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Investigation of ACC-Deaminase and Indole Acetic Acid Producing Bacteria from Rhizospheric Soils in Ağrı Province

Burak ALAYLAR^{1*}, Mehmet KARADAYI²

ABSTRACT: Plant growth promoting rhizobacteria (PGPR) have critical role in promoting plant growth and health with various mechanisms. Production of indole acetic acid (IAA) and 1-aminocyclopropane-1-carboxylate (ACC) deaminase activity are well-known plant growth promoting (PGP) properties. In the current study; A total of 48 morphologically distinct bacterial colonies were chosen during the isolation of bacteria from various rhizospheric wheat, maize and sainfoin soil samples from agricultural areas in Ağrı-Turkey. The bacteria were isolated from soil samples utilizing by culture-dependent techniques. Then, the strains were visualised for PGP activities such as production acetic acid IAA and ACC deaminase activity. The 16S rRNA sequence similarity of potential PGPR rhizospheric strains demonstrated that strains belong to species *Bacillus, Pseudomonas, Microbacterium, Pseudarthrobacter, Arthrobacter* and *Enterobacter*. Totally, 11 of 48 isolates exhibited highly IAA producing and ACC deaminase activity. These results show that IAA producing and ACC deaminase activity of local potential PGPR strains isolated from soil rhizospheric which can appropriate for sustainable agricultural applications in Ağrı province.

Keywords: ACC deaminase activity, IAA production, PGPR, 16S rRNA

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INTRODUCTION

Plant-microbe interactions have beneficial processes plenty of key roles in soil health, crop growth and productivity. Especially, soil microorganisms demonstrated positive effects on numerous agricultural applications. Among these soil microorganisms, plant growth promoting rhizobacteria (PGPR) was initially described as; soil bacteria that located into roots of plants and they have fruitful features when inoculation onto seed and improve to plant growth by Kloepper and Schroth (1978). Therefore, researchers have been attracted attention to microorganisms based fertilizers instead of chemical fertilizers because of their high cost and detrimental impacts on sustainable agricultural applications in past decades (Adesemoye et al., 2009; Beneduzi et al., 2012; Alaylar et al., 2019; Ferreira et al., 2019; Alaylar et al., 2020a). Furthermore; they have been identified and reported PGPR bacteria including; Alcaligens, Arthobacter, Azospirillum, Azotobacter, Bacillus, Burkholderia, Enterobacter, Klebsiella, Pseudomonas, Rhizobium and Serratia (Egamberdieava, 2015; Afzal et al., 2017; Ferreira et al., 2019). Besides, various beneficial mechanisms have shown of numerous type of favorable abilities for plant growth with various mechanisms. For example; atmospheric nitrogen (N_2) fixation, solubilization of phosphorus, improved to solubility of nutrients like synthesizing of siderophores, production of plant growth regulators such as cytokinins, ethylene, gibberellic acid, and indole acetic acid (IAA), and 1-aminocyclopropane-1-carboxylate (ACC) deaminase activity (Ahmad et al., 2008; Alaylar et al., 2020b).

According to the literature, one of the most frequently used mechanisms to explain the distinct effects of PGPR on plants is the production of phytohormones (Duca et al., 2014). Among these phytohormones; auxin is one of the important molecules that play a vital role in plants by regulating many processes in plants directly or indirectly (Bal et al., 2012; Junior et al., 2015). The most well-known and researched auxins in plants is indole-3-acetic acid (IAA) because IAA is a crucial physiologically active phytohormone for plants and secondary metabolite of L-tryptophan which have vital role in numerous biological processes as a regulator in plants like; cell expansion, division, differentiation (Lebrazi et al., 2020). Furthermore, it has been reported that plants have more uptake to soil nutrients as bacterial IAA improves both the root surface region and length. Also, most of rhizobacteria can produce IAA. Thus, PGPR strains can change root structure and promote plant development due to their ability to synthesize and secrete plant hormones like IAA, gibberellins (GAs), cytokinins and certain volatiles, hence they are termed phytostimulators (Mohite, 2013; Montana et al., 2014; Vaikuntabu et al., 2014). Especially, potential PGPR strains isolated from the rhizosphere and rhizoplane from different type of agricultural products have performed more potential of IAA production than from the non-rhizospheric soil (Bal et al., 2012).

