Identification and determination of characteristics of endophytes from rice plants and their role in biocontrol of bacterial blight caused by *Xanthomonas oryzae* pv. *oryzae*

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Summary Endophytic bacteria of rice plants (*Oryza sativa* L.) from eight different cultivars were screened for their ability in inducing disease symptoms, plant growth promotion and antagonistic activity against *Xanthomonas oryzae* pv. *oryzae*. Out of the 63 whole isolates, five were plant pathogens. Based on phenotypic characteristics and 16S rDNA sequence analysis, these were identified as *Pseudomonas oryzihabitans*, *P. fulva*, *Pantoea ananatis*, *Pantoea* sp., *Cellulomonas* sp. Four out of the 63 isolates behaved as potentially good plant growth-promoting and biocontrol agents. These were identified as *Bacillus* sp., *B. subtilis*, *Pseudomonas putida* and *Enterobacter* sp. This is the first report of pathogenic and endophytic bacteria from rice grown in field conditions in North of Iran.

Additional keywords: endophytic bacteria, Oryza sativa L., plant pathogens, PGPR, 16S rRNA gene

Introduction

Beneficial endophytic microorganisms are primarily comprised of fungi and bacteria that colonize internal plant tissues without causing visible damage to their hosts. (Schulz and Boyle, 2006; Rodriguez *et al.*, 2008; Rodriguez *et al.*, 2009; Reinhold-Hurek and Hurek, 2011). These local and systemic colonizers of the internal tissue can be isolated from sterile surface or internal parts of plants (Hallmann *et al.*, 1997) from both monocotyledonous and dicotyledonous plants (Ryan *et al.*, 2008).

Endophytes enter the plant through the natural vents and wounds, or by secretion of hydrolytic enzymes (Hallmann *et al.*, 1997). They subsequently establish closer interaction with the host by increasing access to nutrients such as nitrogen, phosphorus and iron, or by inducing plant defense mechanisms (production of anti-pathogen agents,

suppression of pathogens through competition for the colonization of places and food) and can potentially positively affect plant health and protection (Kang *et al.*, 2007; Reinhold-Hurek and Hurek, 2011; Santoyo *et al.*, 2016; Moronta-Barrios *et al.*, 2017).

Several reports have highlighted the benefits of using endophytic bacteria as biocontrol agents against fungal and bacterial rice pathogens. Burkholderia sp. KJ006, for example, was used against Burkholderia glumae (Cho et al., 2007) and Rhizobium lequminosarum bv. phaseoli against Rhizoctonia solani (Mishra et al., 2006). Similarly, other endophytic bacteria have been used to protect rice cultivars from major fungal infections including Penibacillus spp., Microbacterium spp., Bacillus spp. and Klebsiella spp. against Fusarium oxysporum and Rhizoctonia solani (Ji et al., 2014). Recently certain potential biological control agents such as Bacillus amyloliquefaciens, B. methylotrophicus and B. subtilis were isolated from rice plants showing the highest antimicrobial activities against the two major rice pathogens, Rhizoctonia solani and Burkholderia glumae, which cause sheath blight and bacterial panicle blight, respectively (Shrestha et al., 2016). For these and other reasons, nowadays endophytes are increasingly receiving attention as potential agents in the biologi-

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cal control of plant pathogens as well as for the improvement of plant growth.

Rice is the staple food of more than half of the world's population. It is considered the oldest and the most important crop throughout the world, especially in Asia. Considerable agricultural areas are under cultivation of rice in the world.

Bacterial blight caused by *Xanthomonas* oryzae pv. oryzae (Xoo) is a devastating pathogen in rice, especially in Asian countries. It reduces rice production by 20% in natural conditions and by 70% in epidemic situations, being one of the factors that limit the global production of rice (Ou,1985). For the control of the pathogen, plant disease resistance (R) genes are used to generate new disease-resistant cultivars (Peng *et al.*, 2015) and isolation of potent endophytes as new sources for biological control of plant pathogens is under investigation.

The aim of the present study was to isolate and identify certain endophytic bacteria from commonly used rice cultivars in northern Iran, in order to evaluate their effect as biocontrol agents against rice bacterial blight caused by *Xoo*.

Materials and methods

Isolation of endophytic bacteria from rice plants

Samples were collected from eight rice cultivars cultivated in regions under rice plantation in the north of Iran. They were washed with tap water to separate soil particles from the root surface. Disinfection followed by using 2% hypochlorite sodium for 10 minutes and 70% alcohol for 1 minute. The plant samples were then rinsed four times with 0.02 M phosphate potassium buffer (pH=7). To ensure the effectiveness of disinfectants, a 0.1 ml buffer from the final wash was added to 9.9 ml Tryptic soy agar (TSA) as control. Any growth in the control plates within 48 hours, these were eliminated. The root, stem and leaf samples were then cut into small pieces via a sterile blade. One gram of sample tissue with 9.9 ml phosphate buffer was squashed in a sterile mortar. The solution was diluted serially in the buffer and cultured on a TSA medium in a Petri dish. Single colonies were selected and purified on fresh TSA plates (McInroy and Kloepper, 1995).

