

***Agrobacterium*-MEDIATED TRANSFORMATION OF *Citrus sinensis* AND *Citrus limonia* EPICOTYL SEGMENTS**

Weliton Antonio Bastos de Almeida^{1,4}; Francisco de Assis Alves Mourão Filho^{1,5*}; Beatriz Madalena Januzzi Mendes^{2,5}; Alexandra Pavan^{3,6}; Adriana Pinheiro Martinelli Rodriguez²

¹ Depto. de Produção Vegetal - USP/ESALQ, C.P. 9 - CEP: 13418-900 - Piracicaba, SP.

² Lab. de Biotecnologia Vegetal - USP/CENA, C.P. 96 - CEP: 13400-970 - Piracicaba, SP.

³ UNIMEP - Rod. do Açúcar, Km. 156 - CEP: 13400-911 - Piracicaba, SP.

⁴ CAPES Fellow.

⁵ CNPq Fellow.

⁶ FUNDECITRUS Fellow.

*Corresponding author <famourao@esalq.usp.br>

ABSTRACT: Genetic transformation allows the release of improved cultivars with desirable characteristics in a shorter period of time and therefore may be useful in citrus breeding programs. The objective of this research was to establish a protocol for genetic transformation of Valencia and Natal sweet oranges (*Citrus sinensis* L. Osbeck) and Rangpur lime (*Citrus limonia* L. Osbeck). Epicotyl segments of germinated *in vitro* plantlets (three weeks in darkness and two weeks in a 16-h photoperiod) were used as explants. These were co-cultivated with *Agrobacterium tumefaciens* strain EHA-105 and different experiments were done to evaluate the transformation efficiency: explants were co-cultivated with *Agrobacterium* for one, three or five days; explants were incubated with *Agrobacterium* suspension for 5, 10, 20 or 40 minutes; co-cultivation medium was supplemented with acetosyringone at 0, 100 or 200 $\mu\text{mol L}^{-1}$; Explants ends had a longitudinal terminal incision (2-3 mm); co-cultivation temperatures of 19, 23 or 27°C were imposed. The experimental design was completely randomized in all experiments with five replications, each consisted of a Petri dish (100 x 15 mm) with 30 explants and resulted in a total of 150 explants per treatment. Longitudinal terminal incision in the explant ends did not improve shoot regeneration. However, transgenic plants of all three cultivars were confirmed from explants that had been subjected to inoculation time of 20 minutes, co-culture of three days at 23-27°C, in the absence of acetosyringone.

Key words: genetic transformation, organogenesis, micrografting, improvement

TRANSFORMAÇÃO GENÉTICA EM *Citrus sinensis* E *Citrus limonia* MEDIADA POR *Agrobacterium tumefaciens* A PARTIR DE SEGMENTOS DE EPICÓTILO

RESUMO: A transformação genética permite produzir cultivares com características específicas e pode, dessa forma ser associada a programas de melhoramento de citros. O objetivo deste trabalho foi estabelecer protocolos de transformação genética para as laranjas doce 'Valência' e 'Natal' (*Citrus sinensis* L. Osbeck), bem como para o limão 'Cravo' (*Citrus limonia* L. Osbeck). Segmentos de epicótilo de plântulas germinadas *in vitro* (três semanas no escuro e duas semanas sob fotoperíodo de 16h) foram utilizados como explantes. Estes foram co-cultivados com *Agrobacterium tumefaciens* (EHA-105), realizando-se vários experimentos para avaliar a eficiência do processo de transformação genética: explantes co-cultivados por um, três e cinco dias; tempo de inoculação com a bactéria de 5, 10, 20 e 40 minutos; co-cultivo em meio de cultura contendo 0, 100 e 200 $\mu\text{mol L}^{-1}$ de acetosiringona; Incisão longitudinal (2-3 mm) nas extremidades do explante; temperatura de co-cultivo 19, 23 e 27°C. Todos os experimentos consistiram de cinco repetições por tratamento, sendo cada repetição representada por uma placa de Petri contendo 30 explantes, perfazendo um total de 150 explantes por tratamento. Plântulas transgênicas dos três cultivares foram obtidas utilizando-se tempo de inoculação de 20 minutos, co-cultivo com *Agrobacterium tumefaciens* (EHA-105) por três dias, na ausência de acetosiringona no meio de cultura de co-cultivo e temperatura de co-cultivo de 23-27°C. A incisão longitudinal na extremidade do explante favoreceu à organogênese *in vitro*, mas quando co-cultivado com *Agrobacterium* não houve regeneração de brotações.

