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SIDEROPHORE PRODUCTION FROM *AZOTOBACTER* SP. AND ITS APPLICATION AS BIOCONTROL AGENT

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ABSTRACT

Screening and isolation of siderophore producing bacterial strains were carried out from soil samples collected from plant rhizosphere region. The isolate showed positive for siderophore on Chrome azurol sulfonate agar medium was selected for further studies. The organism was subjected to various biochemical tests and 16S rRNA analysis, which lead to its identification as *Azotobacter* sp. The *Azotobacter* sp. holds optimal growth temperature and pH at 30°C and 8.5 respectively. It was also optimized for carbon and nitrogen with optimal growth in Malic acid and peptone respectively. The presence of magnesium and zinc with iron concentration of 4m concentration also favored the siderophore production. Siderophore production was carried out using optimized media and the harvested siderophore was partially purified by ion exchange chromatography. The antagonist activity of the siderophore producing *Azotobacter* sp. and partially purified siderophore were tested against fungal pathogens such as *Fusarium* sp., *Alternaria* sp., *Phytophthora* sp., *Rhizoctonia* sp., *Colletotrichum* sp. and *Curvularia* sp. This result showed that the *Azotobacter* sp. under study is a good producer of siderophore, which can be beneficial for its antagonistic activity towards fungal pathogens.

Keywords: Chrome azurol sulfonate agar medium, 16S rRNA analysis, Amberlite IR120, Antagonism.

INTRODUCTION

Iron is essential for the growth of almost of the microorganisms. Although iron is abundant in nature, it is not readily available to microorganisms because of its extreme insolubility in aerobic water and soil environments (Neilands, 1972). To sequester iron from the environment, microorganisms excrete high-affinity ferric chelating compounds, termed as siderophores, which solubilize and transport the metal (Neilands, 1974). Nitrogenase and other proteins involved in nitrogen fixation require a high complement of iron (Brill, 1980). Therefore, it is likely that nitrogen fixing organisms evolved efficient iron-acquisition mechanisms. Iron-deficient nitrogen-fixing cultures of *Azotobacter* spp. produce 2,3-dihydroxybenzoic acid (DHB), 2-N,6-N-di-(2,3-

dihydroxybenzoyl)-L-lysine (DHBL), and a yellow-green fluorescent peptide (YGFP) (Carbin and Bulen, 1969). Although, VB the chemical nature of these molecules and their physiological response to iron make them strong candidates for siderophores (Neilands, 1977).

The alternative strategies for disease management include the use of bacteria that show beneficial effects on plants and these bacteria are known as Plant Growth Promoting Rhizobacteria (PGPR). The positive effects of PGPR are normally divided into two categories: growth promotion and biological control (Kleopfer, 1997). Also, certain root colonizing bacteria can protect plants from soil-borne pathogens when used as inoculants (Slininger *et al.*, 1996). *Azotobacter* strain was one of the most active PGPR and dominant bacteria in

the rhizosphere and had been intensively used as nitrogen fixers and bio-control agents. Bio-control by using antagonistic microorganisms is a potential alternative to chemical compounds for crop protection against phytopathogens (El-Katatany *et al.*, 2003). The mechanisms through which *Azotobacter* spp. control plant diseases involve competition for niches and nutrients antibiosis, predation, and induction of plant defense responses. In past years, *Azotobacter* sp. had drawn a worldwide attention because of production of secondary metabolites such as siderophore, antibiotics, enzymes and phytohormones and involving in nitrogen fixation.

It has been observed that the antagonistic activity also conferred through the action of siderophores produced from *Azotobacter* spp. called Azotobactin. The Azotobactin possesses three different types of coordinating moieties, a hydroxamate, an R-hydroxy acid, and a catechol, making it a representative of several classes of siderophores simultaneously. The known interference of pyoverdine-mediated uptake of Fe^{III} provides indirect evidence that azotobactin shows antibacterial effect against root colonizing bacterial and fungal pathogens (Schalk, 2008). Azotobactin is thus a good model compound to test whether the results obtained on the uptake of iron and can be extended as bio-control agent for the plant. The azotobactin seems to be released only under severe Fe limitation (Duhme *et al.*, 1998). Therefore azotobactin is sometimes considered the “true” siderophore of this bacterium (Page *et al.*, 1991). These had been implicated in reduction of plant pathogenic fungi and harmful rhizobacteria (Gupta *et al.*, 2001). *Azotobacter* sp. by itself has utility as an extremely economical and eco-friendly bio-pesticide (Rachin and Ahmed, 2005).

This work interested with the molecular identification through 16S rDNA sequencing of *Azotobacter* sp. isolated from soil, production of siderophore, process optimization (pH, cell mass, iron, sugars, organic acids, nitrogen sources, and

metals), partial purification of siderophore and application study of siderophore, antagonistic activity towards plant fungal pathogens.

