



# Effect of integrated agronomic practices on hybrids and pure-line varieties under different environmental conditions in pigeonpea [*Cajanus cajan* (L.) Millspaugh]

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## ABSTRACT

The field experiment was conducted at two locations during *kharif* 2011 and 2012 at Vasandrao Naik Marathwada Krishi Vidyapeeth (VNMKV), Parbhani, Maharashtra and International Crops Research Institute for the Semi-Arid Tropics (ICRISAT), Patancheru, Andhra Pradesh to study the effect of various integrated agronomic practices on hybrids and pure-line varieties of pigeonpea. The genotypes used were three medium maturity group hybrids (ICPH 2671, ICPH 2740 and ICPH 3762) and three pure-line varieties (BDN 711, BSMR 736 and Asha) laid out in split-plot design with three replications in Vertisols. The experimental material was planted in three treatments *viz.*, T<sub>1</sub> [75 x 30 cm + 100 kg DAP (basal) + no protective irrigation]; T<sub>2</sub> [75 x 60 cm + 50 N:100 P<sub>2</sub>O<sub>5</sub> kg ha<sup>-1</sup> (split application: 50% as basal and 50% at 60 days) + two protective irrigations (during mid-flowering and mid-pod development stage)]; and T<sub>3</sub> [150 x 30 cm + 50 N:100 P<sub>2</sub>O<sub>5</sub> kg ha<sup>-1</sup> (split application: 50% as basal and 50% at 60 days) + two protective irrigations (mid-flowering and mid-pod development stage)]. The results indicated that significantly higher seed yield, biomass and harvest index was recorded in T<sub>2</sub> at 3923.9 kg ha<sup>-1</sup>, 8816.6 kg ha<sup>-1</sup> and 30.8%, respectively as compared to T<sub>3</sub> (3262.5 kg ha<sup>-1</sup>, 7670.1 kg ha<sup>-1</sup> and 29.8%, respectively) and T<sub>1</sub> (2932.3 kg ha<sup>-1</sup>, 7633.3 kg ha<sup>-1</sup> and 27.8%, respectively). Genotype × environment interaction indicated that over the two locations in two years, hybrids recorded higher seed yield as compared to pure-line varieties. However, G<sub>3</sub>T<sub>2</sub> [ICPH 3762 + (75 x 60 cm + 50 kg N:100 P<sub>2</sub>O<sub>5</sub> kg ha<sup>-1</sup> (split application at 50% as basal and 50% at 60 days after sowing) + two irrigations (during mid-flowering and mid-pod development stage))] recorded the highest seed yield (4153.6 kg ha<sup>-1</sup>). Number of pods plant<sup>-1</sup> and biomass weight (kg ha<sup>-1</sup>) showed significant variation in the interaction of environment × genotype × treatment (EGT). E<sub>1</sub>G<sub>3</sub>T<sub>2</sub> recorded significantly higher number of pods plant<sup>-1</sup> (1181.9) followed by E<sub>1</sub>G<sub>3</sub>T<sub>2</sub> (1026.1) which significantly converted into higher seed yield plant<sup>-1</sup> (237.1 g & 240.1 g respectively) while E<sub>2</sub>G<sub>2</sub>T<sub>2</sub> exhibited significantly greater biomass (11977.9 kg ha<sup>-1</sup>). Among all the genotypes tested, Hybrid ICPH 3762 recorded highest number of pods plant<sup>-1</sup> in T<sub>3</sub> in almost all environments.

**Key words :** Agronomic practices, Environment, Hybrid, Pigeonpea varieties, Yield attributes.

## INTRODUCTION

Pigeonpea (*Cajanus cajan* (L.) Millsp.) is an important food legume crop in Indian sustainable agriculture. In India, around 4.04 million hectare of pigeonpea is cultivated with production of 2.65 million tonnes having an average yield of 656 kg ha<sup>-1</sup> (IIPR, 2013). Maharashtra accounts for over 29.92% (1.21 million hectare) area of pigeonpea with production of 0.85 million tonnes and productivity of 704 kg ha<sup>-1</sup> (DAC, 2012). While, in Andhra Pradesh pigeonpea covered 0.48 million hectare (11.93%) with low productivity of 307 kg ha<sup>-1</sup> (DAC, 2012). However, it was observed that in India, pigeonpea area showed positive growth rate but productivity remains unchanged (Sharma *et al.*, 2013).

The World's first commercial hybrid ICPH 2671 (Saxena *et al.*, 2013) and ICPH 2740 has been released for commercial cultivation. In 30 multi-location trials on farmers field, these two promising hybrids exhibited >30-35% yield advantage over high yielding varieties. In earlier studies, it was reported that yield and yield attributes of pigeonpea were significantly increased in wider spacing (Tuppad *et al.*, 2012, Goud *et al.*, 2012) and irrigation (Mula *et al.*, 2013). Meena *et al.* (2013) reported that the low yield of pigeonpea is mainly attributed to inadequate and imbalanced nutrient application particularly with respect to nitrogen and phosphorus. Selection of proper plant spacing (optimum plant density), adequate nutrient supply and irrigation are the most important factors to increase the productivity of pigeonpea (Ali and Kumar, 2000; Sekhon *et al.*, 1996).