Palavras-chave: transformação genética, organogênese, microenxertia, melhoramento

INTRODUCTION

The genus *Citrus* presents limitations for improvement via conventional breeding, and are directly associated with the reproductive biology of *Citrus*, such as nucellar polyembryony, a high level of heterozygosity and a long juvenile period (Grosser &

Gmitter, 1990). Biotechnological techniques such as cell and tissue cultures and molecular biology have helped breeders to overcome these difficulties. In addition, the hybridization through protoplast fusion and genetic transformation may contribute significantly to avoid these limitations (Mendes-da-Glória et al., 2000).

Genetic transformation has become an attractive alternative method for improving *Citrus* and other species because it is possible to maintain cultivar integrity while adding a single trait (Bond & Roose, 1998). However, for efficient transgenic plant production, a previously defined tissue culture system for plant regeneration (Brasileiro & Dusi, 1999) associated with a genetic transformation system for gene introduction is needed (Pérez-Molphe-Balch & Ochoa-Alejo, 1998). For *Citrus*, there are many *in vitro* protocols such as callus and cell suspension cultures (Cabasson et al., 1995), organogenesis induction (Pérez-Molphe-Balch & Ochoa-Alejo, 1997), somatic embryogenesis induction (Tomaz et al., 2001), and protoplast isolation (Mendes et al., 2001) that are viable for use with genetic engineering protocols.

Citrus transformation was initially reported more than a decade ago (Kobayashi & Uchimaya, 1989; Hidaka et al., 1990; Vardi et al., 1990; Moore et al., 1992). However, the success in the recovery of transgenic plants at that time was relatively inefficient due to many factors. Low transformation efficiency (Peña et al., 1995a), the growth of escapes even under selection (Yang et al., 2000), difficulties in rooting of transgenic shoots and genotypic influence (Gutiérrez-E. et al., 1997) are among the limiting factors for the development of transgenic citrus plants.

GUS and *nptII* have been used as reporter and selection genes, respectively on *Citrus* genetic transformation. There are a few reports about the introduction of genes of agronomic importance in *Citrus*, including the gene that encodes for the *Citrus* tristeza virus coat protein (Gutiérrez-E. et al., 1997; Domínguez et al., 2000; Yang et al., 2000), the *HAL2* gene that confers tolerance to salinity (Cervera et al., 2000), *LEAFY* and *APETALA1* genes that promote early flower initiation (Peña et al., 2001), and *CS-ACS1* gene that controls the ethylene biosynthesis in *Citrus* (Wong et al., 2001). Reports on genetic transformation of the main cultivars used in the Brazilian citrus industry are very few (Mendes et al., 2002) and there are no reports of genetic transformation of the mostly used rootstock in Brazil, the Rangpur lime.

The goal of the present work is to establish efficient protocols for genetic transformation of two sweet orange varieties Natal and Valencia and the rootstock, Rangpur lime.

MATERIAL AND METHODS

Plant material: Seeds were extracted from ripe fruits of Valencia and Natal sweet oranges (*Citrus sinensis* L. Osbeck) and Rangpur lime (*Citrus limonia* L. Osbeck) and dried at room temperature for 24-h. The seed integument was removed and desinfestation was done in a 67% commercial sodium hypochlorite solution (2.5% active chlorine) for 15 minutes followed by three rinses in

distilled and sterilized water. The seeds were placed in test tubes (150 x 25 mm) containing 15 mL of MT medium (Murashige & Tucker, 1969) and maintained at $27 \pm 2^\circ\text{C}$ in the dark for three weeks followed by one week in a 16-hour photoperiod ($40 \mu\text{mol m}^{-2}\text{s}^{-1}$). Epicotyl segments approximately 1.0-cm-long were collected for the transformation experiments.

