri dish was filled with the appropriate amount of NaCl solution at the appropriate concentration. Seed germination was recorded at 16:00 h each day, seed germination potential was determined at 4 d of treatment and germination rate was determined after 7 d of treatment and relative germination potential, relative germination rate, salt tolerance index and salt injury index were calculated.

Tolerance test for potting salt outdoors

The study took place at the Tarim University Agronomy experiment station from May-July 2022. 25×25 seedling pots were used. The growing medium was fine sandy soil with a soil weight of 4.5 kg. Maize seeds of each cultivar were sown at a depth of about 3 cm with 10 seeds per pot, selected for uniform and consistent size of 120 grains each. When the seedlings grew to three leaves and one heart, they were treated with nutrient solution containing 1/4 concentration of Hoagland nutrient solution with different concentrations of salt solution, the treatment concentrations were 0 (CK), 60, 120 and 180 mmol/L NaCl solution and each treatment was repeated three times. After 7 days of treatment, the fresh and dry weight of each plant was determined, the root-crown ratio was calculated, conductivity was determined and physiological indices including MDA content, PRO content, SOD activity and POD activity were measured using the kit method.

Calculation formula

Based on the salt tolerance coefficient (salt tolerance coefficient = mean value of salt stress index/mean value of control index) of each identification index of the test varieties, 41 maize varieties were comprehensively evaluated for salt tolerance by using the subordinate function method in fuzzy mathematics, and the calculation formula was as follows

Coefficient of variation (CV) = (standard deviation SD/mean) × 100%.

The formula for calculating the value of the membership function is where X_i is the salt tolerance coefficient of each test material based on the identification index i , $X_{i \max}$ and $X_{i \min}$ are the maximum and minimum values of X_i in the test materials respectively, and $U(X_i)$ is the value of the membership function of each test material X_i .

Inverse membership function formula: $\mu(X_i) = 1 - (X_i - X_{i \min}) / (X_{i \max} - X_{i \min})$,

The weighting formula: $W_i = CV_i / \sum_{i=1}^n CV_i$ ($i = 1, 2, 3, \dots, n$). CV_i is the coefficient of variation of each test material $\mu(X_i)$ and W_i is the ratio of CV_i to the total variation.

Weighted membership function value formula: $D = \sum_{i=1}^n [\mu(X_i) \cdot W_i]$ ($i = 1, 2, 3, \dots, n$). $\mu(X_i)$ is the value of the membership function of each test material X_i and W_i is the weight of each identification index.

Data Analysis

Microsoft Excel was used for data processing, SPSS software for ANOVA and factor analysis, and Origin 2022 software for correlation analysis. The analysis of adaptability of cultivation varieties, discrimination analysis of NaCl concentration, representativeness analysis of NaCl concentration, and analysis of ideal salt-tolerant cultivated varieties were conducted using GenStat software for GGE biplot analysis. The analytical method partially referred to previous studies [25].

Results

Analysis of difference among salt tolerance indexes

An analysis of variance (ANOVA) was conducted on the phenotypic, physiological, and germination indices of 41 maize varieties (refer to Figures 1-3). Plant height, stem thickness, radicle length, leaf area, conductivity, chlorophyll SPAD, germination rate, water content, germination index, salt tolerance index, salt injury index, SOD activity, MDA content, and POD activity exhibited significant difference among the four NaCl concentration treatments. While the root crown ratio and PRO content did not show any significant difference. Germ length, radicle length, leaf area, germination rate, germination index, salt tolerance index, and germination potential decreased with increasing salt concentration, while conductivity and salt injury index increased. The increase in water content with salt concentration could be attributed to the continuous uptake of external water by the root system to maintain physiological metabolism balance under adverse conditions, facilitating the survival and growth of seedlings under salt stress. Plant POD activity was an important indicator of resistance, reflecting the ability of plants to scavenge reactive oxygen radicals. [26-27]. In this study, the POD activity of maize leaves differed significantly from the control group under 60 mMol NaCl treatment, but the changes were minimal as the NaCl concentration increased. This finding contradicted a previous study [28] and may be attributed to variations in POD activity levels among different varieties under NaCl treatment. The SOD activity of antioxidative enzymes in the functional leaves of 41 salt-tolerant maize varieties decreased and the MDA content increased under 120 mM NaCl treatment. However, with higher concentrations of NaCl treatment, the SOD activity increased and the MDA content decreased. This indicates that the ability to remove reactive oxygen species (ROS) generated during normal growth and development could be enhanced at specific salt concentrations. Subsequently, by clearing ROS and reducing lipid peroxidation, it mitigates plant damage and enhances their adaptation to high salt stress at higher concentrations.

Analysis of the correlation among the salt tolerance indicators

Correlation analyses were conducted among the 18 indicators, and the relationships among these indicators were depicted in Figure 4. Within the phenotypic traits, plant height exhibited significant positive correlations with stem thickness, leaf area, germ length, radicle length, chlorophyll SPAD, and salt tolerance index; while displaying negative correlations with water content, root crown ratio, and conductivity. Leaf area displayed a negative correlation with root crown ratio. Water content showed a negative correlation with germ length, radicle length, and SOD activity. The content of MDA was negatively correlated with both SOD activity and chlorophyll SPAD. Additionally, SOD activity exhibited a negative correlation with POD activity. Notably, there were significant positive correlations observed among germination potential, germination percentage, germination index, salt tolerance index, germ length, and radicle length.

There was no significant correlation observed between conductivity and the activities of SOD and POD in the physiological indices. Under adverse stress conditions, plants commonly respond by increasing POD activity in their tissues [29]. Conductivity served as an indicator of leaf damage severity caused by salt stress, and it was expected that as leaf damage increased, POD activity would also increase. However, contrary to expectations, this study found that there was no significant correlation between POD activity and conductivity. This discrepancy could be attributed to the continuous adversity that the plants experienced during this period, leading to a saturation point in POD activity, increased membrane lipid peroxidation, disruption of normal cell metabolism, leaf damage, and inhibition of plant growth. Therefore, our research findings deviated from previous studies.

Principal component analysis and membership function analysis of salt tolerance index

Factorial analyses were conducted on the 18 indicators, as presented in Table 2. Based on a cumulative contribution exceeding 80%, eight principal components were extracted. The eigenvalue of the first principal component was 5.141. The germination index had the highest

loading value and was therefore referred to as the salt tolerance index factor. Principal component 2 had an eigenvalue of 2.825 and was called the germ length factor. The plant height factor was assigned to principal component 3, which had an eigenvalue of 1.541. The SOD activity factor was assigned to principal component 4, which had an eigenvalue of 1.472. Principal component 5, named the chlorophyll SPAD factor, possessed an eigenvalue of 1.067. Similarly, the sixth principal component, with an eigenvalue of 1.006, was associated with the highest loading value of MDA content, called the MDA content factor. Principal component 7 had an eigenvalue of 0.941 and was called the POD activity factor. Principal component 8, known as the PRO content factor, had an eigenvalue of 0.781.

The eight factors obtained from the principal component analysis were comprehensively evaluated by membership function, including the germination index, chlorophyll SPAD, germ length, SOD activity, plant height, POD activity, PRO content and MDA content. The D values for these factors were calculated using the membership function method and further computed for the four NaCl concentrations listed in Table 4. In the absence of treatment, cultivar no. 23 exhibited the highest D value, while under a 60 mM NaCl solution treatment, cultivar no. 16 had the highest D value. Conversely, cultivar no. 12 showed the highest D value in 120 mM NaCl solution treatments. Meanwhile, cultivar no. 18 showed the highest D value in 180 mM NaCl solution treatments. It is important to note that a higher value of the membership function indicates greater salt tolerance. Therefore, among all cultivars, cultivar no. 23 displayed the highest salt tolerance in the 60 mM NaCl solution treatment, cultivar no. 12 exhibited higher salt tolerance in the 120 mM NaCl solution treatment, and cultivar no. 18 showed higher salt tolerance compared to other cultivars in the 180 mM NaCl solution treatment.

