



Agronomía Colombiana

ISSN: 0120-9965

Universidad Nacional de Colombia, Facultad de Agronomía

Rojas, Adriana; Lopez-Pazos, Silvio; Chaparro-Giraldo, Alejandro
Screening of Colombian soybean genotypes for *Agrobacterium*
mediated genetic transformation conferring tolerance to Glyphosate
Agronomía Colombiana, vol. 36, no. 1, 2018, January-April, pp. 24-34
Universidad Nacional de Colombia, Facultad de Agronomía

DOI: <https://doi.org/10.15446/agron.colomb.v36n1.67440>

Available in: <https://www.redalyc.org/articulo.oa?id=180358537004>

- How to cite
- Complete issue
- More information about this article
- Journal's webpage in redalyc.org

UAEH
redalyc.org

Scientific Information System Redalyc

Network of Scientific Journals from Latin America and the Caribbean, Spain and Portugal

Project academic non-profit, developed under the open access initiative

Screening of Colombian soybean genotypes for *Agrobacterium* mediated genetic transformation conferring tolerance to Glyphosate

Tamizaje de genotipos de soya colombianos para transformación genética mediada por *Agrobacterium* confiriendo tolerancia a glifosato

Adriana Rojas¹, Silvio Lopez-Pazos², and Alejandro Chaparro-Giraldo^{1*}

ABSTRACT

Soybean is a very important crop worldwide due to its multiple uses as raw material for industry and to its high nutritional value. Colombia consumes a large amount of imported soybean because domestic production does not supply demand. There are soybean varieties adapted to the environmental conditions in the Colombian territory, but none of them have been enhanced by genetic engineering to confer competitive advantages compared to imported product. In this research, the Colombian soybean varieties SK7, P29 and Soyica P34 ability to be genetically transformed by *Agrobacterium tumefaciens* strains AGL0 and EHA105 using a Glyphosate tolerance cassette was tested. It was found that SK7 variety presented a better regeneration performance from the cotyledonary node, and also had the highest transformation frequency with AGL0 strain. The P29 variety was also transformed, but a lower efficiency was registered. It was not possible to transform Soyica P34 variety under the established parameters. This research is an advance towards the construction of a platform to enhance the generic transgenic crops in Colombia.

Key words: *Glycine max*, herbicide, genetically modified organism, *in vitro* regeneration, transgenic plant.

RESUMEN

La soya es un cultivo muy importante a nivel mundial debido a sus múltiples usos en la industria y a su alto valor nutricional. Colombia consume una gran cantidad de soya, principalmente importada, porque la producción interna no supe la demanda. Existen variedades de soya adaptadas a las condiciones medioambientales del territorio colombiano, pero ninguna de ellas ha sido mejorada por ingeniería genética para conferir ventajas competitivas al cultivo frente a las importaciones. En este trabajo se evaluaron las variedades de soya SK7, P29 y Soyica P34 respecto a su capacidad para ser transformadas genéticamente por las cepas de *Agrobacterium tumefaciens* AGL0 y EHA105, utilizando un casete de tolerancia a glifosato. Se encontró que la variedad SK7 presentó un mejor desempeño en regeneración a partir de nudo cotiledonar, e igualmente tuvo la mayor frecuencia de transformación con la cepa AGL0. La variedad P29 también fue transformada, aunque con una eficiencia menor. No fue posible transformar la variedad Soyica P34 bajo los parámetros establecidos. Este trabajo fue un avance hacia la construcción de una plataforma de generación de cultivos transgénicos genéricos en Colombia.

Palabras clave: *Glycine max*, herbicida, organismo genéticamente modificado, regeneración *in vitro*, planta transgénica.

Introduction

Soybean (*Glycine max*) is a Fabaceae plant, whose seeds contain sugars (~30%), protein (~35%), edible oil (~20%), fiber, vitamins, and minerals. Soybean is a source of protein comparable to meat or eggs. Soybean cake is used as animal feed or industrial substrate (Widholm *et al.*, 2010). The soybean production for 2016 was calculated at 320 million t. United States and Brazil are major soybean producer countries. United States has an estimated soybean production of 108 million t, and their harvested area is estimated at 33 million ha. Brazil has a harvested area of 33 million ha with a production of 100 million t (USDA, 2017).

Colombia with an average production rate of 75,000 t in 2016 is considered a small soybean producer, occupying the 37th position worldwide, and the 6th in South America (Fenálce, 2017). Colombian soybean is developed mainly in the Eastern Plains region, which is a key area for agricultural development due to its plain geography suitable for technification, vast land extensions, and development opportunities, especially in the post-conflict period. Considering that soybean is a plant originated and cultivated in temperate latitudes, and due that Colombia is a tropical country, several Colombian breeding programs have been developed since the 80s producing varieties adapted to local soil and weather conditions, including relevant differences in plant physiology parameters as the plant photoperiod

Received for publication: 4 September, 2017. Accepted for publication: 1 March, 2018

Doi: 10.15446/agron.colomb.v36n1.67440

¹ Plant Genetic Engineering Research Group, Biology Department, Universidad Nacional de Colombia, Bogota (Colombia).

² Sciences Faculty, Universidad Antonio Nariño, Bogota (Colombia).

* Corresponding author: achaparro@unal.edu.co



and flowering processes that are correlated to the production rates (Valencia and Ligarreto, 2010). Currently, there are several soybean varieties with a great adaptation to local conditions, and high production (over 25,000 kg ha⁻¹) which are used by local farmers.

The genetic modification by transgenesis (GM) in commercial plants has been a useful technology providing farmers with a tool to increase crop yield, reduce the quantity of pesticides, and increase the farmer profit (Klümper and Qaim, 2014). Soybean has been bred by transgenesis, including different traits as herbicide tolerance, insect pest resistance, and improvement of oil quality (ILSI Research Foundation, 2017). Particularly, herbicide tolerance has been a very widespread and successful trait on soybean, having adoption rates of 94% in USA, 96.5% in Brazil, and 100% in Argentina in 2016 (ISAAA, 2016). GM soybean enhanced with herbicide tolerance has several advantages compared to the regular varieties. Among them, it allows the use of one herbicide per crop, a longer period of weed control, a lower Glyphosate concentration in soil, less herbicide application events, the use of low toxicity herbicides such as Glyphosate, and higher profits to the producer as an expression of all the above. Also, herbicide tolerant soybean is compositionally equivalent to the conventional genotypes and finally, the expiration of patents protecting herbicide tolerant soybeans can be a base for generic GM crops (Bonny, 2009). In Colombia, the government policies allow growers to produce transgenic plants and there are no restrictions to the consumers. Last reports indicate that in 2016 100,000 ha of GM maize were planted, of which 9,800 ha were GM cotton and 12 ha were GM flowers (ISAAA, 2016). Governments are committed to assess and manage the risks associated with the development and release of genetically modified crops. There is an established regulation of GM soybean specially attending the food safety affairs, including a maximum limit of herbicide, substantial equivalence, and varieties description. Countries have developed protocols by GM detection based on phenotypic or molecular assays (Tillmann *et al.*, 2004).

Agrobacterium tumefaciens is a relevant microorganism used in transgenesis of crops. This process is based on the transfer of a DNA segment (T-DNA) from a tumor-inducing (Ti) plasmid, which is incorporated into the plant genome with its resultant expression. The T-DNA and Vir proteins form a molecular set that delivers a single strand of this T-DNA into the cell (Bourras *et al.*, 2015). Soybean is a recalcitrant plant for *Agrobacterium*-mediated transformation producing low transformation efficiencies and requiring the use of hypervirulent bacterial strains and

specific plant genotypes to allow its transformation (Atif *et al.*, 2013). Some protocols have been applied to improve the transformation efficiency using *A. tumefaciens*; these methodologies include the modification of certain sanitation and infestation procedures toward observing possible differences among genotypes (Liu *et al.*, 2013).

