



Growth promoting effects of endophytic fungus *Piriformospora indica* in small cardamom (*Elettaria cardamomum* Maton)

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Abstract

Piriformospora indica, an endophytic fungus of Sebaciales, colonizes the roots of a wide range of host plants and provides various benefits to the plants. Cardamom (*Elettaria cardamomum* Maton) is an economically valuable spice crop of the tropics. In this work, we describe differentially expressed transcripts responding to *P. indica* root colonization in small cardamom for elucidation of molecular basis of growth and development. During the study, a wild genotype of cardamom was propagated under in vitro conditions using full strength Murashige and Skoog medium supplemented with 1 mg l⁻¹ BAP (for shoot induction) and basal MS liquid medium (for root induction). Cardamom plantlets were co-cultivated with *P. indica*. Microscopic observation confirmed the presence of *P. indica* inside the roots of cardamom plantlets. Growth parameters of control and *P. indica* colonized plantlets were observed for three months at an interval of 15 days. *P. indica* colonization resulted in a significant increase in the morpho-physiological traits of the host plant. The growth enhancement was visible after 15 days of co-culture. There was a significant increase ($p < 0.05$) in the number and length of leaves, height of the plant and chlorophyll content in *P. indica* colonized plants compared to non-colonized control plant. In addition to this, the expression levels of auxin, nitrate reductase, vegetative storage protein and phosphate transporter genes were upregulated by 3.45, 3.26, 1.62 and 1.19 times respectively by the co-cultivation of *P. indica* in cardamom plantlets.

Keywords Cardamom · Real-time PCR · Auxin · Nitrate reductase · Vegetative storage protein · Phosphate transporter

1 Introduction

Small cardamom (*Elettaria cardamomum* (L.) Maton), also known as the “Queen of spices” is one of the most important spice crops in the world. It is the third expensive spice globally after vanilla and saffron (Tangjang and Sharma 2018). It belongs to family Zingiberaceae and has believed to be originated in the moist evergreen forests of the Western Ghats region of Southern India. Cardamom is a shade-loving crop plant cultivated at an altitude of 600–1200 m above mean sea level (MSL) with an annual rainfall of 1500–4000 mm and

a temperature of 10 to 35 °C (Madhusoodanan et al. 2002). Damp, loamy soil rich in humus is well suitable for its cultivation. As it's a shade lover, 40 to 60% shade is requisite for their proper growth and flowering.

Parched ripe fruits (capsules) of cardamom have high economic value with a variety of medicinal properties. The essential oil and other bioactive molecules in the fruits have many curative effects including anti-inflammatory, anticancer, anti-oxidant, antibacterial, antifungal, antiviral, anti-diabetic, gastro-protective and analgesic activities (Al-Zuhair et al. 1996; Verma et al. 2009; Khan et al. 2011; Winarsi et al. 2014). Mishra et al. (1991) found out that antimicrobial activities of cardamom extract are mainly due to the terpenoids. In 2020, Ashokkumar et al. reviewed that terpenoids, flavonoids, anthocyanins, alkaloids and phenolics of capsules are responsible for controlling cardiovascular, pulmonary, kidney and lung associated disorders. Along with this, a variety of cardamom- flavoured food products are available in the markets including tea powder, biscuits, baked food items, confectionaries, etc.

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The axenically cultivable root endophytic beneficial fungus *Piriformospora indica*, isolated from the desert soil of northwest India in the state of Rajasthan, interacts with many plant species and promotes their growth (Weiss et al. 2004; Varma et al. 2012). It also enhances plant resistance to biotic stresses such as bacterial, fungal, viral and nematode diseases and abiotic stresses like heavy metals, salinity and drought (Deshmukh and Kogel 2007; Sherameti et al. 2008a, b; Daneshkhah et al. 2013; Lakshmi Priya et al. 2016; Li et al. 2017; Varkey et al. 2018; Athira and Anith 2020; Paul et al. 2021). Significant increase in growth and yield of many medicinal plant species was recorded on inoculation with *P. indica* (Rai et al. 2001; Kilam et al. 2015). Xu et al. (2017) reported that the co-cultivation with *P. indica* improved the growth of maize plants by enhancing phenotypic traits like plant height and leaf number. The mean number of leaves per plant and the leaf area per plant in the *P. indica* inoculated *Piper nigrum* recorded an increase compared to that of the control plants starting from the first month, to one year after transplanting (Anith et al. 2018). *Bacopa monnieri* co-cultivated with *P. indica* exhibited an enhanced growth, elevated bacoside endogenous level, antioxidant activity and nuclear hypertrophy (Prasad et al. 2013).

Over 100 years ago, Haberlandt conceptualised the idea of plant tissue culture and laid the foundation for the production of plant cells, tissues, and organs in culture. In addition to its utility as a research tool, plant tissue culture techniques have increasingly gained significant industrial importance in the fields of plant improvement, disease eradication, secondary metabolite production, and plant propagation (Hussain et al. 2012). Under controlled conditions, a single explant can be reproduced into many thousand plants in a relatively short amount of time and space, regardless of the season or weather, all year long. Because of the rapid rate of multiplication and low requirements for the initial plant population and available space, endangered, threatened, and unusual species have been successfully cultivated and conserved through micropropagation (Idowu et al. 2009).