Agrobacterium strain: *Agrobacterium tumefaciens* strain EHA-105 carrying the plasmid p35SGUSINT was used. The bacteria were cultivated in solid YEP medium (10 g L⁻¹ peptone, 10 g L⁻¹ yeast extract, 5 g L⁻¹ sodium chloride, 15 g L⁻¹ agar) containing kanamycin (100 mg L⁻¹) and rifampicin (50 mg L⁻¹), for 48-h. After that, a single colony was transferred to a 250 mL Erlenmeyer flask with 50 mL of liquid YEP medium, supplemented with antibiotics and cultivated at 180 rpm at 28°C for 16-h. The bacterial suspension was centrifuged at 4800 rpm (5°C/20 min) and resuspended in liquid MS medium (Murashige & Skoog, 1962).

Transformation, selection, and regeneration: Epicotyl segments were incubated with the bacteria solution for 20 minutes. Following incubation, explants were blotted dry and plated on regeneration EME medium (Grosser & Gmitter, 1990) supplemented with 25 g L⁻¹ sucrose and 1.0 and 2.0 mg L⁻¹ benzylaminopurine (BAP) for sweet orange cultivars and for Rangpur lime, respectively, in the dark at 27°C for a 3-day period. After co-culture, segments were transferred to regeneration medium, supplemented with kanamycin (100 mg L⁻¹) and cefotaxime (500 mg L⁻¹). Explants were subcultured every 2 weeks. Developed shoots were transferred to EME medium supplemented with GA₃ (1.0 mg L⁻¹), kanamycin (100 mg L⁻¹) and cefotaxime (500 mg L⁻¹) for elongation. Shoot basal ends and leaf segments were assayed for GUS activity. GUS⁺ shoots were micrografted onto Carrizo citrange (*C. sinensis* L. Osbeck x *Poncirus trifoliata* L. Raf.) seedlings. After 45 days, well-developed *in vitro* grafted plantlets were transferred to a commercial substrate (Rendmax Citrus™) and kept under high relative humidity for 30 days for acclimatization. In order to investigate the main factors affecting the transformation efficiency, different treatments were performed such as: 1) Co-cultivation of epicotyl segments with *Agrobacterium* for 1, 3 or 5 days; 2) Explant incubation with *Agrobacterium* for 5, 10, 20 or 40 minutes; 3) Addition of acetosyringone at 0, 100 or 200 $\mu\text{mol L}^{-1}$ in the co-cultivation medium; 4) Longitudinal terminal incision (2-3 mm) in the explant; 5) Co-cultivation temperatures of 19, 23 or 27°C.

The experimental design was completely randomized, with five replications, each consisting of one Petri dish (100 x 15 mm) with 30 explants resulting in a total of 150 explants per treatment for all experiments. Percent of explants with adventitious shoots were evaluated. Data were subjected to ANOVA at 1%. Means were compared by Tukey's multiple range test.

Analysis of putatively transformed tissue: leaves and stem segments were excised from 1-2-cm plants. Segments were incubated in the dark at 37°C for 24-h in an X-Gluc solution (Jefferson, 1987) for β -glucuronidase assay (GUS). For histological GUS assay, stem segments were fixed in paraformaldehyde (3% w/v) and glutaraldehyde (2% v/v) in cacodylate buffer (0.2 M, pH 7.2) with the first hour being in low vacuum. Fixed tissues were rinsed in buffer and slowly dehydrated at room temperature in a series of methyl cellosolve, ethanol, propanol and butanol followed by overnight infiltration at 4°C in butanol:infiltration medium (Historesin™/Leica) (1:1). Infiltration was completed with 100% infiltration medium for 24-h or until the samples sank to the bottom of the flask. Polymerization was done at room temperature for 24 to 48-h. Transverse serial sections (5 mol L⁻¹) were prepared in a rotary microtome (Leica RM 2155) with a steel knife, the sections floated in water drops and dried on a hot plate (40°C).