Glyphosate, whose chemical name is N-(phosphonomethyl) glycine (C₃H₈NO₅P), is an odorless white strong acid. It is a crystalline powder with a fusion point of 184.5°C and molecular mass of 169.1 g mol⁻¹. Glyphosate is a systemic herbicide and can be used with practically any type of crop to control weeds worldwide. Enzyme 5-enolpyruvylshikimate-3-phosphate synthase (EPSPS) is the target of Glyphosate, which is directly involved in the synthesis of the aromatic residues: phenylalanine, tyrosine, and tryptophan (Shikimate pathway). In 1996, the first GM soybean with Glyphosate tolerance was introduced in USA, expressing an EPSPS protein from a CP4 strain of *A. tumefaciens*, which has no affinity for Glyphosate and allows the normal perform of the shikimate pathway (Duke and Powles, 2008; Duke and Cerdeira, 2010). Glyphosate tolerance is a successful trait in soybean crop, and it currently offers the possibility to develop generic transgenic crops. Considering that patents that cover the development process of this particular trait recently expired (Jefferson *et al.*, 2015), there is an increase in the freedom to operate (FTO) of the commercial plants related. Colombian soybean varieties have not been bred by transgenesis despite the fact that they are approved as commercial crops with Glyphosate tolerance (ICA, 2010) and that imported GM soybean seeds have low possibilities to grow well in tropical conditions. The main objective of this study was to evaluate the capacity of Colombian soybean varieties to be transformed by *A. tumefaciens*, using a Glyphosate tolerance expression cassette with FTO in Colombian territory, as a stage to produce and commercialize generic GM crops in Colombia.

Materials and methods

Plant genotypes

Three Colombian genotypes of soybean were used in transformation experiments: SK7 and P29 (bred by Kamerun and Panorama companies respectively) and Soyica P-34 (bred by the Colombian Agricultural Institute - ICA). These genotypes were chosen due to its high cultivation rate in the Eastern Plains region. SK7 genotype is adapted to grow between 300 and 1,200 m a.s.l., a vegetative stage of 110-112 d, height of 101 cm, white flowers, brown pubescence, oil content of 20.47%, and a yield rate of 2,600 kg ha⁻¹. P29

genotype is adapted to grow between 300 and 1,200 m a.s.l. as well, has a vegetative stage of 105-115 d, height of 100 cm, purple flowers, brown pubescence, an oil content of 21.37%, and a yield rate of 2,600 kg ha⁻¹. Soyica P-34 is equally adapted to the same altitude (300 to 1,200 m a.s.l.), a vegetative stage of 110 d, height of 69 cm, white flowers, brown pubescence, an oil content of 20%, and a yield rate of 2,700 kg ha⁻¹. As start material field conditions seed were used, without any fungicidal or insecticidal treatment, and a moisture between 11 to 12%. These varieties were chosen because they were the most demanded by farmers at the time in which the experiment was performed.

Agrobacterium strains, expression cassette and vector

A previously reported cassette designated as E-IGP was used; this cassette contains a polyubiquitin promoter from soybean (GmUbi), followed by a transit peptide from *Petunia hybrida*, a codon optimized *cp4 epsps* gene for expression in soybean tissues, and a nopaline synthase (NOS) terminator codon (Jiménez, 2014). The E-IGP cassette was introduced into a pCambia1301 vector on which the GUS reporter gene and the hygromycin selection gene were excised in such a way that only the E-IGP cassette could be transferred to the plant genome (Jiménez, 2014). The vector containing the E-IGP cassette was introduced into *A. tumefaciens* strains AGL0 (bought from an American Type Culture Collection, under ATCC®BAA-100™ denomination) and EHA105 (acquired by a donation from the Cenicaña institution). Recombinant strains were maintained in Luria Bertani (LB) medium containing 50 mg L⁻¹ kanamycin. *Agrobacterium* cultures used for infection of explants were grown in LB medium. The observed optical density at 650 nm (OD₆₅₀) ranged from 0.8 to 1.0, at 28°C and 200 rpm. A bacterial pellet was obtained by centrifugation of 30 ml of bacterial culture at 8000 rpm for 4 min at 20°C. The pellet was resuspended in 25 ml of co-cultivation liquid medium (CCLM) (1X Gamborg vitamins, 0.1X B5 salts (Gamborg *et al.*, 1968), 1.67 mg L⁻¹ benzylaminopurine (BAP), 0.25 mg L⁻¹ gibberellic acid (GA₃), 3% sucrose, 20 mM 2-[N-Morpholino] ethanesulfonic acid (MES), 200 µ macetosyringone, pH 5.7) and then was used as inoculum for the plant tissues.

Explant preparation and *A. tumefaciens* infection

Soybean seeds were selected considering their appearance, choosing those that did not have lacerations or spots, and had a homogeneous size. The seeds surface was sterilized following the chlorine gas technique (Paz *et al.*, 2004; Paz *et al.*, 2006; Song *et al.*, 2013) for 16 h, generating gas from a mix of 4.1 ml 10 N HCl with 100 ml 5% NaClO. Sterilized seeds were germinated on 0.7% Plant Tissue Culture (PTC)

agar medium (water plus agar), pH 5.7, with the hilum proximal to the media for 5 d, and incubated under a 16/8 (light/dark) photoperiod at 26°C.

Once seeds were germinated, the seed coat was eliminated and a cutting was done 5 mm below the cotyledons junction to eliminate the hypocotyl. After that, cotyledons were separated by a longitudinal cut on the remaining piece of hypocotyl. Plumule was eliminated from both cotyledons, and some incisions (7-12) were made on the cotyledonary node. Each cotyledon with its own cotyledonary node was considered as an explant to transformation. Explants were infected with *A. tumefaciens* strains by submerging them in a CCLM solution containing bacterial biomass for 30 min, followed by cultivation on a co-cultivation medium (CCM) (CCLM added with 0.7% PTC agar), with the ad-axial side down, and incubated in the dark for 3 d at 28°C (Zhang *et al.*, 1999).

Regeneration test of soybean varieties

A first trial was performed to observe the behavior of soybean varieties in an *in vitro* system intended for *Agrobacterium*-mediated transformation. For such task, a regeneration ability assay was carried out for every variety without *Agrobacterium* infection. In this essay, 200 seeds of each variety were selected and germinated obtaining 240 explants that were prepared as described above. These explants were cultivated on CCM without *Agrobacterium* strain for 3 d, and later were cultivated on a shoot induction medium (SIM) (1X B5 salts, 1X Gamborg vitamins (Gamborg *et al.*, 1968), 3% sucrose, 1.67 mg L⁻¹ BAP, 3mM MES, 0.7% PTC agar, pH 5.7) for 4 weeks, with a medium replacement at the end of the second week.

After four weeks on SIM, the number of explants producing at least one shoot (regenerating explants) was recorded as well as the number of shoots produced by each regenerating explant.

Transformation, regeneration and selection

Transformation assays were performed following the methodology described by Zhang *et al.* (1999), plus some modifications. To assess the ability of each vegetal variety to be transformed with E-IGP cassette, the assay was divided in two treatments and two controls as follows: Treatment 1: three varieties transformed with an EHA105 strain containing a E-IGP cassette, and selected *in vitro* using the herbicide Glyphosate; Treatment 2: three varieties transformed with an AGL0 strain containing a E-IGP cassette, and selected *in vitro* using the herbicide Glyphosate; Control 1 (relative control): three varieties without transformation,

and selected *in vitro* using Glyphosate; and Control 2 (absolute control): three varieties without transformation, and without *in vitro* selection.