In this context, enhancing production to meet the

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requirement of growing populace, the need to introduce pigeonpea hybrids with appropriate agronomic practices is necessary. The experiment was conducted to understand the effect of integrated agronomic practices in response to wider spacing, fertilizer and irrigation treatments in determining the yield and yield potentials of hybrids and varieties.

## MATERIALS AND METHODS

The experiment was conducted at VNMKV, Parbhani and the International Crops Research Institute for the Semi-Arid Tropics (ICRISAT), Patancheru, A.P. during cropping season 2011-12 and 2012-13 in Vertisols by using six medium duration genotypes of pigeonpea which includes three hybrids [ICPH 2671 (G<sub>1</sub>), ICPH 2740 (G<sub>2</sub>), and ICPH 3762 (G<sub>3</sub>)] and three varieties [BDN 711 (G<sub>4</sub>), BSMR 736 (G<sub>5</sub>) and Asha (G<sub>6</sub>)]. Among varieties, BSMR 736 and Asha were used as standard check. The experimental materials have three treatments in combination with spacing, fertilizer and irrigation which are T<sub>1</sub> control [75 x 30 cm + 100 kg ha<sup>-1</sup> DAP (basal) + No irrigation]; T<sub>2</sub> [75 x 60 cm + 50 kg N:100 P<sub>2</sub>O<sub>5</sub> kg ha<sup>-1</sup> (split application at 50% as basal and 50% at 60 days after sowing) + two irrigations (during mid-flowering and mid-pod development stage)]; and T<sub>3</sub> [150 x 30 cm + 50 kg N:100 P<sub>2</sub>O<sub>5</sub> kg ha<sup>-1</sup> (split application at 50% as basal and 50% at 60 days after sowing) + two irrigations (during mid-flowering and mid-pod development stage)]. The different treatments were laid out in split plot design with three replications. Recommended package of cultural management practices and plant protection measures were adopted to raise a healthy crop.

The data for yield and yield contributing characters were recorded on five competitive plants in each treatment and replication. The data were subjected to statistical analysis to understand the effect of treatment combination on yield and yield contributing characters of pigeonpea hybrids and varieties.

**Details of statistical analysis :** The experiment was laid out in Split Plot design with three treatments of a. treatments (T<sub>1</sub>, T<sub>2</sub>, and T<sub>3</sub>) as whole plot units; b. six genotypes (G<sub>1</sub>, G<sub>2</sub>, G<sub>3</sub>, G<sub>4</sub>, G<sub>5</sub>, G<sub>6</sub>) as subplot units and; c. across four environments (E<sub>1</sub> - 2011 at Parbhani, E<sub>2</sub> - 2012 at Parbhani, E<sub>3</sub> - 2011 at Patancheru, E<sub>4</sub> - 2012 at Patancheru).

For each morphological trait, data were analyzed by using SAS MIXED procedure (SAS Inst. 2002-2008, SAS V 9. 3).

## RESULTS AND DISCUSSION

**Effect of environments and genotypes for various yield and yield attributes for two seasons :** The variation due to environment was found significant. Among two environments, the pigeonpea genotypes tested in Parbhani showed superiority for almost all the characters as compared in Patancheru except plant height at maturity and harvest index. Among four seasons, the genotypes performed well in E<sub>1</sub> (2011 at Parbhani) for yield and yield contributing characters. The pooled mean over two years & across the environments recorded for yield and harvest index was 3372.9 kg ha<sup>-1</sup> & 29.6%, respectively (**Table 2**).

Hybrids required more number of days to 50% flowering (119.1 days) and maturity (173.6 days) as compared to varieties (113.3 and 162.4 days, respectively) (**Table 3**). Likewise, hybrids

**Table 1.** Analysis of variance (F values) for various yield and its associated traits of pigeonpea over two years and across the environments

Source	Df	Days to 50% flowering	Days to 80% maturity	Plant height at maturity (cm)	No. of primary branches plant <sup>-1</sup>	No. of secondary branches plant <sup>-1</sup>	No. of pods plant <sup>-1</sup>
Environment	3	1007.05**	50.49**	43.13**	59.10**	20.49**	9.64**
Genotype	5	1715.17**	2042.39*	130.42**	23.88**	36.42**	13.19**
Treatment	2	2.77	12.70**	0.03	4.50*	9.86**	64.77**
Environment × Treatment	6	1.48	11.21**	1.30	2.15	0.55	3.16*
Genotype × Treatment	10	1.77	1.98*	1.05	1.45	1.32	2.80**
Genotype × Environment	15	40.94**	15.33**	4.99**	1.96*	3.25**	2.79**
Environment × Genotype × Treatment	30	0.96	0.75	0.67	0.84	0.96	1.95**