Salt tolerance or adaptation of different maize varieties on the basis of the analysis of the GGE biplot

The D values of the test varieties under four stress concentrations were analyzed using GGE biplot analysis. The first principal component accounted for 56.62% of the total variation, while the second principal component explained 20.12% of the variation. The GGE biplot was employed to visualize varietal adaptation functions, where sectors represented the growth environments under the stress of the four NaCl solutions. The maize varieties for trial number 8, 38, 18, 33, 23, and 35 located in the apex of the sector area. The polygon was divided into six sectors, with the four environmental NaCl stress concentrations distributed in two sectors. C2, C3, and C4 resided in the first sector, whereas C1 was situated in the second sector. Cultivar no. 23 exhibited closer proximity to the C1 region, suggesting its superior adaptation to the C1 environment compared to other varieties. On the other hand, cultivar no. 18, 16, and 12 demonstrated a higher adaptability to high salt stress conditions, as they were positioned closer to regions C2, C3, and C4 respectively. This observation implies that these three varieties possessed relatively higher salt tolerance under salt stress conditions when compared to the remaining 38 varieties.

Representativeness and discriminatory power of different concentrations of NaCl

An important factor in assessing the suitability of an environment under NaCl stress concentration was its discriminatory power and representativeness. In the GGE biplot analysis, circles connected the mean environment axis and mean environment value. The smaller the circle associated with a stress concentration environment point, the higher the overall level of that particular environment. It was evident that the comprehensive ranking of the four stress environments was $C3 > C2 > C4 > C1$ (Figure 6). This implied that C3 exhibited a stronger ability to identify the salt tolerance of maize germplasm, while C1 had the weakest ability to do so. It should be noted that C1 served as the control environment, which explained its limited capacity to identify the salt tolerance of maize.

The environments under different stress concentrations were analyzed and depicted (Figure 7). The angle among the environment under C1 stress concentration and the environments under C2, C3, and C4 stress concentrations was found to be less than 90° , indicating a strong positive correlation. This suggests that the ranking of varieties was similar across the four stress concentrations. In terms of representativeness, the environment under C2 stress concentration showed the best representation, as it had the smallest angle to the mean axis. On the other hand, the C4 environment had the longest line segment, indicating its strong discriminatory power for varieties.

Furthermore, the analysis of variance for the test maize's identification indexes (Figures 1-3) revealed a significant downward trend. There was also a significant distinction among multiple varieties and concentrations. These results suggest that the measured indexes at 160 mmol/L NaCl concentration can effectively identify the salt tolerance of different maize varieties. Thus, this concentration can be considered an ideal environment for determining maize varieties' salt tolerance, providing strong persuasiveness and credibility. Moreover, the C2 environment indicated less discrimination among varieties, as it had the shortest line segment. Overall, the environments at the four stress concentrations can be divided into two regions: one comprising C2, C3, and C4, and another comprising C1.

Analysis of the salt tolerance of different maize varieties

The central point of the circle on the environmental axis represents the average stress concentration. Combined with this experiment, the varieties closer to the center circle exhibit the best overall performance in terms of salt tolerance and stability. The top six varieties in terms of combined salt tolerance and stability were ranked as follows: 18>12>16>21>32>40 (Figure 8). On the other hand, the bottom six varieties in terms of combined salt tolerance and stability were ranked as follows: 8<35<38<13<23<34. When comparing the variety rankings obtained from the GGE biplot analysis with the membership function values calculated using the affiliation function method, it was found that after conducting the GGE dual standard analysis of the membership function values, the stability of the varieties could be further evaluated based on the assessment of salt tolerance in different maize varieties. This facilitates the identification of salt-tolerant germplasm resources with high potential stability.

Discussion

Effect of salt stress on maize phenotypic and physiological indicators

Salinity stress is a significant abiotic factor that greatly affects maize yields in maize-producing countries. It is crucial to develop and utilize crop varieties adapted to saline soils in order to address this issue [30]. With the increase of soil salinity and the expansion of saline-alkali land, it is urgent to screen and cultivate maize varieties with enhanced salt-tolerance. [17]. Salt stress has negative impacts on crucial plant processes, impedes growth and development, and disrupts cellular structures [31-32].

Previous studies on maize salt tolerance mainly focused on individual growing seasons, without considering multiple reproductive periods. Furthermore, results may be various when diverse germplasm were employed. For instance, Fu et al. [33] found that a concentration of 100 mMol/L NaCl promoted maize seed germination, while treatments with 200 mMol/L NaCl and higher concentrations inhibited it. In our experiment, various indicators related to seed germination, such as germ length, germination rate, and germination index, decreased with increasing salt concentration. The NaCl solution impeded maize seed germination. It is important to note that the results of this study may be biased due to the use of different germplasm. Nevertheless, it aligns with the finding that high concentrations of NaCl solution inhibit maize germination. When we analyzed the combined germination and seedling stages, most of the evaluation indices for all tested maize varieties exhibited a significant declining trend at NaCl concentrations ≥ 120 mMol/L. In conclusion, the index measured at 120 mMol/L NaCl

concentration could effectively identify the salt tolerance of different maize varieties. These findings contradict previous conclusions that reported inconsistent levels of salt tolerance among different maize varieties across different test materials and measurement periods. It was speculated that the different salt tolerance exhibited by the same plant in different growth stages arose from distinct underlying mechanisms [33].

Screening of maize salt-tolerant germplasm by means of the membership function method and the GGE biplot analysis

The GGE biplot analysis method encompassed the assessment of genotype and genotype-by-environment interactions [35]. These scatterplots visually represented bidirectional data, with genotypes as inputs and environments as outputs. This method is frequently used to describe the super-environment, rank genotypes, and determine stable environments. [36]. Furthermore, we found that this method has been widely utilized in assessing the adaptability and productivity of agricultural crops across multiple experimental sites. Previous researchers utilized GGE biplot analysis to investigate heat tolerance in wheat by analyzing late yield and multi-point trials [37, 38]. Similarly, this analysis was commonly utilized in adaptation and stability analyses for crops such as sorghum, sugarcane, and soybean [36, 39, 40]. Most prior studies applied the affiliation function method to calculate affiliation function values, which evaluated the comprehensive performance of plants under abiotic stresses such as drought tolerance, salt tolerance, and cold tolerance. Furthermore, we discovered that numerous studies employed a combination of analytical approaches, including throughput and hierarchical analyses, along with GGE biplot analysis, to evaluate diverse varieties. However, no previous study had evaluated the resistance of varieties by combining the membership function values under the membership function method with GGE biplot analysis [41-42]. In our study, we employed GGE biplot analysis to analyze the membership function values at four concentrations of NaCl solutions. This approach further enhanced the accuracy and reliability of the analyses compared to the observed results. In the GGE biplot polygons (Figure 5), varieties were positioned within specific environments, with genotypes represented at the top. This positioning indicated the superior performance of these genotypes in those environments [43,44]. The angle among environment vectors in GGE biplot analysis reflected their correlation, with a smaller angle indicating a higher correlation [45]. The length of the environment vectors approximated the standard deviation within each environment and served as an indicator of environmental distinctiveness. Consequently, environments with longer vector lengths had larger standard deviations and were more distinguishable [44, 46]. In our study (Figure 7), maize membership function values under NaCl solution stress and their corresponding environments were analyzed. The environment with a concentration of 160 mMol/L NaCl exhibited the longest vector line segment, followed by the environment with a concentration of 180 mMol/L NaCl. The small angle among these two segments indicated a positive correlation with better salt tolerance. Both of these environments were ideal for screening maize varieties for salt tolerance [46]. In summary, the three varieties examined in this study, namely Youqi 909, Xuanhe 8, and Qunze 888, demonstrated excellent adaptability to saline-alkali soil.