Preliminary diagnosis of bacterial pathogenic isolates

To determine the pathogenic properties of 63 isolates, appropriate suspensions of yellow colonies and fluorescent bacteria were prepared and the inocula were infiltrated into tobacco leaves to fulfill HR development. To confirm pathogenicity tests on rice plants cv. Hashemi, two pathogenicity tests were conducted simultaneously (Zou et al., 2006; Shrestha et al., 2016; Jabeen et al., 2011).

In the first experiment, the leaves were cut off and surface-sterilized with sodium hypochlorite solution for 1 minute and then washed with ddH₂O. The sterilized leaves were then placed on a three-layer paper towel in Petri dishes and inoculated with suspensions of representative isolates with 10⁸ cfu/ml concentration using the pin prick method. All treated leaves including water treated control leaves were inoculated at 25°C. The test was conducted in three replications for each treatment.

In the second experiment, one-monthold rice plants were inoculated with bacteria at a concentration of 10⁸ cfu/ml from fresh cultures by the leaf-clipping method (Kauffman *et al.*, 1973). The treated plants were transferred to the greenhouse at 25-30°C with a 12- hr exposure period. They were observed every day within one month for any disease symptoms. The control plants were inoculated with sterile phosphate buffer. Experiments were performed in three replicates for each treatment.

Phenotypic characteristics of the pathogenic isolates

Pathogenic isolates were studied by a series of key phenotypic tests, including physiological and biochemical tests, in terms of the determination of genus and species

(Schaad et al., 2001).

In vitro assay for antagonistic properties of isolates

To carry out the tests for antagonistic properties of isolates, *Xoo*-strain (MT) was used. From a 24-48 hour culture of the strain on a TSA medium, a suspension was prepared in sterile distilled water and its optical density (OD₆₀₀) was adjusted 1. Isolates with colonies other than yellow colours, all positive fluorescent bacteria and saprophytes with no HR reaction, were spot-inocuated on TSA medium and incubated for three days at 25°C. Then, *Xoo* suspension at 108 cfu/ml concentration was sprayed uniformly on TSA agar medium. The plates were incubated at 25°C, and inhibition zones were measured. This was done in triplicates.

Germination and vigour index of inoculated rice seeds with endophytic bacteria

The test was conducted on the basis of the standard method of roll towel (ISTA, 1993). At first, 20 surface sterilized seeds were treated in 3×10⁸ cfu/ml concentration of each isolate and each batch was placed separately between two plies of a wet tissue and the seeds were a slightly compressed. The tissues containing the treated seeds were then placed separately in plastic bags which were rolled and kept in a growth chamber for 14 days. Three replications were considered for each treatment. The percentage of seed germination, and the mean root and shoot length of each seedling were indices evaluated for the growth-inducing activity of isolates. To calculate the vigour index of the seedling, the following method was used (Baki and Anderson, 1973):

Vigour index = % germination \times seedling length

Seedling length = (shoot length + root length)

Evaluation of plant growth promotion of endophytic bacteria on rice in greenhouse conditions

An assessment of the growth improve-

ment activity of 17 selected isolates from the Vigour index test was done according to Chithrashree *et al.* (2011). Seeds treated with fresh suspensions of antagonistic isolates (3×10⁸ cfu/ml) and untreated control seeds were separately planted in pots filled with sterile soil and sand in equal volumes. The pots were irrigated daily and each pot received 25 ml of Hoagland 1/3 (V/V) solution once a week. Within 30 days after planting, the plant's growth promotion activities such as plant height and fresh weight of seedlings were measured. Each treatment consisted of four replicates (Chithrashree *et al.*, 2011).

Evaluation of the effect of endophytic-antagonistic bacteria against *Xoo* in greenhouse conditions

The rice seeds were kept in sterile distilled water for 48 hours after surface disinfection with 2% hypochlorite sodium for 30 seconds. Then, the extra water of the seeds was removed and air-dried in the shade for half an hour. Six seeds were planted in vases containing sterile soil. Each cellular suspension of antagonistic bacteria was used separately in greenhouse during plantation. The one-month plants with suspension of Xoo (108 cells/ml) were inoculated using the leaf clipping method (Kauffman et al., 1973). The test was done in the form of a complete random design in three replications. The control plants were inoculated with sterile distilled water. The length of lesions on leaves of each plant was measured separately after 45 days of inoculation with the pathogen, based on a five-point scale: 0=no sign of diseases and lesion length less than 0.2 cm, 1=lesion length between 0.2 and 1.5 cm, 3=lesion length between 1.5 and 3 cm, 5=between 3 and 5 cm, 7=between 5 and 10 cm, 9=above 10 cm. The percentage disease index was calculated by the following formula (Fang et al., 1999):

Disease index (%) = $\{(1 \times N_1 + 3 \times N_3 + 5 \times N_5 + 7 \times N_7 + 9 \times N_9) / 9Nt\} \times 100$

where N_{1-9} is the number of the leaf index and Nt is the total number of tested leaves.

