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Metabolic profiling, in vitro propagation, and genetic assessment of the endangered rare plant *Anarrhinum pubescens*



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Abstract

Background: Anarrhinum pubescens Fresen. (Plantaginaceae) is a rare plant, endemic to the Saint Catherine area, of South Sinai, Egypt. Earlier studies have reported the isolation of cytotoxic and anti-cholinesterase iridoid glucosides from the aerial parts of the plant. The present study aimed to investigate the chemical profiling of the wild plant shoots as well as establish efficient protocols for in vitro plant regeneration and proliferation with further assessment of the genetic stability of the in vitro regenerated plants.

Results: Twenty-seven metabolites have been identified in wild plant shoots using the Nuclear Magnetic Resonance (NMR) spectroscopy. The metabolites include alkaloids, amino acids, carbohydrates, organic acids, vitamins, and a phenol. In vitro propagation of the plant was carried out through nodal cutting-micropropagation and leaf segment-direct organogenesis. The best results were obtained when nodal cutting explants were cultured on Murashige and Skoog medium with Gamborg B5 vitamins supplemented with 6-benzylaminopurine (BAP) (1.0 mg/L) and naphthaleneacetic acid (NAA) (0.05 mg/L), which gave a shoot formation capacity of 100% and a mean number of shoots of 27.67 \pm 1.4/explant. These shoots were successfully rooted and transferred to the greenhouse and the survival rate was 75%. Genetic fidelity evaluation of the micropropagated clones was carried out using random amplified polymorphic DNA (RAPD) and inter simple sequence repeat (ISSR) molecular markers. Jaccard's similarity coefficient indicated a similarity as high as 98% and 95% from RAPD and ISSR markers, respectively.

Conclusions: This study provides the chemical profiling of the aerial part of *Anarrhinum pubescens*. Moreover, in vitro regeneration through different tissue culture techniques has been established for mass propagation of the plant, and the genetic fidelity of the in vitro regenerated plants was confirmed as well. Our work on the in vitro propagation of *A. pubescens* will be helpful in ex situ conservation and identification of bioactive metabolites.

Keywords: Anarrhinum pubescens, Metabolic profiling, In vitro propagation, Callus culture, Genetic fidelity, NMR spectroscopy

Background

Anarrhinum pubescens Fresen. (Family: Plantaginaceae) is an endemic perennial plant that grows on granite rocks, in the Saint Catherine region of Egypt [1–4]. The plant is described by the International Union for Conservation of Nature (IUCN) as a rare plant (https://doi. org/10.2305/IUCN.UK.2017-3.RLTS.

T84119796A84119800.en). The wild plant is distributed in limited narrow areas in only two localities in the Saint Catherine protectorate (Wadi Gebal, Elgabal Elahmar, and Wadi Abu Twita) [5]. The number of plant individuals sharply decreased due to climatic changes and human activities [5–7]. As the wild plant is used in folk medicine for relieving pain. Phytochemical and biological analysis of the plant showed the presence of iridoid compounds which possess cytotoxic and acetylcholinesterase inhibitory activities [8, 9], suggesting

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the possible application of the plant in developing anticancer and Alzheimer disease relieving drugs. The genus *Anarrhinum* is characterized by the presence of iridoid glycoside compounds. These structurally diverse metabolites are well known for their biological activities, principally, the antimicrobial activities [10, 11]. Iridoid glycosides were isolated from *Anarrhinum orientale* and some of them showed a selective inhibition to hepatitis C virus protease [12, 13].

The application of NMR analysis in metabolic profiling of medicinal plants is a growing field of research. It provides a snapshot for what is included inside the plant tissues and, accordingly, it might help in biomarker discovery in certain plants. In addition, since it gives real information about metabolites included inside the plant cells or tissues, it can be of great importance in distinguishing different plant species sharing the same genus [14].

In vitro propagation strategies for conserving *A. pubescens* have not been studied yet. Traditional methods of propagation via seed germination are unpredictable as it is usually influenced by seasonal changes [15], in the case of *A. pubescens*, only a limited number of plants can be produced due to the rareness of the plant's seeds. However, biotechnological tools, like tissue culture technology, have proven to provide a reliable method for rapid multiplication of the desired plant species and so conserving its germplasm by establishing tissue banks or increasing intervals between subcultures [16, 17]. Despite the large number of studies on the in vitro propagation of Plantaginaceae plants [18, 19], the application of this in vitro technology to propagate any of the species from genus *Anarrhinum* is still not fully established.

Occasionally, establishment of clonal propagation protocols leads to genetic variations due to stress induced by using plant growth regulators, different media ingredients, or callus intermediate stages [20]. Accordingly, investigation of the genetic stability of the resulting progenies should be considered. Molecular markers techniques such as RAPD, ISSR, and restriction fragment length polymorphism (RFLP) have been increasingly utilized to evaluate the genetic fidelity of the propagated plants, and their common use for this purpose has been well documented [21, 22].

