Effects of the Inoculation of Burkholderia vietnamensis and Related Endophytic Diazotrophic Bacteria on Grain Yield of Rice

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Abstract

During a survey of endophytic diazotrophic bacteria associated with different rice varieties in Tamilnadu, some "endophytes" were obtained. Thirteen bacterial isolates from surface-sterilized roots and shoots were obtained in pure culture, which produced indole acetic acid (IAA) and reduced acetylene to ethylene. Polymerase chain reaction (PCR) amplification confirmed the presence of nif-H gene in all the isolates. Morphological, biochemical, and molecular characteristics indicated that all of them belonged to the genus Burkholderia One of them, MGK3, was consistently more active in reducing acetylene, and 16S rDNA sequences of isolate MGK3 confirmed its identification as Burkholderia vietnamiensis. Colonization of rice root was confirmed by strain MGK3 marked with gusA gene. The inoculated roots showed a blue color, which was most intense at the points of lateral root emergence and at the root tip. Transverse sections of roots, 15 days after inoculation, revealed beta-glucuronidase (GUS) activity within many of the cortical intercellular spaces next to the stele and within the aerenchyma. Nitrogen fixation was quantified by using 15N isotope dilution method with two different cultivars grown in pot and field experiments. Higher nitrogen fixation was observed in variety Ponni than in ADT-43, where nearly 42% (field) and 40%

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(pot) of the nitrogen was derived from the atmosphere (% Ndfa). Isolate MGK3 was used to inoculate rice seedlings in a comparison with four other diazotrophs, viz., Gluconacetobacter diazotrophicus LMG7603, Herbaspirillum seropedicae LMG6513, Azospirillum lipoferum 4B LMG4348, and B. vietnamiensis LMG10929. They were used to conduct two pot and four field inoculation experiments. MGK3 alone, and combined with other diazotrophs, performed best under both pot and field conditions: combined inoculation produced yield increases between 9.5 and 23.6%, while MGK3 alone increased yield by 5.6 to 12.16% over the uninoculated control treatment.

Microb

Ecology

Introduction

The high-yielding rice varieties of the "Green Revolution^ have resulted in large increases in rice production but require large amounts of nitrogen fertilizers, which contribute to nitrate contamination of soils and ground water supplies, leading to health hazards and environmental pollution. Moreover, for many farmers, the use of chemicals is too costly, especially nitrogen, which is the most frequent limiting factor of rice production. An alternative is the use of bacterial inoculants in crop plants. Rhizobacteria that establish positive interactions with plant roots have been called plant growth-promoting rhizobacteria (PGPR) and are promising for their potential use in sustainable agriculture [\[18\]](#page-14-0). Application of bacterial inoculants as biofertilizers has resulted in

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improved growth and increased yield of cereal crops [\[36](#page-15-0)]. Nitrogen-fixing bacteria belonging to the general Azospirillum, Gluconacetobacter, Azoarcus, Enterobacter, Herbaspirillum, and Burkholderia appear to be frequent colonizers of important cereal crops and grasses and have been extensively studied [[2,](#page-14-0) [3,](#page-14-0) [6,](#page-14-0) [8,](#page-14-0) [19,](#page-14-0) [21,](#page-14-0) [25](#page-14-0) and [50](#page-15-0)].

During the last two decades, molecular techniques have been widely used to study the endophytic diazotrophs in many cereal crops. Previously, methods used to detect and isolate endophytes mostly involve maceration and centrifugation of surface-sterilized plant tissues. These methods can give estimates of bacterial population and extent of colonization but fail to reveal the site of infection of the colonization process. The development in use of reporter genes, such as the gusA gene from Escherichia coli has facilitated investigations of plant– microbe interactions [[17\]](#page-14-0); on the other hand, ^{15}N dinitrogen incorporation assay make it possible to quantify the biologically fixed nitrogen in the plants precisely [\[11](#page-14-0)].

The genus Burkholderia comprises over 50 species; of which, nine species viz., Burkholderia vietnamiensis, Burkholderia kururiensis, Burkholderia tuberum, Burkholderia phymatum, Burkholderia unamae, Burkholderia tropica, Burkholderia xenovorans, Burkholderia phytofirmans, and Burkholderia terrae were verified as nitrogen fixers [\[13](#page-14-0), [22,](#page-14-0) [26,](#page-14-0) [27,](#page-14-0) [29,](#page-15-0) [52,](#page-15-0) [54](#page-15-0) and [64\]](#page-16-0). Recently, Chen et al. [\[15](#page-14-0) and [16\]](#page-14-0) isolated several Burkholderia strains from legume nodules (including B. phymatum and B. tuberum), and some of these have been confirmed to be N-fixing symbionts of Mimosa spp. Of which, the species B. vietnamiensis TVV75 has been extensively studied in Vietnam, for its PGPR effect on rice, both under pot and field conditions. Inoculation significantly increased several yield components, resulting in a final 13 to 22% increase in grain yield [\[62\]](#page-15-0). Baldani et al. [\[4\]](#page-14-0) observed 42–64% increases in growth of rice plants when Burkholderia "brasilensis" and B. vietnamiensis were inoculated under gnotobiotic conditions. Govindarajan et al. [[28](#page-15-0)] inoculated B. vietnamiensis strain MG43 to micropropagated sugarcane plantlets in a comparison with two other diazoptrophs, viz. Gluconacetobacter diazotrophicus and Herbaspirillum seropedicae. Inoculated plants and uninoculated controls were used in a pot experiment followed by two field experiments under different rates of nitrogen fertilizers. Yield increase due to B. vietnamiensis strain MG43 inoculation reached 20% in the field. Taken together, these examples indicate that these bacteria have an important role in improving plant nutrition, in the context of low input sustainable agriculture.

In this study, we report (1) to isolate endophytic nitrogen-fixing Burkholderia spp. from the different cultivated rice fields of Tamilnadu, India, and (2) to investigate endophytic colonization and in planta nitrogen fixation of a selected diazotroph to verify whether the bacteria function as nitrogen-fixing endophytes, and (3) their performance under pot and field conditions as potential bio-fertilizer inoculants capable of increasing rice grain yield.

Material and Methods

In short, the methods used for identification, growth promoting properties, and studying colonization pattern of Burkholderia sp. are: fatty acid analysis, amplification of nif-H gene, sequencing of 16S rDNA, acetylene reduction (AR) activity, IAA production, ${}^{15}N$ dinitrogen incorporation assay, and GUS assay. To determine inoculation effects, two pot experiments followed by four field trials were conducted using with four different cultivars.

Culture Media and Growth Conditions. Isolation media were PCAT and LGIM [[12,](#page-14-0) [23\]](#page-14-0); BDN [\[23\]](#page-14-0) was used for growing strains before DNA extraction; BMGM media [\[23\]](#page-14-0) was used to analysis physiological, biochem- ical, and acetylene reduction activity. LGI-P [\[51](#page-15-0)] and LGIM were used for Gluconacetobacter and Burkholderia recovery; JNFb [[40](#page-15-0)] was used to recover Herbaspirillum and Azospirillum.