MATERIAL AND METHODS

Isolation of *Azotobacter* sp.

The rhizosphere soil samples from various plant roots were collected from different locations of cultivable lands around Chennai. The enrichment of the culture was made by inoculating one gram of soil sample into sterile Ashby's mannitol broth (G/L: Mannitol – 20, K₂HPO₄ – 0.2, MgSO₄ - 0.2, NaCl - 0.2, K₂SO₄, CaCO₃ - 15, pH-7.4) and incubated at room temperature for three days for the formation of pellicle. The pellicle was collected aseptically and inoculated into same fresh broth incubated further in the same condition. This method was repeated thrice for the enrichment. After enrichment, the pellicle was streaked into sterile Ashby's mannitol agar medium and incubated at 28°C for more than 48 hours.

Screening for Siderophore production

In order to screen the production of siderophore, the isolates were grown in Chrome azurol sulfonate agar medium. The inoculated agar plates were incubated at 37°C for 24 hours. The observation was made for the change of medium color from blue to reddish yellow to determine the siderophore production. The isolate which showed high intensity of color change was considered for further study.

Identification of the bacterial isolate

The selected bacterial isolate was subjected to identification by conventional methods such as Gram's staining to check the Morphology and Gram's reaction. The motility was checked using Hanging drop method. The further identification was done using routine biochemical tests based on the key provided by the Bergey's manual of Systematic bacteriology.

The molecular identification of the positive isolate was carried out through 16S ribosomal coding gene sequencing. DNA extraction was carried out

by alkaline lysis and phenol chloroform extraction method. The PCR amplification of 16S ribosomal RNA coding sequence using universal primers and the reaction conditions as follows: 94°C (1 min) for denaturation, 51°C (30 sec) for annealing and 72°C (1 min) for extension. The amplified products were purified and cycle sequenced using Big Dye terminator cycle sequencing kit. The sequence was determined using ABI PRISM sequence analyzer. The obtained sequences were subjected to BLAST (Basic Local Alignment Research tool) analysis with online tool with the preexisting sequence available in NCBI/Genbank to confirm the sequences. The obtained 16S ribosomal RNA sequences of strain was aligned with various *Azotobacter* spp. obtained from GenBank, NCBI using Clustal-X Ver.2.0 aligning tool. The phylogenetic analysis was carried out using MEGA (Molecular Evolutionary Genetic Analysis) Ver.4.0 (Nei and Kumar, 2000).

Siderophore production

The culture inoculum was prepared using Luria Bertani broth contains 1% Peptone, 1% Sodium chloride and 3% Yeast extract. The isolate was inoculated into sterile LB broth and incubated at 37°C for overnight and 150 rpm agitation. The Siderophore production was carried out using Fe-deficient mannitol medium containing g L⁻¹: K₂HPO₄, 6.0; KH₂PO₄, 3.0; MgSO₄ 0.2; NH₄SO₄, 1.0 and succinic acid, 4.016; pH-7.0 was inoculated with 1% of overnight inoculums and incubated at 30°C for 24-30 hours with constant shaking at 120 rpm. The culture broth was centrifuged at 10,000 rpm for 10 min and the cell free supernatant was subjected to siderophore assay.

Partial purification

Partial purification of siderophore was carried out by ion exchange chromatography using amberlite IR120 (Mayer *et al.*, 1998). After the production of siderophore, the bacterial cells were harvested by centrifugation at 10000 rpm for 5 min and the supernatant was collected. The supernatant was considered as crude siderophore extract and pH

was adjusted to pH 6.0. The crude extract was passed through an ion exchange chromatographic column containing Amberlite IR120 (Na⁺). The siderophore was later eluted with methanol (60%). The obtained partially purified siderophore was proceeded with further studies.

Siderophore assay

Siderophore assay was carried out based on the CAS shuttle assay of Payne (1994). The culture extract (0.5 ml) was mixed with CAS reagent and 0.5ml of chrome azurol S reagent. The color obtained was measured using the spectrophotometer at 630 nm after 20 minutes of incubation. The blank was prepared using uninoculated broth medium by carrying same procedure. The siderophore content in the aliquot was calculated by using following formula:

$$\% \text{ of siderophore unit} = \frac{As - Ar \times 100}{As}$$

Where, As = absorbance of sample at 630 nm (CAS reagent) and Ar = absorbance of the reference

Optimization of Physico-Chemical parameters for Siderophore Production

Siderophore production as a function of time

The siderophore production was carried out by submerged shake flask fermentation in using mannitol medium at 35°C, 100 rpm agitation. Culture was aseptically with drowned periodically at every 6 hours intervals up to 48 hours. The growth rate of the culture was estimated by spectrophotometrically at 360nm and siderophore content of the cells free culture filtrate at different intervals was estimated.

