



## ***Providencia rettgeri* AVR20 with multiple tolerance of Heavy metals from municipal solid waste dump site**

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### Abstract

The presence of toxic heavy metals (HMs) from municipal solid waste (MSW) hoists serious concern about the adverse impact to the environment. These researches focused to isolation and identification of native metal resistant bacteria and assess their efficacy against heavy metals. Metal resistant strains were isolated by pure culture technique. Identification of strains based on morphological, biochemical and molecular characteristics. The bioremediation efficacy tested by enrichment method of different metals. Twenty-five stains were selected, among these *Providencia rettgeri* AVR20 (Genbank accession number MH327529) had the multiple metal resistance. Growth and viability of isolate was measured by enrichment of metals separately, 33.7% of viable cells observed in medium enriched with 100 mg/l of Arsenic. Likewise, 33.3 % of cells in Lead and 63.8% of cells in Nickel. Efficacy of metal degradation measured, the strain AVR20 effectively degrade 51.4 % of Arsenic, 76.5 % of Lead and 80.9 % of Nickel. Effective metal degradation ability of *Providencia rettgeri* AVR20 is suitable for bioremediation of HM pollution.

## 1. Introduction

Waste management is a challenging issue, as a result of rapid urbanization and population explosion. Globally, increasing concentration of heavy metals (HM) in municipal solid waste (MSW) is major threat to the living organism in the ecosystem. Open dumping of MSW is generating leachate by rainwater percolating through the waste layers in a landfill. It contains a large amount of contaminants among the various hazardous substance HM are harmful pollutants always associated with municipal solid waste & contaminate the surrounding environment [14]. Significant amount of HM are enter into the environs through the disposal of waste by way of open dumps and landfills. [10]. Metals like Arsenic are the rich element, present most part of our environment. It is easily attached fine particle in the air and stay for many days. Arsenic mainly used as an herbicide, insecticide, germicide, medicine

and wood preservatives also in electronics and industrial manufacturing industries [24]. When dumping of these wastes to landfill can get into to the environment through the leaching process. Continuing exposure of arsenic induces cancer, skin lacerations and cardiovascular defect, and inhalation of Arsenic affects respiratory and nervous system [22]. Many of the peoples from Argentina, Chile, China, Hungary, Mexico and USA were showing to elevated amount of As through drinking of As polluted ground water [3,17].

The widespread use of lead can contaminate environment for example, use agricultural pesticide of lead arsenate will enter in to the ground water. EPA declared that dump is a site dispose waste without environmental controls and municipal, industrial waste are disposed directly to the landfill but in sanitary landfill waste managed with environmental production standard [21, 13]. Lead is the naturally occurring toxicants, and affects gastrointestinal, cardiovascular, renal, hematological, neurological and immune function [23]. Nickel distributed in water, soil and air, and trace level of Ni essential for various metabolic activities of all organisms, however long term exposure can cause lung cancer, respiratory failure, birth defect and heart disease [5].

Microorganism present in the polluted environment can develop special mechanism to tolerate or remove toxic metals. The occurrence of organic and inorganic substances in landfills are highly attracted by microorganism especially bacteria are greatly reduce the level of HM from the environment. Bioaccumulation, bio sorption and bio reduction are the successful process for biodegradation of metals [15].

Some plants have a vital role in the heavy metal removal mechanism, according to Stofejova et al., 2021, Ar and Pb are not essential for the plant, and they accumulate on plant parts and cause toxic effects. Irrigation of metal-contaminated water or plants grown in polluted soil will increase the level of heavy metals in plants.

Hence the research goal is to examine the capability of native bacteria from landfill involved in reducing the level of HM (**As**, **Pb** and **Ni**). Naturally occurring metal resistant bacteria was isolated and characterized to reduce the level of bacteria. To ensure the level of resistance and degradation is useful for the determination of bacteria is suitable for bioremediation.

