

Regeneration and Transformation *via Agrobacterium tumefaciens* of *Echinacea purpurea* L.

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Abstract

Leaf explants of *Echinacea purpurea* L. taken from aseptically germinated seedlings were inoculated with *A. tumefaciens* strains EHA105, carrying a binary vector conferring herbicide resistant *bar* gene and fungal resistant chitinase gene. Glufosinate ammonium-resistant shoots were regenerated on a medium containing BAP and NAA at a concentration of 4.88 and 0.053 μ M, respectively. A subsequent transfer of shoots to medium containing BAP was necessary for stem elongation and leaf development. Transgenic *Echinacea* plants carrying *bar* and chitinase genes were selected for their resistance to glufosinate ammonium herbicide. Molecular analysis using PCR confirmed the integration of the transgenes into plant genome. This is the first report on genetic transformation of *Echinacea* plant using *bar* gene as a selectable marker.

Introduction

Echinacea purpurea L. is a group of purple coneflowers in the family Asteraceae. It has been used traditionally as an herbal medicine and dietary supplements for hundreds of years (Percival 2000). In recent years, the purple coneflower has gained global attractiveness due to its beneficial effect on human's immune system (Bauer and Wagner 1991). Extracts from the plant have shown anti-oxidative, antibacterial, antiviral and antifungal properties, and are used in the treatment of the common cold, as well as respiratory and urinary diseases (Grimm and Muller 1999, Barrett 2003). Recent technological advances have allowed researchers to analyze some of the medicinally active compounds present in *Echinacea* sp. and to speculate on their modes of action (Choffe et al. 2000). Complex polysaccharides, such as arabinogalactane and xyloglucan, extracted from the roots of different *Echinacea* spp. have been found to stimulate

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mammalian immune systems (Coeugniet and Elek 1987) and to act as anti-inflammatory agents (Tragni et al. 1988). In the middle of the 20th century, *Echinacea* sp. was introduced as a medicinal plant to Europe (Bauer et al. 1991, Bauer 1998).

Many approaches that were unfeasible to implement by customary genetics can now be realized through transgenic techniques. Regeneration through tissue culture is a critical step for efficient transformation of most plants. However, in some species the lack of an efficient regeneration method is a huge impediment to employ the transformation technology (Penna et al. 2002). Developing protocols for efficient genetic transformation of medicinal plants with unique metabolic pathways, is important to understand the molecular basis and regulation of secondary metabolism in plants and to engineer them for specific metabolites (Pandey et al. 2010). Though transformation of a number of agriculturally important plant species has been reported, such efforts on medicinally important plants have been very few (Gómez-Galéra et al. 2007)

Despite the importance of *Echinacea* and abundant pharmacological and clinical studies, information concerning tissue culture and genetic transformation is quite rear. Recently, *Echinacea* species have been regenerated from a range of tissues from *in vitro* seedlings to mature, field-grown plants (Abbasi et al. 2007). Tissue culture of *Echinacea* can play a vital role in the development of novel germplasm, rapid multiplication, and genetic modifications for enhanced potential active compounds production. *In vitro* propagation and regeneration from petiole explants of *E. purpurea* have been established (Choffe et al. 2000). Moreover, axillary buds, adventitious shoots and somatic embryos have been used for *in vitro* mass propagation of four commercially important *Echinacea* species, including *E. angustifolia*, *E. pallida*, *E. paradoxa*, and *E. purpurea* (Lakshmanan et al. 2002).

Recovery of transgenic *Echinacea* plants *via* *Agrobacterium*-mediated transformation using neomycin phosphotransferase II (*nptII* conferring kana-mycin resistance) as a selectable marker has been reported (Koroch et al. 2002; Wang and To 2004). Transformed hairy root cultures of *Echinacea purpurea* was established by infecting different types of explants with three type strains of *Agrobacterium rhizogenes* (Wang et al. 2006). In this study a new protocol for *Agrobacterium*-mediated transformation of *E. purpurea* using *bar* gene which confers tolerance to herbicide BASTA®, as a selectable marker has been developed.

Materials and Methods

Seeds of *Echinacea purpurea* L. were secured from Floriculture Department, Faculty of Agriculture, Cairo University. *Echinacea* seeds were surface sterilized

by a solution containing 2.36% (w/v) sodium hypochlorite for 20 min and washed thoroughly with sterilized distilled water. The sterilized seeds were then germinated on basal medium containing MS salts and 0.7% agar. The pH value of the medium was adjusted to 5.8. The cultures were incubated in 25°C growth chamber under light conditions of 16 hr per day. Within four weeks of cultivation, the *in vitro* growing *Echinacea* plantlets reached about 6 - 8 cm in height and used in further regeneration and transformation experiments.