Source	Df	Seeds pod <sup>-1</sup>	100-seed weight (g)	Seed yield plant <sup>-1</sup> (g)	Seed yield ha <sup>-1</sup> (g)	Biomass ha <sup>-1</sup> (kg)	Harvest index (%)
Environment	3	5.32*	45.51**	10.34**	34.75*	53.53**	10.51**
Genotype	5	32.21**	148.31**	6.25**	11.75**	47.53**	9.53**
Treatment	2	0.78	0.49	56.53**	75.82**	28.96**	17.69**
Environment × Treatment	6	1.97	3.08	2.39	13.67**	8.72**	2.65
Genotype × Treatment	10	0.61	0.80	0.35	0.98	0.55	1.51
Genotype × Environment	15	1.74*	23.00**	1.48	2.11**	6.87**	3.72**
Environment × Genotype × Treatment	30	1.16	1.52	0.96	0.95	1.73*	1.33

**Note :** \* Significant at 5% and \*\* Significant at 1% level.

**Table 2.** *Per se* performance for various yield and yield contributing characters of pigeonpea in each environment

Characters	Environments						Pooled mean
	E <sub>1</sub>	E <sub>2</sub>	Mean	E <sub>3</sub>	E <sub>4</sub>	Mean	
Days to 50% flowering	120.9a	114.7c	117.8	119.1b	110.2d	114.6	116.2
Days to 80 % maturity	170.1a	167.3b	168.7	169.4a	165.3c	167.4	168.0
Plant height at maturity (cm)	189.3b	174.1b	181.7	173.1c	213.6a	193.4	187.5
No. of primary branches plant <sup>-1</sup>	15.8a	9.9b	12.8	15.0a	9.5b	12.3	12.5
No. of secondary branches plant <sup>-1</sup>	56.6a	54.5a	55.5	43.8b	33.6c	38.7	47.1
No. of pods plant <sup>-1</sup>	665.3a	514.9bc	590.1	522.0b	414.2c	468.1	529.1
Seeds pod <sup>-1</sup>	3.5b	3.5b	3.5	3.7ab	3.6b	3.6	3.6
100-seed weight (g)	11.3a	10.7c	11.0	11.2b	10.9c	11.0	11.0
Seed yield plant <sup>-1</sup> (g)	184.3a	127.9b	156.1	129.6b	128.7b	129.2	142.6
Seed yield ha <sup>-1</sup> (kg)	4224.8a	3149.5b	3687.1	2752.9c	3364.5b	3058.7	3372.9
Biomass ha <sup>-1</sup> (kg)	8851.9a	8975.9a	8913.9	6376.9c	7955.2b	7166.1	8040.0
Harvest index (%)	32.3a	26.0b	29.3	30.2a	29.7a	29.9	29.6

**Note :** Means followed by the same letter in a row do not differ significantly at P = 0.05

E<sub>1</sub>- 2011 at Parbhani, E<sub>2</sub> - 2012 at Parbhani, E<sub>3</sub> - 2011 at Patancheru, E<sub>4</sub> - 2012 at Patancheru.

showed greater plant height (197.3 cm) at maturity, number of primary branches plant<sup>-1</sup> (13.8) and higher number of secondary branches plant<sup>-1</sup> (54.1) which is directly correlated to increased dry matter accumulation. The higher dry matter accumulation of hybrids (18%) resulted in increase in seed yield (3513.2 kg ha<sup>-1</sup>) as compared to varieties (3232.9 kg ha<sup>-1</sup>). The seed yield plant<sup>-1</sup> of hybrids was significantly higher (151.4 g) than varieties (133.9 g) due to more number of pods plant<sup>-1</sup> and 100-seed weight (11.4 g). These results indicated that hybrids out performed varieties for seed yield and its related traits which conforms to the findings of Saxena *et al.* (2013), Meena *et al.* (2013) and Tuppad *et al.* (2012).

**Treatments effect :** Days to 50% flowering, plant height at maturity and seed pod<sup>-1</sup> was not significantly influenced by the effect of various treatments over two years and across the environments (Table 4). However, significant number of primary branches plant<sup>-1</sup> (13.2), higher number of pods plant<sup>-1</sup> (624.8)

and higher seed yield plant<sup>-1</sup> (163.9) are observed in T<sub>3</sub> as compared to T<sub>2</sub> (12.5; 549.2; 151.1g, respectively) and T<sub>1</sub> (11.9; 413.3 and 112.8 g, respectively). Among three treatments, T<sub>2</sub> showed the highest in biomass weight (8816.6 kg ha<sup>-1</sup>) which significantly contributed to increase in seed yield (3923.9 kg ha<sup>-1</sup>) and harvest index (30.8%) as compared to T<sub>3</sub> (7670.1 kg ha<sup>-1</sup>; 29.8%, respectively) and T<sub>1</sub> (7633.3 kg ha<sup>-1</sup>; 27.8%, respectively). The results indicated that pigeonpea genotypes with wider spacing, higher fertilizer rate and protective irrigations produced significantly higher biomass and seed yield which corresponds to the findings of Goud *et al.* (2012).

