# **ARTICLE / INVESTIGACIÓN**

# Assessment of genetic fidelity of lacy tree philodendron (Philodendron bipinnatifidum Schott ex Endl.) micro propagated plants

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DOI. 10.21931/RB/2022.07.01.10

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Abstract: Tissue culture is a potent means for producing clonally propagated plants. However, genetically identical regenerants are of great concern due to their economic consequences. Therefore, polymerase chain reaction (PCR)-based molecular markers are employed to detect somaclonal variations. In this study, the genetic fidelity of in vitro raised Philodendron bipinnatifidum clones, and their mother plant was tested using 11 randomly amplified polymorphic DNA (RAPD) markers. The RAPD decamers produced 92 amplicons with 8.4 bands ranging from 260-5000 bp. The bands varied from one to fifteen for primer 8 and primer 7, respectively. The genetic similarity between the micro propagated plantlets and mother plant of P. bipinnatifidum was nearly 100%, assuring uniformity and true-to-type regenerated plantlets for this commercially crucial ornamental plant.

Key words: Araceae, genetic fidelity, micropropagation, polymorphism, RAPD analysis.

### Introduction

Rapid clonal propagation is one of the most extensively used approaches of biotechnology. In vitro plants undergo several subculture cycles before their rooting, and acclimatization and cytogenetic variations may occur in this long process<sup>1</sup>. Somaclonal variations can be a valuable source for new genetic material<sup>2</sup> and might be advantageous for crop improvement<sup>3</sup>. However, its occurrence in micro propagated plants results in undesirable plant off-types, limiting its applications in the tissue culture industry. Therefore, the most critical concern has been the maintenance of genetic uniformity and the production of true-to-type plants. The influence of in vitro culture conditions i.e., culture media, type of explant, and subculture cycles, may lead to genomic variation in the regenerated plantlets. Therefore, assessment of genetic stability is a prerequisite while performing micropropagation<sup>4</sup>.

Several methods were available to assess regenerated plants' genetic fidelity, such as morphological characteristics, cytological and flow cytometry analysis<sup>5</sup>. DNA based molecular markers emerged as the most suitable genetic tools to determine the genetic variation among regenerated plants<sup>6</sup>. Molecular markers such as randomly amplified polymorphic DNA (RAPD)<sup>7</sup>, amplified fragment length polymorphism (AFLP)<sup>8</sup>, simple sequence repeat (SSR)<sup>9</sup> and inter simple sequence repeats (ISSR)<sup>10,11</sup> are used for determining the genetic fidelity of micro propagated plants. Cost-effectiveness is a major checkpoint in micropropagation. Because of the simplicity and cost effectiveness of RAPD technique, it is used for genetic stability analysis, amplifying the different regions of the genome<sup>12</sup> in several plant species<sup>4,6,13,14</sup>.

Lacy tree philodendron (Philodendron bipinnatifidum

Schott Ex Endl.), formerly known as P. selloum K. Koch, is a self-heading ornamental plant species that can reach heights of 4–5 m and possesses deeply cut, green to dark green leaves that can grow up to 1 m in length. Conventional propagation does not encourage the multiplication of the species<sup>15</sup>. Therefore, in vitro propagation methods can be utilized to facilitate the continuous generation of high-quantity and high-quality plant materials compared with traditional vegetative propagation. Our previous report established an efficient micropropagation protocol for lacy tree philodendron through axillary shoot regeneration with a 100% rooting and acclimatization<sup>15</sup>. Therefore, the objective of this study was to assess the genetic fidelity of the regenerants to assure the production of genetically identical plantlets.

### Materials and methods

### Plant material

The mother plant of *P. bipinnatifidum* (Figure 1 A) and ten micro propagated plantlets through axillary shoot proliferation<sup>15</sup> were randomly selected and used as plant material in this study. The axillary shoots were multiplied on MS medium<sup>16</sup> that contained 6-benzylaminopurine (1 mg·L<sup>-1</sup>) and indole-3-butyric acid (0.5 mg·L<sup>-1</sup>) for 6 wks (Figure 1 B). These in vitro shoots were kept in our laboratory for nine subculture cycles, in vitro plantlets rooted on MS medium that contained naphthalene acetic acid  $(1 \text{ mg} \text{L}^{-1})$  for 6 wks (Figure 1 C) and acclimatized in a 1:1 (v:v) mixture of peat moss and perlite for

Citation: Alwahibi MS, Alawaadh AA, Dewir YH, Soliman DA, and Seliem MK. Assessment of genetic fidelity of lacy tree philodendron (Philodendron bipinnatifidum Schott ex Endl.) micro propagated plants. Revis Bionatura 2022;7(1). 10. http://dx.doi.org/10.21931/RB/2022.07.01.10 Received: 17 June 2021 / Accepted: 20 August 2021 / Published: 15 February 2022

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30 d (Figure 1 D) as described by Alawaadh *et al.*<sup>15</sup>. Young fresh leaves of the mother plant and clonal regenerants were thoroughly cleaned, washed with sterile distilled water, and kept in liquid nitrogen until molecular analysis.