Plants serve as a substantial source of bioactive metabolites. Plants belong to Plantaginaceae family are sources for many biologically active compounds, principally, the antimicrobial compounds [23, 24].

Here, we report the metabolic profiling, in vitro propagation, and callus induction for genus *Anarrhinum*, which could be an important basic reference for any further research concerned with chemical analysis and mass propagation of the *A. pubescens* plant or other species sharing the same genus. Our study displays efficient

protocols for in vitro regeneration and callus induction of *A. pubescens* as reliable methods for medium to long-term conservation of this endangered rare plant.

Methods

Plant material

Seeds and aerial parts (leaves and stems) of *Anarrhinum pubescens* were collected from St. Catherine, Egypt in April 2015. The plant was authenticated by the staff members of the St. Catherine protectorate. Plant aerial parts were used for the metabolic profiling of the plant, whereas seeds were used for the tissue culture experiments.

Chemicals

Tissue culture media (Murashige and Skoog medium with Gamborg B5 vitamins) and growth regulators (6-benzylaminopurine, kinetin, naphthaleneacetic acid and 2, 4 dichlorophenoxy acetic acid) were purchased from Phyto-Technology Laboratories (Shawnee Mission, KS, USA). Ethanol, methanol, chloroform, and TMSP buffer components were purchased from Sigma-Aldrich (St. Louis, MO, USA).

Metabolites extraction from the wild plant

Six individuals from the wild plant were selected randomly, air dried, ground, and homogenized under liquid nitrogen. From each sample, 20 mg dry weight was used for the extraction processes. Metabolites were extracted using 2:2:1.8 methanol: chloroform: water (V/V/V) [25] and resulted in separated polar and nonpolar liquid phases upon centrifugation for 10 min at $2000\times g$ and 4 °C. The polar extract (upper phase) was dried using a Centrivap vacuum concentrator for 24 h at 30 °C.

NMR sample preparation and data collection

The dried polar extract was rehydrated in 620 μ L NMR buffer (100 mM sodium phosphate buffer (pH 7.3), 1 mM TMSP (internal standard,3-(trimethylsilyl) propionic-2,2,3,3-d4 acid, CAS: 24493-21-8), and 0.1% sodium azide, in 99.9 atom % D₂O).

All data were collected using a spectral width of 16.0 ppm and 64K points resulting in an acquisition time of 2.9 s. The first increment of the presat-noesy spectra was collected with 120 scans, 4 dummy scans, 3 s relaxation delay, and on-resonance pre-saturation at the residual water frequency for solvent suppression. The 90 ° pulse widths were measured for each sample using the automatic pulse calculation experiment (pulsecal) in TopSpin 2.1.1 (BrukerBioSpin, Billerica, MA). Two dimensional ¹H-¹³C HSQC data were collected at 700 MHz using a Bruker hsqcedetgpsisp2.2 pulse sequence. The ¹H was observed in the F2 channel with a spectral

width of 11 ppm while the ¹³C was observed in the F1 channel with a spectral width of 180 ppm.

Metabolite identification and quantification

Polar metabolites from the aerial part of the plant were identified by comparing the ¹H data with the 700 MHz database of metabolite standards in Chenomx NMR Suite (Chenomx Inc., Edmonton, Alberta, Canada). All identified metabolites were authenticated by comparing the obtained 1D and 2D spectra with the data represented in Human Metabolome Database (HMDB).

In vitro propagation protocol

Culture conditions and seedling development

Anarrhinum pubescens seeds were rinsed three times with distilled H₂O for 5 min each, then surface sterilized by dipping in 95% ethanol for 1 min, followed by disinfection with 20% commercial Clorox® solution containing one drop of Tween 20® for 20 min and finally washed four times with sterile distilled H₂O for 5 min each under aseptic conditions. Seeds were germinated by culturing on full strength Murashige and Skoog medium with Gamborg B5 vitamins (MS-B5) [26, 27] in magenta vessels using six seeds per vessel. The culturing media contained 30 g/L sucrose, adjusted to pH 5.7 and solidified with 2 g/L phytagel (St. Louis, MO, USA) before autoclaving at 121 °C for 20 min. Cultures were incubated at 25 °C under 16 h photoperiod provided by cool white fluorescent lamps for 4 weeks.

Shoot multiplication

Seedlings (4-week-old) were defoliated and cut into 0.8-1.0 cm nodal segments (0.5 cm down from the nodes) prior to being vertically inoculated in culture medium with an immersion length from the bottom 0.3 cm of the bottom internode (nodes were kept above the surface of the medium). Nodes were cultured on fullstrength MS-B5 medium supplemented with 30 g/L sucrose and containing different concentrations of 6benzylaminopurine (BAP) (0.5, 1.0, 2.0, 3.0, and 4.0 mg/ L) or kinetin (Kn) (0.5, 1.0, 2.0, 3.0, and 4.0 mg/L) plus a fixed concentration of naphthaleneacetic acid (NAA) (0.05 mg/L). The medium was solidified using 2.0 g/L phytagel. Nodal cutting explants were placed in magenta vessels using three explants per vessel. At least nine replicates were used per treatment and all experiments were repeated three times. All cultures were incubated for a 16-h photoperiod at 25 °C in a completely randomization design. Six weeks later, the cultures were screened for percentage of shoot formation (%) =

 $\frac{No.~of~explants~that~formed~shoots}{Total~No.of~explants} \times 100~$ and mean number of

shoots = $\frac{Total\ No.\ of\ shoots}{Total\ No.\ of\ explants}$ and the results were statistically analyzed.

