

1 **ASSESSMENT OF GENOTYPE X ENVIRONMENT INTERACTION AND STABILITY**
2 **OF PROMISING SUGARCANE GENOTYPES FOR DIFFERENT AGRONOMIC**
3 **CHARACTERS IN PESHAWAR VALLEY**

4 Mohammad Tahir^{*}, Hidayatur Rahman[†], Mohammad Khalid^{*}, Amjad Ali^{*}, and Sajjad Anwar^{*}
5 Sugar Crops Research Institute, Mardan, NWFP, Pakistan

6 **Abstract**

7 Sugarcane germplasm screening is a regular feature of the breeding
8 program at Sugar Crops Research Institute, Mardan, NWFP, Pakistan.
9 Sixteen genotypes were evaluated in the final stages of selection in three
10 different environments for G x E interaction and stability. Combined
11 analysis of variance showed highly significant variances for Environments
12 (E), Genotypes (G), and their interaction (G x E). The effect of
13 environments was much pronounced for all the characters signifying their
14 importance in the performance of genotypes. None of the genotypes was
15 stable for all characters. However, genotype MS-94-CP-90 showed a
16 relative stability for three characters viz. germination percentage, cane
17 yield (t/ha) and millable canes.

18 Key words: **G x E Interaction, Environments, Stability, Sugarcane**

19 **Introduction**

20 Sugarcane is an important field crop of the North West Frontier Province of Pakistan. It
21 is cultivated on an area of 0.1 million hectare with a production of 4.65 million tones and
22 per hectare cane yield amounting to 46 tones per hectare (Agriculture Statistics of
23 Pakistan, 2006-07). Sugar Crops Research Institute (SCRI), Mardan, is mandated with
24 the development of sugarcane varieties which are high yielding, disease and frost

^{*} Research Officers, Sugar Crops Research Institute, Mardan, KPK, Pakistan

[†] Professor, Department of PB&G, KPK Agriculture University, Peshawar, Pakistan

25 resistant and with better quality. Germplasm is procured both from within the country
26 and from abroad. It is tested in various selection stages and advanced to final stages of
27 selection. Varieties are sought which would interact minimally with the environment so
28 that their performance could be generalized over a range of environments.

29 However, Genotype by environment (G x E) interaction complicates selection and
30 testing of plant genotypes. Measuring G x E is important in order to determine an
31 optimum strategy for selecting genotypes with adaptation to target environments
32 (Romagosa et al., 1993; De Lacy et al., 1994; Annicchiarico, 1997). Productivity
33 stability is shown by some cane varieties in both predictable and unpredictable
34 environments. In a predictable environment (i.e. climatic, soil type, day length and
35 controllable variables such as fertilization, sowing dates and harvesting methods), a
36 high level of genotype and environmental interaction was desirable, so as to ensure a
37 maximum yield or financial return; whereas, in an unpredictable environment (inter and
38 intra-season fluctuation, fluctuation in quantity and distribution of rainfall and prevailing
39 temperature), a low level of interaction is desirable so as to ensure maximum uniformity
40 of performance over a number of locations or seasons (Khan, 1981). However the
41 performance of genotypes in favorable environments does not indicate their adaptability
42 and stability. Hence a breeder is always in a hunt for suitable high yielding genotypes
43 which would interact minimal with the environments and are stable.

44
45 The current study was undertaken to assess genotype by environment interaction and
46 stability of 16 sugarcane genotypes for different plant and yield characters.

47

48

49

50 **Materials and Methods**

51 Three experiments were grown in three environments: 2 at Sugar Crops Research
52 Institute during 2005-06 and 2006-07 and one at Harichand Seed Multiplication Farm
53 during 2005-06. The experimental material comprised 16 advanced lines/varieties
54 including two checks laid out in randomized complete block design. Data were recorded
55 on germination percentage, number of tillers, plant height, cane yield and millable
56 canes.

57 The data were analyzed using MSTATC version 2.01 (Nissen, 1983). Homogeneity of
58 variances was run after Gomez and Gomez (1984). Combined analyses of variance and
59 stability parameters were worked out using PBSTAT online version 1.0 (Willy *et al*
60 2008).