Micropropagation opens up a lot of possibilities for rapid vegetative propagation of elite clones that are free of systemic diseases like viruses. Cardamom elite clones can be selected and rapidly multiplied using tissue culture technique (Hemavathy and Balaji 2010). Micro propagation technology has gained traction in India, reclaiming the global monopoly in cardamom production. The first report of cardamom tissue culture was published in 1982, claiming that callus cultures could regenerate plants (Rao et al. 1982). In vitro methods for the clonal propagation of cardamom from vegetative buds have been standardised (Reghunath and Gopalakrishnan 1991). Many commercial laboratories are using micropropagation techniques for large scale production of cardamom. The field evaluation of tissue cultured cardamom showed that the micropropagated plants

performed on par with suckers (Chandrappa et al. 1997). The cardamom shoot growth patterns were studied on 45 media treatments, with the best shoot proliferation reported on Murashige and Skoog (MS) medium (Murashige and Skoog 1962) supplemented with and 2.32 M kinetin (Kn) for root induction and 4.4 M 6-benzylaminopurine (BAP) for shoot induction and 5.8 shoots regenerated per explant. On full strength MS basal medium, regenerated shoots rooted well, creating 3.5 roots per explant with an average root length of 4.3 cm in 4 weeks (Malhotra et al. 2021). Mixture of garden soil, sand and vermiculite in equal proportions was determined to be the best for planting, with 92% establishment rate (Babu et al. 1999).

A comparative study was undertaken to realise the role of *P. indica* in the growth response of cucumber (*Cucumis sativus*), okra (*Abelmoschus esculentus*), egg plant (*Solanum melongena*) and chilli (*Capsicum annum*) under in vitro condition. These *P. indica*-colonised plants exhibited a considerable improvement ($P < 0.001$) in root lengths, shoot number and shoot length. Additionally, a concomitant rise in chlorophyll concentration was seen, suggesting a potential improvement in photosynthetic efficiency (Shukla et al. 2022). *P. indica* was found to have prospective uses in the banana tissue culture sector. In comparison to controls, *P. indica* colonised banana seedlings had greater plant height, more and longer roots, increased shoot fresh weight, greater leaf breadth and length, and larger stem circumference at 4 weeks following inoculation. The chlorophyll a, chlorophyll b, total chlorophyll, and carotenoid levels of banana leaves were likewise markedly improved by the *P. indica* treatments (Li et al. 2019). Colonization by *P. indica* is expected to improve the overall growth and yield characteristics of cardamom as reported in other studies (Rai et al. 2001; Kilam et al. 2015; Xu et al. 2017; Weiss et al. 2004; Varma et al. 2012). Present study reports for the first time the genetic basis involved in the growth and development of *E. cardamomum* induced by the symbiotic fungus *P. indica*.

Very little is known about the exact mechanisms behind growth promotion in plants upon *P. indica* colonization. High-affinity phosphate (PO_4^{3-}) transporter (PiPT) found in *P. indica* is found to enhance plants' ability to absorb phosphate (Pederson et al. 2013), but without *P. indica* similar enhancement could not be observed (Bakshi et al. 2017). According to some recent research, *P. indica* modifies metabolic pathways during colonisation, such as a rapid rise in auxin levels during early host recognition, which is crucial for reprogramming the root development (Conchillo et al. 2021; Meents et al. 2019).

In this study, differential expression of four growth-related genes namely *AUX1*, *NIA1*, *VSP* and *PHT1* were analyzed in *P. indica* colonized cardamom plants in comparison with the non-colonized control plants. These four genes were selected based on the observations made in previous studies

reported elsewhere. In root and shoot tissues, *AUX1* activity is necessary for both polar and phloem-based IAA transport routes (Leyser 2006). It was previously reported that co-cultivation with *P. indica* resulted in a massive transfer of nitrogen from the agar plates to the plants' aerial parts and this action was linked to the activation of the NADH-dependent nitrate reductase (*NIA1*), a critical enzyme in the uptake of nitrate in plants (Sherameti et al. 2008a, b). Lin et al. (2019) reported that the expression of *VSP* (Vegetative storage protein gene) marked an increment after infection of the roots with *Ralstonia solanacearum* post infection in comparison with that of uncolonized controls. Plants have numerous transporters, including the *PHT1*, *PHT2*, *PHT3*, and *PHT4*, which are involved in inorganic phosphate (Pi) uptake by the root and remobilization in plants for being adapted to low phosphate environments. Among the four major phosphate transporters, *PHT1* is responsible for phosphate uptake from soil by root cells, which is the first and most important stage in the Pi assimilation process (Liu et al. 2011). Considering these findings, genes namely *AUX1*, *NIA1*, *VSP* and *PHT1* were selected for the present study.

The major objective of the study was to identify differentially expressed transcripts responding to *Piriformospora indica* root colonization in small cardamom for elucidation of molecular basis of growth and development.

