

# Genetic transformation of *Paulownia elongata* S. Y. Hu., mediated by *Agrobacterium tumefaciens* and biolistic system

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## ABSTRACT

**Objective:** The most appropriate conditions for genetic transformation through direct (bioballistic) and indirect (*Agrobacterium tumefaciens*) transformation systems in *Paulownia elongata* were established.

**Design/methodology/approach:** Starting from *in vitro* propagation through both direct and indirect organogenesis, internodal stem segments with 0.5 to 1 cm length were determined as the best explant. The optimum dose for selection media was determined to be 15 mg L<sup>-1</sup> of kanamycin. It was possible to obtain transgenic plants under both transformation systems. In the case of *Agrobacterium tumefaciens*, two hours of incubation, 48 h of co-cultivation, and optical density of 0.9 were used; while for bioballistics, the best conditions were 120 PSI of shot pressure, shot height at level 6 (16 cm), and vacuum pressure of 22 Hg mm, with particle inflow gun system (PIG).

**Results:** Both systems produced complete transformants, chimeras, as well as those confirmed by histochemical X-GLUC and PCR analysis, producing a total of 14 positive plants by *A. tumefaciens* transformation from 26 trials and ten positive plants by the bioballistic system from 30 trials; a construction with chitinase and glucanase, NPT II selection gene and the GUS reporter gene were used.

**Findings/conclusions:** So far, this has been the first report including integration of chitinase and glucanase genes.

**Keywords:** (BA) 6-bencil-adenine, (ANA) naphthalenic acid, micro-propagation.

## INTRODUCTION

*Paulownia elongate* (Scrophulariaceae) originates in China, from where it extended to countries like Japan, Australia, Brazil and the U.S.A. in the 20th century, and to Mexico at the end of the 20th century, considered as a species potential for production. It is a forest species of rapid growth and considered an alternative for the establishment of timber-yielding commercial plantations (Tang *et al.*, 1980). *Paulownia* wood is of low density, which provides lightness and allows its use in manufacture of furniture, crafts, musical instruments, and interior decoration. Because of their quick growth all the species of genus *Paulownia* have great potential for reforestation and improvement of poor soils (Zhu *et al.*, 1986, Melhuish *et al.*, 1990), they are drought-tolerant and

adapt to different types of soil, which increases their economic potential (Tang *et al.*, 1980). These qualities, however, make them especially vulnerable for the distribution in plantations under agro-climatic conditions other than those of their original habitat, which may lead to failure, furthermore, susceptibility to soil pathogens such as *Rhizoctonia* has been reported in different *Paulownia* species (Mehrotra, 1994; 1997).

The aforementioned, may be moderated by the improvement of their adaptation qualities through genetic transformation. Genetic transformation is a technique that consists in the integration of strange genes to the genome of an individual through the techniques of recombinant DNA, the incorporated genes can improve plant characteristics by direct methods such as bioballistics, which is the acceleration of particles with DNA of desirable features (Vain *et al.*, 1993), or indirectly by *Agrobacterium tumefaciens* through the *Ti* plasmid (Herrera-Estrella *et al.*, 1983), thanks to advances in *in vitro* propagation by different organogenic pathways in species of the *Paulownia* genus, such as propagation by shoot tips (Song *et al.*, 1990; Sharma *et al.*, 2003), or direct organogenesis from different explants (Song *et al.*, 1991; Castillo *et al.*, 2012); such as by direct embryogenesis (Ipecki and Gozukirmizi, 2003) or indirect embryogenesis (Ipecki and Gozukirmizi, 2004), laying the foundation for the initial processes to carry out research aimed at genetic engineering.

With the purpose of improving the characteristics in *Paulownia elongata* and the adequate conditions for

