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Agriculture and Economic Development of Nepal: A Comparative Study between Two Different Time Period

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Yadav Mani Upadhyaya and Khom Raj Kholel²

Asterisk (*) for the corresponding author

Superscript for indicating affiliation of author and coauthors

¹Department of Economics, Tribhuvan University, Saraswati Multiple Campus, Kathmandu, Nepal

² Departments of Economics, Tribhuvan University, Saraswati Multi Campus, Kathmandu, Nepal

Superscript for indicating affiliation of author and co-authors

Times New Roman, font size 9, Lower case and centred for affiliation of author and co-authors

rresponding author email: yadav.upadhaya@smc.tu.edu.np

Asterisk (*) for the corresponding author

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Abstract

The Nepalese education system is more competitive and harmonious although for the lack of system is still in the transition phase. The study has focused on the impact of public expenditu there is increasing government involvement and how it has continued with the rising education this article is to show the importance of public expenditure on primary education in the context of Nepal in Gorkha district. The data gathering tool used is secondary methods from the publication of A structural simple statistical method is used in which budget on primary education is taken as dependent varia that the government contribution to primary education imbalance, education should provide more incentive to

3-6 key words, separated by semi colon (;)

primary education and try to show how a expenditure. The focus of the objective of cture of Gorkha' and 60 households has been taken. imed at a specific type of information. Government variable in this study. This research article concludes development of education in Nepal. To reduce the

nd resources. The present education

Key words: Education; public expenditure; household income; government budget; positive impact

References to the literature cited for the manuscript should be written as shown here. In case of multiple references, they should be separated by (;) sign

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Introduction

arboxylate (ACC) deaminase, an enzyme capabl 1-Aminocyclopropa afolyzing ACC, the immediate precursor of sorted in 1978, which was isolated from *Pseudor* sp. strain ACP (Honma and Shimomura, 1978). ethylene was firstly The ACC deaming has been detected in the fungus, Penicillium citridum and in a number of bacteria (Ma et al., 2003; Blaha et al., 2006; Madhaiyan et al., 2006; Belimov et al., 2007). The gene responsible for ACC deaminase activity (acdS) has been recently found in Azopirillum., Burkholderia cepacia genomovars (which include PGPR, phytopathogens and opportunistic human pathogens), and Agrobacterium genomovars (Blaha et al., 2006). These microorganisms were identified by their ability to grow on minimal medium containing ACC as its sole nitrogen source (Honma and Shimomura, 1978; Belimov et al., 2007; Ma et al., 2003).

The plant growth-promoting bacterium, Pseudomonas putida GR12-2, which contains the enzyme ACC deaminase, stimulates root elongation (Glick et al., 1994) and significantly reduces the level of ACC in emerging roots and shoots. Three separate mutants of Pseudomonas putida GR12-2, deficient in ACC deaminase activity, were reported to lose the ability to promote canola root elongation under gnotobiotic conditions (Glick et al., 1994). The ACC in the exudates may be taken up by the bacteria and subsequently hydrolyzed by the enzyme, ACC deaminase, to ammonia and α-ketobutyrate. The uptake and cleavage of ACC by plant growth-promoting bacteria decrease the amount of ACC outside the plant. Increasing amounts of ACC are exuded by the plant in order to maintain the equilibrium between internal and external ACC levels. It is proposed that plant growthpromoting bacteria that possess the enzyme ACC deaminase and are bound to seeds or roots of seedlings can reduce the amount of plant ethylene and thus cause inhibition of root elongation. Thus, these plants should have longer roots and possibly longer shoots as well, in as much as stem elongation, except in flooding resistant plants, is also inhibited by ethylene (Abeles et al., 1992). Soil bacteria that have ACC deaminase activity should then have a selective advantage over other soil bacteria in situations in which the main bacterial nutrients are from exudates of plants (Shah et al., 1998). It should also be borne in mind that soil bacteria may acquire ACC deaminase genes by mechanisms other than fortuitous mutation-transfer of such a gene from another soil bacterium is another possible mechanism. The regulation of ethylene production in plants, especially to prevent increased ethylene production and accumulation, may reduce many of the inhibitory effects of this hormone (Jacobson et al., 1994). Many agricultural and horticultural crops are particularly sensitive to ethylene levels which regulate fruit ripening and control the deleterious effects of senescence in vegetables and flowers (Sisler and Serek, 1997). The bacterial enzyme, ACC deaminase, is potentially a valuable tool for controlling the levels, and hence the effects of ethylene in plants. ACC deaminase has already been used to substantially reduce ethylene levels in transgenic tomato plants which have exhibited a prolonged ripening phase (Klee et al., 1991) and to lower stress ethylene levels following infection by bacterial and fungal pathogens (Lund et al., 1998). Strains of plant growth-promoting bacteria that contain ACC deaminase are known to reduce ACC, and hence