Rooting and acclimatization

Developed shoots (3-4 cm) derived from nodal cuttings cultured on full strength MS-B5 medium supplemented with BAP (1.0 mg/L) and NAA (0.05 mg/L) were separated and transplanted into full strength hormone free MS-B5 basal medium with and without 0.1 mg/L IAA. Twenty shoots were transferred to each rooting media. After 30 days of incubation, cultures were examined for root formation percentage of (%)No. of explants formed roots \times 100. Well rooted shoots were removed from the culture vessels, washed thoroughly with tap water, and transferred to 4-cellplastic trays containing the sterilized mixture of Lowes Miracle-Gro All Purpose Potting mix and Vermiculite at 3:1 volumetric ratio (Lowe's Company, USA) and completely enclosed in transparent plastic bags. Afterwards, plantlets were transferred to the greenhouse and maintained at 27 \pm 2 °C under programmed watering.

Statistical analysis

For in vitro propagation, a completely randomized experimental design was performed. All experiments were repeated three times, each time using at least 9 replicates (for in vitro propagation) or 12 replicates (for callus induction) per treatment. Each replicate was represented by 1 plate/4 explants. All replicates came from different seeds. Collected data were subjected to analysis of variance (ANOVA) and means were compared through Fisher pairwise comparison ($p \leq 0.05$). Statistical analyses were accomplished using Minitab* software.

Genetic fidelity assessment

Genomic DNA of mother plants and eight replicates from the in vitro propagated plantlets was isolated using E.Z.N.A. DNA isolation kit as per manufacturer's instructions (Promega, Madison, WI, USA). DNA quality and quantity were determined by recording the ratio of absorbance at 260 to 280 nm using a NanoDrop-1000° Spectrophotometer (NanoDrop Technologies, Wilmington, DE, USA). DNA samples were diluted to 50 ng/µl. A total of 17 RAPD and 25 ISSR primers (Eurofins MWG Operon LLC, Louisville, KY, USA) were initially screened to evaluate the genetic stability of the micropropagated plants; then only 10 RAPD and 10 ISSR primers were chosen based on their reproducibility to generate clear, scorable bands. Amplification reaction was carried out in a 25-µl mixture containing 50 ng of template DNA, 5 µl 5XTaq buffer (Promega, Madison, WI, USA), 2.5 µl dNTPs, 1 U Taq polymerase (Promega,

Madison, WI, USA), 25 mM MgCl₂, and 1 µl primer in a thermal cycler (Eppendorf Mastercycler®, Hamburg, Germany). PCR conditions were programmed as follows: initial denaturation at 94 °C for 4 min followed by 40 cycles of 1 min denaturation at 94 °C, 30 s annealing at the specified temperature for each primer, and 1 min extension at 72 °C and then final extension for 10 min at 72 °C. Amplification products were resolved by gel electrophoresis on 1.5% agarose gel running for 1 h at 105 V in 1X TAE; DNA bands were stained with ethidium bromide. Resulting bands were detected and photographed by Gel Doc XR+ imager® (Bio-Rad, Hercules, CA, USA). Bands sizes were determined by the aid of 1 kb plus DNA ladder (Fisher Scientific, USA). Quantity One Software (BioRad Laboratories, Hercules, CA, USA) was used to analyze the gel profiles. Data matrix was constructed based on scoring clear, reproducible bands into a binary data sheet (1 for presence and 0 for absence). Jaccard's similarity coefficient [28] was calculated using community analysis package 1.14 (CAP) by Henderson and Seaby [29].

Results

Metabolic profiling of the wild plant shoots

The peaks from the ¹H spectra were assigned using Chenomx software (8.6) and its 700 MHz database. For veri-¹H-¹³C HSQC data were analyzed and fication, compared with H-13C HSOC data available online in Human Metabolome Database (HMDB). Twenty-seven metabolites (Table 1) were successfully identified in the polar extract of the wild plant shoots (Fig. 1). The identified compounds include organic acids, vitamins, amines, amino acids, sugars, sugar alcohols, alkaloids, and a phenol. The aliphatic region between δ 0.5 and 3.5 ppm was characterized by the presence of essential and nonessential amino acids like isoleucine, valine, arginine, and betaine. In the sugar region (Fig. 2), δ 3.5–5.5 ppm, we identified the presence of myo-inositol, sucrose, mannitol, fructose, and glucose. Xanthine, gallate, and trigonelline were identified in the aromatic region δ 5.5.5-10.0 ppm. Also, vitamins and organic acids like choline and acetic acid were identified.