2 Materials and methods

2.1 Source of culture

Initiated in vitro cultures of cardamom (wild genotype from the natural forest area in Edamalayar, Ernakulam district of Kerala, India.) were received from the Plant Tissue Culture laboratory of Biotechnology and Bioinformatics Division, Jawaharlal Nehru Tropical Botanic Garden and Research Institute, at the 1st subculture cycle and repeatedly subcultured to produce enough plantlets.

2.2 Propagation of in vitro cardamom culture

Initiated cultures (3 weeks old) were repeatedly subcultured to fresh Murashige and Skoog (MS) supplemented with 1 mg l^{-1} BAP (pH 5.8) for further shoot multiplication and elongation. Each subculture passage lasted for 3 weeks. After 2 weeks, shoot plantlets were transferred to full strength MS basal liquid medium (pH 5.8) for root induction. The cultures were kept in sterile conical flasks at $23 \pm 2^\circ\text{C}$ provided with 16 h/8 h photoperiod, light intensity of $25 \mu\text{mol m}^{-2} \text{ s}^{-1}$ using white fluorescent tubes and 55–65% relative humidity.

2.3 Fungal culture

P. indica was originally gifted by Dr. Ajit Varma; former Professor, School of Life Sciences, Jawaharlal Nehru University, New Delhi, India. It was maintained by sub culturing on Potato Dextrose Agar (PDA) at pH 7.0 and incubated in the dark at 27°C . For mass multiplication, fungal hyphal discs were transferred to Potato Dextrose Broth (PDB) and shaken at 110 rpm in a shaker with maintenance of the same growth conditions as above.

2.4 Co-cultivation

2.4.1 In vitro co-culture

Mycelial discs of 0.8 cm diameter from 20 days old *P. indica* maintained on PDA plates were placed near roots of three weeks old *E. cardamomum* plantlets maintained in MS liquid culture under aseptic condition. It was then incubated for 35 days and provided 16 h: 8 h light/dark photoperiod, $23 \pm 2^\circ\text{C}$ temperature, 55–65% humidity and a light intensity of $25 \mu\text{mol m}^{-2} \text{ s}^{-1}$. In vitro plantlets maintained under identical conditions in MS liquid medium without *P. indica* co-cultivation served as control.

2.4.2 Ex vitro co-culture

Potting mixture containing sand, soil and vermiculite in 1:1:1 ratio (pH 6.5) was mixed with *P. indica* mycelium at the rate of 1% (w/v) (Anith et al. 2015). Primary hardening of 40 days old plantlets was done in small plastic cups (8×6) filled with the above medium. Inoculated plants were kept inside a mist house. After three weeks of primary hardening, the plants were transferred to larger plastic pots (18.5×14) containing sand, soil, and vermiculite in a 2:2:1 ratio and kept for secondary hardening in a mist house with 85% humidity. The plants were observed for growth parameters such as leaf numbers, leaf length and plant height for three months at an interval of every 15 days. Plants maintained under identical conditions in potting mixture containing sand, soil and vermiculite (without *P. indica*) served as control.

2.5 Confirmation of root colonization

2.5.1 Microscopic observation

Roots of *P. indica* inoculated and control in vitro grown plants were collected and were carefully cleaned in tap water before being chopped into one-cm pieces. They were boiled 100°C in made 10% KOH solution for 5 min to soften them. The root pieces were washed in distilled water, then acidified with 1 M HCl for 5 min, followed by staining for 15 min

in 0.02% lactophenol-trypan blue. Before viewing under a compound bright field microscope (Olympus CX43, Japan), the staining was removed using lactophenol solution. Chlamydo spores in the root cortex cells were interpreted as a sign of root colonization.

2.6 Estimation of chlorophyll content

Chlorophyll content of cardamom plantlets maintained under in vitro condition (after 30 days of co-cultivation) and *ex vitro* condition (after 90 days of co-cultivation) was determined. Leaf sample (250 mg) was macerated with 10 mL of 80% acetone using a pestle and mortar. The extract was centrifuged at 3000 rpm for 10 min. The supernatant was transferred into a 25 mL volumetric flask and final volume was made up to 25 mL using 80% acetone. Color intensity of the green pigment was read at 645 nm, 663 nm and 652 nm. Chlorophyll a, chlorophyll b and total chlorophyll were calculated and expressed in mg of pigments/gram of fresh weight (Arnon 1949).

2.7 Gene expression analyses

2.7.1 RNA extraction and cDNA synthesis

RNA was isolated from 100 mg of young leaf (6th leaf) of both control and *P. indica* treated in vitro plantlets (9 weeks old after second subculture) using a modified CTAB approach in combination with the RNeasy Plant Mini Kit method (Nadiya et al. 2015). The integrity and quality of RNA were checked using Agarose gel (1.2%) electrophoresis. The N50 NanoPhotometer (IMPLEN, Germany) was used to determine both the concentration and purity of RNA (A_{260}/A_{280} and A_{260}/A_{230}). Takara PrimeScript™ RT reagent Kit (Takara Bio Inc., Japan) was used to synthesize cDNA from RNA. The first strand cDNA synthesis was carried out

according to the manufacturer's instructions. A total of 10 µL reaction mixture (Prime script buffer (5 X) – 2 µL; Primerscript RT – 0.5 µL; OligodT primer (50 µM) – 0.5 µL; Random hexamer (100 µM) – 0.5 µL; RNA (100 ng/µL) – 2.5 µL; RNase free water – 4 µL) was made and incubated for 15 min at 37 °C.