ethylene, levels in canola seedlings (Penrose and Glick, 2003), promote root elongation in a variety of plants (Hall *et al.*, 1996), decrease the deleterious effects of flooding on tomato plants, and prolong the shelf-life of ethylene sensitive cut flowers (Klee *et al.*, 1991). Therefore, plant growth promoting bacteria containing ACC deaminase activity can be utilized for the improvement of crop yields. Based on our knowledge, there is no any report of *Microbacterium* having ACC deaminase activity present in the rhizosphere of plants, whereas, only a few endophytes has been reported for this activity. The objective of the present study was to evaluate the activity of the ACC deaminase enzyme in rhizospheric plant growth promoting rhizobacteria with special reference to *Microbacterium* in the various conditions. In addition to above objective, we attempted to find out the presence of *acdS* gene in all the strains used in this study.

Materials and Methods

Bacterial strains and growth conditions

Bacterial strains and their culture conditions of rhizospheric plant growth promoting bacteria showing nitrogen fixation, IAA production, siderophore production and P-solubilization properties were previously reported (Shrivastava, 2012). With an objective to optimize of various growth conditions for the estimation of ACC deaminase activity and amplification of *acdS* gene, nine most efficient strains isolated from rice rhizosphere of Indo-Nepal border region (1 *Microbacterium* sp strain ECI-12A; 3 *Klebsiella* sp strains ECI-10A, AF-4C and BN-4A; 2 *Agrobacterium* sp strains AF-1D and BN-2A; 2 *Pseudomonas* sp strains AF-4B and PN-4D and 1 *Serratia* sp strain AF-5A) were taken.

Estimation of 1-Aminocyclopropane-1-Carboxylate Deaminase (ACC Deaminase) Activity

For the measurement of ACC deaminase activity, selected isolates were grown overnight in 10 mL of NB medium and thereafter harvested by centrifugation. The pellet was washed with normal saline and suspended in 7.5 mL of JNFb medium containing 5 mM of 1-aminocyclopropane-1-carboxylate (ACC). Tubes were incubated at 28°C with shaking (120 rpm) for growth. ACC served as the sole source of nitrogen in the medium. After 24h of growth, cells were centrifuged at 8000 rpm at 4°C for 10 min. The pellet was suspended in 1 mL of 0.1M Tris-HCl (pH 7.6) and again centrifuged at 15000 rpm for 15 min. Pellet was collected and supernatant discarded. The pellet was re-suspended in 600 µL of 0.1 M Tris-HCl (pH 8.5). 30 µL of toluene was added to the cell suspension and vortexed at higher setting for 30 s. Tube was kept at 4°C for 1h and then centrifuged at 1200 rpm for 10 min at 4°C. The thin layer of toluene was aspirated by micro-pipette gently. Now, the toluenized cells were equally distributed in two eppendroff tubes. First part was stored at 4°C for protein assay and other part was used for ACC deaminase assay immediately. 200 µL of toluenized cells was transferred in a fresh 1.5 mL microcentrifuge tube and 20 µL of 0.5 M ACC was added to the suspension. It was briefly vortexed and incubated at 30°C for 15 min. 1.0 mL of 0.56 M HCl was added, vortexed and centrifuged for 10 min at 12000 rpm. Now, 1mL of the supernatant was taken in another tube and 800 µL of 0.56 M HCl was added and vortexed briefly. Thereafter 300 µL of 2,4, dinitrophenylhydrazine (2 % w/v) was added to the tube. It was mixed properly by vortexing and incubated at 30°C for 30 min. 2µL of 2M NaOH was added and after mixing absorbance was recorded at 540 nm. The amount of µmol of \alpha-ketobutyrate produced by this reaction was determined and compared with a standard curve of α-ketobutyrate ranging between 0.1 and 1.0 µmol. For the purpose of standard curve generation a sock solution of 100 mM αketobutyrate (Sigma-Aldrich Co., USA) was prepared in 0.1M Tris-HCl pH 8.5 and stored at 4°C. Enzyme activity was expressed as μ mol/mg protein/h.

