

The Green Fluorescent Protein (GFP) is a Vital Visual Marker in Citrus Transgene Research

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Abstract

pGREEN is a flexible Ti binary vector for *Agrobacterium* mediated transformation system [1]. The advantage of this vector is that optional combinations of selection and reporter gene cassettes can be optimally assembled to meet different transformation requirements. *gfp* (green fluorescent protein) gene driven by 35S-CaMV promoter was used as a reporter gene to enable transformant detection in vivo and the REExtract-N-Amp Plant PCR Kit was applied as a high throughput approach for screening large amount of regenerated transformed plants in this report. With the designated vector pGK assembled using pGreen0029 and 35S-GFP, different phenotypes of transformation chimerism were observed and average transformation rates 7.75% and 2.03 % were acquired in term of consistent PCR results between the selectable and the reporter gene for empty vector and inserted recombinants respectively in transformation events of citrus grapefruits.

Key words: *Agrobacterium*, transformation, citrus, GFP (*green fluorescent protein*), PCR

1. Introduction

Genetic transformation is the direct approach for gene function characterization for any important clone selected from constructed bacterial libraries in genomic projects. Citrus is one of the most recalcitrant woody species to transform in gene engineering. For gene function evaluation for a bunch of gene candidates from genomic libraries, high transformation efficiency is required to acquire large amount of regenerated transformed plants and the high throughput reliable techniques are critical for screening those transformed plants without destructivity. We have compared several widely

used expression vectors in our citrus transformation and excellent progress has been made for obtaining high transformation efficiency while in many cases, transformed plants will be damaged during transgene detection procedure and transgene verification won't be done until other plants are grown enough size for sampling.

The green fluorescent protein (GFP) converts blue light emitted from aequorin in jellyfish to green light. The use of GFP marker protein has been described in bacteria [2], yeast [3], worm [4], fly [5] and mammals [6]. In plants, GFP has also been widely used as a visual marker gene that functions as a reporter of gene expression and permits the recovery of transformed plants [7].

pGREENs are a new set of flexible binary vectors for crop transformation newly developed by Roger Hellens et al. [8] and they have been optimised for the improvement of plant transformation via *Agrobacterium*. In pGREEN, the T-DNA region is flanked by *Bgl*II sites, which are useful in post-transformation analysis. The polylinker is based on pBluescript, and all associated plasmids and cassettes have been modified, to remove restriction sites included in this polylinker. Further unique sites *Hpa*I and *Stu*I are located internal to the Left and Right border sequence respectively. These blunt sites are designed to facilitate the cloning of selection and marker cassettes, which are flanked by blunt *Eco*RV sites. The *npt* II gene of pACYC 177 conveys resistance to kanamycin, and was used to select for plasmid transformation. Site-directed mutagenesis was used to remove the sites represented in the multiple cloning site, and an *Nco*I site was incorporated 3' of the gene, to facilitate convenient cloning. The *RepA* gene acts in trans upon the pSa Ori sequence. The *RepA* is therefore resident on a separate plasmid pSoup. This plasmid can be co-electroporated with pGreen vector, or *Agrobacterium* cells with the plasmid can be made competent, for independent electroporation.

2. Materials and methods

2.1 Plant material

Seeds of grapefruit (*Citrus paradisi* Macf. 'Duncan') were sterilized with 5% bleach for 5 min and rinsed four times with sterile water. Sterile seeds were cultivated in G medium, i.e. ½ MS, 03 mg/L NAA, 2% sucrose and 0.8% phytoagar (pH5.7) at 26°C under darkness for 4 weeks. On the transformation day, shoots of etiolated seedlings were collected and cut into fragments in 1 cm long as initial transformation explants.

