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# **Stability Assessment of Groundnut ABLs Using AMMI and GGE Biplot Analysis for Yield Performance under Multi-environments**

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#### *Authors' contributions*

*This work was carried out in collaboration among all authors. All authors read and approved the final manuscript.*

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# **ABSTRACT**

The current study focused on Identification of stable line among the eight groundnut ABLs and two checks were assessed in three different environments in Karnataka during the 2020-2021 Kharif and Rabi seasons with randomized complete block design (RCBD) with three replications. The analysis of variance revealed significant differences (p≤0.01) among the genotypes, environments, and the genotype by environment interaction (G×E) for kernel yield. The AMMI analysis also showed highly significant differences (p≤0.01) for varieties, environments, and their interaction on kernel yield. The IPCA1 and IPCA2 components explained 72.72% and 25.00% of the total G×E sum of squares, respectively. The variations in kernel yield were attributed to the environment (16.44%), ABLs (81.87%), and ABLs by environment interaction (1.68%). ABLs T65, T77, T81, and T82 were identified as stable across all three environments based on ASV and SI for kernel yield

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per plant. These stable lines can be used as parents in breeding programs. The AMMI model and GGE biplots were effective tools for evaluating the adaptability and stability of groundnut genotypes in diverse environments.

*Keywords: Groundnut; ABLs; ASV; SI; AMMI; stability analysis; GGE biplot.*

# **1. INTRODUCTION**

Groundnut, also known as peanut, is a significant legume crop that is ranked  $12<sup>th</sup>$  among the world's food crops [1]. It is a valuable source of vegetable oil and is rich in protein, minerals, vitamins, and carbohydrates. The kernel of groundnut contains approximately 46-52% oil, 25-30% crude protein, and 12-18% carbohydrates [2]. "Additionally, groundnuts can improve soil fertility through nitrogen fixation when used in crop rotation, making them beneficial for agriculture. Groundnut is commonly cultivated in semiarid countries with moist weather, including Africa, America, and Asia, as stated by Singh and Singh" [3]. "Groundnut is commonly grown in semiarid countries with moist weather, such as Africa, America" [4]. "It is cultivated globally on 27.9 million hectares, producing 47 million tonnes with an average productivity of 1685 kg/ha. India contributes 22% to the global production, cultivating it on 6.014 million hectares and yielding 10.02 million tonnes (1703 kg/ha). In Karnataka, the current groundnut productivity is 720 kg per hectare, which is lower than the national average of 999 kg per ha" [5]. Despite the advent of numerous high-yielding crop varieties in recent years, TMV-2, a crop variety developed 82 years ago, continues to be the preferred choice among many farmers for cultivation. This enduring preference for TMV-2 is attributed to its remarkable resilience and adaptability to a wide range of environmental conditions, making it a reliable option for consistent yields.

Despite the availability of high-yielding groundnut varieties, TMV-2, which was developed 82 years ago, remains a preferred choice. However, its removal from the official seed supply list by the government has made it difficult to access. Addressing the productivity gap in groundnut cultivation necessitates the development of new varieties that not only yield higher but also retain the favorable pod and kernel characteristics of TMV-2. The widespread issue of low productivity is exacerbated by the slow adoption of these improved varieties and their variable performance under different environmental conditions. Furthermore, groundnut productivity

is influenced by genotype by environment interactions (GEI), indicating that the relationship between various traits and yield can be either beneficial or detrimental [6].

In order achieve the goal to increase production, it is important to adapt genotypes to changes in the environment over time and location. The interaction between genotype and environment can lead to inconsistent crop yields due to various factors such as unpredictable rainfall, limited resources in farming communities, crop diseases, and the inherent potential of genotypes. However, there is potential to identify or develop stable genotypes that yield well in different environments [7]. The AMMI model [8, 9] and the GGE biplot [10-12] are commonly used methods to analyze genotype-environment interactions and crop attributes. AMMI helps to understand the effects of genotype and environment, while the GGE biplot focuses on the interaction between genotype and environment, providing a comprehensive analysis and assessment of genotypes [13, 12].

Genotype stability analysis is essential for identifying genotypes that can consistently perform well in various environments [14]. "This interaction between genotype and environment is crucial for breeders to enhance breeding programs and minimize negative agro-climatic effects. Unfortunately, the influence of environmental conditions on quantitative traits in groundnut genotypes has not been extensively studied. Stable genotypes are able to adjust their phenotypic responses in order to maintain consistency even in the face of environmental fluctuations" [15]. Therefore, this study aimed to assess the performance of TMV-2 type groundnut advanced breeding lines in different locations and identify adaptable lines that are suitable for cultivation.