genetic transformation of this species, studies have been carried out designed to determine the susceptibility of *P. elongata* to *Agrobacterium tumefaciens* and *Agrobacterium rhizogenes*, which demonstrated that inoculating shoots generated under *in vitro* conditions with the strains 542, A281, and C58 of *A. tumefaciens*, the formation of crown gall is possible with a frequency by 83%. Other plants have promoted quick calli proliferation and root formation in damaged shoot parts, when inoculated with strain R1601 of *A. rhizogenes*. The molecular analyses showed the presence of opinae, the same as Southern blot tests, which confirmed the existence of the transformation plasmid (T-DNA) in callus as well as in roots (Bergmann *et al.*, 1999). In other species like *P. fortunei*, callus was obtained, transformed after co-cultivating petiole segments with *A. tumefaciens* strain LBA 4404 with a binary vector (pB121), that included the GUS reporter genes and those of NPT II selection, resistant to kanamycin, and the transformation was confirmed by histochemical tests of the GUS gene expression and the detection of incorporation of the selection gene to the genome of transformed calli (Mohri *et al.*, 2003). Transformation studies, already approached to gene integration with the purpose of conferring some characteristics have been carried out in a hybrid of *P. tomentosa* × *P. fortunei*, which was transformed via *A. tumefaciens* with vector p43 8PRSI, containing the *shiva-1* gene, which codes a bactericide peptide controlled by the promoter gene called CaMV35S. The molecular analyses like PCR, Southern blot, showed the successful integration of the *shiva-1* gene in the plant genome, confirming the gene transcription through RT-PCR (Tao *et al.*, 2005).

On the other hand, with the objective to generate resistance to root diseases, studies have been conducted with chitinase genes (RCH10) and  $\beta$ -1-3-glucanase (AGLU) genes, having generated genotypes with tolerance to pathogens like *Rhizoctonia solani* in *Oryza sativa* L. (Mao *et al.*, 2003; Li *et al.*, 2004). The aforementioned allows considering the viability of application of these techniques and incorporating them to forest species para generar arboles resistententes a enfermedades como reportaron en *Populus deltoides* (Zou *et al.*, 2006); therefore the objective of this study was integrating chitinase-glucanase genes through genetic transformation by *Agrobacterium*-mediated and bioballistic systems in *Paulownia elongata* (S.Y.Hu.).

## MATERIALS AND METHODS

### Genetic Transformation

The tests of *Agrobacterium tumefaciens*-mediated transformation were established based on the principles created by Horsch *et al.* (1985), utilizing *Agrobacterium tumefaciens* strain LBA4404, which contains the plasmid pCAMBIA 2301-CHIT-GLU with chitinase and glucanase genes, the GUS reporter gen, and the NPTII selection gen (resistance to kanamycin) with a total size of 11621 base pairs. In order to conduct the transformation assays, internodal segments of 0.5 m length were used generated in MS medium (Murashigue and Skoog, 1962), cultivated for 35 days to favor the greatest possible internode elongation. Within the culture recipient, shoots of 8 cm length on average were produced, and the dissected segments were placed in liquid MS medium before their inoculation with the *Agrobacterium*

*tumefaciens* strain. Different transformation conditions were created through variations in bacteria incubation times, bacterium concentration, measurement through culture optical density, and times of co-incubation.

### Transformation by Particle Acceleration

The particle acceleration system utilized was a PIG (Particle Inflow Gun), (Vain et al. 1993) which operates with helium gas and tungsten particles as projectile agents and carrying plasmids at low pressure (40-120 *psi*) utilizing a vacuum pump (25-30 *psi* Hg). Calli were used for transformation by particle acceleration, which were obtained from internodes cultured in MS medium supplemented with 4 mg L<sup>-1</sup> of TDZ, 3% of sucrose, 0.7% agar, and pH of 5.8. Its growth period took until reaching 1 cm<sup>2</sup> at two weeks of 26 °C under conditions of darkness. Different transformation conditions were established, such as: variations in utilized pressure, regulating helium gas outlet and shot speed, keeping constant plasmid concentration in particles, and shot distance.

### Selection of Transformed Segments and Shoot Evaluation

The segments of the transformation assays were transferred to a selection medium made up of MS medium, added with 4 mg L<sup>-1</sup> of BA, 0.2 mg L<sup>-1</sup> of ANA, 15 mg L<sup>-1</sup> of kanamycin, 3% sucrose, 0.7% agar, and pH of 5.8, the stem segments as well as calli remained for a period of 35 days, time when shoot formation was initiated. The shoots longer than 1.5 cm were dissected, taking a 2 mm segment in order to conduct the histochemical XGLUC test and observe the expression of the GUS gene in the tissues, the rest of the shoot was used for DNA extraction required for PCR analysis confirming the integration of the genes of interest for the genome of the transformed shoots.

