



## ISOLATION AND CHARACTERIZATION OF PHOSPHATE SOLUBLIZING BACTERIA FROM ANAND AGRICULTURE SOIL.

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### ABSTRACT

Phosphorus is vital to seed formation and its content is higher in seeds than in any other part of the plant. It helps plants to survive winter rigors and also contributes to disease resistance in some plants. Also known to improve quality of many fruits, vegetables and grain crops. The inoculation of P-solubilizing microorganisms is a promising technique because it can increase P availability in soils fertilized with rock phosphates. In the present work we have made efforts to isolate the phosphate solubilizing microorganism from soil followed by its biochemical and molecular characterization

**Key words:** Phosphorus, PVK media, halo zone, 16S rRNA amplification

### INTRODUCTION

Phosphorus plays an indispensable biochemical role in photosynthesis, respiration, energy storage and transfer, cell division, cell enlargement and several other processes in the living plant. An adequate supply of phosphorus in the early stages of plant growth promotes physiological functions including early root formation, and is important for laying down the primordia for reproductive parts of plants. It is vital to seed formation and its content is higher in seeds than in any other part of the plant. It helps plants to survive winter rigors and also contributes to disease resistance in some plants. Also known to improve quality of many fruits, vegetables and grain crops. Biological Nitrogen Fixation depends appreciably on the available forms of phosphorus. Phosphorus (P) is an important structural constituent of nucleic acids, phytin and phospholipids. Mineral forms of phosphorus constitute the biggest reservoirs of phosphorus, represented primarily by rocks and deposits formed

during geological age. The principal characteristic of these primary minerals (oxyapatite, hydroxyapatite, apatite) is their insolubility. A large portion of soluble inorganic phosphate applied to agricultural soil as chemical fertilizer is rapidly immobilized soon after application and becomes unavailable to plants (Dadarwalet *et al.*, 1997).

PSMs include different groups of microorganisms, which not only assimilate phosphorus from insoluble forms of phosphates, but they also cause a large portion of soluble phosphates to be released in quantities in excess of their requirements. Species of *Aspergillus* and *Penicillium* are among fungal isolates identified to have phosphate solubilizing capabilities. Among the bacterial genera with this capability are *Pseudomonas*, *Azospirillum*, *Bacillus*, *Rhizobium*, *Burkholderia*, *Arthrobacter*, *Alcaligenes*, *Serratia*, *Enterobacter*, *Acinetobacter*, *Flavobacterium* and *Erwinia* (Rodriguez *et al.*, 1996). Seed or soil inoculation with PSMs is known to improve

solubilization of fixed soil phosphorus and applied phosphates resulting in higher crop yields (Jones et al., 1994) PSMs are a low-cost solution that enriches the soil giving a thrust to economic development without disturbing ecological balance. The inoculation of P-solubilizing microorganisms is a promising technique because it can increase P availability in soils fertilized with rock phosphates [Reyes et al. (2002)]. Several authors reported yield increasing on wheat [Whitelaw et al (1997)], onion [Vassilev et. al (1997)], alfalfa [Rodríguez 1999] and soybean [Abd-Alla et. al (2001)] through simple inoculation of P-solubilizing fungi (PSF). Inoculation of phosphate solubilizing fungi and mycorrhizal fungi improves the physio-chemical, biochemical and biological properties of rock-P amended soil [Caravaca, F et .al (2004)].

## MATERIALS AND METHODS

### *Sample collection*

Total of 8 samples were collected from different ecological sources in sterilized bottles. Ecological sources included were mainly agriculture land.

### *Isolation of phosphate solubilizing Bacteria from soil sample*

#### *Preparation of media for isolation of PSM*

Readymade formulation of dehydrated media from the makers of 'HI MEDIA' was used for preparing solid as well as liquid media. 9 cm diameter petri plates were used for routine culture of bacteria.

#### *Preparation of inoculums from soil sample and inoculation*

Acidic soil sample from nearby village of Anand, (Gujarat) was collected and dissolved in sterile distilled water and then serially diluted up to  $10^{-7}$ . Then after 100  $\mu$ l of  $10^{-7}$  serially diluted sample spread on PVK agar and the plates were incubated at 37°C for 24 hrs. After incubation the bacterial colonies having clear zone of phosphate solubilization were selected and streaked on new PVK agar plates to get isolated pure colonies. Further purification was carried out on the same media.