Shoot tips and leaves were cut into small segments and inculcated on MS supplemented with different combinations of NAA and BAP. Cultures were incubated in the same previous conditions for two months. Regenerated shoots (3 - 4 cm in height) were rooted on MS supplemented with different concentrations of IBA in the range of 4.92 - 14.76 μM .

An effective concentration of PPT for the selection of transformed shoots was determined by culturing non-transformed leaf explants (control) on MS supplemented with 4.88 μM BA + 0.053 μM NAA and contains different concentrations of PPT (0, 0.5, 1.0, 1.5, 2.0, 2.5, 3 and 4 mg/l).

The cultures were transferred twice to the same medium added with the same level of herbicide at two weeks intervals and were scored for the number of surviving explants. The herbicide was added to the media after autoclaving.

The dual-binary vector system pGreenII/pSoup was used in the present work (Hellens et al. 2000). The T-DNA contains the *bar* gene fused between *nos* promoter and terminator sequences of *A. tumefaciens* as a selectable marker and a heterologous chitinase gene (Chit30) from *Streptomyces olivaceoviridis* ATTCC11238 (Fig. 1). The chimeric chitinase gene was cloned via PCR based method into pGreenII binary vector 0229 under constitutive double 35S promoter

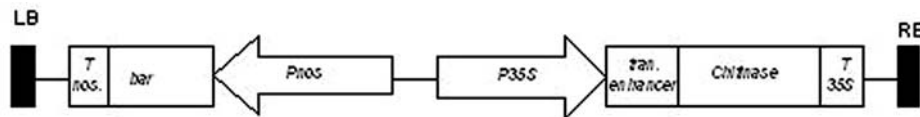


Fig. 1. Schematic structure of the T-DNA region of the transformation vector used for *Echinacea* transformation. *Pnos*, promoter sequence of the nopaline synthase gene; *Tnos*, terminator sequence of the nopaline synthase gene; *P35S*, 2x35S promoter of cauliflower mosaic virus; *tran. enh.* Translation enhancer; *T35S*, terminator sequence of cauliflower mosaic virus; *RB* and *LB*, right and left borders, respectively, of the T-DNA region. The construct is not shown to scale.

from cauliflower mosaic virus (Provided by Hans-Joerg Jacobsen and Fathi Hassan, Leibniz University Hannover, Germany). The *bar* gene encodes a phosphinothricin acetyltransferase (PAT) enzyme which confers resistance to bialaphos and the related compounds phosphinothricin (PPT), the active ingredient of herbicide BASTA® and gulfosinate ammonium, through acetylation. Young leaves were used as the explants for transformation experiments. *A. tumefaciens* strain EHA105 (Hood et al. 1993) harboring the

transformation vector, was cultured in LB liquid medium supplemented with 50 mg/l kanamycin. Bacterial cultures were grown on an orbital shaker at 28°C until OD₆₀₀ = 1.0. *Agrobacterium* cells were harvested by centrifugation at 8000 xg for 10 min and re-suspended in liquid cocultivation MS (4.88 µM BAP and 0.053 µM NAA).

Explants were soaked in *Agrobacterium* suspension for 30 min and blotted dry before culturing on solid cocultivation medium for two days. After cocultivation, the explants were washed thoroughly in sterile distilled water containing 400 mg/l cefotaxime. The explants were then transferred to a selection medium containing MS + 4.88 µM BA, 0.053 µM NAA, 300 mg/l cefotaxime and 2 mg/l glufosinate ammonium and they were sub-cultured at two-week intervals to eliminate the bacteria growth and stimulate shoot regeneration. After about one month of culture, the explants started to regenerate. The shoots were excised and transferred to BA-containing medium for shoot elongation. Well developed shoots were transferred to rooting medium.

The genomic DNA was extracted from transformed and non-transformed *Echinacea* plantlets using a modified CTAB method Sul and Korban (1996). PCR analysis was conducted with the following primers, bar447-F: 5'-GATTTCCGGTGACGGGCAGGA-3', bar447-R: 5' TGCGCTCGGTACGGAAGTT-3', with a predicted product size of 447 bp. For amplification of the *A. tumefaciens* picA chromosomal locus, the following primers were used: picA-1,5'-ATGCGCATGAGGCTCGTCTTCGAG-3' and picA-2, 5'-GACGCAACGCATCCTCGATCAGCT-3' (Rong et al 1991), with a prediction product size of 550 bp.

