

## IDENTIFICATION AND INHERITANCE OF A NEW DWARFING GENE IN PIGEONPEA

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(Received: September 10, 1991; accepted: October 14, 1991)

### ABSTRACT

A spontaneous dwarf ( $D_{11}$ ) mutant was identified in an advanced line ICPL 146. In order to study inheritance of the dwarfness in  $D_{11}$  and its allelic relationship to the  $D_1$  dwarfing gene,  $D_{11}$  was crossed with three tall lines (ICPL 146, ICPL 85024, ICPL 85037) and a  $D_1$  dwarf (ICPL 85059) in 1986. The segregation patterns in  $F_1$ ,  $F_2$ , backcrosses to both the parents and  $F_3$  progenies suggested that  $D_{11}$  dwarfness is governed by a single recessive gene in homozygous condition ( $t_{11}t_{11}$ ). The genes in  $D_{11}$  and  $D_1$  were found to be nonallelic.

**Key words:** *Cajanus cajan*, dwarf mutant, inheritance.

The excessive vegetative growth related to tallness of traditional pigeonpea [*Cajanus cajan* (L.) Millsp.] cultivars leads to reduced harvest index and hinders efficient crop management practices. Delayed plantings can result in reduced height [1]. However, Mohammed and Ariyanayagam [2] argued that the use of genetic dwarfs would be a more desirable approach to reduce plant height.

A bushy dwarf pigeonpea with brittle branches and condensed internodes was reported [3–5]. They found that the dwarfness was controlled by a single recessive gene. Twelve sources of dwarfism ( $D_0$  to  $D_{11}$ ) in pigeonpea are available at ICRISAT Center. Genetic studies of the  $D_0$  indicated that the dwarfness was controlled by two nonallelic recessive genes  $t_1t_1$  and  $t_2t_2$  [6]. Jain [7] found that dwarfing in  $D_1$  was controlled by a single recessive gene ( $t_4t_4$ ). Inheritance of dwarfness  $D_6$ ,  $PD_1$  ( $D_7$ ) and PBNA ( $D_8$ ) indicated that the dwarf phenotype in each of the three lines was controlled by a single recessive gene in homozygous state [8]. They also reported that  $D_6$  and  $PD_1$  had similar alleles ( $t_3t_3$ ) and PBNA had a different allele ( $t_3^h t_3^h$ ) for dwarfness.

During 1986 rainy season a spontaneous dwarf mutant plant was identified at the ICRISAT Sub-Center, Hisar in an advanced short duration pigeonpea line ICPL 146. Its

height at maturity was 35 cm as against the 130 cm of ICPL 146. This dwarf was designated as D<sub>11</sub>. The present study was conducted to study the inheritance pattern of the dwarfing gene in D<sub>11</sub> and its allelic relationship to the gene controlling dwarfness in the D<sub>1</sub> dwarf, an extensively used parent in the crossing program at ICRISAT.

#### MATERIALS AND METHODS

Two dwarf (D<sub>1</sub> and D<sub>11</sub>) and three tall (ICPL 146, ICPL 85024 and ICPL 85037) pigeonpea lines were included in this study. Characteristics of these dwarf and tall parents are summarized in Table 1. The D<sub>11</sub> dwarf was the shortest parent with a mean height of 39.5 cm and ICPL 85037 was the tallest with a mean height of 120 cm. The mean plant height of D<sub>1</sub> dwarf (ICPL 85059) and tall parent ICPL 85024 was about the same (Table 1), however, the branching pattern and the internode length in these two parents were significantly different. ICPL 85024 had on an average 7.2 primary branches per plant at mean internode length of 5.3 cm, while ICPL 85059 (D<sub>1</sub> dwarf) had on an average 12.8 primary branches per plant at mean internode length of 1.9 cm. The internodes in D<sub>1</sub> dwarf are condensed so that acute branches radiate from a narrow region about 10 to 15 cm above the ground level. The main branches are brittle.

Table 1. Characteristics of the parents used in the study on pigeonpea

Parent	Plant height (cm)	No. of primary branches	Internode length (cm)	Days to flowering
D <sub>11</sub> dwarf	39.5 ± 1.7	5.8 ± 0.3	3.0 ± 0.1	61.8 ± 0.4
D <sub>1</sub> dwarf (ICPL 85059)	85.7 ± 1.4	12.8 ± 0.7	1.9 ± 0.1	64.1 ± 0.6
ICPL 146	106.4 ± 0.9	7.9 ± 0.4	7.2 ± 0.2	66.5 ± 0.4
ICPL 85024	85.6 ± 1.0	7.2 ± 0.3	5.3 ± 0.2	58.5 ± 0.5
ICPL 85037	120.0 ± 0.6	9.0 ± 0.4	8.7 ± 0.2	63.6 ± 0.4