### **2. MATERIALS AND METHODS**

Study was conducted to evaluate eight groundnut genotypes and two control varieties across three different environments during 2021- Kharif and Rabi seasons (Table 1). The groundnut varieties used as parents in the crosses and checks were detailed in Table 2 and while the geographical and climatic data for each environment were provided in Table 3. The dibbling method of sowing was done in both Kharif and Rabi seasons at all three locations, with a randomized complete block design with three replications and 30 x 10 cm spacing between rows and plants. The standard crop management practices were adopted except for the spray of fungicides during the crop growth period in all environments. Kernel yield and other yield-related traits were recorded.

To accurately assess the interaction effects between advanced breeding lines (ABLs),

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parents, and different seasons on the growth trait indicated by the Genotypic Stability Index (GSI), data collected across three distinct seasons were analyzed utilizing the Additive Main Effects and Multiplicative Interaction (AMMI) model. This sophisticated statistical model facilitates a comprehensive evaluation of both the additive main effects, which are the direct influences of ABLs and parents, as well as the effects of seasons on the trait of interest and the multiplicative interaction effects which show the combination of ABLs and parents interact with seasonal variations to influence the trait. Initially, the analysis commenced with a univariate.









Locations	Environment label	Geographical position		<b>Altitude</b> (m.a.s.l)
		Latitude	Longitude	
National Seed Project (NSP), University of E1		$13^{0}08"$ N	77034" E	924m
Agricultural Sciences, GKVK, Bengaluru				
Agriculture Research Station, Balajigapade	E <sub>2</sub>	$13^{0}43"$ N	77079"E	915m
Organic Farming Research Station, Mandya	E3	12037"N	76°66"E	678m

**Table 3. Description of three locations used for evaluation of groundnut varieties**

Analysis of Variance (ANOVA), a procedure tailored to rigorously ascertain the additive main effects attributed to the ABLs, parents, and the three seasons under study. This step, documented in Table 5, is crucial for establishing a foundational understanding of the individual contributions of these factors to the trait being examined. Subsequently, the analysis advanced to a more nuanced stage involving the Interaction Principal Component Analysis (IPC), a pivotal component of the AMMI model. This stage specifically targets the  $(ABLs + parents) \times$ season interaction, providing a multidimensional view of how these factors collectively influence the trait in question. The IPC analysis is particularly adept at disentangling the complex interactions that are not readily apparent from the main effects alone, thus offering a more granular insight into the dynamics at play. The employment of the AMMI model in this context is underpinned by its capacity to simultaneously handle additive and multiplicative effects, rendering it an invaluable tool in the field of agricultural research for dissecting and understanding the multifaceted interactions between genetic lines, environmental conditions, and their combined influence on phenotypic outcomes. This methodological approach affords researchers a nuanced perspective on the genetic and environmental factors shaping trait expression, thereby informing targeted breeding and selection strategies to enhance crop performance across varying conditions.

$$
Y_{ij} = \mu + g_i + e_{j+\sum_{k=1}^{n} \lambda_k \alpha_{ik} \gamma_{jk} + \varepsilon_{ij}}
$$

Where.,

μ is the grand mean; αg is the deviation of genotype g from the grand mean, βe is the deviation of the environment e; λn is the singular value for IPCA, γgn is the genotype eigenvector for axis n, and δen is the environment eigenvector; εger is error term and ρge is PCA residual. "Accordingly, genotypes with low magnitude regardless of the sign of interaction principal component analysis scores have general or wider adaptability while genotypes with high magnitude of IPCA scores have specific adaptability" [16-18].

The AMMI stability value (ASV) for the i<sup>th</sup> genotype was computed for each genotype and environment based on the proportion of IPCA1 to IPCA2 in the interaction SS as follows [19].

$$
ASV = \sqrt{\frac{SSIPC1}{SSIPC2}} (IPC1 score)^{2} + (IPC2 score)^{2}
$$

Where,

"The sum of squares (SS) related to the first two IPCs are denoted as SSIPC 1 and SSIPC 2. In theory, ASV is the separation between zero and an IPC 1 vs. IPC 2 scatter plot in two dimensions" [19]. "In order to account for the relative contributions of IPC 1 and IPC 2 scores to the total GSI sum of squares, the IPC 1 score which typically contributes proportionately more to GSI is weighted by the proportional difference between IPC 1 and IPC 2 values". [20] Greater stability is indicated by lower magnitude ASV estimates, whereas lesser stability of genotypes is indicated by higher magnitude ASV estimates [19].

#### **3. RESULTS AND DISCUSSION**

#### **3.1 Combined Analysis of Variance**

Analysis of variance revealed that there were statistically significant differences (P<0.01) among the varieties, across different environments, and also in the interaction between varieties and environments regarding kernel yield, as shown in Table 4. This finding suggests that there is a substantial genetic variation among the groundnut varieties being studied. Such genetic diversity indicates the potential for selecting varieties that not only yield high but are also stable across various environmental conditions. The results further underscore the fact that environmental conditions influence groundnut yield and that the performance of different groundnut varieties can vary significantly depending on the environmental context. This finding aligns with previous research which has similarly identified significant variations in the performance of<br>
across different groundnut varieties across different environments. The implication of these results is significant for breeding programs. It highlights the necessity of considering both the genetic makeup of the varieties and the environmental factors influencing crop yield during the selection process for high-yielding and environmentally resilient groundnut varieties [21,22].