## **2. Methodology**

### **2.1. Location and sample collection**

Municipal solid waste and leachate samples were collected from landfill dumpsite Madurai, Tamil Nadu, India. The samples were kept in pre sterilized bags and bottles and stored at 4 °C for further microbiological analysis.

### **2.2. Isolation and selection of HM resistant bacteria**

In the effective isolation process one gram of solid waste sample and one ml of leachate were mixed with 98 ml of saline water (8 g NaCl in 1000 ml distilled water) for 24 hours, under shaken (180 rpm). Screening and isolation of pure culture done with 0.1 ml of sample inoculated on Luria Bertani (LB) agar plate enriched by HMs (10 mg/l of Pb as PbCl<sub>2</sub>) then incubated on 37° C. After 2days of incubation, plates were witnessed the growth of any culture on plates. After preliminary screening of metal degrading bacteria, pure culture method was done to isolates desired bacteria. Control plate was prepared without HM in LB agar. Colonies were picked and purified then preserved for further analysis.

## 2.3 Biochemical, Molecular and Phylogenetic analysis

The pure culture of metal tolerant strain was characterized by morphologically and biochemically using standard procedure from Bergey's Manual of Systematic Bacteriology 1994 [8,4]. According to Molecular identification of the strain AVR20 was done by using 16Sr DNA analysis. DNA was extracted, and then amplification of bacterial genome carried out with polymerase chain reaction (PCR). Forward (5'-AGAGTTTGATCCTGGCTCAG-3') and reverse primers (5'-ACGGCTACCTTGTTACGACT-3) were used for PCR analysis. Steps involved in PCR amplification are initial denaturation at 94°C for 4 min; 30cycles for amplification of 60S at 94°C, 60s at 55° C, and 90s at 72° C respectively. Sequence was examined on the National Center for Biotechnology Information (NCBI). Phylogenetic tree constructed by Maximum Likelihood method [20] was built to deliver a graphic representation of the patterning of relationship among individuals. MEGA version 7 software is highly useful to construct phylogenetic tree [11].

## 2.4 Multiple metal resistance assay

Totally 25 isolates were selected for multiple metal tolerant assay. Isolates were cultured on( LB broth) enriched by As, Ni and Pb (50mg/l) and incubated 37° C for 48 hours, after incubation the multiple resistance isolate was collected and preserved and stored for further analysis.

## 2.5 Efficiency and growth of metal resistant strain

Among the 25 isolates one strain AVR20 has the ability to tolerate As, Pb and Ni. To measure the efficiency of selected strain was experienced with increasing concentration of metals, and growth measured by OD at 600nm. The bacterial culture (1 ml) inoculated 100 ml of LB broth amended with increasing concentration (0, 25, 50,75and 100mg/l) of HMs (As as AsCl<sub>2</sub>, Pb as PbCl<sub>2</sub> and Ni as NiSo<sub>4</sub>) separately. All the culture was incubated at 37°C for 3 days. Growth of metal resistant bacteria were monitored the absorbance at 600 nm.

## 2.6 Heavy metal degradation assay

One ml of cells was inoculated in LB broth (100 ml) dissolved with 100 mg/l As, Pb and Ni respectively. Culture was incubated at 37°C for 3 days, then culture was centrifuged at 5000 rpm (15 min) then supernatant was collected and digested with concentrated HNO<sub>3</sub>(10ml) at 80°C Allen et al., 1986) [1]. The resulting solution was analyzed for concentrations of HMs using an atomic absorption spectrophotometer (Modal-ELICO, SL173) APHA, 2005 [2]. Heavy metal degradation capacity (%) calculated by :

$$\%HM \text{ utilized} = \frac{HM \text{ utilized mg/L}}{\text{Concentration of HM in medium mg/L}} \times 100$$

