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In vitro polyploidization from shoot tips of *Jatropha curcas* L.: a biodiesel plant

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Abstract Jatropha curcas L. has been considered one of the most promising alternatives for biofuel production and, thus, a relevant economic crop. In this context, in vitro tissue culture techniques such as organogenesis and embryogenesis have been conducted for mass clonal propagation of elite J. curcas lines. However, despite advancements, in vitro induction of polyploids has not yet been related for this crop. In this sense, the present study attempted to induce polyploidy in plantlets generated from shoot tips of J. curcas 'Gonçalo' $(2n = 2 \times = 22$ chromosomes, 2C = 0.85 pg). For this purpose, some criteria were adopted for selection of the most adequate colchicine treatment: (a) survival rate of the explants, and (b) number of tetraploid and (c) mixoploid plantlets. Tetraploid and mixoploid plantlets were obtained from different treatments, with 0.5 mM colchicine for 96 h being the most efficient. The plantlets were recovered and clonally propagated in tissue culture medium supplemented with indole-3-acetic acid and 6-benzylaminopurine. These results show that the tissue culture procedures were adequate for induction, propagation and recovery of tetraploid and mixoploid plantlets. Moreover, DNA ploidy level screening by flow cytometry was a practical and rapid strategy for selection of diploid, mixoploid and tetraploid plantlets. The

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tissue culture system presented here represents a reliable methodology for in vitro polyploid induction of this and other elite lines of *J. curcas*.

Keywords Jatropha curcas · Biofuel plant · In vitro polyploidization · DNA ploidy level · Flow cytometry · Tetraploid plantlets

Introduction

The genus *Jatropha* (family Euphorbiaceae, subfamily Crotonoideae, tribe Joannesieae) comprises over 175 native species, occurring in South to Central America (Mukherjee et al. 2011), Asia and Africa (Kumar and Reddy 2012). In particular, the species *Jatropha curcas* L. has recently been pointed out as a relevant economic crop (Kumar and Reddy 2012). The most important aspect of this species is its large potential for biofuel production, owing to high oil content of the seed, rapid growth and stiffness of the plant (Mukherjee et al. 2011) and the low oil production cost (Jha et al. 2007).

Due to the increasing demand for biofuel, breeding programs of *J. curcas* have been established in distinct countries, for instance Brazil, India, Senegal and Cape Verde (Divakara et al. 2010). In this context, in vitro tissue culture techniques have mainly been performed for mass clonal propagation of *J. curcas* elite lines (Kalimuthu et al. 2007). For this purpose, in vitro regeneration of *J. curcas* plantlets has been achieved mainly through organogenesis (Rajore and Batra 2005; Jha et al. 2007; Kalimuthu et al. 2007; Kumar and Reddy 2012) and embryogenesis procedures (Jha et al. 2007).

For some species, new elite plants have been obtained by in vitro polyploidization, which has received special

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attention as an important tool for plant breeding programs (Zhang et al. 2008). Polyploid plantlets, especially tetraploid ones, have been successfully recovered mainly for vegetable, ornamental and medicinal crops (Dhooghe et al. 2011). Chromosome doubling efficiency, and ultimately polyploidy induction, depends on various factors, such as: (a) species, (b) type of explant, (c) type and concentration of the anti-tubulinic agent, as well as exposure time, and (d) tissue culture medium for polyploidization and propagation (Dhooghe et al. 2011).

An efficient in vitro polyploidization system also requires effective methods for direct screening and monitoring of DNA ploidy level. Flow cytometry (FCM) is the quickest and most reliable method for this purpose, given that the DNA ploidy level can be determined for several plantlets in a short time (Clarindo et al. 2008). According to Ochatt (2008), application of FCM in plant science increased significantly with the use of innovative and interesting techniques for basic research and commercial breeding.

Considering the relevance of protocols for in vitro polyploidization of *J. curcas*, the present study attempted to induce polyploidy from shoot tips of this relevant biofuel species.