As it is known, plants are faced to various biotic and abiotic stresses in nature such as cold, drought, flooding, heat, salinity, exposure to heavy metals and lack of nutrients, attack by phytopathogens and pests. Any of these stress conditions are demonstrated to enhancement of ethylene synthesis in plants. Because, ethylene is a precursor of 1- aminocyclopropane-1-carboxylic acid (ACC) so that the ACC deaminase enzyme hydrolyzes as a biosynthetic precursor of the hormone ethylene in plant tissues to ammonia and α -ketobutyrate instead of ethylene. The main mechanisms used by PGPR strains to decrease the stress contains reducing the level of ethylene through hydrolyzing ACC by the enzyme ACC deaminase. Therefore, most of PGPR strains have ACC deaminase activity utilized for to decline hazardous effects of ethylene under stressed conditions (Abbas-Zadeh et al., 2010; Qin et al., 2014; Kadioglu et al., 2016; Misra and Chauhan 2020).

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PGPR strains and their relations with plants are used commercially nowadays and it has huge potential and promise for sustainable agriculture because of their beneficial effects on plenty of agricultural products. It has been studied and examined various type of agricultural crops containing barley, canola, maize, lentils, pepper, cucumber, rice, soy, cowpea, wheat *etc*. (Adesemoye et al., 2009; Hayat et al., 2010; Sharan and Nehra, 2011).

In this study, we are focused on IAA production and ACC deaminase activity of bacterial strains which were isolated from rhizospheric soil. The main goal of the present study was identified local potential PGPR strains isolated from soil rhizosphere and quantified them *in vitro* production of IAA and ACC deaminase activity and showed to these potential PGPR strains can be used in as local specific strains in agricultural areas in Ağrı province.

MATERIALS AND METHODS

Chemicals

All of the chemicals utilized in the current study were purchased from Merck (Hohenbrunn, Germany), Fluka (Steinheim, Germany), Sigma-Aldrich (St. Louis, USA), Promega (Madison, USA), Oxoid (Hampshire, England), LAB M (Lancashire, England) and Lonza (Rockland, USA).

Collection of Soil Samples and Isolation of Rhizospheric Bacteria

Rhizospheric bacteria were isolated from soils samples in agricultural lands from Ağrı-Turkey as following procedure in the month of August 2020. Sampling was taken from 3 different sites, at depts of 0-7 cm in the region of directly surrounding the roots of the wheat, maize and sainfoin plants rhizosphere in agricultural areas. Then, soil samples were transferred under aseptic conditions to Central Research and Application Laboratory, Agri Ibrahim Cecen University. Then serially diluted soil samples (10⁻¹-10⁻⁷) were spread onto Lauria Bertani (LB) agar incubated for 2 days at 28 °C. With the end of incubation period, distinct bacterial colonies were streaked on LB agar plates to obtain pure and single colonies for further studies.

Determination of the Indol Acedic Acid Production

IAA production is one of the major pivotal natural auxins that promote plant growth. The assay was prepared according to Vaikuntapu et al., (2014) procedure. This procedure was performed as following; bacterial isolates were grown in LB broth medium containing 0.1% tryptophan and incubated for 72 hours at 28 °C. After the incubation, the isolates were centrifuged at 6000 g for 30 minutes. Then, 2 ml of the supernatant was taken and mixed with 2 drops of ortho-phosphoric acid and 4 ml of Salkowski solvent (50 ml of 35% of perchloric acid + 1 ml of 0.5 M FeCl₃ solution). Finally, potential IAA production bacterial isolates calorimetric measurement was carried on at 530 nm.