Effect of pH

The mannitol medium was prepared with different pH values ranging from of 4 to 10 in separate flasks. Each flask was seeded with 3% overnight inoculum of *Azotobacter* sp. and incubated at 35°C for 30 hours under shaking at 120 rpm. After incubation the growth and siderophore production was measured at 410 and 630 nm respectively.

Effect of sugars and organic acids

The sterile mannitol basal medium was prepared without carbohydrate source and aliquoted into different conical flasks. Each flask were amended with different carbon sources like glucose, sucrose, dextrose and lactose, different organic acid like oxalic acid, malic acid and lactic acid, at the concentration of 0.1%. The flasks were inoculated with 3% of overnight inoculum and incubated at 30°C for 36 hours under shaking at 100 rpm. After incubation the growth and siderophore production was estimated.

Effect of nitrogen sources

The sterile basal nitrogen source free mannitol medium was prepared and amended with different concentrations of organic nitrogen sources like peptone and yeast extract, inorganic nitrogen source such as urea, ammonium nitrate and ammonium chloride added at the concentration range of 0.1%. The medium inoculated with 3% of *Azotobacter* sp. overnight inoculum and incubated at 35°C for 30 hours under shaking at 120 rpm. After incubation the siderophore production was estimated.

Effect of iron concentration

The mannitol medium was prepared with different iron concentration ($\text{FeCl}_3 \cdot 6\text{H}_2\text{O}$) in the range of 1-100 μM . The medium inoculated with *Azotobacter* sp. and incubated at 30°C for 36 hours under shaking at 100 rpm. After incubation the growth and siderophore production was estimated.

Effect of metal ions

The sterile mannitol medium was prepared in several flasks and amended with 100 μM /L concentration of different metal ions like Na^{2+} , K^+ , Ca^{2+} , Cu^{2+} , Mg^{2+} , Mn^{2+} , and Zn^{2+} . The flasks were inoculated and incubated at 35°C for 36 hours under shaking at 100 rpm. After incubation, siderophore production was estimated.

Antagonistic activity of *Azotobacter* sp.

Dual culture technique

The antagonistic properties of *Azotobacter* sp. were tested against plant fungal pathogens such as *Fusarium* sp., *Alternaria* sp., *Phytophthora* sp.,

Rhizoctonia sp., *Colletotrichum* sp. and *Curvularia* sp. by dual culture technique (Rabindran and Vidhyasekaran, 1996). *Azotobacter* sp. was streaked on Potato Dextrose Agar (Potato infusion-4 g L⁻¹; Dextrose-10g L⁻¹; Agar-18g L⁻¹) at one side of petri dish, approximately 1 cm away from the edge. Five mm of fungal mycelial agar disc of actively growing 8-15 day-old-culture from PDA was taken and placed at the opposite side of petri dishes perpendicular to the bacterial streak. Petri dishes were then incubated at room temperature for 4 days. Petri dishes inoculated with fungal discs alone were served as control. The experiment was done in triplicates. Observations on width of inhibition zone and mycelial growth of test pathogens were recorded and percent inhibition of pathogen growth was calculated.

Where, I=Inhibition of Mycelial growth, C=Growth of plant pathogen in the control plate (cm) and T=Growth of pathogen in dual cultures (cm).

$$(\%) I = \frac{(C - T) \times 100}{C}$$

Bio control assay of partial purified siderophore

Sterile potato Dextrose Agar was prepared and solidified in petri dishes. The well size 6mm diameter was made aseptically in the agar plate. Forty μl of partially purified siderophore was added to the well and allowed to diffuse radially for 1 hour at low temperature. The inoculum size of 6×10^3 spores of plant pathogenic fungi was swabbed over the surface of the PDA plates and incubated at room temperature for 48 hours. After incubation, the zone of inhibition of the mycelia growth was measured.

RESULTS AND DISCUSSION

There were 26 bacterial isolates that produced siderophore were isolated and siderophore production was confirmed by the color change from blue to reddish yellow (Milagres *et al.*,

1999). In order to obtain nitrogen fixer among the siderophore producers, The isolates were short listed by inoculating the isolates using Derx's medium. Among the short listed isolates which showed double benefit organisms that producing siderophore and nitrogen fixing capacity, potent siderophore producer was used for further study. The siderophore production is quite common phenomenon exhibited by various organisms like *Pseudomonas* spp., *Bacillus* spp., Clinical isolates like *E. coli*, *Klebsiella* etc. (Syed and Vidhale, 2011) for the acquisition of iron complex from the soil. The production of antagonisms against fungal pathogen through siderophore production by nitrogen fixing organisms giving dual benefit for the plants (Frank *et al.*, 1983).