These findings provided significant insights for further investigations into the adaptability of maize in saline-alkali soil. Moreover, our study underscored the efficacy of utilizing GGE biplot analysis to assess the potential salt tolerance of maize varieties. A comprehensive evaluation of maize performance under specific environmental conditions was conducted, taking into account factors such as inter-environmental correlation, standard deviations, and vector lengths. This bore considerable significance in the selection of high-quality maize varieties that were well-suited to saline-alkali soil, ultimately leading to improved productivity and quality within maize cultivation in these regions.

Nevertheless, it was imperative to conduct subsequent research to validate these findings and expand the scope of the sample size. Such endeavors would facilitate a more profound understanding of the underlying mechanisms governing maize's salt tolerance and the genetic traits that contributed to adaptation under diverse saline-alkali soil conditions. These efforts would contribute to the establishment of a robust theoretical foundation for maize cultivating practices, thereby fostering sustainable agricultural development in saline-alkali regions.

Significance of screening salinity tolerant germplasm resources for future cultivating and development of salinity tolerant crops

Land salinization became a prominent issue that hindered ecological and environmental development as well as economic progress. The development of Salt Lake agriculture carried substantial practical significance for Chinese food industry, guaranteeing efficient food supply and maintaining the ecological equilibrium of the region [47]. Therefore, cultivating salt-alkali-tolerant plant varieties and enhancing the salt-alkali tolerance of plants proved to be an effective biological measure in alleviating the impact of saline-alkali land on plants. It also generated favorable ecological and economic benefits, promoting the sustainable development of agriculture [48]. In many countries, plant salt tolerance tests and cultivation of salt-tolerant plants are used to improve saline-alkali land and develop its potential. At that time, over 100 countries grappled with saline soils, including Chinese major grain-producing areas in the northwest, north, and northeast, where grain production and quality were affected. Therefore, apart from soil improvement measures, cultivating salt-tolerant crop varieties became imperative due to their absence, which was the leading cause of low maize yields in saline soils across China [49]. It was worth noting that there was a lack of systematic research on the response of plants in Xinjiang to soil salinity and the stress resistance of unique plant species [50]. Therefore, it has profound significance to evaluate the salt-tolerance of maize varieties comprehensively, screen the salt-tolerant varieties, and cultivate the salt-tolerant germplasm suitable for saline-alkali soil.

Conclusions

Given the increasing soil salinity levels, identifying maize genotypes with high salt tolerance has become crucial for expanding maize cultivation on saline-alkali soil, while also enhancing both its quality and yield potential. The objective of this study was to further validate the reliability of GGE biplot analysis in selecting maize germplasm through the utilization of the Membership Function method combined with GGE biplot analysis. Through our research, we identified five suitable maize genotypes, namely Youqi 909, Xuanhe 8, Qunze 888, Wugu 568, and New Jade 66, which demonstrate promising performance for planting in saline-alkali soil in Xinjiang. These findings provide a theoretical foundation for the cultivation and promotion of maize genotypes in other regions of Xinjiang. This study contributes to the field by confirming the effectiveness of the combined approach of GGE biplot analysis and the Membership Function method in the selection of salt-tolerant maize germplasm. Future studies could focus on exploring the salt tolerance of other maize germplasm resources, improving the GGE biplot analysis method, and investigating alternative cultivation techniques applicable to saline-alkali soil conditions. Such research endeavors will further advance our understanding of maize cultivation in challenging environments and pave the way for sustainable agriculture practices.

Acknowledgements

The authors are grateful to the anonymous reviewers for their valuable comments and suggestions.

References

- Jiang, D.; Lu, B.; Liu, L.T.; Duan, W.J.; Chen, L.; Li, J.; Zhang, K.; Sun, H.C.; Zhang, Y.J.; Dong, H.Z.; Li, C.D.; Bai, Z.Y. Exogenous melatonin improves salt stress adaptation of cotton seedlings by regulating active oxygen metabolism. *PeerJ* **2020**, *8*, e10486-e10486.
- Neupane, D.; Adhikari, P.; Bhattarai, D.; Rana, B.; Ahmed, Z.; Sharma, U.; Adhikari, D. Does Climate Change Affect the Yield of the Top Three Cereals and Food Security in the World? *Earth*. **2022**, *3*, 45-71.
- Li, W.Y.; Wang, C.; Shi, H.H.; Wang, B.; Wang, J.X.; Liu, Y.S.; Ma, J.Y.; Tian, S.Y.; Zhang, Y.W. Genome-wide analysis of ethylene-response factor family in adzuki bean and functional determination of VaERF3 under saline-alkaline stress. *Plant Physiol Biochem*. **2020**, *147*, 215-222.
- Luo, J.Y.; Zhang, S.; Zhu, X.Z.; Ji, J.C.; Zhang, K.X.; Wang, C.Y.; Zhang, L.J.; Wang, L.; Cui, J.J. Effect of NaCl-stressed *Bacillus thuringiensis* (Bt) cotton on the feeding behaviors and nutritional parameters of *Helicoverpa armigera*. *PLoS One*. **2018**, *9*, e0198570.
- Gupta, A.; Rico-Medina, A.; Cano-Delgado, A. I. The physiology of plant responses to drought. *Science (New York, N.Y.)*, **2020**, *368*, 266-269.
- Ghatak, A.; Chaturvedi, P.; Nagler, M.; Roustan, V.; Lyon, D.; Bachmann, G.; Postl, W.; Schröfl, A.; Desai, N.; Varshney, R. K.; Weckwerth, W. Comprehensive tissue-specific proteome analysis of drought stress responses in *Pennisetum glaucum* (L.) R. Br. (Pearl millet). *Journal of proteomics*. **2016**, *143*, 122-135.
- Foolad, M.R.; Hyman, J.R.; Lin, G.Y. Relationships between cold-and salt-tolerance during seed germination in tomato: Analysis of response and correlated response to selection. *Plant Breeding*. **1999**, *118*, 49-52.
- Carpýcý, E.B.; Celýk, N.; & Bayram, G. Effects of salt stress on germination of some maize (*Zea mays* L.) cultivars. *African Journal of Biotechnology*. **2009**, *8*, 4918-4922.
- Song, J.; Fan, H.; Zhao, Y.; Jia, Y.; Du, X.; Wang, B. Effect of salinity on germination, seedling emergence, seedling growth and ion accumulation of a euhalophyte *Suaeda salsa* in an intertidal zone and on saline inland. *Aquatic Botany*. **2008**, *88*, 331-337.
- Wang, J.Z.; Gao, S.R.; Sun, L.F.; Wang, X.; Wang, J.; Hu, K.F.; Deng, J. Physiological responses and tolerance of three maize inbred lines to salt stress. *Agricultural Research in the Arid Areas*. **2017**, *35*, 89-95.
- Rubio, M. C.; Bustos-Sanmamed, P.; Clemente, M. R.; & Becana, M. Effects of salt stress on the expression of antioxidant genes and proteins in the model legume *Lotus japonicus*. *The New phytologist*. **2009**, *181*, 851-859.
- Ben Ahmed, C.; Ben Rouina, B.; Sensoy, S.; Boukhriss, M.; Ben Abdullah, F. Exogenous proline effects on photosynthetic performance and antioxidant defense system of young olive tree. *Journal of agricultural and food chemistry*. **2010**, *58*, 4216-4222.
- Zhao, X. Q.; Peng, Y. L.; Li, J.Y.; Ren, X.W. Comprehensive evaluation of salt tolerance in 16 Maize Inbred Lines. *Agricultural Research in the Arid Areas*. **2014**, *32*, 40-45+51.
- Qi, T.; Huang, J.; Xu, F.; Han, D.X. Effects of Compound Salt Stress on Germination of 7 Maize Varieties. *Xinjiang Agricultural Sciences*. **2022**, *59*, 1855-1863.
- Fu, Y.; Gao, S.R.; Wang, Z.H. Evaluation of Salt Tolerance of Maize Germplasm in Seedling Stage. *Journal of Maize Sciences*. **2009**, *17*, 36-39+50+2.

