

Research Article

Genetical studies of corn crop to exploit heterosis, proportional contribution and gene action at diverse water regimes

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Abstract

The study is based on mating fashion of line \times tester and its relative analysis to exploit the heterosis, proportional contribution of diverse lines, testers, their interaction and gene action in cross combinations of different corn genotypes under different water regimes. Total 12 parents comprised of eight lines and four testers; crossed to produce 32F₁ hybrids. Parents including their hybrids were evaluated in CRD with three water treatments under controlled conditions with three replications in two seasons. Heterosis, percent contribution of lines, testers and line \times tester interaction and gene action were studied for the traits i.e. ear leaf area, number of kernel rows per ear, 100 grain weight, biological yield, vegetative dry matter and anthesis-silking interval. Variable heterosis both in magnitude and direction was observed. Both additive and non-additive gene effects have role in the control of anthesis-silking interval; whereas non-additive gene effects were observed for ear leaf area, number of kernel rows per ear, 100 grain weight, biological yield and vegetative dry matter. However 100 seed weight was mainly affected by dominance effects under this study that helped for the selection of parents to be used for the development of synthetics and hybrids durable for different water regimes.

Keywords: Additive; Hybrid; Line \times tester; Susceptible; Water stress

Introduction

Food security in the world is challenged by increasing food demand and threatened by declining water availability. This water crisis is a severe threat for sustainable agriculture, particularly in Asia, where irrigated agriculture accounts for 90% of total diverted fresh water [1]. Due to rapid increase in

population of developing countries, the demand for food is increasing with every passing day, and this pressure for high food demand will continue for next three coming decades.[1].

There are many constraints in corn production especially in developing countries like Pakistan, including different kinds of

biotic and abiotic stresses. Biotic stresses include pathogens, diseases and bacteria [2] and abiotic stresses include salinity, fluctuating heat pattern, temperature variations, high pH, metal toxicity and water stress having detrimental effects on plant growth and yield [3]. Corn a C4 plant is an efficient user of water and highly responsive to water shortages [4]. Heterosis and combining ability is pre-requisite for developing desirable corn hybrids [5].

Combining ability is a measure of additive gene action [6] while specific combining ability (SCA) is due to genes showing non-additive effects [7]. Genetics use for the development of drought tolerant plants is an important for yield stability on global bases [8]. In conventional breeding methods, genetic variability against drought tolerance can be identified and introduced through different mating techniques to develop cultivars with good agronomic traits [9]. Thus selection of parental lines is essentially based on general and specific combining ability.

Therefore a study was made to investigate different parents and their crosses under different water regimes in Line \times tester analysis, proportional contribution of lines, testers and line \times tester interaction, nature of gene action and direction and magnitude of heterosis to obtain reliable genetic information of grain yield and their related traits in corn breeding program. So this information helps the plant breeder to

construct the breeding strategy under water stress conditions [10].

Materials and methods

Total 44 newly developed genotypes including 12 parents and 32 crosses were sown in the polythene bags (45 \times 30cm) filled with mixture of soil (Soil, Silt and organic matter, in equal amount) having (pH 8.2 and EC.0.4) under three water treatments:

Water treatment1 T1 Eighty percent (80%) of field capacity

Water treatment2 T2 Sixty percent (60%) of field capacity

Water treatment3 T3 Forty percent (40%) of field capacity in wire-house of Department of Plant Breeding and Genetics, University College of Agriculture, Sargodha, Pakistan. Recommended agronomic, cultural and plant protection measures were kept uniform to all the three sets of experimental units except irrigation water. Insecticide was applied to control the insect attack. Two factor factorial triplicate randomized complete designs were followed. Each genotype was sown in three lines per replication per treatment in two seasons during spring and autumn seasons. Three water treatments were applied following [11]. The Average maximum and minimum temperature of 100 days of Season I & II was depicted by (Figure 1 & 2) respectively. The different traits data were recorded at maturity (Table 1).