## RESULTS AND DISCUSSION

### Genetic Transformation

A total of 26 assays were carried out representing the management of 7,984 explants, submitted to *Agrobacterium*- mediated transformation process. Only 37 shoots were obtained from all of them (Table 1), which produces a mean transformation frequency of 1.12 of the total of shoot assays; each shoot-yielding assay, however, had its particular percentage. It is important to emphasize that the highest transformation frequency of independent experiments was obtained with two hours of bacteria incubation, more than 40 h of co-cultivation, and optical density from 0.8 to 1.1, since it is quite

probable that higher optical density coincides with the stationary phase of bacterial growth, leaving less activity to strains and thus lower transformation efficiency; likewise, it is probable that the strain working with low optical density (>0.5) in the phase of growth lag and the time of co-cultivation, may be not sufficient for the bacterium to infect the tissues properly. Therefore, the obtained results partially agree with those reported by Mohri et al. (2003), who after 48 h of co-cultivation found better response in segments of leaves, stem, petiole, and root, but at incubation times not longer than 30 min, and maximum optical density of 0.5 in petioles. Even though in the present research we worked with internodes determined by direct organogenesis response, it is considered that this way is the most appropriate for *P. elongata*, taking into account that the studies previous to trans-formation with species of genus *Paulownia* spp. had led to obtaining of calli and roots, not of shoots or complete plants, this factor being the most important of the expected results. Even when it had been necessary to utilize a great many initial explants, in the total of assays as well as in each of them particularly, in none of the cases fewer than 250 stem segments were used. At first we expected to obtain a larger number of shoots, starting from the selected organogenetic route; however, this did not happen, partly due to the fact that although the kanamycin dose was not higher than 15 mg L<sup>-1</sup> it is probable that it would have indirect effects on diminishing the capacity of forming shoots, even in the transformed tissues.

### Internode and Calli Transformation by Particle Acceleration

Shoot formation were evaluated in each experiment with the possibility of having a transformants in a total of 20 assays, which represented the management of 6,474 explants submitted to the transformation process by particle acceleration through (PIG). Nineteen shoots were obtained (Table 1), generating a mean transformation percentage of 1.19 of the total of shoot assays; however, putting these results on a level with similar studies on *P. elongata* presents difficulties as a species, since so far, genetic transformation through particle acceleration has not been examined yet. Nevertheless, there is information of other forest species of the coniferous family, such as *Pinus roxburghii*, where direct biolistic transformed on mature embryos was made in order to incorporate bar gene, obtaining transformed plants from epicotyls and adventitious shoots. This approaches the system used in the present study, due to the fact that after biolistic

**Table 1.** Response of assays with stem and callus samples of *P. elongata*, in X-GLUC substrate and by PCR analysis.

No. of trials	Terms	No. of SPROUTS	Positive in X-GLUC	Positive by PCR for the chitinase gene	Full or partial expression
6 A	H1=2 H2=40 OD=1.1	4	2	2	partial
9 A	H1=2 H2=48 OD=0.9	3	1	--	partial
10 A	H1=2 H2=48 OD=0.9	1	--	--	----
11 A	H1=1.5 H2=40 OD=0.8	2	2	2	partial
16 A	H1=2 H2=48 OD=0.9	6	3	3	total
17 A	H1=1.5 H2=48 OD=1.1	7	2	2	partial
18 A	H1=2 H2=48 OD=0.8	1	1	1	partial
22 A	H1=2 H2=48 OD=0.8	2	--	--	----
23 A	H1=2 H2=48 OD=0.9	5	4	2	partial
24 A	H1=2 H2=48 OD=0.7	3	--	--	----
25 A	H1=1.5 H2=48 OD=0.8	2	2	2	partial
Subtotal with <i>A. tumefaciens</i>		36	17	14	
8 Bio	PSI= 120/ A=6 /PL 1	1	--	--	----
9 Bio	PSI= 120/ A=6 /PL 1	5	3	3	total
10 Bio	PSI= 120/ A=6 /PL 1	4	4	2	partial
17 Bio	PSI= 120/ A=6 /PL 1	2	1	1	partial
20 Bio	PSI= 120/ A=6 /PL 1	1	1	1	partial
3 Biocall	PSI= 120/ A=7 /PL 1	5	1	1	partial
6 Biocall	PSI= 120/ A=7 /PL 1	7	6	2	partial
7 Biocall	PSI= 120/ A=7 /PL 1	2	2	--	partial
10 Biocall	PSI= 120/ A=7 /PL 1	4	--	--	----
Subtotal with Biobalistics	PSI= 120/ A=7 /PL 1	31	18	10	
TOTAL		67	35	24	