16. Yu, Y.; Zhang, S.Q.; Guo, Y.L.; Yao, Y.B.; Zhou, F.; Zhao, D.S.; Liu, J.Z.; Li, W.H. Comprehensive evaluation of saline - alkaline tolerance of 31 maize inbred lines at germination stage. *Journal of Northeast Agricultural University*. **2018**, *49*, 9-19.
17. Zhang, H.Y.; Zhao, H.J. Comprehensive Evaluation of Salt Tolerance of Different Corn Varieties at the Germination and Seedling Stages. *Journal of Maize Sciences*. **2016**, *24*, 61-67.
18. Oladosu, Y.; Rafii, M. Y.; Abdullah, N.; Magaji, U.; Miah, G., Hussin, G.; Ramli, A. Genotype× Environment interaction and stability analyses of yield and yield components of established and mutant rice genotypes tested in multiple locations in Malaysia. *Acta Agriculturae Scandinavica, Section B—Soil & Plant Science*. **2017**, *67*, 590-606.
19. Yu, Z. E.; Luo, L. X.; Zhang, F.; Hong, M. Y.; Zhang, X.X.; Guo, R. X. Evaluation of yield, stability and adaptability of national winter rapeseed regional trials in the upper Yangtze River region in 2017-2018. *Oil Crop Science*. **2020**, *5*, 121-128.
20. Flores, F.; Hybl, M.; Knudsen, J. C.; Marget, P.; Muel, F.; Nadal, S.; Narits, L.; Raffiot, B.; Sass, O.; Solis, I.; Winkler, J.; Stoddard, F.L.; Rubiales, D. Adaptation of spring faba bean types across European climates. *Field Crops Research*. **2013**, *145*, 1-9.
21. Choudhary, M.; Kumar, B.; Kumar, P.; Guleria, S. K.; Singh, N. K.; Khulbe, R.; Kamboj, M. C.; Vyas, M.; Srivastava, R.K.; Puttaramanaik.; Swain, D.; Mahajan, V.; Rakshit, S. GGE biplot analysis of genotype× environment interaction and identification of mega-environment for baby corn hybrids evaluation in India. *Indian Journal of Genetics and Plant Breeding*. **2019**, *79*, 658-669.
22. Oliveira, T. R. A. D.; Carvalho, H. W. L. D.; Oliveira, G. H. F.; Costa, E. F. N.; Gravina, G. D. A.; Santos, R. D. D.; Carvalho, J. L. S. D. Hybrid maize selection through GGE biplot analysis. *Bragantia*. **2019**, *78*, 166-174.
23. Yan, W. K.; Pageau, D.; Frégeau-Reid J.; Durand, J. Assessing the representativeness and repeatability of test locations for genotype evaluation. *Crop Science*. **2011**, *51*, 1603-1610.
24. Yan, W. K.; Hunt, L. A.; Sheng, Q. L.; Szlavnics, Z. Cultivar evaluation and mega-environment investigation based on the GGE biplot. *Crop Science*. **2000**, *40*, 597-605.
25. Kumar, R.; Dhansu, P.; Kulshreshtha, N.; Meena, M.R.; Kumaraswamy, M.H.; Appunu, C.; Chhabra, M.L.; Pandey, S.K. Identification of Salinity Tolerant Stable Sugarcane Cultivars Using AMMI, GGE and Some Other Stability Parameters under Multi Environments of Salinity Stress. *Sustainability*. **2023**, *15*, 1119.
26. Ashraf, M. A.; Akbar, A.; Parveen, A.; Rasheed, R.; Hussain, I.; Iqbal, M. Phenological application of selenium differentially improves growth, oxidative defense and ion homeostasis in maize under salinity stress. *Plant Physiology and Biochemistry: PPB*. **2017**, *123*, 268–280.
27. Guo, H. J.; Hu, T.; Fu, J.M. Effects of saline sodic stress on growth and physiological responses of *Lolium perenne*[J]. *Acta Prataculturae Sinica*, **2012**, *21*, 118-125.
28. Yuan, H.; He, P.F.; Wu, J.J.; Wu, Y.X.; Li, X.Y.; He, P.B.; Kang, Z.Y.; He, Y.Q. Effects of salt stress on growth and biological traits of salt - tolerant and salt - sensitive maize seedlings. *Jiangsu Agricultural Sciences*. **2019**, *47*, 86-89.
29. Zhang, F.Q.; Wang, Y.S.; Dong, J.D.; Sun, C. C.; Ying, J.P. Effects of wastewater borne heavy metals on some proteceive enzymes and lipid peroxidation od *Bruguiera gymnorhiza* seedling. *Journal of Tropical Oceanography*. **2006**, (02), 66-70.
30. Epstein, E.; Norlyn, J. D.; Rush, D. W.; Kingsbury, R. W.; Kelley, D. B.; Cunningham, G. A.; Wrona, A. F. Saline culture of crops: a genetic approach. *Science (New York, N.Y.)*. **1980**, *210*, 399–404.