In this test, growth factors such as fresh weight of root and shoots were measured.

Bacterial identification using 16S rRNA gene sequencing

DNA Extraction

To extract DNA of isolates, the boiling method was used. Bacterial cultures were obtained in 5 ml Nutrient broth (Nb) medium. After keeping it in 37°C incubator for 20 hours, 1ml from each culture was transfered into sterile Eppendorf tube, and centrifuged in 13000 rpm for 5 minutes. The supernatants were discharged and pellets were resuspended by vortexing with 300 µl lysis buffer containing 0.1 N NaOH and 0.5% SDS (Elboutahiri et al., 2009). The tubes were placed in a boiling bain-marie for 10-13 minutes and immediately were placed on the ice for one minute. The tubes were centrifuged in 13000 rpm for 5 minutes and the supernatants contain suspended DNA was transferred into new tubes for PCR reaction.

PCR amplification

The amplification of 16S rDNA was carried out in a reaction with a final volume of 25 μl containing 2.5 μl of 10x PCR buffer, 1 μl of $MgCl_2$, 0.5 μ l of dNTP (2.5 mM) (Bio-Rad), 1 μl (100 ng/μl) of the P1 primer (5'-CGGGATC-CAGAGTTTGATCCTGGTCAGAACGAACGCT -3'), 1 μl (100 ng/μl) of the P6 primer (5'-CG-GGATCCTACGGCTACCTTGTTACGACTTCAC-CCC -3') (Palacio-Bielsa et al. 2009), 0.3 µl of Taq DNA polymerase, 2 µl of total DNA and 16.7 μl of sterile distilled H₂O. The PCR reaction conditions were as follows: 94°C for 2 min, followed by 25 cycles of denaturation at 94°C for 1 min, annealing at 52°C for 1 min and extension at 72°C for 3 min, before a final elongation at 72°C for 20 min.

Statistical analysis

The results of the experiments were analyzed independently. The treatment means were compared by Duncan's multiple range test. The statistical software SAS version 92.1 was used for the statistical analyses.

Results

Isolation, identification and characterization of endophytic bacteria

Sixty three (63) culturable bacterial endophytes were isolated from internal tissues of the rice plants collected from different parts of North of Iran. Yellow and fluorescent colonies HR positive were tested for HR and pathogenicity tests on rice plants (Table 1).

Pathogenicity test

Of the 63 endophytic bacterial isolates, 6 isolates had positive HR reaction on tobacco leaves. Five isolates namely, OS6, OS14, OS18, OS22 and OS36, caused disease symptoms on inoculated rice plants. Some of these were considered as pathogens with low disease potential. All pathogenic strains in the detached leaf assay showed a typical elongated water soaked lesion which developed into a brownish black necrosis. These were recorded 48-72 h post-inoculation (hpi). In the pot experiment all plants inoculated with pathogenic endophyte strains showed progressive water-soaking lesions in the inoculated point after 2-4 days. The lesions became necrotic and produced a necrotic stripe on the leaf that spread to the full length of the leaf. The isolates with low disease potential produced confined dark brown spots on the inoculated region. Koch's postulates were fulfilled for all pathogenic strains.

Phenotypic properties of the pathogenic isolates

The tested endophytes isolates were divided into two groups; yellow colonies up to fluorescent pigmentation on king's B medium and the non-fluorescent, and the white colonies group. As presented in Table 2, five strains were tentatively identified based on morphological and various biochemical characteristics. Two of the strains namely, OS6 and OS36, were designated as *Pseudomonas* spp. Two other strains (OS14 and OS18) were identified as *Pantoea* spp., and the strain OS22 could not be assigned

Table 1. Overall characteristics of endophytic bacteria isolated from rice cultivars.