**Environment × treatment interactions effect :** The environment × treatment interaction were significantly different for days to maturity, number of pods plant<sup>-1</sup>, seed yield (kg ha<sup>-1</sup>) and biomass weight (kg ha<sup>-1</sup>) as shown in Table 1. Among all environment and treatment combinations, E<sub>4</sub>T<sub>1</sub> (2012 at Patancheru with T<sub>1</sub>) matured early (164.5 days) whereas, E<sub>1</sub>T<sub>1</sub>

**Table 3.** *Per se* performance of hybrids and varieties of pigeonpea for yield and yield contributing characters for two years and across the environments

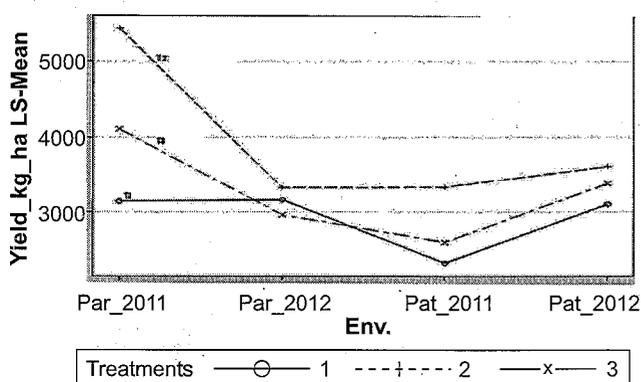
Geno- types	Days to 50% flowering	Days to 80% maturity	Plant height at maturity (cm)	No. of primary branches plant <sup>-1</sup>	No. of secondary branches plant <sup>-1</sup>	No. of pods plant <sup>-1</sup>	Seed pod <sup>-1</sup>	100-seed weight (g)	Seed yield plant <sup>-1</sup> (g)	Seed yield ha <sup>-1</sup> (kg)	Biomass ha <sup>-1</sup> (kg)	Harvest index (%)
<b>Hybrids</b>												
G <sub>1</sub>	114.1d	168.5c	191.5b	14.0a	57.1a	547.5b	3.5c	11.2b	154.9a	3556.9ab	8568.9ab	29.3ad
G <sub>2</sub>	120.1c	174.4b	201.0a	13.4ab	52.2ab	490.9c	3.6b	11.9a	145.8a	3425.2ab	8733.1a	28.2d
G <sub>3</sub>	123.2a	178.0a	199.3a	14.1a	53.0ab	629.4a	3.6b	11.1c	153.5a	3557.5a	8812.9a	28.8ad
Mean	119.1	173.6	197.3	13.8	54.1	555.9	3.6	11.4	151.4	3513.2	8705	28.8
<b>Varieties</b>												
G <sub>4</sub>	103.3e	155.1e	158.6d	8.9d	23.3d	431.9d	3.9a	9.8e	118.3b	2955.4c	6364.5d	31.7a
G <sub>5</sub>	114.5d	163.8d	185.0c	12.1c	45.8bc	545.8b	3.5c	10.7d	143.3a	3336.2b	7544.7c	30.7a
G <sub>6</sub>	122.0b	168.5c	189.8b	12.8bc	51.3bc	529.1bc	3.4c	11.4b	140.0a	3406.3ab	8215.7b	29.3bc
Mean	113.3	162.4	177.8	11.3	40.1	502.3	3.6	10.6	133.9	3232.6	7375	30.5

**Note :** Means followed by the same letter in a column do not differ significantly at P = 0.05

G<sub>1</sub> = ICPH 2671, G<sub>2</sub> = ICPH 2740, G<sub>3</sub> = ICPH 3762, G<sub>4</sub> = BDN711, G<sub>5</sub> = BSMR 736, G<sub>6</sub> = Asha

**Table 4.** Effect of treatments & environment × treatment interaction on various yield and yield attributes of pigeonpea

