Research Article

Establishment of protocol for genetic transformation of carnation with 1-aminocyclopropane-carboxylate deaminase (*acdS*) gene

Hui Yeong Jeong · Aung Htay Naing · Chang Kil Kim

Received: 13 June 2021 / Revised: 28 June 2021 / Accepted: 28 June 2021 © Korean Society for Plant Biotechnology

Abstract This study was conducted to develop an Agrobacterium-mediated genetic transformation protocol for the carnation cv. "Jinju" to counteract its ethylene sensitivity. The new protocol involves the use of an improved shoot regeneration medium, optimized minimal concentrations of the selective agent, a pre-culture period, and co-cultivation periods. Silver nanoparticles (NAg) added at a concentration of 2.0 µM to the Murashige and Skoog (MS) basal shoot regeneration medium supplemented with 0.1 mg/L indole-3butyric-acid (IBA) and 0.2 mg/L thidiazuron (TDZ) improved the shoot regeneration efficiency, number of shoots per explant, and plant growth compared to the control without the addition of NAg. The phosphinothricin (PPT) concentration of 1.0 mg/L was determined to be the minimal and optimal concentration for the selection of putative transgenic plants. When the explants were infected with Agrobacterium cells harboring the acdS gene, the explants that were pre-cultured for three days induced more putative transgenic plants than those that were co-cultivated for four days. Therefore, we expect that the results of this study will benefit researchers who are developing genetic transformations of carnations.

Keywords Carnation, Transformation, 1-aminocyclopropane-1-carboxylate deaminase, Ethylene, Vitrification

Introduction

Around the world, carnations are one of the three most popular cut flowers in floristry due to their high attractiveness to consumers. Currently, the market value of carnations is steadily increasing and is expected to reach 2,950

H. Y. Jeong · A. H. Naing · C. K. Kim (⊠) Department of Horticultural Science, Kyungpook National University, Daegu 41566, Korea e-mail: ckkim@knu.ac.kr million USD in 2024 (Naing et al. 2021). However, as carnation flowers are highly sensitive to ethylene, their postharvest flower quality is negatively affected by the ethylene that is produced during long-term transportation through domestic or global markets before arriving in consumers' hands, which means that low-quality flowers are being sold on the market. Low-quality cut flowers are poor market competitors for high-quality flowers, and this results in significant economic losses for florists and floriculturists. To overcome the problem, many attempts have been made to reduce the sensitivity of carnations to ethylene. Metabolic engineering, conventional breeding, and ethylene inhibitors are some of the approaches that have been used to reduce ethylene sensitivity. Although ethylene inhibitors have been proven to reduce ethylene production, they create environmental pollution and pose a risk to public health. Although the conventional breeding technique has long been used to produce novel cultivars, but it is difficult to obtain a cultivar with desirable traits due to the large number of genes that exist in its genome.

Plants produce ethylene due to the accumulation of the 1-aminocyclopropane-1-carboxylic acid (ACC) enzyme, which is an ethylene precursor found in plant tissues. ACC is converted into ethylene by ACC oxidase. The ACC deaminase enzyme present in plant growth-promoting bacteria (PGPB) is encoded by the *acdS* gene, and ACC deaminase can break down ACC into ammonia and a-ketobutyrate (Glick et al. 2014). A promising approach for reducing ethylene production is to break down ACC accumulation in plant tissues with overexpression of the acdS gene isolated from PGPB. Overexpression of the acdS gene via Agrobacterium-mediated genetic transformation has been demonstrated to reduce ACC accumulation and ethylene production in horticultural plants such as tomato, canola, Camelina sativa, and petunia (Grichko 2000; Grichko and Glick 2001; Heydarian et al. 2016; Klee 1991; Nie 2002; Sergeeva 2006; Stearns 2005). In their studies, transgenic

plants overexpressing the *acdS* gene produced lower levels of ethylene, which improved flower longevity, shelf life, and abiotic stress tolerance. However, genetic transformation of carnations is limited by a low shoot regeneration rate and other adverse factors, such as an inefficient pre-culture period and co-cultivation period. Thus, the genetic transformation process might impede the efficiency of genetic transformation efficiency (Naing et al. 2016). To date, the overexpression of the *acdS* gene has not been achieved in carnations. Therefore, it would be a breakthrough to develop a genetic transformation protocol for carnations that overexpresses the *acdS* gene if it produces an ethylene-resistant carnation cultivar.