61 **Results and Discussion**

62 **Mean Squares for individual environments:**

63 The mean squares for individual environments are given in table 1. The range of CV for
64 all the characters over the three environments was less than 20 (Gomez and Gomez
65 1984) and hence were forwarded for combined analysis of variance. Genotypic
66 variances were significant for the characters under study however showing a non
67 significant effect for Number of tillers.

68

69

70

71 **Table 1: Mean squares for the characters over individual environment**

Environments	Source of Variation	D.F	Germination %age	No. of Tillers	Plant Height	Cane Yield	Millable cane
E1	Reps	2	12.771 ^{ns}	2541.396 ^{ns}	1563.271 ^{**}	50.333 ^{ns}	20.813 ^{ns}
	Genotypes	15	221.022 ^{**}	8645.106 ^{**}	577.654 ^{**}	312.706 ^{**}	177.321 ^{**}
	Error	30	43.726	639.418	138.538	111.556	22.79
	CV		12.1	10.61	8.1	12.62	4.94
E2	Reps	2	134.021 [*]	4497.646 [*]	280.750 ^{ns}	180.063 ^{ns}	21 ^{ns}
	Genotypes	15	120.465 ^{**}	4456.800 ^{**}	453.222 ^{**}	389.343 ^{**}	941.443 ^{**}
	Error	30	31.932	1015.646	148.106	91.351	38.822
	CV		11.16	13.08	7.52	16.18	7.03
E3	Reps	2	6.083 ^{ns}	446.333 ^{ns}	63.521 ^{ns}	59.313 ^{ns}	95.063 ^{**}
	Genotypes	15	59.194 ^{**}	642.706 ^{ns}	1626.376 ^{**}	90.154 ^{**}	47.699 ^{**}
	Error	30	2.61	378.156	395.876	20.913	15.507
	CV		4.74	14.25	11.43	8.74	9.17

72 ns: non-significant ** : Highly Significant * : Significant

73 **Mean performance of the genotypes over environments:**

74 Genotypic means are given in table 2. Mean performance of the genotypes for
75 germination percentage showed that MS-94-CP-90, MS-92-CP-1100, and MS-91-CP-
76 965 performed better than the rest with a mean range of 51 to 54. In number of tillers,
77 genotypes Mardan 93, MS-94-CP 90, and MS-91-CP 965 outperformed the rest of the
78 genotypes. MS-91-CP-288, Malakand 17, MS-94-CP-90 and MS-92-CP-623 were taller
79 than the rest of the genotypes. Regarding cane yield (t/ha) MS-91-CP-920, MS 92-Cp-
80 623, MS-91-CP-623, and CP 77/400 performed well. Higher Millable canes were given
81 by MS-92-CP-623, Mardan 93, and MS-94-CP-90, respectively.