Treatments	Days to 50% flowering	Days to 80% maturity	Plant height at maturity (cm)	No. of primary branches plant <sup>-1</sup>	No. of secondary branches plant <sup>-1</sup>	No. of pods plant <sup>-1</sup>	Seed pod <sup>-1</sup>	100-seed weight (g)	Seed yield plant <sup>-1</sup> (g)	Seed yield ha <sup>-1</sup> (kg)	Biomass ha <sup>-1</sup> (kg)	Harvest index (%)
<b>Treatments effect</b>												
T <sub>1</sub>	115.9a	167.8b	187.7a	11.9b	39.1b	413.3c	3.6a	11.0a	112.8c	2932.3c	7633.3b	27.8c
T <sub>2</sub>	116.2a	167.8b	187.6a	12.5ab	51.9a	549.2b	3.6a	11.0a	151.1b	3923.9a	8816.6a	30.8a
T <sub>3</sub>	116.4a	168.5a	187.3a	13.2a	50.3a	624.8a	3.6a	10.9a	163.9a	3262.5b	7670.1b	29.8b
<b>Environment x Treatment interactions effect</b>												
E <sub>1</sub> T <sub>1</sub>	120.6	170.5a	188.8	15.8	50.3	538.2cd	3.5	11.4	152.8	3138.7ed	7586.5d	29.3d
E <sub>1</sub> T <sub>2</sub>	120.9	169.9a	187.3	14.9	57.1	675.9b	3.6	11.4	186.5	5436.0a	10566.9a	34.0a
E <sub>1</sub> T <sub>3</sub>	121.3	169.8ab	191.7	16.7	62.3	782.0a	3.5	11.2	213.6	4099.6b	8402.2c	32.8b
E <sub>2</sub> T <sub>1</sub>	114.4	166.5d	175.8	9.8	45.7	447.4de	3.5	10.8	111.3	3164.9ed	8801.2b	26.4f
E <sub>2</sub> T <sub>2</sub>	114.7	167.1c	174	10.4	62.2	558.1c	3.5	10.6	139.2	3321.8d	9701.4a	25.5g
E <sub>2</sub> T <sub>3</sub>	114.9	168.4ab	172.5	9.5	55.6	539.1cd	3.5	10.8	133.3	2961.9f	8425.2c	26.0ef
E <sub>3</sub> T <sub>1</sub>	118.7	169.8ab	175.5	13.9	36.4	385.1e	3.7	11	94.3	2323.3h	6013.0e	27.9e
E <sub>3</sub> T <sub>2</sub>	118.8	169.2a	173.5	15.3	50.9	507.8cd	3.7	11.3	136.9	3331.4d	7114.1d	31.9b
E <sub>3</sub> T <sub>3</sub>	119.7	169.3a	170.3	15.6	44.2	673.3b	3.6	11.2	157.5	2604.0g	6003.8e	30.3d
E <sub>4</sub> T <sub>1</sub>	110.2	164.5f	210.7	8.3	24.2	282.5f	3.6	11	92.8	3102.2ed	8132.4c	27.6e
E <sub>4</sub> T <sub>2</sub>	110.4	165.1e	215.5	9.6	37.3	455.0de	3.7	10.8	141.8	3606.8c	7884.1dc	31.4c
E <sub>4</sub> T <sub>3</sub>	109.8	166.4d	214.6	10.7	39.4	504.9cd	3.6	10.7	151.6	3384.6d	7849.1dc	30.1d



**Fig. 1.** The graph showing the result of environment × treatment interaction for the trait seed yield (kg ha<sup>-1</sup>)

(2011 at Parbhani with T<sub>1</sub>) matured late (170.5 days) (Table 4). High seed yield (5436.0 kg ha<sup>-1</sup>), biomass weight (10566.9 kg ha<sup>-1</sup>) and harvest index (34%) were obtained in E<sub>1</sub>T<sub>2</sub> (2011 at Parbhani with T<sub>2</sub>) interaction. However, the highest number of pods plant<sup>-1</sup> (782.0), number of secondary branches plant<sup>-1</sup> (62.3) and primary branches plant<sup>-1</sup> (16.7) were recorded in E<sub>1</sub>T<sub>3</sub> (2011 at Parbhani with T<sub>3</sub>). The graphical presentation of environment × treatment interaction (**Fig. 1**) clearly indicated that E<sub>1</sub>T<sub>2</sub> (2011 at Patancheru with T<sub>2</sub>) interacted best for recording the highest seed yield (kg ha<sup>-1</sup>) followed by E<sub>1</sub>T<sub>3</sub> (2011 at Parbhani with T<sub>3</sub>).

**Genotype × treatment interactions effect :** Genotype × treatment interaction showed significant variation for days to 80% maturity and number of pods plant<sup>-1</sup> (Table 1). For days to 80% maturity, G<sub>2</sub>T<sub>2</sub> (BDN 711 + T<sub>2</sub>) interaction matured early

(154.9) whereas, G<sub>3</sub>T<sub>3</sub> (ICPH 3762 + T<sub>3</sub>) matured late (179.0) than rest of the interactions (Table 5). The highest number of pods plant<sup>-1</sup> was recorded in the G<sub>3</sub>T<sub>3</sub> (804.5) however, the lowest number of pods plant<sup>-1</sup> was obtained in G<sub>4</sub>T<sub>1</sub> (BDN 711 + T<sub>1</sub>) (378.4). Moreover, ICPH 3762 (G<sub>3</sub>) recorded higher pods plant<sup>-1</sup> in T<sub>1</sub> (459.6), T<sub>2</sub> (630.9) & T<sub>3</sub> (804.5) as compared to other genotypes. Among all the interactions (Table 5), G<sub>3</sub>T<sub>2</sub> (ICPH 3762 + T<sub>2</sub>) recorded highest seed yield (4153.6 kg ha<sup>-1</sup>) followed by G<sub>1</sub>T<sub>2</sub> (4113.7 kg ha<sup>-1</sup>) & G<sub>5</sub>T<sub>2</sub> (4073.2 kg ha<sup>-1</sup>) while G<sub>4</sub>T<sub>1</sub> (BDN 711 + T<sub>1</sub>) exhibited lowest seed yield (2557.9 kg ha<sup>-1</sup>).