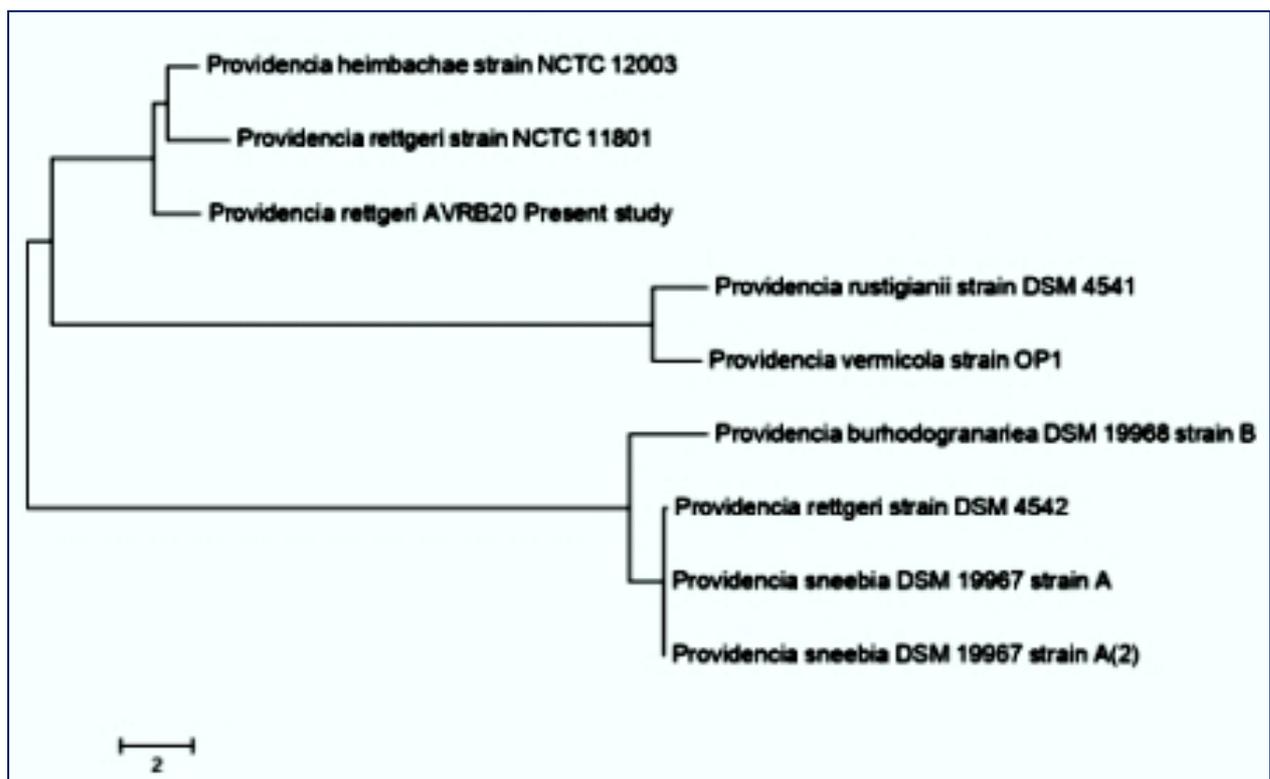
HM utilized mg/l = Initial concentration of HM in broth mg/l-Final concentration of HM in broth mg/l

## 3 Result and discussion

### 3.3 Isolation and molecular identification of HM resistant bacteria

Twenty-five isolates were obtained from municipal solid waste and leachate mixed sample. All the isolated have the ability to grow in 10 mg/l of Pb containing LB agar medium. Further the isolates were involved for multiple metal tolerance tests with 50 mg/l of As, Pb and Ni respectively. Based on the study the strain AVR20 was more suitable for the bioremediation of heavy metal. The chosen