Materials and methods

Plant material

Seeds of *J. curcas* 'Gonçalo' were collected at a crop area located in Resende Costa, Minas Gerais, Brazil. *Solanum lycopersicum* L. 'Stupické' (primary standard for FCM, 2C = 2.00 pg; Praça-Fontes et al. 2011b) were kindly supplied by Dr. Jaroslav Doležel (Experimental Institute of Botany, Czech Republic).

2C-value and karyotype of J. curcas

The genome size of *J. curcas* was measured from nuclei suspensions extracted and stained according to procedure described by Carvalho et al. (2008). The suspensions were analyzed with a Partec PAS[®] cytometer (Partec[®] GmbH, Munster, Germany), equipped with a laser source (488 nm). Flow cytometer parameters (i.e. gain and channel) were determined before each measurement, based on external FCM analyses of primary standard (*S. lycopersicum* 'Stupické') and sample (*J. curcas*). Six independent repetitions, accounting for more than 10,000 nuclei, were carried out at each analysis.

2C-value of *J. curcas* was calculated by dividing the mean channel of the G_0/G_1 fluorescence peak from the primary standard by the mean channel of the G_0/G_1 peak

from each sample. Genome size mean value in picograms (pg) was converted to base pairs (bp), considering that 1 pg DNA corresponds to 0.978×10^9 bp (Doležel et al. 2003).

For karyotype characterization of *J. curcas* explant donor, metaphasic chromosomes were obtained from root tips treated according to Carvalho et al. (2008). The slides were prepared using the cell dissociation method, followed by air-drying technique (Carvalho et al. 2008). Subsequently, the slides were stained with a 5 % Giemsa solution (Merck[®]) in phosphate buffer (pH 6.8) for 5 min, washed twice in distilled water (dH₂O), air-dried and placed on a hot plate at 50 °C for 3 min.

Images of metaphase chromosomes were captured with a Media Cybernetics[®] Camera EvolutionTM charge-coupled device (CCD) video camera, mounted on a Nikon 80i microscope (Nikon, Japan).

In vitro recovery of J. curcas shoot tips

For shoot tip recovery, *J. curcas* seeds were decontaminated by immersion in 70 % ethanol (Merck[®]) for 1 min, followed by 2.5 % sodium hypochlorite for 20 min and rinsing three times with autoclaved dH₂O. After pericarp removal under laminar flow hood, the seeds were decontaminated again for 20 min with 2.5 % sodium hypochlorite solution containing one drop of Tween (Merck[®]) per 100 ml. Next, the seeds were washed with autoclaved dH₂O and inoculated into medium M1 (Table 1).

In vitro multiplication and assessment of DNA ploidy level

After 30 days in medium M1, shoot tips of the seedlings germinated in vitro were excised and inoculated into bottles containing 100 ml of medium M2 (Table 1). These cultures were maintained at 25 ± 1 °C under 16/8 h light/ dark regime, with 36 µmol m⁻² s⁻¹ light radiation.

As medium M2 (Table 1) was supplemented with indole-3-acetic acid (IAA) and 6-benzylaminopurine (BAP) (Rajore and Batra 2005; Mukherjee et al. 2011), the shoot tips were used to verify the occurrence of numeric chromosomal aberration, eu- and aneuploidy, associated with somaclonal variation.

For this purpose, leaf samples were collected from *J. curcas* plants seed-raised in greenhouse (control) and from plantlets propagated in medium M2 for 120 days (samples). Nuclei suspensions of control and samples were prepared as described by Clarindo et al. (2008) and CyStain UV Ploidy Partec[®] protocols. These suspensions were analyzed with a Partec-PAS[®] flow cytometer, equipped with an UV lamp emitting at 388 nm and a TK 420 filter. The FlowMax[®] software (Partec[®]) was used for data analyses. More than 10,000 nuclei were analyzed, and three

