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Genotype × environment interaction and stability analysis for yield and quality traits in little millet (*Panicum sumatrense* L.)

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Abstract

The present study was conducted to evaluate Genotype × Environment interaction and stability analysis for yield and quality parameters in 50 little millet genotypes under three environments *i.e.* Waghai, Vanarasi and Navsari locations under Gujarat, India during the year *Kharif*-2020. Stability analysis revealed that G x E interaction was significantly differed for all the characters except calcium content (mg/100 g) and ash content (mg/100 g) indicated that different genotypes reacted differently to different environmental conditions. Estimates of environmental indices indicated that Waghai location was favourable for yield contributing characters along with quality parameters followed by Navsari and Vanarasi. The results of present study revealed that none of the genotypes exhibited average stability for all the characters. Among the genotypes, WV 262, WV 258, WV 256, WV 293 and WV 273 were found average stable over environments for grain yield per plant with quality parameters. So, these genotypes may be used in further breeding programme in little millet.

Keywords: Little millet, stability, genotype × environment interaction, grain yield

Introduction

Little millet (*Panicum sumatrense* L.) is one of the coarse cereals consumed in the form of rice. It is self-pollinated crop with a chromosome number of 2n=4x=36. Little millet belongs to the family Poaceae, sub-family Panicoideae and the tribe Paniceae (Rachie, 1975)^[14]. Little millet's inflorescence is a panicle, contracted or thyrsiform and 15-45 cm long and 1-5 cm in wide (Seetharam *et al.*, 2003)^[16]. The spikelet is persistent and 2-3.5 mm long. Panicle branches are scabrous and drooping at the time of maturity. Spikelets were produced on unequal pedicels but solitary at the end of the branches. Each spikelet consisted of two-minute flowers. The lower one is sterile; the upper one is fertile or bisexual without rachilla extension (Sundararaj and Thulasidas, 1976)^[17]. The lateral vein is absent in lower glume and its apex is acute. The upper glume is ovate and without keel but larger than lower glume (Nanda and Agrawal, 2008)^[9]. The flowering progressed from the top to the bottom of the panicle. The anthesis occurred between 9.30 to 10.30 a.m. (Jayaraman *et al.* 1997)^[5]. The glumes open for a short while and self-pollination is the rule. The whole process of the anthesis is very rapid and is completed within 2-5 min.

Little millet is grown in India under various agro-ecological situations and commonly known as *samai, samo, moraio, vari and kutki.* Little millet is an important crop grown in the tribal belt of Madhya Pradesh, Chhattisgarh, Gujarat, Maharashtra, Odisha and Andhra Pradesh in India. In India, little millet having 1.42 lakh tones of production. In Gujarat, little millet is cultivated in an area of 10,634 hectares with 9,526 tonnes of production having the productivity of 896 kg/ha (Anonymous, 2021)^[1]. The area under this crop is mainly concentrated in the districts of Dangs, Valsad and Narmada of South Gujarat and Panchmahal of middle Gujarat.

Little millet is better as comparable to other cereals in terms of fiber, fat, carbohydrates, protein, calcium, iron and rich in phytochemicals included phenolic acids, flavonoids, tannins and phytate (Patil *et al.*, 2019) ^[11]. Therefore, it could address nutritional sensitive agriculture, which aimed at nutritional enhancement to combat the present scenario of micronutrient malnutrition. Little millet is known for its drought tolerance and considered as one of the least waters demanding crop. Crop improvement work carried out so far in this crop has thrown some success.

In the recent past some improved cultivars were developed but have limited yield potential. The potentiality of little millet has not been exploited in India and the yield levels were very low there by indicated a greater scope for exploitation of little millet under Indian condition.

Phenotype is defined as a linear function of Genotype (G), Environment (E) and G x E interaction effects. Relative importance of main and interaction effects might vary from genotype to genotype (Eberhart and Russell, 1966; Perkins and Jinks, 1968) ^[3, 13]. The study of G x E interaction served as a guide for various environmental niches. It is possible to identify genotypes with stability for high yield, through the stability for yield character as well as for quality traits.

Materials and methods

The experiment was conducted during Kharif-2020 having 50 little millet genotypes, viz., WV 254, WV 255, WV 256, WV 257, WV 258, WV 259, WV 260, WV 261, WV 262, WV 263, WV 264, WV 265, WV 266, WV 267, WV 268, WV 269, WV 270, WV 271, WV 272, WV 273, WV 274, WV 275, WV 276, WV 277, WV 278, WV 279, WV 280, WV 281, WV 282, WV 283, WV 284, WV 285, WV 286, WV 287. WV 288. WV 289. WV 290. WV 291. WV 292. WV 293. WV 294. WV 295. WV 296. WV 297. WV 298. WV 299, WV 300, WV 301, WV 302 and WV 303 were evaluated in randomized block design at Hill Millet Research Station, Navsari Agricultural University, Waghai, Gujarat, India; Niger Research Station, Navsari Agricultural University, Vanarasi, Gujarat, India and College Farm, N. M. College of Agriculture, Navsari Agricultural University, Navsari, Gujarat, India during Kharif-2020. The seedlings were planted at 22.5 x 10 cm² spacing. All recommended practices were followed and timely plant protection measures were taken to avoid damage through insect-pests and diseases.

The observations on five randomly selected plants were recorded for grain yield per plant and quality characters *viz.*, hulling (%), chlorophyll content (mg/100 g fresh weight), leaf area (cm²), protein content (%), crude fiber (%), mineral matter (mg/100 g), iron content (mg/100 g), calcium content (mg/100 g) and ash content (mg/100 g). Estimation of stability parameters evaluated by the Eberhart and Russell (1966) ^[3] model.