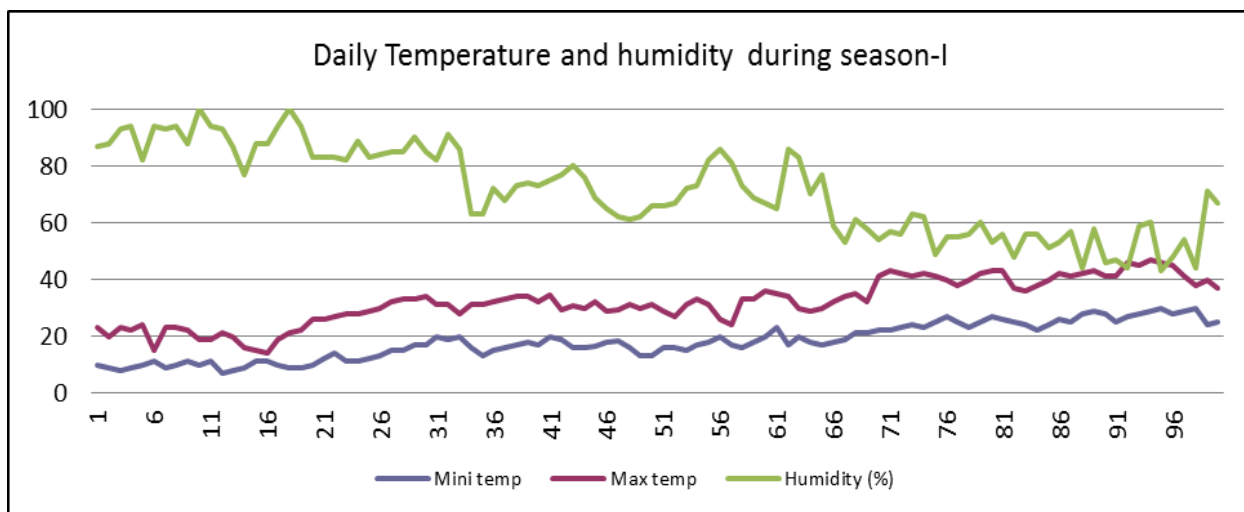


Figure 1. Average maximum and minimum temperature of 100 days of Season 1

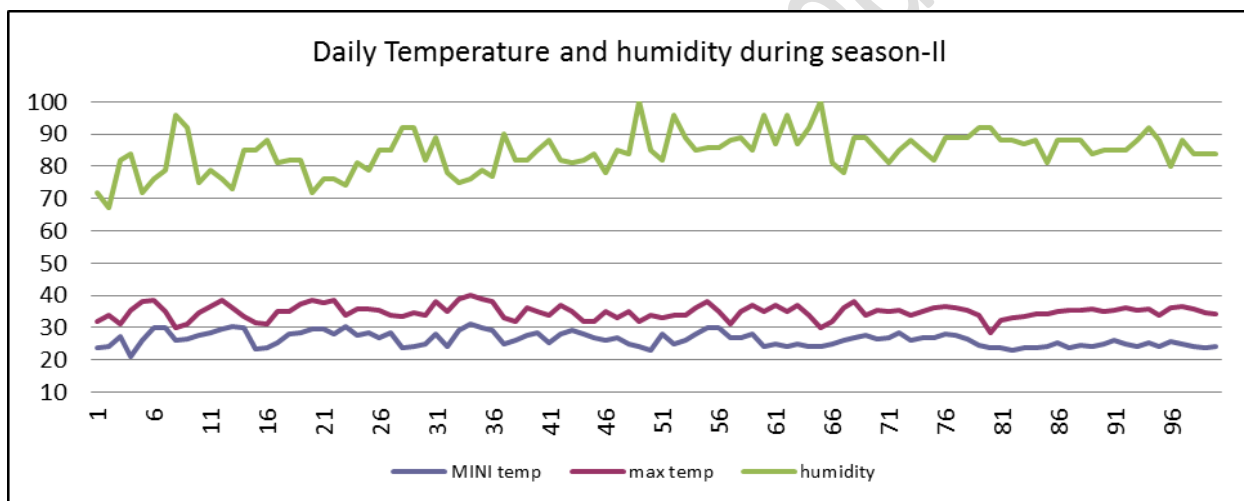


Figure 2. Average maximum and minimum temperature of 100 days of season II

Table 1. The data was recorded on appropriate time for the following traits

1	Ear Leaf Area	Five randomly selected guarded plants were measured in cm ² , using leaf area meter (ΔT -MK2, England) and then average was computed
2	Number of Rows per Ear	Five randomly selected guarded plants of each treatment was counted and then average was calculated.
3	100-grain Weight	Measured in grams with an electronic balance (OHAUS-GT 4000, USA) from the bulk produce of five randomly selected plants in each treatment.
4	Biological Yield	Average weight of five randomly selected plants separately in grams
5	Vegetative Dry Matter	Five randomly selected plants in grams by using electronic balance (OHAUS-GT 4000, USA) and then average was determined
6	Anthesis-Silking Interval	Anthesis was started before the emergence of silk on the cob of the respective plants. Dates of anthesis and silking of all corn accessions were recorded. The difference between dates of anthesis-silking intervals was calculated for all the genotypes separately.

Statistical analysis

The recorded data were statistically analyzed [12] and were also subjected to line \times tester analysis of variance (ANOVA) as outlined by [13].

1. Heterosis

Heterosis over better parent was estimated as followed by [14].

T-test was calculated [15] to compare significance of heterosis.

$$\text{Cov H.S. (lines)} = M_l - M_{lt} / rt ; \text{Cov H.S. (tester)} = M_t - M_{lt} / rl$$

$$\text{Cov H.S. (average)} = [l / r(2lt - l - t)] [\{ (l-1)(M_l) + (t-1)(M_t) \} / (1+t-2) - M_{lt}]$$

Results

Analyses of variance for per cent contribution of lines, testers and lines \times testers interaction and gene action for number of leaves of maize genotypes in 8x4 line \times tester cross in contrasting water regimes under controlled conditions for season-I and II. (Table 2)

In season-I, heterotic manifestations in 32 corn genotypes for ear leaf area, number of kernel rows per area, 100 grain weight, biological yield, vegetative dry matter and anthesis-silking interval were measured under water treatment 1, 2 and 3, (Table 3) heterosis value depicts different in magnitude and direction. Under water treatment 1, Crosses DR-187 \times Pak Afgoeefor ear leaf area, DR-189 \times Sadaaf for number of kernels rows per ear, DR-158 \times Pak Afgoeefor 100 grain weight, DR-198 \times Pak Afgoeefor biological yield, DR-177 \times EV-6098 for vegetative dry matter and DR-198 \times EV-6098for anthesis-silking had maximum positive and significant heterosis while Crosses DR-159 \times Ev-1098 for ear leaf area, DR-185 \times Ev-1098for number of kernels per row, DR-177 \times EV-6098 and DR-177 \times Pak Afgoeefor 100 grain weight, DR-198 \times EV-6098for biological yield, DR-159 \times Ev-6098 and DR-159 \times Sadaaf, DR-198 \times EV-1098 for anthesis-silking had minimum positive and significant