Amplification of acdS gene

The primers 5'- GGCAAGGTCGACATCTATGC-3' and 5'-GGCTTGCCATTCAGCTATG-3' (Duan et al., 2009) were used to amplify acdS gene. The thermal profile for amplification was 2-min initial denaturation at 94°C, 35 cycles of 1-min denaturation at 92°C, 50-s primer annealing at 58°C, and 1 min of elongation at 72°C. The amplified products were visualised with ethidium bromide stained agarose gel electrophoresis.

Result and Discussions

Estimation of ACCD activity in selected isolates

ACCD activity is indirectly responsible for growth promotion in plants; therefore, its activity was measured. Out of nine isolates tested highest activity was found in *Klebsiella* sp strain ECI-10A followed by *Pseudomonas* sp strain AF-4B. All the isolates showed activity in the range of 122 - 539.1 nmol α -ketobutyrate/mg protein /h (Table 1).

Time course assay of ACCD activity

Since all the isolates tested showed appreciable level of ACCD activity, it was desirable to test time course appearance of activity. Accordingly the *Microbacterium* sp strain ECI-12A was incubated in JNFb⁻ medium containing 5mM ACC and ACCD activity was measured at desired time intervals (0, 4, 8, 12, 24 and 48h). It is evident from the data of Fig.1 that there was presence of some basal activity at 0 h which increased with time of incubation. Activity showed linear increase from 8 h, the maximum level was attained at 24 h (316.0 nmol α -keto butyrate/mg protein/h). Beyond 24 h there was no significant increase; it was almost constant after 24 h (Fig. 1)

Do varying concentrations of ACC affect activity?

Effect of varying concentrations of ACC was tested so as to find out the optimal concentration required for the ACCD activity. It is evident from the result of Fig. 2 that there was negligible activity without ACC addition to culture, but activity started appearing with the addition of as low as 0.1mM ACC in the culture medium. The level of enzyme activity increased with the increasing concentrations of ACC and maximum increase was attained with 5mM ACC. Further increase in ACC concentration did not show any increase in activity (data not shown).

Effect of varying temperature on ACCD activity

Since temperature plays important role in the regulation of plant growth promoting features of any bacteria, it was desirable to test ACCD activity at varying temperature. Keeping this objective in mind, ACCD activity at varying temperature was tested in *Microbacterium* sp strain ECI-12A. It is evident from the data of Fig. 3 that maximum ACCD activity was observed at 30°C and thereafter there was decrease. Only 33.2% activity was left at 35°C (Fig. 3).

Amplification of acdS gene

The PCR amplification of acdS gene responsible for ACCD activity was done in all the nine isolates. It is evident from the gel photograph (Fig. 4) that all the isolates showed amplicon of acdS gene (~ 1.0 kb).

Amelioration of salt stress has been reported in canola (Brassica napus L.) growth by the ACCD containing fluorescent pseudomonads (Jalili et al., 2009). Prevalence of ACC deaminase activity in various PGPR including Enteobacter cloacae (Belimov et al., 2005), Pseudomonas sp ACP (Sheehy et al., 1991), Serratia quinivorans SUD165 (Belimov et al., 2005), Pseudomonas putida strain UW4 (Shah et al., 1998) and Klebsiella pneumonie strain Kp 342 (Iniguez et al., 2005) as well as certain yeast and fungi (Yao et al., 1995) has been reported. Results of the present finding are in agreement with above reports since isolates reported here in also belong to the genus Klebsiella sp (AF-4C and BN-4A), Pseudomonas sp (AF-4B and PN-4D), Serratia sp (AF-5A) and Agrobacterium sp (AF-1D and BN-2A). However presence of ACC deaminase activity in the Microbacterium species has not been reported from rhizosphere of any plant except a few reports from endophytic bacteria. Endophytic Microbacterium sp G16 isolated from rape (Brassica napus) roots showed ACC deaminase activity (Sheng et al., 2009), whereas ACC deaminase activity showing Microbacterium sp ECI-12A isolated from rice rhizosphere is a first report. Time course study of the activity suggests that the presence of ACC in the medium is prerequisite for optimal activity of the enzyme. ACC deaminase activity plays vital role in maintaining ethylene level in any plants. ACC released exudates by roots of plants may be utilized as nitrogen source by root associated bacteria if they possess ACC deaminase enzyme. It has been reported that plants treated with ACC deaminase containing bacteria have longer roots and can show resistance to inhibitory effects of ethylene stress on plant growth. Several stresses such as flooding, heavy metals, salinity as well as pathogens are known to induce ethylene stress. Under such stresses PGPR possessing ACC deaminase activity would be useful to counteract the inhibitory effects imposed by ethylene. Furthermore, results of this study clearly show that all the nine isolates could be exploited to manage the ethylene stress if they establish colonization in root region of any plants.