Results and Discussions

The analysis of variance for stability (Table 1) revealed that, the differences among the genotypes and environments were also significant for all the traits when tested against pooled deviation as well as pooled error. The environments + (genotypes x environments) interaction was observed to be significant for all traits when tested either against pooled deviation or pooled error. Further partitioning of environments + (genotypes x environments) component of variation revealed that the environments (linear) components of variation as well as genotypes x environments (linear) component except for calcium content (mg/100 g) were observed to be significant for all the characters under study. The G x E interaction was significant for all characters except calcium content (mg/100 g) and ash content (mg/100 g). So, these traits were not considered for further analysis. The variance due to pooled deviation was found significant for hulling (%), chlorophyll content (mg/100 g fresh weight), leaf area (cm²), protein content (%), crude fiber (%), mineral matter (mg/100 g), iron content (mg/100 g), calcium content (mg/100 g) and ash content (mg/100 g). Highly significant

differences among genotypes, environments and G x E interaction were reported by Fentie *et al.* (2013) ^[4] and Kandel *et al.* (2020) ^[6].

The environmental indices computed for the quality characters studied were presented in Table 2 indicating both the favourable and unfavourable environments for all the component characters. Estimates of environmental indices indicated that Waghai location was favourable for most of the yield contributing characters along with quality parameters followed by Navsari and Vanarasi. It was also realized that among all the characters, leaf area (cm²) was the most vulnerable to environmental fluctuations.

The environmental indices calculated as the deviation of the mean of all the genotypes at a particular environment from the grand mean of all the genotypes revealed that in E_1 (Waghai) increased values in the environmental index for traits *viz.*, grain yield per plant (g), harvest index (%), hulling (%), chlorophyll content (mg/100 g fresh weight), leaf area (cm²), protein content (%), crude fiber (%), mineral matter (mg/100 g) and iron content (mg/100 g). The environmental index was observed to be congenial as well as poorest in environment E_2 (Vanarasi) for none of the traits. In E_3 (Navsari) environment index obtained poorest value for hulling (%), chlorophyll content (mg/100 g fresh weight), leaf area (cm²), protein content (%), crude fiber (%), mineral matter (mg/100 g) and iron content (%), mineral matter (mg/100 g) and iron content (%), crude fiber (%), mineral matter (mg/100 g) and iron content (mg/100 g).

When environmental indices of the different characters are studied, the most fluctuating traits observed were hulling (%), chlorophyll content (mg/100 g fresh weight) and leaf area (cm²). It indicated the vulnerability of these traits to the variation in the environment. It was also realized that among all the characters, leaf area (cm²) was the most vulnerable to environmental fluctuations. The protein content (%) was moderately affected by environmental changes. While, the grain yield per plant (g), crude fiber (%), mineral matter (mg/100 g) and iron content (mg/100 g) were less influenced by environmental fluctuations as compared to those listed before.

Patel *et al.* (2019) ^[10] reported the G × E interaction was significant for iron content. Patil (2007) ^[12] reported the genotypes *viz.*, RPSP 742, EC 138375 and RPSP 732 were high yielder with average stability of genotypes. Kandel *et al.* (2020) ^[6] reported significant genotypes and genotypes and their interaction for plant height along with genotype CO-4656 which had mean yield that was higher than the overall mean (0.429 t/ha) with parameter of response (b_i)=1.16 and parameter of stability (S²di)=0.05.

When genotypes with higher mean performance and nonsignificant deviation from regression (S²d_i=0) were tested for the significance of regression coefficient from unity, six genotypes viz., WV 262, WV 258, WV 256, WV 293, WV 294 and WV 273 for grain yield per plant (g); ten genotypes viz., WV 256, WV 291, WV 263, WV 286, WV 288, WV 299, WV 293, WV 259, WV 282 and WV 296 for hulling (%); three genotypes viz., WV 289, WV 286 and WV 302 for chlorophyll content (mg/100 g fresh weight); eight genotypes viz., WV 289, WV 272, WV 288, WV 286, WV 263, WV 303, WV 296 and WV 282 for leaf area (cm²); one genotype WV 286 for protein content (%); three genotypes viz., WV 263, WV 286 and WV 303 for crude fiber (%); none of genotype for mineral matter (mg/100 g) and three genotypes viz., WV 263, WV 287 and WV 303 for iron content (mg/100 g) showed a regression coefficient nearly equal to unity (bi=1), which demonstrated good general adaptation of character under various environments (Table 3, 4, 5,6 and 7). Four genotype *viz.*, WV 302, WV 303, WV 301 and WV 272 for grain yield per plant (g); one genotype WV 303 for hulling (%); three genotypes *viz.*, WV 297, WV 265 and WV 263 for chlorophyll content (mg/100 g fresh weight); one genotype WV 297 for leaf area (cm²); two genotypes *viz.*, WV 297 and WV 273 for protein content (%); two genotypes *viz.*, WV 297 and WV 302 for crude fiber (%); two genotypes *viz.*, WV 297 and WV 303 for mineral matter (mg/100 g) and one genotype WV 274 for iron content (mg/100 g) which had a higher mean value, regression coefficients below unity (b_i<1) and nonsignificant deviation from regression (S²d_i=0) was considered as only adapted to poor environment.