Cultivar	Tissue origin	Pathogenicity in rice	Strain number	Colony color on TSA
Hashemi	10 ⁻³ Leaf	-+	OS4, OS18	yellow
	10⁻⁴ Leaf	++-	OS6, OS14, OS46	yellow
	10⁻⁵ Leaf		OS15, OS16	yellow
	10 ⁻³ Root	-	OS2, OS19	yellow
Tarom amrollahi	10⁻⁵ Leaf	-	OS7	colorless
	10 ⁻⁶ Leaf	-	OS17	yellow
Kouhsar	10 ⁻² Leaf	+	OS22	yellow
	10 ⁻⁴ Leaf	-	OS38	yellow
	10 ⁻² Root	-	OS20	colorless
Binam	10 ⁻³ Leaf	+	OS36	yellow
	10⁴Leaf	-	OS25	yellow
	10 ⁻² Root	-	OS28	white
	10⁻⁴ Root	-	OS49	creamy
Amol	10 ⁻² Stem	-	OS31	colorless
	10 ⁻² Root	-	OS29	yellow
	10 ⁻³ Root	-	OS30	yellow
Fajr	10⁻³ Root		OS32, OS34	yellow
	10⁻⁴ Leaf	-	OS33	yellow
Khazar	10 ⁻³ Root	-	OS42	yellow
	10 ⁻³ Root	-	OS43	brown
Tarom hashemi	10 ⁻³ Root	-	OS53	colorless
	10 ⁻³ Leaf	-	OS60	yellow
Tarom mahalli	10⁻⁴Root		OS61, OS62	yellow
	10⁻⁵ Root	-	OS63	yellow

Table 2. The phenotypic characteristics of five main endophytic bacteria with low disease potential isolated from rice plants.

Test	OS6	OS14	OS18	OS22	OS36
Yellow pigment	+	+	+	+	+
Gram staining	-	_	_	+	-
Oxidase test	-	-	-	+	-
Fermentative	No	Yes	Yes	No	No
Fluorescence on King's B medium	-	-	-	-	-
Starch hydrolysis	-	-	_	-	_
Catalase hydrolysis	+	+	+	+	+
Litmus milk	ALK	ND	ND	-	ALK
Levan from sucrose	+	-	_	-	+
H₂S from cysteine	-	-	_	-	_
Xanthomondin	-	-	-	-	-
Mucoid growth on YDC	-	+	+	+	-
Growth on 0.1% TTC	+	+	+	+	+
Growth at 40°C	-	+	+		-
HR on Tobacco	+	+	+	+	+
Pathogenicity on rice	+	+	+	+	+

⁺⁼ positive; = negative; O= oxidative; v= variable; ALK= alkaline; NC= no change; ND= not determined.

to a known species (Table 2).

In vitro assay for antagonistic activities of isolates to *X. oryzae* pv. *oryzae*

In the initial screening of the antagonistic activity of 39 bacterial isolates against *Xoo* strain, 21 bacterial isolates exhibited the most potent antagonistic activity against the *Xoo*. The maximum inhibitory activity was recorded for OS59 with an inhibition zone of 32.67 mm, whereas the isolates OS58, OS43, OS20 and OS40 could produce inhibition zones of 32.33 mm, 26.65 mm, 18 mm and 16.7mm, respectively (Fig.1). The selected isolates were consequently used for further experimentation.

Germination and vigour index of inoculated rice seeds with endophytic bacteria

In total, 17 bacterial strains selected from the in vitro assays, namely, OS3, OS10, OS12, OS20, OS21, OS23, OS28, OS31, OS40, OS43, OS44, OS49, OS52, OS53, OS55, OS58 and OS59, were tested. The treatment of seeds with some of these strains significantly increased the seed germination and seedling vigour in comparison to the untreated control. Of these, 6 isolates namely, OS40, OS23, OS43, OS52, OS31 and OS53, had statistically significant effects compared to the control (Table 3). Further tests were carried out to evaluate their potential in promoting plant growth.

Effect of endophytic bacteria on growth promotion of rice in greenhouse conditions

Under greenhouse conditions, of the 17 isolates tested, 41.176% of the isolates caused a significant difference in growth parameters compared to the control plants. Seed treatments with OS3, OS23, OS31, OS40, OS53, OS58 and OS59 isolates increased seedling height and fresh weight in comparison with the control. The highest heights of 23.7, 22.81, 21.9 cm were obtained from seeds treated with the OS58, OS40 and OS31 isolates. The application of the OS53, OS31, and OS59 isolates also resulted in the highest seedling weights of 0.893, 0.806

and 0.6662 grams (Fig. 2).

Evaluation of the effect of endophytic-antagonistic bacteria against *Xoo* in greenhouse condition

Seventeen (17) representative antagonistic bacteria with potential plant growth promotion activities were assessed for the control of bacterial blight disease 60 days after the treatment of rice seeds with endophytic bacteria. As shown in Fig. 3, the plants with bacterial inoculants of OS52, OS40, OS23 and OS53, reduced the incidence of the disease by 28.5%, 33.25%, 34.25% and 37.25%, respectively (Fig. 3). Also in presence of the pathogen Xoo, root and shoot fresh weight of the rice plants inoculated with the mentioned isolates showed significantly increased values of these parameters compared to the control treatment (Fig. 4). This indicates their function as biological control agents of Xoo causing bacterial blight in rice.