 Table 1 Composition of the tissue culture media

Compound	M1	M2	M3
MS salts (Sigma [®])	4.3 g l ⁻¹	4.3 g 1 ⁻¹	2.15 g l ⁻¹
MS vitamins ^a	$10 \text{ ml } l^{-1}$	$10 \text{ ml } l^{-1}$	$10 \text{ ml } l^{-1}$
Adenine sulfate (Fluka®)	_	$0.025 \text{ g } 1^{-1}$	-
Glutamine (Sigma [®])	-	$0.1 \text{ g } 1^{-1}$	-
BAP (Sigma [®])	-	$0.002 \text{ g } 1^{-1}$	-
IAA (Vetec [®])	-	$0.0005 \text{ g } \text{l}^{-1}$	-
Sucrose (Sigma [®])	$30 \text{ g } 1^{-1}$	$30 \text{ g } 1^{-1}$	$30 \text{ g} \text{ l}^{-1}$
Agar (Sigma [®])	$7 \text{ g } l^{-1}$	$7 \text{ g } 1^{-1}$	-
pH	5.7	5.7	5.7

M1 germination medium, M2 proliferation medium, M3 polyploidization liquid medium

^a Stock solution composed of glycine (0.20 g l^{-1}), nicotinic acid (0.05 g l^{-1}), pyridoxine (0.05 g l^{-1}) and thiamine (0.01 g l^{-1})

independent replications were performed for determination of DNA ploidy level in each *J. curcas* plantlet.

In vitro polyploidization

Shoot tip explants were excised from the plantlets obtained in medium M1 and placed in Erlenmeyer flasks containing 10 ml of liquid polyploidization medium (Praça et al. 2009) M3 (Table 1). This medium was supplemented with different concentrations (Table 2) of filter-sterilized colchicine (Sigma[®]). The flasks were shaken (40 rpm, at 25 ± 1 °C) in a growth room for different time periods (Table 2). Each flask contained three shoot tip explants, corresponding to three replicates for each treatment. After the set times, the explants were rinsed five times with autoclaved dH₂O and inoculated into medium M2 (Table 1). Cultures were maintained at 25 °C under a 16/8 h light/dark regime, with 36 µmol m⁻² s⁻¹ light radiation provided by two fluorescent lamps (20 W, Osram[®]).

The shoot tips were propagated for 4 months, being subcultured every 30 days into fresh medium M2. During each subculture, the leaves were excised and the new shoots isolated. For polyploidy screening, leaves were excised from all *J. curcas* plantlets (samples; Table 2) recovered in medium M2. *J. curcas* plants seed-raised in greenhouse were used as diploid standards (control). Nuclei suspensions of controls and samples were prepared and analyzed by FCM, as described above.

Polyploidization efficiency was statistically analyzed by the *t* test method, using the Assistat 7.6 beta statistical software (Silva 2012). Based on statistical comparisons, a second in vitro polyploidization procedure was performed using the most efficient colchicine concentration (Table 2).

Results

2C-value and karyotype of J. curcas

The fluorescence peaks of G_0/G_1 nuclei of *J. curcas* and *S. lycopersicum* showed coefficient of variation (CV) lower than 5 %. With the fluorescence peaks of G_0/G_1 nuclei from the standard *S. lycopersicum* (2C = 2.00 pg; Praça-Fontes et al. 2011b) being turned to channel 200 (data not shown), the mean 2C value for all *J. curcas* 'Gonçalo' plants was equivalent to 0.85 pg \pm 0.01 (0.83 \times 10⁹ bp).

Jatropha curcas root tips treated with 4 μ M amiprophos-methyl (APM, microtubule inhibitor) and macerated in 1:10 pectinase solution for 2.5 h provided metaphases adequate for morphometric analysis. The metaphases presented well-spread chromosomes with well-defined constrictions, without cytoplasmic background and chromatin damage (data not shown). Morphometric analysis showed that J. curcas 'Gonçalo' has 2n = 22 chromosomes, with total length ranging from 1.60 to 1.14 μ m.