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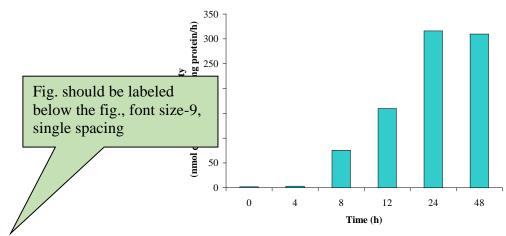


Fig. 1: Time course induction of ACCD activity in *Microbacterium* sp strain ECI-12A. Culture was grown with 5 mM ACC and activity was measured at desired time intervals. Data shown is the average of two independent experiments performed separately in identical condition.

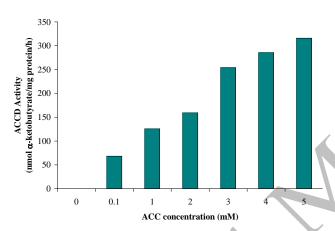


Fig. 2: ACCD activity in *Microbacterium* sp strain ECI-12A in the presence of varying concentrations of ACC. Data shown is the average of two independent experiments performed separately in identical condition.

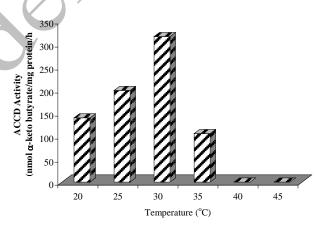


Fig. 3: ACCD activities at varying temperature in the Microbacterium sp strain ECI-12A.

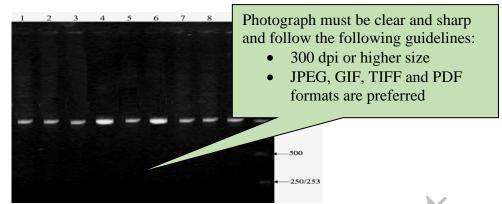


Fig. 4: Agarose gel photograph showing acdS gene fragment (1.0 kb) in nine selected isolates. Lane 1 to 9: acdS gene fragment of various strains (Klebsiella sp strain ECI-10A, Microbacterium sp strain ECI-12A, Agrobacterium sp strain AF-1D, Pseudomonas sp strain AF-4B, Serratia sp strain AF-5A, Klebsiella sp strain AF-4C, Pseudomonas sp strain PN-4D, Agrobacterium sp strain BN-2A, and Klebsiella sp strain BN-4A); M: 1.0 kb DNA marker.

Table 1: ACC deaminase activity in PGP strains

Bacterial strains	ACC deaminase activity
	(nmol α-keto butyrate/ mg protein/h)
Klebsiella sp strain ECI-10A	539.1
Microbacterium sp strain ECI-12A	122.0
Agrobacterium sp strain AF-1D	237.3
Pseudomonas sp strain AF-4B	435.2
Klebsiella sp strain AF-4C	171.9
Serratia sp strain AF-5A	305.7
Pseudomonas sp strain PN-4D	358.4
Agrobacterium sp strain BN-2A	316.0
Klebsiella sp strain BN-4A	261.9

Data shown is the average of two independent experiments performed in identical conditions. The induction of ACCD activity was tested with 5mM ACC

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