



Assessment of somaclonal variations in two lines of pearl millet [*Pennisetum glaucum* (L.) R. Br.]

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Abstract

Somaclonal variation has great potential to enhance the advantages of conventional plant breeding, and increase the production and productivity of crops to meet the increasing demand for food and food products in the future. In pearl millet [*Pennisetum glaucum* (L.) R. Br.], plant morphogenesis has been achieved via somatic embryogenesis. Somatic embryos were isolated and regenerated into whole green plants on MS medium supplemented with indole-3-acetic acid (1 mg l^{-1}) and kinetin (0.5 mg l^{-1}). Ten R_0 plants of each cultivar were selected on the basis of 13 quantitative characters. Transmission of somaclonal variation was studied from callus derived (R_0) plants to R_1 and R_2 generations and were compared with those obtained from embryo culture without any callus formation (E_0 - E_2). Significant variation was observed among regenerants for number of leaves/plant, days to heading, length of panicle, average width of panicle and weight of panicle.

Key words: Pearl millet, agronomic performance, somatic embryogenesis, tissue culture

Introduction

It is well known that genetic changes occur in plant tissue culture and these changes expressed as variant traits, are transmitted to regenerated plants and their progeny through sexual or vegetative propagation [1, 2]. Genetic variation is an essential component of any conventional crop breeding programme. Plant cell and tissue culture provides increased genetic variability relatively rapidly. Somaclonal variation has been reported in a number of crops including rice [3], maize [4], wheat [5], barley [6] and pearl millet [7, 8]. There may be modifications in copy number of genes [9], activation of transposable elements [10] and changes in DNA methylation patterns [4] which may lead to variation for quantitatively inherited characters. The present study reports on the agronomic performance of plants derived from callus cultures of pearl millet and their progeny for thirteen quantitative characters.

Material and methods

Plants regenerated from embryos and callus derived

from immature embryo culture of pearl millet [*Pennisetum glaucum* (L.) R. Br.] var. RIB 3135 and var. RIB 20K86 were investigated for somaclonal variation and its transmission in three successive generations.

Somatic tissue culture: Entire panicle containing immature embryos (10-15 days after pollination) of pearl millet var. RIB 20K86 and var. RIB 3135 were taken from plants grown in the fields of Agricultural Research Station, Durgapura, Jaipur, India. Initially the entire panicle was washed with 20% (w/v) liquid detergent (Extran) (Merck) and rinsed 3-4 times thoroughly with water. Panicle containing immature embryos was surface sterilized with 0.1% (w/v) HgCl_2 solution for 3 min and rinsed thoroughly 3 times with sterile distilled water. Immature embryos were cultured on MS basal medium (11) with varying concentrations of 2,4-D ($1-5 \text{ mg l}^{-1}$), 3% sucrose (w/v) and 0.9% (w/v) agar for somatic embryogenesis.

Plant regeneration: Approximately 200 mg embryogenic callus of each variety was transferred to regeneration medium consisting of MS medium supplemented with IAA (1 mg l^{-1}) and Kn (0.5 mg l^{-1}). All the cultures were incubated in a growth chamber. For rooting, plantlets were kept in tubes containing half strength MS medium. Regenerated wellrooted plantlets were transferred to pots containing 1:1 ratio of soil and compost at the Department of Botany, University of Rajasthan, Jaipur. The regenerated plants were referred to as the R_0 generation. Nineteen regenerants were obtained in var. RIB 20K86 and twenty three regenerants were obtained in var. RIB 3135. Plants were also numbered separately in both the varieties. Ten control plants (E_0) were also raised for each variety.

Morphological analysis: Ten R_0 plants of each variety were selected on the basis of 13 characters. From each of 20 R_0 plants 10 seeds of individual plants were collected and sown to raise 200 R_1 plants. All the plants were allowed to open pollinate. 200 R_1 plants progeny of 20 R_0 plants were compared with 20 E_1 plants progeny of ten control plants (E_0) of both varieties.

Seeds of 200R₁ plants were individually bulked and sown to raise 200R₂ plants. Ten plants have been used per progeny for recording data on morphological traits. R₁ and R₂ generations were evaluated in different seasons.

Data of each R₀, R₁ and R₂ progeny plants of each variety were recorded on maturity and after harvesting. Plants were evaluated for Plant height (cm), tillers / plant (nu.), internodes / plant (nu.), internode length (cm), number of leaves / plant, flag leaf length (cm), flag leaf width (cm), days to heading (nu.), number of panicles / plant (nu.), length of panicle (cm), average width of panicle (cm), weight of panicle (g) and average weight of 100 seeds (g).