For the detection of specific T-DNA sequences, DNA was extracted from leaves of *in vitro* plantlets (Doyle & Doyle, 1990). PCR amplification was performed using 50-100 ng of DNA, 200 μ mol L⁻¹ of dNTP, 2.5 mM MgCl₂, 2U taq DNA polymerase (Promega) and 0.1 mol L⁻¹ *nptII* primer. The primers 5'GAG GCT ATT CGG CTA TGA CTG'3 and 5'ATC GGG AGC GGC GAT ACC GAT A'3 (Bond & Roose, 1998) were used to amplify a 700 bp specific fragment of *nptII* gene. Samples were heated to 94°C for 4 min followed by 30 cycles of 2 min at 96°C, 2 min at 50°C and 3 min at 72°C.

RESULTS AND DISCUSSION

Citrus genetic transformation has been preferentially obtained from explants collected from juvenile tissue. *Agrobacterium* has been the most frequently used genetic transformation system in *Citrus* with explants collected from seedlings germinated *in vitro* or under greenhouse conditions. Some authors used as explants, epicotyl segments excised from seedlings germinated in the dark for 3-6 weeks (Kaneyoshi et al., 1994; Bonde & Roose, 1998; Luth & Moore, 1999; Yang et al., 2000) and then transferred to a 16 h photoperiod

that varied from 1 to 3 weeks (Cervera et al., 1998a; Peña et al., 1995a). Explants have been collected from seedlings germinated under light conditions for 2-4 months (Moore et al., 1992; Pérez-Molphe-Balch & Ochoa-Alejo, 1998). Other authors have used internodal segments excised from seedlings cultivated in the greenhouse for 6-12 weeks (Peña et al., 1995b; 1997; Dominguez et al., 2000). In our work, the explants used were epicotyl segments excised from seedlings germinated in the dark for 3 weeks followed by 10-15 days under in a 16-h photoperiod. This explant type was chosen due to the high efficiency in shoot regeneration and efficient control of contamination as observed in an earlier study (Almeida et al., 2002).

The co-culture period has been reported as an important factor that may influence the genetic transformation process. Our results indicate that GUS⁺ shoots were recovered only when co-cultivation was done for 3 days (Table 1). An overgrowth of the bacteria after five days of co-cultivation was detected similar to that reported by Cervera et al. (1998b).

Another important factor in the genetic transformation efficiency is the period of inoculation of explants with *Agrobacterium*. In this work, a 20-min incubation time resulted in higher number of GUS⁺ shoot recovered for all varieties (Table 2). Yang et al. (2000) also used 20 minutes of inoculation but did not compare different periods of inoculation.

Several authors have reported the supplementation of the co-culture medium with substances that stimulate the infection with *Agrobacterium*. Acetosyringone has been the most used for this propose at a concentration of 100 μ mol L⁻¹. In our study, acetosyringone at either 100 or 200 μ mol L⁻¹ did not favor genetic transformation and the best results were observed in the absence of acetosyringone (Table 3). The liberation of phenolic compounds by the explants was probably enough to favor *Agrobacterium* infection. Therefore, the exogenous supply of acetosyringone when combined with endogenous phenolic compounds may have contributed negatively to the infection. Similar results were observed by Peña et al. (1995a) in Carrizo citrange when GUS⁺ shoots were regenerated in the

Table 1 - Co-culture period of epicotyl segments with *Agrobacterium* (time of inoculation = 15 minutes at 27°C) versus bud differentiation and regeneration of GUS⁺ plants of Natal and Valencia sweet oranges and Rangpur lime.