2.2 Plasmid construction

Recombinants containing genomic fragments of *Poncirus* were constructed by Dr. C.X. Chen by inserting selected DNA fragments into between *Xba*I and *Spe*I cloning site of the designed vector pGK, which contained the pGreenII backbone, the nos-Kan (*nptII* under the nos promoter for kanamycin resistance), the *lacZ* gene with the multiple cloning sites from pBlueScript, and GFP as the reporter, assembled with pGreen0029 and 35S-GFP (the GFP reporter gene driven by 35S-CaMV promoter) provided by Roger Hellens and Phil Mullineaux (Figure 1).

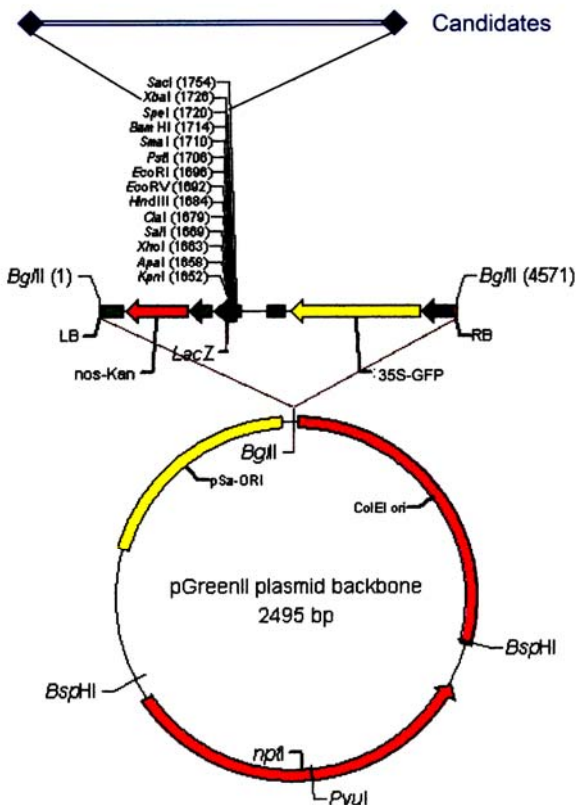


Figure 1. Scheme of Recombinant used in this experiment.

2.3 Agrobacterim transformation

1 µL target clone DNA (0.1-0.5 µg) was laid in TE buffer (pH 8.0) on top of 40 µL thawed frozen competent *Agrobacterium* AGL1 cells, on ice 1 min, DNA-competent cell mixture was pipetted in an ice cold 0.2 cm electroporation cuvette and electroporated in Bio-Rad Gene Pulser system. Parameters are 25 µF, 400Ω, 2.49 kv followed by a 8-9 ms delay. Immediately rinsed the cuvette with 0.8 mL LB antibiotic free medium and transferred the mixture to a sterile culture tube followed by incubating with shaking at 240 rpm, 28°C for 2.5 hrs. The electroporated cells were collected by centrifuge at 12,000 g for 2 min and the pellet was resuspended in 200 µL LB broth. Spread 25-50 µL of the culture on LB plate containing 50µg/mL of rifamycin, carbenicillin and kanamycin. The plates were incubated at 28°C for 2 days. Temperature should not be over 30°C which will result in loss of plasmid.

2.4 Plant transformation

Citrus transformation procedure is modified according to Yu et al. [9]. Streak *Agrobacterium* strain harboring recombinants or empty vector only on LB agar plates contained 50µg/mL of rifamycin, carbenicillin and kanamycin and incubate the above plates in incubator at 28°C for 2 days. Single colonies of *A. tumefaciens* was grown overnight at 28°C in 3 mL liquid LB medium (Invitrogen life technologies) supplemented with the same antibiotics to OD 1.0. Dilute bacterium to 1/100 vol/vol in 10 mL fresh LB broth with same antibiotics and 100 µM of the *vir* gene inducer acetosyringone (AS), and inoculate overnight to obtain culture of OD600 = 0.5-1.0. The pellet of *A. tumefaciens* was collected through centrifuge at 3000 g 10 minutes at room temperature. Re-suspend the bacterium pellet with 10 mL resuspended medium RM and centrifuge as above. Liquid bacterium resuspension medium (RM) is MS supplemented with 6-benzylaminopurine (6-BA) 3 mg/L, α-naphthaleneacetic acid (NAA) 0.1 mg/L and sucrose 50 g/L, pH=5.5. Recentrifuge and Resuspend the washed pellet with another 10 mL RM for transformation infection. Around 1 cm long cut epicotyl fragments from four weeks old etiolated seedlings were immersed in RM bacterium liquid for about 5 minutes, shaking slowly and frequently by forceps and then extra liquid was removed by autoclaved filter paper. The infected explants were placed horizontally on co-cultivation medium (CM) which is RM supplemented with 100 µM AS. Spare some non-infected and infected explants for setting up systematic controls. All co-cultivated plates were sealed with parafilm and placed at 26°C in the continuous darkness for 2.5 days. Explants were then transferred onto selection medium (SM) for selection and regeneration. Two steps were employed in selection phase, first in continue darkness for two weeks and then in the fluorescent