Identification of Bacteria

The potent isolate was Gram negative short oval shape rod in nature and motile. The biochemical test showed that the isolate could be considered as *Azotobacter* sp. (Table 1). For the confirmation of identification, molecular identification was carried out 16S ribosomal coding gene identification. The amplified gene product showed 1 Kb in size in agarose gel electrophoresis whereas showed 995 nucleotide base pair in size after sequencing. The Blast result revealed that obtained nucleotide base pairs showed 100% identify with *Azotobacter chroococcum* (Accession AB742373) 16S ribosomal genes sequence (Table 2). Furthermore, the sequence was aligned and phylogenetic tree was constructed using related sequences. The sequence paired with *Azotobacter chroococcum* of Indian isolate with bootstrap value of 99% (Fig. 1). So the isolate was confirmed as *Azotobacter chroococcum* and designated as IMS. Widely, the biochemical test of identification is found to be preliminary identification in which the genus will be identified and species can be identified certain extent. Whereas the molecular identifications through 16S ribosomal RNA coding sequence analysis is widely accepted method among the

modern techniques for prokaryotes. The prokaryotic 16rRNA sequence consisted of maximum of 1.6 kb in size and unique sequence distributed among the prokaryotes facilitates to identify the species. The complete sequence of 16S rRNA comprises of approximately 1.5 kb has been so far reported (Gimmestad *et al.*, 2006).

Growth rate and siderophore production

The organism did not exhibit any traces of siderophore synthesis during beginning hours of incubation whereas the bud in the vegetative growth was observed. The strain started siderophore production after 10 hours of inoculation (Fig. 2). The maximum of siderophore production was observed (65% of siderophore units) at 30 hours of incubation and started decreasing beyond 42 hours of incubation. The highest production of siderophores occurred in Nitrogen fixing *Azotobacter* sp. at a period of 18 hours coinciding with the stationary growth phase (Frank *et al.*, 1983).

Production optimization

Of the different level of pH tested, production of siderophore by *Azotobacter* sp. was optimum at pH 7.0 with the yield of 62. In the pH values below and above the neutral the production was also somewhat significant (Table.3). The most of the previous reports of *Azotobacter* sp. have been stated that the siderophore has been produced in large quantity at neutral pH (Frank *et al.*, 1983). The other nitrogen fixing bacteria *Rhizobium* strains isolated from root nodules of *Sesbania sesban* (L.) Merr. produced siderophore in culture after 4 h of incubation and at neutral pH (Sridevi and Malalialah, 2007) Since the production of siderophore depended largely on the pH of the medium. It was very important to maintain the pH of the fermentation broth between 7.0-7.5, which is the optimum for siderophore production in *Azotobacter* sp. The pH of the medium increased from 7 to 8.5 during the growth period in accordance with the siderophore concentration

which suggest that alkalinity is important to avoid siderophore destruction -

The mannitol medium with different organic acids oxalic acid, malic acid and lactic acid, at the concentration of 4.0 g L⁻¹ showed considerable amount of siderophore production by *Azotobacter* sp. in which the quantity was high in malic acid with the yield of 71% (Fig. 3). The presence of sugars such as sucrose increased the growth of the organism but not siderophore production. Similar results were obtained in *Azotobacter* sp. (Anne *et al.*, 2000) and in *Pseudomonas fluorescens*, *Pseudomonas putida* and *Alcaligenes faecalis* (Sayyed *et al.*, 2005). In contrast *Nocardia levis* MK-VL_113 utilized sucrose and tryptone as good carbon and nitrogen sources for the elaboration of bioactive metabolites (Kavitha and Vijayalakshmi, 2009). Mannitol and sucrose at the concentration 2% stimulated growth and siderophore production in *Rhizobium* strains (Sridevi and Malalialah, 2007). Nitrogen sources such as peptone, yeast extract, urea, ammonium nitrate and ammonium chloride was added to the mannitol medium at the concentration of 1 g L⁻¹. Among these, the organic nitrogen peptone and urea enhanced the siderophore of *Azotobacter* sp. with the yield of maximum 70% respectively. In other nitrogen sources siderophore production was almost same as control (Fig. 4). Some of the amino acids as nitrogen sources tested stimulated the siderophorogenesis in *Azotobacter* sp. (Anne *et al.*, 2000). Arginine and praline stimulated growth and siderophore production in *Rhizobium* strains (Sridevi and Malalialah, 2007). The *Nocardia levis* MK-VL113 utilized tryptone as a good nitrogen source for the elaboration of bioactive metabolites. The presence of urea metabolism is known to increase the pH due to hydrolysis of urea into ammonium ion intracellularly (Sama *et al.*, 2009).