2.7.2 Primers used in the study

Out of the four selected genes studied, primer sequences of three genes such as *NIA1* and *PHT1* (Bandyopadhyay et al. 2022) and *VSP* (Lin et al. 2019) were taken from previous studies (Table 1). *UBCE* and *AUX1* gene sequences were developed from the transcriptome sequences of cardamom deposited in NCBI (Accession ID SRX1141276 and SRX1141272). These sequences were aligned using the software 'MEGAX' (Kumar et al. 2018). Primers for the real time PCR analyses were designed using IDT's primer quest tool (<http://www.idtdna.com>). Then, using Net Primer and NCBI Primer BLAST, the quality and specificity of the primers were tested. Running a normal PCR with cardamom cDNA as template allowed for the preliminary selection of primers for qPCR experiments. Using gradient PCR, the annealing temperature for all of these primers was standardised at 60 °C.

2.7.3 Real-Time PCR analysis

Using the StepOnePlus™ Real Time PCR equipment (Applied Biosystems, USA), the expression patterns of *AUX1* (Auxin responsive gene), *NIA1* (gene involved in nitrate transport-Nitrate reductase 1), *PHT1* (gene involved in phosphate uptake - Phosphate transporter1) and *VSP* (Vegetative storage protein gene) were studied. The relative change in gene expression was calculated using the Comparative Cq method utilising the Ct values

Table 1 Primer sequences used for this study

Sl. No.	Gene name	Direction	Sequence (5'-3')	Source
1	UBC	F	CCAGTGTGGCAATGTTCTC	Primers developed from transcriptome of cardamom
		R	GAGTCCAGCCCTCCAAATC	
2	AUX1	F	CAGCTCGATAACCTCGACTTC	Lin et al. 2019
		R	TACCGCTAAAGAGGGCAAAC	
3	VSP	F	CATAGACTTCGACACGGTGC	Bandyopadhyay et al. 2022
		R	AGGAGGGTATCATCTAGGTCA	
4	NIA1	F	CGGAGAGCTACTACCATTAC	Bandyopadhyay et al. 2022
		R	GTTTATCATGTACTCCGGCT	
5	PHT1	F	GTACTCAAACAAGAAGACGC	Bandyopadhyay et al. 2022
		R	AGGAAGATGCTAGAGACGAT	

F: Forward primer; R: Reverse primer

UBC - ubiquitin C, AUX1 – Auxin, VSP - Vegetative Storage Protein, NIA1 - Nitrate reductase, PHT1 - Control-Phosphate Transporter

obtained from Real time analysis. The real time gene expression data was normalized using the house keeping gene ubiquitin C (UBC).

2.8 Statistical analysis

For analysis of growth parameters, experiments were laid out in Completely Randomized Design (CRD). The mean values were compared using Two Sample *Student's t* test ($p < 0.05$), and analysis of data was carried out using the Grapes 1.0.0 web application designed by Kerala Agricultural University (<https://www.kaugrapes.com/>).

3 Results

3.1 Colonization of *P. indica* in cardamom root

Intracellular hyphae were observed at 10 days and sporulation at 20 days after inoculation. Morphologically, the hyphae were straight and hyaline, and the surface of the hyphal walls was smooth. Pear shaped chlamydospores were observed within the cortical cells of root tissues of the inoculated in vitro grown plants when detection was carried out at 20 days after inoculation (Fig. 1). No colonization was observed in the uninoculated control plants.

3.2 Growth performance of *P. indica* co-cultured cardamom plants

The symbiotic association of cardamom with *P. indica* resulted in a significant increase in the morpho-physiological traits of the host plant. The growth enhancement was visible even from 15th day of co-culture. There was a significant increase ($p < 0.05$) in the number of leaves, length of leaves and height of the plant of colonized plants compared to control plant (Table 2; Figs. 2 and 3). Three months after transplanting, the *P. indica* inoculated plants

recorded a mean leaf number of 20.6 per plant; average leaf length of 21.9 cm and mean plant height of 38.1 cm whereas that of the control plants were 12.8, 13.9 and 27.7 cm respectively.