The findings of present investigation revealed that hybrids with different treatments performed better than the control varieties for plant height, number of branches plant<sup>-1</sup>, no. of pods plant<sup>-1</sup> & seed yield (kg ha<sup>-1</sup>). The results are in conformity of the finding of Meena *et al.* (2013) & Goud *et al.* (2012).

**Environment × genotype × treatment interactions effect**

: The number of pods plant<sup>-1</sup> and biomass weight (kg ha<sup>-1</sup>) showed significant variation due to environment × genotype × treatment interaction (Table 1). The highest number of pods plant<sup>-1</sup> was recorded in E<sub>1</sub>G<sub>3</sub>T<sub>3</sub> (2011 at Parbhani by ICPH 3762) at 1181.9 (Table 6) followed by E<sub>1</sub>G<sub>3</sub>T<sub>2</sub> (2011 at Parbhani by ICPH 2740) at 1026.1. The higher number of pods plant<sup>-1</sup> recorded by E<sub>1</sub>G<sub>3</sub>T<sub>3</sub> and E<sub>1</sub>G<sub>3</sub>T<sub>2</sub> produced significantly highest seed yield plant<sup>-1</sup> (237.1 g and 240.1 g respectively). Among all interactions, E<sub>2</sub>G<sub>2</sub>T<sub>2</sub> (2012 at Parbhani with ICPH 2740) revealed the highest biomass weight (11977.9 kg ha<sup>-1</sup>) followed by the interaction of E<sub>2</sub>G<sub>3</sub>T<sub>2</sub> (2012 at Parbhani with ICPH 3762) at 11360.7 kg ha<sup>-1</sup> and E<sub>2</sub>G<sub>1</sub>T<sub>2</sub> (2012 at Parbhani with ICPH 2740) at 11165.0 kg ha<sup>-1</sup> (Table 6).

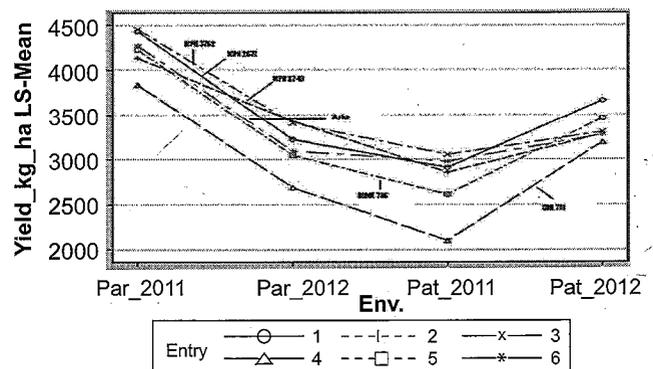
**Table 5.** Interaction effect of genotype × treatment on various yield and its traits of pigeonpea for (two years)

Treat-ments	Days to 50% flowering	Days to 80% maturity	Plant height at maturity (cm)	No. of primary branches plant <sup>-1</sup>	No. of secondary branches plant <sup>-1</sup>	No. of pods plant <sup>-1</sup>	Seed pod <sup>-1</sup>	100-seed weight (g)	Seed yield plant <sup>-1</sup> (g)	Seed yield ha <sup>-1</sup> (kg)	Biomass ha <sup>-1</sup> (kg)	Harvest index (%)
G <sub>1</sub> T <sub>1</sub>	114	168.4f	192.2	13.4	46.4	452.6fg	3.5	11.1	126.6	3125	8070.2	29.9
G <sub>2</sub> T <sub>1</sub>	119.9	174.7c	202.9	12.9	43.1	386.4g	3.6	12	112.6	2913	8337.9	25.9
G <sub>3</sub> T <sub>1</sub>	123	177.5b	196.8	13.6	42.3	459.6f	3.6	11.1	125	3022.3	8641.9	25.9
G <sub>4</sub> T <sub>1</sub>	103.3	155.1i	156.7	7.7	21.1	378.4g	3.9	9.8	88.4	2557.9	5837.4	30.5
G <sub>5</sub> T <sub>1</sub>	114.1	163.6h	186.8	11.5	37.6	402.2fg	3.4	10.7	111.6	2983.9	7197.4	29.3
G <sub>6</sub> T <sub>1</sub>	121.3	167.8g	190.8	12.5	44.3	400.7fg	3.4	11.5	112.6	2991.6	7714.7	27.9
G <sub>1</sub> T <sub>2</sub>	114.1	168.0f	191.5	15	65.7	598.9cb	3.5	11.3	163.8	4113.7	9445.1	30.3
G <sub>2</sub> T <sub>2</sub>	119.6	173.8d	200	13	52.6	495.7f	3.6	11.8	151.8	3860.6	9698.3	28.5
G <sub>3</sub> T <sub>2</sub>	123.1	177.5b	202.7	12.8	57.4	630.9b	3.6	11.1	154.3	4153.6	9373.4	30.7
G <sub>4</sub> T <sub>2</sub>	103.3	154.9i	159.9	9	24.7	450.2fg	3.9	9.8	133.8	3537.6	7201.9	32.9
G <sub>5</sub> T <sub>2</sub>	115.1	163.7h	181	12.3	53	548.2ed	3.5	10.7	152.6	3805.1	8105.5	31.9
G <sub>6</sub> T <sub>2</sub>	122.2	168.9f	190.3	13.1	57.8	571.2dc	3.5	11.4	150.4	4073.2	9075.7	31.0
G <sub>1</sub> T <sub>3</sub>	114.2	169.1e	190.8	13.5	59.2	584.0dc	3.5	11.3	174.2	3432	8191.5	29.5
G <sub>2</sub> T <sub>3</sub>	120.9	174.9c	200	14.3	60.8	590.7dc	3.6	11.9	173	3502	8163.2	30.0
G <sub>3</sub> T <sub>3</sub>	123.6	179.0d	198.3	15.9	59.3	804.5a	3.6	11	181.3	3496.7	8423.5	29.3
G <sub>4</sub> T <sub>3</sub>	103.3	155.2i	159.3	10	24.2	467.2fg	3.8	9.8	132.7	2770.7	6054.2	31.4
G <sub>5</sub> T <sub>3</sub>	114.3	164.2h	187.1	12.4	46.9	687.1b	3.5	10.6	165.6	3219.5	7331.1	30.5
G <sub>6</sub> T <sub>3</sub>	122.4	168.8f	188.3	12.9	51.7	615.4b	3.5	11.3	157	3154.2	7856.8	28.6