Reference Strains. The reference strains used in this study were G. diazotrophicus [[67\]](#page-16-0) strain LMG7603 has been isolated from sugarcane root tissues in Brazil as strain PAL 5 of Acetobacter diazotrophicus [[25](#page-14-0)]. H. seropedicae [[3\]](#page-14-0) strain LMG6513 has been isolated from surface sterilized rice roots in Brazil. B. vietnamiensis [[26\]](#page-14-0) strain LMG10929 has been isolated by Tran Van et al. [\[61\]](#page-15-0), from rice roots in Vietnam. Azospirillum lipoferum [\[6\]](#page-14-0) strain 4B (LMG4348) has been isolated from a rice rhizosphere in France.

Sampling, Isolation, and Enumeration. In ten different fields of Tamilnadu, the roots, stems, and leaves of rice plants were sampled. Cultivars were IR20, Ponni, ADT-37, ADT-43, and Co-47. Isolation followed the method described by Estrada et al. [\[23\]](#page-14-0). Tenfold serial dilutions were used to inoculate (in triplicate) N-free semisolid LGIM tubes. After 96–120 h of incubation, vials were assayed for AR activity following Mascarua-Esparza et al. [\[45](#page-15-0)]. Bacteria growing in nitrogenase-positive vials as a white or yellowish pellicle at a depth of 1 to 4 mm were streaked on LGIM agar plates. All isolates were checked for purity and their ability to grow on modified PCAT agar (0.05 g/l yeast extract powder added) plates [[12](#page-14-0)]. Most Probable Numbers (MPN) of PCAT-growing and AR activity positive isolates were calculated using the McGrady tables. Isolates were maintained in semisolid BMGM for further studies.

Fatty Acid Analysis. The MIDI-FAME technique was used to determine the cellular fatty acid profiles of the isolate MGK3 (most active strain, isolated from surface sterilized stems of cv. Ponni) and the B. vietnamiensis LMG10929. Isolates were grown overnight on trypticase soy agar plates. One loopful of fresh cells was harvested and transferred to a screw cap culture tube. One milliliter of saponification reagent was added. Tubes were tightly sealed with a Teflon-lined screw cap and vortexed for 5– 10 s. The tubes were placed in water bath at $100 \pm 2^{\circ}$ C for 5 min. They were then removed from the boiling water bath and cooled slightly, vortexed for 10 s, and incubated in a water bath for an additional 25 min. After a total of 30 min of saponification in the water bath, the samples were placed under tap water. Each tube received 2.0 ml of methylation reagent, was tightly capped and vortexed for 10 s. The tubes were placed in a water bath at 80 ± 1 °C for 10 min. After that, samples were cooled at room temperature. Then, 1.25 ml of extraction reagent was added into the tubes. The tubes were centrifuged, and the bottom phase was removed using a pipette. Finally, 3.0 ml of base reagent was added, and this was placed again for 5 min in a laboratory centrifuge. The upper solvent phase was removed and transferred to vials for fatty acid analysis.

Physiological and Biochemical Characterization. Pigment production was monitored on nutrient agar and King "B" media. Colony morphology was examined on PCAT and BMGM agar plates. Morphology and Gram type were determined using a Trinocular Phase Contrast Fluorescent Microscope (Olympus AX 80T). Bacterial motility was tested by growth in a semisolid 0.3% mannitol motility test medium. Oxidase and catalase tests were determined using commercially available discs (Hi media, Bombay, India). Growth and acid production were tested using the BMGM medium in which the carbon source was replaced by individual carbon substrates (5 g/l) such as D-glucose, sorbitol, meso-inositol, mannose, glycerol, L-rhamnose, lactose, fructose, L-arabinose, trehalose, L-raffinose, meso-erythritol, galactose, mannitol, cellobiose, xylose, sucrose, starch, sodium acetate, maltose; organic acids were also tested including adipic, malonic, succinic, oxalic, valeric, fumaric, hippuric, malic, tartaric, keto glutaric, and citric acids. Growth on the following amino acids was tested (in the presence of sorbitol as a carbon source): cysteine, glutamic acid, proline, trytophane, leucine, threonine, histidine, lysine, tyrosine, and valine.

Isolation of Total DNA. Isolate MGK3 and B. vietnamiensis LMG10929 were grown in BDN medium at 28°C for 24 h and centrifuged at 12,300 \times g. The pellet was washed with Tris–ethylenediaminetetraacetic acid (EDTA) (TE) buffer, then resuspended in 10 ml of TE $(1\times)$ with 3 ml of 5% sodium dodecyl sulfate (SDS) in TE $(1\times)$ and 3 ml of proteinase K 2.5 mg/ml. This was

then incubated at 37° C for 1 h. The cleaned lysates were extracted with phenol is to chloroform is to isoamylic alcohol (25:24:1). DNA was precipitated by adding onetenth volume of 3 M sodium acetate (pH 5.2) and 2.5 volume of ethanol to the supernatant. Dried pellets were dissolved in $1 \times$ TE buffer.

PCR Amplification of the nif-H Genes. PCR amplification was performed to determine the presence of nif-H gene using specific primers described by Ueda et al [[63\]](#page-15-0). Amplification reactions were performed in a total volume of 25 µl. The reaction mixture contained: 2.5 μ l 10 \times PCR buffer, 2.5 μ l of 2 mM each of dATP, dCTP, dTTT, and dGTP; 3 µl of each forward and reverse primer (30 ng), 1 µl of template DNA (10 ng) and 0.3 μ l of (3 U/ μ l) Taq polymerase; final volume was made into 25 µl using milli-Q water. The step-up PCR procedure included denaturation at 95°C for 3 min, 52°C for 1 min, and 72° C for 1 min, followed by 30 cycles of 95°C for 1 min, 54°C for 30 s, and 72°C for 1 min, with a final extension at 72° C for 10 min. Amplification products were electrophoresed on 1.5% agarose gel in 1 × Tris–borate–EDTA (TBE) buffer.

PCR Amplification and 16S rDNA Sequencing. The 16S rDNA gene sequences were determined by PCR amplification [\[43](#page-15-0)] and direct sequencing [[31\]](#page-15-0). For the phylogenetic analyses, related 16S rDNA gene sequences within the genus Burkholderia were downloaded from GenBank. 16S rDNA sequences were aligned by using the MEGALIGN program of DNASTAR. An evolutionary distance matrix was generated as described by Jukes and Cantor [[38\]](#page-15-0). The evolutionary tree for the dataset was inferred from the neighbor-joining method of Saitou and Nei [\[53](#page-15-0)] using the neighbor-joining program of MEGA version 2.1 [\[42](#page-15-0)]. The stability of relationships was assessed by performing bootstrap analyses of the neighbor-joining data based on 1,000 resamplings.

Nucleotide Sequence Accession Numbers. The nucleotide sequences of diazotrophs isolated in this work have been deposited under the accession numbers to NCBI: AY789926-MGK3.