light on the photoperiod of 16h/8h light/darkness for four weeks. Solid selection medium is MS supplemented with 6-BA 3 mg/L, NAA 0.4 mg/L, sucrose 50 g/L, cefotaxim (Cef) 450 mg/L, 75 mg/L of kanamycin as a selection reagent and phytoagar 8g/L, pH=5.8. After four weeks of selection, agar was carefully removed from regenerated shoots in selection and regeneration medium by rinsing in tap water and they were grafted onto Carrizo rootstock grown in trays on chamber or in tubes for speeding up growth. MS basal salt Mixture (Sigma #M5519) was substituted for MS in this research.

2.5 GFP observation

Explants with small regenerated shoots grown on the parafilm-sealed culture plates containing selection medium were observed and images were captured under the digital fluorescence microscope. Shoots were then grafted onto rootstocks and marked accordingly for further growth and detection.

2.6 Identification of transgenic plants through PCR technique

All marked putative transformants were screened initially through PCR with the REDExtract-N-Amp Plant PCR Kits following manufacturer's instruction (Sigma, Saint Louis, Missouri, USA). Briefly, a 0.5 cm leaf disc was punched from each plant using a

standard one-hole paper punch and collected in a 2-ml collection tube followed by adding 100 μ l of Extraction Solution. Close the tube and vortex briefly. Make sure the disk is covered by the Extraction Solution. Incubate samples at 95 $^{\circ}$ C for 10 minutes. Add 100 μ l of Dilution Solution and vortex to mix. Store the diluted leaf extract at 2-8 $^{\circ}$ C until use. 4 μ l of the diluted extract was used for each PCR reaction.

Selectable marker gene *npt II* and reporter gene *gfp* were checked in all marked transformed plants. *NptII* gene-derived primers were 5'- TCGGCTATGA CTGGGCACAACAGA-3' (24-mer) (forward), and 5'- AAGAAGGCGATAGAAGGCGATGCG-3' (24-mer) (reverse), which are within *nptII* (conferring kanamycin resistance) coding region of *npt II* gene. PCR product is 722 bp. Primers derived from *gfp* coding region are available upon request.

Reaction parameters are at 94 $^{\circ}$ C, 3 min and then 94 $^{\circ}$ C, 2 min; 93 $^{\circ}$ C, 30 sec, 55 $^{\circ}$ C, 90 sec, 72 $^{\circ}$ C, 1.5 min, for 35 cycles; 72 $^{\circ}$ C, 7 min for final extension. 15 μ L PCR products were loaded in 1.0 % agarose gel containing 0.1 μ g/mL ethidium bromide for each sample and gel was run in 1 \times TAE at 7 v/cm. Images were captured with digital gel imaging system.

