

Non rhizobial endophytic bacteria from chickpea (*Cicer arietinum* L.) tissues and their antagonistic traits

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Abstract

Bacteria that colonize plant tissues other than rhizobia and are beneficial for plant growth referred to non rhizobial plant growth-promoting endophytic bacteria (PGPEB). This study was designed to assay the biocontrol activity of plant growth promoting endophytic bacterial isolates those found positive for P. solubilization, ACC deaminase, Indole acetic acid and Gibberelic acid production. These bacterial isolates were obtained from chickpea (*Cicer arietinum* L.) tissues (roots and nodules). In a previous study a total of 263 non rhizobial endophytic bacterial isolates were isolated. Out of 263 isolates, 64.5% and 34.5% were Gram positive and negative, respectively. Further for biochemical characterization, catalase, oxidase, citrate utilization, nitrate reduction, methyl red and Voges Proskauer's tests, were performed. On the basis of P solubilization, ACC deaminase, Indole acetic acid and Gibberelic acid production 75 potential isolates were selected and screened for their biocontrol activity viz. (production of cell wall degrading enzymes, production of HCN and fluorescent pigment). Out of 75 isolates, only 29 isolates produced cellulase, 64 isolates were able to produce protease and 28 were positive for both cellulase and protease. Of 75 endophytic isolates 12 isolates (7 from root tissue and 5 from nodules tissue, respectively) were positive for HCN production and 16 isolates were found to be fluorescent pigment producer under μ v light. As chemical fertilizers and pesticides have detrimental effects on the environment. So these bacterial endophytic isolates will be used not only as a biofertilizer because of their plant growth promotional activities but also used as an alternative of synthetic chemicals for control of several plant diseases.

Keywords: Biocontrol, Cellulase, *Cicer arietinum* L., Non rhizobial endophytic bacteria, Protease

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INTRODUCTION

Bacterial endophytes colonize the interior tissue of plant exhibiting no apparent sign of infection or harmful impact (Kusari *et al* 2014). Approximately 3,00,000 existing plant species are in relation with endophytic inhabitants vary from a small to massive numbers (Dudeja and Giri 2014). Many promising endophytic bacteria like *Azoarcus sp.*, *Gluconoacetobacter diazotrophicus*, *Burkholderia sp.*, *Herbaspirillum sp.*, *Enterobacter* and genus *Serratia* are reported to reinforce yield in various agricultural crops (Vessey 2003). Plant growth promoting endophytic bacteria (PGPEB) are known to effect plant growth by decreasing plant disease and might be useful for sustainable agriculture as an alternative of chemical pesticides for improving the quality and yield of crop (Lugtenberg and Ka-

milova, 2009). Biological control has been described as eco friendly approach to reduce crop damage due to plant pathogens as compared to the use of chemical control of plant diseases (Wang *et al* 2013).

Endophytic microorganism can enhance plant establishment under stressful conditions in legume and non legume plants and preventing disease via antifungal and outcompeting pathogens for nutrients with siderophore production and better plant general resistance. Some bacterial endophytes show biocontrol activity (antibacterial and antifungal) by producing cell wall degrading enzymes or allelochemicals (antibiotics). Endophytic bacteria can get entry into the root tissues by two ways: actively, by production of hydrolytic enzymes (e.g. endoglucanase, exoglucanase and endopolygalacturonase) and these enzymes in-

involved in cell wall degradation of plant (Compant *et al* 2005) and passively, by penetrating the lateral roots emergence sites or junction of adjacent epidermal cells (Govindasamy *et al* 2008). Production level of these cell wall degrading enzymes differentiated between phytopathogens (deleteriously high levels) and root-colonizing bacterial endophytes (low levels) (Elbeltagy *et al* 2001) and contributed in endophytic bacteria entry into host and their spread inside plant tissue. Endophytic bacteria have the ability to protect their plant host from harmful microorganisms and pests by competition for space and nutrients and antagonism or by initiating the defence mechanisms of plant to respond immediately and efficiently against the pathogens. Antagonism against plant pathogens can be achieved directly by the production of fungal growth inhibitors, antibiotics and antibacterial secondary metabolites.

Most commonly strains of actinobacteria and bacteria viz. *Pseudomonas*, *Bacillus* and *Paenibacillus* spp. are reported as antagonistic for fungal pathogens and have been assayed for control of disease in a wide range of plants, e.g. wheat, potato and black pepper (Aravind *et al* 2009). The *Pseudomonas* spp. (fluorescent) form a long time for their biocontrol activity on plant soil borne pathogens for suppression of diseases. Bacteria with multiple biocontrol mechanisms antibiotics, chitinolytic enzymes, siderophores, HCN are being used widely (Saharan and Nehra 2011).