### *Microscopic study of Bacteria*

Size, shape, arrangement and gram's nature of the isolates were studied. For grams staining smear was prepared from the isolated culture on clean grease free microscopic glass slide, heat fixed and stained. The stained smear was observed under microscope (oil immersion lance-100X).

### *Identification of Bacterial isolates through Biochemical tests.*

Biochemical test were performed as suggested by Garrity et al., 2004, which include following tests Gram staining, IMVIC reaction, Catalase test, Starch hydrolysis test, Oxidation fermentation test, Phenyl alanine deamination test, Lead acetate paper strip test, Ammonia production test, Nitrate reduction test, Gelatin hydrolysis test, Urea hydrolysis test, Dehydrogenase test, Casein hydrolysis test, Citrate utilization test, Indole production test, Triple sugar iron (TSI) test, Carbohydrate fermentation test (Glucose, Fructose, Xylose, and Sucrose), Motility test, Endospore staining and Capsule staining.

### *Molecular characterization of Bacterial isolates*

#### *Isolation of genomic DNA*

Pellet of bacterial cell was suspended in 2 ml Tris-EDTA (pH 8.0), 250  $\mu$ l 10% SDS, 50  $\mu$ l Lysozyme (20 mg/ml), and incubated at 37°C for 1 hr. then 500  $\mu$ l 5M NaCl and 100  $\mu$ l CTAB were added and incubated for 10 min at 65°C. Then phenol, chloroform and isoamyl alcohol were added and centrifuged for 10 minute. Supernatant was extracted with chloroform:isoamyl alcohol and finally precipitated with 1/10 volume of 7.5 M Ammonium acetate (pH-7) and double the volume of chilled ethanol. Pallet was suspended in 30 ml of sterile water.

#### *Amplification and automated fluorescent nucleotide sequencing of the 16S ribosomal DNA fragment*

Bacterial ribosomal gene analysis was performed as described by Weisberg et al. (1991), with minor modifications. The PCR products were purified by using a nucleotide removal kit (Qiagen, Germany)

and the sequencing products were purified according to the manufacturer's protocol. Base calling was performed with Applied Biosystems Sequence Analysis Software (version 5.2) and quality value bar above 40 was considered to be acceptable for further analysis DNA sequence similarity search

was performed using the BLAST algorithm (Altschul et al. 1994, 1997).

#### ***16srRNA PCR of isolated bacterial DNA***

100 ng of the genomic DNA was used for performing 16s PCR and the composition of reaction mixture is a below

Components	Quantity (μl)
2 X PCR Master Mix	15.0
Forward Primer (10pmole/μl)	1.0
Reverse Primer(10pmole/μl)	1.0
Nuclease Free Water	9.0
DNA Template	4.0
Grand Total	30.0

Primers (forward reverse) and	Cycling conditions				
	Initial denaturation	Denaturation	Annealing	Extension	Final extension
16S rRNA	94oC 3min	94oC 40sec	55oC 45sec	72oC 75sec	72oC 3min

#### ***Agarose Gel Electrophoresis***

To confirm the targeted PCR amplification, 8μl of PCR product from each tube was mixed with 2μl of 6X gel loading buffer from each tube and electrophoresed on 1.0 % agarose gel along with 100 bp DNA Ladder (Bangalore genie) and stained with ethidium bromide at constant 80 V for 45 minutes in 0.5X TBE buffer. The amplified product was visualized as a single compact band of expected size under UV light and documented by gel documentation system (Alpha imager, Alpha innotech). .

#### ***ARDRA***

For ARDRA 16S rDNA was amplified and restricted digested by the enzyme ECO-RV. Steps involved in this method are amplification of 16S r-DNA was performed using 30ng DNA in 25μL of total volume reaction. A 10μL aliquot of each PCR reaction was incubated with 1-2 units of one restriction enzyme and the mixture was incubated at

370C for 3 hours. The restriction products were then analyzed by electrophoresis on 1.5% agarose in 1X-TBE buffer at 100 V and the gel was visualized in gel documentation system (Alpha Innotech).

