

28 INTRODUCTION

29 Sugarcane is an important field crop of the Khyber Pakhtunkhwa Province of Pakistan. It is
30 cultivated on an area of 0.1 million hectare with a production of 4.65 million tones and cane
31 yield amounting to 46 tones per hectare^[1]. Sugar Crops Research Institute (SCRI), Mardan,
32 is mandated with the development of sugarcane varieties with high yield, disease and frost
33 resistance and accompanied with better quality. Germplasm is procured from within the
34 country and abroad as well. It is tested in various selection stages and advanced to final
35 stages of selection. Varieties are sought which would interact minimally with the environment
36 so that their performance could be generalized over a wide range of environments.

37 Genotype by environment (G x E) interactions considerably complicates selection and
38 testing of plant genotypes, particularly when exposed to diverse set environments.
39 Measuring G x E is important in order to determine an optimum strategy for selecting
40 genotypes with adaptation to target environments^[2,3]. Productivity stability is shown by some
41 cane varieties in both predictable and unpredictable environments. In a predictable
42 environment (i.e. climatic, soil type, day length and controllable variables such as
43 fertilization, sowing dates and harvesting methods), a high level of genotype and
44 environmental interaction was desirable, so as to ensure a maximum yield and financial
45 return; whereas, in an unpredictable environment (inter and intra-season fluctuation,
46 fluctuation in quantity and distribution of rainfall and prevailing temperature), a low level of
47 interaction is desirable so as to ensure maximum uniformity of performance over a number
48 of locations or seasons^[4]. However, the performance of genotypes in favorable environments
49 does not indicate their adaptability and stability. Hence, a breeder is always in a hunt for
50 suitable high yielding genotypes which would interact minimal with the environments and are
51 stable over a series of environments.

52
53 The current study was undertaken to assess genotype by environment interaction and
54 stability of 16 sugarcane genotypes for different plant and yield characters.

55 MATERIALS AND METHODS

56 Three experiments were grown in three environments: two at Sugar Crops Research
57 Institute during 2005-06 and 2006-07 and one at Harichand Seed Multiplication Farm during
58 2005-06. The experimental material comprised of 16 advanced lines/varieties mostly of CP
59 (Canal Point, Florida) origin, including two checks (Mardan 93, and CP 77/400), laid out in
60 randomized complete block design. Data were recorded on germination percentage, number
61 of tillers, plant height, cane yield and millable canes.

62 The data were analyzed using MSTATC version 2.01^[6]. Combined analyses of variance and
63 stability parameters were worked out using PBSTAT online version 1.0^[6]. It calculates
64 regression coefficients (bi) values by regressing individual variety means on the mean yield
65 of all varieties for each environment.

66 RESULTS AND DISCUSSION

67 Mean Squares for individual environments:

68 The mean squares for individual environments are given in Table 1. The range of cv for all
69 the characters studied over the three environments was less than 20^[7] and hence were
70 forwarded for combined analysis of variance. Genotypic variances were significant for the
71 characters under study except a no-significant effect for number of tillers only.

72

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

Environments	Source of Variation	D.F	Germination %	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

74 ns: non-significant ** : Singinificant at P=0.01 * : Significant at P=0.05

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

76 Genotypic means are given in Table 2. Mean performance of the genotypes for germination
77 percentage showed that MS-94-CP-90, MS-92-CP-1100, and MS-91-CP-965 performed
78 better than the rest with a mean range of 51 to 54. For number of tillers, genotypes Mardan
79 93, MS-94-CP 90, and MS-91-CP 965 outperformed the rest of the genotypes. MS-91-CP-
80 288, Malakand 17, MS-94-CP-90 and MS-92-CP-623 were taller than the rest of the
81 genotypes. Regarding cane yield (t/ha) MS-91-CP-920, MS 92-Cp-623, MS-91-CP-623, and
82 CP 77/400 performed well above average. Higher Millable canes were given by MS-92-CP-
83 623, Mardan 93, and MS-94-CP-90, respectively.