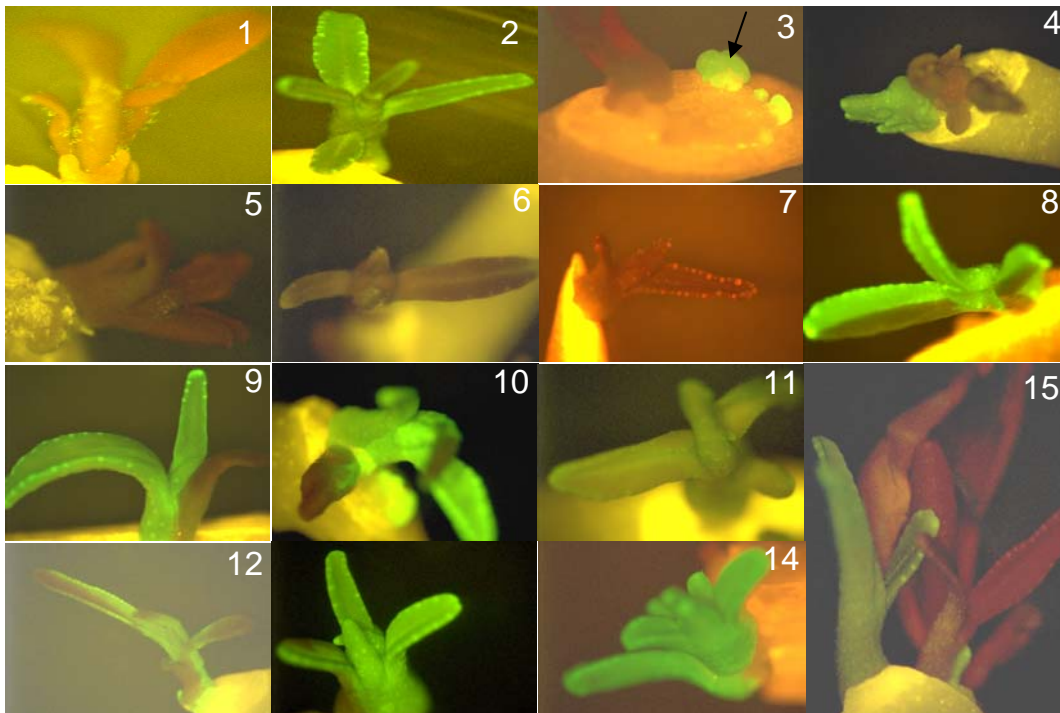


Figure 2. GFP expression of transgenic shoots/tissue of grapefruit under stereomicroscope. 1: nontransgenic regenerated shoots on explants, clearly appeared red, negative control; 2-15: Transgenic shoots being verified by PCR, showing different phenotypes of expression of *gfp* gene, arrow points at callus containing GFP protein. Plants 7-12 showed sectorial expression of *gfp* in leaves from chimeric regenerated shoots. Plants 13, 14 showed transgenic plants displaying bright green fluorescence. Plant 15 showed different types of transformants regenerated lined closely on the same cutting end of an explant.

3. Results and Discussion

3.1 Types of transgenic plants observed under the digital fluorescence microscope

GFP expression observation showed that there are some different phenotypes over the expression of *gfp* gene in regenerated shoots of transgenic grapefruit; some are chimeric (Figure 2). transformants appeared fully bright green fluorescence under the fluorescence stereomicroscope were selected from others for future functional testing of the target genes.

3.2 Foreign gene identification in tranformants and estimation of transformation efficiency through REExtract-N-Amp Plant PCR Kit

nptII gene and *gus A* gene were clearly detectable consistently in 66.67% (12 out of 18) of regenerated plants. *nptII* gene was detectable in another 16.67% (3 out of 18) of regenerated plants, while *gus A*

gene was not. There were 16.67% of regenerated plants that were considered as escapes (Fig. 3). In term of consistent PCR analysis with both selectable marker and the reporter gene, transformation rate are approximately 7.75% and 2.03% for empty vector and recombinant containing genomic DNA inserts respectively (Table 1). Target genomic gene (s) can be detectable in only a portion of those plants that contained both selectable and reporter genes (data not shown). The REExtract-N-Amp Plant PCR Kit enables us to look into each portion of the plant and up to one hundred transformants at a time in a day. More leaves discs from different leaves are recommended if the chimeric plants appear in the transformation events. The corresponding results depend on the type of the transgenic plants or on which portion of the chimeric leaves was sampled for PCR experiment. Expected target transgenic plants should be selected excluding the chimeric for accurate functional characterization of the candidate gene(s).