Determination of the ACC Deaminase Enzyme Production

ACC deaminase enzyme activity was analysed according to the method of Penrose and Glick (2003). In this present protocol, bacterial isolates were inoculated in LB broth medium and incubated at 28 °C for 24 hours. After an overnight incubation period, 2 ml of each bacterial culture was taken and centrifuged at 8000 g for 5 minutes. Then, pellets of each sample was washed two times with Dworkin and Foster minimal salts medium. Furthermore, 2 ml of Foster minimal salts medium supplemented with ACC as only nitrogen source (DF-ACC) were transferred to bacterial cultures. The cultures were incubated at 28 °C for 24 h. Moreover, non-inoculated DF-ACC medium was used as a control group of the experiment and blank sample for the spectrophotometric measurement. Then, cultures were centrifuged at 8000 g for 5 minutes. 1 ml of each sample were taken from the cultures and centrifuged at 8000 g for 5 minutes. 100 µl of the supernatant was diluted with 1 ml of DF medium. 60 µl of each

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working solution was used in the standard ninhydrin experiment (60 μ l working solution +120 μ l ninhydrin solution). The solution was mixed and situated in a boiling water bath for 30 minutes at 65°C. After boiling, the solution turned into a purple color. The samples were cooled at the room temperature for 20 minutes. When compared to the control, the opaque visibility in color can be evaluated as a positive result. Finally, the calorimetric measurement was performed at 570 nm.

Molecular Identification of Potential IAA and ACC Deaminase Enzyme Production Capability of PGPR strains

Genomic DNA isolation of the selected potential IAA and ACC deaminase enzyme producer PGPR isolates was used according to method defined by Wilson in 1997. The isolates were identified through 16S rRNA gene sequencing. The partial16S rRNA gene regions were amplified utilizing polymerase chain reaction (PCR) for molecular identification of the bacterial isolates. The primers were 27F (5'-AGAGTTTGATCMTGGCTCAG-3') and 1492R (5'-CGGTTACCTTGTTACGACTT-3'). The reaction mixtures (30 µl) reaction included 50 ng of extracted as a template, 1.5 mM MgCl₂, 0.2 mM each dNTP, 1.2 µl of dimethyl sulfoxide (DMSO), 25 pmoles of forward primer and reverse primer, and 5 U Taq DNA polymerase along with reaction buffer. The amplification procedure was contained following reactions: The initial step at 95 °C for 2 min, and 36 cycles of 1 min at 94 °C, 1 min at 53 °C, 2 min at 72 °C, followed by a final 5 min extension step at 72 °C, then brought down to 4 °C. Amplified 16S rRNA gene PCR products were analysed by QIAxcel® advanced analysis system and sequenced by Macrogen Inc. BLAST Netherlands. The nucleotide BLAST (Basic Local Alignment Search Tool) search program of NCBI was exploited to determine the nucleotide sequence homology and relation to other bacterial sequences existing in the GenBank® (http://blast.ncbi.nlm.nih.gov./blast.cgi). Evolutionary analyses were conducted in MEGA X (Kumar et al., 2018)

RESULTS AND DISCUSSION

A total of 48 morphologically distinct bacterial isolates were isolated on August from agricultural areas from rhizospheric soil samples from 3 different localities in agricultural lands in Ağrı-Turkey. The rhizospheric bacteria isolated and identified utilizing by 16S rRNA gene analysis. Totally, 11 potential bacterial species were shown plant growth promoting activities not only IAA producing but also ACC deaminase avtivities from rhizospheric soil samples. The isolates were on 100% identical to the closest relatives assigned to GenBank. Determination of nucleotide sequence homology was established via nucleotide BLAST search program of NCBI. The 16S rRNA sequences were deposited to GenBank® with provided accession numbers from MW426255 to MW426265 respectively. Detailed information was given in Table 1.

On the basis of 16S rRNA sequences of the isolated strains and the closest relatives from GenBank the neighbour joining phylogenetic tree was constructed (Figure 1). As it's shown in Table 2; 11 isolated species belonging to 3 distinct phyla and 5 order: Proteobacteria (BM2, BM9, BM10, and BM11), Firmicutes (BM1, BM4, BM5, and BM6), Actinobacteria (BM3, BM7, and BM8).