Micropropagation via nodal cutting

Nodal cuttings were excised from seedlings (Fig. 3a) and cultured on MS-B $_5$ medium supplemented with BAP or Kn, at different concentrations (0.5, 1.0, 2.0, 3.0, and 4.0 mg/L). Statistical analysis of data collected after 6 weeks of incubation revealed that, BAP concentrations were superior in adventitious shoots induction and proliferation (Fig. 3). Different BAP concentrations showed 100% capacity rate (all explants produced shoots). On the other hand, Kn treatments showed an induction rate range of 71.1–100%. A higher mean number of shoots (27.67 \pm

1.4, 23.19 ± 0.5 shoots/explant) were recorded with 1.0 and 2.0 mg/L BAP, respectively (Table 2). Mean number of shoots decreased from two to five times by using Kn instead of BAP.

Micropropagation via leaf explant

In vitro regeneration using leaf explants was performed by culturing the seedling-derived leaf segments (Fig. 3a) on the same plant growth regulator types and concentrations as used in nodal cutting culture. BAP was significantly superior to Kn in adventitious shoots induction and proliferation. The maximum percentage rate (88.8–81.5 %) of adventitious shoots induction was produced at a wide range of BAP (0.5, 1.0, and 4.0 mg/L), but only at one concentration from Kn (4.0 mg/L). The highest mean number of shoots (9.30 \pm 1.4 shoots/explant) was obtained by using 0.5 mg/L BAP (Fig. 3b, c, Table 2). A lower mean number of shoots was recorded with 1.0 mg/L Kn. Within Kn concentrations, 4.0 mg/L was superior in shoots induction (90.3%) and multiplication (6.5 \pm 1.4 shoots/explant).

Rooting and acclimatization

Healthy regenerated shoots derived from nodal cuttings and induced on MS- B_5 medium with 1 mg/L BAP were examined for rooting capacity. Rooting capacity test was carried out on MS- B_5 hormone-free medium or MS- B_5 medium supplemented with 0.1 IAA mg/L. Rooting response (a percentage of root formation) was higher (85%) in IAA-treated samples (Fig. 3d and Fig. 4). IAA-treated plantlets were transferred to the greenhouse and showed 75% survival after 10 weeks of acclimatization (Fig. 3e, f).

Genetic fidelity assessment

As a rule of thumb, maintaining the genetic uniformity of the propagated plants is an essential prerequisite for most of the mass propagation experimentation. So, in this study, RAPD and ISSR molecular markers were used to evaluate the genetic fidelity of the micropropagated clones. Genomic DNA was isolated from both mother plants and those obtained from the best propagation medium (MS-B $_5$ media with 1 mg/L + 0.05 mg/L NAA) and analyzed by PCR-based molecular markers to trace the occurrence of any polymorphism. Ten out of 17 RAPD and 10 out of 25 ISSR primers were selected to accomplish the study.

The 10 RAPD primers generated a total of 63 clear, distinct, and reproducible bands with an average of 6.3bands/primer and range size 3860–408 PB (Table 3). Eight out of the chosen 10 RAPD primers showed 100% monomorphism. The highest percentage of polymorphism was 25% recorded by the primer OPI-07 (Fig. 5b). The highest number of bands was generated by primer OPA-04 which gave a total of 13

Table 1 list of metabolites identified in the polar extract of *A. pubescens* shoot