2. Proportional Contribution of lines, Testers and their interaction to total variance

Contribution of lines =

$$\{ss(l) / ss(\text{crosses})\} \times 100 ;$$

Contribution of tester

$$\{ss(t) / ss(\text{crosses})\} \times 100 ;$$

Contribution of (lxt) = $\{ss((lxt) / ss(\text{crosses}))\} \times 100$

3. Genetic Components

heterosis whereas Crosses DR-158 \times Ev-1098 for ear leaf area, DR-194 \times EV-1098for number of kernel rows per ear, DR-194 \times EV-1098 for 100 grain weight, DR-159 \times Sadaaf for biological yield, DR-194 \times Ev-6098for vegetative dry matter and DR-185 \times Pak Afgoeefor anthesis-silking had maximum negative and significant heterosis while Crosses DR-159 \times Sadaaffor ear leaf area, DR-198 \times EV-1098for number of kernel rows per ear, DR-159 \times Ev-1098 for 100 grain weight, DR-158 \times Pak Afgoeefor biological yield, DR-198 \times Pak Afgoeefor vegetative dry matter and DR-158 \times Sadaaf for anthesis-silking had maximum negative and significant heterotic value. Under water treatment 2 Crosses DR-187 \times Pak Afgoeefor ear leaf area, DR-189 \times Sadaaf for number of kernels per row, DR-159 \times Ev-1098 for 100 grain weight, DR-185 \times Pak Afgoeefor biological yield, DR-177 \times Pak Afgoeefor vegetative dry matter and DR-187 \times Ev-6098 for anthesis-silking interval had maximum positive and significant heterosis while Crosses DR-185 \times Ev-6098for ear leaf area, DR-177 \times EV-6098 for number of kernels per row, DR-189 \times Pak Afgoeefor 100 grain weight, DR-198 \times EV-6098 for biological yield, DR-198 \times EV-6098for vegetative dry matter and DR-159 \times Ev-6098 for anthesis-silking interval had minimum positive and significant heterosis

whereas Crosses DR-159 × Ev-1098 for ear leaf area, DR-159 × Sadaaf and DR-159 × Ev-6098 for number of kernels per row, DR-194 × Sadaaf for grain weight, DR-159 × Sadaaf for biological yield, DR-194 × Ev-6098 for vegetative dry matter and DR-194 × Pak Afgooe for anthesis-silking had maximum negative and significant heterosis while crosses DR-158 × Ev-1098 for ear leaf area, DR-187 × Ev-1098 for number of kernels per row, DR-194 × Ev-6098 for 100 grain weight, DR-159 × Ev-1098 for biological yield, DR-187 × Ev-6098 for vegetative dry matter and DR-189 × Ev-1098 and DR-159 × Ev-1098 for anthesis-silking had minimum negative and significant heterosis effects. Under water treatment 3, Crosses DR-187 × Pak Afgooe for ear leaf area, DR-159 × Ev-1098 for 100 grain weight, DR-198 × Pak Afgooe for biological yield, DR-185 × Pak Afgooe for vegetative dry matter and DR-177 × Sadaaf for anthesis-silking interval had maximum positive and significant heterosis while crosses DR-159 × Sadaaf for ear leaf area, DR-189 × Sadaaf for number of kernel rows per ear, DR-159 × Sadaaf for 100 grain weight, DR-194 × Ev-6098 for 100 grain weight, DR-198 × Ev-6098 for biological yield, DR-194 × Ev-6098 for vegetative dry matter and DR-177 × Ev-6098, DR-158 × Ev-6098, DR-189 × Pak Afgooe and DR-159 × Sadaaf for anthesis-silking interval had minimum positive and significant heterosis whereas Crosses DR-159 × Ev-1098 for flag leaf area, DR-159 × Ev-6098 for number of kernels rows per ear, DR-198 × Pak Afgooe for 100 grain weight, DR-159 × Sadaaf biological yield, DR-189 × Pak Afgooe for vegetative dry matter and DR-185 × Sadaaf for anthesis-silking interval had maximum negative and significant heterosis while Crosses DR-158 × Ev-1098 for ear leaf area, DR-177 × Sadaaf for number of kernels rows per ear, DR-177 × Ev-6098 for 100 grain weight, DR-158 × Pak Afgooe for biological yield, DR-194 × Pak Afgooe for

vegetative dry matter and DR-177 × Pak Afgooe, DR-194 × Ev-6098, DR-185 × Pak Afgooe for anthesis-silking interval had minimum negative and significant heterosis. In season-II under water treatment 1, (Table 3) The crosses DR-187 × Pak Afgooe for ear leaf area, DR-158 × Pak Afgooe for 100 grain weight, DR-185 × Ev-6098 for biological yield and DR-177 × Ev-1098, DR-194 × Ev-1098 anthesis-silking interval had maximum positive and significant heterosis while Crosses DR-159 × Ev-6098 for ear leaf area, DR-177 × Ev-6098 for number of kernels per ear, DR-158 × Ev-1098 for 100 grain weight, DR-194 × Ev-6098 for biological yield and DR-177 × Ev-6098, DR-185 × Ev-1098 anthesis-silking interval had minimum positive and significant heterosis whereas Crosses DR-185 × Sadaaf for ear leaf area, DR-159 × Ev-1098 for number of kernel rows per ear, DR-198 × Ev-1098 for 100 grain weight, DR-158 × Ev-1098 for biological yield, DR-185 × Pak Afgooe for vegetative dry matter and DR-159 × Pak Afgooe for anthesis-silking interval had maximum negative and significant heterosis whereas Crosses DR-194 × Sadaaf for ear leaf area, DR-189 × Pak Afgooe, DR-189 × Ev-1098 for number of kernel rows per ear, DR-189 × Ev-1098 for 100 grain weight, DR-185 × Sadaaf for biological yield, DR-187 × Ev-6098 for vegetative dry matter and DR-177 × Sadaaf, DR-198 × Sadaaf for anthesis-silking interval had minimum negative and significant heterosis. Under water treatment 2, Crosses DR-198 × Sadaaf for ear leaf area, DR-159 × Ev-1098 for 100 grain weight, DR-185 × Ev-6098 for biological yield, DR-159 × Ev-6098 for vegetative dry matter and DR-189 × Ev-1098 for anthesis-silking interval had maximum positive and significant heterosis while Crosses DR-159 × Ev-1098, DR-159 × Ev-6098 for ear leaf area, DR-159 × Sadaaf for 100 grain weight, DR-189 × Ev-1098 for biological yield, DR-189 × Ev-6098 for