### **Genomic DNA isolation**

Genomic DNA was isolated from fresh leaves of both mother plants and micro propagated plants by using the cetyltrimethylammonium bromide (CTAB) method<sup>17</sup>. Approximately 100-150 mg of leaves were ground using 600 µL of preheated (65°C) extraction buffer (2% CTAB, 20 mM EDTA, 100 mM Tris-HCl, 1.4 M NaCl, 2% polyvinylpyrrolidone (PVP), 0.2% mercaptoethanol), transferred to a centrifuge tube (2 mL) and incubated for 30 min in a 65°C water bath, the samples were inverted every 5 min. 600 µL of chloroform-isoamyl alcohol (24: 1) was added and mixed by inverting the tubes carefully 10 times, and the cells were centrifuged at 12000 rpm for 10 min at room temperature. The supernatant was collected and carefully mixed with a two-thirds volume of ice-cold Isopropanol, and the DNA samples were collected by centrifuging for 10 min. RNaseA (10  $\mu$ g/ml) was added to the 50  $\mu$ l of TE buffer (10 mM Tris and 0.1 mM EDTA) before dissolving the DNA to remove any RNA in the preparation, and the mixture was incubated at 37 °C for 30 min. 100  $\mu L$  and 750  $\mu L$  of 3 M sodium acetate and absolute ice-cold ethanol were added. The DNA was collected by high-speed centrifugation for 10 min, and then carefully washed with ice-cold absolute and 70% ethanol and centrifuged at 12000 rpm for 10 min. Finally, the samples were dried at room temperature and dissolved in 50-100  $\mu$ L of TE buffer. The quality and concentration of DNA were determined by EMPLEN photometer P330.

### PCR conditions

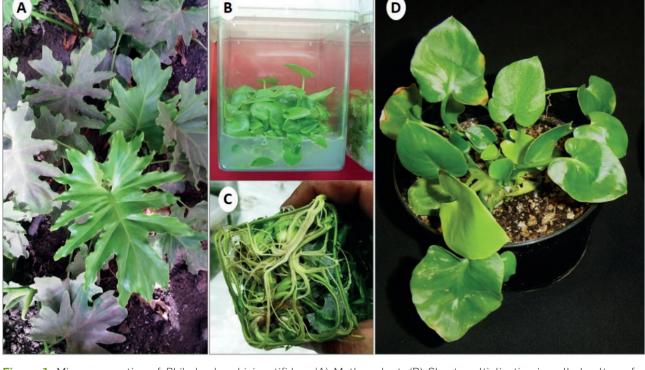
11 decamer RAPD primers (Macrogen Korea) were used for DNA amplification. PCR reaction was carried out in a volume of 20  $\mu$ L. PCR was performed as follows: 94°C for 5 min; followed by 35 cycles of 94°C for 1 min, specific annealing temperature (Ta) according to the primer sequence for 30 sec and 72°C for 3 min and the final extension step at 72°C for 10 min. Amplification was carried out in MJ Mini Bio RAD, thermal cycler in 25  $\mu$ L reaction volume containing the following reagents: 1.0  $\mu$ L of dNTPs (10 mM), 1.0  $\mu$ L of MgCl<sub>2</sub> (25 mM), 5  $\mu$ L of 10x buffer, 1.0  $\mu$ L of primer (10 pmol), 1.0  $\mu$ L of DNA (25 ng  $\mu$ L<sup>-1</sup>), 0.3  $\mu$ L of taq polymerase (5 u  $\mu$ L<sup>-1</sup>) and 15.7 dd.H<sub>2</sub>O.

### Gel electrophoresis and genetic analysis

The RAPD products were electrophoresed in 1.5 % agarose gel containing Red safe dye in TAE buffer (40 mM Tris-acetate, 20 mM glacial acetic acid, 1 mM EDTA, pH 7) at 75 V. The gels were documented using a gel documentation system (UVITEC, UK) and according to an analysis by Phoretix program 1D gel analysis software version 4.01). Two repeats were performed to confirm the results. The bands obtained by scoring the RAPD profiles were treated as binary characters and coded accordingly (presence = 1, absence = 0). The genotypes showing similarity in their RAPD characteristics were grouped using UPGMA (Unweighted Paired Group with Arithmetic Average) The SPSS-10 package was used for statistical analysis.