While four genotypes *viz.*, WV 259, WV 288, WV 269 and WV 296 for grain yield per plant (g); two genotypes *viz.*, WV 260 and WV 273 for hulling (%); three genotypes *viz.*, WV 288, WV 294 and WV 260 for chlorophyll content (mg/100 g fresh weight); five genotypes *viz.*, WV 256, WV 260, WV 294, WV 273 and WV 291 for leaf area (cm²); three genotypes *viz.*, WV 263, WV 256 and WV 260 for protein content (%); two genotypes *viz.*, WV 256 and WV 260 for crude fiber (%); two genotypes *viz.*, WV 286 and WV 256 for mineral matter (mg/100 g) and one genotype WV 286 for iron content (mg/100 g) were regarded as specifically adapted to a favourable environment because they had a higher mean value, a regression coefficient above unity (b_i>1), and a nonsignificant deviation from regression (S²d_i=0).

The Table 8 indicates the classification of genotypes by number based on their adaptation in different environments in little millet while, Table 9 indicates the classification of genotypes by name based on their adaptation in different environments in little millet. In general, the numbers of genotypes identified for average stability and wide/general adaptability were higher as compared to stable and adapted to poor environment or stable and adapted to better environment. Patel *et al.* (2019) ^[10] noted significant G × E interaction for yield and quality traits in pearl millet. Madhavilatha *et al.* (2020) ^[8] reported among the tested genotypes that PR-1041 recorded average stability for grain yield indicated the wide adoptability of this genotype for important traits. Also, found out significant G × E interaction for grain yield per plant. Kandel *et al.* (2020) ^[6] reported that the genotype CO-4656 had mean yield which was higher than the overall mean (0.429 t/ha), parameter of response (b) = 1.16 and parameter of stability (S2di) =0.05. Madhavilatha *et al.* (2020) ^[8] reported average stability for grain yield was found in VR 990 which revealed the wide adaptability of the genotype across different locations. Kandel *et al.* (2022) ^[7] studied genotypes *viz.*, GE-0382, KLE-216, NE-94 and KLE-559 that were found environmentally sensitive producing higher grain yield throughout the environments.

Patel *et al.* (2019) ^[10] reported significant $G \times E$ interaction for leaf area when tested against pooled error in pearl millet. Chavan *et al.* (2018) ^[2] recorded average stability for protein content (%) for genotypes *viz.*, GE-1680, Kanika Reddy, IVT-25, Nagli Dapoli 1 which indicated wider adoptability of these genotypes under all environments. Chavan *et al.* (2018) ^[2] found out general stability for iron content (mg/100 g) in the genotypes *viz.*, MR-6, PEH-1201 and IVT-11. Also, recorded average stability for protein content (%) for genotypes *viz.*, GE-1680, Kanika Reddy, IVT-25, Nagli Dapoli 1 which indicated wider adoptability of these genotypes under all environments.

Saritha *et al.* (2018) ^[15] noted that the genotypes *viz.*, VR-1034, GPU-71, DHWFM 11-3, OUAT-2 and JWM-1 were consistently stable across the environments whereas VR-936, GE-728, GE-6834-1, WFM-10, KMR-344, DHWFM 2-3 and GPU-67 were poorly adapted across the environments for their grain iron content. In general, the numbers of genotypes identified for average stability and wide/general adaptability were higher as compared to stable and adapted to poor environment or stable and adapted to better environment.

Source of variation	DF	Grain yield per plant (g)	Hilling	Chlorophyll content (mg/100 g fresh weight)	Leaf area (cm ²)	Protein content (%)	Crude fiber (%)	Mineral matter (mg/100 g)	Iron content (mg/100 g)	Calcium content (mg/100 g)	Ash content (mg/100 g)
Genotype (G)	49	6.01***	56.43***	72.77***	6074.21***	2.05***	0.71***	0.17***	1.53***	0.84***	0.11***
Environment (E)	2	17.29***	1224.21***	1651.92***	139918.90***	45.44***	16.96***	4.01***	30.98***	16.23***	2.48***
Env. + (Gen. x Env.)	100	0.73***	36.24***	51.00***	4156.08***	1.39***	0.51***	0.12***	0.97***	0.51***	0.08***
G x E	98	0.39**	12.00*	18.33*	1385.41*	0.49*	0.17*	0.04*	0.36*	0.19	0.03
Environment (Linear)	1	34.59***	2448.42***	3303.85***	279837.90***	90.88***	33.93***	8.03***	61.96***	32.46***	4.96***
G x E (Linear)	49	0.61***	16.02**	25.12**	1946.74**	0.69**	0.24**	0.06***	0.50**	0.23	0.04*
Pooled deviation	50	0.18	7.81*	11.30***	807.60***	0.30***	0.11***	0.02***	0.21***	0.15***	0.02***
Pooled error	294	0.16	5.03	1.36	316.91	0.02	0.008	0.002	0.01	0.06	0.003

Table 1: Analysis of variance for stability parameters with regards to different quality characters in little millet

*, ** and *** significant at 5, 1 and 0.1 per cent levels, respectively.