Statistical analysis: Data analysis was performed using ANOVA. Somaclones (R₀, R₁ and R₂) and control plants (E₀, E₁ and E₂) were analyzed as independent experimental treatments. Each set of data was subjected to standard analysis of variance (ANOVA).

Results and discussion

R₀ generation: In var. RIB 3135 and RIB 20K86, R₀ generation showed significant variation for most of the 13 characters studied. Except for plant height in var. RIB 3135 (Table 1). In var. RIB 20K86, tillers/plant, number of panicles/plant and average weight of 100 seeds were significant only at p = 0.05.

Table 1. Mean performance of R₀ (Tissue culture raised plants) and E₀ (Seed grown plants) generation in pearl millet var. RIB 3135 & var. RIB 20K86

Character	Unit	Var. RIB 3135		Var. RIB 20K86	
		R ₀	E ₀	R ₀	E ₀
Plant height	(cm)	93.3 ^a	88.5 ^a	88.7 ^b	134.5 ^a
Tillers / plant	(nu.)	1.36 ^b	1.0 ^a	2.2 ^b	1.0 ^a
Internode number / plant	(nu.)	6.0 ^b	5.0 ^a	6.3 ^b	7.0 ^a
Internode length	(cm)	14.6 ^b	15.8 ^a	12.7 ^b	16.3 ^a
Number of leaves / plant	(nu.)	7.7 ^b	7 ^a	15.5 ^b	9.5 ^a
Flag leaf length	(cm)	26.4 ^b	29 ^a	19.8 ^b	31 ^a
Flag leaf width	(cm)	1.63 ^b	1.3 ^a	1.64 ^b	1.60 ^a
Days to heading	(nu.)	34.8 ^b	38.5 ^a	38 ^b	45 ^a
Number of panicles/ plant	(nu.)	1.36 ^b	1.0 ^a	2 ^b	1.0 ^a
Length of panicle	(cm)	11.45 ^b	9.1 ^a	10.7 ^b	15.5 ^a
Average width of panicle	(cm)	0.92 ^b	0.86 ^a	1.18 ^b	3.33 ^a
Weight of panicle	(g)	3.89 ^b	2.89 ^a	3.42 ^b	7.56 ^a
Average weight of 100 seeds	(g)	0.58 ^b	0.68 ^a	1.71 ^b	0.76 ^a

Means followed by the same letters are not significantly different at the 0.05 level of confidence (ANOVA)

R₁ and R₂ generation: In var. 3135, plant no. 2, 3, 4, 6, 9 performing better than control for five characters were selected for raising R₁ progeny and plant no. 10, 11, 12, 13 and 16 were selected on the basis of characters which were poorer than control

(Table 2). Significant variations were observed for characters like plant height, number of leaves/plant, days to heading, flag leaf width, average weight of 100 seeds at p < 0.05.

In var. 20K86 plant no. 5,6,8,15 and 22 were selected on the basis of any five dominant characters and plant no. 4, 12, 13, 18 and 23 were selected on the basis of characters which were poorer. Significant variation were observed for characters like internode length, day of heading, length of panicle, average width of panicle and average weight of 100 seeds at p < 0.05. In R₂ generation, for var. RIB 3135 non-significant differences were observed for all the characters studied (Table 3) while for var. RIB 20K86 significant differences were observed for average width of panicle. As somaclonal variations are genotype dependent, these may account for differential response of var. RIB 3135 and var. RIB 20K86.

These data suggest that *in vitro* environment is mutagenic to pearl millet and that significant variations can be generated for several characters in the plant. This corroborates previous reports on wheat also (12). Retaining this variation through R₂ generation in both varieties suggests that the variation is genetic. Morrish *et al.* (13) in their studies on quantitative characters suggested the involvement of epigenetic factors and concluded that genetic fidelity of the progeny of embryogenic tissue culture is dependent on the genotype of the explant.

This work supports previous studies indicating that variation generated during tissue culture is heritable. Regardless of the mechanisms by which genetic variation is produced, evidence has accumulated that somaclonal variation includes few agronomically useful types [14, 15, 6]. Genetic variability in tissue culture can be perceived as a new source of variability or as an undesirable consequence of *in vitro* culture. Further research is needed on the nature and causes of this variation.

The application of this culture technique to adapted varieties in conjugation with appropriate screening and testing may provide useful variation to plant breeders.

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Table 2. Mean performance of R₁ (progeny of successively numbered R₀ plants) and E₁ (progeny of E₀ plants) generation in pearl millet (var. RIB 3135) (A) and (var. RIB20K86) (B).