vegetative dry matter and DR-177 × Pak Afgoe, DR-189 × Sadaaf for anthesis-silking interval had minimum positive and significant heterosis. Whereas Crosses DR-159 × Sadaaf for ear leaf area, DR-159 × Ev-1098 for number of kernel rows per ear, DR-194 × Ev-1098 for 100 grain weight, DR-158 × Ev-1098 for biological yield, DR-187 × Pak Afgoe for vegetative dry matter, DR-159 × Ev-1098 anthesis-silking interval had maximum negative and significant heterosis while Crosses DR-159 × Pak Afgoe for ear leaf area, DR-189 × Ev-1098 for number of kernel rows per ear, DR-198 × Pak Afgoe for 100 grain weight, DR-185 × Sadaaf for biological yield, DR-159 × Ev-1098 for vegetative dry matter and DR-158 × Pak Afgoe, DR-158 × Sadaaf for anthesis-silking interval had minimum negative and significant heterosis. Under water treatment 3, Crosses DR-177 × Pak Afgoe for ear leaf area, DR-159 × Ev-1098 for 100 grain weight, DR-185 × Ev-6098 for biological yield and DR-194 × Ev-1098 for anthesis-silking interval had maximum positive and significant heterosis while Crosses DR-159 × Ev-6098 for ear leaf area, DR-177 × Ev-6098 for number of kernel rows per ear, DR-158 × Ev-1098 for 100 grain weight, DR-194 × Ev-6098 for biological yield and DR-198 × Ev-6098, DR-158 × Sadaaf and DR-189 × Sadaaf for anthesis-silking interval had minimum positive and significant heterosis whereas Crosses DR-159 × Ev-1098 for ear leaf area, DR-158 × Ev-1098 for number of kernel rows per ear, DR-198 × Pak Afgoe for 100 grain weight, DR-158 × Ev-1098 for biological yield and DR-187 × Ev-1098 for anthesis-silking interval had maximum negative and significant heterosis while Crosses DR-158 × Ev-1098 for ear leaf area, DR-185 × Sadaaf for number of kernel rows per ear, DR-177 × Ev-6098 for 100 grain weight, DR-185 × Sadaaf for biological yield and DR-158 × Pak Afgoe for anthesis-

silking interval had minimum negative and significant heterosis.

Under water treatment 1 in season-I and II (Table 4), proportional contribution of lines for the traits ear leaf area, lines for number of kernel rows per ear, Line × tester interaction for 100 grain weight. Lines for biological yield except in season-II where Line × tester interaction for the biological yield, Lines for vegetative dry matter except in season-II where Line × tester interaction for vegetative dry matter and line × tester interaction for anthesis-silking interval and in season-II, lines for anthesis-silking interval contribute maximum in the total phenotype.

Under water treatment 2 and 3 in season-I, proportional contribution of lines for ear leaf area, lines for number of kernel rows per ear, Line × tester interaction for 100 grain weight, Line × tester interaction for biological yield, lines for vegetative dry matter except in water treatment 3, Line × tester interaction for vegetative dry matter and Line × tester interaction for anthesis-silking interval had more contribution under water treatment 2 and 3. In season-II, proportional contribution of lines for ear leaf area, Line × tester interaction for number of kernel rows per ear, Line × tester interaction for 100 grain weight, Line × tester interaction for biological yield, lines for vegetative dry matter and lines for anthesis-silking interval shared maximum in total phenotypic variance.

Dominance variance, additive variance and potence ratio for the traits ear leaf area, 100 grain weight, biological yield, vegetative dry matter and anthesis-silking interval (Table 4) under three water treatments in season-I and II, over dominance was observed for all these traits under these water conditions in both seasons except for anthesis-silking interval in season-II where over dominance was observed under water treatment 1 and 2 but partial dominance observed for anthesis-silking under water treatment 3.

Table 2 . Analyses of variance, for per cent contribution of lines, testers and lines × testers interaction and gene action for number of leaves of maize genotypes in 8x4 line × tester cross in contrasting water regimes under controlled conditions for season-I and II

S.O.V	d.f	Season-I			Season-II		
		Normal	WS-I	WS-II	Normal	WS-I	WS-II
Genotype	43	4.85**	2.42**	2.58**	3.81**	4.94**	2.66**
Parents	11	2.51**	2.16**	3.79**	6.18**	9.35**	5.66**
Crosses	31	5.30**	2.24**	2.02**	2.66**	2.95**	1.49**
Parents Vs Crosses	1	16.55**	11.10**	6.73**	13.14**	18.03**	5.67**
Lines	7	8.68	3.25	2.99**	7.31**	7.33**	3.83**
Testers	3	0.79	0.34	0.13	0.25	1.01	0.74
Line × tester	21	4.82**	2.17**	1.97**	1.46**	1.76**	0.82**
Residual	86	0.65	0.25	0.53	0.48	0.32	0.41
Contribution percentage		Normal	WS-I	WS-II	Normal	WS-I	WS-II
Lines		36.96	32.82	33.47	62.02	56.15	57.88
Testers		1.44	1.48	0.60	0.91	3.32	4.77
Line × tester		61.60	65.70	65.94	37.07	40.54	37.35
Dominance δ		1.39	0.64	0.48	0.33	0.48	0.14
Additive δ		0.01	0.00	0.00	0.02	0.01	0.01
Potence ratio (Degree of dominance)		481.22	1577.35	1480.39	44.89	67.68	34.16

Table 3. Heterosis effects in 32 corn genotypes for ear leaf area, number of kernel rows per ear, 100 grain weight, biological yield, vegetative dry matter and anthesis-silking interval in contrasting water regimes under controlled conditions in season-I and II