For each treatment and control, 130 explants of each variety as described above were prepared. Treatments were inoculated with *Agrobacterium* strains and co-cultivated as described above. Controls were cultivated on CCM in the same way as treatments, but without bacteria. After co-cultivation, all explants were rinsed in sterile water added with 50 mg L⁻¹ cabenecillin, on a rotary shaker at 410 rpm for 40 min, three times. After this rinse, the explants were transferred to SIM mixed with antibiotics (250 mg L⁻¹ cefotaxime, 100 mg L⁻¹ timentin), and they were incubated for two weeks, under a 16/8 (light/dark) photoperiod at 26°C. After this procedure, SIM was renewed with a fresh SIM plus antibiotics solution and mixed with a 148 µM Glyphosate reagent grade (Phytotechnology Laboratories®, Lenexa, KS, USA). This methodology was followed in Treatments 1, 2 and in Control 1; Glyphosate was not added in Control 2. During the process to transfer explants to fresh SIM, the remaining hypocotyl of each explant was cut to allow the fresh tissue to directly contact the growth medium. Growing process was performed for two additional weeks under the same conditions.

Once the shoot induction period was finished, it was followed by a shoot elongation period (SEP). All explants were cut to eliminate remaining cotyledon and thus allowing fresh tissue to be in contact with the medium. Explants developing at least one shoot were transferred to Shoot Elongation Medium (SEM) (1X B5 salts, 1X Gamborg vitamins (Gamborg *et al.*, 1968), 3% sucrose, 0.5 mg L⁻¹ GA₃, 0.1 mg L⁻¹ indole acetic acid (IAA), 0.7 mg L⁻¹ BAP, 50 mg L⁻¹ glutamine, 50 mg L⁻¹ asparagine, 3 mM MES, 250 mg L⁻¹ cefotaxime, 100 mg L⁻¹ timentin, 0.7% PTC agar, pH 5.7) added with a 35 µM Glyphosate reagent grade (Phytotechnology Laboratories®, Lenexa, KS, USA) in Treatments 1, 2 and in Control 1, and without Glyphosate in Control 2. The SEM solution was replaced in the explants every two weeks for fresh SEM. In treatments and control containing Glyphosate, the herbicide was added during four weeks, removing it between the fifth week and until the end of SEP. The explants were allowed to grow until shoots reached a height of 3 cm, for a maximum SEP of 10 weeks, under 16/8 (light/dark) photoperiod to 26°C.

Shoots that reached the required height (3 cm) were individualized, labeled as “KJ” for SK7 variety, “PJ” for P29 variety and “SJ” for Soyica P34 variety, and marked by a number to indicate a consecutive individualization, thus discriminating the lines obtained from different treatments. Each

individual line was transferred to a Propagation Medium (PM) (MS salts 0.66X (Murashige and Skoog, 1962), (1X Gamborg vitamins (Gamborg *et al.*, 1968), 3% sucrose, 0.7% PTC agar, pH 5.7), and it was propagated to obtain biomass to consume in molecular and phenotypical analysis.

Molecular and phenotypical analysis

Individualized lines that were successfully propagated were subjected to the polymerase chain reaction (PCR) analysis to detect transgene insertion. Genomic DNA was extracted from each line following CTAB buffer methodology (Doyle, 1991), and it was quantified spectrophotometrically using NanoDrop equipment (Thermo Fisher Scientific, Waltham, MA USA). A 205 bp region of transgene was amplified using the pair of primers 5'-CTTTGCTGAA-GGAGCTACCG-3' and 5'-GTGATCGAGATGCGTAG-CAA-3' along with reagents included in Kapa3G Plant PCR kit (Sigma-Aldrich Corp. St. Louis, MO, USA) following manufacturer's instructions. PCR products were separated by electrophoresis on a 1% agarose gel. Transformation frequency was calculated as number of positive PCR lines / total number of transformed explants × 100.

To detect EPSPS expression, the vegetal biomass of propagated lines was subjected to the commercial Enzyme Linked Immunosorbent Assay (ELISA) Roundup Ready CP4 EPSPS (Agdia Inc. Elkhart, IN, USA), following manufacturer's instructions. Absorbance of each sample was measured using iMark™ Microplate Reader (Biorad, Hercules, CA, USA) at 655 nm wavelength. Percentage of functional transformants was calculated as number of positive ELISA lines / total number of transformed explants × 100.

On each positive PCR sample, a Southern Blotting assay was performed following recommendations of digoxigenin (DIG) applications platform for filter hybridization (Sigma-Aldrich Corp. St. Louis, MO, USA) (Eisel *et al.*, 2008). A DIG-labeled probe was synthesized using the PCR DIG Probe Synthesis kit (Sigma-Aldrich Corp., St. Louis, MO, USA) with the same primers described above for PCR. Twenty µg of genomic DNA were digested with *PvuII* and *NdeI* enzymes (New England Biolabs® Inc., Ipswich, MS, USA) in a double digest reaction. Restriction fragments were separated on 0.7% agarose gel by electrophoresis and transferred onto a positively charged nylon membrane (Sigma-Aldrich Corp., St. Louis, MO, USA). Prehybridization, hybridization, membrane washing, and detection were conducted following the platform instruction manual (Eisel *et al.*, 2008). The detection of probe-target hybrids was done by chemiluminescence using CDP-Star substrate (Sigma-Aldrich Corp., St. Louis, MO, USA).

Only a small amount of possible transgenic lines were allowed to grow on soil-type substrate. On hardened lines, a “plant painting” was performed using a 0.5% dilution of commercial Glyphosate (Victorius® 48 SL, Sodiak SA, Bogota, Colombia), over the shoot of the plant. Herbicide affectation was recorded after 10 d based on plant survival. Parallel to application on transformed lines, the same Glyphosate solution (0.5%) was applied on a non-transformed plant as positive control for test herbicide activity. In contrast, solvent (tap water) was applied on another non-transformed plant as a negative control.

Statistical analysis

A completely randomized design was used in biological assays, non-parametric comparison tests were used by statistical differences determination using a *P* value of 0.05. R software (R Development Core Team, 2008) was applied for the calculation.

Results

Regeneration capacity of soybean varieties

Three Colombian soybean varieties were evaluated regarding their regeneration capacity as a preliminary clue of how their behavior on an *Agrobacterium*-mediated transformation system will be. Starting from 240 explants of each variety, 212 explants of Soyica P34 (88.3%), 181 explants of SK7 (75.4%) and 231 explants of P29 (96.2%) developed regenerated shoots. After counting the number of shoots per explant (on regenerative explants) in each variety, it was found that, on average, Soyica P34 regenerates 11.5 shoots per explant (± 3.9 shoots), SK7 regenerates 17.7 shoots per explant (± 7.73 shoots) and P29 regenerates 12.7 shoots per explant (± 5.2 shoots) (Fig.1).

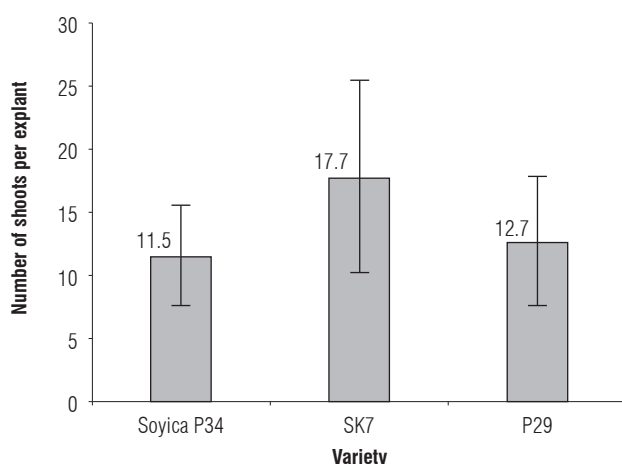


FIGURE 1. Regeneration capacity of vegetal varieties as number of shoots per explant.