84

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

S.No.	Genotype	Germination %	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 ^{bcd}	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 ^{bcd}	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 ^{bcd}	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 ^{bcd}	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}

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

87 **Genotype x Environment Analysis**

88 G x E analysis in Table 3 revealed highly significant variances for Environments (E),
89 Genotypes (G), as well as their interaction (G x E). The effect of environments was much
90 pronounced for all the characters signifying its importance in the performance of genotypes.
91 Mean square differences were also significant for genotypes showing that the differences
92 among the genotypes were persistent over the environments. These were higher than G x E
93 interaction mean squares, indicating the varied response of the genotypes was a permanent
94 characteristic for locations. Similar results were reported by Tai *et al.*^[8] wherein they found
95 significant cultivar differences over interactions. Variance components analyses exhibited
96 that interaction variance was larger for all characters except germination percentage. Higher

97 phenotypic variance revealed the impact of environmental factors on the genotypes. Similar
 98 results have also been reported by Singh and Singh^[9], wherein they found significant mean
 99 squares for environments, **genotypes** and their interaction for various characters studied
 100 sugarcane.

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
Variances						
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 _{GxL}		10.96	943.81	126.73	56.85	85.22
h ² _{bs}		79.2	47.83	57.98	18.81	53.45

103 ns: non-significant **: Significant at P=0.01 *: Significant at P=0.05

104 V_G= Genotypic Variance V_{GxL}= Interaction Variance V_p= Phenotypic variance h²_{bs}= Broad Sense Heritability.

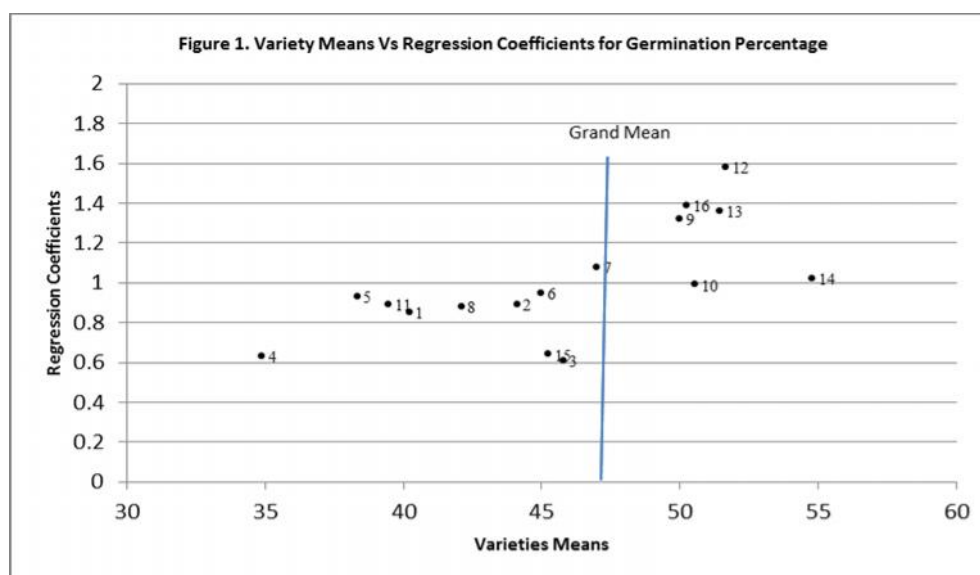
105 Stability Analysis

106 A cultivar with 'b' value less than 1.0 has above average stability and is anticipated to
 107 perform well under unfavorable environments, while a cultivar with 'b' value greater than 1.0
 108 has below average stability and is specially adapted to improved environments. On the other
 109 hand a cultivar with 'b' value equal to 1.0 has average stability and is expected to be well
 110 adapted to all environments accompanied with high mean performance^[10].