Nitrogenase Activity. Nitrogen fixation by the isolate MGK3 was determined in semisolid BMGM medium by the acetylene reduction method. The vials containing 10 ml of the BMGM semisolid media were inoculated with single colonies and incubated at 28°C for 3 days. Acetylene (10% v/v) was injected into the inoculated vials, and again, they were incubated at 28° C for 24 h. To determine the ability of reduced acetylene on different carbon sources, isolates were incubated on BMGM medium in which the carbon sources was replaced by the individual carbon sources (5 g/l). Eight important carbon sources were used: azelaic acid, fructose, glycerol, succinic acid, sucrose, mannitol, malic acid, and glucose. Acetylene reduction activity was measured using a flame ionization gas chromatograph (Systronic) equipped with a Porapak N column. Uninoculated vials were used as negative controls. The AR activity of inoculated plants roots was determined according to Ladha et al. [[44](#page-15-0)]. Ten seedlings from each treatment were taken at panicle initiation and grain filling stages, and roots were separated and washed twice with sterile water to remove loosely associated bacteria. The roots were then transferred to fresh, N-free, liquid Jensen's medium [\[58](#page-15-0)]. The tubes containing the roots were sealed with a rubber seal, and 10% of the headspace volume was replaced with acetylene. Uninoculated plant roots and tubes not injected with acetylene served as controls. Tubes were maintained at 25 ± 2 °C with a relative humidity of 75% and 16 h light $(60 \text{ m }\text{lum }\text{m}^{-2})$ and 8 h dark. AR activity was determined as mentioned above.

IAA Production. To quantify the production of IAA by the isolate MGK3 and B. vietnamiensis LMG10929, bacteria were grown in CCM for 1 week, and the cells were pelleted by centrifugation at 10,000 \times g for 15 min. The pH of the supernatant was adjusted to 2.8 with HCl and then extracted three times with equal volumes of ethyl acetate [[60\]](#page-15-0). The extract was evaporated to dryness and resuspended in 1 ml of ethanol. The samples were analyzed on high performance liquid chromatography (HPLC) (Shimadzu SPE 10A, 10AD) using an ultraviolet (UV) detector and a Techsphere C-18 column. Pure IAA was used as a standard. Methanol is to acetic acid is to water (30:1:70 $\nu / \nu / \nu$) was used as a mobile phase at the rate of 1.2 ml/min [[49\]](#page-15-0).

GUS Labeling of Burkholderia sp. Strain MGK3. E. coli (S17.1 pir) containing the constitutive transposons (mTn5ssgusA11) in plasmid pCAM111 [\[66\]](#page-16-0), which was used as the donor, was grown in Luria agar medium containing 100 mg I^{-1} of ampicillin at 37°C with shaking overnight. pCAM111 was transferred to MGK3 by conjugation using a filter-mating technique [\[57\]](#page-15-0). The transconjugants7 were isolated on LGIM medium containing spectinomycin $(100 \text{ mg } l^{-1})$ and gusA substrate 5-bromo-4-chloro-3-indolyl-b-D-glucuronate (X-Gluc; Biosynth AG, Switzerland, at 500 mg I^{-1}). The transconjugants were selected based on the phenotypic expression in the presence of substrate containing medium. The marker strains were further analyzed for the presence of marker gusA gene by PCR technique [\[41\]](#page-15-0).

Plant Inoculation and Gus Staining. Dehulled seeds of rice (Oryza sativa) cultivars, ADT-43, ADT-37, Co-47, and Ponni were surface-sterilized with 95% (ν / ν) ethanol for 2 min and 2.5% (w/v) NaClO₃ for 30 min, followed by washing seven times with sterile water. They were then transferred to plates containing 0.75% (w/v) agar for germination. The germinated seeds were transferred aseptically on to filter paper and placed in 50 ml glass tubes with 15 ml modified Jensen's N-free medium [[58\]](#page-15-0) and maintained at 25 ± 2 °C with a relative humidity of 75% and 16 h light (60 m) lum m⁻²) and 8 h dark. The broth culture, 1 ml, of transconjugant (isolate tagged with gusA) was diluted to approximately 10^8 cells ml^{-1} and used for inoculation. For GUS staining, three plant samples (inoculated and uninoculated) were collected at intervals, washed with sterile 50 mM PBS, pH 7, and stained with 500 µg 5-bromo-4-chloro-3indolyl-â-p-glucuronate(X-Gluc) ml⁻¹, 0.1% (w/v) SDS in 1% (ν/ν) Triton X-100, and incubated overnight at 30-C. Samples were then cut into small pieces and fixed in 4% (ν/ν) glutaraldehyde in 50 mM phosphate buffer (pH 7) containing 0.1% (v/v) Triton X-100 under vacuum for 15 min and then incubated overnight at atmospheric pressure. Fixed samples were rinsed in 50 mM phosphate buffer and taken up for sectioning and viewing under the light microscopy.

Enumeration of Transconjugant MGK3 Colonizing the Inoculated Plants. The roots were washed in excess sterile water and then immersed in sterile water and vortexed for 30 s. The resulting solution was serially diluted and placed on LGIM agar plates containing spectinomycin (100 µg m^{-1}) and X-Gluc (40 µg m^{-1}) . Blue colonies were then enumerated. In another set, the roots were surface sterilized by immersion in 95% (ν/ν) ethanol for 5 min followed by treatment with 2.5% (w / v) NaClO₃ for 20 min followed by sterile water wash seven times. It was then macerated in saline and the homogenate was serially diluted and plated on LGIM agar as described above.

¹⁵N Dinitrogen Incorporation. Thirty-day-old inoculated and uninoculated rice (cv. Ponni and ADT-43) seedlings were grown in pot and field experiments and were transplanted in Pyrex tubes $(25.5 \times 2.5 \text{ cm})$ containing sterilized vermiculite. All the plants, including the uninoculated control plants were approximately 10 cm tall. For each treatment, eight plants were used. Half strength of N-free Jensen's medium was used as nutrient source. 15 N-labeled ammonium sulfate (100 mg N/tube) of 10% atom excess was added as a tracer to quantify nitrogen fixation. Plants were kept at $30 \pm 2^{\circ}$ C during the day and 25 ± 2 ^oC at night for 4 weeks. At the time of harvesting, root area of the plant was measured with the Root Image Analysis Program (Washington State University Research Foundation, USA). The plants were dried in an oven at 70°C until no change in weight was noted. The dried plant samples were ground to a fine powder, and total N in these samples was determined by using a semi micro-Kjeldahl

method based on wet combustion in a Rapid Kjeldahl (Labconco, Kansas city, Missouri). The analysis for ^{15}N excess was carried out on a double inlet mass spectrometer (Varian MAT GD 150). Quantification of nitrogen fixation based on isotope dilution was calculated by the formula of Fried and Middleboe [\[24\]](#page-14-0) which is:

% N fixed =
$$
1 \frac{(^{15}N \text{ atom } \% \text{ excess}) \text{fs}}{(^{15}N \text{ atom } \% \text{ excess}) \text{nfs}}.100
$$

Where fs is fixing system and nfs is non-fixing system.