At first the non-parametric Kruskal-Wallis test was applied using R software, to compare if varieties outputs have equality in the middle range or if at least one of them is different (Hollander *et al.*, 2014). In statistical terms, some of the following hypothesis systems were assessed:

$$H_0: \mu_{\text{Soyica P34}} = \mu_{\text{SK7}} = \mu_{\text{P29}} \quad \text{vs} \quad H_a: \mu_{\text{Soyica P34}} \neq \mu_{\text{SK7}} = \mu_{\text{P29}}$$

$$H_0: \mu_{\text{Soyica P34}} = \mu_{\text{SK7}} = \mu_{\text{P29}} \quad \text{vs} \quad H_a: \mu_{\text{Soyica P34}} = \mu_{\text{SK7}} \neq \mu_{\text{P29}}$$

$$H_0: \mu_{\text{Soyica P34}} = \mu_{\text{SK7}} = \mu_{\text{P29}} \quad \text{vs} \quad H_a: \mu_{\text{Soyica P34}} \neq \mu_{\text{P29}} = \mu_{\text{SK7}}$$

$$H_0: \mu_{\text{Soyica P34}} = \mu_{\text{SK7}} = \mu_{\text{P29}} \quad \text{vs} \quad H_a: \mu_{\text{Soyica P34}} \neq \mu_{\text{SK7}} \neq \mu_{\text{P29}}$$

When the test was carried out, a *P*-value of 0 was obtained, and the null hypothesis was rejected, suggesting that at least one of these average ranges is different from the others. Therefore, the non-parametric Wilcoxon test was used, in order to compare the differences on the average ranges of two samples, and to identify which is higher (Hollander *et al.*, 2014). The test was carried out two by two, with $\alpha = 0.1$, as follows:

$$H_1: \mu_{\text{Soyica P34}} = \mu_{\text{SK7}} \quad \text{vs} \quad H_a: \mu_{\text{Soyica P34}} < \mu_{\text{SK7}}$$

$$H_2: \mu_{\text{P29}} = \mu_{\text{Soyica P34}} \quad \text{vs} \quad H_a: \mu_{\text{P29}} < \mu_{\text{Soyica P34}}$$

$$H_3: \mu_{\text{P29}} = \mu_{\text{SK7}} \quad \text{vs} \quad H_a: \mu_{\text{P29}} < \mu_{\text{SK7}}$$

The *p*-values associated with the tests performed were 0.00 for *H*₁, 0.99 for *H*₂ and 0.00 for *H*₃. For any *p*-value smaller than α , the null hypothesis is rejected, indicating that the mean range of shoots of the SK7 variety is greater than the mean range of the Soyica P34 and P29 varieties.

Agrobacterium-mediated transformation

The three Colombian soybean varieties described above were subjected to a genetic modification. Such methodology consisted in an insertion of a transgene designed to express a CP4 EPSPS protein to confer tolerance to the herbicide Glyphosate. This process was performed using the *A. tumefaciens* bacteria as a transgene's vehicle, assessing two bacterial strains, AGL0 and EHA105.

To assess the behavior of treatments and controls in each variety, the explant number in each process stage prior line individualization was recorded. Each treatment and control had initially 130 explants, and this number was decreasing in successive stages (Fig. 2). It was observed that in all three plant varieties, addition of Glyphosate to the culture medium reduced the explants regeneration compared to the control without herbicide addition. Taking as a reference the explants subjected to absolute control that reached the shoot elongation 3 stage (SE-3) in each

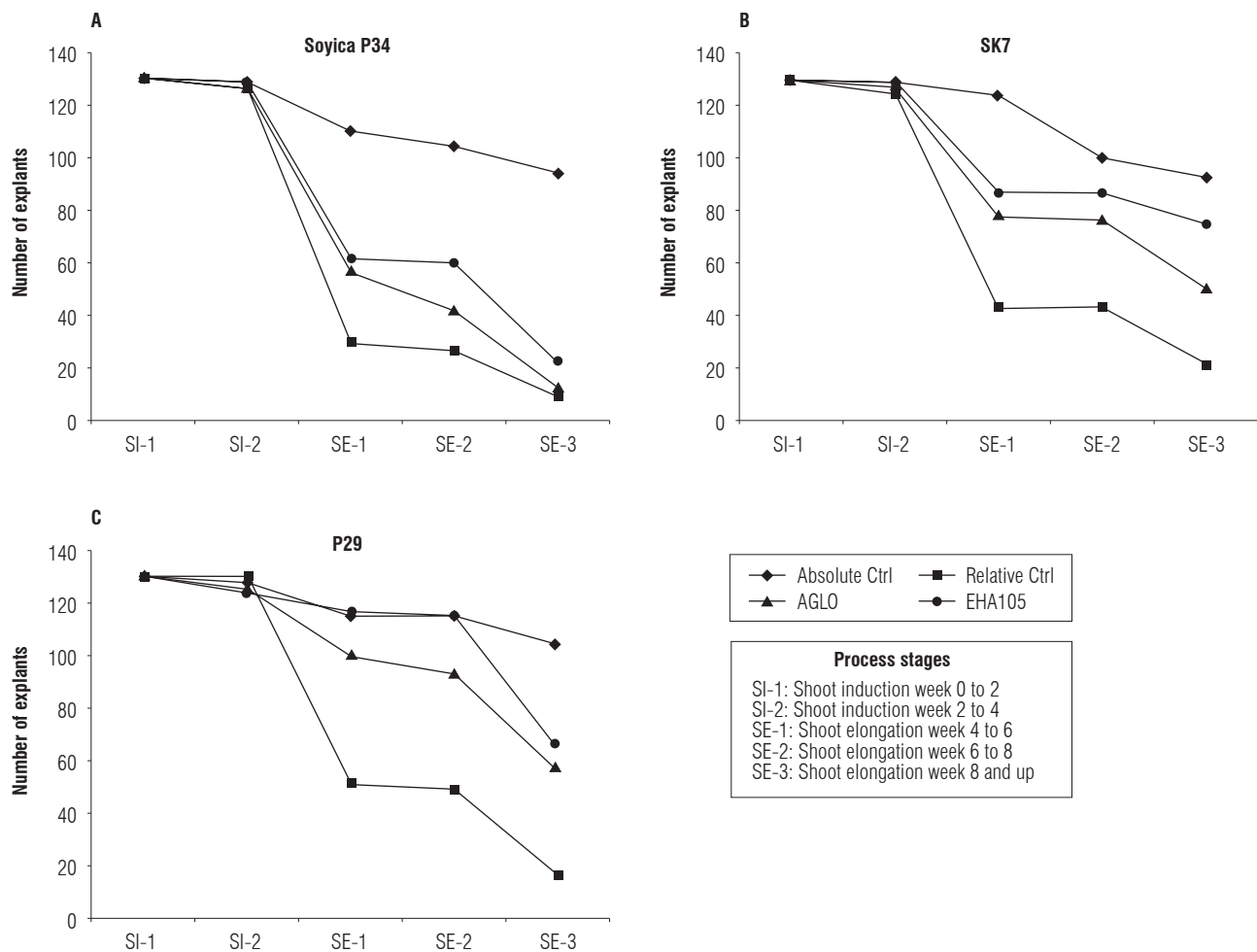


FIGURE 2. Recovered explants per variety on different stages of transformation process.

variety (100%), it was observed that the highest damage by herbicide was presented in Soyica P34. For this variety, 8.5% of explants in relative control, 12.8% in transformation with strain AGL0 and 22.3% in transformation with strain EHA105 reached the SE-3 stage. Furthermore, for P29 16% of explants in relative control, 55.8% in transformation with strain AGL0 and 64.4% in transformation with strain EHA105 reached the SE-3 stage. The most tolerant variety to herbicide addition was SK7, in which 23.7% of explants in relative control, 54.8% in transformation with AGL0 and 80.6% in transformation with strain EHA105 reached SE-3 stage (Fig. 2).