Co-culture period	Varieties					
	Natal		Valencia		Rangpur	
	N° expl. with buds/total expl.	GUS ⁺ /analyzed shoots	N° expl. with buds/total expl.	GUS ⁺ /analyzed shoots	N° expl. with buds/total expl.	GUS ⁺ /analyzed shoots
days	----- % -----					
1	21/150(14.0) c	00/16(0.0)	26/150(17.3) b	00/11(0.0)	16/150(10.6) b	00/07(0.0)
3	52/150(34.6) a	02/37(5.4)	59/150(39.3) a	01/21(4.7)	34/150(22.6) a	00/19(0.0)
5	32/150(21.3) b	00/24(0.0)	33/150(22.0) b	00/19(0.0)	15/150(10.0) b	00/09(0.0)

Values for percent followed by the same letter do not differ (Tukey, 0.01).

absence of acetosyringone in the culture medium. However, other authors report transgenic plant recovery using acetosyringone at 100 $\mu\text{mol L}^{-1}$ in Pineapple sweet orange (*Citrus sinensis* L. Osbeck) (Cervera et al., 1998a) and Duncan grapefruit (*Citrus paradisi*) (Luth & Moore, 1999) and 200 $\mu\text{mol L}^{-1}$ in Washington navel sweet orange (*Citrus sinensis* L. Osbeck) (Bond & Roose, 1998). The response to acetosyringone could possibly be genotype-dependent.

Formation of meristematic regions or cell proliferation can be induced by cuts in the explant (Escudero & Hohn, 1997). Anatomical characterization of organogenesis from epicotyl segments indicated that meristematic regions formed in the cambium were responsible for the formation of buds (data submitted for publication). With the goal of exposing the cambial region for a better contact with *Agrobacterium*, the explant ends

were cut longitudinally (2-3 mm). This cut significantly favored the *in vitro* organogenesis and induced more buds per responsive explant when compared to the control (Table 4, Figures 1a-b). However, when the cut explants were co-cultivated with *Agrobacterium*, an overgrowth of bacteria did not allow shoot regeneration (Figure 1c). The contact of *Agrobacterium* with a larger number of exposed cells at the concentration used in this experiment (5×10^8 UFC mL^{-1}) may have been too high, thereby favoring the overgrowth of bacteria and restricting shoot regeneration. A study with different concentrations of *Agrobacterium* may increase the transformation efficiency.

The temperature during co-cultivation had a significant effect on the stimulation of the *vir* region of the bacterial plasmid. This effect has also been reported in genetic transformation of several species where the best

Table 2 - Inoculation time of epicotyl segments with *Agrobacterium* (period of co-culture = 3 days at 27°C) versus bud differentiation and regeneration of GUS⁺ plants of Natal and Valencia sweet oranges and Rangpur lime.

Inoculation time	Varieties					
	Natal		Valencia		Rangpur	
	N° expl. with buds/total expl.	GUS ⁺ /analyzed shoots	N° expl. with buds/total expl.	GUS ⁺ /analyzed shoots	N° expl. with buds/total expl.	GUS ⁺ /analyzed shoots
minutes	----- % -----					
5	16/150(10.6) c	00/09(0.0)	14/150(9.3) d	00/06(0.0)	05/150(3.3) c	00/03(0.0)
10	32/150(21.3) b	00/23(0.0)	28/150(18.6) c	00/19(0.0)	22/150(14.6) b	00/15(0.0)
20	98/150 (65.3) a	10/86(11.6)	79/150(52.6) a	09/68(13.2)	53/150(35.3) a	00/36(0.0)
40	42/150(28.0) b	00/29(0.0)	46/150(30.6) b	00/31(0.0)	27/150(18.0) b	00/19(0.0)

Values for percent followed by the same letter do not differ (Tukey, 0.01).

Table 3 - Acetosyringone concentration in the culture medium at co-culture (time of inoculation = 20 minutes at 27°C, and period of co-culture = 3 days) versus bud differentiation and regeneration of GUS⁺ plants of Natal and Valencia sweet oranges and Rangpur lime.