Plant generation and Number	Plant height (cm)		Tiller/plant	Internode number/plant		Internode length (cm)		Flag leaf length (cm)		Flag leaf width (cm)		Days to heading		No. of panicles/plant		Length of panicle (cm)		Av. width of panicle (cm)		Weight of panicle (g)		Av. weight of 100 seeds (g)			
	A	B		A	B	A	B	A	B	A	B	A	B	A	B	A	B	A	B	A	B	A	B	A	B
E ₁	90.5 ^a	104.6 ^c	1 ^a	1 ^a	8 ^a	6.5 ^a	15.5 ^a	15.9 ^a	28.5 ^a	30.6 ^a	1.4 ^a	1.55 ^a	38 ^a	43.5 ^a	1 ^a	1 ^a	8.8 ^a	15.9 ^a	0.93 ^a	3.4 ^a	2.7 ^a	5 ^a	0.68 ^a	0.7 ^a	
2R ₁	155.9 ^b	95.1 ^a	1 ^a	1 ^a	8.7 ^b	6.9 ^a	15.6 ^a	12.8 ^b	35.6 ^b	30.5 ^a	2.5 ^b	2.29 ^b	33.7 ^b	36.9 ^b	1 ^a	1 ^a	15.7 ^b	10.8 ^b	1.21 ^b	1.1 ^b	7 ^{ab}	3.3 ^b	0.56 ^b	0.4 ^b	
3R ₁	108 ^c	76.8 ^b	1 ^a	1.1 ^b	8.9 ^{ab}	6.8 ^a	12.3 ^b	10 ^c	21.5 ^c	24.8 ^b	1.6 ^c	1.7 ^a	30 ^c	33.9 ^c	1 ^a	1 ^a	10.5 ^c	9.7 ^c	1.06 ^c	0.9 ^c	3.2 ^b	2.2 ^c	0.6 ^{cb}	0.3 ^{cb}	
4R ₁	101.1 ^{dc}	85.1 ^{cb}	1 ^a	1 ^a	8.4 ^{cb}	7.7 ^b	11.8 ^{cb}	10.6 ^{dc}	36.6 ^{cb}	24.1 ^{cb}	2.6 ^{cb}	1.6 ^a	27.4 ^d	39.2 ^d	1 ^a	1 ^a	11.3 ^{dc}	8.9 ^{dc}	1.03 ^{dc}	1.1 ^{db}	3.8 ^c	2.4 ^{dc}	0.51 ^d	0.3 ^{db}	
6R ₁	110.3 ^{ec}	95 ^a	1 ^a	1 ^a	9 ^{db}	6 ^a	12.2 ^{db}	12 ^e	7.3 ^{cb}	27.7 ^e	2.2 ^e	1.6 ^a	30.4 ^{ec}	30.3 ^e	1 ^a	1 ^a	11.3 ^{ec}	11.1 ^{eb}	1.01 ^{ed}	1.3 ^e	2.2 ^a	5.9 ^e	0.36 ^e	0.5 ^e	
9R ₁	29 ^f	93.3 ^{dec}	1 ^a	1.33 ^c	5.9 ^e	6.6 ^a	7.6 ^e	9.2 ^c	10.6 ^d	15.1 ^f	31.3 ^a	0.8 ^f	2.1 ^c	32.8 ^b	41 ^f	1 ^a	1 ^a	12.8 ^f	11.1 ^{fb}	1.23 ^{fb}	1.2 ^f	4.2 ^{dc}	6.9 ^f	0.44 ^f	0.5 ^f
10R ₁	42.6 ^g	125 ^e	1 ^a	1.16 ^d	5.2 ^f	10 ^c	7.4 ^{le}	12.4 ^{gde}	8.1 ^{gd}	14.5 ^e	18.8 ^g	0.9 ^{gf}	1.9 ^{da}	46.9 ^g	32.3 ^g	1 ^a	1.16 ^b	10.4 ^{gde}	12.7 ^g	0.96 ^a	1.4 ^{ge}	1.5 ^e	3.9 ^{gb}	0.32 ^{ge}	0.4 ^{ge}
11R ₁	54.7 ^h	73.4 ^{lb}	1 ^a	1 ^a	8 ^{bc}	7.7 ^{db}	9.6 ^g	10.3 ^{bc}	9.4 ^{hbc}	9.4 ^a	21.9 ^{hc}	1.1 ^h	1.6 ^a	41.2 ^h	33.5 ^{hc}	1 ^a	9.3 ^{ai}	7.8 ^h	1.14 ^g	1.1 ^{hb}	1.4 ^f	1.6 ^{hc}	0.28 ^{hg}	0.3 ^{hc}	
12R ₁	57.0 ^h	110 ^a	1 ^a	1 ^a	7.4 ^g	7.2 ^{eb}	9.3 ^{hg}	13.7 ^b	10 ^{ad}	22.3 ^{ic}	29 ^{ae}	0.9 ^{fh}	1.8 ^{ea}	49.1 ⁱ	37.8 ⁱ	1 ^a	10.3 ^{hc}	10.1 ^{bc}	1.09 ^{hc}	1.2 ^f	2.1 ^{ae}	3.4 ^{ib}	0.31 ^{ieg}	0.2 ^f	
13R ₁	102 ^{ic}	127.2 ^{ge}	1 ^a	1 ^a	11.3 ^h	8.9 ^f	12.2 ^b	14.2 ^l	10.5 ^{le}	11.1 ^{ld}	30.3 ^{ak}	2.7 ^b	2.7 ^f	42.5 ^h	37.1 ^l	1 ^a	9.9 ^c	11.7 ^{hg}	1.12 ^{gh}	1.3 ^f	1.6 ^{ge}	5 ^a	0.25 ^h	0.5 ^{ief}	
16R ₁	109.1 ^{kc}	121.1 ^{he}	1 ^a	1 ^a	10.2 ⁱ	8.9 ^{gf}	11.2 ^c	13.8 ^k	10.6 ^{ke}	11.5 ^{gd}	31.7 ^k	2.9 ^{ae}	2.9 ^g	42.7 ^h	47.8 ^k	1 ^a	14.7 ⁱ	10.8 ^{db}	1.39 ⁱ	1.4 ^k	6.7 ^h	3.9 ^{jb}	0.46 ^{kd}	0.5 ^{kef}	