PCR test for identification of representative bacteria

To identify the four most efficient antagonistic endophytes (OS52, OS40, OS23 and OS53), the extracted DNA of these isolates were amplified using universal 16S rDNA primers (P1 and P6) and a PCR product size of 1500 bp was obtained. The sequencing samples were sent to Macrogene Inc. (Seoul, South Korea). The sequence results were compared with those available in the NCBI GenBank. It was shown that the isolates OS23, OS40, OS52 and OS53 with 97%, 90%, 95% and 98% identities belonged to Bacillus sp., B. subtilis, Enterobacter sp. and Pseudomonas putida, respectively (Fig. 5). However, 2 species were indistinguishable by 16S rRNA gene sequences.

Discussion

Eight cultivars of rice were screened for beneficial endophytic bacteria in the provinces of Mazandaran and Gilan in northern Iran during two seasons in the year 2016. Sixty three endophytes were isolated from differ-

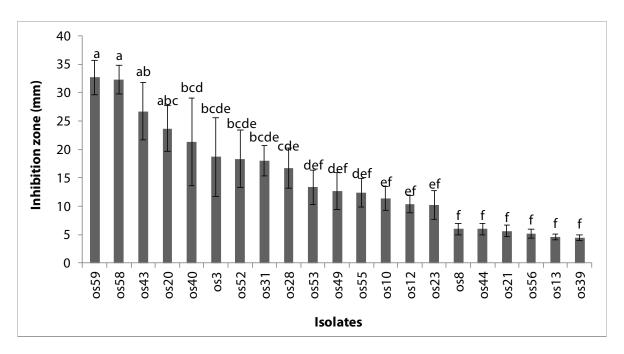


Figure 1. Comparative antagonistic potential of 21 endophytic bacteria against *Xanthomonas oryzae* pv. *oryzae* in rice. *Data are mean of three replications; in a column, means followed by a common letter are not significantly different at 5% level by Duncan's Multiple Range Test.

Table 3. Plant growth promoting activity of endophytic bacterial isolates in rice plants.

Isolates	MRL(cm)	MSL(cm)	Germination (%)	VI*
OS3	2.72 ± 0.60 ^{bcd}	5.57 ± 0.51 ^{ef}	100	8.16 ± 0.44 ^{fg}
OS10	2.59 ± 0.13 ^{cde}	4.69 ± 0.45 ⁹	96	7.12 ± 1.00^{h}
OS12	2.40 ± 0.26^{de}	5.78 ± 0.62 ^{cdef}	98	8.01 ± 0.77^{fgh}
OS20	3.12 ± 0.21 ^b	5.41 ± 0.25 ^f	100	7.55 ± 0.22 ^{gh}
OS21	2.17 ± 0.40^{de}	4.00 ± 0.27 ^{gh}	100	4.90 ± 0.36^{i}
OS23	4.19 ± 0.32 ^a	6.45 ± 0.47 ^{abcd}	100	11.01 ± 0.77^{ab}
OS28	2.4067 ± 0.1^{de}	3.37 ± 0.21 ^h	94	5.43 ± 0.21^{i}
OS31	2.52 ± 0.12^{de}	6.18 ± 0.13 ^{abcde}	100	9.64 ± 0.32^{bcd}
OS40	4.31 ± 0.25 ^a	6.51 ± 0.58 ^{abc}	100	11.28 ± 0.56^{a}
OS43	4.01 ± 0.10^{a}	6.73 ± 0.39^{a}	100	10.51 ± 0.38^{ab}
OS44	2.03 ± 0.17^{e}	3.63 ± 0.27 ^h	96	5.43 ± 0.44^{i}
OS49	2.56 ± 0.16^{cde}	6.13 ± 0.25 ^{abcdef}	100	8.66 ± 0.11^{def}
OS52	4.03 ± 0.25^{a}	5.95 ± 0.58 ^{bcdef}	100	10.03 ± 0.18 ^{bc}
OS53	3.15 ± 0.37 ^b	6.64 ± 0.41ab	100	9.45 ± 0.66 ^{cd}
OS55	3.07 ± 0.26 ^{bc}	6.57 ± 0.53^{ab}	100	9.20 ± 0.27^{cde}
OS58	3.82 ± 0.26^{a}	5.70 ± 0.40^{def}	100	7.78 ± 0.42^{fgh}
OS59	2.55 ± 0.40^{cde}	5.40 ± 0.35 ^f	100	8.36 ± 0.94^{efg}
Со	2.02 ± 0.33^{e}	4.00 ± 0.21gh	100	5.65 ± 0.64^{i}

MRL= Mean root length, MSL= Mean shoot length, VI= Vigour Index .

^{*} Data are mean of three replications; in a column, means followed by a common letter are not significantly different at 5% level by Duncan's Multiple Range Test.

