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Additive Main-Effect and Multiplicative Interaction (AMMI) Model and Genotype Plus Genotype-Environment Interaction (GGE) Biplot Analysis of Tillering-Associated Traits for Climate-Smart Dual-Purpose Sorghum Selection in Lesotho

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# Introduction

#### Importance of Sorghum:

- Climate-resilient cereal crop vital for food security.
- Dual-purpose varieties address food and livestock feed demands; daptable to diverse environments;
- Tillering traits influence sorghum productivity and adaptability.

#### **Problem Statement:**

- Climate change impacts sorghum farming in Lesotho due to unpredictable rainfall, drought, and shrinking arable land.
- Lack of varietal recommendations tailored to local genotypes hinders productivity during climatic variability.

#### **Study Objectives:**

- Examine genotype-by-environment interaction (G×E) on tillering traits.
- Identify stable, high-performing sorghum genotypes.





# **Study site and Experimental Design**

#### Locations:

- Siloe (South): 1634m elevation, Arenosols.
- Nyakosoba (Central): 2034m elevation, Cambisols.
- Mahobong (North): 1682m elevation, fertile Luvisols.

#### **Climatic Conditions:**

• Variation in temperature and rainfall influencing sorghum tillering **Materials**:

- Evaluated 33 sorghum accessions from diverse origins.
- **Experimental Layout:**
- Lattice design with three replications.

### Measured Traits:

• Number of nodes, number of secondary tillers, internode length





# **Data Analysis Methods**

- Statistical Tools: GenStat software.
- AMMI Analysis: Partitioning genotype, environment, and G×E interaction effects; stability assessment using AMMI Stability Value (ASV).
- GGE Biplot Analysis: Visualizing genotype performance and stability across environments; identifying "winner" genotypes.



## **Results and Discussion:**

## **AMMI ANOVA**

Source of	D.f.	Numbe	er nodes		n	umber o	f	Internode l	ength	
variance					second	ary tillers	5			
		Ss	M.s.	%tv	Ss	M.s.	%tv	Ss	M.s.	%tv
Treatment	98	644.5	6.58***	64.4	89.59	0.914***	76.2	1611.8	16.45***	59.0
Genotype	32	125.4	3.92***	12.5	9.16	0.286**	7.8	594.3	18.57***	36.9
Environme nt	2	205.5	102.76***	20.5	63.36	31.68***	53.9	244.3	122.15** *	8.9
GE interaction	64	313.6	4.9***	31.3	17.07	0.267***	14.5	773.2	12.08***	28.3
IPCA 1	33	198.8	6.03***	63.4	15.60	0.473***	91.4	466.4	14.13***	60.3
IPCA 2	31	114.8	3.7**	36.6	1.47	0.047ns	8.6	306.8	9.9**	39.7
Error	192	340.3	1.77		26.27	0.137		1051.3	5.48	
Total	A 2nd	1000.7	3.38		117.57	0.397	_	2730.3	9.22	
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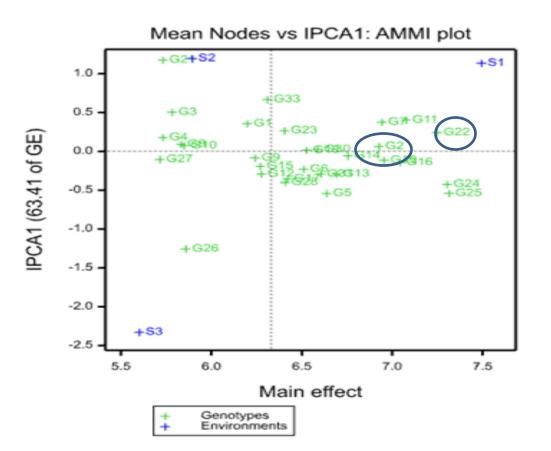
### **AMMI** Biplot

Genotypes on the right side of the grand mean level suggests high mean performance and near to the IPCA=0 line suggest no GE interaction

The environment points are more scattered than genotypes, indicating that environment variability is greater than genotype variability.

Traits	High- performing genotypes	More stable
Number of nodes	G22, G24 and G2	G2 and G14
Number of	G19, G2 and	G2 and G18
secondary tillers	G15	
Internode <sub>APPS</sub> length	A G15, G7 and	G6, G7 and G6
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#### AMMI Stability Value (ASV) & Genotype Selection Index (GSI)

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- ASV indicates stability of genotypes across environments.
- GSI combines ASV rank and mean performance rank for the genotype selection

Traits	Genotypes	RASV	RGM	GSI
Number of	G2	3	8	11
nodes	G19	6	6	12
	G22	10	3	13
Number of	G4	1	4	5
secondary tiller	G12	2	5	7
	G2	7	2	9
Internode	G14	2	2	4
length	G22	8	1	9
	G2	1	14	15