**Note :** Means followed by the same letter in a column do not differ significantly at P = 0.05

G<sub>1</sub> = ICPH 2671, G<sub>2</sub> = ICPH 2740, G<sub>3</sub> = ICPH 3762, G<sub>4</sub> = BDN711, G<sub>5</sub> = BSMR 736, G<sub>6</sub> = Asha

The present investigation revealed that pigeonpea hybrids recorded significantly higher number of pods plant<sup>-1</sup> and biomass (kg ha<sup>-1</sup>), which contributes into higher seed yield as compared to pure-line cultivars. Various treatments significantly influenced the seed yield plant<sup>-1</sup>, seed yield (kg ha<sup>-1</sup>), biomass (kg ha<sup>-1</sup>), and harvest index (%). The genotype × environment interaction over four environments (Fig. 2) indicated that hybrids obtained greater seed yield than inbred cultivars. Likewise, the genotypes showed better performance in wider spacing, higher fertilizer rate and two protective irrigations (T<sub>2</sub> and T<sub>3</sub>) due to greater availability of space, light, nutrients, and moisture as compared to closer spacing, normal fertilizer rate and no protective irrigation. Hybrid ICPH 3762 recorded higher number of pods plant<sup>-1</sup> in T<sub>1</sub> (459.6), T<sub>2</sub> (630.9) and T<sub>3</sub> (804.5) as compared to other genotypes and may have the potential to produced more in intensive cultivation (T<sub>2</sub> and T<sub>3</sub>). However, in environment × genotype × treatment interaction, E<sub>1</sub>G<sub>3</sub>T<sub>3</sub> exhibited significantly higher number of pods plant<sup>-1</sup> (1181.9) followed by E<sub>1</sub>G<sub>3</sub>T<sub>2</sub> (1026.1) that significantly converted into higher seed yield per plant<sup>-1</sup> (237.1g and 240.1g respectively). While E<sub>2</sub>G<sub>2</sub>T<sub>2</sub> obtained significantly greater biomass (11977.9 kg ha<sup>-1</sup>). However, the findings of the present study showed that among the genotypes and treatment interactions, hybrid ICPH 3762 and ICPH 2671 produced the highest seed yield (4153.6 kg ha<sup>-1</sup> and 4113.7 kg ha<sup>-1</sup> respectively) following wider spacing of 75 × 60 cm applied with higher fertilizer rate (50 kg N:100 P<sub>2</sub>O<sub>5</sub> kg ha<sup>-1</sup> (split application at 50% as basal and 50% at 60 days after sowing) and 2 protective irrigations during flowering & pod development.



**Fig. 2.** The graph showing the result of genotype × environment interaction for the trait seed yield (kg ha<sup>-1</sup>)

This data set indicated that the cultivation of high yielding hybrids in wider spacing, higher fertilizer rate and two protective irrigations (like T<sub>2</sub> and T<sub>3</sub>) could be help to increase the productivity of pigeonpea upto some extent.