After reaching the SE-3 stage, the shoots that grew above 3 cm were labeled and marked. These Lines were propagated, and only those that were successful were subjected to molecular and phenotypical analysis. A total of 91 possible primary transformants of all varieties and treatments were labeled (Tab. 1), but only 53 lines were effectively propagated, so DNA extraction was performed exclusively

on those 53 lines. PCR was done on extracted DNA and separated by electrophoresis (Fig. 3).

SK7 showed the highest transformation frequency (10.8%) when it was combined with AGL0 strain (Tab. 1). P29 was also efficiently transformed in a lower proportion, (5.4%) when it was combined with AGL0 strain, and 6.1% when it was combined with EHA105 strain. It was not possible to transform Soyica P34 variety under the procedures followed in this research (Tab. 1).

To assess the protein expression, an ELISA test on biomass of the propagated plants from possible primary transformants was performed. In this test positive results by 12 lines of SK7 and P29 varieties were obtained (Tab. 1). Thus, the highest percentage of functional transformants was obtained for SK7 when it was combined with AGL0 (3.8%). For P29 variety, the highest percentage of functional transformants was obtained also by using AGL0 strain (2.3%) (Tab. 1).

TABLE 1. Results of transgene presence / absence test and CP4 EPSPS protein detection on generated lines.

| Plant genotype | Bacterial strain | Primary transformants | PCR (+) | ELISA (+) | Transformation frequency (PCR (+)) | Functional transformants (ELISA (+)) |
|----------------|------------------|-----------------------|---------|-----------|------------------------------------|--------------------------------------|
| SK7 | AGL0 | 22 | 14 | 5 | 10.8% | 3.8% |
| | EHA105 | 31 | 6 | 3 | 4.6% | 2.3% |
| P29 | AGL0 | 14 | 7 | 3 | 5.4% | 2.3% |
| | EHA105 | 23 | 8 | 1 | 6.1% | 0.8% |
| Soyica P34 | AGL0 | 1 | 0 | 0 | 0% | 0% |
| | EHA105 | 0 | 0 | 0 | 0% | 0% |



FIGURE 3. Detection of the E-IGP cassette on possible primary transformants. The name of each line is indicated at the top of each lane. PCtrl: Non transformed P29 variety, KCtrl: Non transformed SK7 variety, H2O: absolute control with primers, Ctrl (+): positive control, corresponding to plasmid vector E-IGP extracted by miniprep. The molecular weight marker corresponds to 50 bp DNA Ladder (New England Biolabs).

A Southern blot was performed on positive PCR samples. All of them showed a single insertion band, ranging between 6.5 to 24 kbp at a different size in every DNA sample (Fig. 4).

Some lines allow their hardening on soil-type substrate, supplying material to perform some greenhouse phenotypical tests. After applying a 0.5% solution of the commercial Glyphosate dose in a time period of 10 d, tolerance was assumed as the survival of treated plants. In contrast, susceptibility was considered as death of treated plants. The KJ7, KJ8, KJ15, KJ18 and PJ32 lines were found to effectively tolerate the herbicide, as there was no plant death after 10 d (Fig. 5C), whereas lines KJ24, KJ47 and SJ1 did not tolerate the herbicide application and died (Fig. 5D).

Discussion

Transformation efficiency in soybean using *A. tumefaciens* is, in general terms low, mainly due to difficulties in the T-DNA transfer efficiency. To obtain transformed plants in regeneration systems, it is necessary to use hypervirulent

strains of *Agrobacterium* (Atif *et al.*, 2013). In the present study, three soybean varieties adapted to cropping conditions in Colombian territory were used, and considering that it is the first research carried out in the country on soybean genetic transformation, it was absolutely necessary to establish a transformation platform for this species, evaluating diverse factors that could affect the process, as the predisposition of local varieties to be genetically transformed.

The transformation efficiency in soybean is highly genotype-dependent (Atif *et al.*, 2013), which may be linked to its regeneration capacity, among other factors. Paz *et al.* (2004) reported that Williams variety had a higher rate of regeneration within a set of 10 different soybean varieties with 100% regeneration. Within this set, the lowest rate of regeneration was obtained by the Harosoy variety, with 68%. In this study, the selected plant genotypes had different performances in each of the developed tests. In evaluation of regeneration capacity, it was observed that the P29 variety had the highest percentage of regeneration (96.2%), while SK7 had the lowest percentage (75.4%).

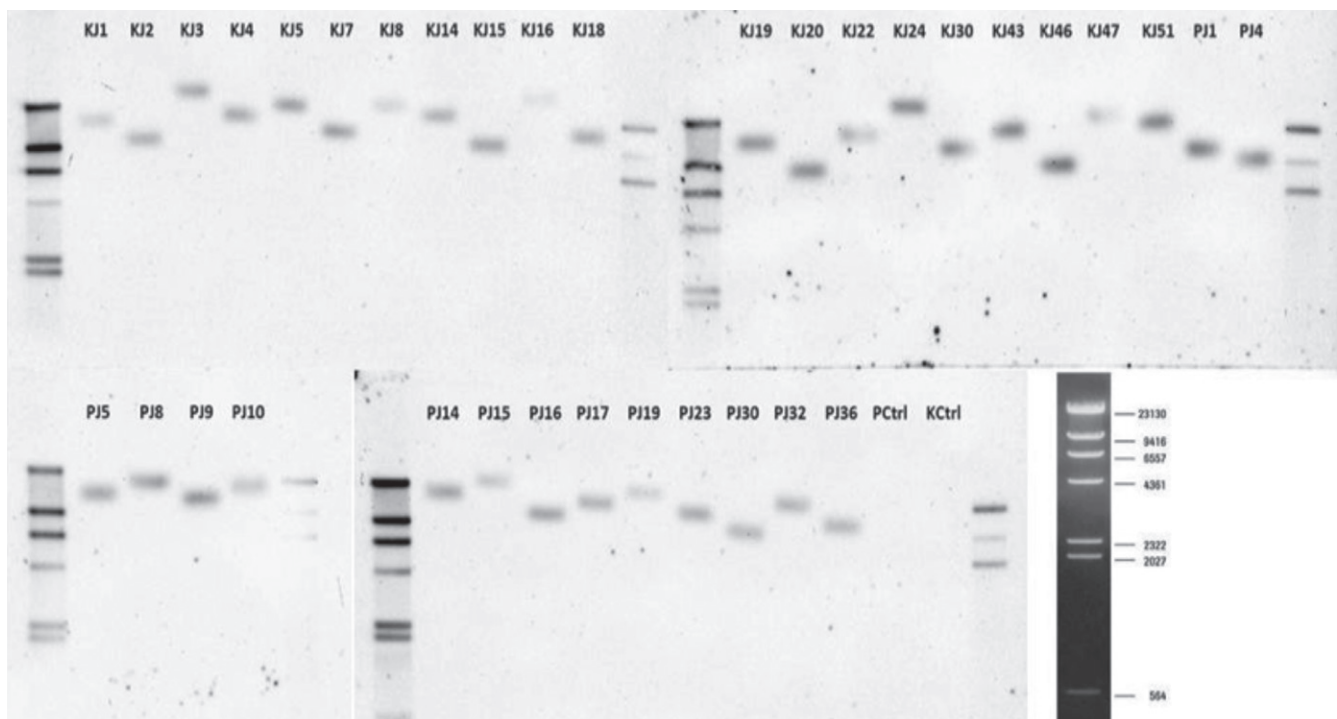


FIGURE 4. Southern blotting on positive PCR samples. The name of each transformant is indicated on the top of each lane. PCtrl: Non transformed P29 variety, KCtrl: Non transformed SK7 variety. Each membrane has, on its left side, a molecular marker, corresponding to DNA Molecular Weight Marker II, DIG-labeled (Sigma-Aldrich Corp. St. Louis, MO, USA), and, on its right side, a positive control, corresponding to E-IGP cassette integrated into pCambia1301 non-linearized vector.