Pot Experiments. Two pot experiments were conducted using two different rice cultivars (ADT-43 and Ponni) differing in length of growth cycle. Dehulled seeds were germinated as mentioned above. Five-day-old uncontaminated seedlings were treated with a suspension of respective cultures containing 10^8 cfu ml⁻¹. After 30 min contact, seedlings were transplanted into pots. Pots were 60 cm in diameter, 45 cm high, and contained 30 kg of soil. Experiments comprised seven levels of inoculation viz. (1) control pots treated with an autoclaved mixture of bacterial strains, (2) G. diazotrophicus LMG7603, (3) H. seropedicae, (4) A. lipoferum LMG4348, (5) B. vietnamiensis LMG10929, (6) isolate MGK3, and (7) a combination of all five strains. In each pot, eight hills were maintained, with three seedlings per hill. Irrigation was natural rainfall and borewell water. G. diazotrophicus LMG7603 counts were done on medium LGIP where they form a typical subsurface pellicle; isolates were obtained from this pellicle by streaking on LGI agar plates supplemented with 50 mg/l yeast extract [\[50](#page-15-0)]. Burkholderia were counted on LGIM [[22\]](#page-14-0) followed by streaking on PCAT [\[11](#page-14-0)] for confirmation. H. seropedicae and A. lipoferum LMG4348 were counted on JNFb agar plates [[40\]](#page-15-0). Leaf N was determined by the microKjeldahl method [\[32](#page-15-0)]. Shoot height, shoot weight, root weight, and tiller numbers were done at two dates (60th and 110th day for ADT-43 and 60th and 150th day for Ponni), and grain weight was measured at harvest.

Field Experiments. Four commercially important cultivars were selected for conducting field trial experiments at different places of Tamilnadu: ADT-43, ADT-37, Co-47, and Ponni. To germinate, rice seeds were tied in gunny bags, dipped overnight in water tanks, and kept in the shade for germination (36 to 48 h). After germination, the first inoculation was performed by adding approximately 10^8 cells per seed (based on the optical density at 600 nm); water was then added to keep the seeds completely immersed overnight to allow bacterial cells to adhere to seeds and roots. Treatments were: (1) control pots treated with an autoclaved mixture of bacterial strains, (2) G. diazotrophicus LMG7603, (3) H. seropedicae LMG6513, (4)

A. lipoferum LMG4348, (5) B. vietnamiensis LMG10929, (6) isolate MGK3, and (7) a combination of all five strains. After incubation, inoculated pre-germinated seeds were sown in nursery beds, which had a surface representing 10% of the experiment area. The nurseries were flooded to 2 cm water above the soil surface after seedling emergence. At the three-leaf stage (25 to 30 days), rice seedlings were transplanted into the field plots. At transplanting, young plantlets were inoculated again by dipping them in the respective bacterial suspension for 30 min. The total area of each experiment was $5,600 \text{ m}^2$, divided into seven 800 m² areas. Control and inoculated plots were separated by mud levees. Irrigation was natural rainfall and bore well water; sub-canals for individual treatments were maintained. Each hill contained three seedlings. For dwarf varieties (ADT-43, ADT-37, and Co-47), the distance between hills was 5 cm and between rows 7 cm. For the tall variety (Ponni), the distance between hills was 7 cm and between rows, 12 cm. Potassium and phosphorus were applied at the rate of 50 and 100 kg ha^{-1} for all treatments and no N fertilization was applied. For estimation of yield components, five subplots per treatments were randomly selected, and five hills per subplot were sampled. Each sample contained plants and the adhering clod. All samples were kept in plastic bags and processed in the laboratory to determine the plant height (from soil surface to the tip of the upper leaf), root weight, number of fertile tillers, and the dry weight of grains.

Statistical Analysis. The data for each treatment were subjected to a variance analysis using the Statgraphics software (Release 5.0, Uniware STSC, Inc.). When analysis of variance showed significant treatment effects, the least significant difference (LSD, $p < 0.05$) test was applied to make comparisons between treatments.

Results

Isolation and Characterization of Nitrogen-Fixing Bacteria from Rice. The inoculation of N-free semisolid LGIM medium with samples from surface sterilized roots, stems, and leaves of rice, followed by subsequent streaking on the PCAT media, allowed the recovery of 13 N2-fixing isolates. Colonies on BMGM agar plates were large yellow with round entire margins. Colonies on nutrient agar and King's B medium were white, smooth without pigment. After transfer to PCAT agar plates, all colonies were small, white, round with entire margins. The most active strain was MGK3, isolated from surface sterilized stems of tall Ponni variety at Ulundurpet, Tamilnadu.

Identification and Growth Promoting Properties. The presence of 16:0 3-OH in fatty acids suggested that isolate MGK3 belonged to the Burkholderia genus. The 16S rDNA gene sequence of isolate MGK3 gave a 100% match with B.

vietnamiensis LMG10929 (Fig. 1). PCR amplification of nif-H showed that isolate MGK3 produced the expected 390 bp amplification products (data not shown). Isolate MGK3 produced more indole acetic acid in the presence of tryptophan (100 mg/l) than in its absence. With tryptophan, isolate MGK3 produced the highest amount (16.4 mg/ml); whereas B. vietnamiensis LMG10929 produced 10.5 μ g ml⁻¹.

Physiological and Biochemical Characterization. Isolate MGK3 was Gram-negative, rod-shaped, aerobic, and motile. Isolate MGK3 showed oxidase and catalase positivity. All grew well at pH 4.0 to pH 7.0 at room temperature (30°C). Isolate MGK3 grew in different carbon sources such as glucose, sorbitol, meso-inositol, mannose, glycerol, fructose, arabinose, trehalose, raffinose, galactose, mannitol, cellobiose, xylose, and sucrose. But isolate MGK3 did not grow on L-rhamnose, mesoerythritol, starch, sodium acetate, and maltose. Isolate MGK3 grew well in the following organic acids: azelaic, succinic, valeric, fumaric, hippuric, malic, tartaric, aketoglutaric and citric acids, whereas isolate could grow on the oxalic, malonic, and adipic acids. Isolate MGK3 and type strains grew well in the presence of the following

Fig. 1 Phylogenetic position of strain MGK3 within the genus Burkholderia on the basis of 16S rDNA gene sequences. The phylogenetic tree was constructed by the neighbor-joining method [\[50\]](#page-15-0), and the 16S rDNA gene sequence of Pelistega europaea LMG10982 was used as the out group. The numbers at nodes indicate the levels of the bootstrap support based on a neighbor-joining analysis of 1,000 resampled data sets.

amino acids: L-cysteine, L-threonine, L-glutamic acid, Lproline, L-tryptophane, L-leucine, L-lysine, L-histidine, L-tyrosine, and L-valine with sorbitol as a carbon source.

Acetylene Reduction Activity. Isolate MGK3 was capable of N_2 fixation with different carbon sources viz., fructose, mannitol, malate, azelaic, sucrose, glucose, succinic, and glycerol as single carbon sources. However, B. vietnamiensis LMG10929 showed negative AR activity with glucose as single carbon source (Table 1). AR activity also was determined at the rice panicle initiation and grain filling stages. The AR activity was approximately three times higher at the grain-filling stage as compared with the panicle-initiation stage. Roots of cultivars ADT-43 and Ponni produced more ethylene compared to other cultivars of both the panicle initiation and grain filling stages (Table [2](#page-7-0)). The bacterial populations in roots, as estimated by ARA-based MPN counts, also were higher at the grain-filling stage than at the panicleinitiation stage.