In this study, we optimized factors that are involved in the *Agrobacterium*-mediated genetic transformation process to develop transgenic carnations overexpressing *acdS* to evaluate the effects on ethylene production and the quality of cut carnations.

Materials and Methods

Plant material

In a previous study, *in vitro* vitrified leaves exhibited superior shoot regeneration efficiency compared with normal leaves (Thu et al. 2020). In addition, utilization of *in vitro* vitrified carnation leaves as explants produced a transgenic carnation (Firoozabady et al. 1995). Therefore, in this study, induction of vitrified leaves was performed by culturing 4-week-old carnation shoot tips (cv. Jinju) in a Murashige and Skoog (MS) basal medium (pH 5.7) with 2% sucrose (LPS solution, Korea), 80 mg/L adenine hemisulfate (MB cell, Korea), and 85 mg/L sodium phosphate monobasic (Sigma, Germany). The culture bottles were placed at $25^{\circ}C \pm 2^{\circ}C$ under a 16 h photoperiod with 70 µmol m⁻² s⁻¹ light intensity. After four weeks of culture, vitrified leaves were used as materials for shoot regeneration and genetic transformation.

Effect of sodium nitroprusside (SNP) and sliver nanoparticles (NAgPs) on shoot regeneration

The combination of indole-3-butyric-acid (IBA 0.1 mg/L) (Duchefa, The Netherlands) and thidiazuron (TDZ 0.2 mg/L) (Duchefa, The Netherlands) induced a reasonable number of shoots per leaf explant (approximately 0.5 cm in size) in the carnation cv. Jinju (Thu et al. 2020). To verify whether the addition of sliver nanoparticles (Sigma-Aldrich, Germany) or sodium nitroprusside (Sigma-Aldrich, Germany) to the

shoot regeneration media would improve shoot regeneration efficiency and plant growth, different concentrations (0, 0.25, 0.5, 1.0, 1.5, 2.0 μ M sliver nanoparticle or 0, 5, 10, 15, 20, 30 μ M sodium nitroprusside) were added to the shoot regeneration media. Leaf segments (approximately 0.5 ~ 1.0 cm) excised from the 4-week-old *in vitro* vitrified leaves were cultured on regeneration media containing different concentration of nano silver particles or sodium nitroprusside. The plates were placed at the same culture condition as described above. One treatment contained ten explants, and there were three replicates for each treatment. Data on the percentage of shoot induction and number of shoots per explant were collected after four weeks of culture.

Optimization of the PPT concentration for the selection of putative transgenic explants

To reduce the risk of damage to putative transgenic plants during the selection period, different concentrations of the selective agent phosphinothricin (0, 0.3, 0.5, 1.0, and 1.5 mg/L PPT) were added to the shoot regeneration medium. Then, four-week-old *in vitro* vitrified leaves were segmented cut into $0.5 \sim 1.0$ cm pieces and used as sources of explants. The segments were cultured in shoot regeneration media containing PPT. Each treatment contained ten explants with three replications. The plates were placed in the same incubation room as described above. After four weeks of culture, the minimum concentration that suppressed shoot generation was determined and recorded.

Plasmid construction

Agrobacterium tumefaciens strains GV3101 harboring the plasmid pCB302-3 were used in this study. The plasmid was constructed with the *acdS* gene isolated from *Pseudomonas veronii*-KJ (Jung et al. 2018), in which the *acdS* gene was placed under the control of the cauliflower mosaic virus 35S (CaMV 35S) promoter, and the bar gene conferring PPT resistance was used as the selection marker.

Effect of the pre-culture period on shoot regeneration

Whole leaves (approximately 0.5 to 1.0 cm) were excised from the vitrified plants of 4-week-old *in vitro* plants. Explants were initially pre-cultured in the shoot regeneration medium for 1, 2, and 3 days under dark conditions. The shoot regeneration medium (pH 5.8) supplemented with 2.5% gelrite, 3% sucrose, 590 mg/L MES monohydrate (Duchefa, The Netherlands), 0.1 mg/L indole-3-butyric acid (Duchefa, The Netherlands), and 0.2 mg/L thidiazuron (TDZ; Duchefa, The Netherlands). After that, the plates were placed in the same incubation room as described above. Each treatment contained ten explants with three replications. After four weeks of culture, the percentage of shoot regeneration and number of shoots per explant were recorded.