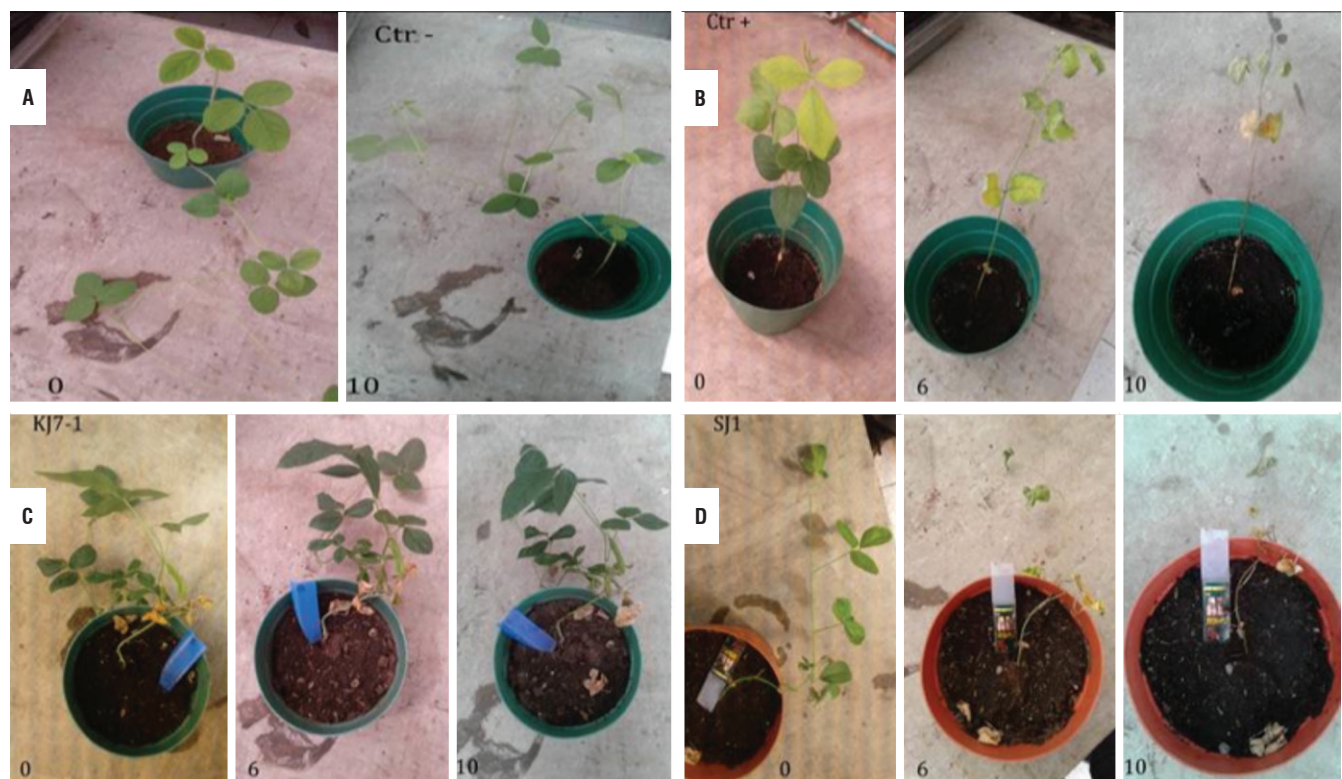


FIGURE 5. Greenhouse test for Glyphosate tolerance. A. Negative control, corresponding to a non-transformed plant painted with the herbicide solvent (tap water). B. Positive control, corresponding to a non-transformed plant painted with 0.5% commercial Glyphosate. C. Tolerant line, corresponding to KJ7 line. D. Susceptible line, corresponding to SJ1 line. C and D were painted with 0.5% commercial Glyphosate. Labels regarding the name of the lines are located at the top of each picture. The next day after the application event is located at the bottom of each picture.

However, SK7 compensates its low regeneration percentage with a high rate of shoot production per explant, with 17.7 shoots on average, being the highest value among the evaluated genotypes. In general, regeneration of the three plant varieties was good with the tested system, considering that percentage values of explants with shoots and number of shoots per explant are adequate to take as baseline on a process of genetic transformation.

Regeneration differences between genotypes could be related to the balance of growth regulators present in SIM and its interaction with tissues and endogenous factors of each variety. Concentration of growth regulators must be optimized for each variety separately (Paz *et al.*, 2006; Soto *et al.*, 2013; Li *et al.*, 2017).

It is important to include a selector agent in SIM to avoid the generation of chimerisms, or escapes from the transformation system. Most soybean transformation trials follow an in-vitro and greenhouse selection using glufosinate ammonium (Zhang *et al.*, 1999; Paz *et al.*, 2004; Zeng *et al.*, 2004; Paz *et al.*, 2006; Xue *et al.*, 2006; Song *et al.*, 2013; Jia *et al.*, 2015; Yang *et al.*, 2016; Li *et al.*, 2017), Kanamycin (Liu *et al.*, 2004) or Hygromycin (Olhoft *et al.*, 2003; Arun *et al.*, 2015; Kuma *et al.*, 2015). There are fewer reports on Glyphosate selection (Clemente *et al.*, 2000; Soto *et al.*, 2016). In this report, a selection scheme was used to produce adequate plants further individualized and propagated as primary transformants. It is important to emphasize that Glyphosate selection is based on phenotypical criteria (shoot height and appearance) instead of the categorical criteria used for other selector agents as live/dead shoots.

The maximum transformation frequencies obtained in other reports, with different soybean varieties, are generally ranging between 4 and 15.8%, with an average of 8.5% (Olhoft *et al.*, 2003; Liu *et al.*, 2004; Paz *et al.*, 2004; Zeng *et al.*, 2004; Paz *et al.*, 2006; Yukawa *et al.*, 2008; Jia *et al.*, 2015; Yang *et al.*, 2016; Li *et al.*, 2017). In this report, a maximum transformation frequency on SK7 variety of 10.8% was obtained, which is above the average, reported for other varieties. P29 variety was also transformed with a frequency of 6.1%. This indicates that there are Colombian soybean varieties with potential to be genetically transformed, in this case, with special success on SK7 variety which had the highest regeneration capacity and also the highest transformation frequency.

Differences between varieties regarding their capacity to be transformed can be associated with defensive responses of plant tissues against bacterial infection and the capacity of cellular division, which varies between plant genotypes

(Jia *et al.*, 2015). There was a slight difference between both bacterial strains in relation to their infective capacity, showing AGL0 a higher infective ability than EHA105 on SK7. Both strains are derivatives from EHA101 strain, which in turn is derived from hypervirulent A281 strain, with a C58 chromosomal background (a nopaline type strain) (Lazo *et al.*, 1991; Hood *et al.*, 1993). So, considering that both strains have the same chromosomal background and vir helper vector, similar results are expected.

Recently, genome editing has been developed by addition, removal, or alteration on specific bases in the plant genome. A relevant technology of genome edition is CRISPR-Cas9 (clustered regularly interspaced short palindromic repeats associated to protein Cas9). The CRISPR-Cas9 technology has their origin in a bacterial defense system from viruses using RNA segments to target viral DNA, so as a result, Cas9 protein cut the parasite DNA. Basically, CRISPR-Cas9 is based on a guide RNA segment that binds to a specific sequence of DNA, using a Cas9 enzyme. This enzyme cuts the target DNA and repairs the cell machinery to add or remove the desired sequences. CRISPR-Cas9 system leaves no traces on the genome, which has been postulated as a form of genome modification different to transgenesis (Hussain *et al.*, 2018). This affair was assessed by the European Union on July 25, 2018 claiming that genome edition should be subject to the same regulations as the conventional GM crops. CRISPR-Cas9 system has been applied in soybean by stress tolerance (edition of Sucrose non-fermenting related protein kinases), multiple loci edition, flowering delay (by edition of GmFT2a gene in a specific base or short deletion), and promoter edition with relevance in expression, among others (Du *et al.*, 2016; Cai *et al.*, 2018; Li *et al.*, 2018; Kanazashi *et al.*, 2018).