GUS Staining and Plant Colonization. Blue colonies were observed in the transconjugants grown on LGIM agar containing spectinomycin $(100 \mu g \text{ m}^{-1})$ and X-Gluc (40 μg cm^{-1}) but not in the wild strain because of the absence of gusA gene. Furthermore, specific PCR analysis of the gusA gene revealed a fragment of 1,200 bp in the transconjugant, but no amplification was observed in wild-type isolate MGK3 (data not shown). Inoculated plants roots after 24 h incubation showed that the blue color cells were distributed mainly on the root surface. After 5 days, more intensive colonization was observed in the lateral roots emergence (Fig. [2A](#page-7-0)). The Ponni rhizosphere populations of MGK3 increased up to 15 days after inoculation with the maximum being after 10 days (Table [3](#page-8-0)). Although bacteria could be isolated from surface of the roots at the first day, they could be re-isolated from roots as endophytes only after 5 days $(2.5 \times 10^7 \text{ cells g}^{-1} \text{ fresh weight})$. The same trend was observed in ADT-43 with rhizosphere populations increasing up to 15 days (Table [3](#page-8-0)), while in the endophytic populations from the roots, re-isolation was possible after 5 days $(1.5 \times 10^6 \text{ cells g}^{-1}$ fresh weight) with the maximum at 15 days (1.5×10^8) . In Co-47 and ADT-47, the rhizosphere populations were increased for 15 days (Table [3](#page-8-0)). However, they could be re-isolated from roots as endophytes only after

10 days $(1.5 \times 10^7 \text{ cells g}^{-1}$ fresh weight). The occurrence of a blue color at the root tips after MGK3 gusA inoculation also supports the possibility of entry via root tips. Transverse sections of roots 15 days after inoculation revealed GUS activity within many of the cortical intercellular spaces next to the stele and within the aerenchyma.

 15 N Dinitrogen Incorporation Assay. The acetylene reduction assay is an indirect method to verify nitrogen fixation in planta of endophytes, and the possibility of ethylene emission from plants could not be completely excluded. Therefore, we designed a ^{15}N dinitrogen incorporation experiment to follow $15N_2$ gas into rice plants through endophytic B. vietnamiensis MGK3. Quantification of nitrogen fixation by $15N$ isotopic dilution method showed that the effect of B. vietnamiensis MGK3 inoculation was more prominent in cultivar Ponni as compared to ADT-43 in field and in pot trial plants (Table [2](#page-7-0)). In this rice cultivar, field and pot trial plants showed maximum fixation with 42 and 40% Nfda recorded, respectively. In cultivar ADT-43, field and pot trial plants showed maximum fixation with 39 and 36% Nfda recorded, respectively. Type strain B. vietnamiensis LMG10929 also showed comparable fixation in both cultivars (Table [4](#page-8-0)).

Rice Inoculation Experiments. In both pot trials, the uninoculated control rice hosted a mixture of diazotrophs in its roots, but in low numbers: 21.2 and 40.9×10^3 CFU/g fresh weight at days 50 and 110, respectively. In both pot experiments, inoculation raised the diazotroph population level up to 1.15×10^5 CFU/g fresh weight at 60 days. It stayed at this level onto 110 days, except for G. diazotrophicus LMG7603, whose level decreased down to 73.5×10^3 CFU/g fresh weight. In the fields, uninoculated control plant roots also contained a mixture of diazotrophs but at a low level, similar to what was observed in the controls of pot experiments (data not shown). Differences between inoculated treatments and the controls were visible in all experiments. In all assays, the mixed inoculum performed best. For all measured parameters, it produced a significant increase over the uninoculated control. Increases of shoot biomass over control was 63.8, 29.0, 60.3, 32.7, 12.2, and 32.3% (mean 38.4%) at 60 days after inoculation

Table 1. AR activity of isolate MGK3 and type strain with different carbon sources^a [means (SD), $n = 4$ replicates]

		AR activity with different carbon sources ^b								
Isolate	Fructose	Azelaic	Glvcerol	Succinic acid	<i>Sucrose</i>	Mannitol	Malic acid	Glucose	BMGM	
MGK3 $TVV75^c$	466.32 3.94	166.15 84.11	3009.09 127.67	5.17 191.39	821.1 l.86	840.1 25.65	95.91 96.26	466.32	112.94 241.17	

^aBMGM medium carbon sources were replaced with individual carbon source.

bandar meanin carbon sources were replaced with marvious carbon source.
⁴Values represent nanomoles of $C_2H_4/h/c$ ulture and the means of three replicate cultures; - Negative activity ϵ ^T

Type strain B. vietnamiensis LMG10929

Cultivars	ARA at panicle initiation ^a	MPN counts $(\times 10^6)^b$	ARA at grain filling ^a	MPN counts $(x10^6)^b$
$ADT-43$	17.36 ± 3.1	2.5 ± 0.5	66.43 ± 9.2	4.5 ± 1.4
ADT-37	15.26 ± 2.6	1.2 ± 0.1	52.45 ± 6.4	2.5 ± 1.0
$Co-47$	13.28 ± 2.3	1.5 ± 0.3	58.53 ± 8.6	1.5 ± 0.2
Ponni	24.32 ± 4.0	4.5 ± 1.2	95.51 ± 13.1	11.5 ± 2.1

Table 2. Detection of AR activity and bacterial populations associated with roots at pannicle initiation and grain filling stage of rice varieties ADT-43, ADT-37, Co-47, and Ponni

^anmol C₂H₄/g dry weight/day

 b AR activity based MPN counts

followed by 46.8, 21.6, 39.9, 90.4, 19.2, and 20.4% (mean 39.71%) at 110 days after inoculation, and root biomass over control was 130.0, 28.6, 88.2, 34.8, 50.0, and 166.7% (mean 83.05%) at 60 days after inoculation followed by 90.9, 98.6, 69.2, 95.2, 51.0, and 110.5% (mean 85.9%) at 110 days after inoculation. Number of tiller increases over control was 49.7, 51.0, 45.6, 29.6, 47.2, and 93.1 (mean 52.7%) at 60 days after inoculation followed by 39.1, 65.4, 57.9, 35.6, 53.4, and 94.6 (mean 57.7%) at 110 days after inoculation. Yield increase over control was 10.1, 13.2, 9.5, 23.6, 14.4, and 15.80% (mean 14.4%) in the two pot and four field experiments respectively (Tables [5,](#page-9-0) [6,](#page-9-0) [7,](#page-10-0) [8,](#page-10-0) [9,](#page-11-0) [10\)](#page-11-0).

Inoculation of B. vietnamiensis MGK3 alone produced a significant increase over the control for all measured parameters. Increases of shoot biomass over control was 39.6, 14.3, 32.4, 25.0, 13.3, and 22.6% (mean 24.53%) at 60 days after inoculation followed by 23.4, 8.4, 21.0, 8.9, 14.5, and 11.8 (mean 14.66%) at 110 days after inoculation, and root biomass over control was 72.8, 12.7, 8.8, 25.8, 36.6, and 133.3% (mean 48.3%) at 60 days after inoculation followed by 36.4, 20.0, 30.7, 9.5, 36.7, and 57.9 (mean 31.86%) at 110 days after inoculation. Tiller increase over control was 36.5, 25.9, 36.5, 19.0, 25.5, and 27.3 (mean 28.5%) at 60 days after

inoculation followed by 25.8, 43.8, 44.0, 35.9, 46.0, and 33.8 (mean 38.2%) at 110 days after inoculation. Yield increase over control was 8.2, 12.8, 5.6, 10.7, 6.7, and 12.16% (mean 9.36%) in the two pot and four field experiments, respectively (Tables [5](#page-9-0), [6](#page-9-0), [7,](#page-10-0) [8](#page-10-0), [9,](#page-11-0) [10\)](#page-11-0). B. vietnamiensis LMG10929 gave a significant effect on growth parameters as compared to the uninoculated control except in the field trial with cv. Ponni at 150 days (Table [10\)](#page-11-0). It had little or no effect on N leaf content. Yield increase over control was 3.7, 2.8, 3.9, 8.9, 4.2, and 8.9% (mean 5.4%) in the two pot and four field experiments, respectively (Tables [5](#page-9-0), [6](#page-9-0), [7](#page-10-0), [8](#page-10-0), [9](#page-11-0), [10](#page-11-0)).