31. Frukht, A.; Siddiqi, T. O.; Khan, M. I. R.; Ahmad, A. Modulation in growth, biochemical attributes and proteome profile of rice cultivars under salt stress. *Plant physiology and biochemistry: PPB*. **2020**, *146*, 55–70.
32. Truong, H. A.; Lee, W. J.; Jeong, C. Y.; Trinh, C. S.; Lee, S.; Kang, C. S.; Cheong, Y. K.; Hong, S. W.; Lee, H. Enhanced anthocyanin accumulation confers increased growth performance in plants under low nitrate and high salt stress conditions owing to active modulation of nitrate metabolism. *Journal of plant physiology*. **2018**, *231*, 41–48.
33. Fu, C.F.; Zhang, H.Y.; Effects of Salt Stress on Seed Germination and Seedling Cgilorophyll Content and Osmotic Potential of Maize. *Shandong Agricultural Sciences*. **2015**, *47*, 27–30.
34. Shahid, M. A.; Balal, R. M.; Pervez, M. A.; Abbas, T.; Mattson, N. S. Differential response of pea (*Pisum sativum* L.) genotypes to salt stress in relation to the growth, physiological attributes antioxidant activity and organic solutes. *Australian Journal of Crop Science*. **2012**, *6*, 828–838.
35. Batista, P. S. C.; Menezes, C. B.; Carvalho, A. J.; Portugal, A. F.; Bastos, E. A.; Cardoso, M. J.; Santos, C. V.; Julio, M. P. M. Performance of grain sorghum hybrids under drought stress using GGE biplot analyses. *Genetics and molecular research: GMR*. **2017**, *16*, 10.4238.
36. Yan, W.K. Singular-Value Partitioning in Biplot Analysis of Multi-environment Trial Data. *Agronomy Journal*. **2002**, *94*, 990–996.
37. Gupta, V.; Mehta, G.; Kumar, S.; Ramadas, S.; Tiwari, R.; Singh, G. P.; Sharma, P. AMMI and GGE biplot analysis of yield under terminal heat tolerance in wheat. *Molecular biology reports*. **2023**, *50*, 3459–3467.
38. Saeidnia, F.; Taherian, M.; Nazeri, S. M. Graphical analysis of multi-environmental trials for wheat grain yield based on GGE-biplot analysis under diverse sowing dates. *BMC plant biology*. **2023**, *23*, 198.
39. Luo, J.; Zhang, H.; Deng, Z. H.; Que, Y. X. Ying yong sheng tai xue bao = The journal of applied ecology. **2012**, *23*, 1319–1325.
40. Silva, W. J. D. S.; Alcântara Neto, F.; Al-Qahtani, W. H.; Okla, M. K.; Al-Hashimi, A.; Vieira, P. F. M. J.; Gravina, G. A.; Zuffo, A. M.; Dutra, A. F.; Carvalho, L. C. B.; Sousa, R. S.; Pereira, A. P. A.; Leite, W. S.; Silva Júnior, G. B. D.; Silva, A. C. D.; Leite, M. R. L.; Lustosa Sobrinho, R.; AbdElgawad, H. Yield of soybean genotypes identified through GGE biplot and path analysis. *PloS one*. **2022**, *17*, e0274726.
41. Deng, L.; Guo, M.J.; Miao, J.L.; Ren, L. Comprehensive analysis of peanut cultivars with big grain based on path coefficient and GGE biplot. *Jiangsu Agricultural Sciences*. **2021**, *49*, 129–133.
42. Wang, X.B.; Wang, H.; Hu, K.M.; Li, Y.J.; Qing, T.Y.; Zeng, W.J.; Li, X.; Zhang, K.L.; Zhang, J.L.; Bai, J.P. Comprehensive Evaluation of Introduced Potato Germplasm Resources Based on the Analytical Hierarchy Process and GGE - biplot. *Journal of Plant Genetic Resources*. **2017**, *18*, 1067–1078.
43. Yan, W.; Kang, M. S.; Ma, B.; Woods, S.; Cornelius, P. L. Gge biplot vs. ammi analysis of genotype-by-environment data. *Crop Science*. **2007**, *47*, 641–653.
44. Saremirad, A.; Taleghani, D. Utilization of Univariate Parametric and non-Parametric Methods in the Stability Analysis of Sugar Yield in Sugar Beet (*Beta vulgaris* L.) Hybrids. *Journal of Crop Breeding*. **2022**, *14*, 49–63.
45. Taleghani, D.; Rajabi, A.; Saremirad, A.; Fasahat, P. Stability analysis and selection of sugar beet (*Beta vulgaris* L.) genotypes using AMMI, BLUP, GGE biplot and MTSI. *Scientific reports*. **2023**, *13*, 10019.

46. Yan, W.; Kang, M. S. GGE Biplot Analysis: A Graphical Tool for Breeders, Geneticists, and Agronomists. CRC PRESS :Boca Raton London New York Washington, D.C, 2002; pp. 63-65.
47. Kong, F.J.; Zheng, M.P.; Zhang, H.X.; Li, Z.; Wang, L.W. Salt Lake Agriculture and Ints Development Strategy. Strategic Study of CAE. **2019**, *21*, 1478-152.
48. Wang, Q. Z.; Liu, Q.; Gao, Y. N.; Liu, X. Review on the mechanisms of the response to salinity-alkalinity stress in plants. Acta Ecologica Sinica. **2017**, *37*, 5565-5577.
49. Ye, X.X.; Wang, H.; Cao, X.L.; Jin, X.J.; Cui, F.Q.; Bu, Y.Y.; Liu, H.; Wu, W.W; Takano, T.; Liu, S.K. Transcriptome profiling of *Puccinellia tenuiflora* during seed germination under a long-term saline-alkali stress. BMC genomics. **2019**, *20*, 589.
50. Tian, C. Y.; Zhou, H. F.; Liu, G. Q. THE PROPOSAL ON CONTROL OF SOIL SALINIZING AND ARGICULTURAL SUSTAINING DEVELOPMENTN IN 21'S CENTURY IN XINJIANG. Arid Land Geography. **2000**,177-181.

Figure 1

Analysis of difference in various morphological indicators under salinity stress.

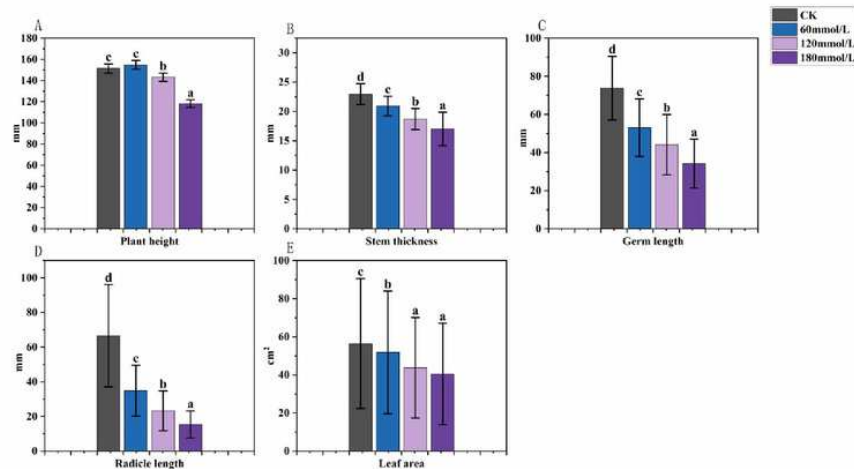


Figure 1. Analysis of difference in various morphological indicators under salinity stress. A: Analysis of the difference in plant height among four NaCl solutions; B: Analysis of the difference in stem thickness among four NaCl solutions; C: Analysis of the difference in germ length among four NaCl solutions; D: Analysis of the difference in radicle length among four NaCl solutions; E: Analysis of the difference in leaf area among four NaCl solutions; a, b, c and d are markers of significance of differences at 0.05 level.

Figure 2

Analysis of difference in various physiological indicators under salinity stress.

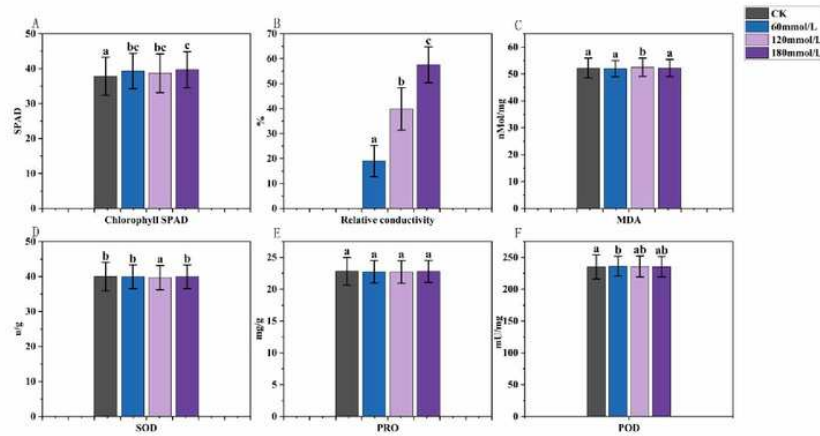


Figure 2. Analysis of difference in various physiological indicators under salinity stress. A: Analysis of the difference in chlorophyll content among four NaCl solutions; Analysis of the difference in conductivity among four NaCl solutions; Analysis of the difference in MDA content among four NaCl solutions; Analysis of the difference in SOD activity among four NaCl solutions; Analysis of the difference in PRO content among four NaCl solutions; Analysis of the difference in POD activity among four NaCl solutions; a, b, c and d are significant difference markers at 0.05 level.