82

83

84 **Table 2: Genotypic means of the 16 genotypes combined over environments.**

S.No.	Genotype	Germination %age	No. of Tillers	Plant Height	Cane Yield (t/ha)	Millable Cane*
1	Malakand 17	40.22 ^{de}	159.67 ^{ef}	177.22 ^{ab}	53.00 ^d	47.00 ^d
2	MS-92-CP-623	44.11 ^{bcd}	225.67 ^{abc}	168.44 ^{abc}	71.11 ^{ab}	83.89 ^a
3	MS-92-CP-624	45.78 ^{bcd}	198.89 ^{bcdef}	163.11 ^{abc}	67.67 ^{abcd}	77.78 ^{abc}
4	MS-91-CP-611	34.89 ^e	189.22 ^{cdef}	149.56 ^{cd}	60.33 ^{abcd}	73.22 ^c
5	MS-91-CP-572	38.33 ^{de}	210.67 ^{bcd}	157.67 ^{bcd}	65.22 ^{abcd}	76.33 ^{abc}
6	MS-91-CP-288	45.00 ^{bcd}	204.56 ^{bcd}	183.44 ^a	68.11 ^{abc}	77.56 ^{abc}
7	AEC-86-347	47.00 ^{abcd}	202.00 ^{bcde}	166.33 ^{abc}	66.89 ^{abcd}	76.33 ^{abc}
8	Mardan 93	42.11 ^{cde}	258.22 ^a	155.33 ^{bcd}	69.11 ^{abc}	82.11 ^{ab}
9	MS-91-CP-471	50.00 ^{abc}	211.00 ^{bcd}	132.33 ^d	63.22 ^{abcd}	76.78 ^{abc}
10	MS-91-CP-623	50.56 ^{abc}	203.33 ^{bcde}	154.00 ^{bcd}	70.89 ^{ab}	77.44 ^{abc}
11	MS-91-CP-920	39.44 ^{de}	177.33 ^{def}	144.44 ^{cd}	72.22 ^a	79.33 ^{abc}
12	MS-91-CP-965	51.67 ^{ab}	234.56 ^{ab}	160.56 ^{abc}	57.00 ^{bcd}	74.00 ^{bc}
13	MS-92-CP-1100	51.44 ^{ab}	200.22 ^{bcdef}	149.22 ^{cd}	66.00 ^{abcd}	78.78 ^{abc}
14	MS-94-CP-90	54.78 ^a	234.67 ^{ab}	168.56 ^{abc}	63.67 ^{abcd}	81.11 ^{abc}
15	CPF-236	45.22 ^{bcd}	156.89 ^f	166.44 ^{abc}	56.00 ^{cd}	74.67 ^{bc}
16	CP 77/400	50.22 ^{abc}	231.33 ^{abc}	165.33 ^{abc}	70.00 ^{abc}	75.33 ^{abc}

85 * Means followed by the same letters do not differ significantly.

86 **Genotype – Environment Analysis**

87 G x E analysis in table 3 revealed highly significant variances for Environments (E),
88 Genotypes (G), and their interaction (G x E). The effect of environments was much
89 pronounced for all the characters signifying their importance in the performance of
90 genotypes. However, mean square differences were also significant for genotypes
91 showing that the differences among the genotypes were persistent along the
92 environments. These were higher than G x E interaction mean squares indicating the
93 varied response of the genotypes was a permanent characteristic for locations. Similar
94 results were found by Tai et al (1982) wherein they found significant cultivars

95 differences over interactions. However, variance components analyses showed that
 96 interaction variance was larger for the characters except germination percentage.
 97 Higher phenotypic variance revealed the control of environmental factors in the
 98 genotypes. Similar results were found by Singh and Singh, 1987, wherein they found
 99 significant mean squares for environments, Genotypes and their interaction for all the
 100 characters studied.

101 **Table 3: Mean Squares for environments and genotypes in Combined analysis of**
 102 **variance**

Source	df	Germination	Tillering	Plant Height	Cane Yield	Millable Canes
Environments(E)	2	7215.05**	175140.36**	9559.15**	13109.42**	39277.75**
REP*E	6	50.96 ^{ns}	2495.13**	635.85*	96.57 ^{ns}	45.63 ^{ns}
Genotypes (G)	15	282.97**	6726.29**	1443.88**	301.92**	604.07**
G*E	30	58.86**	3509.16**	606.69**	245.14**	281.19**
Error	90	25.97	677.74	226.51	74.61	25.54
CV		11.16	12.63	9.4	13.28	6.67

Variationes

V_p	31.44	747.37	160.43	33.55	67.12
V_G	24.9	357.46	93.02	6.31	35.88
$V_{G \times L}$	10.96	943.81	126.73	56.85	85.22
h^2_{bs}	79.2	47.83	57.98	18.81	53.45

103

104 V_G = Genotypic Variance $V_{G \times L}$ = Interaction Variance V_p = Phenotypic variance h^2_{bs} = Broad Sense
 105 Heritability.

106 Stability Analysis

107 A cultivar with 'b' value less than 1.0 has above average stability and is specially
 108 adapted to low-performing environments, a cultivar with 'b' value greater than 1.0 has
 109 below average stability and is specially adapted to high performing environments and a
 110 cultivar with 'b' value equal to 1.0 has average stability and is well or poorly adapted to

111 all environments depending on having a high or low mean performance (Finlay &
112 Wilkinson 1963) but a cultivar with $b = 1.00$ and $S^2d = 0.00$ may be defined as stable
113 (Eberhart & Russell 1966).