Effect of iron concentration

The iron concentration is an essential parameter which determines the production of siderophore. The mannitol medium was prepared with different iron concentration (FeCl₃.6H₂O) in the range of 1-

100 µm. The presence of low concentration (1-3µM) of iron enhanced the siderophore production of *Azotobacter* sp. up to 82%. The iron concentration between 5µM to 30 µM decreased the production and beyond this level the production was inhibited though growth occurred (Fig. 5). Increase in growth with increase in iron concentration reflects the iron requirement for the growth of the organism for cellular process (Syed, *et al.*, 2005). Siderophores are iron-specific compounds which are secreted under low iron stress and found that production of siderophore in the medium employed was inversely proportional to the iron concentration in the medium (Budzikkiewicz, 1993). The addition of limited iron in the media can increase the growth of the culture, which can lead to an increased production of siderophore in *Rhizobium* strains (Sridevi and Malalialah, 2007). The increase of Fe (III) concentration had a negative effect in siderophore production, especially above 10 µM (Villigas *et al.*, 2002).

Metal ions

Inorder to find out the effect of other minerals towards siderophore, the production was carried out in mannitol medium amended with different mineral ions. The metal ions Mg²⁺ and Zn²⁺ in the medium enhanced the siderophore production. The other metal ions tested inhibitory effect on siderophore production (Fig.6). The siderophore and cofactors produced by *Azotobacter* sp. is an iron chelating substance with the ability to bind to different metal ions was increased, Molybdenum and Cadmium (Neilands, 1974). Whereas, synthesis of metabolites azotichelin and aminochelin was inhibited by Mn²⁺ and Zn²⁺ in *Azotobacter vinelandii* (Page, 1995). Lead increased the siderophore production in *Pseudomonas fluorescens*, *Pseudomonas putida* while Mn²⁺, Hg²⁺ and Co²⁺ decreased the production (Sayyed *et al.*, 2005).

Antagonistic activity

Azotobacter sp. was evaluated for its ability to control plant root fungal pathogens such as

Fusarium sp., *Alternaria* sp., *Phytophthora* sp., *Rhizoctonia* sp., *Colletotrichum* sp. and *Curvularia* sp. by dual culture technique and assay of cell free supernatant. It was found that the *Azotobacter* sp. Lp1 inhibited all the tested pathogens in both the methods of evaluation. The percentage of inhibition was increased from 40% to 50.5% in partially purified siderophore (Fig. 7). This proved the ability of siderophore as biocontrol agent in our study. *Azotobacter* sp. was found to effectively inhibited 20-40% of the tested fungi in dual culture technique and 25-50% in purified siderophore assay (Table.4). *Azotobacter* sp. produces azotobactin type siderophore under iron starving conditions with high stability and affinity for iron that restricts growth of microorganism with low iron competition ability such as phytopathogenic fungi (Kloepper, 1977). Purified siderophore of *A. calcoaceticus* at 500 µg/mL concentrations inhibited the growth of phyto-pathogens up to 30.00%, suggested that both siderophore rich supernatant as well as pure siderophore has the inhibitory potential against phyto-pathogenic fungi (Prashanth *et al.*, 2009). The isolates of *Azotobacter* spp. were found to effectively inhibit the mycelia growth of both fungal pathogen in dual cultures with rhizospheric bacteria and soil borne pathogens (Sapna *et al.*, 2012).

Siderophore mediated antagonism was observed against common phytopathogens viz., *A. flavus*, *A. niger*, *C. capsicum* and *F. oxysporum* by *A. calcoaceticus* isolated from wheat rhizosphere (Sapna *et al.*, 2012).

CONCLUSION

The production of microbial metabolites and their applications in various fields were gaining attention. Siderophores facilitates the bacteria to take up iron under iron-limited environment. In the present study, siderophore producing bacteria was isolated and the production conditions were optimized for their better production. A further

study in this field can bring about their applications in various fields.

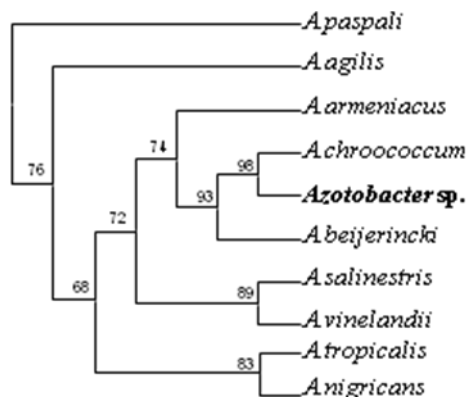
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Fig. 1 Taxonomic Position of *Azotobacter* sp. IMS



GenBank Accession: *Azotobacter vinelandii* (JX472999.1), *Azotobacter chroococcum* (AB742373), *Azotobacter salinestrus* (JX680337), *Azotobacter beijerinckii* (NR_041037), *Azotobacter paspali* (AF035210), *Azotobacter tropicalis* (JX262169), *Azotobacter nigricans* (NR_041033), *Azotobacter agilis* (NR_041034), *Azotobacter armeniacus* (NR_041037).