bacteria is a gram negative, motile and rod shaped bacteria. The biochemical analysis showed that this belongs to the family *Enterobacteriaceae* and genus *Providencia*. Molecular identification by gene (16S rRNA) amplification and sequencing showed that the isolate had 100% similarity with *Providencia rettgeri*; isolated strain was submitted in Genbank (Accession number MH327529). Phylogenetic analysis done with BLAST similarity search, and tree constructed with Neighbor joining method. (Figure 1) the present isolate designated as *Providencia rettgeri* AVR20. Similarly a study conducted by Shardendu et al., 2017, they isolated arsenic resistant bacteria *Paracoccus denitrificans*, *Alcaligenes faecalis* and *Stenotrophomonas maltophilia* from waste water. In the same way a study isolated arsenic resistant *Rhizobacteria* FB4 and FB9 with maximum tolerance of arsenic 30-1000 ppm [19].



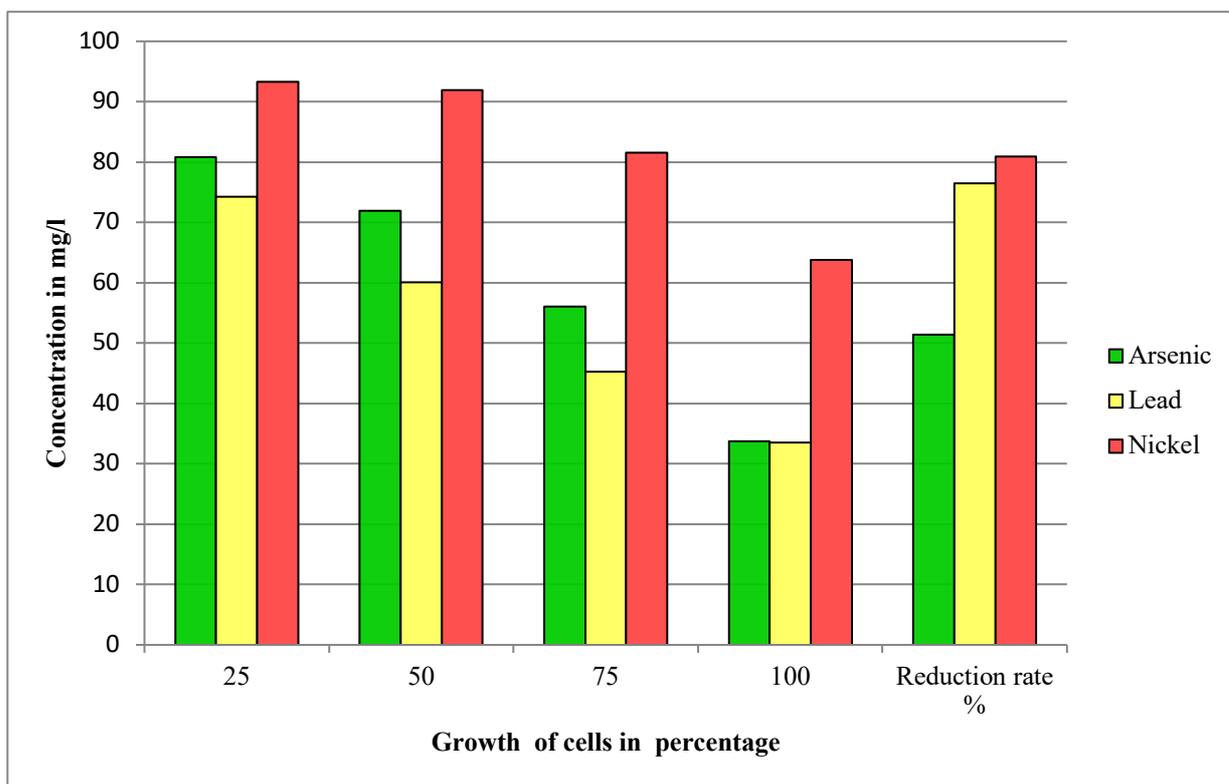
**Figure 1.** Phylogenetic tree of isolate *Providencia rettgeri* AVR20