Effect of the co-cultivation period on shoot regeneration from leaf explants inoculated with *Agrobacterium*

Based on the above preculture period result, the explants were pre-cultured for three days in the dark. Agrobacterium tumefaciens strains GV3101 were cultured for two days on solid Luria Broth containing antibiotics (50 mg/ml kanamycin and 25 mg/ml rifampicin) and were inoculated in liquid Luria Broth containing antibiotics (50 mg/ml kanamycin and 25 mg/ml rifampicin) for $15 \sim 16$ hours on a rotary shaker (at 28°C, 200 rpm, dark). The Agrobacterium suspension was pelleted using a centrifuge at 3,500 rpm for 20 minutes. The upper solution was removed using a pipette, and a liquid solution (pH 5.4) containing 1% sucrose, 1/2 MS, and 100 µM acetosyringone were used to reveal its effects on the transformation efficiency. Following this method, the pre-cultured explants were infected with Agro*bacterium* suspension, which was adjusted to an $OD_{600} =$ $0.6 \sim 0.7$ and incubated at 28°C on a rotary shaker at 50 rpm for 15 minutes. After inoculation, the explants were blotted dry on sterilized tissue and placed on the pretreatment medium again. To optimize the co-culture period, the infected leaf explants were plated onto the medium for 3, 4, or 5 days. Each treatment contained ten explants with three replicates. From pre-treatment to co-cultivation, the petri dishes were plated in the dark. After that, they were transferred to light condition. The plates were placed in the same incubation room as described above.

Regeneration of putative transgenic explants

After co-cultivation, explants were washed three times with distilled water (pH 5.8) containing 300 mg/L ticarcillin disodium/ clavulanate potassium (Duchefa, The Netherlands). After blotting dry using sterilized tissue paper, leaf segments were transferred on MS medium (pH 5.8) supplemented with 2.5% gelrite, 3% sucrose, 590 mg/L MES monohydrate, 0.1 mg/L IBA, 0.2 mg/L TDZ, 1.0 mg/L PPT, and 300 mg/L ticarcillin disodium/clavulanate potassium.

After three weeks, the shoots were regenerated and transferred to a new medium that had the same composition for another three weeks. If small shoots clusters had been formed, they were separated into $2 \sim 3$ sections. After another three weeks, the shoot clusters were harvested and moved to the rooting medium (pH5.7) supplemented with MS media, 2% sucrose, 80 mg/L adenine hemisulfate, 85 mg/L sodium phosphate monobasic, and 0.8% plant agar for three weeks.

To normalize the vitrified shoots, the elongated shoots were selected and transferred to a conversion medium (pH5.7) that has the same composition as the rooting medium with 1% plant agar instead of 0.8% for four weeks. Healthy normal shoots were harvested two to three times in the subcultures.

Statistical analysis

A statistical analysis was conducted using SPSS (version 25.0). All data were presented as mean and bar in the figure shown the standard deviation. ANOVA was conducted to show differences between groups by Duncan's multiple range test (DMRT, $p \le 0.05$). Each treatment was performed in triplicate.

Results and Discussion

Effect of sodium nitroprusside (SNP) and sliver nanoparticles (NAgPs) on shoot regeneration

The addition of different concentrations of SNP to the shoot regeneration medium could neither improve shoot regeneration and the number of shoots per explant nor shoot growth or size. Similar results were observed when different concentrations of NAgPs $(0.25 \sim 1.5 \mu M)$ were added to the shoot regeneration media, whereas the percentage of shoot regeneration observed in 1.5 µM NAgPs was significantly better than the control (Table 1). In contrast to other treatments, 2.0 µM NAgPs significantly improved the percentage of shoot regeneration and number of shoots per explant as well as plant growth and size compared to other treatments, including the control. Therefore, it can be ruled out that SNP is not applicable for improving the shoot regeneration of this cultivar, whereas 2.0 mg/L NAgPs were applicable. A positive effect of NAg was also observed in the in vitro shoot growth of Hosta capitata (Pe et al. 2020), whereas lower NAg concentrations did not promote plant growth or shoot regeneration, as was observed in this study, yet a higher NAg concentration promoted plant growth and shoot regeneration. In our previous study (Thu et al. 2020), the combination of 0.1 mg/L