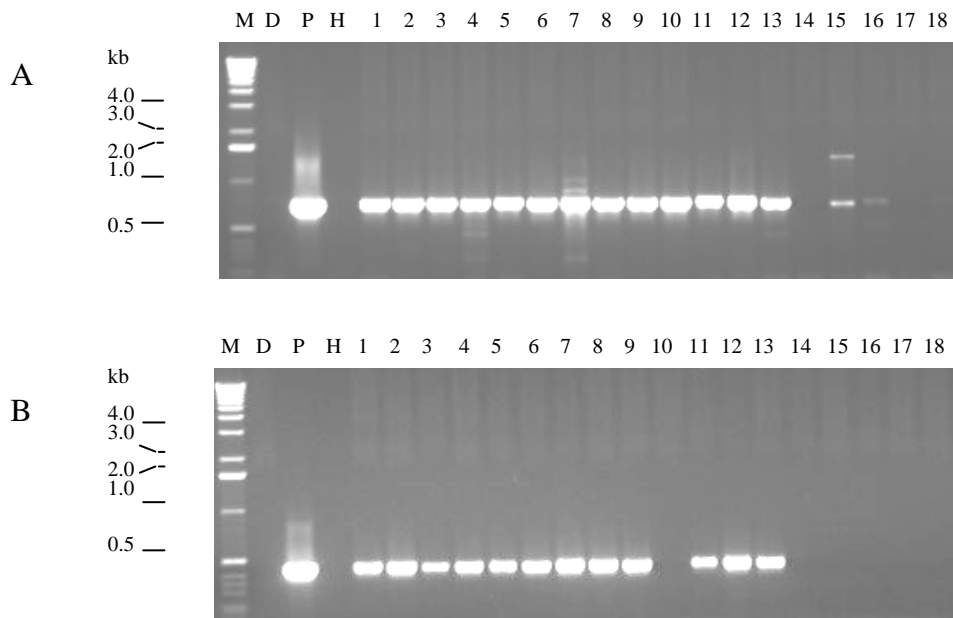


Figure 3. Electrophoresis images for PCR results amplified with a selectable gene *npt II* derived primers and a reporter gene *gfp* derived primers for transformed grapefruit plants. M: 1 kb DNA marker, the sizes of the DNA standards are indicated on left side of images. D: Duncan nontransformant, receptor plant background check, negative control P: plasmid pGREEN, PCR systematic check, positive control, H: DNA free water to check possible contamination of components for PCR 1-18: Regenerated plants showing consistent positive PCR results between selectable and reporter genes (espected transgenic plants, plants 1-9, 11-13) and inconsistent PCR results (*nptII* gene was detectable and *gus A* gene was undetectable in plant 10,15,16). Plants 16,17,18 were considered as escapes.

Table 1 Efficiency of transformation with pGREEN in Duncan grapefruit *

Plasmid	No. explants	No. plants regenerated	Regeneration Rate (%)	PCR <i>gfp</i> & <i>nptII</i> positive plants	Transformation Rate (%)
Empty vector	142	12	8.45	11	7.75
pK1	186	4	2.15	4	2.15
pK2	315	9	2.86	6	1.90

* Regeneration rate (%)=(No. of plants produced/ No. of explants)×100, Transformation rate (%)= (No. of PCR positive/ No. of explants) ×100, pK1 and pK2 are two different recombinant clones containing different genomic inserts. Average transformation rate of these two recombinants is 2.03%.

Acknowledgement

The authors would like to thank Ms Marjorie K. Wendell for her excellent technical assistance in this research. Thank Dr. Vladimir Orbovic, manager of Core Citrus Transformation Facility, at UF/IFAS CREC for his their helpful assistance in GFP observation.

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