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**Table 6.** Environment × genotype × treatment interaction effect on various yield and yield attributes of pigeonpea

Genotypes	No. of pods plant <sup>-1</sup>											
	Environments											
	E <sub>1</sub>			E <sub>2</sub>			E <sub>3</sub>			E <sub>4</sub>		
	Treatments											
	T <sub>1</sub>	T <sub>2</sub>	T <sub>3</sub>	T <sub>1</sub>	T <sub>2</sub>	T <sub>3</sub>	T <sub>1</sub>	T <sub>2</sub>	T <sub>3</sub>	T <sub>1</sub>	T <sub>2</sub>	T <sub>3</sub>
G <sub>1</sub>	550.0a	614.7a	654.5a	544.4a	683.3a	575.9a	436.5a	523.7a	604.9a	307.4c	573.8ab	500.7b
G <sub>2</sub>	495.2a	520.4a	647.5a	468.0a	577.2a	641.8a	361.1b	524.3a	585.1a	221.1c	360.9ab	488.4a
G <sub>3</sub>	523.5c	1026.1b	1181.9ab	410.2a	634.1a	579.6a	481.9b	447.4b	866.7a	394.8cb	416.1b	589.9a
G <sub>4</sub>	663.9a	493.0a	490.2a	372.9a	432.5a	393.9a	261.9cb	421.8ba	562.3a	214.7c	453.5ab	422.4b
G <sub>5</sub>	491.2cb	692.7b	992.6a	386.5a	449.8a	533.1a	406.1cb	568.5ba	699.3a	324.7c	481.9b	523.3ab
G <sub>6</sub>	505.1b	708.3ab	725.1a	502.3a	571.8a	510.3a	363.0c	560.7ba	721.3a	232.5c	444.1b	504.9ab

Contd...

Genotypes	Seed yield plant <sup>-1</sup>											
	Environments											
	E <sub>1</sub>			E <sub>2</sub>			E <sub>3</sub>			E <sub>4</sub>		
	Treatments											
	T <sub>1</sub>	T <sub>2</sub>	T <sub>3</sub>	T <sub>1</sub>	T <sub>2</sub>	T <sub>3</sub>	T <sub>1</sub>	T <sub>2</sub>	T <sub>3</sub>	T <sub>1</sub>	T <sub>2</sub>	T <sub>3</sub>
G <sub>1</sub>	175.1	187.3	215.5	112.9	158.9	147.7	120.1	140.7	162.7	98.1	168.5	148.2
G <sub>2</sub>	166.6	189.6	217.5	116.5	161.2	155.6	89.9	143.9	168.6	77.5	112.5	150.4
G <sub>3</sub>	157.6	240.1	237.1	110.0	157.8	151.9	116.0	103.6	181.1	116.5	116.6	176.8
G <sub>4</sub>	112.9	135.2	170.2	102.1	105.5	99.6	60.0	137.1	121.6	78.6	157.4	139.4
G <sub>5</sub>	140.6	180.1	231.7	105.5	109.9	125.0	90.7	152.7	149.4	109.7	167.9	156.3
G <sub>6</sub>	164.1	187.6	208.5	120.8	142.2	119.8	89.2	143.6	161.3	76.3	128.2	138.4

Contd...

Genotypes	Biomass ha <sup>-1</sup> (kg)											
	Environments											
	E <sub>1</sub>			E <sub>2</sub>			E <sub>3</sub>			E <sub>4</sub>		
	Treatments											
	T <sub>1</sub>	T <sub>2</sub>	T <sub>3</sub>	T <sub>1</sub>	T <sub>2</sub>	T <sub>3</sub>	T <sub>1</sub>	T <sub>2</sub>	T <sub>3</sub>	T <sub>1</sub>	T <sub>2</sub>	T <sub>3</sub>
G <sub>1</sub>	8025.2cb	11165.0a	8687.1b	8797.0a	9351.2a	8433.6a	6681.7bc	8777.3a	6586.5c	8776.9a	8487.0a	9058.9a
G <sub>2</sub>	8395.5cb	10504.3a	8199.3b	10263.2ab	11977.9a	9060.2b	6573.9a	6857.1a	6842.1a	8119.1cb	9453.6a	8551.4ba
G <sub>3</sub>	8423.6cb	10991.6a	9006.5b	11140.4ab	11360.7a	10225.6b	7051.4ab	7848.7a	6726.8b	7952.4a	7292.5a	7735.0a
G <sub>4</sub>	5778.3c	9648.0ab	8250.6b	5526.3a	6750.7a	5137.8a	4644.1ab	5722.4a	4230.6b	7401.0a	6686.4ba	6597.8cb
G <sub>5</sub>	7610.3cb	10195.7a	7664.2b	8170.4a	8930.9a	8884.7a	5325.8a	6334.4a	5773.2a	7683.0a	6960.9a	7002.5a
G <sub>6</sub>	7286.1c	10896.9a	8605.3b	8909.8a	9837.1a	8809.5a	5800.8cb	7144.7a	5863.4b	8862.2a	8424.0a	8149.1a

**Note :** Means followed by the same letter in a row in individual environment do not differ significantly at P = 0.05

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