		Shoot induction (%)	Mean number of shoots/explants
SNP (µM)	NAgPs (µM)		
-	-	70.00bcd	3.56ab
5	-	63.33cd	2.51b
10	-	63.33cd	2.50b
15	-	60.cd	3.05ab
20	-	53.33d	3.33ab
30	-	63.33cd	2.51b
-	0.25	73.33abc	3.05ab
-	0.5	60cd	3.06ab
-	1.0	60cd	3.17ab
-	1.5	83.33ab	2.87ab
-	2.0	86.67a	4.07a

Table 1 Effects of different NAgPs and SNP concentrations on shoot regeneration of carnation

Different letters denote significant differences at $p \leq 0.05$

of IBA and 0.2 mg/L of TDZ provided a reasonable percentage of shoot regeneration (66.67%) and 3.46 number of shoots per explant in the carnation cv. Jinju. This was determined to be the optimal combination for Agrobacteriummediated genetic transformation. However, the shoots induced from the leaf explants showed slow growth and were small in size. Therefore, we tried to improve the regeneration media by adding various concentrations of SNP and NAgPs. SNP and NAgPs promoted shoot regeneration efficiency and in vitro growth in previous studies (Arun 2017; Castro-Gonzalez et al. 2019; Kim 2017; Subiramani 2019). In contrast to those studies, SNP at any concentration did not promote the number of shoots per explant and even suppressed them in some cases. This was not in agreement with the results of Arun et al. (2017), who claimed that 0.83 µM SNP could promote shoot regeneration and plant growth in chrysanthemums. This discrepancy could be due to the difference in plant species and/or plant growth regulators used in the regeneration medium; BA 4.44 µM was used in the study by Arun et al. (2017). In fact, the concentrations used in our study and that of Pe et al. (2020) were relatively different, despite their positive effects. This discrepancy may be due to the different plant species that were used. Considered together, the media comprising IBA (0.1 mg/L), TDZ (0.2 mg/L), and NAgPs (2.0 µM) was selected as the optimal media composition for Agrobacteriummediated genetic transformation of carnation cv. Jinju.

Optimization of the minimal PPT concentration on the suppression of shoot regeneration

To obtain an efficient transformation protocol, optimization

of the minimal concentration of a selective agent was an essential step during the selection period. Because a high concentration of a selective agent such as PPT can kill transgenic plants, a lower concentration can give both escape and putative transgenic plants, which will be laborious in the further selection of putative transgenic plants.

In this study, the addition of different concentrations of PPT $(0.3 \sim 0.5 \text{ mg/L})$ to the shoot regeneration medium did not completely suppress shoot regeneration, whereas complete suppression of shoot regeneration was observed in media containing PPT (1.0 or 1.5 mg/L) (Table 2). Based on this result, 1.0 mg/L PPT is optimal minimal concentration that can kill escape plants (non-transgenic plants).

Effect of the pre-culture period on shoot regeneration

In this study, pre-culturing the explants in the shoot regeneration media for three days prior to *Agrobacterium* infection promoted high transformation efficiency compared to that after one or two days. The explants were pre-cultured in the shoot regeneration medium and were placed in a dark condition for 1, 2, or 3 days, followed by infection with *Agrobacterium* cell suspension and co-cultivation for 3 days.

On the selection medium containing 1.0 mg/L PPT, the induction of shoots was observed from the cut-edge of the explants. However, the explants pre-cultured in a dark condition for 3 days exhibited a higher percentage of shoot regeneration, although the number of shoots per explant was not significantly different (Fig. 1). Therefore, it was suspected that a pre-culture of the explants for three days would improve the transformation efficiency of carnation cv. Jinju.