Associative, endophytic diazotrophic and non rhizobial endophytic bacteria were characterized from different plant species in last couple of years have raised their prospects to be used as biofertilizer (Akhtar and Siddiqui 2009). In recent decades, interest in endophytic microorganisms has been increased, as they have important role in sustainable agriculture. Knowing and understanding the negative impact of artificial fertilizers in agriculture, novel approaches such as the application of endophytic bacteria as biopesticides which are associated with plants, may help to improve plant health and increase productivity.

This study was designed *in vitro* screening of non rhizobial endophytic bacterial isolates for their biocontrol traits. These isolates were previously found positive for their plant growth promotional traits.

MATERIALS AND METHODS

Isolation of endophytic bacteria from chickpea: Healthy plants of chickpea (*Cicer arietinum* L.) were carefully taken out then washed with running tap water to wash off soil from undamaged tissue samples of root and nodules. Soaked in distilled water in a separate beaker and drained. Sample sterilization was done by using HgCl_2 (0.1%) for 30 seconds and ethanol (70%) for 3 min

for. After sterilization the tissue sample was washed thrice with sterilized water.

Aseptically surface sterilized tissues were homogenized and macerated. After maceration serial dilution of the tissue up to 10^{-6} was prepared and appropriate dilutions (100 μ l) were used and plates were incubated at 37°C. Further, the isolation of endophytic bacteria was done by streak plate method on Nutrient, Jensen's and *Pseudomonas* agar media. Sterility test was performed by placing the washed tissue on same medium and incubation was done at 28 \pm 2°C for 2-3 days. Carefully bacterial colonies were isolated and streaked over the plate containing their specific medium viz. Nutrient agar for *Bacillus* sp., *Klebsiella* sp., Jensen's agar for *Azotobacter* and *Pseudomonas* agar for *Pseudomonas* sp. Further, these isolates were maintained at 4°C on specific medium slants for future use.

Production of HCN: Nutrient agar medium supplemented with glycine (4.4g/litre) was streaked with exponentially grown selected endophytic bacterial isolates with simultaneously keep a filter paper soaked in picric acid (0.5%) in Na_2CO_3 (5%) in the upper lid of Petri dish. Incubation was done at 28 \pm 2°C for 2 to 3 days. Change in filter paper colour from yellow to light brown or strong (reddish-brown) represented as positive test for HCN production (Bakker and Schippers 1987).

Cell wall degrading enzyme production

Protease: Skimmed milk agar plates were prepared and spot inoculated with pure culture of test bacteria and incubated at 28 \pm 2°C for 2-5 days. Presence of halo clear zone around the growth indicated as positive test for protease production (Chaiarn *et al* 2008).

Cellulase: Cellulase activity of pure cultures was assayed by plating on Carboxy Methyl Cellulose (CMC) agar according to Ariffin *et al* (2006). Spot inoculation with test organism was done and plates were incubated at 28 \pm 2°C for 5 days. Appearance of halo zone around the bacterial growth was considered as positive test for cellulase production. Five days of incubation was done to allow the activity of cellulase on CMC agar plates at 28 \pm 2°C. After incubation, the agar plates were flooded with an aqueous solution of Congo red (1% w/v) for 15 minutes then solution was poured off, and further plates were flooding with 1M NaCl for 15 minutes. Appearance of a clear zone of hydrolysis confirmed cellulose degradation. Isolates with high cellulase activity was selected on the basis of clear zone diameter.

Production of Fluorescent pigment: Non rhizobial endophytic bacterial isolates were spot inoculated on King's B medium and plates were incubated at 28°C for 2 days. Observation of plates was done for yellowish green colour under μ v light. Fluorescence ability considered as positive for fluorescent pigment production at 400 nm.

Table 1. Biochemical characterization of non rhizobial endophytic bacteria of chickpea.

Characteristics Biochemical test	Endophytic bacterial isolates (%)	
	Positive	Negative
Gram's staining	64.5	34.5
Oxidase	87.4	12.6
Catalase	74.4	25.6
Citrate utilization	12.6	87.4
Methyl red (MR)	37.4	62.6
VogesProskauer (VP)	55.7	44.3
Nitrate reduction (NR)	36.6	63.4
Carbohydrate utilization Test	Different sources of sugar	Acid producers
	Dextrose	38.6
	Fructose	29.6
	Sucrose	48
	Sorbitol	18.7
		Gas producers
		6.7
		5.2
		13
		2.7

RESULTS AND DISCUSSION

Morphological characteristics of non-rhizobial endophytic bacteria and their biochemical characterization:

On the basis of morphological studies, out of 263 endophytic bacterial isolates 124 on nutrient agar medium produced large sized, irregular shaped, off-white and rough colonies, whereas 6 isolates showed rhizoid growth and were tentatively identified as *Bacillus* sp. Further, 71 isolates on *Pseudomonas* agar medium produced medium sized, round shaped and raised colonies with smooth margin and light yellow to off white in colour and were provisionally identified *Pseudomonas*. Further, few isolates also produced a fluorescent green pigment on King's B medium. 53 isolates were streaked on nutrient agar medium produced medium, round shaped and raised colonies having entire margin, mucoid and cream in colour were assigned as a *Klebsiella* sp., *Enterobacter* sp. and *Enterococcus* sp. Nine isolates produced yellow colour pigment on nutrient agar with circular, pinhead colonies and were convex with entire margins. On the basis of Gram's reaction, out of 263 non rhizobial endophytic bacterial isolates, 64.5% and 34.5% were Gram positive and negative, respectively. On the basis of biochemical characterization, out of 263 non rhizobial endophytic bacterial isolates 74.4%,

87.4%, 12.6%, 36.6%, 37.4% and 55.7% were found to be positive for catalase, oxidase, citrate utilization, nitrate reduction, methyl red and Voges Proskauer's tests, respectively (Table 1). In an another *in vitro* study, further these isolates were screened qualitatively and quantitatively for their plant growth promotional traits *viz.* P solubilisation, IAA and ACC deaminase production. On the basis of plant growth promotion trait only 75 potential isolates were selected for *in vitro* assay of biocontrol traits. Our results were in accordance with Saini *et al* (2015) who isolated 166 endophytic bacteria from root of legumes, chickpea (*Cicer arietinum*), pea (*Pisum sativum*), and lucerne (*Medicago sativa*) and non-legumes wheat (*Triticum aestivum*) and oat (*Avena sativa*) and from nodules of chickpea. Similarly, Zaghoul *et al* (2016) total of 167 endophytic bacterial isolates were isolated from roots, nodules, leaves and stems of faba bean (*Vicia faba*), pea (*Pisum sativum*), fenugreek (*Trigonella foenum-gracum*), lupine (*Lupinus spp.*), common bean (*Phaseolus vulgaris*) and rice (*Oryza sativa*) at flowering stage.

Biocontrol activity: Out of 75 non rhizobial endophytic bacterial isolates 38.70 %, 82.70%, 16.0% and 21.33% were cellulose, protease, HCN and fluorescent pigment producers, respectively (Fig 1).

Production of cell wall degrading enzymes: All the 75 non rhizobial endophytic bacterial isolates screened for cellulase and protease production (Table 2). Out of 75 isolates, only 29 (38.7%) were cellulose producers and 62 (82.7%) were protease producers. Maximum diameter of zone around bacterial colonies was observed for RBR 34, RBR40 and RBR139 (2.2 cm) on CMC and RBR 155 (2.8) on Skim milk agar media. Our results are well supported by Geetha *et al* (2014) who had also reported from mungbean rhizosphere out of 6 potential bacterial isolates, only 4 (WG-57, TG-60, BG-72 and KG-50) were able to produce cellulase whereas 3 showed protease activity. Similarly, Etesami *et al* (2015) procured

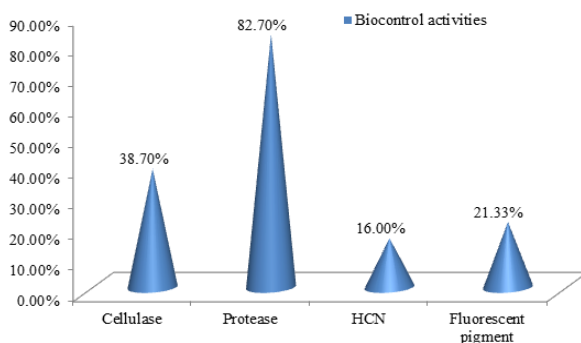


Fig. 1. Cellulase, protease, HCN and fluorescent pigment production by potential non rhizobial endophytic bacteria of chickpea.

Table 2. Cell wall degrading enzyme production by non rhizobial endophytic bacteria of chickpea.