A. lipoferum LMG4348 gave a significant effect on growth parameters as compared to the uninoculated control except for tiller numbers and root weight in the first two field experiments (Tables [7](#page-10-0) and [8](#page-10-0)). It had little effect on N leaf content. Yield increase over control was 3.7, 7.5, 4.4, 11.7, 3.7, and 10.05% (mean 6.8%) in the two pot and four field experiments, respectively (Tables [5](#page-9-0), [6,](#page-9-0) [7](#page-10-0), [8,](#page-10-0) [9](#page-11-0), [10\)](#page-11-0). Its overall effect was higher than B. vietnamiensis LMG10929. H. seropedicae LMG6513 sometimes gave a significant effect on growth parameters as compared to the uninoculated control in three field experiments (Tables [8,](#page-10-0) [9,](#page-11-0) [10\)](#page-11-0). Nevertheless, yield increase over control was very weak: 3.1, 0.8, 2.2, 0.8, 2.1, and 6.8%

Fig. 2 X-Gluc stained rice roots inoculated with MGK3 tagged with the gusA gene. A Intense colonization at lateral root emergence (100 X); B-glucuronidase activity is revealed in situ by the blue color $(200\times)$. Bars: 100 µm.

Table 3. Root surface colonization by isolate MGK3, expressed rable 5. Noot surface colonization by isolate MON.5, expresse
as colony-forming U g^{-1} fresh weight in four varieties of rice

Days after inoculation	$ADT-43$	$ADT-37$	$Co-47$	Ponni
	2.5×10^{3}	2.5×10^{3}	1.5×10^{2}	1.5×10^{4}
5	1.5×10^{6}	4.5×10^{5}	1.5×10^{5}	2.5×10^{7}
10	1.5×10^{7}	1.5×10^{7}	1.5×10^{7}	1.5×10^{8}
15	1.5×10^{8}	1.5×10^{8}	1.5×10^{8}	1.5×10^{9}

All values are expressed as means derived from triplicate experiments.

(mean 2.6%) in the two pot and four field, experiments respectively. G. diazotrophicus LMG7603 gave a significant effect on growth parameters as compared to the uninoculated control for all vegetative parameters. The effect on N leaf content was inconsistent. Yield increase over control was 6.3, 10.2, 3.3, 7.0, 2.7, and 10.05% (mean 6.6%) in the two pot and four field experiments, respectively (Tables [5,](#page-9-0) [6](#page-9-0), [7](#page-10-0), [8,](#page-10-0) [9,](#page-11-0) [10\)](#page-11-0). Shoot is to root ratios were significantly varied among the cultivars used in pot and field experiments. Cultivar ADT-43 produced shoot is to root ratio almost 10:1 (Tables [5](#page-9-0) and [7\)](#page-10-0), and cultivar Ponni ratio was 3:1 (Tables [6](#page-9-0) and [10\)](#page-11-0) in both pot and field experiments. Field experiments with cultivar Co-47 produced shoot is to root ratio nearly 7:1 (Table [8](#page-10-0)) and cultivar ADT-37 shoot is to root ratio was 3:1 (Table [9\)](#page-11-0).

The weight of 1,000 grains is a late yield component. It is increased by all inoculants (Table [11\)](#page-12-0). The combined inoculum ranks first, followed by B. vietnamiensis MGK3; the controls consistently ranked last, just after H. seropedicae LMG6513, A. lipoferum 4B LMG4348, G. diazotrophicus LMG7603, and B. vietnamiensis LMG10929 rank between MGK3 and H. seropedicae LMG6513. In terms of overall effect on yield, treatments can be ranked using the mean effect on yield in the six experiments: the combined inoculum ranks first (14.4% increase over control), followed by B. vietnamiensis MGK3 (9.36%), A. lipoferum 4B LMG4348 (6.8%), G. diazotrophicus LMG7603 (6.6%), B. vietnamiensis LMG10929 (5.4%), and H. seropedicae LMG6513 (2.6%).

Discussion

The present study was undertaken to isolate and enumerate new nitrogen-fixing Burkholderia endophytes from different rice cultivars of Tamilnadu State, India. This concept of Biological Nitrogen Fixation by endo-phytes has been introduced by Döbereiner [[19\]](#page-14-0) and mostly tested with graminaceous plants. Boddey and Döbereiner [\[10\]](#page-14-0) suggested that endophytic bacteria better express their nitrogen fixation potential inside plant tissues due to the lower competition for nutrients and protection against high levels of $O₂$ present on the root surface. This habitat has already been identified as an important source of endophytic PGPR that do not induce obvious symptoms of disease. Examples include Azospirillum strains isolated from "inside" host roots (after surface sterilization), which increase yield when inoculated on that homologous host [[9](#page-14-0)], the diazotrophic endophytes of Azoarcus inside Kallar grass [[7,](#page-14-0) [33](#page-15-0)], and Acetobacter diazotrophicus inside sugar cane [\[18](#page-14-0)]. Nature appears to select endophytes that are competitively fit to occupy compatible niches within this nutritionally enriched and protected habitat of the root interior without causing pathological stress on the host plant. Furthermore, Yanni et al [\[68](#page-16-0)] reported that field inoculation with the rhizobial endo-colonizer did not qualitatively alter rice grain protein composition and that nutritionally important proteins are all present in treated and control samples in similar ratios.

Identification and Growth-Promoting Properties. From surface-sterilized plant tissues, we obtained 13 isolates

	Pot grown plants			Field grown plants		
Treatments	Plant hiomass Root area (cm^2) (mg/plant)		Percent Ndfa $(%)^a$	Root area $(cm2)$	Plant biomass (mg/plant)	Percent Ndfa (%)
Cultivar-Ponni						
$Control^b$ B. vietnamiensis MGK3 B. vietnamiensis LMG10929 Cultivar-ADT-43	36.12 c 58.43 a 53.48 a	321c 456 a 424a	- 40.4 36.8	41.32 c 62.45a 58.63 a	363c 478 a 452 h	42.2 39.6
$Control^b$ B. vietnamiensis MGK3 B. vietnamiensis LMG10929	28.52c 42.12a 39.45 a	256c 378 a 359 a	36.4 35.6	36.42c 53.23 a 47.25a	264c 483 a 462a	39.2 36.6

Table 4. Effect of B. vietnamiensis strain MGK3 on root area, plant biomass, and nitrogen fixation of rice varieties: Ponni and ADT-43

Plants were grown under microbiologically controlled conditions in sterile vermiculite for 4 weeks. Eight plants were used for each treatment. a Percentage of nitrogen derived from air. Means followed by the same letter are not statistically different at 5% level according to Duncan's