Fig. 2 Growth and Siderophore production relationship

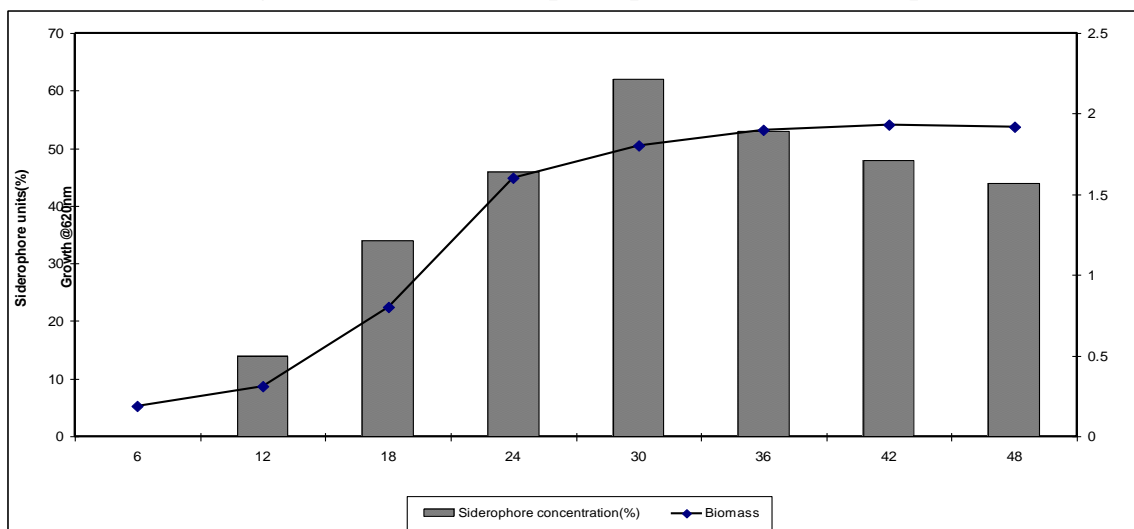


Fig.3 Effect of Carbon sources on Siderophore production.