#### ***SDS-PAGE:-***

##### ***Gel preparation and sample preparation:***

A 10% resolving gel and a 5% stacking gel was formed in the Biorad SDS –PAGE unit. 2 X SDS-PAGE sample buffer was added to the loosened pellet, the protein lysate so obtained was centrifuged at 12,000 rpm for 12 minutes, supernatant was collected in a new vial and heated at 950C for two minutes followed by cooling of the samples at room temperature and giving a short spin, the samples thus prepared were then loaded in the gel along with a mid-range protein marker of 14KDa-95KDa (SRL BIOLITTM).The samples were allowed to run at 100 V for 5 hours followed by overnight staining of the gel in 1% Commassie

Brilliant Blue. The destaining of gel was carried out the next day and the bands were visualized and documented by gel documentation system (Alpha imager, Alpha innotech).

## RESULT

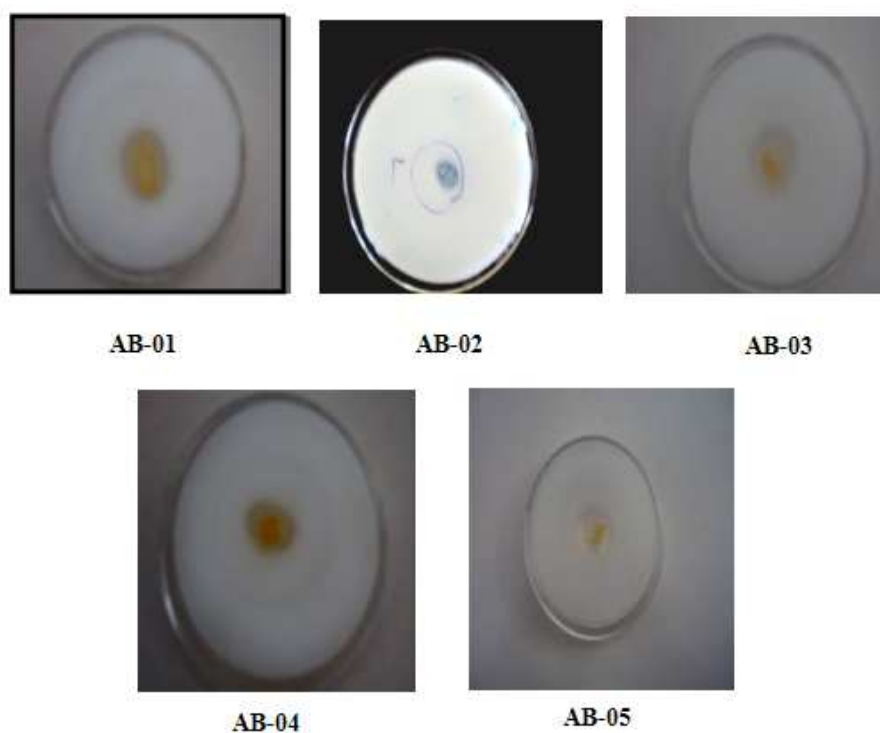
### *Screening of Ecological Sources:*

Location of soil sample collection was chosen because of the possibility of occurrence of phosphate solubilizing microbes. Sampling was done at various sites, in order to maintain uniform representation of the micro flora in and around the collection area. For initial growth of micro flora PVK medium was used embedded with tricalcium phosphate as phosphate source. Preliminary

investigation of these cultures in PVK media embedded with tricalcium together with agitation and aeration for 5 day allowed microbial solubilization of P with fall in pH. This was followed by the dilution plating in order to isolate the single colonies. The concentration of 1.5% agar adequately maintained the desired texture of the solid medium, while simultaneously retained enough moisture to promote microbial growth.

Its been reported that organisms capable of doing phosphate solubilization give clear zone around the colony by which it can be concluded that they are phosphate solubilizing microorganisms. Five colonies gave clear zone by which it was confer that they can solubilize phosphate. **Fig 1.**