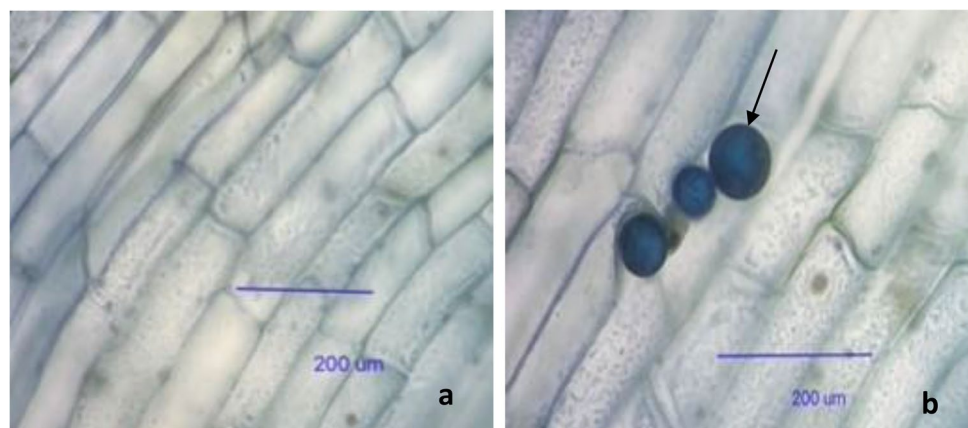
3.3 Effect of *P. indica* on chlorophyll content

A significant enhancement ($p < 0.05$) was obtained in the chlorophyll a, chlorophyll b and total chlorophyll contents in *P. indica* colonized plants when compared to non-colonized control plants (Table 3). The percent increase in photosynthetic pigment content in *P. indica* colonised plants above control was 19% (chlorophyll a), 60% (chlorophyll b), and 22% (total chlorophyll) in in vitro conditions. Similarly, the increase in photosynthetic pigment content in *P. indica* colonised plants compared to control was 92% (chlorophyll a), 86% (chlorophyll b), and 84% (total chlorophyll) in *ex vitro* conditions.

3.4 Real-time PCR analyses for the validation of gene expression

The relative enhancement in the transcript levels of the aforementioned genes in both the control and *P. indica* co-cultured samples is shown in Fig. 3. The most over-expressed genes were *NIA1*, which plays an irreplaceable role in nitrogen assimilation in plant cells as well as serving as a key enzymatic source of nitric oxide, which governs plant growth and *AUX1* which is involved in the auxin transport. Samples treated with *P. indica* showed a 3.45 and 3.26 times increases in the transcript levels of *NIA1* and *AUX1*. The gene which encodes *VSP* marked a slight up-regulation by 1.19 times in transcript level. Similarly, *PHT1*, which is involved in Pi uptake by the root and remobilization in plants, marked a 1.16 times expression at transcript level.

Fig. 1 **a** Stained root of uninoculated control cardamom plant. **b** Chlamydospores (black arrow) of *P. indica* within the cortex cells of cardamom root (400 x)