Table 2: Estimation of environment index (Ij) for various quality characters under different environments in little millet

Sr. No.	Characters	Environmental index							
51. 10.	Characters	Waghai (E1)	Vanarasi (E ₂)	Navsari (E ₃)					
1.	Grain yield per plant (g)	0.62	-0.08	-0.55					
2.	Hulling (%)	5.54	-1.57	-3.98					
3.	Chlorophyll content (mg/100 g fresh weight)	6.35	-1.51	-4.84					
4.	Leaf area (cm ²)	59.01	-15.85	-43.17					
5.	Protein content (%)	1.06	-0.27	-0.79					
6.	Crude fiber (%)	0.65	-0.16	-0.49					
7.	Mineral matter (mg/100 g)	0.32	-0.10	-0.22					
8.	Iron content (mg/100 g)	0.88	-0.24	-0.64					

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Table 3: Estimation of mean and stability parameter for	r hulling (%) and chlorophyll con	tent (mg/100 g fresh weight) in little millet

Sn No	Construes	Hulling (%)						Chlorophyll content (mg/100 g fresh weight)						
Sr. No.	Genotypes	Mean		bi 🛛		S ² d _i		Mean		bi		S ² d	i	
1	WV 254	52.83	0.51	**	+	-3.09		23.54	0.43	**	++	-1.15		
2	WV 255	53.71	0.90	*		0.99		24.56	0.79	*		5.61	*	
3	WV 256	59.34	1.38	**		0.30		29.85	1.56	**	++	0.98		
4	WV 257	57.91	1.19	**		-1.22		27.38	1.45	**		4.34	*	
5	WV 258	58.80	1.37			21.36	*	29.17	1.49	*		23.33	***	
6	WV 259	66.12	1.63	**		7.13		39.27	1.74	**		12.84	**	
7	WV 260	58.77	1.90	**	++	-4.89		30.00	1.85	**	++	-0.88		
8	WV 261	65.53	0.90			40.96	**	38.24	0.94			65.67	***	
9	WV 262	60.57	1.13			22.79	*	31.63	1.29			49.24	***	
10	WV 263	59.97	0.93	**		-4.23		32.12	0.84	**	++	-1.11		
11	WV 264	52.58	0.72	**	++	-4.76		24.89	0.30	**	++	-1.36		
12	WV 265	56.85	0.79	**	+	-4.49		30.27	0.47	**	++	-1.32		
13	WV 266	55.96	1.13	**	·	-3.37		26.28	1.12	**		1.81		
14	WV 267	54.60	0.85	*		2.66		25.12	0.83	*		8.24	**	
15	WV 268	64.40	0.03			6.72		36.57	0.17			25.23	***	
16	WV 269	57.56	1.91	**		5.74		28.41	2.01	**	+	14.22	***	
10	WV 270	56.22	1.26	**	++	-5.04		27.63	1.29	**	++	-1.27		
18	WV 271	52.26	0.05		++	-0.24		22.44	-0.12		++	2.52		
10	WV 271 WV 272	58.38	1.49	**	TT	-1.82		29.61	1.65	**	+	4.37	*	
20	WV 272 WV 273	58.76	2.00	**	++	-3.49		29.90	2.07	**	++	8.96	**	
20	WV 274	58.39	0.94	**	++	-3.49		29.90	0.78	**	++			
21										**		0.70		
22	WV 275	50.64	-0.05	**	++	-4.53		22.10	-0.11 1.09	**	++	-1.37	*	
	WV 276	55.00	1.16	~~		-2.56		25.58		ጥጥ		5.11	**	
24	WV 277	57.72	-0.08	**	++	2.82		29.11	-0.05	**	+	10.31	~~	
25	WV 278	51.69	0.48	**	++	-3.77		23.51	0.28	**	++	-0.97		
26	WV 279	55.23	1.24	**		-2.17		27.20	1.25	**		3.01		
27	WV 280	54.23	0.93			-4.98		24.82	0.75		++	-1.00	-	
28	WV 281	53.41	0.43	**	++	-4.67		23.47	0.49	**	++	-0.63		
29	WV 282	66.33	1.57	**		5.24		39.22	1.57	**		13.06	**	
30	WV 283	56.36	1.27	**	++	-5.04		27.01	1.26	**		0.24		
31	WV 284	59.68	1.11			22.62	*	31.75	1.05			23.56	***	
32	WV 285	57.11	1.38	**	++	-4.69		28.14	1.46	**	++	-1.30		
33	WV 286	60.64	1.07	**		-4.91		32.38	1.03	**		-0.64		
34	WV 287	56.50	1.67	**		13.64		28.18	1.56	*		28.57	***	
35	WV 288	60.67	1.68	**		3.33		32.87	1.58	**	+	3.31		
36	WV 289	58.31	1.06	**		-0.44		31.55	1.19	**		1.18		
37	WV 290	57.47	0.50			41.66	**	29.60	0.31			53.73	***	
38	WV 291	59.37	1.75	**		3.77		30.09	1.92	**	+	12.69	**	
39	WV 292	54.53	1.04	**		-2.43		27.59	1.00	**		-0.23		
40	WV 293	64.55	1.39	**		3.42		37.84	1.30	**		13.25	**	
41	WV 294	57.89	2.04	**	++	-3.78		30.64	1.79	**	++	-0.97		
42	WV 295	56.28	0.41	**	++	-4.80		25.59	0.38			19.30	***	
43	WV 296	67.39	1.46	**		7.02		41.22	1.47	**		17.14	***	
44	WV 297	58.17	-0.43	**	++	-4.91		32.19	-0.28	**	++	-1.35		
45	WV 298	61.41	0.95			22.70	*	34.30	1.00			34.74	***	
46	WV 299	61.99	0.72	*		0.01		31.97	-0.06		++	0.76		
47	WV 300	55.96	0.72			8.23		27.12	0.61			12.05	**	
48	WV 301	55.79	0.55	**	+	-3.32		27.06	0.51	**	++	-0.96	1	
49	WV 302	66.15	0.21	1	++	-3.34		39.82	1.30	**		1.66	1	
50	WV 303	69.59	0.68	**	++	-4.37		40.01	1.42	*		31.73	***	
	General mean	58.39			<u>ــــــــــــــــــــــــــــــــــــ</u>		-	29.96		1			1	
	Contrai intenni	20.07							I					