Acetosyringone concentration	Varieties					
	Natal		Valencia		Rangpur	
	N° expl. with buds/total expl.	GUS ⁺ /analyzed shoots	N° expl. with buds/total expl.	GUS ⁺ /analyzed shoots	N° expl. with buds/total expl.	GUS ⁺ /analyzed shoots
μM	----- % -----					
0	94/150(62.6) a	9/84(10.7)	83/150(55.3) a	09/65(13.8)	50/150(33.3) a	00/33(0.0)
100	89/150(59.3) a	00/45(0.0)	68/150(45.3) b	00/42(0.0)	49/150(32.6) a	00/32(0.0)
200	93/150(62.0) a	00/38(0.0)	72/150(48.0) ab	00/47(0.0)	41/150(27.3) a	00/29(0.0)

Values for percent followed by the same letter do not differ (Tukey, 0.01).

Table 4 - Effect of the longitudinal cut (2-3 mm) in the explant ends on *in vitro* organogenesis of Natal and Valencia sweet oranges and Rangpur lime.

Treatment	Varieties					
	Natal		Valencia		Rangpur	
	Expl. with buds	Buds/expl.	Expl. with buds	Buds/expl.	Expl. with buds	Buds/expl.
	%	mean	%	mean	%	mean
Control	92.6 a	286/139 (2.1) b	88.0 a	272/132 (2.1) b	77.0 a	212/116 (1.8) b
Cut Explants	94.0 a	435/141 (3.1) a	91.0 a	421/137 (3.1) a	82.0 a	321/123 (2.6) a

Values for percent followed by the same letter do not differ (Tukey, 0.01).

temperature was 22°C for *Phaseolus acutifolius* and *Nicotiana tabacum* (Dillen et al., 1997) and was of 22-24°C for *Lycopersicon sculentum* (Costa et al., 2000). Fullner & Nester (1996) studied the T-DNA transfer ability of several *Agrobacterium tumefaciens* strains under different temperatures (15 to 31°C). In that case, the frequency of T-DNA transfer decreased with the increase in temperature suggesting that the T-DNA transfer is sensitive to higher temperatures. For *Citrus* genetic transformation, temperatures between 26-28°C have been used for co-cultivation. In this work, we evaluated the temperatures of 19, 23 and 27°C. The temperature of 23°C induced the best genetic transformation efficiency in Natal sweet orange and the best temperature for Valencia sweet orange genetic transformation was 27°C. For Rangpur lime, GUS⁺ shoots were strongly dependent on the co-cultivation temperature and were obtained only when the co-cultivation temperature was 23°C. However, these results suggest that T-DNA transfer is not only dependent on the temperature of co-cultivation but also upon a combination of different factors such as: temperature, genotype, co-cultivation period and inoculation time.

Genetic transformation in the same *Citrus* species has been considered as recalcitrant due to the low transformation efficiency (Moore et al., 1992; Peña et al., 1995b). Our studies confirm this fact. The highest percents of efficiency (GUS⁺/total explants) were 11.6% (Table 2), 13.8% (Table 3) and 19.3% (Table 5) for Natal

sweet orange, Valencia sweet orange and Rangpur lime, respectively. These numbers are relatively low, especially in sweet orange cultivars, when compared to our previous work with Hamlin orange where 81.5% of transformation efficiency was reported (Mendes et al., 2002).

The occurrence of shoot escapes was high. The regeneration of non-transformed plants can be explained by non-efficient selection due to the protection of non-transformed cells (Ghorbel et al., 1999). In addition, two chimeric plants of Natal sweet orange were obtained (Figures 2c, d). The regeneration of chimeric plants can be associated with the fact that adventitious buds have multicellular origin. Citrus chimeric plants were also described by other authors (Peña et al., 1997; Gutiérrez-E. et al., 1997; Cervera et al., 1998b).

GUS⁺ shoots (Figure 2e, f, g) were *in vitro* grafted (Figure 2h) with 80% of bud-take efficiency. Histological sections (Figure 2g) showed the expression of GUS gene in the stem tissue, from the epidermis to the pith, confirming the GUS gene transfer to the cells responsible for shoot formation. Genetic transformation was also confirmed by PCR analysis with the presence of the 700 bp DNA fragments, corresponding to the *nptII* gene (Figure 2i). After they were grafted, the plants were acclimatized and transferred to the greenhouse (Figure 2j).