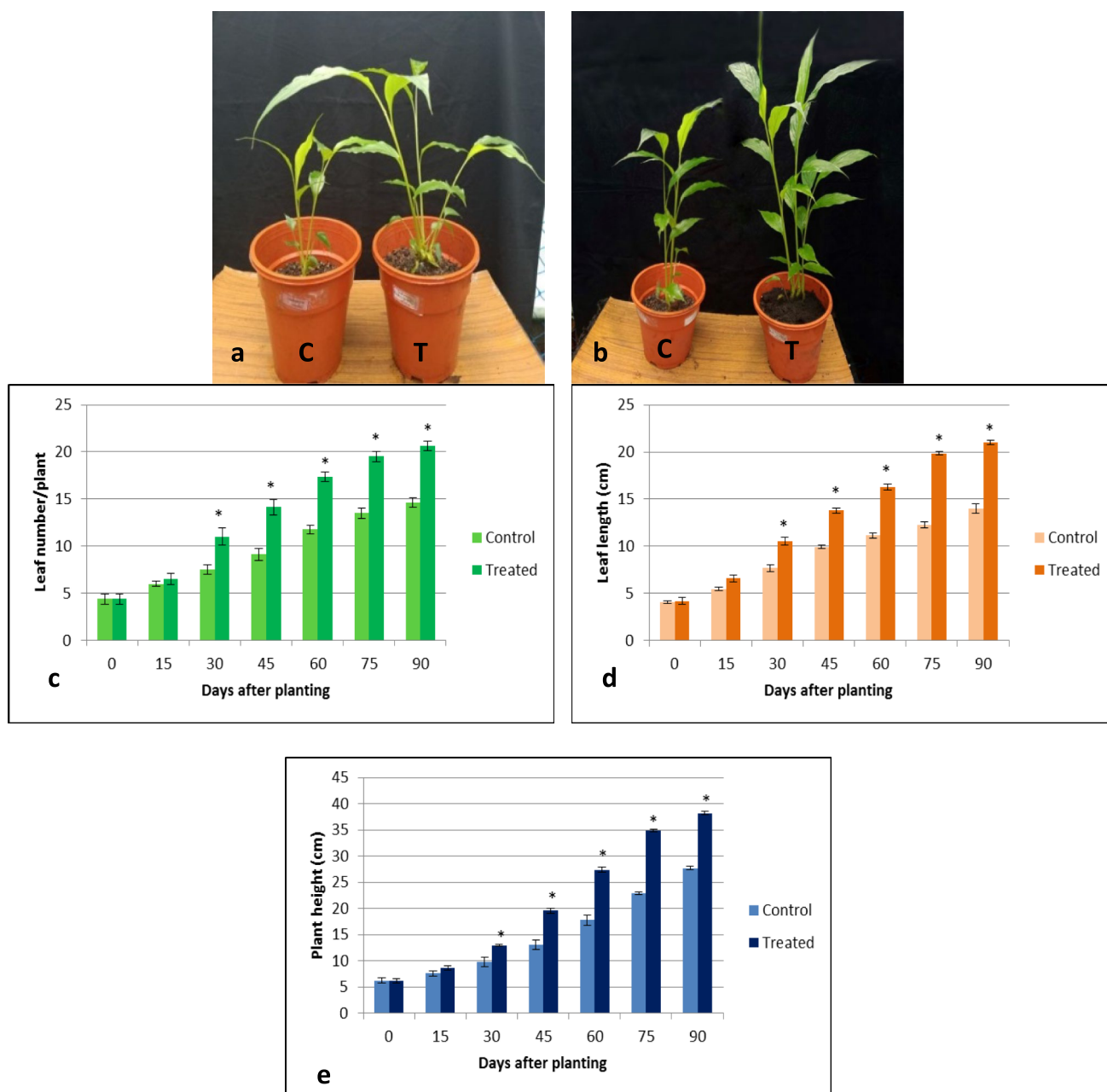


Fig. 2 Plant growth promoting effect of *P. indica* on cardamom plants (C- Control; T- Treated) – **a** Day 45, **b** Day 90, **c** Effect of *P. indica* colonization on the total number of leaves (**d**) leaf length (**e**) plant height. The error bars indicate standard deviation values

4 Discussion

Endophytes are ubiquitous which grow in plant tissues and offers many growth promoting effects to the host plants (Ali-kulov et al. 2022; Choudhury et al. 2023; Urumbil and Anilkumar 2021). The effects of root colonization by *P. indica* on growth have been reported in spice crops like black pepper, chilli (Anith et al. 2011; Nandana and Anith 2020; Lekshmi et al. 2022). However, in cardamom, there is no report on the co-cultivation with *P. indica* and its effects on growth

promotion. In the present study we used in vitro raised cardamom plantlets as the planting material for growth evaluation studies, and colonization by the root endophyte was confirmed through microscopic detection method. All the planting materials used were clonally propagated from a single mother plant and thus it was ensured that the genetic base of the plants remained the same, but due to the microbial treatment, growth pattern of the control and treated plants differed. Assessment of plant growth promotion was done by taking observations on the number of leaves, length of leaves

Fig. 3 Accumulation of different growth-related gene transcripts in cardamom as a result of *P. indica* co-cultivation. ***C-NIA1 - Control-Nitrate reductase, T-NIA1 - Treated-Nitrate reductase, C-VSP - Control-Vegetative Storage Protein, T-VSP - Treated-Vegetative Storage Protein, C-AUX1 - Control-Auxin, T-AUX1 - Treated-Auxin, C-PHT1 - Control-Phosphate Transporter, T-PHT1 - Treated-Phosphate Transporter. The error bars indicate ΔC_T standard error values

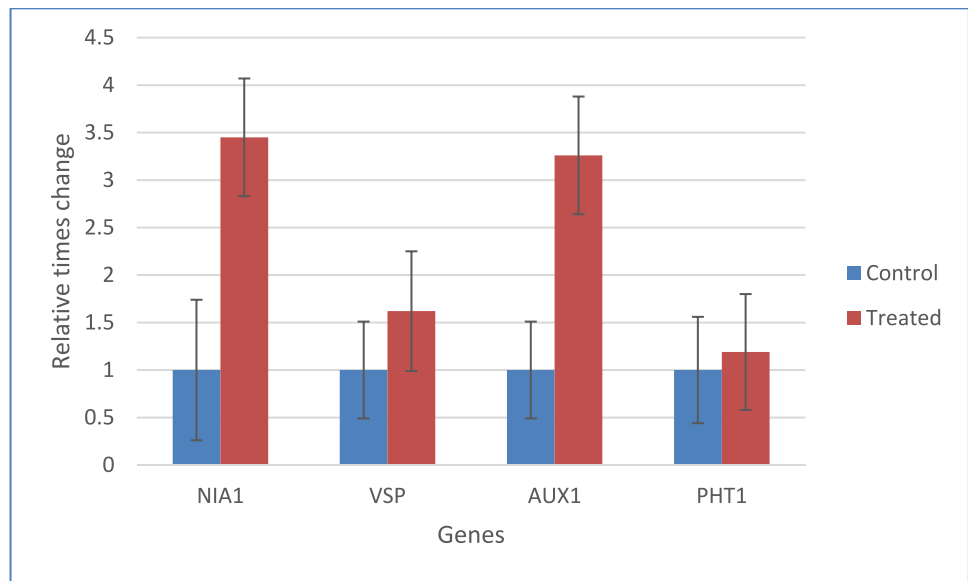


Table 2 Effect of *P. indica* colonization on chlorophyll content of cardamom under in vitro conditions