Denaturation at 94°C for 4 min followed by 30 amplification cycles (94°C/60s, 60°C/60s (bar) or 65°C/60s (picA), 72°C/60 s) and final extension step at 72°C for 10 min. The PCR products were visualized by running the completed reaction on a 1% agarose gel containing ethidium bromide. Pictures were taken under UV light. Meanwhile the plasmid was used as positive control.

Results and Discussion

Leaf and shoot tip explants prepared from *in vitro* germinated seedlings of *Echinacea* were assessed for multiple shoot regeneration on cytokinin-containing medium. Since BAP was found to be the most effective and widely used cytokinin in several plants species including *Echinacea*, alone or in combination with auxin (Koroch et al. 2002, Mechanda et al. 2003, Koroch et al. 2003, Sauve et al. 2004, Gockel et al. 1992, Harbage 2001), it was chosen to induce multiple shoot formation from shoot tip and leaf explants on MS supplemented with BAP alone at a concentration of 4.4 and 8.87 µM. After 4 weeks of cultivation the explants produced adventitious shoot (Fig. 2). Addition of NAA (0.053 µM) and

BAP (4.88 μM) to MS was the most effective, providing shoot regeneration for 93.5% of leaf explants and the highest number of shoots per explant (2.685). On the other hand, the same medium providing 52.75% of shoot regeneration for shoot tip explants with 1.52 shoots per explant (Table 1). For length of shoots, it was found that shoots produced from the shoot tip explants were longer than ones comes from leaf explants (4.86 and 3.81 cm) respectively. Increasing NAA concentration resulted in increased callus production and low shoots initiation (data not shown). It is clear however that leaf explants were more competent for regeneration than shoot tip explants.

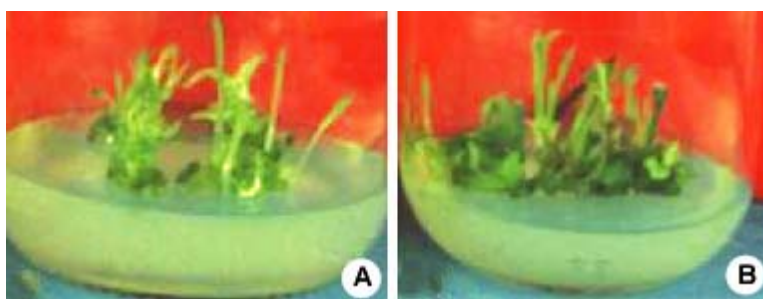


Fig. 2. Direct regenerated plantlets derived from leaf (A) and shoot tip (B) explants of *E. purpurea* cultured for two months on MS supplemented with 0.053 μM NAA and 4.88 μM BAP.

Table 1. Morphogenetic response of *Echinacea purpurea* explants after two months cultivation on MS fortified with 0.053 μM NAA and 4.88 μM BA.

Explant	Frequency of shoot formation (%)	Number of shoots per explant	Shoot length (cm)
Shoot tip	52.75 \pm 9.71	1.52 \pm 0.066	4.86 \pm 0.355
Leaf	93.5 \pm 11.84	2.685 \pm 0.133	3.81 \pm 0.282

Each value represents the mean \pm SE of four replicates.

Root formation is an obligatory phase for micropropagation of plants produced *in vitro*. Some of them initiate roots without special treatments while others require a medium supplemented with different growth regulators essentially of an auxin nature. Different plant species might vary in their requirement of auxin type for adventitious root formation. Shoots produced from either leaf or shoot tip explants of *Echinacea* were cultured on MS supplemented with different concentration of IBA (0.0, 4.92, 9.84 and 14.76 μM). Data showed that, shoots come from both leaf and shoot tip explants gave roots on MS-basal medium after one month of incubation (Table 3). Increasing IBA concentrations resulted an increasing in root length (Table 2). Shoots come from leaf explants

had roots longer than ones come from shoot tip. Using IBA at a concentration of 14.76 μM gave the best result (5.0 cm). In previous reports, plant

Table 2. Effect of different concentrations of IBA on root development of *in vitro* regenerated shoots derived from shoot tip (ST) and leaves (L), of *Echinacea purpurea*.