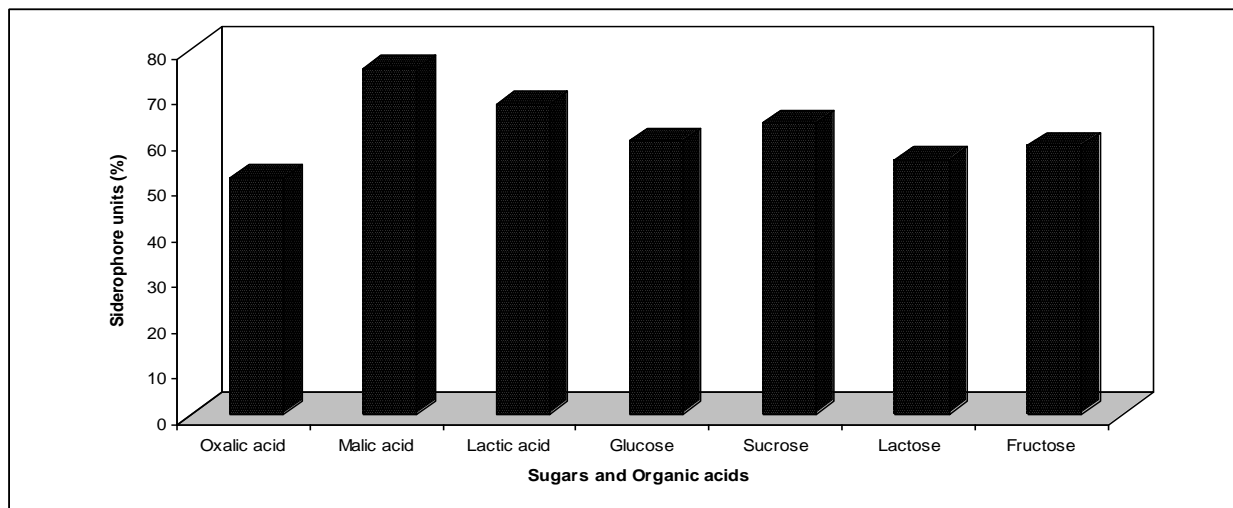


Fig.4. Effect of Nitrogen sources on Siderophore production

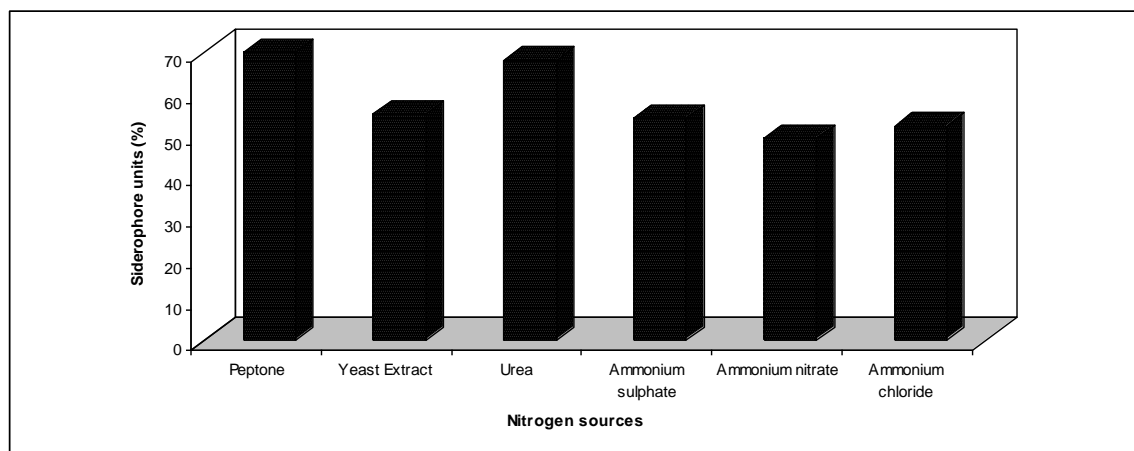


Fig.5 Effect of Iron concentration on Siderophore production

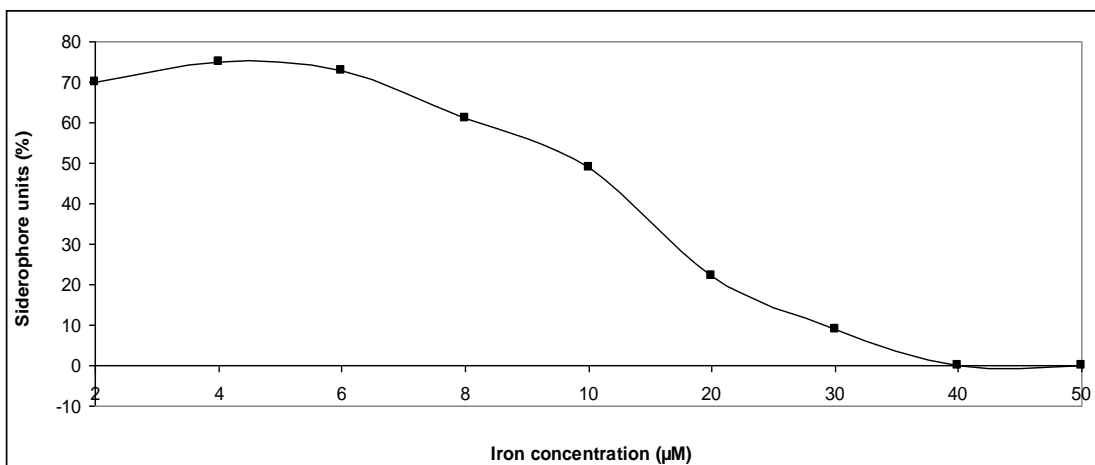


Fig.6 Effect of metal ions on Siderophore production

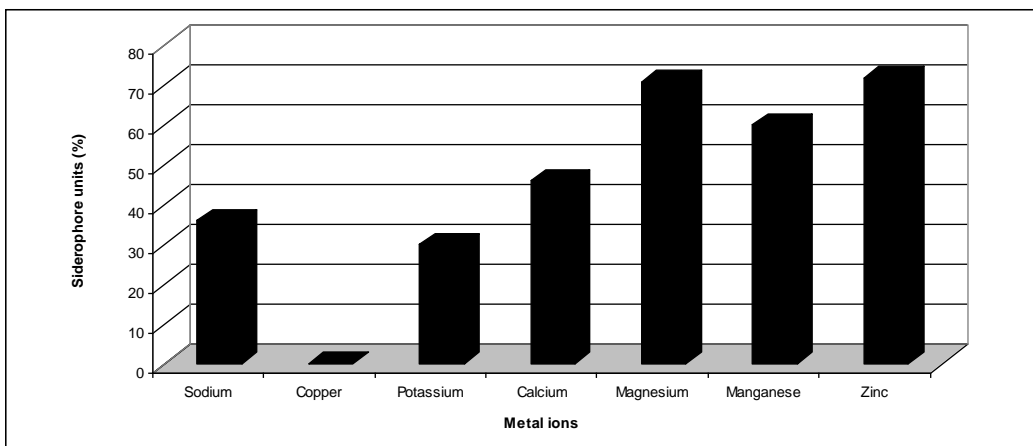


Fig.7 Antogostic activity of siderophore against *Fusarium oxysporum*

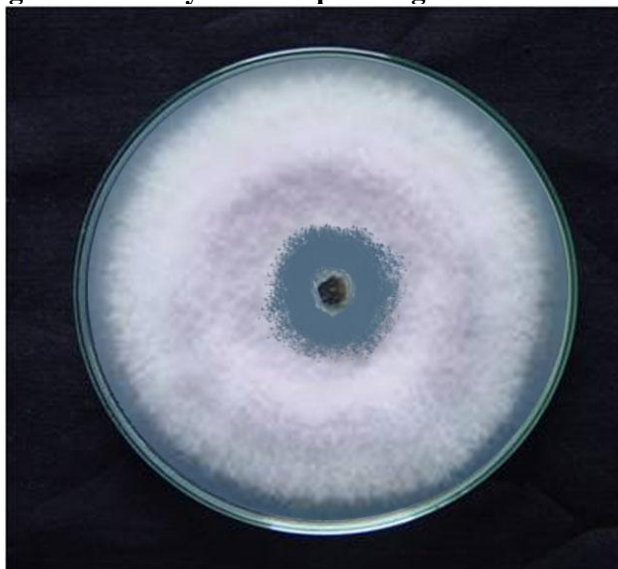


Table 1 Biochemical characters of *Azotobacter* sp. IMS

Colony Morphology		Cell Morphology	Gram's reaction	Motility	Capsule	Metachromatic Granules	Sugar Utilization	
3 Days Growth	7 Days Growth						Rhamnose	Mannitol
Soft, mucoid, flat colonies	Yellow white colonies	Short ovoid Bacilli	Negative	Motile	Present	Present	Negative	Positive

Table 2 Blast result of 16S rRNA coding sequence of *Azotobacter* sp. IMS

Sequences producing significant alignments:					
Description	Max score	Total score	Query cover	Max ident	Accession
<i>Azotobacter chroococcum</i> gene for 16S ribosomal RNA, partial sequence, strain: SUP-III	1245	1245	100%	100%	AB742373.1
<i>Azotobacter chroococcum</i> small subunit ribosomal RNA gene, partial sequence	592	738	62%	100%	AF035211.1
<i>Azotobacter chroococcum</i> strain LMG 8756 16S ribosomal RNA gene, partial sequence	588	730	61%	100%	EU401906.1