In vitro recovery of J. curcas shoot tips

Owing to the decontamination procedure and handling of seeds under aseptic conditions, no contamination was detected during the in vitro process. After 8 days in medium M1 (Table 1), *J. curcas* seeds germinated with a rate of 86.66 % and speed of germination (SG) of 4.86. Consequently, plantlets were regenerated after 30 days, providing enough shoot tips for propagation and multiplication in medium M2 and in vitro polyploidization in medium M3.

In vitro multiplication and assessment of DNA ploidy level

Shoot tips were transferred to medium M2 and cultivated for 120 days, with monthly subcultures. After four subcultures in M2, the shoot tips showed a mean multiplication rate of 4.5 shoots per explant.

Leaves from the plantlets were collected for analysis of DNA ploidy level stability by FCM. In vitro (samples) and greenhouse (control; 2C = 0.85 pg; $2 \times = 22$ chromosomes) plantlets exhibited G_0/G_1 peaks in the same channel (Fig. 1a). This way, it could be verified that numeric chromosomal aberration (eu- and/or aneuploidy) has not occurred during propagation and multiplication of *J. curcas* plantlets in medium M2.

In vitro polyploidization

In vitro polyploidization was conducted in two approaches. The first was performed to select the most adequate

Table 2 Survival rate, and number of mixoploid and tetraploid J. curcas plantlets obtained after polyploidization treatments

Colchicine concentration (mM)	Treatment duration (h)	No of individuals treated	Survival rate (%)	No of regenerated plantlets*	No diploids (%)***	No tetraploids (%)***	No mixoploids (%)***
First polyploidization	approach						
0	24	9	100 ^a	27 ^a	27 (100)	0 ^b (0)	0 ^b (0)
	48	9	100 ^a	25 ^{a,b}	25 (100)	0 ^b (0)	0 ^b (0)
	72	9	100 ^a	25 ^{a,b}	25 (100)	0 ^b (0)	0 ^b (0)
	96	9	100 ^a	26 ^{a,b}	26 (100)	0 ^b (0)	0 ^b (0)
0.5	24	9	100 ^a	24 ^{a,b}	24 (100)	0 ^b (0)	0 ^b (0)
	48	9	66.66 ^b	22 ^{b,c}	22 (100)	$0^{b}(0)$	$0^{b}(0)$
	72	9	22.22 ^d	10 ^e	9 (90)	0 ^b (0)	1 ^b (10)
	96	9	66.66 ^b	21 ^c	12 (57.15)	3 ^a (14.28)	6 ^a (28.57)
1.5	24	9	66.66 ^b	12 ^{d,e}	12 (100)	$0^{b}(0)$	$0^{b}(0)$
	48	9	66.66 ^b	12 ^{d,e}	10 (83.34)	2 ^a (16.66)	$0^{b}(0)$
	72	9	33.33 ^c	15 ^d	15 (100)	$0^{b}(0)$	$0^{b}(0)$
	96	9	0.0^{d}	0^{f}	0 (0)	0 ^b (0)	0 ^b (0)
2.5	24-96**	9	0.0^{d}	0^{f}	0 (0)	$0^{b}(0)$	$0^{b}(0)$
Second polyploidization	on approach						
0	96	20	100 ^a	20^{a}	20 (100)	0 ^b (0)	0 ^b (0)
	120	20	100 ^a	20^{a}	20 (100)	0 ^b (0)	0 ^b (0)
	144	20	100 ^a	20^{a}	20 (100)	0 ^b (0)	0 ^b (0)
	168	20	100 ^a	20^{a}	20 (100)	$0^{b}(0)$	$0^{b}(0)$
0.5	96	20	65 ^b	13 ^b	0	7 ^a (53.85)	6 ^a (46.15)
	120	20	35 ^c	7 [°]	3 (42.85)	3 ^b (42.85)	1 ^b (14.3)
	144	20	35 ^c	7 ^c	5 (71.42)	0 ^b (0)	2 ^b (28.58)
	168	20	20°	$4^{\rm c}$	4 (100)	0 ^b (0)	0 ^b (0)

In the columns for each polyploidization approach, means followed by the same letter are not significantly different at the $p \le 0.05$ by t test * Plantlets regenerated after four subsequent subcultures totalizing 120 days

** As all explants treated with 2.5 mM colchicine concentration died, the data were summarized

*** Number (%) of diploid, tetraploid, and mixoploid plantlets

colchicine concentration. For this purpose, some criteria were adopted: (a) survival rate of the explants, and (b) number of tetraploid and (c) mixoploid plantlets.