IBA (μM)	Frequency of root formation (%)		Number of roots		Root length (cm)	
	ST	L	ST	L	ST	L
Control	14.25 \pm 2.85	18.25 \pm 4.27	1.5 \pm 1.55	2.0 \pm 0.19	0.2 \pm 0.02	0.3 \pm 0.04
4.92	38.25 \pm 7.27	52.5 \pm 9.01	2.2 \pm 0.64	4.5 \pm 0.58	0.5 \pm 0.04	1.1 \pm 0.09
9.84	47.0 \pm 6.98	73.75 \pm 9.55	3.0 \pm 0.91	6.75 \pm 0.64	0.8 \pm 0.06	2.7 \pm 0.41
14.76	51.0 \pm 7.71	89.0 \pm 12.72	4.15 \pm 1.49	7.5 \pm 0.92	1.0 \pm 0.09	5.0 \pm 0.91

Each value represents the mean \pm SE of four replicates.

regeneration from petiole explants of *E. purpurea* was achieved by using only a small amount of BAP (Choffe et al. 2000), whereas, in the present study, BAP in combination with NAA was most effective in inducing adventitious shoot regeneration from leaf explants. Response of leaf explant to BAP and NAA concentrations in the media could be a reflection of probable differences of endogenous hormonal levels in the explant sources or different tissue sensitivities to these plant growth regulators (Lisowska and Wysonkiska 2000). All shoots longer than 1.5 cm were transferred to rooting medium for root development. The survival rate of regenerated plantlets transferred to soil was 95%.

In view of the fact that the selection of transformed cells is a prerequisite to facilitate shoot regeneration, the choice of selectable marker gene, selective agent, and timing of application is a key step in this process. The selectable marker *bar* gene of *Streptomyces hygrosopicus* encodes phosphinothricin acetyl-transferase (PAT), which inactivates phosphinothricin (PPT); It is the ammonium salt of glufosinate, the active component of BASTA by acetylation (Thompson et al. 1987). Therefore, glufosinate ammonium (PPT) was used to select the *Echinacea* transformed plants during tissue culture.

A gradual decrease in survival explants was observed in leaf explants cultured on increasing concentration of PPT (data not shown). PPT was found to be lethal at concentration of 16.56 μM , as it completely inhibited regeneration as well as survival of the explants; hence, this concentration was applied for the selection of transformed shoots. Herbicide-based selection of transformants has led to the successful recovery of transgenic plants such as *Pisum sativum* (Schroeder et al. 1993, Bean et al. 1997), *Glycine max* (Zhang et al. 1999), *Lupinus*

species (Pigeaire et al. 1997) and *Trifolium subterranean* (Khan et al. 1994), faba bean (Hanafy et al. 2005), rice (Wenefrida et al. 2007), sweet potato (Yi et al. 2007), *Sedum erythrostichum* (Yoon et al. 2002).