_	Compound name	¹³ C (ppm) from HSQC	¹ H (ppm) (functional group or specific H, multiplicity)	Coupling constant J (Hz)	
1	Acetate	26.14 (CH ₃)	1.90(CH ₃ , s)	_	
2	Alanine	18.93 (C ^β)	1.46 (H ^β , d)	7.25	
		53.5 (C ^α)	$3.77(H^{\alpha}, q)$	7.23	
3	Arginine	26.6(C ^V)	1.64 (H ^Y , m)	_	
		26.5(C ^β)	1.88 (H ^β , m)	_	
		57.03(C ^α)	3.75 (H ^a , m)	=	
4	Asparagine	37.5 (C ^β)	$2.94 (H^{\beta}, dd)$	4.16,16.8	
			$2.85 (H^{\beta}, dd)$	7.74,16.8	
5	Betaine	56.23 (CH ₃)	3.24 (CH ₃ , s)	=	
		69.19 (CH ₂)	3.89 (CH ₂ , s)	=	
6	Choline	56.81 (C ^Y)	3.19 (H ^Y , s)	=	
		70.38 (C ^β)	3.50 (H^{β} , m)	=	
7	Dimethylamine	37.0 (CH ₃)	2.71(CH ₃ , s)	_	
8	Formate	=	8.44 (CH, s)	_	
9	Fructose	71.95	3.98 (⁵ CH, m)	_	
		78.20	4.12 (³ CH, m)	-	
10	Fumarate	138.12 (CH)	6.51 (CH, s)	_	
11	Gallic acid	7.02 (CH)	112.119 (CH, s)	_	
12	Glucose	74.41 (^{2α} CH)	3.52 (^{2a} CH, m)	_	
		72.58 (⁴ CH)	3.40 (⁴ CH, m)	_	
		63.70 (⁶ CH)	3.74 (^{5β} CH, m	=	
		98.92 (^{1α} CH)	4.66 (^{1a} CH, d)	7.86	
13	Glutamine	29.08 (C ^β)	2.14 (H ^β , m)	_	
		57.09 (C ^a)	3.76 (H ^a , t)	6.40	
14	Glycine	44.34 (C ^a)	3.55 (H ^a , s)	_	
15	Homoserine	35.12 (C ^β)	2.02 (H ^β , m)	_	
16	Isoleucine	17.54 (C ^Y)	1.00 (H ^Y , d)	7.01	
		26.83 (C ^Y)	1.24 (H ^Y , m)	-	
17	myo-Inositol	77.0	3.3 (CH,t)		
		73.8	3.6(CH, t)		
18	Mannitol	65.98	3.85 (CH ₂ , dd)		
		73.48	3.76(CH, d)		
19	Pyroglutamate	27.9 (C ^β)	2.4 (H ^Y ,m)	_	
		60.9 (C ^α)	$4.2(H^{\alpha}, dd)$	9.0, 5.9	
20	Pyruvate	29.33 (CH ₃)	2.36 (CH ₃ , s)	_	
21	Succinate	37.14 (CH ₂)	2.39 (CH ₂ , s)	_	
22	Sucrose	72.1 (³ CH)	3.48 (³ CH, m)	-	
		64.2 (¹ CH ₂)	3.69 (¹ CH ₂ , s)	_	
		79.8 (³ CH)	4.22 (³ CH, d)	8.82	
		95.1 (¹ CH)	5.42 (¹ CH, d)	3.90	
23	trans-Aconitate	40.4 (CH ₂)	3.43 (CH ₂ , s)	-	
		134.0 (CH)	6.56 (CH, s)	_	
24	Trigonelline	148.2 (² CH)	9.10 (CH, s)	=	

Table 1 list of metabolites identified in the polar extract of A. pubescens shoot (Continued)

- Compound name		¹³ C (ppm) from HSQC	¹ H (ppm) (functional group or specific H, multiplicity)	Coupling constant J (Hz)	
		147.4 (^{4,6} CH)	8.84 (CH, t)		
		130.5 (⁵ CH)	8.08 (CH, t)	8.80 7.37	
		51.05 (¹ CH ₃)	4.4 (CH ₃ , s)		
25	Trimethylamine	47.3 (CH ₃)	2.87 (CH ₃ , s)	_	
26	Valine	20.83 (C ^v)	0.97 (H ^y , d)	7.02	
		19.53 (C ^Y)	1.03 (H ^Y , d)	7.00	
27	Xanthine	140.3 (CH)	7.90 (CH, s)	-	

bands. Pair-wise analysis indicated a similarity index ranged from 0.97 to 0.98 between the mother and the in vitro raised plants (Table 4).

The ISSR-based genomic analysis gave a total of 66 bands where 59 of them were monomorphic (Table 3). Band sizes ranged from 4220 to 464 PB. The maximum number of amplified fragments (13) was recorded by UBC-849C primer, while primers UBC-834T and UBC-848A produced only 3 bands each. Jaccard's similarity coefficient analysis indicated a similarity index of 0.92–0.95 between the mother and the in vitro regenerated plants (Table 5).

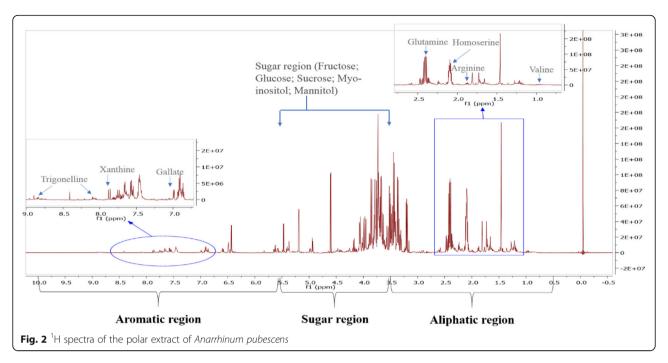
Discussion

Metabolic profiling of the wild plant shoots

In this study, we successfully identified 27 metabolites in the polar extract of the wild plant shoots which possess nutritional and medicinal values. Here, amino acids like isoleucine, valine, alanine, arginine, glutamine, and betaine have been identified. These amino acids have been previously reported in other members of the family Plantaginaceae, e.g., alanine, arginine, glutamine, isoleucine, and valine have been identified in *Plantago ovata* leaves [30]. Also, arginine, glutamine, isoleucine, and glycine were the most abundant amino acids in *Veronica*



Fig. 1 Anarrhinum pubescens wild plant



teucrium [31]. Isoleucine and valine are essential branched amino acids, which cannot be de novo synthesized by the human body and should be supplied in food. Isoleucine and valine have been reported to stimulate muscle protein synthesis. Isoleucine has been proved to encourage the uptake of glucose in muscle and

glucose oxidation in the whole body; moreover, it can decrease the gluconeogenesis process in the liver [32]. Arginine has been used for regulation of blood pressure and the immune system [33]. Betaine has a role in the treatment of many human diseases including obesity, inflammation, diabetes, cancer, and Alzheimer's [34].