Hybrids		Season-I						Season-II					
		E.L.A	R.N	100 g.wt.	B.Y	V.D.M	ASI	E.L.A	R.N	100 g.wt.	B.Y	V.D.M	ASI
DR-187 × Pak Afgoee	T1	27.94	-15.79	-4.92	-18.29	-6.98	-22.73	27.40	-16.67	-4.20	-18.08	-7.84	-11.76
	T2	32.93	-17.39	-12.44	-11.64	-13.73	-13.33	18.8	-17.39	-4.53	-18.36	-7.95	-8.00
	T3	37.98	-22.00	-22.99	-19.91	-12.78	7.69	21.43	-21.74	-23.24	-18.64	-12.24	1.79
DR-187 × Sadaaf	T1	17.04	-14.56	-20.49	-29.67	11.05	-15.22	16.86	-4.35	-18.49	-28.29	9.39	-12.5
	T2	20.99	-4.55	-22.49	-28.46	3.75	-4.26	17.49	-9.09	-7.79	-28.3	4.01	-10.87

	T3	25.38	-7.5	-25.29	-31.51	-1.61	-13.79	9.63	-4.76	-25.70	-28.31	-1.65	-22.22
DR-187 × Ev-1098	T1	13.33	-22.94	-9.84	-17.02	-2.28	-11.11	13.62	-4.17	-6.72	-22.04	-4.75	-26.67
	T2	14.47	-4.35	-13.88	-15.11	-8.79	-8.16	15.25	-4.35	-4.50	-23.92	0.85	-2.38
	T3	15.72	-4.15	-19.54	-19.85	-5.14	-11.54	8.79	-8.34	-14.68	-25.84	-0.55	-38.89
DR-187 × EV-6098	T1	15.78	-20.81	-10.66	-23.15	-0.27	21.62	19.88	-9.52	-8.40	-24.93	-6.33	-26.67
	T2	17.62	-10.00	-18.18	-24.86	-2.62	8.89	19.01	-4.18	-6.13	-24.8	-3.58	-11.32
	T3	19.52	-3.72	-28.74	-24.38	-6.83	-3.85	21.1	-4.39	-28.4	-24.67	-6.92	-33.33
DR-177 × Pak Afgoe	T1	22.62	-20.88	0.00	-20.75	-1.69	-25	22.67	-16.67	0.88	-15.41	-7.32	-11.76
	T2	29.12	-17.39	-13.3	-16.31	-2.42	-18.75	15.68	-15.96	-3.45	-15.87	-4.11	0.00
	T3	33.92	-22.73	-30.68	-25.62	-2.54	-7.69	30.64	-20.3	-26.22	-16.33	-1.96	-8.93
DR-177 × Sadaaf	T1	4.62	-25.00	5.22	6.38	10.52	-21.74	4.25	-21.74	5.31	-26.46	52.92	-6.25
	T2	7.14	-13.64	-10.10	9.28	-11.47	-18.75	8.26	-16.68	-3.28	-28.02	29.58	2.17
	T3	7.19	-20.00	-30.11	5.23	-1.22	48.28	5.33	-18.26	-26.68	-29.59	2.82	1.85
DR-177 × Ev-1098	T1	12.32	-19.63	-9.57	-20.23	6.11	-13.33	11.35	-20.83	-7.96	8.96	-6.78	18.18
	T2	11.26	-21.74	-17.49	-14.51	-0.62	-10.2	12.1	-15.48	-6.33	9.12	-2.11	9.52
	T3	9.54	-12.53	-27.84	-23.09	-1.31	19.23	14.31	-15.93	-27.15	-13.68	-1.21	16.67
DR-177 × EV-6098	T1	9.18	-10.44	13.14	19.3	-2.43	-21.95	7.53	0.00	14.78	-24.27	42.96	7.14
	T2	8.84	0.00	-8.87	6.25	10.45	-31.25	14.39	0.55	-5.37	-24.75	52.75	-11.32
	T3	8.28	1.83	-15.91	17.36	55.43	0.00	13.42	0.00	-10.9	-25.25	56.73	1.85
DR-198 × Pak Afgoe	T1	17.12	-8.38	19.81	29.52	-6.18	-38.64	16.99	-12.5	20.00	-22.84	52.31	-11.76
	T2	19.45	-13.04	-2.76	-1.15	1.09	-31.25	20.16	-11.11	-1.95	-23.73	29.37	-8.00
	T3	22.01	-19.73	-34.67	28.1	-5.36	21.43	24.1	-16.67	-37.92	-24.63	-3.49	-10.53
DR-198 × Sadaaf	T1	21.72	-7.13	-2.83	-11.00	5.03	-41.3	21.05	-13.04	-2.86	-20.9	11.05	-6.25
	T2	23.96	-13.64	1.66	-17.25	5.58	-18.75	22.63	-13.14	0.97	-21.09	10.81	-12.00
	T3	27.13	-19.18	8.00	-14.98	-2.18	44.83	25.99	-11.14	7.05	-21.28	10.57	-10.53
DR-198 × EV-1098	T1	18.91	-8.33	1.77	20.26	-1.06	0.00	18.42	-26.00	2.70	20.91	-9.92	-31.82
	T2	19.89	-21.74	16.85	17.19	2.22	-2.04	20.97	-20.74	4.52	20.47	-1.51	-16
	T3	19.18	-24.18	28.67	21.47	-0.21	7.14	17.58	-21.23	27.15	20.03	-0.39	-21.05
DR-198 × EV-6098	T1	14.5	-14.63	-7.63	5.84	10.41	7.14	13.35	4.35	-5.22	17.96	-9.13	-4.55
	T2	16.66	4.55	8.29	4.96	10.89	6.25	16.95	4.55	2.14	17.83	-3.01	-13.21
	T3	19.14	1.68	16.00	5.90	-1.73	14.29	19.23	4.76	8.34	17.69	-2.76	0.00
DR-194 × Pak Afgoe	T1	9.55	-29.61	-4.51	7.77	-2.65	-38.64	8.53	-16.67	-4.65	15.06	-8.32	-9.8
	T2	13.55	-17.39	-9.78	3.67	4.48	-40.00	5.73	-15.52	-3.22	15.79	-6.19	-2.00
	T3	18.84	-20.48	-17.39	7.76	-1.73	7.69	19.17	-20.74	-18.77	16.54	-3.83	-14.29
	T1	2.48	-28.56	-24.81	-30.8	8.17	-41.3	-0.17	-13.04	-21.71	-18.71	-10.28	-6.25