### **3.2 Additive Main Effects and Multiple Interaction (AMMI) Model**

The results of the combined analysis of variance showed highly significant variations among environments, genotype x environment interaction, IPCA-1, and IPCA-2 (Table 5) [23]. This finding indicates differences in yield performance among groundnut varieties across various testing environments, underscoring a strong genotype by environment interaction. The GEI value represent a significant effect on the kernel yield of groundnuts and other yield attributing traits and highlighted significant variations in how varieties responded to changes in growing environments and the distinguishing characteristics of the test environments. Except for days to 50% flowering, locational contexts displayed significant mean squares for all traits, demonstrating the potential of temporal environments to differentiate the ABLs under

study. The noteworthy mean squares associated with ABLs further underscored the variations among the ABLs for each trait (Table 4). Similar outcomes as reported by Yayis *et al.,* [24] and Akande *et al.,* [25]. The interaction between genotype and environment  $(G \times E)$  was predominantly influenced by IPCA-1 (72.72%) and IPCA-2 (25.00%), which were utilized to construct a two-dimensional GGE biplot. Gauch and Zobel [26] along with Amare and Tamado [21] proposed that the optimal model for AMMI could be forecasted by considering the initial two IPCA components.

### **3.3 To Identify Specifically Stable Adaptable or Widely Adaptable TMV-2 Type Groundnut Advanced Breeding Lines**

### **3.3.1 GGE biplot analysis of GEI patterns**

The GGE bi-plot graph, which displays the distribution of ABLs based on their IPCs, offers a qualitative assessment of the stability and adaptability of ABLs across different spatial settings [25]. The traditional GGE bi-plot, also referred to as the SREG (sites regression) model, was initially introduced by Yan et al. [27]. This model incorporates genotype (G) and genotype × environment (GE) data, providing a comprehensive analytical approach that effectively visualizes the interactions between each ABL and its corresponding location environment.

<b>Source</b> variation	ofDegrees οf freedom	<b>Plant</b> height (cm)	<b>Primary</b> <b>branches</b> plant <sup>-1</sup>	<b>Days</b> 50% flowering	to	<b>Pods</b> plant <sup>-1</sup>		Pod yield plant <sup>-1</sup> (g)	
Replication	6	2.51	0.61	8.64		4.19	3.29		
ABLs	8	66.71**	$5.17***$	48.45**		82.60**		$13.29**$	
Location	$\overline{2}$	$1.67**$	$0.16**$	25.64**		14.75**		$8.14***$	
ABLsxLocation16		11.09**	$0.52**$	$8.46**$		$1.73**$		$1.23**$	
Poolederror	48	8.08	0.63	6.50		2.94	3.11		
<b>Source</b> variation	ofDegrees freedom	of Kernel	plant <sup>-1</sup> (g)	yield Shelling per Sound cent		kernel	mature Test	weight (g)	
Replication	6	1.53		0.25		26.90		2.63	
ABL <sub>s</sub>	8	$3.74***$		182.73**		195.44**		534.54**	
Location	2	$2.16***$		$9.17***$		$8.21***$		35.18**	

**Table 4. The pooled ANOVA estimation of groundnut ABLs across three locations for yield and its contributing traits during** *Rabi* **2021-22**

*Significant at P =0.05; \*\*Significant at P=0.01*

ABLsxLocation 16  $0.44**$  1.09\*\* 23.65\*\* 17.30\*\* Poolederror 48 0.81 0.18 24.70 17.86



Which Won Where/What







# **pattern of ABLs and locational environments for kernel yield plant-1 (g) during** *Kharif* **2021**

The graph depicting the results for Kernel yield plant-1 in the Kharif season of 2021 indicated that T72 and T82 emerged as the top-performing genotypes in environment 1 (GKVK), while T77, T65, and T61 were the winning genotypes in environment 2 (Mandya). Additionally, T89 was identified as the winning genotype in environment 3 (Balajigapade) as illustrated in Fig 1.

During *Rabi* 2021, as per our graph for kernel yield plant<sup>-1</sup> T72 was found to be the winning genotype in environment 1 (GKVK), T65 and T79 were discovered to be the winning genotype in environment 2 (Mandya), and T89 were shown to be the winning genotype in environment3 (Fig. 2).