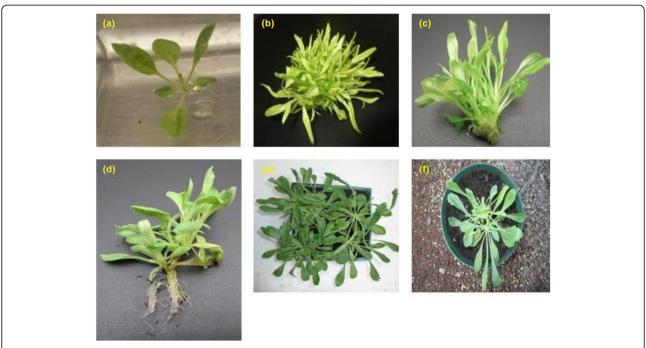


Fig. 3 Developmental stages of the in vitro propagation of *Anarrhinum pubescens*. A 4-week-old seedling (a), micropropagated shoots using BAP 1 mg/L + NAA 0.05 mg/L (b), organogenic shoots using BAP 0.05 mg/L + NAA 0.05 mg/L (c), rooted-shoots using MS-B₅ with 0.1 IAA mg/L (d), 10-week-old greenhouse-acclimatized plantlets (e, f)

Table 2 Effect of different cytokinin (BAP and Kn) concentrations in combination with 0.05 mg/L NAA on shoot multiplication from different explant sources of *A. pubescens*

Growth regulators (mg /L)			Explant source						
BAP	Kn	NAA	Nodal cuttings		Leaf segments				
			Shoot response* (%)	No. of shoots*	Shoot response* (%)	No. of shoots*			
0.5	-	0.05	100 ^a	13.85 ± 0.7 ^c	81.5 ^{ab}	9.3 ± 0.6 ^a			
1	-	0.05	100 ^a	27.67 ± 1.4^{a}	88.8 ^{ab}	7.1 ± 0.7^{b}			
2	-	0.05	100 ^a	23.19 ± 0.5 ^b	77.8 ^b	7.4 ± 0.2^{b}			
3	-	0.05	100 ^a	$13.33 \pm 1.6^{\circ}$	48.2 ^c	$3.7 \pm 0.6^{\circ}$			
4	-	0.05	100 ^a	12.67 ± 1.7 ^c	85.2 ^{ab}	8.2 ± 0.2^{ab}			
-	0.5	0.05	96.3 ^a	3.6 ± 0.2^{e}	74 ^c	1.2 ± 0.6^{d}			
-	1	0.05	100 ^a	5.8 ± 0.6^{de}	7.4 ^d	0.4 ± 0.3^{d}			
-	2	0.05	100 ^a	8.3 ± 1.7^{d}	66.7 ^c	2.3 ± 0.2^{cd}			
-	3	0.05	74.1 ^b	5.8 ± 2.4^{de}	70.4 ^c	1.2 ± 0.4^{d}			
-	4	0.05	92.6 ^a	5.1 ± 0.8^{de}	90.3 ^a	6.5 ± 1.4^{b}			

*Values represent mean ± SEM; numbers in columns followed by same letters are not significantly different according to Fisher pairwise comparison (p ≤ 0.05)

In our study, we identified the presence of myoinositol, sucrose, mannitol, fructose, and glucose in the polar plant extract. Glucose, fructose, sucrose, and mannitol have been reported in other species in the family Plantaginaceae [35]. Mannitol is a sugar alcohol that increases the tolerance of the plant to biotic and abiotic stress [36, 37]. We attribute the presence of this compound in *Anarrhinum pubescens* polar extract because the plant is growing in a stressful climatic condition.

The phenolic compound, gallic acid, and the alkaloids trigonelline and xanthine were identified in the aromatic region between $\delta 5.0$ and $\delta 10.0$. The presence of phenolic and alkaloid compounds in the family Plantaginaceae has been

widely reported [23, 38]. Gallic acid was reported in other Plantaginaceae species like *Plantago coronopus* L. subsp. *coronopus* [39] and in *Plantago lanceolata* [40]. Gallic acid has been reported to have antioxidant and neuroprotective activities [41]. It was shown to improve gut microbe activity and positively modify the immune response [42], along with anti-cancer and anti-inflammatory properties [43]. Trigonelline is well known for its medicinal activity against diabetes and nervous system diseases [44, 45].