Sample		Chlorophyll a (mg/g FW) [#]	Chlorophyll b (mg/g FW) [#]	Total chlorophyll (mg/g FW) [#]
-Pi	Mean ± S.D	0.569 ± 0.003	0.159 ± 0.003	0.818 ± 0.006
	SE (mean)	0.002	0.001	0.004
+Pi	Mean ± S.D	0.680 ± 0.002*	0.269 ± 0.003*	1.001 ± 0.004*
	SE (mean)	0.001	0.005	0.002
P value		0.0001	0.0001	0.0001
% increase over control		19%	60%	22%

[#]After 30 days of co-cultivation

*Significant at $p < 0.05$, mean of 8 observations

Table 3 Effect of *P. indica* colonization on chlorophyll content of cardamom under ex vitro conditions

Sample		Chlorophyll a (mg/g FW) [#]	Chlorophyll b (mg/g FW) [#]	Total chlorophyll (mg/g FW) [#]
-Pi	Mean ± S.D	0.802 ± 0.03	0.338 ± 0.03	1.200 ± 0.08
	SE (mean)	0.002	0.001	0.005
+Pi	Mean ± S.D	1.540 ± 0.02*	0.629 ± 0.04*	2.269 ± 0.02*
	SE (mean)	0.001	0.002	0.007
P value		0.0001	0.0001	0.0001
% increase over control		92%	86%	84%

[#]After 90 days of co-cultivation

*Significant at $p < 0.05$, mean of 8 observations

and the plant height. Plant growth promotion was evidenced by the production of more number of leaves, increased leaf length and plant height in the *P. indica* inoculated plants compared to the control plants. Chlorophyll (Chl) is a crucial photosynthetic pigment for the plant, and it plays a significant role in determining photosynthetic capacity and, consequently, plant growth (Li et al. 2018). Plant metabolism is directly impacted by the amount of chlorophyll in leaves

through photosynthesis, growth, and development (Dai et al. 2009). Furthermore, formation and growth of plant structures in higher plants is influenced by light harvesting such as accumulation of food reserves in leaf base cells (Olive et al. 2007). With this perspective, it can be concluded that the increased leaf chlorophyll content may facilitate more harvesting of light thereby directly promote plant growth. The chlorophyll content in the leaves of *P. indica* inoculated

cardamom plants recorded an increase over the control plants under both *in vitro* and *ex vitro* conditions. Similar observations have been reported earlier on *Arabidopsis* (Sherameti et al. 2008a, b); cucumber, okra, chilli (Jisha et al. 2019); black pepper (Anith et al. 2018) by the co-cultivation of *P. indica*. Decrease in chlorophyll content in plant fresh leaves is considered as an indicator of photosynthetic incapability of plant machinery (Sakuraba et al. 2018). Primarily, chlorophyll a and b are necessary for the primary photosynthetic reaction (Croft et al. 2017). Abdelaziz et al. (2021) reported that *P. indica* supports chlorophyll metabolism by inducing biosynthesis genes (protochlorophyllide oxidoreductase [POR], chlorophyll a oxygenase [CAO]) and downregulating selected chlorophyll degradation genes (PPH, pheophorbide a oxygenase [PAO], and red chlorophyll catabolite reductase [RCCR]). These improvements in chlorophyll content and leaf greening indicates that *P. indica* colonization indirectly slows leaf senescence.

The growth enhancing capacity of *P. indica* has been demonstrated in many crop plants (Kundu et al. 2022). Previous reports suggest that *P. indica* enhanced the nutrient uptake in plants, especially nitrogen, phosphorous, sulphur, magnesium (Sherameti et al. 2005; Yadav et al. 2010; Kumar et al. 2011; Cruz et al. 2013; Bakshi et al. 2017; Narayan et al. 2021; Kushwaha et al. 2022). The broad impression of *P. indica* colonisation suggests numerous beneficial interactions with the host plants, implicating their role in general recognition and triggering of intracellular signalling pathways (Jogawat et al. 2020). Understanding the molecular mechanisms behind the growth enhancement would be useful for developing genetic markers which remains a research gap in cardamom. Here, we have observed the differential expression of four growth-related gene namely *AUX1*, *NIA1*, *VSP* and *PHT1* at the transcript level. It was previously reported that co-cultivation with *P. indica* resulted in a massive transfer of nitrogen from the agar plates to the plants' aerial parts and this action is linked to the activation of the NADH-dependent nitrate reductase (*NIA1*), a critical enzyme in the uptake of nitrate in plants (Sherameti et al. 2008a, b). In this study, we found out that the relative expression of the *NIA1* gene got upregulated by 3.45 times in cardamom plants colonized by *P. indica* than that of control plants. Numerous studies have reported the positive influence of *P. indica* on auxin transport. Sirenberg et al. (2007) demonstrated the production of IAA in a liquid culture of *P. indica*, which may induce growth promotion, as shown by other plant beneficial fungi such as *Trichoderma virens*, which increases biomass production in *Arabidopsis thaliana* through an auxin-dependent mechanism (Contreras-Cornejo et al. 2009). In this work, the expression of auxin (*AUX1*) was significantly upregulated by 3.26 times in *P. indica* colonized cardamom plants compared with the non-colonized controls which points out

the hormone dependent growth of cardamom plants on co-cultivation with *P. indica*. In root and shoot tissues, *AUX1* activity is necessary for both polar and phloem-based IAA transport routes (Leyser 2006).