This work is a preliminary approach to establish a stable transformation platform for crops of economic relevance such as soybean in Colombia. It was possible to produce entire plants with evident Glyphosate tolerance in greenhouse conditions, but the process should be subjected to further optimization to get a totally refined platform. This is an important step to initiate the production of generic transgenics in this country, specifically designed to develop Glyphosate tolerance as a transgenic trait in soybean varieties, while the patents protecting this technology expired in recent years (Jefferson *et al.*, 2015).

Acknowledgments

The authors would like to thank S. Leal, Y. Rodríguez and J. Mora for their technical assistance and to Kamerun and Panorama seed companies for their collaboration

as seed donors. This work was funded by the Colombian National Federation of Cereals and Legumes Farmers (Fenalce - Research and Transfer Program of National Fund for Soybean). The authors express their gratitude to Fenalce for its commitment to develop science and technology in our country. Rojas A.C. express her gratefulness to the National Department of Science and Technology (Colciencias, Call for Scholarships No. 528/2011) for her Ph.D. fellowship.

Literature cited

- Arun, M., K. Subramanyam, T.S. Mariashibu, J. Thebora, G. Shivanandhan, M. Manickavasagam, and A. Ganapathi. 2015. Application of sonication in combination with vacuum infiltration enhances the *Agrobacterium*-mediated genetic transformation in Indian soybean cultivars. *Appl. Biochem. Biotechnol.* 175, 2266-2287. Doi: 10.1007/s12010-014-1360-x
- Atif, R.M., E.M. Patat-Ochatt, L. Svabova, V. Ondrej, H. Klenotcova, L. Jacas, M. Griga, and S.J. Ochatt. 2013. Gene transfer in legumes. pp. 37-100. In: Lüttge, U., W. Beyschlag, D. Francis, and J. Cushman (eds.). *Progress in Botany* 74. Springer-Verlag, Berlin Heidelberg. Doi: 10.1007/978-3-642-30967-0_2
- Bonny, S. 2008. Genetically modified Glyphosate-tolerant soybean in the USA: adoption factors, impacts and prospects. A review. *Agron. Sustain. Dev.* 28, 21-32. Doi: 10.1051/agro:2007044
- Bourras, S., T. Rouxel, and M. Meyer. 2015. *Agrobacterium tumefaciens* gene transfer: how a plant pathogen hacks the nuclei of plant and nonplant organisms. *Phytopathol.* 105, 1288-1301. Doi: 10.1094/PHYTO-12-14-0380-RVW
- Cai, Y., L. Chen, X. Liu, C. Guo, S. Sun, C. Wu, B. Jiang, T. Han, and W. Hou. 2018. CRISPR/Cas9-mediated targeted mutagenesis of *GmFT2a* delays flowering time in soya bean. *Plant Biotechnol. J.* 16, 176-185. Doi: 10.1111/pbi.12758
- Clemente, T.E., B.J. LaVallee, A.R. Howe, D. Conner-Ward, R.J. Rozman, P.E. Hunter, D.L. Broyles, D.S. Kasten, and M.A. Hinchee. 2000. Progeny analysis of Glyphosate selected transgenic soybeans derived from *Agrobacterium*-mediated transformation. *Crop Sci.* 40, 797-803. Doi: 10.2135/cropsci2000.403797x
- Doyle, J. 1991. DNA protocols for plants. pp. 283-293. In: Hewitt G.M., A.W.B. Johnston, and J.P.W. Young (eds.). *Molecular techniques in taxonomy*. Vol 57. Springer, Berlin, Heidelberg. Doi: 10.1007/978-3-642-83962-7_18
- Du, H., X. Zeng, M. Zhao, X. Cui, Q. Wang, H. Yang, H. Cheng, and D. Yu. 2016. Efficient targeted mutagenesis in soybean by TALENs and CRISPR/Cas9. *J. Biotechnol.* 217, 90-97. Doi: 10.1016/j.jbiotec.2015.11.005
- Duke, S.O. and A.L. Cerdeira. 2010. Transgenic crops for herbicide resistance. pp. 133-166. In: Kole C., C.H. Michler, A.G. Abbott, and T.C. Hall (eds.). *Transgenic crop plants volume 2: Utilization and biosafety*. Springer, Heidelberg, Germany. Doi: 10.1007/978-3-642-04812-8_3
- Duke, S.O. and S.B. Powles. 2008. Glyphosate: a once-in-a-century herbicide. *Pest Manag. Sci.* 64, 319-325. Doi: 10.1002/ps.1518
- Eisel, D., O. Seth, S. Grünewald-Janho, B. Kruchen, and B. Rüger (eds.). 2008. *DIG Application Manual for Filter Hybridization*. Editorial Roche Diagnostics GmbH.
- Fenalce - Federación Nacional de Cultivadores de Cereales y Leguminosas. 2017. Situación del sector soya. Informe del Departamento de Información Económica y Estadística. URL: http://www.fenalce.org/nueva/plantillas/arch_web/Situacion_del_Sector_II_Trimestre_2017_-_Soya.pdf (accessed August 2017).
- Gamborg, O.L., R.A. Miller, and K. Ojima. 1968. Nutrient requirements of suspension cultures of soybean root cells. *Exp. Cell Res.* 50, 151-158. Doi: 10.1016/0014-4827(68)90403-5
- Hood, E.E., S.B. Gelvin, L.S. Melchers, and A. Hoekema. 1993. New *Agrobacterium* helper plasmids for gene transfer to plants. *Transgenic Res.* 2, 208-218. Doi: 10.1007/BF01977351
- Hollander, M., D.A. Wolfe, and E. Chicken. 2014. *Nonparametric statistical methods*. John Wiley & Sons, Hoboken, New Jersey, USA. Doi: 10.1002/9781119196037
- Hussain, B., S.J. Lucas, and H. Budak. 2018. CRISPR/Cas9 in plants: at play in the genome and at work for crop improvement. *Brief. Funct. Genomics* 1-10. Doi: 10.1093/bfpg/ely016
- ICA - Instituto Colombiano Agropecuario. 2010. Resolución No 2404 del 19 de Julio de 2010, Por la cual se autoriza siembras comerciales de soya Roundup Ready® (MON-Ø4Ø32-6) para la subregión natural de la Orinoquía.
- ILSI Research Foundation. 2017. GM crop database. ILSI Research Foundation, Washington D.C. URL: <http://cera-gmc.org/gmcropdatabase> (accessed August 2017).
- ISAAA. 2016. Global status of commercialized Biotech/GM crops: 2016. ISAAA Brief No. 52. Ithaca, NY, USA.
- Jefferson, D.J., G.D. Graff, C.L. Chi-Ham, and A.B. Bennett. 2015. The emergence of agbiogenics. *Nature Biotechnol.* 33, 819-823. Doi: 10.1038/nbt.3306
- Jia, Y., X. Yao, M. Zhao, Q. Zhao, Y. Du, C. Yu, and F. Xie. 2015. Comparison of soybean transformation efficiency and plant factors affecting transformation during the *Agrobacterium* infection process. *Int. J. Mol. Sci.* 16, 18522-18543. Doi: 10.3390/ijms160818522
- Jiménez, J.P. 2014. Diseño de genes semi-sintéticos que confieran tolerancia a herbicidas en soya. MSc. thesis. Universidad Nacional de Colombia, Bogota.
- Kanazashi, Y., A. Hirose, I. Takahashi, M. Mikami, M. Endo, S. Hirose, S. Toki, A. Kaga, K. Naito, M. Ishimoto, J. Abe, and T. Yamada. 2018. Simultaneous site-directed mutagenesis of duplicated loci in soybean using a single guide RNA. *Plant Cell Rep.* 37, 553-563. Doi: 10.1007/s00299-018-2251-3
- Klümper, W. and M. Qaim. 2014. A meta-analysis of the impacts of genetically modified crops. *PLOS One.* 9, e111629. Doi: 10.1371/journal.pone.0111629
- Kuma, K.M., V.S. Lopes-Caitar, C.C.T. Romero, S.M.H. Silva, M.K. Kuwahara, M.C.C.G. Carvalho, R.V. Abdelnoor, W.P. Dias, and F.C. Marcelino-Guimarães. 2015. A high efficient protocol for soybean root transformation by *Agrobacterium rhizogenes* and most stable reference genes for RT-qPCR analysis. *Plant Cell Rep.* 34, 1987-2000. Doi: 10.1007/s00299-015-1845-2
- Lazo, G.R., P.A. Stein, and R.A. Ludwig. 1991. A DNA Transformation-competent *Arabidopsis* genomic library in *Agrobacterium*. *Nature Biotechnol.* 9, 963-967. Doi: 10.1038/nbt1091-963
- Li, H.Q., C. Chen, R.R. Chen, X.W. Song, J.N. Li, Y.M. Zhu, and X.D. Ding. 2018. Preliminary analysis of the role of GmSnRK1.1 and GmSnRK1.2 in the ABA and alkaline stress response of the