Each of the two dwarf lines was crossed to all the three tall parents and also among themselves to study allelic relationship. The F<sub>1</sub>s were grown during 1987 at Hisar to produce F<sub>2</sub> seed and to backcross with both the parents. The parents, F<sub>1</sub>, F<sub>2</sub> and backcross to both the parents were grown during 1988 at Hisar. The parents, F<sub>1</sub>, and the backcrosses were planted in one row and F<sub>2</sub> populations were grown in 20 row plots of 9 m length. The rows were spaced 60 cm apart with intra-row spacing of 15–20 cm. The number of dwarf and tall plants in each generation for each of the four crosses were recorded. In each of the three F<sub>2</sub> populations involving crosses between D<sub>11</sub> dwarf and the three tall parents, 20–50 and 52–231 tall plants were selected randomly to study the segregation pattern in the F<sub>3</sub> generation. In the 1989 rainy season F<sub>2</sub>-derived F<sub>3</sub> progenies were grown at Hisar, along

with their respective parents, in 9 m long one row plots. The observation on segregation of tall and dwarf plants in each single plant progeny was recorded separately for each of the three crosses. The chi-square test was applied to test the significance of segregation ratios.

## RESULTS AND DISCUSSION

### INHERITANCE

The F<sub>1</sub> plants of all the three crosses involving D<sub>11</sub> dwarf and the three tall parents resembled their tall parents, suggesting that D<sub>11</sub> dwarf is inherited as a recessive trait. In the F<sub>2</sub> populations obtained by crossing D<sub>11</sub> dwarf with tall parents, the observed segregation of tall and dwarf plants fitted the expected 3 tall : 1 dwarf ratio indicating that the D<sub>11</sub> dwarf phenotype was controlled by a single recessive gene in homozygous state (Table 2). This was further confirmed by the phenotypic segregation patterns in the backcrosses (Table 2) and F<sub>3</sub> progenies (Table 3). The backcross of F<sub>1</sub> to tall parent produced only tall progenies. Segregation in the test cross (F<sub>1</sub> × D<sub>11</sub> dwarf) progenies of all the three crosses showed a good fit to the expected ratio of 1 tall : 1 D<sub>11</sub> dwarf (Table 2). As expected within each cross, all the F<sub>3</sub> progenies of D<sub>11</sub> dwarf F<sub>2</sub> plants bred true for dwarfness. However, two-thirds of F<sub>3</sub> progenies of tall F<sub>2</sub> plants segregated for D<sub>11</sub> dwarf and tall plants and the remaining one-third bred true for tallness (Table 3). Within each segregating progeny, good fit for 3 tall : 1 D<sub>11</sub> dwarf was found. The data pooled over the segregating F<sub>3</sub> progenies in each of the three crosses (Table 4) also showed a good fit for the expected 3 tall : 1 D<sub>11</sub> dwarf ratio. These observations confirmed that D<sub>11</sub> dwarfness was governed by a single recessive gene which we designate as *t5t5*. The dwarf stature in pigeonpea has been reported to be controlled by a single recessive gene [3–5, 7–9].

Table 2. Phenotypic classification of F<sub>2</sub> and test cross progenies between D<sub>11</sub> dwarf and three tall pigeonpea lines

Generation and cross	total	Number of plants				Ratio	$\chi^2$	P
		observed		expected				
		tall	dwarf	tall	dwarf			
F <sub>2</sub> : D <sub>11</sub> × ICPL 146	1211	909	302	908.25	302.7	3:1	0.003	0.90–0.95
BC : F <sub>1</sub> × D <sub>11</sub>	21	11	10	10.50	10.5	1:1	0.047	0.80–0.90
F <sub>2</sub> : D <sub>11</sub> × ICPL 85024	1257	952	305	942.75	314.2	3:1	0.362	0.50–0.60
BC : F <sub>1</sub> × D <sub>11</sub>	23	13	10	11.50	11.5	1:1	0.391	0.50–0.60
F <sub>2</sub> : D <sub>11</sub> × ICPL 85037	1661	1262	399	1245.75	415.2	3:1	0.848	0.30–0.40
BC : F <sub>1</sub> × D <sub>11</sub>	19	10	9	9.50	9.5	1:1	0.052	0.80–0.90
Pooled : F <sub>2</sub>	4129	3123	1006	3096.75	1032.2	3:1	0.889	0.30–0.40
F <sub>1</sub> × D <sub>11</sub>	63	34	29	31.50	31.5	1:1	0.397	0.50–0.60

**Table 3. Segregation in F<sub>3</sub> progenies grown from random tall F<sub>2</sub> plants of the crosses between D<sub>11</sub> dwarf and three tall parents of pigeonpea**