DR-194 × Sadaaf	T2	5.54	-13.64	-24.44	-30.51	9.68	-23.4	8.27	-12.59	-8.33	-18.41	-7.43	2.17
	T3	10.53	-12.53	-23.91	-33.64	-4.45	-10.34	5.24	-12.19	-19.13	-18.1	-4.26	-5.56
DR-194 × EV-1098	T1	11.68	-30.22	-26.32	-34.38	10.81	2.22	10.21	-20.83	-24.03	9.09	-36.32	18.18
	T2	10.71	-13.04	-22.67	-33.56	34.71	-2.04	16.23	-21.26	-8.64	9.19	-20.24	9.52
	T3	9.13	-10.00	-17.39	-35.26	-1.15	-11.54	14.94	-21.98	-14.4	9.29	-3.56	21.43
DR-194 × Ev-6098	T1	20.76	-18.50	-10.53	-28.93	5.72	-14.29	18.55	-21.74	-9.30	7.37	-33.13	0.00
	T2	22.41	-13.64	-6.22	-26.17	22.53	-6.67	21.83	-20.7	-2.31	7.98	-18.27	-5.66
	T3	25.15	-18.33	0.00	-28.43	-0.15	-7.69	22.89	-16.69	-3.46	8.61	-3.24	-11.11
DR-158 × Pak Afgooe	T1	12.16	-7.13	20.39	-9.04	-2.87	-25.00	13.28	-25.00	20.39	11.76	-20.84	-17.65
	T2	16.9	-21.74	29.61	-6.84	10.11	-6.67	9.46	-24.65	9.99	12.25	-9.58	-8.00
	T3	21.16	-24.27	42.11	-7.6	1.14	11.54	23.77	-25.61	43.13	12.75	-1.67	-5.26
DR-158 × Sadaaf	T1	1.81	-11.06	2.91	-49.89	67.54	-2.17	1.56	-20.83	2.91	-27.77	16.67	-4.17
	T2	3.27	-18.18	5.59	-33.73	17.31	-4.26	0.40	-19.8	2.66	-27.33	6.04	-6.38
	T3	3.02	-17.53	9.21	-50.84	-4.3	-10.34	6.69	-20.7	9.37	-26.88	-5.11	0.00
DR-158 × Ev-1098	T1	-2.31	-6.94	7.08	-42.09	46.54	-40.00	-2.39	-16.67	9.46	-43.45	38.6	-37.5
	T2	-3.43	-17.39	12.01	-43.04	22.99	-38.78	1.54	-20.54	3.52	-43.02	22.79	-6.38
	T3	-5.26	-14.18	4.61	-44.09	-4.26	-19.23	-1.97	-40.41	5.77	-42.59	-4.8	-36.84
DR-158 × Ev-6098	T1	8.48	-17.38	-20.34	-16.5	-0.56	-23.08	7.52	-4.17	-16.96	-29.12	8.78	-31.25
	T2	8.34	6.07	-9.50	-22.81	-0.63	-17.78	10.69	-4.35	-3.27	-29.38	4.66	-11.32
	T3	7.74	-4.15	-10.53	-18.28	-4.94	0.00	12.59	-4.05	-9.12	-29.63	-5.37	-26.32
DR-185 × Pak Afgooe	T1	6.82	-22.13	6.48	23.18	-4.71	-42.55	6.25	-9.34	10.00	-26.56	60.76	-23.53
	T2	10.37	-8.70	-1.22	26.14	-22.53	-28.26	5.37	-8.22	-0.09	-26.41	31.11	-16.00
	T3	16.85	-16.68	-16.07	20.29	-1.79	-7.69	14.01	-12.33	-13.31	-26.26	-3.49	-19.64
DR-185 × Sadaaf	T1	1.00	-21.00	-16.67	-12.85	-4.93	-42.55	-1.6	-8.70	-14.29	-7.95	-13.69	-43.75
	T2	3.42	-9.09	2.44	-14.85	3.67	-29.79	3.39	-9.09	-6.26	-7.58	-10.00	-10.87
	T3	7.79	-10.00	39.29	-15.88	-2.29	-31.03	0.22	-7.95	-4.86	-7.21	-5.99	-27.78
DR-185 × Ev-1098	T1	7.46	2.06	-6.19	-10.99	-4.32	-4.26	8.25	-20.83	-4.50	-28.79	13.5	7.14
	T2	6.97	-17.39	-9.76	-11.29	-12.81	-8.16	6.32	-21.02	-2.77	-28.4	14.48	16.67
	T3	6.19	-11.70	-25.00	-12.6	-3.47	-19.23	12.84	-16.68	-29.85	-28.01	10.66	-5.56
DR-185 × Ev-6098	T1	2.07	-4.17	1.69	20.72	-0.45	-4.26	-0.36	-4.76	4.35	23.69	-7.33	-7.14
	T2	2.05	-5.00	-0.91	23.22	-1.30	-2.17	6.34	-7.95	-0.44	23.91	-2.53	-11.32
	T3	2.04	-8.81	-21.43	21.61	-0.39	-19.23	4.34	-3.33	-22.03	24.14	-1.64	-16.67
DR-189 × Pak Afgooe	T1	7.46	-4.88	-7.96	-19.21	-4.07	-11.36	5.99	-8.33	-6.76	-11.59	-9.48	-11.76
	T2	11.11	-8.7	7.36	-17.2	3.20	-2.17	8.18	-8.70	2.68	-11.15	-5.74	4.00