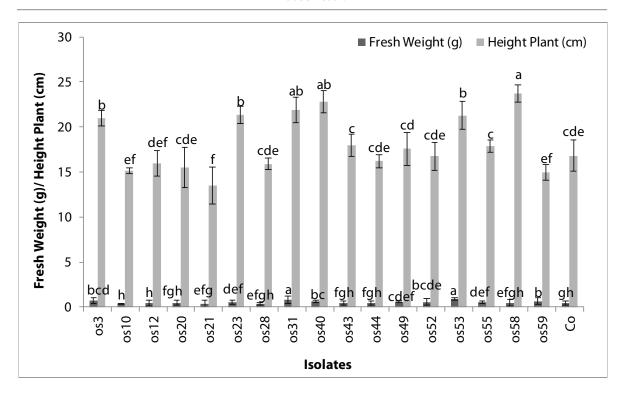


Figure 2. Effect of endophytic bacteria on growth promotion of rice seedlings 30 days after application in greenhouse conditions.

*Data are mean of four replications; in a column, means followed by a common letter are not significantly different at 5% level by Duncan's Multiple Range Test.

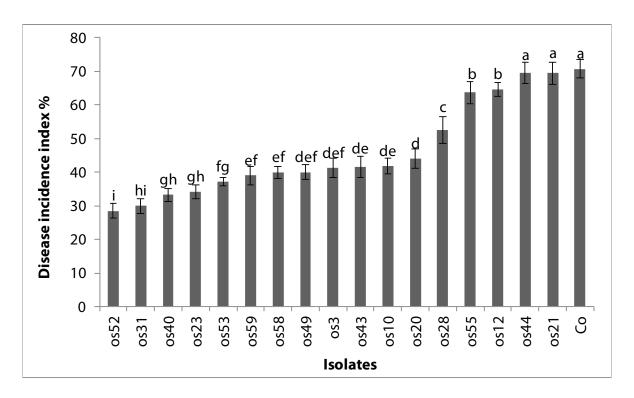


Figure 3. Control of bacterial blight of rice by endophytic bacteria under greenhouse conditions 60 days after treatment. *Data are mean of four replications; in a column, means followed by a common letter are not significantly different at 5% level by Duncan's Multiple Range Test.

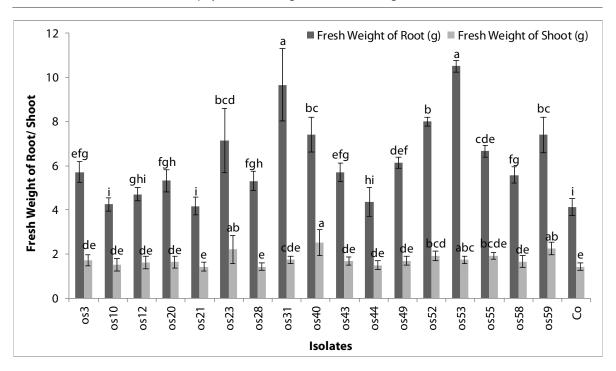


Figure 4. Different plant growth promoting activity of the representative endophytic bacteria on rice in greenhouse conditions and in presence of the pathogen *Xanthomonas oryzae* pv. *oryzae* in rice.

*Data are mean of four replications; in a column, means followed by a common letter are not significantly different at 5% level by Duncan's Multiple Range Test.

ent parts of the plants including the leaves, stem and roots.

Based on pathogenicity tests, five of the isolates were considered as potential pathogens of rice, which after 16s rDNA sequencing were identified as *Pseudomonas oryzihabitans* (99%), *Pantoea ananatis* (96%), *Pantoea* sp. (96%), *Cellulomonas* sp. (97%) and *Pseudomonas fulva* (97%). It is worth mentioning that neither of these bacteria have been previously reported as rice pathogens in Iran. The remaining isolates were considered non-pathogenic isolates.

Of the pathogenic isolates found on rice, some have already been reported on rice plants. *Pantoea ananatis* from Cambodia, and *Pantoea* spp. and *Pantoea stewartii* from the Philippines have been identified as pathogens with different pathogenic potential (Cother et al., 2010; Mano et al., 2006; Mano et al., 2007; Okunishi et al., 2005; Cottyn et al., 2001; Cottyn et al., 2009). *Pseudomonas oryzihabitans, Pseudomonas* spp. and *Cellulomonas flavigena* have been reported as seed saprophytes in irrigated areas with low pathogen-

ic potential trait from the Philippines (Cottyn et al., 2001; Cottyn et al., 2009). Moreover, *Pseudomonas fulva* has been identified as a pathogen in pepper from China and as a diazotrophic endophyte in rice seeds (Qiang et al., 2017; Verma et al., 2001).