PPT concentration (mg/L)	Shoot induction (%)	Mean number of shoots/explants
-	66.67a	3.46a
0.3	36.67b	1.08b
0.5	40b	1.00b
1.0	-	-
15	_	_

Table 2 Effects on different PPT concentrations on carnation shoot regeneration (DMRT, $p \le 0.05$)





Fig. 1 Effects of pre-cultivation period on regeneration and number of shoots/explants of carnations (DMRT, $p \leq 0.05$)

During the genetic transformation process, before explants were infected with an Agrobacterium cell suspension, the pre-culture of the explants in a dark condition was important for improving the transformation efficiency (M.Marutani-Hert et al. 2012). Pre-culturing the explants in shoot regeneration media for an optimum period increased the competency of Agrobacterium, as cell division and dedifferentiation were initially started (Chateau et al. 2000). Although the precise mechanism of action is not yet known, the thin walls of cells were newly formed after vigorous cell division due to the effects of pre-culturing on shoot regeneration, changes in microtubule composition, and increases in plant cells in the S-phase, which make it easy to insert T-DNA (Sangwan et al. 1992). However, the longer the pre-treatment period, the longer the wound recovery period, which makes it difficult for bacteria, get through to the plant cell.

Effect of the co-cultivation period on shoot regeneration

According to the results shown in Figure 1, a pre-culture of the explants in the dark condition for three days exhibited improved regeneration efficiency.

In this study, when the co-cultivation periods were extended three to five days, a variation in shoot regeneration efficiencies was observed, specially the explants that were

Fig. 2 Effect of co-culture period on regeneration and number of shoots/explants of carnation (DMRT, $p \le 0.05$)

co-cultivated with *Agrobacterium* cells for four days significantly improved the shoot regeneration of the PPTcontaining media compared to those co-cultivated for three or five days, despite no difference in the number of shoots per explant (Fig. 2). This indicated that the application of the pre-culture period (three days) and co-cultivation period (four days) would have more potential to improve the transformation efficiency of this carnation. *Agrobacterium* cells transfer their T-DNA into the plant genome during the co-cultivation period (Thomashow et al. 1980). If the co-cultivation period is too short, bacteria are not able to effectively transfer their T-DNA into the plant genome. Similarly, if the co-cultivation period is too long, over-growth of bacteria can occur in the explants, and which will negatively affect the survival of the explants.

In the above experiments, shoots were induced from the edge of the explants in the selection medium. Most of the shoots were vitrified and likely to have low chlorophyll contents in their leaves (Fig. 3A). The shoots produced roots in the rooting media and converted to normal shoots when they were further cultured on hormone-free MS media containing 1.0% plant agar (Fig. 3B). Subsequent transfer of the normalized plants to the same media containing 1.0% plant agar produced completely normalized plants (Fig. 3C).



Fig. 3 Transformation of *Dianthus caryophyllus L*. cv. Jinju. (A) Shoot regenerated after four weeks in the selection medium (bar = 1 mm). (B) Shoots gradually changed to normal in the conversion medium. (C) Shoots normalized and rooted in the conversion medium

Conclusion

This study demonstrated that the addition of SNP to the shoot regeneration medium suppressed shoot regeneration in the carnation cv. "Jinju". However, when NAg was added to the media, the concentration of 2.0 µM was found to promote shoot regeneration and plant growth, which indicated its potential as a positive promoter of shoot regeneration and plant growth compared to SNP. Considering PPT as a selective agent, 1.0 mg/L was the minimal and optimal concentration for the selection of putative transgenic plants. When the explants were infected with Agrobacterium cells harboring the acdS gene, the explants that were pre-cultured for three days induced a greater number of putative transgenic plants with a co-cultivation period of four days. Therefore, we expect that this study will provide advantages to researchers who are developing genetic transformations of carnations.

Acknowledgements

This work was supported by a grant from Rural Development Administration in Korea (Project no. PJ014858).

References

- Arun M, Naing AH, Jeon SM, Ai TN, Aye T, Kim CK (2017) Sodium nitroprusside stimulates growth and shoot regeneration in chrysanthemum. Horticulture, Environment, and Biotechnology 58(1):78-84
- Castro-González CG, Sánchez-Segura L, Gómez-Merino FC, Bello-Bello JJ (2019) Exposure of stevia (Stevia rebaudiana B.) to silver nanoparticles in vitro: Transport and accumulation. Scientific reports 9(1):1-10
- Chateau S, Sangwan RS, Sangwan-Norreel BS (2000) Competence of Arabidopsis thaliana genotypes and mutants for Agrobacterium tumefaciens-mediated gene transfer: role of phyto-