In terms of plant growth and development, phosphorus (P), which makes up 0.5% of plant cell biomass, is the second most important limiting macronutrient after nitrogen (N) (Schachtman et al. 1998). Functional characterization of *P. indica* roots revealed a high affinity PiPT transporter (Pht1 family) gene that transports phosphate (Pi) from the fungus to the host plant. In P-deficient conditions, PiPT are activated, increasing the absorption of Pi against the gradient in concentration between inner plant tissue and soil solution. These results suggest that PiPT plays a crucial role in P transfer at the *P. indica*-plant root interface, which enhances overall plant development and yield (Muchhal et al. 1996). *P. indica* is now considered as one of the important contributors in defining alternate P absorption mechanisms by terrestrial plants. This symbiotic endophyte benefits plants by enhancing P absorption and translocation and facilitating the movement of other nutrients. Plants have numerous transporters, including the *PHT1*, *PHT2*, *PHT3*, and *PHT4*, which are involved in Pi uptake by the root and remobilization in plants for being adapted to low phosphate environments. Among the four major phosphate transporters, *PHT1* is responsible for phosphate uptake from soil by root cells, which is the first and most important stage in the Pi assimilation process (Liu et al. 2011). In this study, *E. cardamomum* on *P. indica* co-cultivation did not show significant up-regulation in *PHT1* transcript level expression when compared with the non-colonized controls. In fact, there are conflicting reports on the role of *P. indica* in phosphate transfer and host plant enhancement. *P. indica* increases phosphate uptake 2–3 times in *Arabidopsis* seedlings, according to Shahollari et al. (2005), implying that *P. indica* stimulates *Arabidopsis* growth in a similar way to mycorrhizal fungi. On the other hand, it has been found that *P. indica* does not cause a considerable increase in leaf P and N, and that phosphate has no function in improved growth of *Nicotiana attenuata* (Barazani et al. 2005).

As expected, the expression of *VSP* did not show significant up-regulation in transcript level under the influence of *P. indica*. Plants benefit from the accumulation of store protein because it helps them survive (Fujiwara et al. 2002). Lin et al. (2019) reported that the expression of JA-responsive genes like *VSP* (Vegetative storage protein gene) marked an increment after infection of the roots with *R. solanacearum* at 12 h, 1, 2 days, and this increase was stronger for the colonised *Anthurium* plants than that of uncolonized controls. Considering these findings, it can be speculated that the accumulation of VSPs take place in conjunction with some exogenous abiotic or biotic stress. Based on the coherent

results, it can be concluded that the growth enhancement in cardamom plants under the influence of *P. indica* is directly correlated with the elevated expressions of *NIA1* and *AUX1*, whereas an upregulated transcript levels of *PHT1* and *VSP* might be seen in the later stages of growth of the cardamom plant.

To the best of our knowledge, this work represents the first report on the co-cultivation of *in vitro* *E. cardamomum* with beneficial root endophyte *P. indica* and the expression of essential genes participating in the process of growth enhancement in plants. This could be further extended to find out whether *P. indica* promotes early flowering and thereby induce early fruiting. Earliness in flowering on *P. indica* inoculation was reported in barley (Achatz et al. 2010), *Coleus forskohlii* (Das et al. 2012), *Arabidopsis* (Kim et al. 2017; Pan et al. 2017), black pepper (Anith et al. 2018). Varma et al. (2014) reported that *P. indica* culture filtrate promotes plant growth and seed germination in *Helianthus annuus* and *Phaseolus vulgaris*. Considering this, cardamom plantlets primed with *P. indica* can be used in fields for faster cultivation, better performance and improved yield. Amendment of *P. indica* during the hardening of tissue cultured cardamom plants would result in better acclimatization and survival under normal climatic conditions. It is known that the growth of cardamom plant is altitude specific. Thus our study is extending to evaluate the performance of cardamom under the influence of *P. indica* on higher and lower altitudes. We have already initiated research on this line and hope to get early results. Early flowering and fruiting, if occur in field grown cardamom, will definitely be a breakthrough, as first flowering in normal cardamom plants occurs only after 2–3 years of planting.

5 Conclusion and future prospects

The study recommends the use of *P. indica* as a potential means for enhancing the growth and overall performance of cardamom and other crop plants. The study's findings also suggest that this symbiotic association could be used as a model for further research into the molecular and physiological pathways involved in symbiotic association and plant growth promotion. *P. indica* serves as a model organism for studying beneficial plant microbe interactions as well as a novel tool for improving plant production systems due to its simplicity of cultivation. Growth enhancement caused by *P. indica* colonisation minimises the need for fertiliser in the soil, lowering the risk of fertiliser overuse and contamination of the environment.

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Data Availability Data availability The datasets generated during and analyzed during the current study are available from the corresponding author on reasonable request.

Declarations

Conflicts of interest/competing interests The authors declare that they have no conflict of interest.

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