Where, b_i and S^2d_i were regression coefficient and deviation from regression, respectively * and ** significant at 5 and 1 per cent levels, respectively when Ho: $b_i = 0$ + and ++ significant at 5 and 1 per cent levels, respectively when Ho: $b_i = 1$

Sr. No.	Construnce	Leaf area (cm ²)						Protein content (%)					
Sr. No.	Genotypes	Mean	ł	Di		S ² d _i		Mean	bi		S ² d	i	
1	WV 254	481.49	0.48	**	++	-228.62		7.91	0.52	**	++	0.01	
2	WV 255	491.52	0.82	*		348.32		8.02	0.92	*		0.20	**
3	WV 256	544.21	1.53	**	++	-166.23		9.04	1.62	**	++	0.03	
4	WV 257	516.56	1.53	**	++	-185.08		8.53	1.61	**	++	0.04	

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5	WV 258	527.43	1.69	**		471.09		8.76	1.83	**	+	0.18	**
6	WV 259	625.27	1.57	**		1029.65	*	10.63	1.73	**		0.31	***
7	WV 260	540.16	1.87	**	++	-309.03		9.04	1.96	**	++	-0.01	
8	WV 261	617.36	0.87			5083.20	***	10.46	0.94			1.90	***
9	WV 262	563.23	1.08			2678.83	**	9.44	1.19			1.06	***
10	WV 263	564.49	1.08	**		-276.51		9.61	1.28	**	++	-0.02	
11	WV 264	483.21	0.64	**	++	-276.93		7.85	0.67	**	++	-0.01	
12	WV 265	522.86	0.74	**	+	-236.00		8.70	0.80	**		0.07	
13	WV 266	516.26	1.12	**		-67.97		8.36	1.31	**	++	-0.01	
14	WV 267	500.84	0.89	**		243.47		8.25	0.84	*		0.31	***
15	WV 268	604.18	0.02			1904.76	**	10.18	0.12			0.66	***
16	WV 269	525.52	2.02	**	++	111.35		8.76	2.13	**	++	0.30	***
17	WV 270	522.20	1.14	**	++	-309.80		8.53	1.36	**	++	-0.01	
18	WV 271	472.44	0.01		++	229.21		7.63	-0.10		++	0.27	***
19	WV 272	540.21	1.61	**		295.49		9.16	1.46	**		0.17	**
20	WV 273	543.57	2.05	**	++	139.11		10.74	0.67	**	++	-0.01	
21	WV 274	535.27	0.79	**		-25.06		8.93	0.87	**		0.06	
22	WV 275	472.59	-0.08		++	-305.40		7.46	-0.07		++	-0.01	
23	WV 276	504.84	1.06	**		-52.69		8.31	1.17	**		0.08	*
24	WV 277	533.19	-0.10		+	1284.54	*	8.87	-0.12		+	0.43	***
25	WV 278	474.88	0.43	**	++	-244.04		7.68	0.50	**	++	0.01	
26	WV 279	505.82	1.34	**		70.67		8.53	1.26	**		0.11	*
27	WV 280	498.20	0.77	**	++	-279.48		8.48	0.57	**	++	-0.02	
28	WV 281	481.26	0.49	**	++	-245.30		7.91	0.52	**	++	0.01	
29	WV 282	654.31	0.87	**		-78.30		10.56	1.62	**		0.37	***
30	WV 283	509.54	1.37	**	++	-316.87		8.48	1.39	**	++	-0.02	
31	WV 284	555.49	1.03			3076.11	**	9.27	1.19			1.07	***
32	WV 285	531.14	1.39	**	++	-291.85		8.87	1.19	**	++	-0.02	
33	WV 286	562.84	1.06	**		-302.63		9.55	1.02	**		-0.02	
34	WV 287	514.86	1.70	**		1860.26	**	8.59	1.74	*		0.82	***
35	WV 288	560.14	1.66	**		883.99		9.44	1.73	**		0.31	***
36	WV 289	539.79	1.13	**		-12.66		9.04	1.09	**		0.15	**
37	WV 290	524.00	0.72	**	++	-269.80		8.87	0.42			1.64	***
38	WV 291	536.86	2.11	**	++	163.79		9.04	2.03	**	++	0.26	**
39	WV 292	500.53	0.83	**		-254.48		8.36	0.97	**		0.04	
40	WV 293	611.96	1.36	**		1207.67	*	10.35	1.36	**		0.34	***
41	WV 294	540.37	1.91	**	++	41.14		8.99	1.93	**	+	0.22	**
42	WV 295	496.23	0.65	1		672.41		8.42	0.37			0.24	**
43	WV 296	640.86	1.47	**		813.78		10.86	1.54	**		0.41	***
44	WV 297	543.93	0.08	**	++	-316.54		9.21	-0.10	**	++	-0.02	
45	WV 298	576.80	0.81	1		3115.11	**	9.61	0.89			1.32	***
46	WV 299	549.95	0.01	1	++	-238.04		9.21	0.02	1	++	-0.01	1
47	WV 300	517.60	0.52	1		2550.98	**	8.53	0.70			0.57	***
48	WV 300	514.23	0.52	*	<u> </u>	81.55		8.48	0.52	*	+	0.08	*
49	WV 301	507.99	0.19	+		1751.35	*	8.48	0.32		++	0.00	+
50	WV 302	634.63	1.03	**		-293.06	1	8.53	0.22	**	+	0.02	+
50	General mean	536.66	1.05	1	1	275.00	1	8.93	0.57	I	· ·	0.05	1
	±SEbi	550.00	0.40					0.75	0.41				
		I	I		ι	0.10	l	l		0.41			