The survival rate of explants not treated with the antimitotic agent colchicine, or treated with 0.5 mM for 24 h, was 100 %. With increasing concentration, the survival rate statistically decreased, reaching 0 % when the explants were treated with a colchicine concentration of 1.5 mM for 72 h (Table 2).

Shoot tips treated with colchicine showed a lower growth and multiplication rate, yielding few regenerated plantlets in comparison to explants not treated with colchicine. In consideration of the vegetative development, DNA ploidy level of the plantlets was assessed by FCM only after 4 subcultures in medium M2, totaling 120 days.

Jatropha curcas plantlets cultured in medium M3 without colchicine, as well as greenhouse plants (control; 2C = 0.85 pg; $2 \times = 22$ chromosomes), exhibited G₀/G₁ peaks in the same channel (Fig. 1a). Therefore, these

plantlets (Fig. 1b) showed the same DNA ploidy level. Similarly, some explants treated with colchicine generated seedlings that provided FCM histograms with the same profile in comparison to the control (Table 2; Fig. 1a). Based on this, these plantlets were also considered diploid.

Mixoploid plantlets of *J. curcas* (Fig. 1c, d) were obtained in colchicine concentration of 0.5 mM for 72 h (10.00 %) and 0.5 mM for 96 h (28.57 %). These plantlets exhibited G_0/G_1 peaks equivalent to nuclei 2C = 2X and 2C = 4X (Fig. 1c).

FCM analysis also evidenced tetraploid plantlets of *J. curcas* (Fig. 1e, f). In comparison to control and plantlets cultured in medium M3 without colchicine, these plantlets exhibited G_0/G_1 peaks equivalent to nuclei 2C = 4X (Fig. 1f). Tetraploids were observed in the treatment with 0.5 mM colchicine for 96 h (14.28 %), and 1.5 mM with exposure time of 48 h (16.66 %) (Table 2).

In regard of the three criteria (number of tetraploid and mixoploid plantlets, and survival rate), the treatment with



Fig. 1 *J. curcas* plantlets obtained by in vitro polyploidization, and cultivated in M2 medium (Table 1), and respective FCM *histograms*. **a** *Histogram* showing G_0/G_1 *peak* (channel 200) of the diploid samples (2C = 2X). **b** Diploid plantlet representing the regenerants that showed the same DNA ploidy level of the control plants (2C = 2X). **c** *Histogram* showing G_1/G_0 *peak* (channel 200 and 400)

0.5 mM colchicine for 96 h was considered the most adequate for polyploidy induction in *J. curcas* shoot tips. Considering this result, a second polyploidization approach was set (Table 2), increasing the treatment time (96, 120, 144, 168 h) and maintaining the colchicine concentration (0.5 mM).

As in the first polyploidization procedure, approximately 65 % of the explants survived in the 0.5 mM/96 h

of the mixoploid samples (2C = 2X and 2C = 4X, respectively). **d** Mixoploid plantlet representing the regenerants that showed DNA ploidy level 2C = 2X and 2C = 4X. **e** *Histogram* showing G_1/G_0 *peak* (channel 400) of the tetraploid samples (2C = 4X). **f** Tetraploid plantlet representing the regenerants that showed DNA ploidy level 2C = 4X. *Bar* 1 cm

treatment. Survival rate statistically decreased with the increase in exposure time (Table 2). The second polyploidization experiment confirmed that the 0.5 mM/96 h treatment is efficient for the production of tetraploids (Table 2). Analysis of DNA ploidy level by FCM evidenced that this treatment provided 53.85 % of tetraploids. This treatment also generated 46.15 % of mixoploid regenerants. Tetraploids were also obtained in the exposure time of 120 h, though with smaller percentage (42.85 %) (Table 2). The mixoploid and tetraploid plantlets have been in vitro multiplied, and some ex vitro acclimatized.