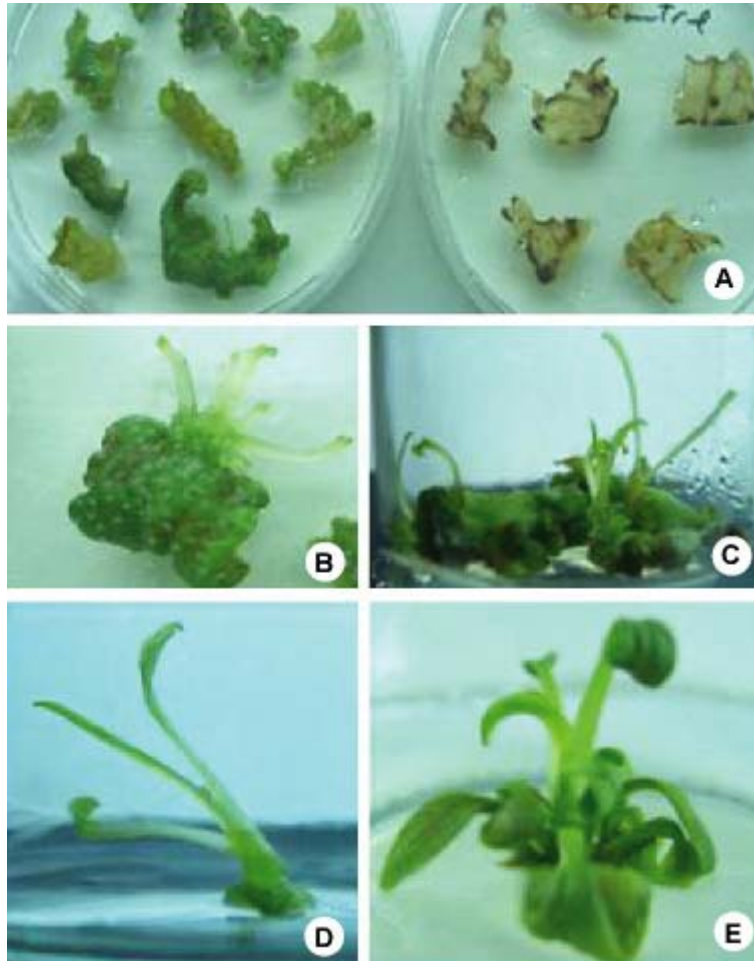


Fig. 3. Plant regeneration of *Echinacea purpurea* at different stages of the transformation procedure:

A. leaf explants were excised from *in vitro* grown plantlets, transformed by *A. tumefaciens* and cultured on Petri dishes containing selection medium supplemented with 16.56 μM PPT and 828.33 μM cefotaxim (right plate is untreated (control) explants). B. A late stage of shoot bud formation from leaf explants. C. A typical culture showing formation of calli and differentiated shoots. D. Elongated shoots cultured on PPT free medium. E. A typical regenerated transgenic shoots.

In series of transformation experiments with *Echinacea* the explants (leaf) were inoculated with *Agrobacterium* strain EHA105 containing a transformation vector harboring *Chit* and *bar* genes, selection was done by 16.56 μM PPT. After three - four weeks of culturing on selective medium, all control explants had died. On the other hand, a number of *Agrobacterium* treated explants started to regenerate (*via* organogenesis) and about 3 - 4 shoots appeared from each explant

on the regeneration medium. A higher percentage of regenerated shoots was obtained from leaf explants cocultured three days in the dark with *Agrobacterium* and then further cultured in the selection medium for up to three months. Subsequently the regenerated shoots were transferred onto MS medium containing BAP at a concentration of 4.88 μM for stem elongation and leaf development. The shoots selected for 2 - 3 months were transferred to rooting medium. Several independent transformants have been regenerated but due to the low growth of the regenerated shoots we recovered only 5 independent clones and further analyzed by PCR. Schematic representation for *in vitro* regeneration and *Agrobacterium*-mediated transformation of *E. purpurea* from leaf explants is shown in Fig. 3.

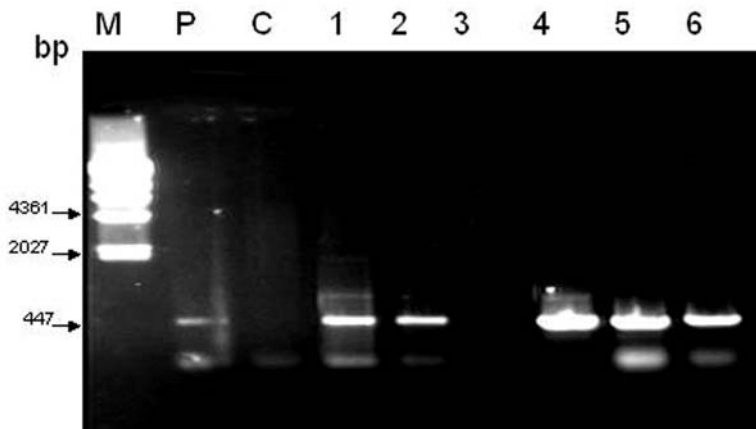


Fig. 4. PCR analysis of putative transgenic plants, DNA primed with oligonucleotides specific to the *bar* gene. Lane M: DNA molecular weight marker; Lane P: Positive control (plasmid). Lane C: DNA from non-transformed (control) plant. Lanes 1-6: DNA from different primary transformants. Lane 3: Escaped plant.

PCR analysis showed amplifications of 447 bp corresponding to the *bar* gene, indicating the presence of transgenes in 5 out of 6 putatively transformed plants recovered (Fig. 4). The negative results could be due to non-transformed shoots surviving in the selection medium (Hess et al. 1990, Langridge et al. 1992). When *Agrobacterium* chromosomal-specific primers were used, no amplification was detected in any of the transgenic materials analyzed (data not shown). This indicated that no residual agrobacteria were present in the analyzed material.

In conclusion, we have developed a simple and reliable genetic transformation system for *Echinacea purpurea* with *bar* as selectable marker for the first time. Transformation procedure, involving direct shoot organogenesis, therefore, is rapid to obtain rooted plants and can be of routine use in *Echinacea purpurea* transformation for studying gene manipulation in this crop and for transferring desirable agronomic traits.

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