Mean values with the same letters are not significantly different at the 5% level using ANOVA

Table 3. Mean performance of R₂ (progeny of R₁ plants) and E₂ (progeny of E₁ plants) generation in pearl millet (Var. RIB 3135) (A) (Var. RIB20K86) (B)

Plant generation and Number	Plant height (cm)		Tiller/plant	Internode number/plant		Internode length (cm)		Flag leaf length (cm)		Flag leaf width (cm)		Days to heading		No. of panicles/plant		Length of panicle (cm)		Av. width of panicle (cm)		Weight of panicle (g)		Av. weight of 100 seeds (g)		
	A	B		A	B	A	B	A	B	A	B	A	B	A	B	A	B	A	B	A	B	A	B	A
E ₂	85.8 ^a	105 ^a	1 ^a	7.4 ^a	10 ^a	12.8 ^a	17.8 ^a	10.5 ^b	26.9 ^a	31 ^a	1.1 ^a	1.2 ^a	37 ^a	42.8 ^a	1 ^a	8.1 ^a	13.5 ^a	0.8 ^a	3.4 ^a	2.2 ^a	4.3 ^a	0.7 ^a	0.6 ^a	
2R ₂	152.2 ^b	53.6 ^b	1 ^a	8.6 ^a	5.8 ^b	12.7 ^a	10.5 ^b	11.4 ^a	38.9 ^a	21.8 ^b	2.8 ^a	1.4 ^a	32.4 ^b	36.6 ^b	1 ^a	19.7 ^a	12.2 ^a	2.1 ^a	1.6 ^b	7.6 ^a	4.7 ^a	0.4 ^b	0.5 ^b	
3R ₂	123.4 ^c	80.6 ^{cb}	1 ^a	0.9 ^a	6.8 ^{cb}	13 ^a	8.5 ^{cb}	9.6 ^{ab}	11.4 ^a	20.9 ^b	1.3 ^{ad}	1.4 ^a	29.2 ^c	33.4 ^c	1 ^a	10.7 ^{ab}	11.1 ^{ba}	1.7 ^a	1.6 ^{cb}	5.2 ^{ab}	4.9 ^a	0.8 ^{ab}	0.5 ^{cb}	
4R ₂	100.8 ^a	98 ^{abc}	1 ^a	8.6 ^a	9.8 ^{ab}	11.9 ^a	11.2 ^{db}	8.6 ^{ac}	11.6 ^a	39.8 ^{ab}	23.1 ^{dbc}	2.6 ^a	2.1 ^a	26.8 ^d	38.6 ^{dbc}	1 ^a	12.7 ^{ac}	9.5 ^c	1.7 ^a	1.3 ^{db}	4.5 ^{ac}	5.7 ^a	0.6 ^{cb}	0.4 ^d
6R ₂	120.4 ^{ac}	109.5 ^{abc}	1 ^a	9 ^a	8 ^{ab}	10.7 ^b	11.2 ^{eb}	10.8 ^a	11 ^{ac}	30.1 ^{fgb}	16.1 ^e	2.2 ^{ae}	1.3 ^{ac}	28.8 ^{ecd}	30.6 ^e	1 ^a	11.3 ^{ad}	9 ^{dc}	1.9 ^a	1.7 ^{eb}	5.8 ^{ad}	15.6 ^a	0.2 ^d	0.6 ^{ebd}