Order I – Bacillales is one the most plenty of order and contains 4 species: *Bacillus safensis* strain BM-1, *Bacillus simplex* strain BM-4, *Bacillus pumilus* strain BM-5, and *Bacillus* sp. strain BM-6. Junior et al. were reported (2015) production of indole-3-acetic acid by *Bacillus* isolated from different soils. Cruz-Martín and coauthors were reported (2015) *B. pumilus* CCIBPC5 grow in culture media without nitrogen, and produced indole acetic acid. Furthermore, *B. pumilus* enhanced dramatically the height and stem thickness and improves dry and fresh weight of plants. On the other study, Misra and Chauhan (2020) were shown ACC deaminase-producing rhizosphere competent of *B. safensis* reduce salt stress and promote *Zea mays* L. growth by modulating ethylene metabolism. Moreover; there are numerous

studies were reported production of IAA and ACC-deaminase activity as PGP properties of as *Bacillus* sp. (Rodriquez–Romero et al., 2005; Shim et al., 2014; Gupta and Pandey, 2019).



Figure 1. Phylogenetic tree on the basis of 16S rRNA were created using the Neighbor-Joining method (Felsenstein, 1985; Saitou and Nei 1987; Tamura et al., 2004). Evolutionary analyses were carried on MEGA X (Kumar et al., 2018).

Order II–Pseudomonadales is second major order of the current study and it is including 3 various species; *Pseudomonas brassicacearum* strain BM-2, *Pseudomonas kilonensis* strain BM-10, and *Pseudomonas* sp. strain BM-11. According to the literature, it is also reported more studies about their PGP. Therefore, *Pseudomonas* sp. is known one of the common genus of PGPR strains which are isolated from rhizospheric soil samples. For instance; Qin et al. (2014) were found activity of ACC deaminase and IAA producing bacteria by 16S rRNA gene sequence *P. brassicacearum* subsp. *brassicacearum* from coastal halophytic medicinal plant. Moreover, *Pseudomonas* species are known not only IAA production and ACC- deaminase activity, but also other PGP properties such as; nitrogen fixation, phosphate solubilization and siderophore production and *Pseudomonas* species have been reported intensively utilized in agricultural applications according to the literature (Gu et al., 2020).

Order III– Actinomycetales contains 2 species: *Microbacterium* sp. strain BM-3 and *Arthrobacter* sp. strain BM-8. A study by Samayoa et al. (2020) were isolated and screened of PGP properties of numerous distinct type of bacterial species from soil rhizosphere and root endophyte of onion. *Microbacterium hydrocarbonoxydans* and *Microbacterium laevaniforman* have shown almost all PGPR

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properties including IAA production and ACC deaminase activity and also *Microbacterium* sp. was shown IAA production according to their study.

Order IV– Micrococcales presented by single species *Pseudarthrobacter oxydans* strain BM-7. Mesa-Marín with coauthors (2019) were reported effect of PGPR on *Salicornia ramosissima* seed germination under salinity and *Pseudarthrobacter oxydans* SRT15 was showed PGP properties such as; nitrogen-fixation, phosphate solubilization and IAA production.

Order V–Enterobacterales also presented by one species *Enterobacter ludwigii* strain BM-9. Singh (2015) studied some of the PGP properties of *Enterobacter ludwigii* PGP 19 isolated from soil rhizosphere of *Pennisetum glacum* (pearl millet). The strain *E. ludwigii* PGP 19 was able to active phosphate solubilization, IAA production, and ACC deaminase activity. According to a report by Aarab et al. (2015) *E. ludwigii* isolated rhizospheric soils of rice demonstrated IAA production.

Strain Code	Strain Name	Accession number	Percentage identity
BM1	Bacillus safensis	MW426255	100
BM2	Pseudomonas brassicacearum	MW426256	100
BM3	Microbacterium sp.	MW426257	100
BM4	Bacillus simplex	MW426258	100
BM5	Bacillus pumilus	MW426259	100
BM6	Bacillus sp.	MW426260	100
BM7	Pseudarthrobacter oxydans	MW426261	100
BM8	Arthrobacter sp.	MW426262	100
BM9	Enterobacter ludwigii	MW426263	100
BM10	Pseudomonas kilonensis	MW426264	100
BM11	Pseudomonas sp.	MW426265	100

Table 1. Taxonomic affiliation of the bacterial isolates and their GenBank® accession numbers

Table 2. Detailed information of IAA production and ACC deaminase activity of potential PGPR

 strains (+: slight positive, ++: moderately positive, +++: strong positive)