According to Cottyn et al. (2009), it can be assumed that under certain conditions of rice cultivation, these endophytes can eventually appear as pathogens. Generally, some endophytes may turn into pathogens depending on several factors such as: the host and endophyte growth stage, plant defensive responses, environmental changes such as CO₂ accumulation or O₂ depletion, the production of specific metabolites in another host, the presence of other microorganisms interacting with it (Lund and Wyatt, 1972; van Peer et al., 1990; Sturz et al., 1997; Schulz and Boyle, 2005; Rosenblueth and Martínez-Romero, 2006). Nevertheless, in artificial inoculation, factors such as inoculation procedure and high inoculum concentrations may increase pathogenicity (Cottyn et al., 2009).

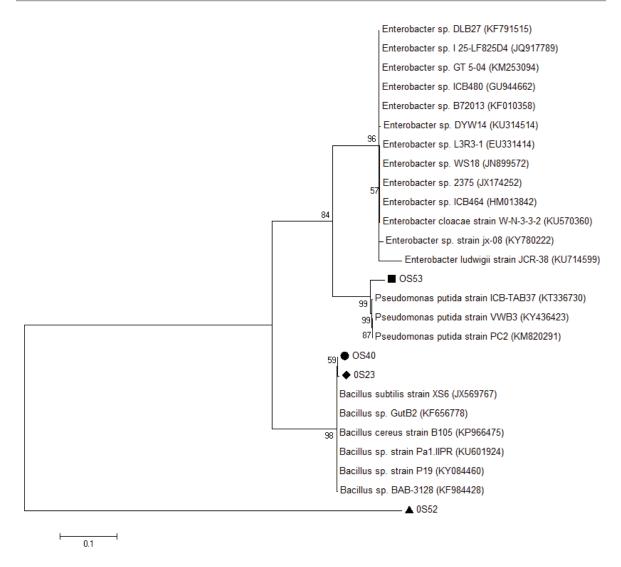


Figure 5. Phylogenetic tree based on the 16S rRNA gene of 4 beneficial endophytic bacteria and related bacterial species. The evolutionary history was inferred by using the Maximum Likelihood method based on the Kimura 2-parameter model. The numbers at the nodes are boot strap values based on 1000 replications. The analysis involved 26 nucleotide sequences. Evolutionary analyses were conducted in MEG 7 (Tamura *et al.*, 2013; Kumar *et al.*, 2016).

To select the best antagonists, screening of isolates was based on their *in vitro* antagonistic activity to *Xoo*, their effect on seed germination rate, seedling growth promotion and the reduction of disease intensity in greenhouse conditions. Four isolates, OS52, OS40, OS23 and OS53, which were identified by nucleotide sequence analysis as *Enterobacter* sp., *B. subtilis, Bacillus* sp. and *Pseudomonas putida*, respectively, could increase plant growth and decrease *Xoo* infection under greenhouse conditions. To the best of our knowledge, this is the first report of isolation of endophytes from rice in Iran. Seed

treatment with these antagonists caused disease to cease in over 60% of plants. These species have been found as endophytes in rice among many other species including Coryneform spp., Pantoea spp., Burkholderia glumae, Xanthomonas spp., Staphylococcus spp., Paenibacillus polymyxa, Methylobacterium spp., Curtobacterium spp., Azospirillum amazonense, Caulobacter crescentus, Kocuria palustris, Micrococcus luteus, Microbacterium spp., Klebsiella spp., Azospirillum spp., Herbaspirillum spp., Rhizobium spp., Sphingomonas spp. and Brevibacillus spp. (Elbeltagy et al., 2000; Okunishi et al., 2005; Mano et

al., 2007; Mano and Morisaki, 2008; Cottyn *et al.*, 2009; Kaga *et al.*, 2009; Ji *et al.*, 2014; Nhu and Diep, 2014).

Several endophytic bacteria such as Enterobacter sp. CNPSo 2480, Bacillus sp. CNPSo 2481, Pseudomonas rhodesiae, Pantoea ananatis, Pseudomonas sp. PsJN, H. rubrisubalbicans, G. diazotrophicus, A.amazonense and Burkholderia sp. induce growth and increase in plants like corn, pepper, grape and sugarcane (Barka et al., 2002; Oliveira et al., 2003; Kang et al., 2007; Szilagyi-Zecchin et al., 2014). In addition, endophytic bacteria have been reported to be involved with other mechanisms such as competition for place and food, and antibiosis (Sturz et al., 2000; Reiter et al., 2002; Compant et al., 2005; Aravind et al., 2009; Ji et al., 2013). Especially in rice, Ji et al. (2013) found that treatment of rice seeds with the diazotrophic endophytes Paenibacillus, Microbacterium, Bacillus and Klebsiella, induced systemic resistance against Fusarium oxysporum and Rhizoctonia solani in addition to the growth induction of the rice plant.