9R ₂	30.8 ^d	90.4 ^{dbc}	1 ^a	4.8 ^b	8.6 ^{cb}	7.7 ^c	6.7 ^c	8.8 ^{ad}	13.6 ^a	22.8 ^{cb}	19.5 ^{lc}	0.9 ^b	1.5 ^{ad}	31.8 ^{fbcd}	41.2 ^{abccc}	1 ^a	10.3 ^{ae}	9.3 ^{ec}	1.4 ^{ab}	1.3 ^{fb}	5.6 ^{ae}	6.6 ^{ad}	0.4 ^{ebd}	0.9 ^{abd}
10R ₂	42.4 ^e	64.4 ^{abd}	1 ^a	5.2 ^{cb}	6 ^b	7 ^{cd}	11.4 ^{gbl}	7.6 ^{ae}	10.6 ^{abd}	19.1 ^d	16 ^g	1 ^{ab}	0.8 ^b	47.8 ^{abcd}	38 ^{be}	1 ^a	9.2 ^{af}	9 ^{lc}	1.4 ^{acb}	1.4 ^{gbe}	5.8 ^{af}	7.8 ^{ae}	0.2 ^d	0.4 ^{ld}
11R ₂	59.8 ^{le}	188.4 ^{abc}	1 ^a	8.6 ^{ab}	10.6 ^{ab}	10.5 ^{ebc}	17.9 ^{abf}	9 ^{af}	15.8 ^a	21.7 ^{abcd}	36 ^{abccg}	1.2 ^{agb}	3.3 ^{ab}	40.4 ^{abcdg}	29.6 ^{ge}	1 ^a	11 ^{ag}	8.8 ^{gc}	1.4 ^{db}	1.5 ^{hbe}	8.3 ^a	5.9 ^{af}	0.4 ^{gbd}	0.9 ^{abd}
12R ₂	57.9 ^{pe}	14.6 ^{ec}	1 ^a	7.6 ^{ab}	6.6 ^{ab}	9.4 ^c	12.6 ^{hfl}	9.2 ^{ag}	14.6 ^{af}	23.1 ^{bd}	20.5 ^{hbcg}	0.9 ^{cb}	1.6 ^{ab}	48.6 ^{abcd}	36.4 ^{hbccc}	1 ^a	10.9 ^{ah}	9.6 ^{hbc}	1.3 ^{abe}	1.4 ^{ibe}	9.4 ^a	6.4 ^{agf}	0.7 ^{abid}	0.3 ^g
13R ₂	108 ^{ake}	158 ^{abcd}	1 ^a	11.2 ^{ab}	10 ^{ab}	13.1 ^{abc}	11.3 ^{bl}	11.4 ^a	14.2 ^a	31.8 ^{hbd}	31 ^{abccj}	2.9 ^{ab}	1.9 ^{abf}	42.6 ^{abcdh}	35.2 ^{bccc}	1 ^a	9.3 ^{ai}	8.1 ^k	1.4 ^{atb}	1.8 ^{ibe}	8.2 ^a	3.4 ^{ab}	0.4 ^{hbd}	0.5 ^{hbdg}
16R ₂	113.8 ^{he}	168 ^{abcd}	1 ^a	10.6 ^{ab}	10.4 ^{ab}	11.6 ^{gabc}	13.2 ^{bl}	11.6 ^a	12.8 ^{ae}	32.4 ^{abd}	28.9 ^{ibccg}	1.5 ^{alb}	2.2 ^{abg}	42.8 ^{abcdi}	48 ^{abccc}	1 ^a	8 ^{aj}	8 ^{ic}	1.3 ^{ag}	1.5 ^{kbe}	9.4 ^a	4.5 ^{abc}	0.5 ^{ibd}	0.7 ^{abidg}

Mean values with the same letters are not significantly different at the 5% level using ANOVA

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