### **Results**

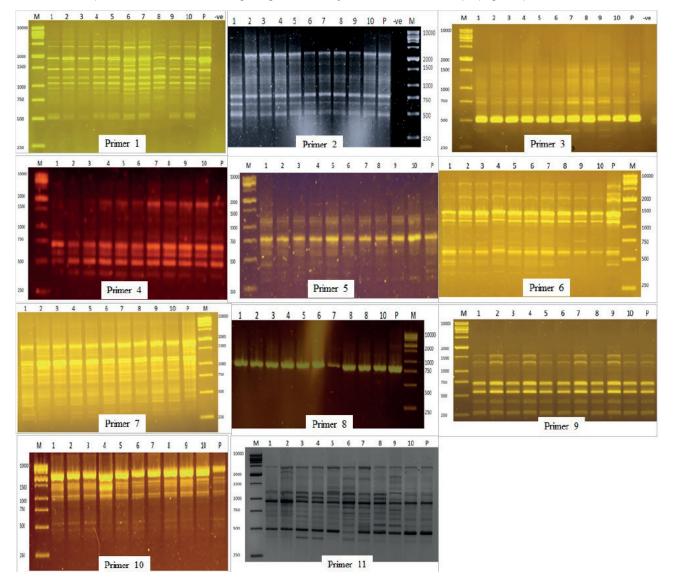
Assessment of genetic fidelity was performed on the *P. bipinnatifidum* micro propagated plantlets using RAPD analysis



**Figure 1.** Micropropagation of *Philodendron bipinnatifidum* (A) Mother plant, (B) Shoot multiplication in gelled culture, full-strength Murashige and Skoog medium that contained 6-benzylaminopurine (1 mg·L<sup>-1</sup>) and indole-3-butyric acid (0.5 mg·L<sup>-1</sup>), after 6 wks of culture, (C) In vitro plantlets rooted on Murashige and Skoog medium that contained naphthalene acetic acid (1 mg·L<sup>-1</sup>), after 6 wks of culture, (D) Acclimatized plantlet in a 1:1 (v:v) mixture of peat moss and perlite, after 30 d. (Unpublished photographs).

Primers code	Primer sequences (5'- 3'	No. of scorable band per primer	Size range for amplified products	No. of monomorphic bands	Polymorphism (%)	
1	CAGGCCTTTC	11	500-3000	11	0	
2	AGTCAGCCAC	8	500-2500	8	0	
3	AATCGGCTG	4	500-1500	4	0	
4	AGGGGTCTTG	5	400-1900	5	0	
5	GGGTAACGCC	8	300-1500	8	0	
6	TCGGCGATAG	11	350-5000	10	9.1	
7	CAGCACCCAC	15	270-1800	15	0	
8	TTCCGAACCC	1	1000	1	0	
9	AGCCAGCGAA	7	260-1500	7	0	
10	GACCGCTTGT	8	500-2000	8	0	
11	GTGCAACGTG	14	400-3000	12	14.3	
Total		92		89	3.3	

**Table 1.** RAPD primers were used for testing the genetic fidelity of *Philodendron* micro propagated plants.



**Figure 2.** DNA Amplification pattern obtained with RAPD primers. Lane M- DNA ladder; Lane-P D.N.A. from mother plant; Lane 1-10 DNA from micropropagated plants.

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	Mother plant	Clone 1	Clone 2	Clone 3	Clone 4	Clone 5	Clone 6	Clone 7	Clone 8	Clone 9	Clone 10
Clone 1	1.00	1.00	0.99	0.99	0.99	0.99	0.99	0.99	0.99	0.99	1.00
Clone 2	0.99	0.99	1.00	0.98	0.98	0.98	0.98	0.98	0.98	0.98	0.99
Clone 3	0.99	0.99	0.98	1.00	1.00	1.00	1.00	1.00	1.00	0.98	0.99
Clone 4	0.99	0.99	0.98	1.00	1.00	1.00	1.00	1.00	1.00	0.98	0.99
Clone 5	0.99	0.99	0.98	1.00	1.00	1.00	1.00	1.00	1.00	0.98	0.99
Clone 6	0.99	0.99	0.98	1.00	1.00	1.00	1.00	1.00	1.00	0.98	0.99
Clone 7	0.99	0.99	0.98	1.00	1.00	1.00	1.00	1.00	1.00	0.98	0.99
Clone 8	0.99	0.99	0.98	1.00	1.00	1.00	1.00	1.00	1.00	0.98	0.99
Clone 9	0.99	0.99	0.98	0.98	0.98	0.98	0.98	0.98	0.98	1.00	0.99

Table 2. Similarity coefficient among the screened plants based on RAPD analysis.

with 11 RAPD primers (Table 1; Figure 2). The selected primers gave 92 scorable bands with an average of 5.8 bands ranging from 260-5000 bp. The bands varied from 1 to 15 for primer 8 and primer 7, respectively.