Figure 3

Analysis of difference in various germination indicators under salt stress.

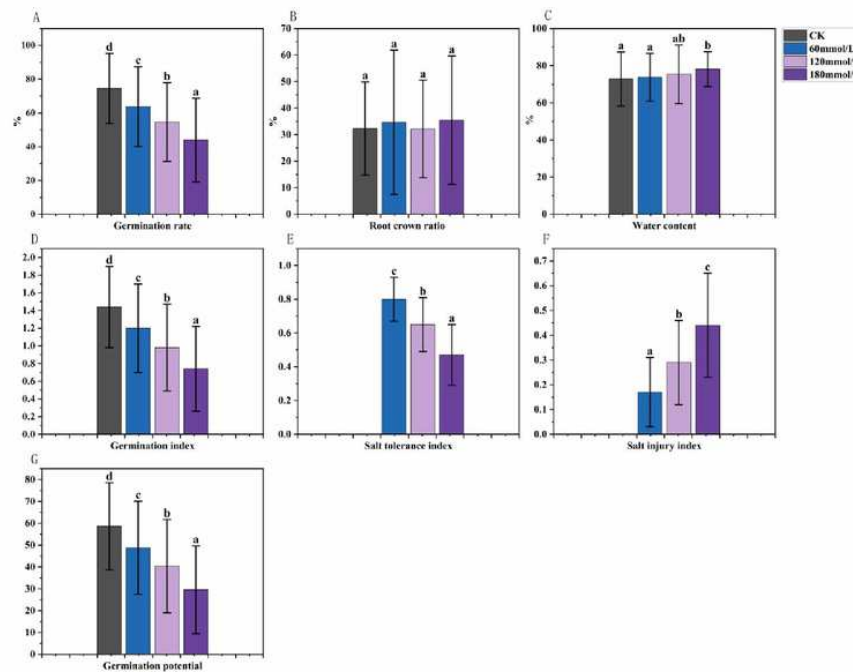


Figure 3. Analysis of difference in various germination indicators under salt stress. Analysis of the difference in germination percentage among four NaCl solutions; Analysis of the difference in root-crown ratio among four NaCl solutions; Analysis of the difference in water content ratio among four NaCl solutions; Analysis of the difference in germination index among four NaCl solutions; Analysis of the difference in salt tolerance index among four NaCl solutions; Analysis of the difference in salt damage index among four NaCl solutions; Analysis of the difference in germination potential among four NaCl solutions; a, b, c and d are significant markers of difference at the 0.05 level.

Figure 4

Correlation of agronomic traits with physiological indicators.

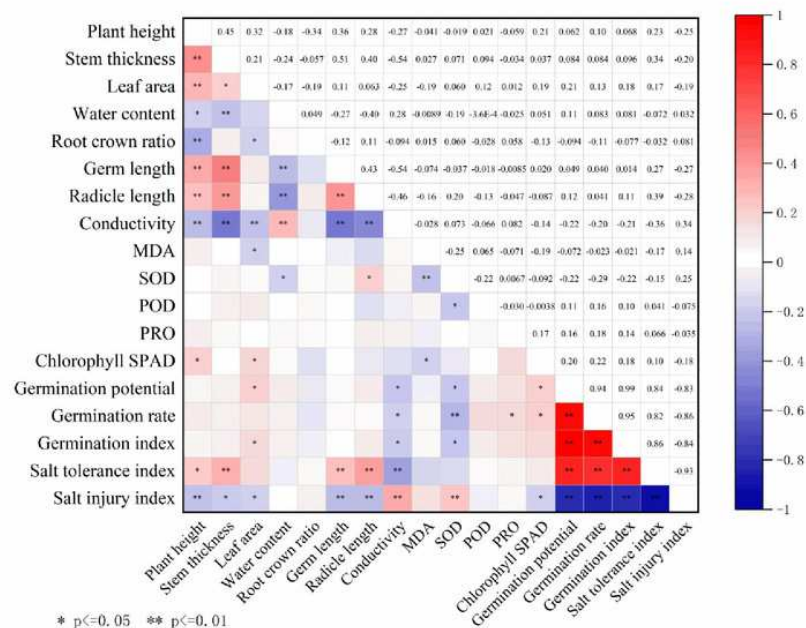


Figure 4. Correlation of agronomic traits with physiological indicators.

Figure 5

Adaptation analysis of maize varieties based on GGE biplot analysis.

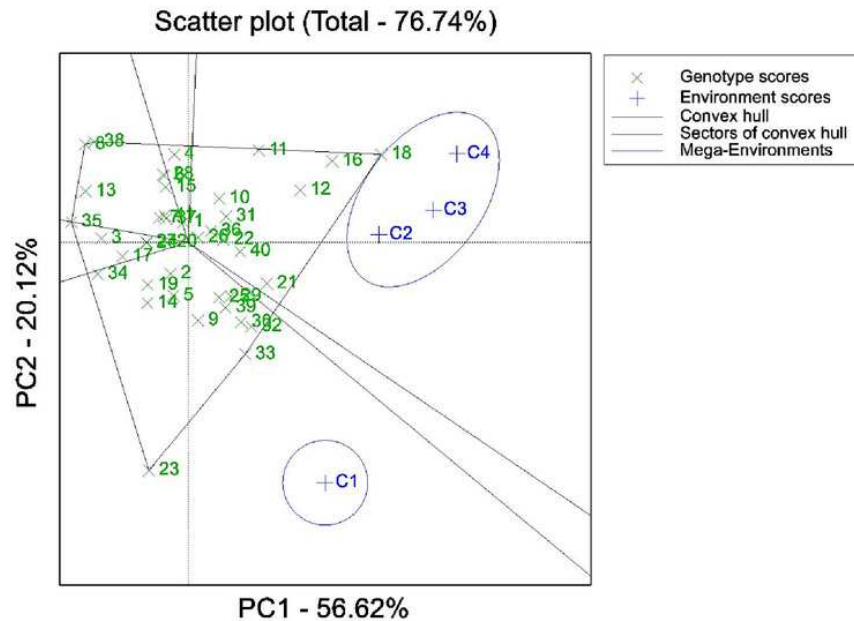


Figure 5. Adaptation analysis of maize varieties based on GGE biplot analysis. 1, 2, 3, etc. are the cultivar numbers of 41 maize varieties; C1, C2, C3 and C4 are the environments under stress of 0 mMol/L NaCl solution, 60 mMol/L NaCl solution, 120 mMol/L NaCl solution and 180 mMol/L NaCl solution, respectively.

Figure 6

Analysis of optimal NaCl stress solutions based on GGE biplots.