Where, b_i and S^2d_i were regression coefficient and deviation from regression, respectively * and ** significant at 5 and 1 per cent levels, respectively when Ho: $b_i = 0$

+ and ++ significant at 5 and 1 per cent levels, respectively when Ho: $b_i = 1$

Sr.	Constynes		Cru	ide fi	ber (%	6)		Mineral matter (mg/100 g)						
No.	Genotypes	Mean]	bi		S ² d _i		Mean	bi			S ² d _i		
1	WV 254	4.70	0.50	**	++	0.002		1.66	0.48	**	++	0.002		
2	WV 255	4.80	0.85	**		0.035	*	1.68	0.92	**		0.013	**	
3	WV 256	5.37	1.54	**	++	0.015		1.98	1.60	**	++	0.001		
4	WV 257	5.07	1.54	**	++	0.015		1.85	1.54	**	++	0.004	*	
5	WV 258	5.33	1.47	*		0.246	***	1.97	1.55	**		0.045	***	
6	WV 259	6.30	1.66	**		0.103	***	2.48	1.88	**		0.035	***	
7	WV 260	5.37	1.88	**	++	-0.001		1.92	2.05	**	++	0.022	***	
8	WV 261	6.20	0.91			0.651	***	2.42	1.00			0.172	***	
9	WV 262	5.60	1.14			0.360	***	2.12	1.05			0.088	***	
10	WV 263	5.57	0.86	**		-0.001		2.08	0.87	**		0.003	*	
11	WV 264	4.67	0.64	**	++	-0.002		1.67	0.57	**	++	-0.002		
12	WV 265	5.17	0.76	**		0.026	*	1.88	0.79	**		0.004	*	

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10		1.0-				0.004		1.00	1.00				
13	WV 266	4.97	1.26	**	+	-0.001		1.80	1.23	**	+	0.002	
14	WV 267	4.90	0.81			0.109	***	1.75	0.85	*		0.022	**
15	WV 268	6.03	0.12			0.228	***	2.32	0.09			0.059	***
16	WV 269	5.20	2.04	**	+	0.113	***	1.96	2.02	**	++	0.021	***
17	WV 270	5.06	1.27	**	++	-0.005		1.83	1.31	**	++	-0.002	
18	WV 271	4.53	-0.09		++	0.093	***	1.63	-0.23		++	0.002	
19	WV 272	5.70	0.97	**		0.088	***	1.95	1.66	**	+	0.012	**
20	WV 273	5.40	2.09	**	++	0.037	*	2.02	2.05	**	++	0.022	***
21	WV 274	5.30	0.83	**		0.020		1.97	0.77	**		0.004	*
22	WV 275	4.42	-0.06		++	-0.001		1.63	-0.10	**	++	-0.002	
23	WV 276	4.93	1.11	**		0.031	*	1.79	1.08	**		0.006	**
24	WV 277	5.27	-0.12		+	0.149	***	1.93	-0.15		+	0.037	***
25	WV 278	4.57	0.47	**	++	0.005		1.58	0.50	**	++	0.002	
26	WV 279	5.09	1.16	**		0.058	**	1.77	1.46	**		0.008	**
27	WV 280	4.90	0.78	**	++	-0.007		1.72	0.90	**		-0.001	
28	WV 281	4.70	0.50	**	++	0.002		1.67	0.44	**	++	-0.001	
29	WV 282	6.27	1.57	**		0.119	***	2.45	1.68	**		0.042	***
30	WV 283	5.10	1.19	**		0.012		1.83	1.30	**	++	-0.002	
31	WV 284	5.50	1.14			0.360	***	2.12	0.85			0.044	***
32	WV 285	5.20	1.40	**	++	-0.007		1.93	1.30	**	++	-0.002	
33	WV 286	5.60	1.07	**		-0.007		2.10	1.10	**	++	-0.001	
34	WV 287	5.10	1.66	*		0.294	***	2.05	1.28	*		0.050	***
35	WV 288	5.60	1.66	**		0.103	***	2.10	1.68	**		0.042	***
36	WV 289	5.40	0.97	**		0.088	***	1.98	1.09	**		0.009	**
37	WV 290	5.21	0.47			0.635	***	1.93	0.47			0.135	***
38	WV 291	5.50	2.21	**	++	0.091	**	1.97	2.05	**	++	0.022	***
39	WV 292	5.00	0.95	**		-0.004		1.75	1.06	**		0.003	
40	WV 293	6.10	1.38	**		0.154	***	2.38	1.29	*		0.059	***
41	WV 294	5.27	1.99	**	++	0.027	*	1.98	1.85	**	+	0.017	***
42	WV 295	4.83	0.62			0.080	**	1.83	0.25			0.041	***
43	WV 296	6.43	1.47	**		0.136	***	2.57	1.61	**		0.031	***
44	WV 297	5.47	-0.09	**	++	-0.008		2.03	-0.10	**	++	-0.002	
45	WV 298	5.70	0.86			0.449	***	2.20	0.66	1		0.113	***
46	WV 299	5.47	0.02		++	-0.002		2.03	0.03		++	0.001	
47	WV 300	5.10	0.59			0.254	***	1.83	0.72	1		0.043	***
48	WV 301	5.17	0.40	**	++	0.007		1.81	0.57	**	++	0.003	1
49	WV 302	6.00	0.62	**	++	-0.008		1.78	0.37	**	++	-0.001	1
50	WV 303	6.20	0.95	**		-0.004		2.38	0.70	**	++	0.002	1
	General mean	5.33		1	1			1.96					4
	±SEbi		0.41						0.40				
33.71	1 1 0 2 1		1 1						1				