111
 112

113 **Germination Percentage**

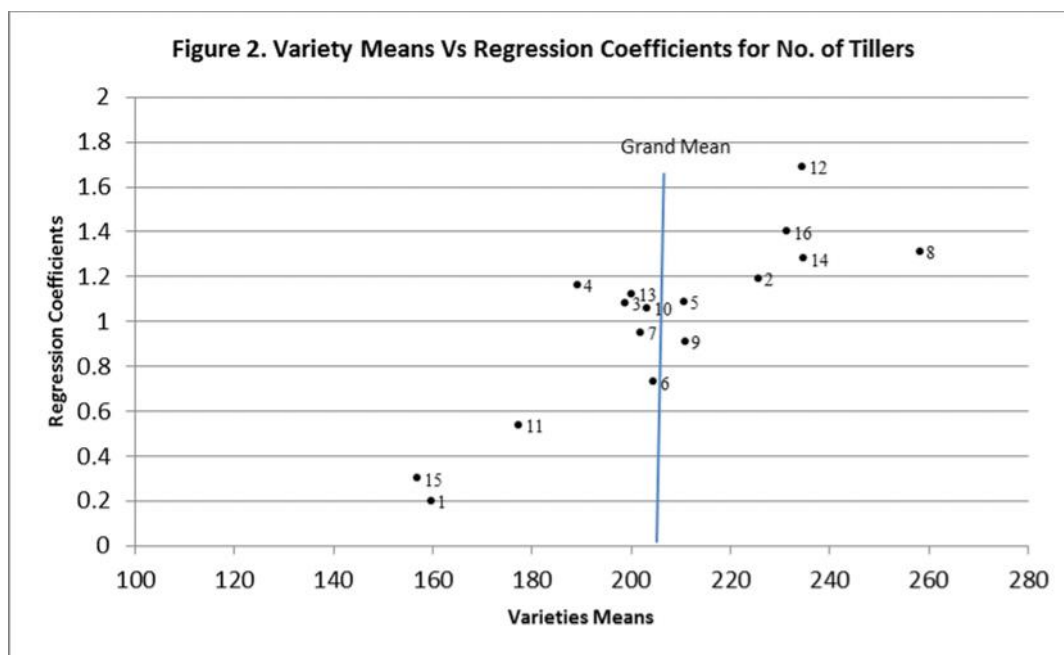
114 Regression values for germination percentage (Table 4) indicated that genotypes MS-91-
 115 CP-623 and MS-94-CP-90 were having regression coefficient value close to unity, showed
 116 average stability for this character with means higher than grand mean and were therefore,
 117 well adapted to all environments. Genotypes MS-91-CP-471, MS-91-CP-965, MS-92-
 118 CP1100 and CP 77/400 (Figure 1) showed regression values above unity indicating that they
 119 had below average stability and were expected to perform better under favorable
 120 environments. The rest of the genotypes exhibited a slope value less than 1 indicating that
 121 they were comparatively better performing under unfavorable conditions.



122

123 **Number of tillers**

124 For number of tillers, genotypes MS-91-CP-572 and MS-91-CP-471 exhibited regression
 125 coefficient values closer to unity accompanied with higher mean values. This indicated that
 126 these genotypes performed well under all tested environments. Figure 2 shows that
 127 genotypes MS-91-CP-288, MS-91-CP471, MS-91-CP-920 and CPF-236 had values
 128 regression coefficient values below 1 and hence were expected to perform well under
 129 unfavorable environments. The rest of the genotypes had values more than 1 and were
 130 supposed to be specifically adapted to favorable environments.



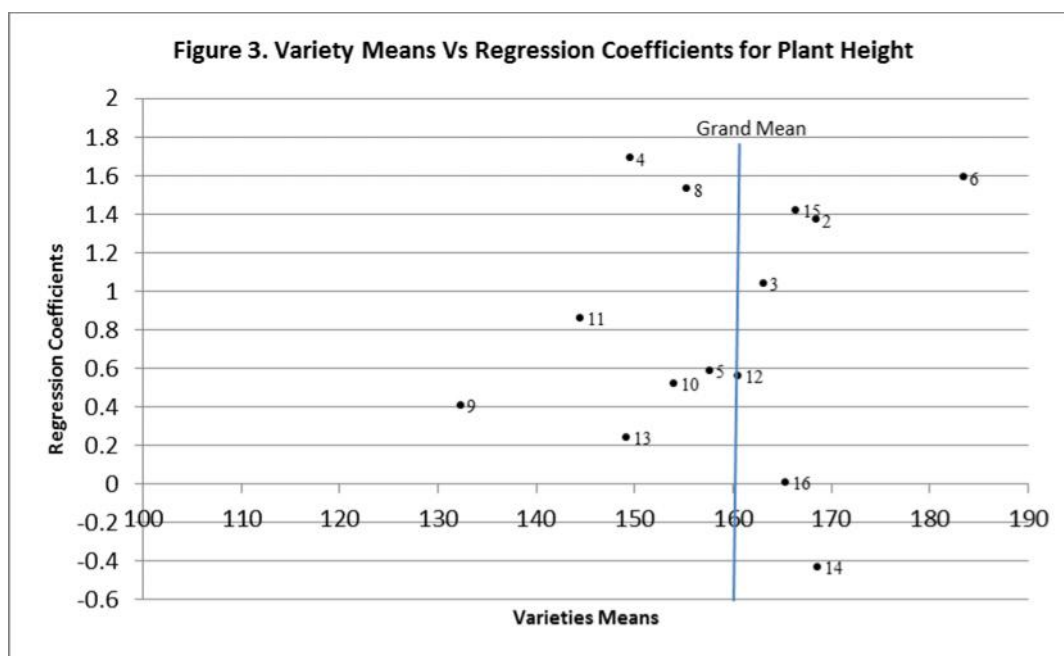
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132 **Plant height**