Species Name	IAA	ACC		
	Production	Deaminase Production	Order	Phylum
Bacillus safensis BM-1	+++	+++	Bacillales	Firmicutes
Pseudomonas brassicacearum BM2	+++	+++	Pseudomonadales	Proteobacteria
Microbacterium sp. BM-3	+++	+++	Actinomycetales	Actinobacteria
Bacillus simplex BM-4	+++	+	Bacillales	Firmicutes
Bacillus pumilus BM-5	+++	+++	Bacillales	Firmicutes
Bacillus sp. BM-6	+++	+++	Bacillales	Firmicutes
Pseudarthrobacter oxydans BM-7	+++	+++	Micrococcales	Actinobacteria
Arthrobacter sp. BM-8	+++	++	Actinomycetales	Actinobacteria
Enterobacter ludwigii BM-9	+++	+++	Enterobacterales	Proteobacteria
Pseudomonas kilonensis BM-10	+++	+	Pseudomonadales	Proteobacteria
Pseudomonas sp. BM-11	+++	+	Pseudomonadales	Proteobacteria

It was also shown by previous studies; rhizospheric bacteria from soil samples have capability to fix atmospheric nitrogen, phosphate solubilization, synthesing siderophores, ACC deaminase activity and type of phytohormones such as auxins for provide it to plants as a plant growth property (Alaylar et al., 2018; Ferchichi et al., 2019). IAA production is one of the most effective PGP mechanism (Pawlik et al., 2017). It is promoting to plant growth directly by enhancing length and root surfaces by stimulating

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plant cell elongation for providing to soil nutrient from plants (Phetcharat and Duangpaeng 2012; Vinayarani and Prakash 2018). There are many reports have been shown IAA production by rhizospheric and endophytic bacteria (Ray et al., 2016; Chowdhury et al., 2017; Vinayarani and Prakash 2018; Wozniak et al., 2019). Various rhizospheric bacteria have potential to produce the auxin phytohormone IAA, these kind of rhizospheric microorganisms produce auxins in the presence of a favourable precursor like L-tryptophane (Phetcharat and Duangpaeng 2012). On the other hand, most of the plant growth promoting rhizospheric strains are well-known with ACC deaminase activity which have role in decrease to level of ethylene stress and help their host plants deal with numerous stress factors like abiotic and biotic factors.

In the line with this purpose, we observed IAA production and ACC deaminase activity as PGP by rhizospheric soil samples in agricultural areas from Ağrı-Turkey. When the compare with our results with 25 of 48 strains were demonstrated IAA production. 11 of 48 the best IAA production potential stains were chosen. On the other hand, 11 of 48 the rhizospheric bacteria were shown ACC deaminase activity.

In summary, as it mentioned it; PGPR have been revealed to beneficial effects of plant development and health. For sustainable agricultural applications, microorganism based fertilizers can be used instead of chemical fertilizers because of their eco-friendly low cost usage. Moreover, potential local PGPR strains can be more effective usage on agricultural products. Therefore, the results showed that, plant growth promoting property IAA production and ACC deaminase activity from rhizospheric bacteria with the selected from soil rhizosphere. According to our knowledge, the present study reveals for the first time to the isolation and identification of rhizospheric bacteria from soil samples in Ağrı province which have IAA production and ACC deaminase activity. It can be use in agricultural application for further.

CONCLUSION

It can be concluded that soil rhizosphere as the huge potential source for obtaining new bacterial strains and their metabolites. In this regards; It can be understood from the present study that rhizospheric bacteria isolated from agricultural areas have crucial PGP such as IAA production and ACC deaminase activity. It is important to note that unknown rhizospheric bacteria community in soil rhizosphere in agricultural areas can be vital role in agricultural products. Rhizospheric bacteria deal with agricultural products such as wheat, barley, canola, oat, cucumber *etc*. have unique strategies that enhance growth and yield productivity of agricultural products. Taking into consideration of its economic value and vital roles of PGPR strains in plant health, enhance productivity, growth, such knowledge gap needs to be filled for active utilization of PGPR strains and detection of novel PGPR strains from local strains in agricultural areas.

Conflict of Interest

The article authors declare that there is no conflict of interest between them.

Author's Contilbutions

The authors declare that they have contributed equally to the article.

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