Where, b_i and S^2d_i were regression coefficient and deviation from regression, respectively * and ** significant at 5 and 1 per cent levels, respectively when Ho: $b_i = 0$ + and ++ significant at 5 and 1 per cent levels, respectively when Ho: $b_i = 1$

Table 6: Estimation of mean and stability parameter for iron content (mg/100 g) in little millet

C. N.	Constant of	Iron content (mg/100 g)									
Sr. No.	Genotypes	Mean		bi	0 0	S ² d	i				
1	WV 254	8.86	0.51	**	++	0.01					
2	WV 255	8.95	0.92	**		0.12	**				
3	WV 256	9.79	1.60	**	++	0.01					
4	WV 257	9.37	1.60	**	++	0.01					
5	WV 258	9.75	1.54	**		0.41	***				
6	WV 259	11.10	1.71	**		0.26	***				
7	WV 260	9.79	1.94	**	++	-0.01					
8	WV 261	10.96	0.91			1.35	***				
9	WV 262	10.21	1.02			0.51	***				
10	WV 263	10.07	0.88	**		0.01					
11	WV 264	8.91	0.57	**	++	-0.01					
12	WV 265	9.51	0.79	**		0.04					
13	WV 266	9.23	1.31	**	++	-0.01					
14	WV 267	9.23	0.70			0.36	***				
15	WV 268	10.73	0.11			0.45	***				
16	WV 269	9.61	2.05	**	+	0.23	***				
17	WV 270	9.37	1.35	**	++	0.01					
18	WV 271	9.05	-0.76	**	++	0.07	*				
19	WV 272	9.89	1.46	**		0.09	*				
20	WV 273	9.81	2.05	**	++	0.14	**				

21	WV 274	9.81	0.68	**	++	-0.01	
21 22	WV 274 WV 275	8.49	-0.07		++	-0.01	+
23	WV 275	9.15	1.18	**	++	0.08	*
23	WV 270	9.15	-0.14			0.08	***
24	WV 277 WV 278	8.77		**	+		
-			0.40	**	++	0.01	*
26	WV 279	9.28	1.36	**		0.07	*
27	WV 280	9.09	0.84	**	++	-0.01	
28	WV 281	9.05	0.23		++	0.01	
29	WV 282	11.05	1.61	**		0.29	***
30	WV 283	9.47	1.21	**	++	-0.01	
31	WV 284	9.98	1.16			0.78	***
32	WV 285	9.75	1.38	**	++	-0.02	
33	WV 286	10.12	1.11	**	++	-0.02	
34	WV 287	10.24	1.18	**		0.03	
35	WV 288	10.12	1.71	**		0.26	**:
36	WV 289	10.11	1.42	*		0.36	**:
37	WV 290	9.61	0.47			1.33	**:
38	WV 291	9.89	1.93	**	++	0.11	**
39	WV 292	9.23	0.96	**		0.02	
40	WV 293	10.82	1.41	*		0.36	**:
41	WV 294	9.71	1.98	**	++	0.07	*
42	WV 295	9.05	0.60			0.33	**:
43	WV 296	11.29	1.51	**		0.33	**:
44	WV 297	9.70	0.17	**	++	-0.01	
45	WV 298	10.31	0.96			0.88	**:
46	WV 299	9.93	0.03		++	0.01	
47	WV 300	9.37	0.70			0.36	**:
48	WV 301	9.51	0.32		++	0.02	1
49	WV 302	11.33	0.76	**		0.07	*
50	WV 303	11.52	0.68	**		0.03	
	General mean	9.79	0.00	-1	II	0.02	1
	±SEbi	,,,,,	0.41				
	_5201		0				

Sr.	Genotypes		Grain yield per plant (g)					
No.		Mean		bi		S ² d _i		
1	WV 254	7.84	0.45	**	++	-0.15		
2	WV 255	7.29	0.76	**		-0.13		
3	WV 256	10.23	0.87	**		-0.14		
4	WV 257	8.74	0.95	**		-0.08		
5	WV 258	9.95	1.01	**		-0.16		
6	WV 259	12.30	1.46	**	++	-0.14		
7	WV 260	10.14	0.22			0.30		
8	WV 261	10.66	0.34			1.14	**	
9	WV 262	9.23	1.08	**		-0.16		
10	WV 263	8.65	1.63	**	++	-0.13		
11	WV 264	8.09	0.87	**	+	-0.16		
12	WV 265	8.13	0.25		+	-0.07		
13	WV 266	8.47	0.50			0.01		
14	WV 267	8.03	-0.35			0.61	*	
15	WV 268	10.22	-1.05		++	0.19		
16	WV 269	10.08	1.96	**	+	-0.04		
17	WV 270	8.36	1.20	**		-0.10		
18	WV 271	7.66	0.69	**	++	-0.16		
19	WV 272	9.80	0.54	*	+	-0.12		
20	WV 273	11.15	0.95	**		-0.13		
21	WV 274	8.04	1.38	**	+	-0.14		
22	WV 275	7.52	1.09	**		-0.13		
23	WV 276	7.61	1.00			0.19		
24	WV 277	8.49	1.60	**	++	-0.16		
25	WV 278	7.14	0.67	**	++	-0.16		
26	WV 279	8.45	1.30	**		-0.01		
27	WV 280	7.42	0.68	**	++	-0.16		
28	WV 281	7.16	0.63	**	++	-0.16		