#### **3.4 AMMI Model-Based Stability Parameters**

### **3.4.1 AMMI Stability value (ASV)**

ASV provides an impartial assessment of stability, facilitating the identification of ABLs that exhibit consistency across all three seasonal conditions. ASV is determined by measuring the distance from zero on a two-dimensional scatter plot of IPCA 1 (Interaction Principal Component Analysis Axis 1) scores against IPCA 2 (Interaction Principal Component Analysis Axis 2) scores. The IPC scores and ASV values for the Kharif 2021 and Rabi 2021 seasons are presented in table 6. The stable genotypes for the Kharif 2021 and Rabi 2021 seasons are

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#### **Table 6. Estimates of ASV and SI to assess stability of groundnut ABLs across three locations during** *Kharif* **and** *Rabi* **2021-22**





listed in (Table 7). In a similar vein, Ajay et al [26] conducted a stability analysis on fifty-two peanut genotypes over a span of two years, considering two phosphorous levels. They employed the AMMI model to investigate the interaction between genotype and environment (GEI) [28].

This study holds significant relevance in addressing the pressing challenges faced by agricultural systems globally, particularly in<br>regions characterized by diverse agroregions characterized by diverse agroenvironments like Karnataka, India. Understanding the performance and stability of crop genotypes across different environments is crucial for developing resilient and high-yielding varieties that can withstand environmental stresses [29, 30]. The findings of this study provide valuable insights into the adaptability of groundnut ABLs to varying soil quality, precipitation patterns, and other environmental

variables prevalent in the study area. By elucidating genotype-environment interactions, the research aids in tailoring crop management practices and breeding strategies to optimize productivity and mitigate the impact of environmental fluctuations on crop yields [31-33].

Soil quality plays a pivotal role in shaping crop performance and stability, as it directly influences nutrient availability [34], water retention [35], and overall plant health [36, 37]. By evaluating groundnut ABLs across diverse soil types present in Karnataka, this study sheds light on the genotype-specific responses to soil characteristics such as fertility, texture, and pH [38, 39]. Understanding how different genotypes interact with soil conditions allows for the selection of varieties with enhanced adaptability and resilience to specific soil types, thereby maximizing yield potential and ensuring sustainable crop production [40, 41].



Which Won Where/What



Furthermore, the influence of precipitation patterns and other environmental variables on groundnut performance underscores the need for climate-resilient crop varieties [42, 43]. As climate change continues to manifest through shifts in rainfall patterns [44, 45], temperature extremes, and unpredictable weather events [46, 47], the development of genotypes capable of thriving under varying climatic conditions becomes imperative [48, 49]. This study elucidates the differential responses of groundnut ABLs to precipitation regimes, highlighting the importance of incorporating genotype-specific adaptive traits in breeding programs [50, 51]. By identifying stable genotypes that exhibit consistent performance across different environmental conditions, this research contributes to the development of climate-smart agricultural practices aimed at enhancing food security and resilience in the face of climate variability [52, 53].

#### **4. CONCLUSION**

The investigation of G×E interactions in multienvironment trials was conducted using the wellestablished AMMI and biplot models. To better recommend stable and high-yielding groundnut ABLs, it is more suitable to simultaneously consider stability measures and yield. In this particular research, the advantages of both AMMI and biplot models were combined to enhance the reliability of the analysis of trials

analysis of variance for kernel yield was performed on the ABLs, revealing significant (p≤0.01) differences among genotypes, environments, and the G×E interaction. This highlighted the substantial variability attributed to the ABLs and their interaction with spatial environments for all traits examined in the study. The AMMI analysis was utilized to identify and characterize GSI, while the GGE bi-plot was employed to interpret GSI patterns of ABLs and pinpoint those with specific or wide adaptation. The AMMI analysis demonstrated highly significant (p≤0.01) differences for varieties, environments, and their interaction on kernel yield. The first and second interaction principal component axes (IPCA1 and IPCA2) were also found to be highly significant (p≤0.01), explaining 72.72% and 25.00% of the total G×E sum of squares, respectively. The environment, ABLs, and ABLs by environment interaction accounted for 16.44%, 81.87%, and 1.68% of the variations, respectively, indicating substantial differences in response among ABLs to changes in growing environments and the discriminating ability of the test environments. AMMI Stability Value (ASV) and Stability Index (SI) were estimated and were used to assess relative stability of ABLs. The results showed that ABLs T65, T77, T81, and T82 exhibited stability across all three locations and in both seasons, based on their ASV and SI values for kernel yield plant-1. These stable lines will undergo further evaluation in future years to

conducted across multiple locations. A pooled

determine their kernel yield potential. If validated, these stable lines can be released as a new variety or used as parents in the development of segregating populations. The use of the AMMI model and GGE biplot proved to be valuable methodologies in evaluating the adaptability and stability of groundnut genotypes in different environmental conditions. These two approaches complemented each other in providing a comprehensive assessment.

# **COMPETING INTERESTS**

Authors have declared that no competing interests exist.

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