RAPD amplification pattern obtained with all primers except for primers 6 and 11 revealed that all bands produced by micro propagated plants were monomorphic and similar to that of the mother plant. Primer 6 amplified 10 monomorphic bands and one unique polymorphic band (750 bp) produced by *in vitro* propagated clone 9. Primer 11 produced 14 scorable bands, out of which 12 were monomorphic while two were polymorphic. A unique band with a molecular size of 2400 bp was produced only in propagated clone 2, whereas micro propagated clones 3,4,5,6,7,8 produced a band with a molecular size of 1000 bp. Of the 11 primers analyzed, nine primers showed 0% polymorphism, while primers 6 and 11 showed 9.1% and 14.3% polymorphism, respectively. The genetic similarity between the micro propagated plantlets and the mother plants ranged from 0.98 to 1.00, with an average of 0.99 (Table 2).

### Discussion

High genetic fidelity and true-to-type clones are critical for commercial micropropagation to maintain the essential characteristics of the mother plant<sup>1</sup>. Phenotypic variations such as abnormal leaf shape and dwarf growth indicate off-type clones. However, the absence of these morphological abnormalities does not guarantee true-to-type regenerated plantlets; therefore, molecular markers should be utilized to assure their clonal fidelity. It has been reported that although in vitro regenerated Philodendron micans plantlets phenotypically resembled mother plants, they showed a high frequency of ploidy change suggests the occurrence of somaclonal variation<sup>18</sup>. The author indicated that analysis of 20 randomly selected Philodendron micans plantlets using DNA flow cytometry showed that two were mixoploid. RAPD profiles of the randomly selected clones of *Philodendron* 'Xanadu' different ages compared to the mother plant were identical, thus assuring a genetic fidelity-maintained protocol for this commercially important plant<sup>19</sup>. RAPD molecular marker was used to assess genetic fidelity in other micro propagated plant species belonging to the Araceae family, such as Aglaonema' Valentine'<sup>7</sup> where all tested plantlets were highly uniform or identical to the mother plant. Conversely, for Anthurium Andreanum Bicolour 'Agnihotri', RAPD analysis revealed variations in band pattern of the regenerated plantlets<sup>20</sup>.

Genetic variation is a common phenomenon observed in

plants regenerated from tissue cultures<sup>3</sup>. Various factors such as genotype, explant type, culture periods and growth regulators combinations and concentrations may disturb the internal polarity and physiology of the explants<sup>21,22</sup> and could also influence the stability of tissue cultured plants<sup>22,23</sup>. Therefore, limiting the number of subculture cycles could maintain clonal characteristics. For example, DNA polymorphism (20%) was reported in protocorm like-bodies of *Phalaenopsis gigantea* after 20 wks of culture on a medium containing 4.5  $\mu$ M thidiazuron and 65.5  $\mu$ M chitosan while reducing the culture period to 16 wks resulted in no variations<sup>24</sup>. However, the amenability to *in vitro* induced mutagenic effect is species-dependent. Borsai *et al.*<sup>25</sup> reported no polymorphism in micro propagated blackberry plants (*Rubus fruticosus*) for 12 subcultures during 30 months of *in vitro* multiplication.

# Conclusions

The genetic fidelity of micropropagated *P. bipinnatifidum* clones was confirmed by using PCR-based RAPD analysis. Polymorphism detected among these clones was negligible. The micropropagation protocol can successfully be employed for commercial application without any risk of genetic instability.

### Author Contributions

Conceptualization, MSA, AAA and YHD; methodology, MSA and AAA; software, MSA and AAA; validation, YHD, DAS and MKS; formal analysis, MSA, AAA and YHD; investigation, MSA and AAA; resources, MSA and YHD; data curation, MSA and AAA; writing—original draft preparation, MSA, AAA and YHD; writing—review and editing, MSA, AAA, YHD; DAS and MKS; visualization, DAS and MKS; project administration, MSA and YHD; funding acquisition, MSA. All authors have read and agreed to the published version of the manuscript.

### Funding

This research was funded by the Deanship of Scientific Research at King Saud University for funding this work through research group NO (RGP-1435-086).

#### Institutional Review Board Statement

Not applicable.

### **Informed Consent Statement**

Not applicable.

### **Data Availability Statement**

All data are presented within the article.

#### Acknowledgments

The authors extend their appreciation to the Deanship of Scientific Research at King Saud University for funding this work through research group NO (RGP-1435-086).

#### **Conflicts of Interest**

The authors declare no conflict of interest. The funders had no role in the design of the study; in the collection, analyses, or interpretation of data; in the writing of the manuscript, or in the decision to publish the results.

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