<i>Azotobacter chroococcum</i> gene for 16S rRNA, partial sequence, strain: NBRC 102613	588	730	61%	100%	AB681887.1
<i>Azotobacter chroococcum</i> strain REG18 16S ribosomal RNA gene, partial sequence	588	730	61%	100%	EU647694.1
<i>Azotobacter chroococcum</i> strain ICMP 15214 16S ribosomal RNA gene, partial sequence	588	730	61%	100%	EF100154.1
<i>Azotobacter chroococcum</i> gene for 16S rRNA, partial sequence	588	730	61%	100%	NR_041035.1
<i>Azotobacter chroococcum</i> strain GD5 16S ribosomal RNA gene, partial sequence	582	725	61%	100%	JX913866.1
<i>Azotobacter beijerinckii</i> gene for 16S rRNA, partial sequence, strain: G7	582	725	61%	100%	AB429527.1
<i>Azotobacter chroococcum</i> strain 10006 16S ribosomal RNA gene, partial sequence	582	725	61%	100%	EU930421.1

Table 3 Siderophore production at various pH Level

S.No.	PH	Siderophore unit
1	4	Trace
2	5	29
3	6	56
4	7	62.8
5	8	58
6	9	28
7	10	Trace

Table 4 Biocontrol efficiency against fungal phytopathogens

Plant phyto pathogens	Dual culture technique		Bio control assay partial purified siderophore
	Radial growth (cm)	Inhibition over Control (%)	Zone of inhibition in diameter (mm)
<i>Fusarium</i> sp.	6.5	18.75	29
<i>Curvularia</i> sp.	6.2	22.5	18
<i>Alternaria</i> sp	4.3	46.25	19
<i>Phytophthora</i> sp.	5.8	27.5	12
<i>Rhizoctonia</i> sp.	3.3	18.6	13
<i>Colletotrichum</i> sp.	3.2	17.5	16