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29	WV 282	11.25	1.93			0.92	**
30	WV 283	8.37	1.32	**		-0.12	
31	WV 284	8.46	2.38	*		0.45	
32	WV 285	8.03	2.28	**		0.15	
33	WV 286	8.56	2.32	**		0.16	
34	WV 287	8.58	2.58	**	++	-0.10	
35	WV 288	10.01	1.84	**	+	-0.08	
36	WV 289	8.55	1.59	**		-0.07	
37	WV 290	8.08	0.54			0.18	
38	WV 291	10.52	2.08			0.89	*
39	WV 292	7.28	0.87	**		-0.15	
40	WV 293	10.53	2.13	**		0.13	
41	WV 294	10.71	1.11	*		0.04	
42	WV 295	7.56	0.89	**		-0.15	
43	WV 296	10.97	3.57	**	++	-0.08	
44	WV 297	8.23	0.23	**	++	-0.16	
45	WV 298	9.24	1.11			0.19	
46	WV 299	8.07	0.00	**	+	-0.16	
47	WV 300	7.79	0.86	**		-0.15	
48	WV 301	10.97	0.52	**	++	-0.16	
49	WV 302	10.90	-1.98	**	++	-0.16	
50	WV 303	12.51	-0.83	**	++	-0.15	
	General mean	9.04		•	•		
	±SEbi		0.51				

Where, b_i and S²d_i were regression coefficient and deviation from regression, respectively

 \ast and $\ast\ast$ significant at 5 and 1 per cent levels, respectively when Ho: $b_i=0$

+ and ++ significant at 5 and 1 per cent levels, respectively when Ho: $b_i = 1$

Table 8: Classification of genotypes by number based on their adaptation in different environments in little millet

SI. No.		Number of suitable genotypes				
	Quality Characters	Average stability and wide/ general adaptability	Stable and adapted to poor environment	Stable and adapted to better environment		
1.	Grain yield per plant (g)	6	4	4		
2.	Hulling (%)	10	1	2		
3.	Chlorophyll content (mg/100 g fresh weight)	3	3	3		
4.	Leaf area (cm ²)	8	1	5		
5.	Protein content (%)	1	2	3		
6.	Crude fiber (%)	3	2	2		
7.	Mineral matter (mg/100 g)	-	2	2		
8.	Iron content (mg/100 g)	3	1	1		

Table 9: Classification of genotypes by name based on their adaptation in different environments in little millet

Sr.		Name of genotypes suitable for				
No.	Quality Characters	Average stability and wide/ general adaptability	Stable and adapted to poor environment	Stable and adapted to better environment		
1.	Grain yield per plant (g)	WV 262, WV 258, WV 256, WV 293, WV 294 and WV 273	WV 302, WV 303, WV 301 and WV 272	WV 259, WV 288, WV 269 and WV 296		
2.	Hulling (%)	WV 256, WV 291, WV 263, WV 286, WV 288, WV 299, WV 293, WV 259, WV 282 and WV 296	WV 303	WV 260 and WV 273		
3.	Chlorophyll content (mg/100 g fresh weight)	WV 289, WV 286 and WV 302	WV 297, WV 265 and WV 263	WV 288, WV 294 and WV 260		
4.	Leaf area (cm ²)	WV 289, WV 272, WV 288, WV 286, WV 263, WV 303, WV 296 and WV 282	WV 297	WV 256, WV 260, WV 294, WV 273 and WV 291		
5.	Protein content (%)	WV 286	WV 297 and WV 273	WV 263, WV 256 and WV 260		
6.	Crude fiber (%)	WV 263, WV 286 and WV 303	WV 297 and WV 302	WV 256 and WV 260		
7.	Mineral matter (mg/100 g)	-	WV 297 and WV 303	WV 286 and WV 256		
8.	Iron content (mg/100 g)	WV 263, WV 287 and WV 303	WV 274	WV 286		

Conclusion

The overall picture of stability of genotypes to different characters, it could be concluded that, genotypes viz., WV 262, WV 258, WV 256, WV 293 and WV 273 were found to be average stable over environments for grain yield per plant with one or more yield contributing characters. The protein content (%) was moderately affected by environmental

changes. While, the grain yield per plant (g), crude fiber (%), mineral matter (mg/100 g) and iron content (mg/100 g) were less influenced by environmental fluctuations. As the genotype WV 294 was found to be stable over environment for grain yield per plant but with none of the yield contributing characters. Hence, it was suggested that in order to identify stable genotypes, actual testing under variable environments including favourable and unfavourable would be advantageous. During selection, the attention should be paid to the phenotypic stability of characters directly related to grain yield per plant in little millet. Estimates of environmental indices indicated that Waghai location was favourable for most of the yield contributing characters along with quality parameters followed by Navsari and Vanarasi. Estimates of environmental indices indicated that Waghai location was favourable for most of the yield contributing characters along with quality parameters followed by Navsari and Vanarasi.

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