114
115 Regression values for germination percentage (Table 4) indicated that genotypes MS-
116 91-CP-288, AEC-86-347, MS-91-CP-623 and MS-94-CP-90 were close to 1 and
117 deviation from regression smaller than the others showing a relative stability for this
118 character. Genotypes MS-91-CP-471, MS-91-CP-965, MS-92-CP-1100 and CP 77/400
119 had b_i values greater than 1 showing these genotypes performed better under favorable
120 environments while the rest of the genotypes exhibited a slope value less than 1
121 indicating that they were better performing under unfavorable conditions. For number of
122 tillers genotypes MS-92-CP-624, MS-91-CP-572, AEC-86-347 and MS-91-CP-623 had
123 a b_i value close to 1 while genotypes Malakand 17, MS-91-CP-288, MS-91-CP-920 and
124 CPF-236 had less than 1. For plant height only genotype MS-92-CP-624 had a value
125 close to 1, 8 genotypes had a value less than 1 while remaining genotypes exhibited
126 slope value more than 1 and 5 genotypes gave a value less than 1 and remaining more
127 than 1. For cane yield (t/ha) genotypes Mardan 93, MS-92-CP-1100, MS-94-CP-90 and
128 CP 77/400 had a value close to 1 and MS-91-CP-288 genotypes gave a value less than
129 1. For millable canes genotypes MS-91-CP-288, AEC-86-347, MS-91-CP-471, MS-94-
130 CP-90, CPF-236, and CP 77/400 exhibited a b_i values close to 1. Among the genotypes
131 studied none showed stability for all characters. However, genotype MS-94-CP-90
132 showed a relative stability for three characters viz. germination percentage, cane yield
133 (t/ha) and millable canes. It can be concluded that G x E interaction and stability

134 analysis should become a regular feature of any breeding program so that genotypes
 135 could be sorted out which perform better under multiple environments.

136 **Table 4: Regression slope, and deviation from regression for 16 genotypes**

S. No.	Genotype	Germination %age			No. of Tillers			Plant Height		
		Mean	b_i^*	SD_i^{**}	Mean	b_i	SD_i	Mean	b_i	SD_i
1	Malakand 17	40.22	0.85	1.86	159.67	0.2	14.56	177.22	2.17	24.71
2	MS-92-CP-623	44.11	0.89	4.61	225.67	1.19	57.36	168.44	1.37	9.86
3	MS-92-CP-624	45.78	0.61	2	198.89	1.08	4.81	163.11	1.04	3.85
4	MS-91-CP-611	34.89	0.63	3.48	189.22	1.16	7.23	149.56	1.69	15.55
5	MS-91-CP-572	38.33	0.93	2.68	210.67	1.09	16.89	157.67	0.59	0.88
6	MS-91-CP-288	45	0.95	2.51	204.56	0.73	25.2	183.44	1.59	17.26
7	AEC-86-347	47	1.08	1.68	202	0.95	96.17	166.33	2.45	0.03
8	Mardan 93	42.11	0.88	1.52	258.22	1.31	0.45	155.33	1.53	5.92
9	MS-91-CP-471	50	1.32	7.14	211	0.91	31.69	132.33	0.41	5.75
10	MS-91-CP-623	50.56	0.99	0.7	203.33	1.06	2.84	154	0.52	2.28
11	MS-91-CP-920	39.44	0.89	2.8	177.33	0.54	26.51	144.44	0.86	12.1
12	MS-91-CP-965	51.67	1.58	5.95	234.56	1.69	31.18	160.56	0.56	4.76
13	MS-92-CP-1100	51.44	1.36	3.74	200.22	1.12	0.97	149.22	0.24	16.89
14	MS-94-CP-90	54.78	1.02	5.9	234.67	1.28	3.61	168.56	-0.43	11.66
15	CPF-236	45.22	0.64	2.7	156.89	0.3	12.56	166.44	1.42	15.89
16	CP 77/400	50.22	1.39	3.3	231.33	1.4	39.07	165.33	0.01	11.47

137 Regression Slope ** Deviation from Regression

138

139

140

141

142

143

144

145

146

147 **Table 4 (Contd.): Regression slope, and deviation from regression for 16**
 148 **genotypes.**