Vitamins and organic acids like choline and acetic acid have been identified in the plant extract. Choline is an essential nutrient for humans and its deficiency was shown to cause liver damage [46]. It is involved in lipoproteins,

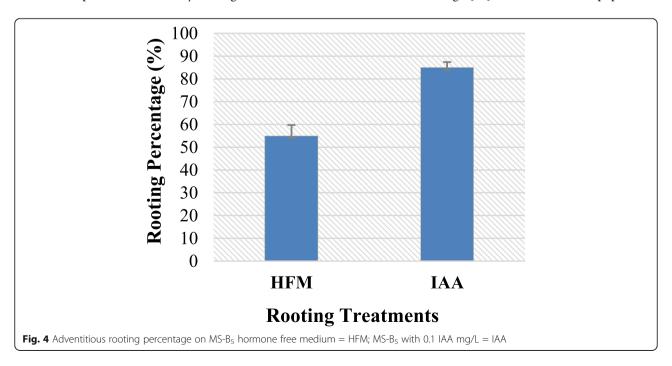


Table 3 List of RAPD and ISSR primers' names, sequences, annealing temperatures, bands sizes range, and number of bands amplified including monomorphic (MB) and polymorphic bands (PB)

Molecular marker	Primer name	Primer sequence 5' to 3'	Temp. ([°] C)	Size range (bp)	Total amplified bands	MB	РВ
RAPD	OPA-03	AGTCAGCCAC	39.5	1787–921	4	4 12 6 3 11 7 4 6 4 61 12–4 6.1 7 7 3 12 8 4 3 3 4 8 59 12–3	0
	OPA-04	AATCGGGCTG	39.5	2667–408	13	12	1
	OPH-07	CTGCATCGTG	39.5	2200-613	6	6	0
	OPI-07	CAGCGACAAG	39.5	1926–617	4	3	1
	OPK-07	AGCGAGCAAG	39.5	1788–690	11	11	0
	OPM-07	CCGTGACTCA	39.5	3860-840	7	7	0
	OPO-07	CAGCACTGAC	39.5	1642–538	4	4	0
	OPS-07	TCCGATGCTG	39.5	1813–516	4	4	0
	OPW-07	CTGGACGTCA	39.5	2114–625	6	6	0
	OPY-07	AGAGCCGTCA	39.5	1919–621	4	4	0
	Total				63	61	2
	Range			3860-408	13–4	12-4	2-0
	Average				6.3	6.1	0.2
ISSR	UBC-842C	GAGAGAGAGAGAGACG	59.9	2013-510	7	7	0
	UBC-842T	GAGAGAGAGAGAGATG	57.6	1617–567	9	7	2
	UBC-846A	CACACACACACACACAAT	55.3	1717–520	4	3	1
	UBC-849C	GTGTGTGTGTGTCA	57.6	2682–678	13	12	1
	R(CA)7	GCACACACACACA	53.4	1410–580	8	8	0
	UBC-834C	AGAGAGAGAGAGAGCT	57.6	4220-734	5	4	1
	UBC-834T	AGAGAGAGAGAGAGTT	55.3	1111–464	3	3	0
	UBC-848A	CACACACACACACACAAG	57.6	1450–687	3	3	0
	UBC-860A	TGTGTGTGTGTGAA	55.3	2170-730	5	4	1
	(AC)8TG	ACACACACACACACTC	57.6	2738-1540	9	8	1
	Total				66	59	7
	Range			4220–464	13–3	12-3	1-0
	Average				6.6	5.9	0.7

blood, and membrane lipid structure, and enhances liver function, neurodevelopment, and muscle movements [47].

In vitro regeneration

Plant tissue culture techniques are ideal tools for conservation and propagation of rare and threatened plants. Many endangered plants have been conserved using different techniques like micropropagation and organogenesis through callus culture [48]. In the present work, the effect of different concentrations of BAP or Kn, in combination with NAA on micropropagation via nodal cuttings and leaf explants was investigated, and the data showed that BAP-treated cultures were significantly superior in de novo shoot induction and proliferation. BAP was the best cytokinin in shoot induction and proliferation from different plants and explants, e.g., *Bambusa balcooa* nodal cuttings [49], *Punica granatum* leaf explant [50], and in plant regeneration from *Bauhinia variegate* seeds [51]. In the family Plantaginaceae, BAP was more effective

than Kn for shoot multiplication from *Bacopa monnieri* and *Plantago asiatica* in vitro cultures [52, 53]. A combination of BAP and NAA was more effective in organogenic shoots proliferation of *Mondia whitei* [54].

In this study, $MS-B_5$ medium supplemented with IAA was the best in adventitious roots induction (85%) compared to $MS-B_5$ hormone free medium. IAA has been found to play a central role in adventitious root formation in plants [55, 56]. Genes responsible for adventitious root formation in lotus seedling was regulated by IAA [57]. IAA was the most effective auxin in adventitious roots formation in apples [58].