	T3	17.36	-12.14	42.00	-17.43	0.33	0.00	14.24	-11.61	46.46	-10.71	-2.91	-8.93
DR-189 × Sadaaf	T1	8.67	5.78	-8.85	-21.32	1.98	-2.17	6.03	4.35	-5.86	-19.60	-1.53	-8.33
	T2	11.99	4.55	-2.45	-16.4	-3.92	-4.26	10.47	4.55	-0.93	-18.99	-3.98	0.00
	T3	17.94	0.00	12.00	-21.12	-2.30	-20.69	11.44	4.76	15.32	-18.36	-6.55	0.00
DR-189 × Ev-1098	T1	11.00	-8.91	-5.31	-24.91	48.76	6.67	10.15	-8.33	-3.60	-15.59	25.89	0.00
	T2	9.62	-8.7	0.61	-20.99	19.63	-6.12	10.88	-7.98	-0.34	4.96	10.87	18.18
	T3	7.37	-5.00	14.00	-24.63	-10.7	-30.77	13.45	-8.59	14.86	-16.5	-11.74	-6.67
DR-189 × Ev-6098	T1	7.71	-7.26	-6.78	-41.51	38.81	-4.55	6.63	-4.76	-4.35	17.91	-35.17	14.29
	T2	16.24	-5.00	29.91	-41.12	68.34	-4.35	7.37	-2.78	8.93	18.09	-15.18	-11.32
	T3	30.39	-5.56	110.00	-42.67	1.01	-3.85	18.54	-29.26	112.39	18.29	-1.42	-1.85
DR-159 × Pak Afgoee	T1	0.15	-23.63	-4.24	-47.59	42.78	4.55	-0.66	-23.08	-4.22	-29.00	4.21	-35.29
	T2	3.79	-20.00	15.19	-44.78	30.04	-6.67	-0.24	-23.56	3.63	-28.15	-4.24	-14.00
	T3	10.44	-15.95	64.29	-49.48	-3.61	-25.00	9.24	-15.72	63.25	-27.27	-7.93	-35.00
DR-159 × Sadaaf	T1	-1.25	-19	-9.32	-61.27	174.05	0.00	-0.99	-29.74	-9.28	-18.19	28.98	-20.83
	T2	0.80	-28.00	25.32	-58.8	126.68	2.13	-0.57	-31.08	6.93	-18.19	-0.82	2.17
	T3	4.50	-26.54	97.83	-65.54	-3.11	0.00	3.14	-25.37	95.47	-18.36	-4.92	-18.33
DR-159 × Ev-1098	T1	1.25	1.45	-3.39	-16.51	-10.04	2.22	0.50	-30.77	-5.06	14.84	-37.21	-13.04
	T2	-82.67	-24.00	40.25	-16.62	13.98	-6.12	2.04	-31.34	10.90	14.87	-16.17	6.82
	T3	-26.06	-27.29	136.96	-17.82	-0.69	-10.71	-6.50	-29.00	141.33	14.9	-2.23	-25.00
DR-159 × Ev-6098	T1	3.10	-4.59	-0.85	-10.9	-11.54	2.78	2.02	-26.92	-1.69	-20.52	-4.72	-21.74
	T2	-79.81	-28.00	24.47	-5.8	-16.87	0.00	2.04	-27.34	8.20	-19.92	-0.16	-16.98
	T3	1.15	-31.83	87.37	-10.73	-2.19	-7.14	1.36	-26.09	107.62	-19.3	-5.24	-25.00

T1: watertreatment 1, T2: water treatment2, T3: water treatment 3, E.L.A: ear leaf area, R.N: number of kernel rows per ear, 100 g.wt:100 grain weight, B.Y: biological yield, V.D.M: vegetative dry matter and ASI: anthesis-silking interval

Table 4. Per cent contribution of lines, testers and line × tester interaction and gene action for ear leaf area, number of kernel rows per ear, 100 grain weight, biological yield, vegetative dry matter and anthesis-silking interval of corn genotypes under controlled conditions for season-I and II

Contribution percentage		Season-I						Season-II					
		E.L.A	R.N.	100 G.wt.	B.Y.	V.D.M	ASI	E.L.A	R.N.	100 G.wt.	B.Y.	V.D.M	ASI
Lines	T1	62.75	34.63	13.45	43.97	54.1	19.35	61.76	32.03	12.62	25.98	31.22	48.1
	T2	63.2	46.14	39.24	41.26	52.43	16.56	54.96	35.79	41.72	26.58	59.55	44.82
	T3	57.49	34.72	40.66	42.07	38.3	4.62	52.74	32.47	46.53	27.21	51.9	67.66

Testers	T1	10.46	2.81	28.54	13.61	8.83	24.21	12.34	2.66	28.45	9.28	4.42	18.33
	T2	11	9.16	5.01	10.59	11.86	0.97	11.72	1.92	6.94	9.11	10.43	26.14
	T3	11.7	0.66	3.5	14.45	9.18	0.71	6.18	3.71	4.24	8.93	11.77	9.26
Line × tester	T1	26.79	62.55	58.01	42.42	37.07	56.44	25.9	65.31	58.93	64.74	64.36	33.57
	T2	25.81	44.7	55.75	48.15	35.72	82.46	33.32	62.3	51.34	64.31	30.02	29.03
	T3	30.81	64.62	55.84	43.49	52.52	94.67	41.08	63.82	49.23	63.86	36.33	23.08
Dominance δ	T1	701.4 3	1.34	9.35	9095.26	23.67	0.16	702.75	1.07	8.43	7466.7 4	76.16	0.3
	T2	571.6 2	1.22	8.74	7243.1	16.72	0.99	461.9	0.82	8.79	7353.4 2	22.71	-0.07
	T3	642.8 3	0.5	26.84	8894.32	38.07	3.07	574.36	0.74	24.26	7248.8 9	7.41	0.02
Additive δ	T1	19.59	0.02	0.03	98.04	0.42	0.01	21.76	0.01	0.02	6.24	0.08	0.06
	T2	17.1	0.01	0.04	54.03	0.31	-0.03	9.71	0.02	0.06	7.08	0.55	0.02
	T3	14.71	0.01	0.11	89.52	0.22	-0.16	9.06	0.01	0.18	7.95	0.14	0.09
Potence ratio (Degree of dominance)	T1	35.81	588.13	318.33	92.77	56.42	28.19	32.29	1184.45	356.13	1195.8 2	940.95	5.4
	T2	33.44	160.37	232.44	134.07	53.79	-38.23	47.55	467.48	158.44	1039.2 2	41.03	-4.44
	T3	43.71	795.91	243.08	99.36	169.7 1	-19.07	63.37	711.15	137.64	911.41	53.56	0.23