Sr.No	Biochemical Test	AB- 01	AB-02	AB-03	AB-04	AB-05
1.	Grams Staining	+	-	+	+	-
2.	Capsule staining	+	-	+	+	-
3.	Motility	-	-	-	+	-
4.	Methyl Read Test	+	-	+	+	-
5.	VogesProskau (VP) Test	+	-	+	+	-
6.	Catalase Test	-	+	-	-	+
7.	Starch Hydrolysis Test	-	-	-	+	-
8.	Oxidation	+	+	-	-	+
	Fermentation Test	+	-	-	-	+
9.	Phenylalanine deamination test	+	-	-	+	+
10.	Lead acetate paper strip test	-	-	+	+	-
11.	Ammonium production test	+	-	+	-	+
12.	Nitrate reduction test	+	+	+	-	+
13.	Gelatin hydrolysis test	-	+	-	-	-
14.	Urea hydrolysis test	-	+	-	-	-
15.	Dehydrogenase test	+	-	-	-	+
16.	Casein hydrolysis test	-	-	-	+	+
17.	Citrate utilization test	+	-	+	-	+
18.	Indole production test	-	+	-	-	+
19.	TSI test(Triple sugar test)	+	-	-	+	+
20.	Carbohydrate fermentation test					
	Glucose					
	Fructose	+	+	+	+	+
	Xylose	+	+	+	+	+
	Sucrose	-	-	+	+	-



**Figure 1: Halo Zone around the colony on PVK media confirm Phosphate solubilizing bacteria**

**Identification of Bacterial isolates through Biochemical tests:**

All the strain was subjected to the biochemical test for their identification. The results are summarized in **Table 2**.

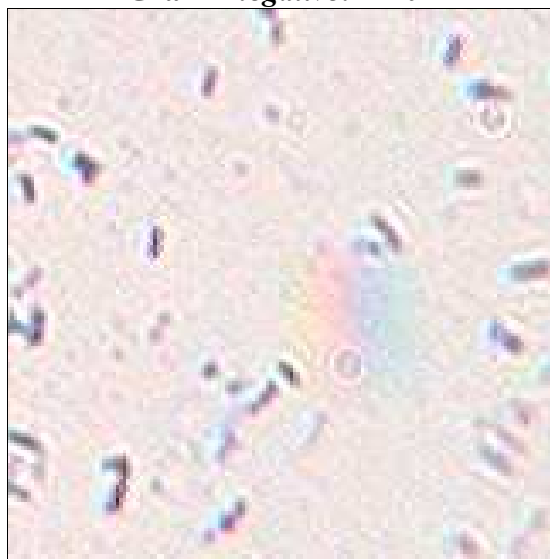
**Table- Colony characteristics of the individual isolates**

**Table- Biochemical characterization of the isolates**

Test	AB-01	AB-02	AB-03	AB-04	AB05
Size	Small	Large	Big	Small	Large
Shape	Round	Round	Round	Punctiform	Round
Margin	Entire	Entire	Entire	Entire	Entire
Elevation	Raised	Flat	Raised	Flat	Flat
Surface texture	Smooth	Smooth	Smooth	Smooth	Contoured
Consistency	Moist/Butyrous	Moist/ Butyrous	Moist/ Butyrous	Moist/ Butyrous	Viscous
Optical Character	Translucent	Translucent	Opaque	Translucent	Translucent
Pigmentation	Orangish	White	Yellowish	Yellowish	Yellowish



**Gram –negative:-AB02**



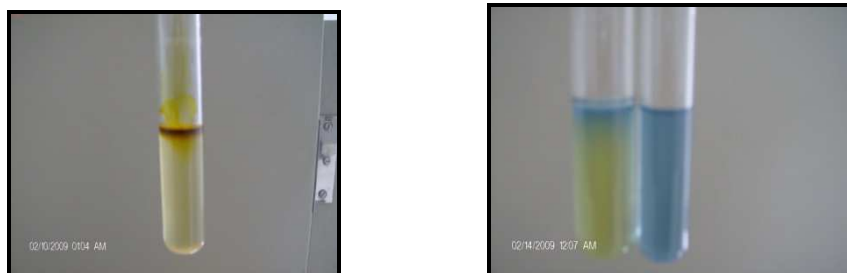
**Gram-positive:-AB01**

**Figure-8:- Gram staining of culture**



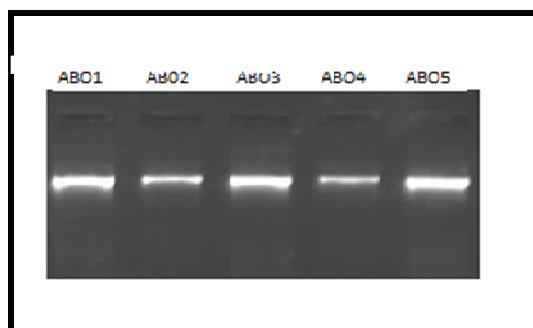
***Positive glucose fermentationPositive citrate utilization***