hormones. Journal of experimental botany 51(353): 1961-1968

- Firoozabady E, Moy Y, Tucker W, Robinson K, Gutterson N (1995) Efficient transformation and regeneration of carnation cultivars using Agrobacterium. Molecular breeding 1(3): 283-293
- Glick BR (2014) Bacteria with ACC deaminase can promote plant growth and help to feed the world. Microbiological research 169(1):30-39
- Grichko VP, Glick BR (2001) Flooding tolerance of transgenic tomato plants expressing the bacterial enzyme ACC deaminase controlledby the 35S, rolD or PRB-1b promoter. Plant Physiology and Biochemistry 39(1):19-25
- Heydarian Z, Yu M, Gruber M, Glick BR, Zhou R, Hegedus DD (2016) Inoculation of soil with plant growth promoting bacteria producing 1-aminocyclopropane-1-carboxylate deaminase or expression of the corresponding acdS gene in transgenic plants increases salinity tolerance in Camelina sativa. Frontiers in microbiology 7:1966
- Jung H, Ali S, Kim JY, Kim W-C (2018) Transgenic Arabidopsis expressing acdS gene of Pseudomonas veronii-KJ alleviate the adverse effects of salt and water-logging stress. Plant Breeding and Biotechnology 6(3):221-232
- Kim DH, Gopal J, Sivanesan I (2017) Nanomaterials in plant tissue culture: the disclosed and undisclosed. RSC advances 7(58): 36492-36505
- Klee HJ, Hayford MB, Kretzmer KA, Barry GF, Kishore GM (1991) Control of ethylene synthesis by expression of a bacterial enzyme in transgenic tomato plants. The Plant Cell 3(11):1187-1193
- Marutani-Hert M, Bowman KD, McCollum GT, Mirkov TE, Evens TJ, Niedz RP (2012) A dark incubation period is important for Agrobacterium-mediated transformation of mature internode explants of sweet orange, grapefruit, citron, and a citrange rootstock. PloS one 7(10):e47426
- Naing AH, Ai TN, Jeon SM, Lim SH, Kim CK (2016) An efficient protocol for Agrobacterium-mediated genetic transformation of recalcitrant chrysanthemum cultivar Shinma. Acta physiologiae plantarum 38(2):38
- Nie L, Shah S, Rashid A, Burd GI, Dixon DG, Glick BR (2002) Phytoremediation of arsenate contaminated soil by transgenic canola and the plant growth-promoting bacterium Enterobacter cloacae CAL2. Plant Physiology and Biochemistry 40(4):

355-361

- Pe PPW, Naing AH, Soe MT, Kang H, Park KI, Kim CK (2020) Establishment of meristem culture for virus-free and genetically stable production of the endangered plant Hosta capitata. Scientia Horticulturae 272:109591
- Sangwan RS, Bourgeois Y, Brown S, Vasseur G, Sangwan-Norreel B (1992) Characterization of competent cells and early events of Agrobacterium-mediated genetic transformation in Arabidopsis thaliana. Planta 188(3):439-456
- Sergeeva E, Shah S, Glick BR (2006) Growth of transgenic canola (Brassica napus cv. Westar) expressing a bacterial 1-aminocyclopropane-1-carboxylate (ACC) deaminase gene on high concentrations of salt. World Journal of Microbiology and Biotechnology 22(3):277-282
- Stearns JC, Shah S, Greenberg BM, Dixon DG, Glick BR (2005) Tolerance of transgenic canola expressing 1-aminocyclopropane-1-carboxylic acid deaminase to growth inhibition by nickel. Plant Physiology and Biochemistry 43(7):701-708
- Subiramani S, Sundararajan S, Sivakumar HP, Rajendran V, Ramalingam S (2019) Sodium nitroprusside enhances callus induction and shoot regeneration in high value medicinal plant Canscora diffusa. Plant Cell, Tissue and Organ Culture (PCTOC) 139(1):65-75
- Thomashow MF, Nutter R, Montoya AL, Gordon MP, Nester EW (1980) Integration and organization of Ti plasmid sequences in crown gall tumors. Cell 19(3):729-739
- Thu HTM, Naing AH, Jeong HY, Kim CK (2020) Regeneration of genetically stable plants from in vitro vitrified leaves of different carnation cultivars. Plants 9(8):950