133 For plant height only genotype MS-92-CP-624 had a value close to unity (Figure 3) and

134 higher mean yield (Table 4), 8 genotypes had a value less than 1 while remaining genotypes

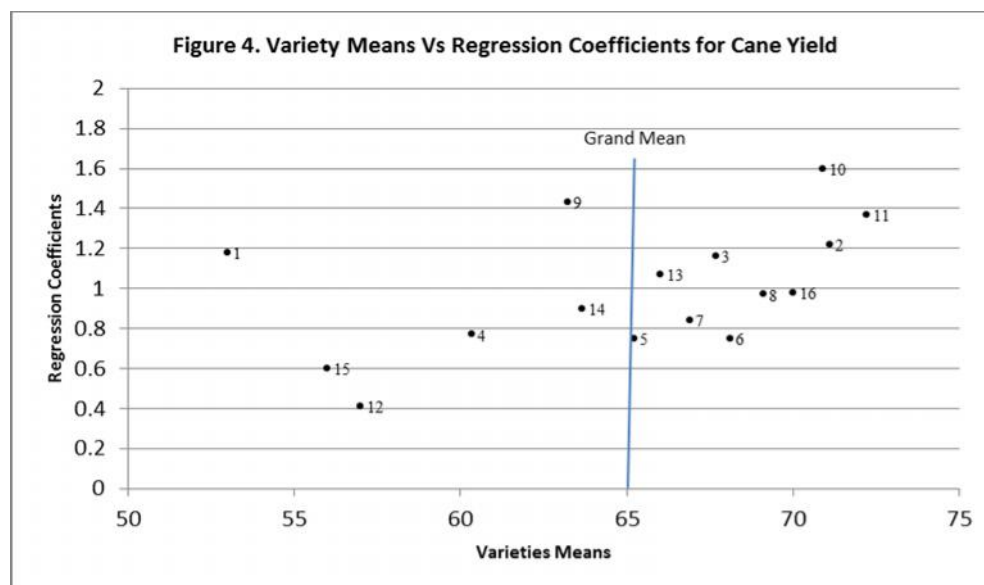
135 exhibited slope value more than 1.



136

137 **Cane yield**

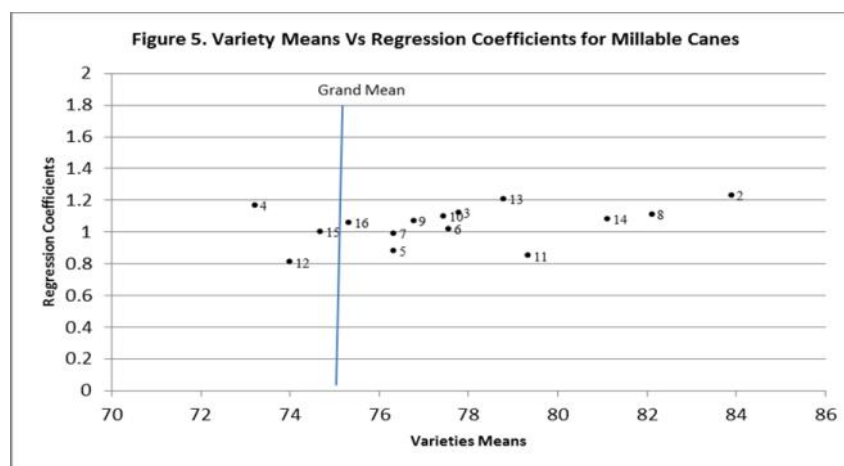
138 For this character, genotypes Mardan 93 and CP 77/400 showed values close to unity and
 139 had higher mean yields (Figure 1). Seven genotypes showed regression values lesser than
 140 1 while rest of the genotypes were having regression coefficient values above 1.