In the present study, B. subtilis seemed to work well against *Xoo* as well as the other Bacillus spp., which need to be investigated further. Recent studies by Chung et al. (2015) and Hossain et al. (2016) showed that Bacillus oryzicolaYC7007, a new endophytic bacterium isolated from rice roots, improves plant growth and controls bacterial blight, cluster blight and bakanae disease via the production of antibiotic and induction of systemic resistance. The application of an endophytic strain Bacillus spp. of tomato and potato with Silwet polysilicon surfactant caused colonization of cocoa leaves (for more than 68 days) and reduction of black pod rot disease via the induction of systemic resistance (Melnick et al., 2008).

The Enterobacter sp. isolate also has a potential beneficial effect on the control of bacterial blight. Of Enterobacter spp., Enterobacter radicincitans in cereals (Witzel et al., 2012), Enterobacter sp. SP1 in sugar cane (Zhu et al., 2012) and Enterobacter cloacae MSR1 in alfalfa (Khalifa et al., 2016) have shown an ability to increase production in plants. The

production of plant hormones by the Enterobacter sp. 638 isolated from the stem of a type of poplar hybrid has led to a 40% increase in poplar growth (Taghavi et al., 2010). Enterobacter cloacae and Enterobacter aerogenes strains have been reported to control Fusarium oxysporum f. sp. spinaciae in spinach (Tsuda et al., 2001) and Setosphaeria turcica, the Northern corn leaf blight fungus, in maize (D'Alessandro et al., 2014). Biological control action of E. cloacae has also been reported by Roberts et al. (1994). Enterobacter isolates increased nodule occupancy of bradyrhizobial strains at nitrogen fixation on legumes (Gulpa et al., 1998).

The increase in the growth of rice plants by the activity of *P. putida* coincides with previous findings on endophytic *Pseudomonas* spp. e.g. nitrogen fixation by *Pseudomonas* stuzeri A1501 endophyte promoted plant growth (Yan et al., 2008). Sheoran et al. (2015) proved that the endophytic *P. putida* BP25, isolated from the black pepper root endosphere, could by internal colonization in Ginger and Arabidopsis plants induce production of volatile compounds that inhibit fungal pathogens and the plant parasitic nematode *Radopholus similis* (Sheoran et al., 2015).

Despite the interesting profile of *Enterobacter* sp. and *P. putida* as antagonists of *Xoo* on rice, these bacteria are potentially dangerous and harmful to human health (Brenner *et al.*, 1986; Bouallegue *et al.*, 2004; Berg *et al.*, 2005; Flores-Tena *et al.*, 2007; Molina *et al.*, 2011; Thomas *et al.*, 2013; Molina *et al.*, 2014). Hence the hazardous properties of the bacteria and human safety issues should also be taken into account when considering their practical use as biological control agents in plant protection.

In conclusion, this is the first record of the pathogens *P. oryzihabitans*, *P. ananatis*, *Pantoea* sp., *Cellulomonas* sp. and *P. fulva* and the endophytic bacteria/ antagonists *Enterobacter* sp., *B. subtilis*, *Bacillus* sp. and *P. putida* against *Xoo* in rice plants from the northern parts of Iran, where rice is mostly cultivated. The beneficial bacteria showed promising efficacy and plant growth pro-

motion activities against *Xoo* in rice both *in vitro* and in greenhouse conditions. Further research is required in the perspective of application of the present findings to the field.

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Ταυτοποίηση και προσδιορισμός χαρακτηριστικών ενδοφύτων σε φυτά ρυζιού και ο ρόλος τους στη βιολογική αντιμετώπιση του βακτηρίου *Xanthomonas oryzae* pv. *oryzae*

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Περίληψη Εξετάστηκαν απομονώσεις ενδοφυτικών βακτηρίων από φυτά ρυζιού (*Oryza sativa* L.) οκτώ διαφορετικών ποικιλιών ως προς την ικανότητά τους να επάγουν συμπτώματα ασθένειας στα φυτά, να ενισχύουν την ανάπτυξη των φυτών και την ανταγωνιστική δράση τους έναντι του βακτηρίου *Xanthomonas oryzae* ρν. *oryzae*. Από τα 63 στελέχη, πέντε ήταν φυτοπαθογόνα. Με βάση τα φαινοτυπικά χαρακτηριστικά και την ανάλυση αλληλουχίας 16S rDNA, τα στελέχη αυτά ταυτοποιήθηκαν ως *Pseudomonas oryzihabitans*, *P. fulva*, *Pantoea ananatis*, *Pantoea* sp., *Cellulomonas* sp. Τέσσερις από τις 63 απομονώσεις έδειξαν ότι είναι δυνητικά καλοί παράγοντες επαγωγής της ανάπτυξης των φυτών και βιολογικής αντιμετώπισης. Τα στελέχη αυτά αναγνωρίστηκαν ως *Bacillus* sp., *B. subtilis*, *Pseudomonas putida* και *Enterobacter* sp. Αυτή είναι η πρώτη αναφορά παθογόνων ενδοφυτικών βακτηρίων σε καλλιεργούμενο ρύζι στο Βόρειο Ιράν.

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