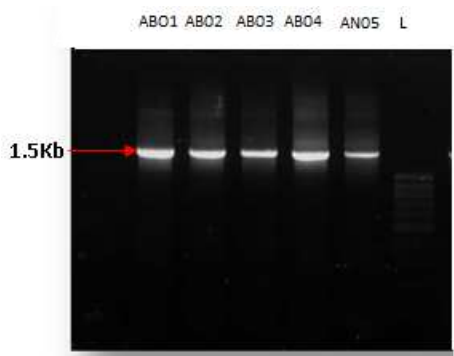
***Casein hydrolysis test******Positive indole test Positive dehydrogenase production test*****Figure-9:- Some of the positive Biochemical Tests.*****Isolation of genomic DNA and Estimation***

DNA of all the strain was isolated and was visualized by running it on 1.0% (w/v) agarose gel. The gel was visualized in UV transilluminator and photographed subsequently. Isolated DNA, after

organic extraction, appeared a thick high molecular weight band with a faint back ground. Optical density was taken at 260/280 nm. All the DNA were found to be pure.

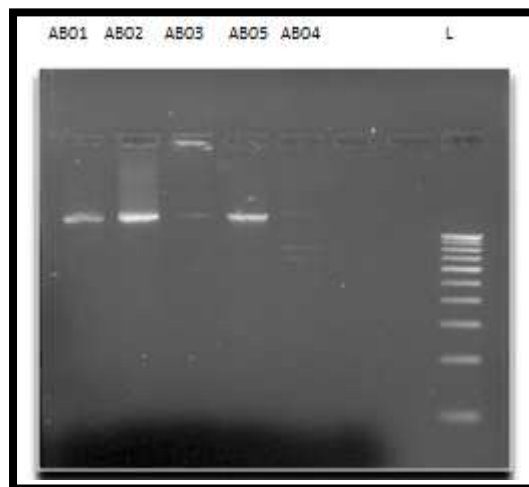
**Figure 10: DNA of AB-01, AB-02, AB-03, AB04 and AB05*****16s rRNA PCR of isolated bacterial DNA and Sequencing***

All the strain gave 1.5 Kb band upon amplifying with 16s rRNA universal primer. Upon sequencing AB02 is *Burkholderia cenocepacia* and AB04 can be a *Bacillus sp.*

**Figure 11: 16s RNA PCR**

***Amplified r-DNA Restriction Analysis (ARDRA):***

16S rPCR product was digested with Eco RV in order to see the polymorphism. Results show that four samples (AN-01, AN-02, AN-03 and AN-05) had no polymorphism while one sample (AN-04) showed different banding pattern. This showed that AN-04 had had Eco RV recognition site in the 16s rRNA amplified product.



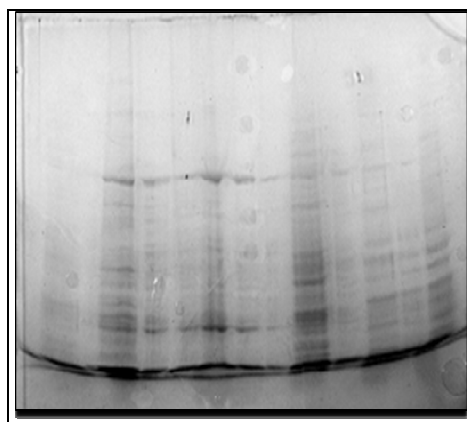
***Figure Eco R V digested 16s RNA PCR product***

***16s RNA sequencing:***

16s rRNA PCR products of all isolates were sent for sequencing. The sequences obtained were studied using BLAST in NCBI database to find out maximum percentage homology with known bacterial DNA sequences to identify the isolates. It was found that AB-02 has 90% homology with *Burkholderia cenocepacia*. While AB-04 had 82% homology with *Bacillus sp.*

***Protein profiling of the Isolates:***

Upon performing protein profiling it was found that all the isolates did not give same banding pattern. There were few bands which were common in isolates but rest showed polymorphism. This results help to interpret that which isolates may be belonging to same group and which are from the divergent group.



***Figure 13: SDS-PAGE of all the isolates***



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