Discussion

2C-value and karyotype of J. curcas

Considering that some authors have reported the occurrence of *J. curcas* tetraploid plants (Soontornchainaksaeng and Jenjittikul 2003; Mukherjee et al. 2011), the first concern was to measure the 2C-value and characterize the karyotype of *J. curcas* 'Gonçalo' explant donors.

As reported by Carvalho et al. (2008), the FCM procedure provided fluorescence peaks of G_0/G_1 nuclei showing CVs below 5 %. This value has been considered adequate for FCM assessments in plants (Praça-Fontes et al. 2011a). In accordance with Praça-Fontes et al. (2011a), an appropriate preparation of nuclei suspensions is imperative to provide stoichiometrically stained nuclei and, consequently, low CV values, as obtained in the present work.

Solanum lycopersicum 'Stupické' was chosen as primary standard for 2C-value measurement. This standard has been regarded as suitable for analyses involving plants rich in phenolic compounds (Clarindo et al. 2012), such as *J. curcas*. Using FCM, Praça-Fontes et al. (2011a) revisited the DNA C-value of seven species often used as primary standard. In a cascade-like manner, from *Arabidopsis thaliana* to *Allium cepa*, these authors demonstrated that *S. lycopersicum* is an ideal primary reference standard.

The mean nuclear DNA content value of 2C = 0.85 pg found here for *J. curcas* was identical to the value reported by Carvalho et al. (2008). This result confirms that the nuclear genome size of *J. curcas*, as well as that of *Euphorbia peplus* L. (2C = 0.7 pg), is relatively small compared to other Euphorbiaceae species (Bennett and Leitch 2011). Corroborating with the FCM data, karyotype analysis showed that *J. curcas* 'Gonçalo' had $2n = 2 \times = 22$ chromosomes, which is considered relatively small (Carvalho et al. 2008).

The previous FCM and cytogenetic approaches showed that *J. curcas* 'Gonçalo' has a stable genome $(2 \times = 22$ chromosomes, 2C = 0.85 pg). Since no pre-existing ploidy variation was evidenced, this line was considered adequate for in vitro polyploidy induction.

In vitro recovery of J. curcas shoot tips

The decontamination process was efficient to provide aseptic cultures of *J. curcas*. Seeds of this species have a high incidence of fungi (e.g. *Alternaria* sp., *Macrophomina* sp., *Cladosporium* sp., *Fusarium* spp.), even after a surface

decontamination process (Vanzolini et al. 2010). Considering this, the removal of the pericarp, followed by decontamination in laminar flow, was crucial to ensure totally aseptic seeds.

As attested by the high germination rate (86.66 %), medium M1 yielded sufficient amount of seedlings for subsequent excision of shoot tips. Distinct authors (Kalimuthu et al. 2007; Kaewpoo and Te-chato 2010) have also considered M1 an efficient medium for germination of zygotic embryos and seeds of *J. curcas*.

Besides the high germination rate, *J. curcas* seeds showed a SG of 4.86 after 8 days in medium M1. This data reflects the efficiency of medium M1 to promote healthy seedlings. Gairola et al. (2011) reported a germination rate of 62.50 % and a SG of 4.85 in *J. curcas* seeds sown in vermiculite substrate. Thus, the in vitro conditions were adequate for seed germination of this species.

In vitro multiplication and assessment of DNA ploidy level

In accordance with Rajore and Batra (2005), medium M2 was ideal for clonal propagation and multiplication of *J. curcas* shoot tips. After four subcultures (120 days) in M2, the mean multiplication rate was 4.5 shoots per explant. This result indicates that the BAP/IAA combination in medium M2 increased shoot frequency, being crucial for the success of in vitro propagation and multiplication of *J. curcas*.