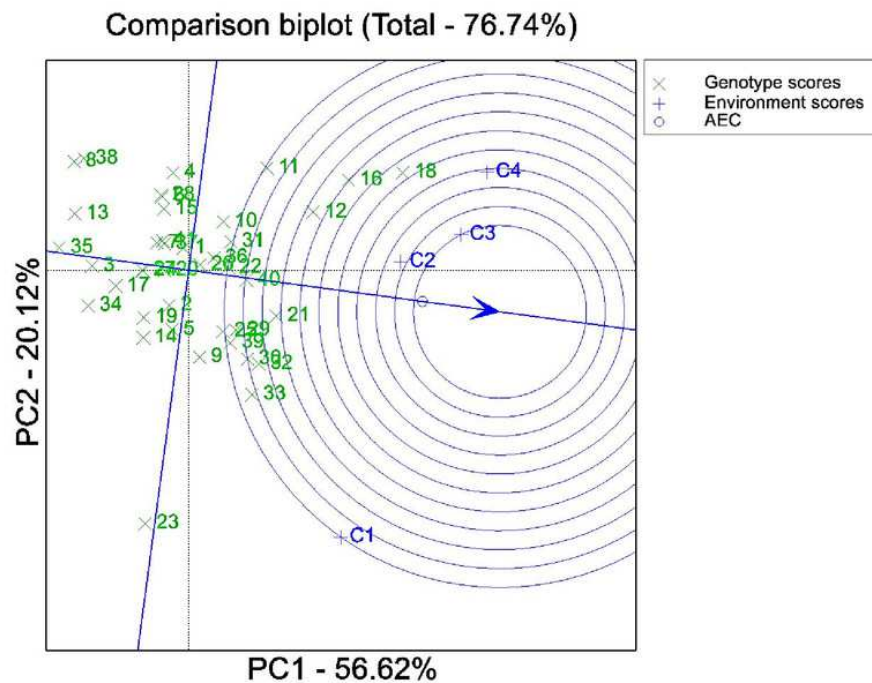


Figure 6. Analysis of optimal NaCl stress solutions based on GGE biplots. 1, 2, 3, etc. are the cultivar numbers of 41 maize varieties; C1, C2, C3 and C4 are the environments under stress of 0 mMol/L NaCl solution, 60 mMol/L NaCl solution, 120 mMol/L NaCl solution and 180 mMol/L NaCl solution, respectively.

Figure 7

Representative analysis of NaCl stressed solutions based on GGE biplot.

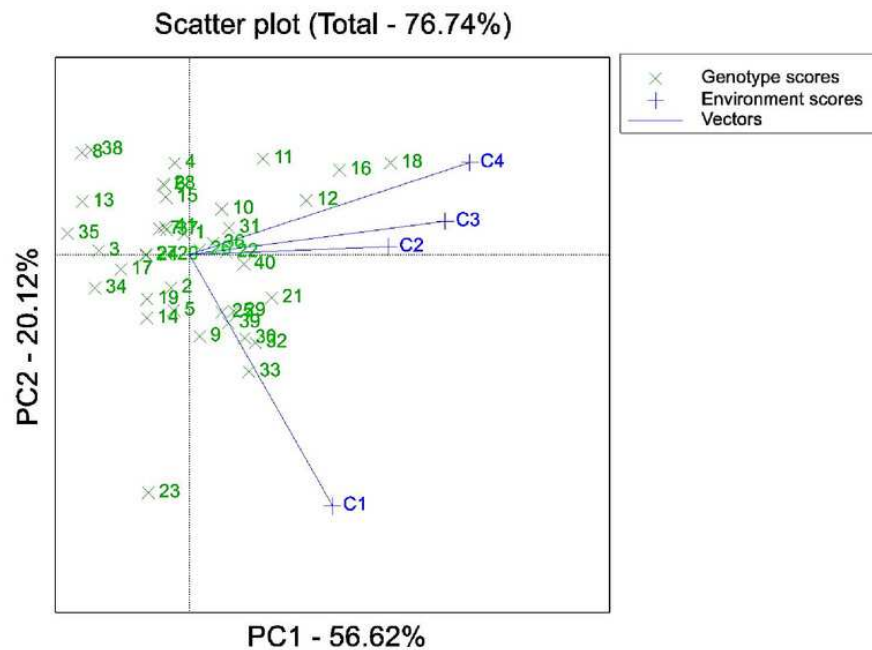


Figure 7. Representative analysis of NaCl stressed solutions based on GGE biplot. 1, 2, 3, etc. are the cultivar numbers of 41 maize varieties; C1, C2, C3 and C4 are the environments under stress of 0 mMol/L NaCl solution, 60 mMol/L NaCl solution, 120 mMol/L NaCl solution and 180 mMol/L NaCl solution, respectively.

Figure 8

Varietal analysis of salt tolerance stability based on GGE biplot.

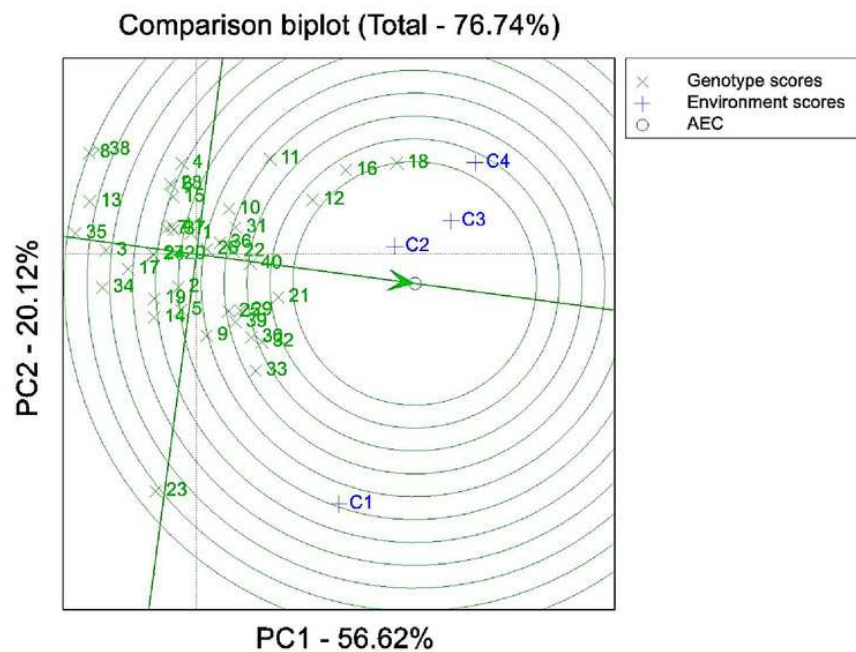


Figure 8. Varietal analysis of salt tolerance stability based on GGE biplot. 1, 2, 3, etc. are the cultivar numbers of 41 maize cultivars; C1, C2, C3 and C4 are the environments under stress of 0 mMol/L NaCl solution, 60 mMol/L NaCl solution, 120 mMol/L NaCl solution and 180 mMol/L NaCl solution, respectively.

Table 1 (on next page)

Maize materials tested.

1

Table 1. Maize materials tested.