Multiple Range Test (DMRT). ^b

Plants inoculated with autoclaved cells

Table 5. Pot trial with rice cultivar ADT-43

Table 5. Pot trial with rice cultivar ADT-43

yield. Ten replicates. Different letters indicate treatments differing at the 5% level (Tukey test). Control: seedlings inoculated with autoclaved cells "Mixture of G. diazo*trophicus*", *H. seropedicae*", A. lipoferum, B. , H. seropedicae T , A. lipoferum, B. vietnamiensis^T, MGK3

Effect of inoculation on different parameters: shoot height (cm), shoot and root weights (g/plant), tiller numbers per plant. Effect of inoculation on leaf N content (at days 60 and 150) and final grain Effect of inoculation on different parameters: shoot height (cm), shoot and root weights (g/plant), tiller numbers per plant. Effect of inoculation on leaf N content (at days 60 and 150) and final grain 6.44 4.47 Coeff. of variation

F-value 184.07 156.20 51.61 20.56 131.64 318.34 1699.26 20.0 5.59 1.59 – – Coeff. of variation 3.56 3.56 8.93 8.93 18.16 4.47 6.44 2.6.57 2.130 7.95 – –

yield; ten replicates
"Mixture of G. diazotrophicus^T $H.$ seropedicae^T A. lipoferum, B. vietnamiensisT, MGK3

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Table 7. Paddy field trial with rice cultivar ADT-43 at Edaiyathur

Table 7. Paddy field trial with rice cultivar ADT-43 at Edaiyathur

and 110) and final grain yield; 25 replicates
"Mixture of G. diazotrophicus", H. serope

, H. seropedicae T , A. lipoferum, B. vietnamiensis^T, MGK3

Table 8. Paddy field trial with cultivar Co-47 at Kavithandalam Table 8. Paddy field trial with cultivar Co-47 at Kavithandalam days 60 and 110) and final grain yield; 25 replicates ^aMixture of G. diazotrophicus^T, H. seropedicae^T, A $H.$ seropedicae T A. lipoferum, B. vietnamiensisT, MGK3

Table 9. Paddy field trial with cultivar ADT-37 at Orakatpet Table 9. Paddy field trial with cultivar ADT-37 at Orakatpet

(at days 60 and 110) and final grain yield; 25 replicates a Mixture of G. diazotrophicus^T, H. seropedicae^T, A. li Effect of seed moculation on unitertit parameters, surve use the section \cdots is \cdots in the set of and 110) and final grain yield; 25 replicates $\binom{m}{k}$ internamients $\binom{m}{k}$, MGK3 , H. seropedicae^T, A. lipoferum, B. vietnamiensis^T, MGK3

days 60, 110, and 150) and final grain yield; 25 replicates aMixture of G. diazotrophicus^T, H. seropedicae^T, A. lipo $H.$ seropedicae T A. lipoferum, B. vietnamiensisT, MGK3

	1,000 grain weight (g)							
<i>Treatments</i>	Pot trial-1 $(ADT-43)$	Pot trial-II (Ponni)	Field trial—I (ADT-43)	Field trial—II $(Co-47)$	Field trial—III (ADT-37)	Field trial—IV (Ponni)		
Control	15.80	17.45	17.80	19.70	20.80	18.91		
G. diazotrophicus	16.82	19.24	18.41	21.55	21.38	20.82		
H. seropedicae	16.34	17.60	18.23	20.30	21.24	20.21		
A. lipoferum	16.41	18.76	18.65	23.70	21.58	20.83		
B. vietnamiensis	16.42	17.95	18.53	21.93	21.68	20.61		
MGK3	17.11	19.70	18.82	22.60	22.20	21.20		
Combination ^a	17.43	19.76	19.5	24.90	23.80	21.91		

Table 11. Effect of inoculation on the weight of 1,000 grains

^aMixture of G. diazotrophicus^T, H. seropedicae^T, A. lipoferum, B. vietnamiensis^T, MGK3

able to grow on PCAT. In planta densities were diverse; the most active strain was MGK3. Isolate MGK3 grew well on different carbon sources. In particular, it grew on sorbitol, mannose, and mannitol which differentiate Burkholderia spp. from the genus Ralstonia [[26\]](#page-14-0). Their 16S rDNA sequences and their fatty acid profiles placed them in the species B. vietnamiensis. Isolate MGK3 showed positive for acetylene reduction, and PCR amplification of nif-H confirmed the presence of this structural nitrogenase gene in isolate MGK3.

We have found large differences in nitrogenase activity when isolate grown in different carbon sources (Table [1\)](#page-6-0) and also rice growth stages (Table [2](#page-7-0)). Similar variation in AR activity with the plant growth stage has been reported by Watanabe et al. [\[65](#page-16-0)] in two rice varieties, IR36 and IR26, where maximum AR activity was detected at the grain-filling stage. Higher AR activity at a particular growth stage may be due to a reduction in inhibitory nitrogen concentrations in the soil or overproduction of root exudates that are conducive to diazotroph growth and activity [[35\]](#page-15-0).

The root surface of rice was intensively colonized at the end of 24 h of inoculation (Fig. [2](#page-7-0)A) and colonized many of the cortical intercellular spaces after 15 days of inoculation (Fig. [2B](#page-7-0)). Primer specific PCR analysis of the gusA gene revealed a fragment of 1,200 bp in the transconjugant [[39\]](#page-15-0). The xylem could be a suitable nonnodular niche for N_2 fixation because it could provide the low pO2 required for the expression and function of nitrogenase and also allow the exchange of fixed N_2 [\[37](#page-15-0)]. Nitrogen fixation by ¹⁵N isotopic dilution method showed that the effect of B. vietnamiensis MGK3 inoculation was higher in cultivar Ponni than in ADT-43, where nearly 42% (field) and 40% (pot) of the nitrogen was derived from atmosphere (% Ndfa), respectively. In both cultivars, native isolate B. vietnamiensis MGK3 showed increased fixation than type strain B. vietnamiensis LMG10929 (Table [4](#page-8-0)).

Rice Inoculation Experiments. Two pot experiments followed by four field experiments were performed at four

different places using four commercially important rice cultivars (ADT-43, Co-47, ADT-37, and Ponni) to evaluate and compare isolate MGK3 to: (1) the type strain of B. vietnamiensis LMG10929, (2) other rice diazotrophs (H. seropedicae LMG6513, A. lipoferum LMG4348), (3) a sugarcane diazotroph (G. diazotrophicus LMG7603), and (4) a mixture of all these strains. In terms of overall effect on yield, the ranking is (1) combined inoculum (2) B. vietnamiensis MGK3, (3) A. lipoferum 4B LMG4348, (4) G. diazotrophicus LMG7603, (5) B. vietnamiensis LMG10929, and (6) H. seropedicae LMG6513. Significant increases of shoot and root biomass due to combined inoculation reached 38.4 and 83.0% (mean of two pot and four field experiments), respectively, at 60 days after inoculation. This is indicative of a very early effect of bacteria on their host plant. Jacoud et al. [\[34](#page-15-0)] reported that a very short and early contact of maize with PGPR bacteria was enough to ensure a significant effect on subsequent growth. Tillering is the next important step in rice development, and the tiller number per plant was significantly increased (+28%, mean of two pot and four field experiments) 60 days after combined inoculation. Tran Van et al. [[62\]](#page-15-0) noticed a similar effect when B. vietnamiensis was inoculated with rice. Root biomass was drastically low compared to shoot biomass among different pot and field experiments; this variation may be difficult to extract all roots in pot and field experiments. The effect on the weight of 1,000 grains somewhat parallels the effect on final yield, suggesting that the effect of inoculated bacteria persists throughout the plant growth cycle.