Cross	total	Number of F <sub>3</sub> progenies				Ratio tested	$\chi^2$	P
		observed		expected				
		segregating	non-segregating (tall)	segregating	non-segregating (tall)			
D <sub>11</sub> x ICPL 146	98	63	35	65.3	32.7	2:1	0.252	0.60-0.70
D <sub>11</sub> x ICPL 85024	231	149	82	154.0	77.0	2:1	0.486	0.40-0.50
D <sub>11</sub> x ICPL 85037	52	32	20	34.7	17.3	2:1	0.616	0.30-0.40
Pooled	381	244	137	254.0	127.0	2:1	1.180	0.20-0.30

ALLELIC RELATIONSHIP WITH D<sub>1</sub> DWARF

The allelic relationship of D<sub>11</sub> and D<sub>1</sub> (ICPL 85059) dwarfs was studied in F<sub>1</sub>, F<sub>2</sub> and backcrosses to both the dwarf parents. All the plants in F<sub>1</sub> between D<sub>1</sub> and D<sub>11</sub> dwarfs were tall, indicating that they have separate genes controlling their dwarfness designated as t<sub>4</sub>t<sub>4</sub> and t<sub>5</sub>t<sub>5</sub>, respectively. Out of 1482 plants studied in F<sub>2</sub>, 830 were tall, 386 were of D<sub>1</sub> dwarf type and 266 were of D<sub>11</sub> dwarf type (Table 5) fitting the expected segregation ratio of 9:3:4. Presence of both the dominant genes (T<sub>4</sub> - and T<sub>5</sub> -) resulted in tall plants. Plants having t<sub>5</sub>t<sub>5</sub> in recessive homozygous form in the absence of t<sub>4</sub>t<sub>4</sub> (T<sub>4</sub>-t<sub>5</sub>t<sub>5</sub>) were of D<sub>11</sub> dwarf types and the plants having t<sub>4</sub>t<sub>4</sub> in recessive homozygous form (t<sub>4</sub>t<sub>4</sub>T<sub>5</sub>- and t<sub>4</sub>t<sub>4</sub>t<sub>5</sub>t<sub>5</sub>) were of D<sub>1</sub> dwarf types. In double homozygous recessive plants (t<sub>4</sub>t<sub>4</sub>t<sub>5</sub>t<sub>5</sub>), t<sub>4</sub>t<sub>4</sub> masked the effect of t<sub>5</sub>t<sub>5</sub> resulting in D<sub>1</sub> dwarfs. As expected backcross of F<sub>1</sub> with D<sub>1</sub> dwarf segregated into 1 tall : 1 D<sub>1</sub> dwarf and with D<sub>11</sub> dwarf into 1 tall : 1 D<sub>11</sub> dwarf, respectively (Table 5). These observations confirmed that the D<sub>1</sub> and D<sub>11</sub> dwarfness in pigeonpea was controlled by two different recessive genes t<sub>4</sub>t<sub>4</sub> and t<sub>5</sub>t<sub>5</sub>, respectively in homozygous state.

**Table 4. Pooled segregation for tall and D<sub>11</sub> dwarf types within the tall F<sub>3</sub> segregating progenies from the crosses between D<sub>11</sub> dwarf and three tall parents of pigeonpea**

Cross	No. of F <sub>3</sub> progenies	total	Number of plants				Ratio tested	$\chi^2$	P
			observed		expected				
			tall	dwarf	tall	dwarf			
D <sub>11</sub> x ICPL 146	63	2216	1680	536	1662.0	554.0	3:1	0.779	0.30-0.40
D <sub>11</sub> x ICPL 85024	149	5523	4161	1362	4142.3	1380.7	3:1	0.339	0.50-0.60
D <sub>11</sub> x ICPL 85037	32	1097	833	264	822.7	274.3	3:1	0.510	0.40-0.50
Pooled	244	8836	6674	2162	6627.0	2209.0	3:1	1.333	0.20-0.30

Table 5. Segregation pattern in F<sub>1</sub>, F<sub>2</sub> and backcross between D<sub>1</sub> and D<sub>11</sub> dwarfs of pigeonpea

Generation and cross	Number of plants						Ratio	$\chi^2$	Probability	
	total	observed			expected					
		tall	D <sub>11</sub> dwarf	D <sub>1</sub> dwarf	tall	D <sub>11</sub> dwarf				D <sub>1</sub> dwarf
F <sub>2</sub> : D <sub>1</sub> × D <sub>11</sub>	1482	830	266	386	833.6	277.9	370.5	9:3:4	1.166	0.60-0.70
BC: F <sub>1</sub> × D <sub>1</sub>	27	15	—	12	13.5	—	13.5	1:1	0.333	0.50-0.60
BC: F <sub>1</sub> × D <sub>11</sub>	23	13	10	—	11.5	11.5	—	1:1	0.391	0.50-0.60

The d<sub>11</sub> dwarf provides an additional source of dwarfness in pigeonpea. Unlike ICPL 85059 (D<sub>1</sub> dwarf) its branches are not brittle. However, its usefulness and linkages with other characteristics has yet to be studied.

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