Our results confirm that transgenic plants of Natal and Valencia sweet orange and Rangpur lime can be obtained using a co-cultivation period of three days, inoculation time of 20 minutes, absence of

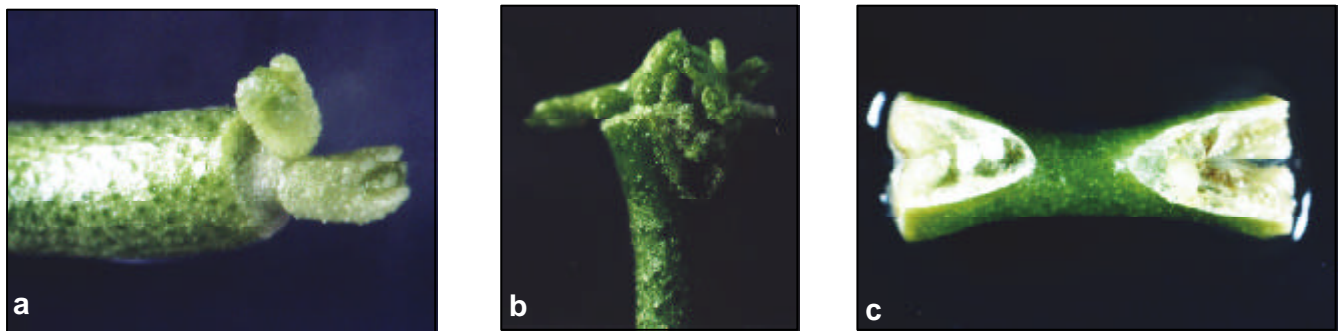


Figure 1 - Influence of explant cut on *in vitro* organogenesis and genetic transformation of Natal and Valencia sweet oranges and Rangpur lime. (a) control; (b) explant end with longitudinal incision; (c) explant cut longitudinally and co-cultivated with *Agrobacterium*.

Table 5 - Temperature of co-culture with *Agrobacterium* (time of inoculation = 20 minutes and period of co-culture = 3 days) versus bud differentiation and regeneration of GUS⁺ plants of Natal and Valencia sweet oranges and Rangpur lime.

Temperature of co-culture	Varieties					
	Natal		Valencia		Rangpur	
	N° expl. with buds/total expl.	GUS ⁺ /analyzed shoots	N° expl. with buds/total expl.	GUS ⁺ /analyzed shoots	N° expl. with buds/total expl.	GUS ⁺ /analyzed shoots
°C	----- % -----					
19	48/150(32.0) b	00/33(0.0)	45/150(30.0) c	00/36(0.0)	27/150(18.0) b	00/21(0.0)
23	52/150(34.6) b	04/35(11.5)	58/150(38.6) b	00/42(0.0)	42/150(28.0) a	06/31(19.3)
27	93/150(62.0) a	7/86(8.1)	74/150(49.3) a	08/61(13.1)	53/150(35.0) a	00/39(0.0)

Values for percent followed by the same letter do not differ (Tukey, 0.01).