The type and concentration of growth regulators, as well as subculture frequency, can promote the occurrence of numeric chromosomal aberrations (eu- and/or aneuploidy) associated to somaclonal variation. Auxins and cytokinins, such as BAP and IAA, are the main growth regulators that act to control cell division and tissue differentiation (Fehér et al. 2003). These regulators interfere in cell cycle control and may lead to genetic variability (Bairu et al. 2011).

As the medium M2 was supplemented with auxin and cytokinin regulators, leaves from plantlets propagated in this medium were used to assess DNA ploidy level. The nuclei suspension of these plantlets provided G_0/G_1 peaks in the same channel as the control plants (Fig. 1a). In this sense, medium M2 (Rajore and Batra 2005) did not promote any numeric chromosomal aberration (eu- and/or aneuploidy) during propagation and multiplication of *J. curcas* plantlets. Therefore, the in vitro tissue culture conditions conserved the nuclear genome stability and homogeneity of propagated and multiplied *J. curcas* plantlets up to the fourth subculture. As related by Fiuk et al. (2010), ploidy maintenance during in vitro condition is a relevant prerequisite for clonal propagation.

Kour et al. (2009) mentioned that variability can manifest at cytological level. For this reason, FCM is commonly used for detection of DNA ploidy level variation (Clarindo et al. 2008). Besides, FCM analysis is the most practical, reliable and efficient method for this purpose (Clarindo et al. 2008; Praça et al. 2009). Therefore, DNA ploidy level screening of *J. curcas* plantlet samples could be done rapidly, in a working day, without the requirement of dividing cells.

In vitro polyploidization

The polyploidization experiments were conducted to find colchicine concentrations with limited toxic effect. The effect of colchicine on the survival rate depended upon concentrations of this compound and duration of treatment (Samala and Te-chato 2012).

Previous experiments performed in our laboratory showed that colchicine concentrations of 3.5, 5.0 and 6.5 mM at treatment periods of 24–72 h were highly toxic to shoot tips (data not shown). These explants died in the first week in medium M2. Thereafter, other concentrations and exposure times were tested (Table 2) so as to obtain surviving explants, as well as tetraploid and mixoploid plantlets.

Considering the three adopted criteria, treatment with 0.5 mM colchicine for 96 h was considered the most adequate for polyploidization (Table 2). Based on this result, a new polyploidization approach was performed, in which the colchicine concentration was maintained (Table 2).

The survival rate of *J. curcas* explants decreased with increasing colchicine concentration and exposure time (Table 2). This result was also found in other polyploidization approaches from woody plants (Gu et al. 2005; Stanys et al. 2006; Xi-Ling et al. 2011; Samala and Te-chato 2012).

Tetraploid plantlets were obtained from different treatments, with 0.5 mM colchicine for 96 h being statistically the most efficient (Table 2). From this treatment, 53.85 % of *J. curcas* tetraploid plantlets were recovered (Fig. 1f; Table 2). Tetraploid plantlets of the woody species *Phlox subulata* L. were regenerated by Zhang et al. (2008) also using 0.5 mM colchicine.

Most of the mixoploid plantlets were also obtained in the treatment with 0.5 mM colchicine for 96 h (Table 2). Though in vitro polyploidization generally yields mixoploid plantlets (Chen and Gao 2007), the mixoploid state has been considered reversible, since some plants return to the diploid condition or become tetraploid plants (Väinölä 2000). Due to the simultaneous occurrence of cells with varying ploidy, embryogenesis systems can be established to enable recovery of plantlets showing a single and stable ploidy level (Chen and Gao 2007).

In this study, a tissue culture procedure was adapted for induction, propagation and recovery of tetraploid plantlets of *J. curcas*. Moreover, nuclear DNA ploidy screening by FCM was a practical and rapid strategy for selection of diploid, mixoploid and tetraploid plantlets. The method presented here is also reliable for routine in vitro production of polyploids of other *J. curcas* elite lines.

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