Isolates	Cellulase Zone Dia (cm)	Protease Zone Dia (cm)
RBR11	-	2.2
RBR14	0.6	2.2
RBR17	2.1	2.4
RBR 19	1.1	2.1
RBR20	2.1	2.3
RBR25	-	1.8
RBR34	2.2	2.5
RBR38	-	2.6
RBR40	2.2	2.3
RBR49	0.7	0.9
RBR57	-	2.6
RBR61	-	2.5
RBR75	-	1.4
RBR80	-	2.2
RBR83	-	1.5
RBR89	-	2.5
RBR116	1.4	2.2
RBR119	1.1	2.3
RBR121	-	1.8
RBR127	-	2.2
RBR128	-	2.5
RBR136	1.6	1.6
RBR139	2.2	2.6
RBR144	-	1.9
RBR146	1.4	1.4
RBR155	1.8	2.8
RBR164	1.4	2
RBR165	-	2.5
RBR167	1.3	1.9
RBN2	-	2.6
RBN4	1.4	2
RBN16	-	2.1
RBN17	1.1	1.8
RBN20	-	2.1
RBN25	-	1.2
RBN27	1.3	1.7
RBN28	2.1	-
RBN30	-	2.3
RBN31	1.4	1.9
RBN32	-	2.4
RBN36	-	2.1
RBN38	-	2.3
RBN41	1.6	2.4
RBN44	-	1.7
RBN49	-	1.2
RBN54	-	2.5
RBN59	-	1.5
RBN61	1.1	1.4
RBN63	-	2.3
RBN64	-	2.3
RBN71	-	2.3
RBN75	1	2.1
RBN83	1.6	2.3
RBN86	1.3	1.6
RBN87	1.7	1.5
RBN88	-	2.2
RBN89	0.6	2.1
RBN91	1.4	2
RBN96	-	2.5
LCNE6	-	2.3
LCNE8	1.8	2
LCRE8	-	2
LCRE9	-	2.5
LGR 33	-	2
RB1	-	1.5

(-) No zone

Table 3. HCN and fluorescent pigment production by non rhizobial endophytic bacteria of chickpea.

PGP traits	Isolates
HCN production	RBR89, RBR112, RBR119, RBR127, RBR128, RBR136, RBR139, RBN20, RBN25, RBN30, RBN31, RBN54
Fluorescent pigment production	RBR11, RBR14, RBR17, RBR19, RBR34, RBR40, RBR75, RBR116, RBN17, RBN20, RBN25, RBN28, RBN54, RBN59, LCNE6, LCRE9

34 isolates of bacteria from two varieties of rice and all isolates were able to produce IAA, 27 isolates were positive for siderophore, 7 were producing HCN, 16 for ACC deaminase, 4 were chitinase, 22 for pectinase and 3 isolates were cellulose producers. Out of 9 isolates from clover plants all produced IAA whereas production of siderophore (6), HCN (2), ACC deaminase (4), chitinase (1), pectinase (6) and cellulose (1) and only 4 were phosphate solubilizers. Egamberdieva *et al* (2016) obtained (40) isolates of bacteria from chickpea root tissues. Isolates EB10 and EB2 were able to produced cellulose, lipase, protease, and chitinase.

HCN production: Data in Table 3 revealed bacterial endophytic isolates (7 and 5 from root and nodules, respectively) were positive for HCN production in chickpea. The present findings are well supported by Thirumal *et al* (2017), where out of 15 *Pseudomonas* isolates five viz., PVP1, DBP, DMuP, PGuP and RGP detected as moderate (++) whereas ten bacterial cultures (viz., DGP1, PSmP, PKP, PRP2, PSP2, DMP2, RGP2, MP1, MP2, SFP1) were scored as weak (+) for HCN production. Similarly, Ahmad *et al* (2008) reported of 72 bacterial isolates 42.85% *Bacillus* and 62.5% *Pseudomonas* spp. were able to produce HCN. Our results are well supported by Geetha *et al* (2014) where out of 6 potential bacterial isolates, only 2 were able to produce HCN. Sharma and Dubey (2017) observed that *Pseudomonas putida* CRN-09 isolated from the rhizosphere of *Vigna radiata* producing hydrogen cyanide (HCN) and also solubilizing phosphate.

Fluorescent pigment production: All the 75 endophytic bacterial isolates grown on King's B medium. Only 16 isolates were putative fluorescent pigment producer under μ v light (Table 3). Present investigation is well supported with Boiu-sicua *et al* (2017) where 20 endophytic bacteria from potato tubers only 6T2 isolate was positive for fluorescent pigment production on King's B agar medium. Similarly, Damodaran *et al* (2013) reported, of 16 rhizobacteria from saline sodic soil only G8, P1 and P2 were positive for fluorescent pigment production.

Conclusion

Two potential isolates RBN20 and RBN25 were found positive for protease, HCN, Fluorescence pigment production. Further, study will be planned to study the biocontrol activity of RBN20 and RBN25 *in vitro* and *in vivo*. These finding will be helpful to design the bacterial biofertilizer with both plant growth promotion and biocontrol activity and be used as an alternative of chemical based fertilizer and pesticides.

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