Some of these strains had already been assayed in the field. B. vietnamiensis LMG10929, for instance, has been inoculated to rice in Vietnam [\[62](#page-15-0)] in three pot and four field experiments in three different locations. Yield increases caused by inoculation ranged from 13 to 22% over control, larger than what was observed in Tamilnadu. In our experiment, B. vietnamiensis LMG10929 gave a significant effect on growth parameters as compared to the uninoculated control except in the field trial with cv. Ponni at the 150th day (Table [9\)](#page-11-0). Yield increase over control was the mean of 5.4% in the two pot and four

field experiments. This strain is not an Indian but a Vietnamese isolate [\[59](#page-15-0)] obtained from an acid sulfate soil, very different from soils used in the present paper. The ability of B. vietnamiensis LMG4348 to internally colonize rice tissues is unknown; however, this species has been recovered from internal tissues of maize [\[22](#page-14-0)].

G. diazotrophicus LMG7603 gave a significant effect on growth parameters as compared to the uninoculated control for all vegetative parameters. Yield increase over control was mean of 6.6% in the two pot and four field experiments. G. diazotrophicus LMG7603 has proved beneficial, where the rice seedlings inoculated with the G. diazotrophicus grew to be significantly taller 30 days after inoculation than plants inoculated with the nif- mutant or uninoculated plants under N-deficient conditions [\[56](#page-15-0)]. Inoculation of G. diazotrophicus LMG7603 to sugarcane has been proven beneficial, where the plant height [\[55\]](#page-15-0) and yield [\[46\]](#page-15-0) of the inoculated plants were higher than the control. Field trials conducted in the sugarcane system revealed the usefulness of G. diazotrophicus LMG7603 with other diazotrophs, which have contributed to the yield equal to that of the control $(280 \text{ kg N} \text{ ha}^{-1})$. Mixed inoculation of vesicular–arbuscular mycorrhizal (VAM) spores and G. diazotrophicus LMG7603 also proved beneficial in improving the yield of different sugarcane varieties. The yield was also not reduced even under 50– 100% reduction from the recommended dose of chemical N compared to the control, attributing the role of inoculated G. diazotrophicus LMG7603 in N contribution [\[46\]](#page-15-0). It has been reported that inoculation of micropropagated sugarcane seedlings would make the plants not only grow faster, but also ensure efficient N-fixing plants in fields.

A. lipoferum LMG4348 gave a significant effect on growth parameters as compared to the uninoculated control except for tiller numbers and root weight in the first two field experiments (Tables [7](#page-10-0) and [8\)](#page-10-0). Yield increase over control was the mean of 6.8% in the two pot and four field experiments. Strain 4B of A. lipoferum LMG4348 has been isolated from a French rice field [\[59](#page-15-0)], proved to be a very efficient diazotroph under gnotobiotic conditions [[30\]](#page-15-0), and for that reason, it was used for three field inoculation experiments [\[14\]](#page-14-0). Yield increases following inoculation were 20.7, 21.3, and 15.9% over uninoculated controls. Its rather poor performance in the Indian context is surprising. H. seropedicae LMG6513 sometimes gave a significant effect on growth parameters as compared to the uninoculated control especially in three experiments (Tables [8,](#page-10-0) [9,](#page-11-0) [10\)](#page-11-0). Nevertheless, yield increase over control was the mean of 2.6% in the two pot and four field experiments. The less performance of H. seropedicae LMG6513 is also surprising because this particular strain was isolated from rice and aggressively colonizes rice tissues [\[35](#page-15-0)] and stimulates rice growth in gnotobiotic conditions [[5\]](#page-14-0). This bacterial species is often found in rice and its wild counterparts

[\[20\]](#page-14-0). There is no explanation for its poor efficiency under Indian conditions. Baldani et al. [\[4](#page-14-0)] reported inoculation of rice under gnotobiotic conditions with Herbaspirillum seropedicae LMG6513, B. "brasilensis" and B. vietnamiensis LMG10929. Total plant dry weight increases were observed after the inoculation viz: 42% for B. vietnamiensis LMG10929, 64% for B. "brasilensis" and 71.5% for H. seropedicae LMG6513 (59% for strain LMG6513). In pots on 4-month-old plants, these figures were 48 and 27% for H. seropedicae LMG6513 and B. "brasilensis", respectively, confirming the former as the best inoculum. This difficulty in comparing inoculation experiments is further increased by the use of different cultivars in different parts of the world. The use of endophytic diazotrophs as a substitute for nitrogen fertilizers [\[1](#page-14-0)] is still a debatable matter. The beneficial results reported in the present article are only one more argument in favor of this new practice. Other arguments are: low cost, absence of pollution, similarity to the natural process, and foreseeable sustainability.

The mixed inoculant, in the present work, displays an important potential in spite of the poor performance of some of its components. Muthukumarasamy et al. [\[46\]](#page-15-0) inoculated a mixture of diazotrophs and mycorrhizal fungi to micropropagated sugarcane and obtained an effect equivalent to half the recommended rate of nitrogen fertilizers under pot conditions. Oliveira et al. [[47\]](#page-15-0) also reported a pot experiment in which the combined inoculation of five strains (G. diazotrophicus LMG7603, H. seropedicae LMG6513, H. rubrisubalbicans, A. amazonense, and Burkholderia sp.) gave higher contribution to plant growth, followed by the treatment with a mixture of Herbaspirillum spp. However, the contribution was much lower when the plants were inoculated with a mixture of G. diazotrophicus LMG7603 with A. amazonense and Burkholderia sp. Govindarajan et al. [\[28\]](#page-15-0) reported that individual inoculation B. vietnamiensis MG43 and G. diazotrophicus LMG7603 gave a higher sugarcane yield as compared to combined inoculation. One can speculate that the different bacteria occupy different niches in the plant, replacing expected competition by a cooperative effect, as proposed by Ueda et al. [[63\]](#page-15-0). Next to the mixed inoculum, strain MGK3 ranks first among single inoculum assays and is a good candidate for further inoculation trials on a larger scale.

Nevertheless, due to its relatedness to Burkholderia species (B. cenocepacia, B. multivorans, and B. stabilis) causing severe pneumonia in cystic fibrosis-affected people, a prerequisite [\[48](#page-15-0)] will be to demonstrate that MGK3 is devoid of any pathogenesis genes and is unable to acquire them through horizontal transfer. Several human pathogenic Burkholderia species (including B. cenocepacia, B. pseudomallei, and B. mallei) have been completely sequenced and other environmental species sequencing projects are in progress (B. vietnamiensis G4,

B. phytofirmans PsJN, and B. phymatum STM815). It can be expected that knowledge derived from these genome sequencing projects will allow us to gain further insights into functional diversity, evolution, and pathogenicity mechanism. This may provide the scientific basis for important decisions regarding the biotechnological use of Burkholderia species.

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