T1: watertreatment 1, T2: water treatment2, T3: water treatment 3, Dominance δ : Dominance Variance, Additive δ : Additive variance, E.L.A: ear leaf area, R.N: number of kernel rows per ear, 100 G.wt:100 grain weight, B.Y: biological yield, V.D.M: vegetative dry matter and ASI: anthesis-silking interval

Discussion

In cereals, photosynthesis contribution of flag leaf varies from 30 to 50% of the assimilates for grain yield [16] and initiation of grain filling coincides with the onset of senescence, therefore photosynthesis in the flag leaf and rate of senescence are most important factors for determining grain yield [17]. It is generally accepted that genotypes that are able to sustain photosynthesis in the flag leaf for a longer time tend to yield more. The reduction in yield was more severe when the stress occurred suddenly rather than gradually and at early stages of grain filling rather than at later stages [18]. With the increase in water stress, flag leaf area reduced and this is the first attempt of any plant to save moisture loss from its body. In the present study, different corn genotypes were studied to know the effect of flag leaf on grain yield. A significant variability for leaf area under low water stress was seen while the biomass accumulation and drought avoidance results in reduction of water transpiration because assimilation of photosynthesis is totally depend on leaf and stomatal size which increase the water use [19].

Water stress at flowering reduces the number of kernels and kernel rows per ear but at the grain filling it reduces the kernel size [20]. With the increase of water stress in corn, results in reduction of grain yield. The similar results were also reported by independent work of [21].

Biological yield and vegetative dry matter are the major component of yield and are directly relates to the amount of water use [19] these are the products of photosynthesis. Underwater shortage, Photosynthesis can decrease and results in poor production of biomass. However biomass production can be enhanced by increasing the soil moisture uptake, results decrease in soil moisture through evaporation [19]. Similar results are concluded by earlier research works of [22]

who reported that corn grain yield, were under the control of non-additive (SCA effect) type of gene action.

Flowering is the critical stage of plant that has greater impact on yield production, very sensitive to low water stress in corn [23]. Silk is the female part of the plant played very important role in pollination and finally results in fertilization. With the increase in water stress at the reproductive stage slow down the silk growth rate, decreases the time of opening the silk to pollinating anther in corn. Slow growth rate of silks increases the anthesis-silking interval, decreases the chancing of pollination and seed setting. For successful seed setting, time of pollen shading must be coinciding with the silk exertion [24]. The time of anthesis is largely independent of water stress and alleles for leaf elongation rate confirmed a low anthesis-silking interval [25]. This open pollinated variety could be regarded as the most desirable parent for this trait. The similar report is made by [26] who independently reported that GCA effects were significant for anthesis-silking interval and additive gene action played a significant role in the inheritance of the character.

For the evaluation of developed hybrids, evaluation and selection of parents was one of the main objects of this study. Combining abilities and gene effects play a significant role in plant breeding programs [27]. General and specific combining ability effects give important genetic information in parent selection on the bases of their hybrids performance [28]. High GCA value including positive and negative indicate that the parental mean is superior or inferior to the general mean and gave information about the concentration of dominant gene with additive effects whereas SCA gives non-additive interactions. The selection of the parents was made on the combination of SCA, GCA and hybrid mean. The frequency of favorable genes increases with best combination was

hybrid mean, favorable SCA and high GCA effects [29] and populations with high frequency of favorable alleles are good source for plant selection. Heterosis is the accumulation of dominant or partially dominant alleles help the Breeders to develop a product with hidden genetic potential which is expressed in the form of heterotic effects [30] and can be exploited for good yield production [31].

Both additive and non-additive gene effects play role in the control of anthesis-silking interval; whereas non-additive gene effects were observed for ear leaf area, number of kernel rows per ear, 100 grain weight, biological yield and vegetative dry matter [32] reported that prominent role of additive gene action in controlling number of kernel rows. However, 100 seed weight mainly affected by dominance effects. Additive and non-additive gene effects were reported by [32, 33]. It was evident that dominant gene has the most important effect on traits such as, kernel per rows, 100 kernel weights and kernel rows per ear controlled by additive genes.(Table 4)

In present study, positive highest and significant heterosis revealed by hybrid DR-158 × Pak Afgoe and DR-198 × Pak Afgoe for 100 grain weight under water treatment 1 in season-I and II. Under water treatment 2 and 3, hybrids DR-158 × Pak Afgoe and DR-159 × Ev-1098 had positive significant heterosis for 100 grain weight in both seasons. The classification of corn germplasm in to heterotic group may help for exploitation of heterosis in corn [34].

Conclusion

The present study showed that positive highest significant heterosis revealed by crosses DR-158 × Pak Afgoe and DR-198 × Pak Afgoe for 100 grain weight under water treatment 1 in season-I and II and under water treatment 2 and 3 crosses DR-158 × Pak Afgoe and DR-159 × Ev-1098 had positive significant heterosis for 100 grain weight in

season-I and II and these are the hybrids with outstanding heterotic values.

Authors' contributions

Conceived and designed the experiments: IR Noorka & JS (Pat) Heslop-Harrison, Performed the experiments: T Ullah, Analyzed the data: T Ullah & FS Awan, Contributed materials/ analysis/ tools: WN Mhret, Wrote the paper: T Ullah & IR Noorka.

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