S.No.	Genotype	Cane Yield (T/ha)			Millable Canes		
		Mean	b_i^*	SD_i^{**}	Mean	b_i	SD_i
1	Malakand 17	53	1.18	33.45	47	0.3	34.4
2	MS-92-CP-623	71.11	1.22	2.35	83.89	1.23	6.95
3	MS-92-CP-624	67.67	1.16	2.27	77.78	1.12	2.59
4	MS-91-CP-611	60.33	0.77	6.82	73.22	1.17	3.85
5	MS-91-CP-572	65.22	0.75	9.81	76.33	0.88	0.29
6	MS-91-CP-288	68.11	0.75	10.33	77.56	1.02	15.43
7	AEC-86-347	66.89	0.84	0.25	76.33	0.99	5.56
8	Mardan 93	69.11	0.97	3.03	82.11	1.11	1.94
9	MS-91-CP-471	63.22	1.43	7.71	76.78	1.07	2.64
10	MS-91-CP-623	70.89	1.6	0.4	77.44	1.1	2.69
11	MS-91-CP-920	72.22	1.37	1.37	79.33	0.85	1.78
12	MS-91-CP-965	57	0.41	7.41	74	0.81	1.58
13	MS-92-CP-1100	66	1.07	7.79	78.78	1.21	2.38
14	MS-94-CP-90	63.67	0.9	7.27	81.11	1.08	2.93
15	CPF-236	56	0.6	1.63	74.67	1	5.37
16	CP 77/400	70	0.98	1.9	75.33	1.06	1.17

149 Regression Slope ** Deviation from Regression

150

151 References

152 Annicchiarico, P. 1997. Joint regression vs. AMMI analysis of genotype-environment
 153 interactions for cereals in Italy. *Euphytica* 94: 53-62.

154 Anonymous. 2006-07. Agriculture Statistics of Pakistan, Ministry of Food, Agriculture
 155 and Livestock Division (MINFAL). Government of Pakistan.

156 De Lacy IH, Fox PN, Corbett JD, Crossa J, Raaram S, Fischer RA, Van Ginkel M.

157 1994. Long-term associations of locations for testing spring wheat. *Euphytica*

158 72: 95-106.

- 159 Ebberhart, S.A., and W.A. Russell. 1966. Stability parameters for comparing
160 varieties. *Crop Science*, 6: 36-40.
- 161 Finlay, R.W. and G.N. Wilkinson. 1963. The analysis of adaptiveness in a breeding
162 programme. *Australian Journal of Agricultural Research*, 14: 742-754.
- 163 Gomez, K.A. and A. A. Gomez. 1984. Analysis over sites. *Statistical Procedures for*
164 *Agriculture Research*. John Willy and Sons, Inc. pp: 332-339.
- 165 Khan, A.Q. 1981. Varietal buffering in sugarcane. *Indian Sugar*, 31: 409-411.
- 166 Nisson, O. 1983. MSTATC: A micro computer program for the design, management
167 and analysis of agronomic research experiments. Michigan State University.
168 Copyright 1988. Distribution- June 1991.
- 169 Romagosa I, Fox PN, García del Moral LF, Ramos JM, García del Moral B, Roca de
170 Togores F, Molina-cano JL (1993) Integration of statistical and physiological
171 analyses of adaptation of near-isogenic barley lines. *Theor. Appl. Genet.*86:
172 822-826.
- 173 Singh, H.N.; Singh, T.K. 1987. Genotype-environment interaction for yield and yield
174 attributes in sugarcane. *Indian Journal of Agricultural Science*. v. 57(5) p.
175 309-313
- 176 Suwarno, W. B., Sobir, H. Aswidinnoor, and M. Syukur. 2008. PBSTAT: A web-
177 based statistical analysis software for participatory plant breeding. 3rd
178 International Conference on Mathematics and Statistics (ICoMS-3). Bogor
179 Agriculture University, Indonesia, August 5-6.

- 180 Tai, P.Y.P., E.R. Rice, V. Chew and J.D. Miller. 1982. Phenotypic stability analyses
181 of sugarcane cultivar performance tests. *Crop Science*, 22: 1179-1184.

UNDER PEER REVIEW