Footnote: A: Transformed by *Agrobacterium*  
 Bio: Transformed by biobalistics  
 Biocall: Callus transformed by biobalistics  
 H1: Hours of co-incubation with bacterium in liquid medium  
 H2: Hours of co-culture in solid medium  
 OD: Optical density of bacterial culture  
 PSI: Shot pressure  
 A7: Shot height  
 PL1: Plasmid with genes of interest **contained plasmid**  
 -- Without response

even when starting from sexual tissue in the beginning, finally the transformants are recovered from organogenesis; however, the usually preferred way for transformation by particle acceleration in conifers by 70% has been through somatic embryogenesis (Tang & Newton 2003). The aforesaid is mainly due to an increment of bombed cells in calli of embryogenic type, which increases the possibility of transformation, once the somatic embryos derived from

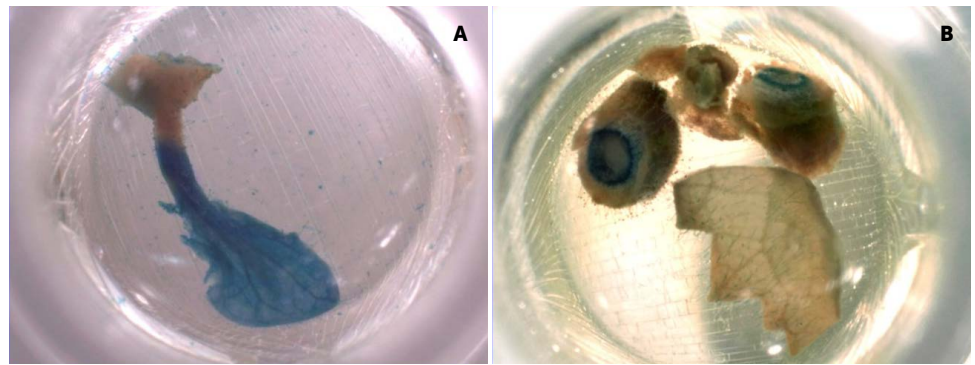
the transformed cells are being produced, since the organogenic way is associated to genetic changes by somaclonal variation. This would be in this particular case of high risk, starting from the fact that the original material is a hybrid of high plasticity and yield, for which it is not desirable to alter these characteristics; since the objective and the point only was integrating the genes of interest without changing the original condition of *P. elongate*.

The calli were transferred at half strength recuperation media without antibiotics for seven days were sub cultivated in selection medium (MSSEL) during 30 days, and shoot production was evaluated in each experiment. The total number of conducted assays was 10 with an average of 24 calli for each; but with the highest transformation percentage to the order of 21.21, which is related to the fact that even though this has to do with a reduced number of

calli each callus had 1.5 cm diameter on an average with greater impact surface for the tungsten particles, which in turn represented a larger number of possibly transformed cells, even when the way of indirect organogenesis was followed. It is important to emphasize that in the case of transformation by particle inflow gun system in forest species such as conifers, somatic embryogenesis is of 70% (Tang & Newton 2003). The use of calli and genetic transformation in *P. elongata* with *Agrobacterium*-mediated systems has been reported (Mohri et al., 2003) achieving transformed tissue and deriving in callus lines. On the other hand, Bergmann et al. (1999) report callus and root formation, after transforming with *Agrobacterium rhizogenes*; however, despite these advances, up to date, complete and stable transformation in *P. elongata* by bioballistic systems have not been reported, since there are only records of preliminary assays with temporary expression, like those carried out by Mohri and Shinohara (1996), who utilized a transformation system by particle inflow gun, in order to insert the luciferase gene in *P. fortunei*; however, they do not report stable expression neither complete plant development.

### Histochemical Evaluation and PCR Analysis of Shoots

Tissue samples in X-GLUC substrate allowed evaluating GUS gene expression, blue coloration in tissues being observed with positive response to the expression of this gene. As a disadvantage that this may be a destructive test. At the end of the evaluation, it could be observed that only 35 out of the 67 shoots obtained and kept in selection medium resulted positive for GUS, 6 of these with total expression and 29 partially (Table 1, Figure 1); the fact of achieving shoots with partially transformed tissue indicates the majority presence of chimeras, whose transformed tissues can be identified—even though they are not fully transformed individuals - and starting from them, complete individuals can be generated. Mohri et al. (2003) report calli with total expression for the case of *P. fortunei*, whereas in the present study, all the calli obtained had partial expression. It must yet be clarified that *P. fortunei* was transformed with *Agrobacterium* (species) and in the present case *P. elongata* was transformed by particle