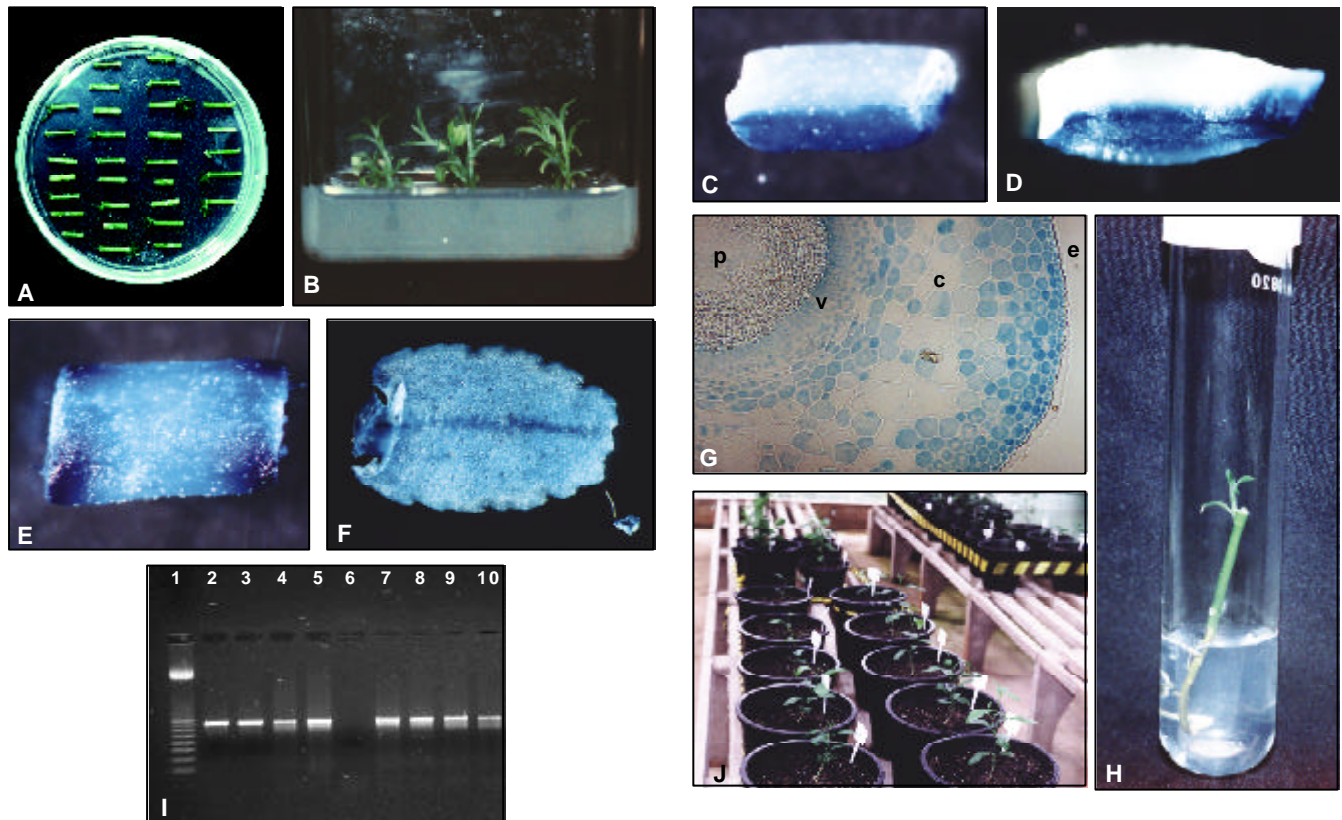


Figure 2 - Genetic transformation of Natal and Valencia sweet oranges and Rangpur lime using epicotyl segments as explants. (a) epicotyl segment after co-cultivation with *Agrobacterium*; (b) regenerated shoots cultivated in elongation medium; (c) stem segment; (d) stem segment showing possible chimeric plants after histochemical GUS assay; (e) stem; (f) and leaf segments; (g) histological section showing GUS⁺ shoots after histochemical and histological GUS assay. e = epidermis; c = cortex; v = vascular cylinder; p = pith; (h) GUS⁺ shoots *in vitro* grafted on Carrizo citrange; (i) analysis of PCR of the GUS⁺ DNA plants. Lane 1 = ladder 100 bp. Lane 2 = positive control consisting of the plasmid. Lane 6 = negative control consisting of non-transformed plant DNA. Lanes 3, 4, 5, 7, 8, 9 and 10 = represent *npdI*⁺ plants; (j) transgenic plants in the greenhouse.

acetosyringone in the co-cultivation medium, and temperature of co-cultivation of 23°C and 27°C, and 23°C, respectively. In addition, the longitudinal cut (2-3 mm) on the explant ends favored *in vitro* organogenesis but did not allow shoot regeneration after being co-cultivated with *Agrobacterium tumefaciens*.

ACKNOWLEDGEMENTS

To Fundação de Amparo à Pesquisa do Estado de São Paulo (FAPESP) and Fundo Paulista de Defesa da Citricultura (Fundecitrus) for financial support.

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Received June 4, 2002