Genetic fidelity assessment of regenerated plants

Genetic variation is expected in in vitro regenerated plants, i.e., somaclonal variation [59]; so, it is important to assess the genetic stability of regenerated plants. In this work, RAPD and ISSR molecular markers were used to evaluate the genetic fidelity of the micropropagated

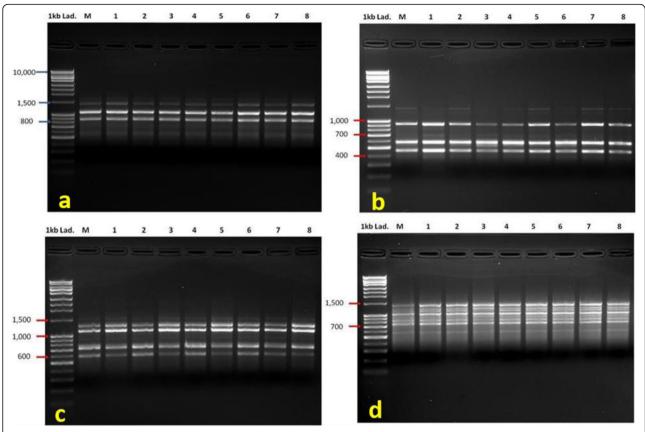


Fig. 5 DNA amplification profile of mother and in vitro propagated plants of *A. pubescens* generated by RAPD and ISSR primers. RAPD primer OPA-03 (**a**), RAPD primer OPI-07 (**b**), ISSR primer UBC-860A (**c**), and ISSR primer R(CA)7 (**d**). 1st lane is 1 kb DNA ladder, lane M-mother plant, lane 1–8 in vitro propagated plants

clones. RAPD primers showed 0.97 to 0.98 similarity between the mother and the in vitro regenerated plants, while ISSR-based genomic analysis indicated a similarity index of 0.92–0.95 between the mother and the in vitro regenerated plants. Assessment of genetic stability using DNA molecular markers is a rapid and reliable method for determining genetic variation

Table 4 Jaccard's similarity coefficient for mother and in vitro regenerated plants of *Anarrhinum pubescens* based on RAPD markers

	М	l1	12	13	14	15	16	17	18
М	1								
11	0.98	1							
12	0.97	0.98	1						
13	0.97	0.98	1	1					
14	0.98	1	0.98	0.98	1				
15	0.98	1	0.98	0.98	1	1			
16	0.97	0.98	1	1	0.98	0.98	1		
17	0.98	1	0.98	0.98	1	1	0.98	1	
18	0.98	1	0.98	0.98	1	1	0.98	1	1

among and within plant species [60, 61]. RAPD and ISSR molecular markers were efficient in assessment of the genetic diversity in other members of the family Plantaginaceae, e.g., *Plantago ovata* [55] and *Plantago major* [62]. RAPD analysis has been used to evaluate genetic diversity within the wild *Anarrhinum pubescens* population [63].

Table 5 Jaccard's similarity coefficient for wild and in vitro regenerated plants of *A. pubescens* based on ISSR markers

	М	l1	12	13	14	15	16	17	18
М	1								
11	0.95	1							
12	0.92	0.94	1						
13	0.94	0.98	0.95	1					
14	0.94	0.95	0.98	0.94	1				
15	0.94	0.98	0.92	0.97	0.94	1			
16	0.92	0.94	0.97	0.92	0.98	0.95	1		
17	0.94	0.98	0.92	0.97	0.94	1	0.95	1	
18	0.94	0.95	0.98	0.94	1	0.94	0.98	0.94	1

Conclusions

Metabolic profiling of the *A. pubescens* wild plant shoot has been performed in this study which can help in chemotaxonomy and in discovering new medicinal/nutritional uses of the plant shoots. We also developed an efficient, reproducible, and rapid protocol for the in vitro regeneration of the plant for the purpose of conserving its genetic resources as the resulting clones exhibited a high degree of genetic similarity to the mother plant.

Abbreviations

NMR: Nuclear Magnetic Resonance; HSQC: Heteronuclear Single Quantum Coherence; BAP: 6-Benzylaminopurine; NAA: Naphthaleneacetic acid; Kn: Kinetin; 2,4-D: 2,4 dichlorophenoxy acetic acid; RAPD: Random Amplified Polymorphic DNA; ISSR: Inter Simple Sequence Repeat; MS-B5: Murashige and Skoog Medium with B5 vitamins

Acknowledgements

Authors are grateful for St. Catherine Protectorate staff members for helping in collecting and identifying the plant material.

Authors' contributions

All authors contributed to the study conception and design. AA and EM performed the experiments, collected the data, analyzed the results, and wrote the first draft of the manuscript. AB and KC revised the collected data and the final manuscript. All authors read and approved the final manuscript.

Funding

Authors are gratefully acknowledged SC-INBRE (2 P20 GM103499) and NSF HBCU-UP (HRD-1332516) for the financial support of the NMR material.

Availability of data and materials

All data generated or analyzed during this study are included in this published article.

Declarations

Ethics approval and consent to participate

Not applicable.

Consent for publication

Not applicable.

Competing interests

The authors declare that they have no competing interests.

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Received: 28 April 2021 Accepted: 16 July 2021 Published online: 26 July 2021

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