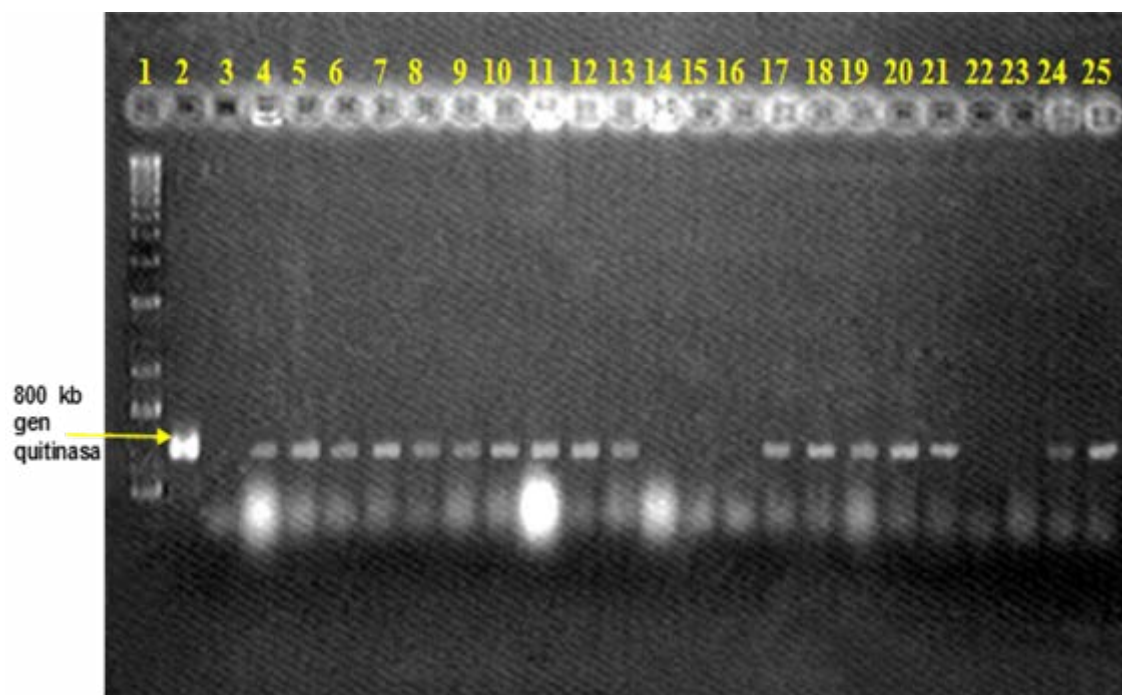
**Figure 1:** GUS expression in X-GLUC substrate of transformed tissues (A) total response, (B) Partial response

inflow gun system. Most of the calli kept in selection medium resulted positive, at least partially. In this sense, mature *Pinus roxburghii* embryos, for example, were transformed by particle inflow gun system, generally obtaining partial transformants, from which complete individuals were identified and generated from such transformed tissues, inducing adventitious shoots (Parasharami et al., 2006). Therefore, shoots and calli from *P. elongata* were kept in selection medium, in order to subsequently identify the transformed tissues and generate complete individuals, which could be identified through histochemical X-GLUC analysis.

Molecular analysis by PCR allowed assessing the incorporation of the genes of interest in the transformants, presence of bands being observed in the shoots with positive response or band absence in non-transformed shoots. We emphasize the fact that 24 out of the 67 assessed elements showed positive response (Table 1, Figure 2), 35% of the total. Interesting situations could be observed, like shoots having had partially positive response for GUS and negative response for PCR; or on the contrary, materials negative for GUS and positive for PCR. One of the main causes of chimeric effects is the fact that mostly partially transformed materials (chimeras) were obtained by *Agrobacterium* as well as by bioballistics, which brings about that tissue parts are taken for the tests, that may contain or not the transformed portions, which produces such responses in the moment of analysis.

### CONCLUSIONS

It was possible to genetically transform *P. elongata* by the direct method of ballistics or indirectly by *Agrobacterium tumefaciens*-mediated transformation in order to incorporate genes of interest. Complete as well as partial transformants could be achieved in both



**Figure 2:** PCR test for chitinase gene integration to transformed *P. elongata* materials by direct and indirect transformation systems, in agar gel of 1.5%, showing the results of Rail 1, stairs 1Kb, rail 2 positive control for chitinase gene, rail 3, negative control, rail 4-25 DNA tissue samples obtained in selection media.

systems, via direct organogenesis and indirectly for the case of transformed calli.

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