### 3.4 Multiple metal resistance assay

Naturally the bacteria present in contaminated area, are developed their resistance mechanism against the pollutant. For example, number of genes responsible for their resistance mechanism *Paracoccus denitrificans* gene encoding arsenical resistant protein arsenate reductase, *Alcaligenes faecalis* gene encoding arsenite reductase, arsenite oxidase and arsenical pump proteins for As(V) and *Stenotrophomonas maltophilia* have genes for arsenite transmembrane efflux pump transporter protein, arsenate reductase and arsenic resistant transmembrane for As (III) [17]. Nickel, lead and zinc resistance observed on *Staphylococcus epidermidis*, *Proteus mirabilis* and *Escherichia coli* [12]. *Bacillus* is a spore producing bacteria and habitat in extreme environment, previous study isolated heavy metal resistant *Bacillus thuringiensis* and *Bacillus cereus* [6,25].

Municipal solid waste and leachate harbors numerous microorganisms which have evolved number of methods to survive under toxic environment. Figure 2 shows growth and viability of isolates under various trial conditions. The growth and viability of isolates were determined by turbidity in the culture

medium by measuring absorbance at 600nm spectrophotometrically. The metal resistant isolate was cultured in LB medium amended with HM. The cells were cultivated in increasing concentration of **As**, **Pb** and **Ni** separately 0, 25, 50, 75 and 100 mg/l. AVR20 highly resistant against Ni with 98.3% of cells observed in medium enriched with Ni (25 mg/l concentration) 91.9, 81.6 and 63.8% of cells observed in 50, 75 and 100mg/l of Ni enriched medium. Similarly, 80.8, 71.9, 56.1 and 33.7 % of cells witnessed in **As** enriched medium and 74.3, 60.1, 45.3 and 33.3 % cells in **Pb** containing medium of 25, 50, 75 and 100mg/l respectively. The overall investigation reported that AVR20 isolate grow well in less concentration of HMs however growth slowly decreased when concentration increased.



**Figure 2** Shows growth and viability of isolates under various trial conditions

### 3.5 Growth and metal degradation efficiency of metal resistant strain

In this study the degradation of HMs (As, Pb and Ni) using *Providencia rettgeri* AVR20 was analyzed by determining concentration of HMs in the culture growth medium. Hundred mg/l concentration of HMs were supplemented in 100 ml LB broth medium separately. One ml of culture was inoculated HM enriched medium and incubated for 3 days. Then cells were centrifuged and separated and the concentration of metals detected using atomic absorption spectroscopy. After incubation period 51.4 % of As, 76.5% of Pb and 80.9% of Ni were utilized by bacteria. Likewise the Bacteria *Chryseobacterium indoltheticum*, *Cupriavidus oxalaticus*, *Pseudomonas helmanticensis*, *Pseudomonas helmanticensis*, *Bacillus mycoides*, *Bacillus mycoides*, *Bacillus almalaya* and *Acinetobacter tjernbergiae* were isolated from soil and were able to grow in maximum of 1600 mg/ L of Pb [21]. Likewise, *Pseudomonas aeruginosa* HF5 had the ability to grow on cobalt (590 mg/l), zinc (1310 mg/l), chromium (26mg/l), lead (570 mg/l), cadmium (843mg/l) and copper (222 mg/l) [7,26-28].

## Conclusion

Municipal solid waste contains huge number of microorganisms. Generally, microbes have the capability of developing resistance mechanism when they continuously exposed in particular pollution. Metal resistance strains were isolated from municipal solid waste and leachate mixed sample. Among these *Providencia rettgeri* AVR20 had the multiple metal (As, Pb and Ni) tolerance and high efficiency of degradation. The strong degradation and multiple tolerance of the *Providencia rettgeri* AVR20 is suitable for the bioremediation of polluted area. It has the potential to become a commercially important strain for toxic compound bioremediation. This study recommended the combined action of microbes; microbial consortium with plant effectively reduces the bioavailability of metals on polluted sources.

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