- soybean using the CRISPR/Cas9-based gene double-knockout system. *Yi Chuan* 40, 496-507. Doi: 10.16288/j.ycz.17-424
- Li, S., Y. Cong, Y. Liu, T. Wang, Q. Shuai, N. Chen, J. Gai, and Y. Li. 2017. Optimization of *Agrobacterium*-mediated transformation in soybean. *Front. Plant Sci.* 8, 246. Doi: 10.3389/fpls.2017.00246
- Liu, Z., H. Xiong, H. Xie, Y. Tao Qin, Z. Ran, and B. Guo. 2013. A technique for *Agrobacterium*-mediated transformation via germinating seeds of soybean. *Adv. Mater. Res.* 749, 413-417. Doi: 10.4028/www.scientific.net/AMR.749.413
- Liu, H.K., C. Yang, and Z.M. Wei, 2004. Efficient *Agrobacterium tumefaciens*-mediated transformation of soybeans using an embryonic tip regeneration system. *Planta* 219, 1042-1049. Doi: 10.1007/s00425-004-1310-x
- Murashige, T. and F. Skoog. 1962. A revised medium for rapid growth and bio assays with tobacco tissue cultures. *Physiol. Plant.* 15, 473-497. Doi: 10.1111/j.1399-3054.1962.tb08052.x
- Olhoft, P.M., L.E. Flagel, C.M. Donovan, and D.A. Somers. 2003. Efficient soybean transformation using hygromycin B selection in the cotyledonary-node method. *Planta* 216, 723-735.
- Paz, M.M., H. Shou, Z. Guo, Z. Zhang, A.K. Banerjee, and K. Wang. 2004. Assessment of conditions affecting *Agrobacterium*-mediated soybean transformation using the cotyledonary node explant. *Euphytica* 136, 167-179. Doi: 10.1023/B:EUPH.0000030669.75809.dc
- Paz, M.M., J.C. Martinez, A.B. Kalvig, T.M. Fonger, and K. Wang. 2006. Improved cotyledonary node method using an alternative explant derived from mature seed for efficient *Agrobacterium*-mediated soybean transformation. *Plant Cell Rep.* 25, 206-213. Doi: 10.1007/s00299-005-0048-7
- R Development Core Team. 2008. R: A language and environment for statistical computing. R Foundation for Statistical Computing, Vienna. URL: <http://www.R-project.org>
- Song, Z., J. Tian, W. Fu, L. Li, L. Lu, L. Zhou, Z. Shan, G. Tang, and H. Shou. 2013. Screening Chinese soybean genotypes for *Agrobacterium*-mediated genetic transformation suitability. *J. Zhejiang Univ. Sci. B* 14, 289-298. Doi: 10.1631/jzus.B1200278
- Soto, N., C. Delgado, Y. Hernández, Y. Rosabal, A. Ferreira, M. Pujol, F. Aragão, and G.A. Enríquez. 2016. Efficient particle bombardment-mediated transformation of Cuban soybean (INCA-Soy-36) using Glyphosate as a selective agent. *Plant Cell Tissue Organ Cult.* 128, 187-196. Doi: 10.1007/s11240-016-1099-x
- Soto, N., A. Ferreira, C. Delgado, and G.A. Enríquez. 2013. *In vitro* regeneration of soybean plants of the Cuban Incasoy-36 variety. *Biotechnol. Aplicada* 30, 34-38.
- Tillmann, M. and S. West. 2004. Identification of genetically modified soybean seeds resistant to Glyphosate. *Sci. Agric.* 61, 336-341. Doi: 10.1590/S0103-90162004000300017
- USDA - United States Department of Agriculture. 2017. Oilseeds: world markets and trade. Informative circular from the foreign agricultural service. URL: <https://apps.fas.usda.gov/psdonline/circulars/oilseeds.pdf> (accessed August 2017).
- Valencia, R.A. and G.A. Ligarreto. 2010. Mejoramiento genético de la soya (*Glycinemax* [L.] Merrill) para su cultivo en la altillanura colombiana: una visión conceptual prospectiva. *Agron. Colomb.* 28, 155-163.
- Widholm, J.M., J.J. Finer, L.O. Vodkin, H.N. Trick, P. LaFayette, J. Li, and W. Parrott. 2010. Soybean. pp. 473-498. In: Kempken F. and C. Jung (eds.). Genetic modification of plants, biotechnology in agriculture and forestry. vol. 64, Springer-Verlag, Berlin, Heidelberg. Doi: 10.1007/978-3-642-02391-0_24
- Xue, R.G., H.F. Xie, and B. Zhang. 2006. A multi-needle-assisted transformation of soybean cotyledonary node cells. *Biotechnol. Lett.* 28, 1551-1557. Doi: 10.1007/s10529-006-9123-6
- Yang, X.F., X.Q. Yu, Z. Zhou, W.J. Ma, and G.X. Tang. 2016. A high-efficiency *Agrobacterium tumefaciens* mediated transformation system using cotyledonary node as explants in soybean (*Glycine max* L.). *Acta Physiol. Plant.* 38, 60. Doi: 10.1007/s11738-016-2081-2
- Yukawa, K., H. Kaku, H. Tanaka, Y. Koga-Ban, and M. Fukuda. 2008. Enhanced soybean infection by the legume "Super-Virulent" *Agrobacterium tumefaciens* strain KAT23. *Biosci. Biotechnol. Biochem.* 72, 1809-1816. Doi: 10.1271/bbb.80080
- Zeng, P., D.A. Vадnais, Z. Zhang, and J.C. Polacco. 2004. Refined glufosinate selection in *Agrobacterium*-mediated transformation of soybean [*Glycine max* (L.) Merrill]. *Plant Cell Rep.* 22, 478-482. Doi: 10.1007/s00299-003-0712-8
- Zhang, Z., A. Xing, P. Staswick, and T.E. Clemente. 1999. The use of glufosinate as a selective agent in *Agrobacterium*-mediated transformation of soybean. *Plant Cell Tissue Organ Cult.* 56, 37-46. Doi: 10.1023/A:1006298622969