## **GGE Biplot Analysis: Mean Performance and Stability**

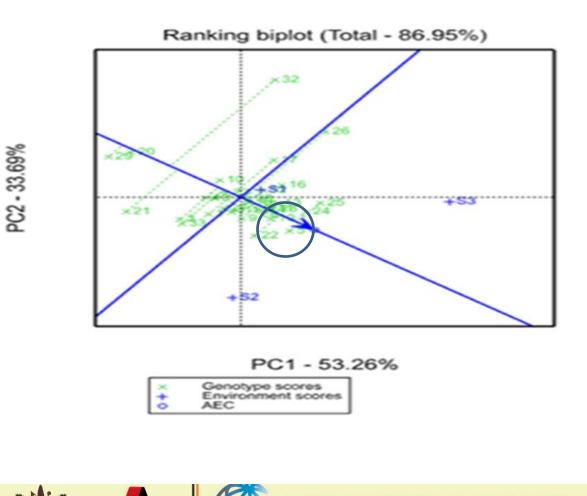
Abscissa line which is aligned with the AEA, shows the mean performance of the genotypes.

Genotypes close to the AEA in the direction of arrow have higher average performance. Genotypes closer to this axis are considered stable.

Traits	High- performing genotypes	Most stable genotypes
Number of nodes	G5, G25 and G24	G5
Number of secondary tiller	G22 and G28	G22
Internode length APPS	G31 and G20	G1 and G22
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## Number of nodes ranking biplot



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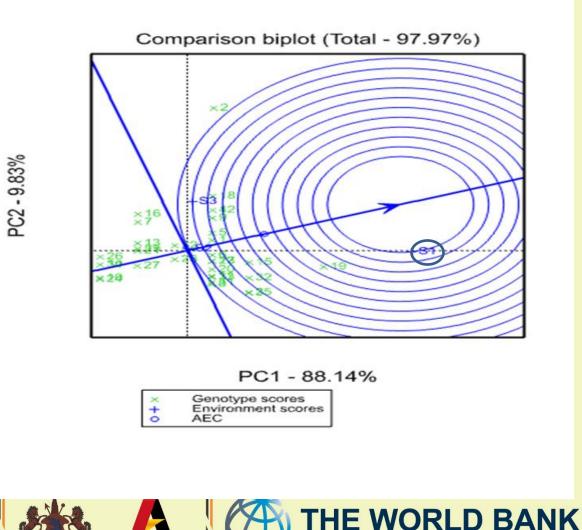
## **GGE Biplot: Comparison of** Test Environments

The ideal environment is positioned closer to the first concentric circle of the AEC and have longer vectors from the biplot origin.

Environment with long vectors are more discriminative.

Traits	Discriminating		
Number of podes	environment		
Number of nodes	Mahobong and		
	Nyakosoba		
Number of	Siloe		
secondary tiller			
Internode length	Mahobong		
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Number of Secondary tillers: Environments comparison biplot



# Genotype ranking: best genotype assessment

Genotypes closer to the center of concentric circle are considered high performing and stable across environments (winning genotypes) Traits High performing

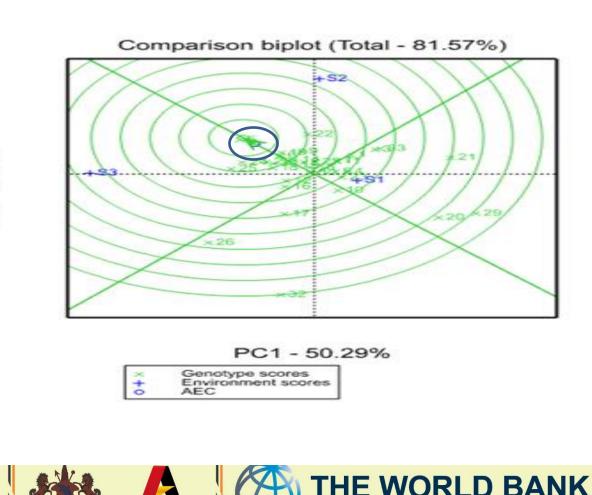
Number of nodes G5, G15, G22 and G24

Number ofG22, G25 and G28secondary tillers

Internode length G19 and G15

# Number of nodes: ranking genotypes assessment biplot

PC2 - 31.28%





# **Conclusion and recommendations**

- **Key Findings**: Genotypes G2 and G22 are stable and highperforming across environments in Lesotho. Siloe and Mahobong are ideal test sites for future trials.
- Recommendations: Promote genotypes G22 and G2 for cultivation; conduct further testing under diverse environmental conditions; focus breeding programs on multi-environment trials to enhance adaptability under climate change scenarios.
  - Limitations: Limited test environments may not generalize findings across all regions in Lesotho.
  - Future Research Areas: Explore long-term adaptability under climate change scenarios; assess additional sorghum traits like grain quality or biomass yield.





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