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142 **Millable Canes**

143 For millable canes genotypes MS-91-CP-288 and MS-91-CP-471 showed regression values
 144 close to unity and had higher mean yields. Genotype 15 though showed a unit regression,
 145 had a lower mean yield than the grand mean. Genotypes MS-91-CP-572, MS-91-CP-920,
 146 and MS-91-CP-965 (Figure 5) had regression values less than 1 and hence exhibited above
 147 average stability. The rest of the genotypes showed their adaptability to favorable
 148 environments.



149

150 **Conclusion**

151 The present study indicated that none of the genotypes performed well under all
 152 environments with respect to all characters. However, genotypes Mardan 93 and CP 77/400
 153 showed average stability with higher mean cane yield (t/ha). It means that they can yield
 154 better under all environments. It can be concluded that G x E interaction and stability
 155 analysis/testing of advanced breeding material needs to be an integral part of sugarcane
 156 breeding program so that sugarcane genotypes with superior cane yield and other desirable
 157 attributes could be identified for multiple environments.

158 **Table 4: Means and regression slope for 16 genotypes**

S. No.	Genotype	Germination %		No. of tillers		Plant height		Cane yield		Millable canes	
		Mean	bi*	Mean	bi	Mean	bi	Mean	bi	Mean	bi
1	Malakand 17	40.22	0.85	159.67	0.2	177.22	2.17	53	1.18	47	0.3
2	MS-92-CP-623	44.11	0.89	225.67	1.19	168.44	1.37	71.11	1.22	83.89	1.23
3	MS-92-CP-624	45.78	0.61	198.89	1.08	163.11	1.04	67.67	1.16	77.78	1.12
4	MS-91-CP-611	34.89	0.63	189.22	1.16	149.56	1.69	60.33	0.77	73.22	1.17
5	MS-91-CP-572	38.33	0.93	210.67	1.09	157.67	0.59	65.22	0.75	76.33	0.88
6	MS-91-CP-288	45	0.95	204.56	0.73	183.44	1.59	68.11	0.75	77.56	1.02
7	AEC-86-347	47	1.08	202	0.95	166.33	2.45	66.89	0.84	76.33	0.99
8	Mardan 93	42.11	0.88	258.22	1.31	155.33	1.53	69.11	0.97	82.11	1.11
9	MS-91-CP-471	50	1.32	211	0.91	132.33	0.41	63.22	1.43	76.78	1.07
10	MS-91-CP-623	50.56	0.99	203.33	1.06	154	0.52	70.89	1.6	77.44	1.1
11	MS-91-CP-920	39.44	0.89	177.33	0.54	144.44	0.86	72.22	1.37	79.33	0.85
12	MS-91-CP-965	51.67	1.58	234.56	1.69	160.56	0.56	57	0.41	74	0.81
13	MS-92-CP-1100	51.44	1.36	200.22	1.12	149.22	0.24	66	1.07	78.78	1.21
14	MS-94-CP-90	54.78	1.02	234.67	1.28	168.56	0.43	63.67	0.9	81.11	1.08
15	CPF-236	45.22	0.64	156.89	0.3	166.44	1.42	56	0.6	74.67	1
16	CP 77/400	50.22	1.39	231.33	1.4	165.33	0.01	70	0.98	75.33	1.06
	Grand Mean	45.67		206.14		160.12		65.03		75.73	

159 *Regression Slope

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161

162 **AUTHORS' CONTRIBUTION**

- 163 1. Mohammad Tahir: Designed, and laid out the experiment; compiled the study results,
 164 followed by statistical analyses; wrote the first draft.
 165 2. Dr Hidayatur Rahman: Critically reviewed the first draft.
 166 3. Amjad Ali and Sajjad Anwar: Helped in relevant literature search.
 167 4. Muhammad Khalid: Helped a lot during field work and compilation of the data.

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