Number	Name	Source	Number	Name	Source
1	Hongxing 990	Jilin Hongxing Seed Industry Co	22	Zhengshan 958	Henan Academy of Agricultural Sciences
2	Ping'an 1523	Jilin Ping'an Seed Industry Co	23	Hengyu 369	Jilin Hengyu Seed Industry Co.
3	Huxin 712	Huludao Agriculture Co	24	Xianghe 9918	Changtu Dingsheng Agricultural Co.
4	Seedstar 618	Inner Mongolia Seedstar Co	25	Xingnong No.1	Zhongyan Seed Industry Co.
5	Simon 6	Inner Mongolia Simon Co	26	Bixiang 809	Beijing Huanong Weiye Seed Industry Co.
6	Hongxing 528	Jilin Hongxing Seed Industry Co	27	Jinfengjie 607	Zhengzhou Fengjie Seed Co.
7	Zengyu 157	Jilin Hongxing Seed Industry Co	28	Jinongyu 309	College of Agriculture, Jilin Agricultural University
8	Wofeng 188	Shanxi Wodafeng Co	29	Xinyu 81	Xinjiang Pioneer Weiye Seed Co.
9	Yuhe 536	Henan Yuyu Seed Industry Co	30	Xinyu 24	Xinjiang Xinshi Seed Co.
10	Wofeng No.9	Shanxi Vodafone Co.	31	Denghai 3672	Shandong Denghai Seed Industry Co.
11	Nongfu 99	Inner Mongolia Zhongnong Co	32	Xinyu 66	Urumqi Shengyang Agricultural Company
12	Xuanhui 8	Xuanhui Agriculture Co	33	Xinnong 008	Inner Mongolia Lanhai Xinnong Agricultural Company
13	Simon 668	Inner Mongolia Simon Co	34	Xianyu 335	Tieling Pioneer Seed Co.
14	Linyu 1339	Yunnan Linpeng Agriculture Co	35	Toyota 14	Chifeng Fengtian Science and Technology Seed Industry Co.
15	Yuanyuan 1	Weishan Jiuyan Agricultural Co.	36	Fuyu 109	Zhongyan Seed Industry Co.

16	Qunze 888	Sichuan Qunze Seed Industry Co	37	Simon 3358	Inner Mongolia Simon Co.
17	Shann ing 23	Ningxia Jinyu Seed Co	38	Jixing2 18	Jilin Xingnong Seed Industry Co.
18	Youqi 909	Jilin Hongxiang Seed Industry Co	39	Sanme ng 9599	Changtu Zewei Agricultural Science Research Institute
19	Jin'ai 588	Inner Mongolia Jin'ai Ailite Co	40	Simon 208	Inner Mongolia Simon Co.
20	Ganxi n 2818	Gansu Province Wuwei Agricultural Research Institute	41	Huxin 338	Huludao Agriculture Co.
21	Wugu 568	Gansu Wugu Seed Industry Co			

Table 2(on next page)

Maize trait contribution eigenvalues. Extraction method: Principal component analysis.

Table 2. Maize trait contribution eigenvalues. Extraction method: Principal component analysis.

Factor	Initial Eigenvalues			Sum of squared rotating loads		
	Total	Percentage variance	Cumulative %	Total	Percentage variance	Cumulative %
1	5.141	28.561	28.561	5.141	28.561	28.561
2	2.824	15.688	44.250	2.824	15.688	44.250
3	1.541	8.560	52.810	1.541	8.560	52.810
4	1.472	8.181	60.990	1.472	8.181	60.990
5	1.067	5.929	66.919	1.067	5.929	66.919
6	1.006	5.589	72.508	1.006	5.589	72.508
7	0.941	5.227	77.735	0.941	5.227	77.735
8	0.781	4.337	82.072	0.781	4.337	82.072
9	0.715	3.971	86.043			
10	0.637	3.541	89.584			
11	0.562	3.122	92.706			
12	0.399	2.214	94.920			
13	0.374	2.080	97.000			
14	0.298	1.656	98.656			
15	0.150	0.831	99.487			
16	0.066	0.365	99.852			
17	0.019	0.104	99.956			
18	0.008	0.044	100.000			

Table 3(on next page)

Matrix of factor loadings after rotation.

Table 3. Matrix of factor loadings after rotation.

Characters	Ingredients							
	1	2	3	4	5	6	7	8
Plant height	0.064	0.503	0.568	0.094	0.266	0.008	0.021	0.091
Stem thickness	0.074	0.759	0.098	0.139	0.057	0.030	0.113	0.034
Leaf area	0.159	0.066	0.187	0.528	0.548	0.125	0.261	0.190
Water content	0.062	0.358	0.050	0.718	0.103	0.122	0.005	0.147
Root crown ratio	0.063	0.035	0.919	0.017	0.029	0.012	0.018	0.013
Germ length	0.032	0.832	0.157	0.082	0.057	0.067	0.010	0.079
Radicle length	0.170	0.656	0.134	0.312	0.215	0.201	0.233	0.045
Conductivity	0.205	0.747	0.232	0.134	0.257	0.141	0.055	0.139
MDA	0.070	0.038	0.007	0.021	0.140	0.933	0.011	0.037
SOD	0.228	0.051	0.055	0.545	0.124	0.475	0.280	0.007
POD	0.071	0.026	0.001	0.028	0.038	0.031	0.952	0.004
PRO	0.103	0.057	0.047	0.063	0.112	0.035	0.004	0.947
Chlorophyll SPAD	0.118	0.046	0.074	0.161	0.834	0.090	0.096	0.187
Germination potential	0.968	0.012	0.013	0.012	0.099	0.000	0.047	0.052
Germination rate	0.956	0.010	0.055	0.069	0.088	0.068	0.088	0.101
Germination index	0.979	0.024	0.008	0.008	0.069	0.051	0.031	0.041
Salt tolerance index	0.909	0.312	0.010	0.030	0.038	0.113	0.026	0.016
Salt injury index	0.906	0.254	0.061	0.052	0.052	0.062	0.004	0.042

Table 4(on next page)

D-values of eight indices such as germination index and chlorophyll SPAD under four NaCl concentration stresses.

1 Table 4. D-values of eight indices such as germination index and chlorophyll SPAD under four
2 NaCl concentration stresses.

Cultivar number	CKD value	60mMol/LD value	120mMol/LD value	180mMol/LD value
1	1.82	2.03	1.43	1.86
2	2.11	1.65	1.57	1.63
3	1.54	2.02	0.99	0.94
4	1.41	1.17	2.14	1.87
5	2.23	1.30	2.12	1.36
6	1.51	1.97	1.07	2.05
7	1.65	1.73	1.84	1.36
8	0.90	1.44	1.57	0.86
9	2.40	2.35	1.61	1.44
10	1.76	2.03	2.13	1.90
11	1.49	2.52	2.74	1.84
12	1.91	2.73	2.78	2.18
13	1.30	1.24	1.12	1.22
14	2.21	1.21	1.75	1.31
15	1.51	1.63	1.90	1.56
16	1.90	2.92	2.64	2.75
17	1.71	1.78	1.59	0.86
18	2.25	2.40	2.53	3.92
19	2.07	1.79	1.28	1.40
20	1.85	1.59	1.92	1.41
21	2.42	2.22	2.58	1.87
22	2.00	2.36	1.93	1.80
23	3.21	1.83	1.12	0.96
24	1.89	1.51	1.16	1.79
25	2.39	1.70	2.27	1.68
26	1.86	2.31	1.90	1.48
27	1.78	1.75	1.48	1.37
28	1.41	1.83	1.74	1.57
29	2.46	1.87	2.01	1.97
30	2.57	2.22	2.22	1.66
31	1.88	2.09	2.20	1.85
32	2.69	2.42	1.77	2.05
33	2.86	1.98	2.05	1.91
34	1.78	1.69	1.10	0.89
35	1.46	1.10	0.96	1.13
36	1.88	2.50	1.76	1.69
37	1.75	1.84	1.37	1.78
38	0.97	1.83	0.91	1.31
39	2.43	1.93	2.35	1.54
40	2.16	2.48	1.80	2.05

3	41	1.74	1.17	2.04	1.62
4					
5					