INTRODUCTION

Along with industrial progress, environmental pollutants like toxic heavy metals are widely spreading throughout the world. This is especially true for developing countries like China and India (Raja et al., 2008). The uncontrolled discharges of large quantity of heavy metal-containing wastes create huge economical and health care burden particularly for the people living near that area (since the effluents of the industries excreted into the environment and through food chain, affect humans and animals from various anthropogenic sources such as industrial wastes, automobile emissions, mining activity and agricultural practices as well). The important toxic metal pollutants like cadmium, nickel and lead enter to the water bodies through industrial wastewater treatment plants (Denise et al., 1989; Ajmal et al., 1998). Cadmium is the most dangerous metal ion characterized by high stability and toxicity. It is not degradable in nature and will thus, once released to the environment, stay in circulation. Cadmium is known to bind with essential respiratory enzymes (Nies, 2003) causing oxidative stress and cancer (Banjerdkji et al., 2005). High concentrations of cadmium is highly corrosion resistant and is widely used to plate metal parts in general industrial hardware as well as in automobiles, electronics, marine and aerospace industries (Herrero et al., 2005).
Cadmium contamination has been also reported particularly in soils containing waste materials from zinc mines and in sludge amended soils fertilized with cadmium rich phosphate fertilizers (Raskin and Ensley, 2000). The current low world market price of cadmium motivates the development of new applications that by time may develop into new sources of emissions to the environment not covered by existing regulation. Therefore, decontamination of these pollutants through bioremediation process and other biotechnological means are prerequisite for any future decision by the governments.

The potential use of metal-resistant microorganisms in the treatment of heavy metal contaminated wastewater plants has become more important (Shakibaie et al., 2008). Different biomass types, such as bacteria, fungi and algae, have been screened and studied extensively by many authors over the past decades with the aim of identifying highly efficient metal removal biological systems (Viraraghavan, 1995; Vieira and Volesky, 2000; Kapoor and Herrero et al., 2005).

Many efforts have been devoted to the isolation of heavy metal-resistant bacterial strains during the past years. *Staphylococcus aureus* (Novick and Roth, 1968) *Escherichia coli* (Mitra et al., 1975) were found to exclude cadmium ion (Cd²⁺) from cell surface. Katarina *et al.* (2004) studied cadmium resistant bacterial community isolated from sewage sludge contaminated by cadmium ions. Among bacteria from bacterial community short cadmium resistant gram-negative rods were predominated. Biochemical tests assigned the eight isolates to six bacterial species, *Alcaligenes xylosoxidans*, *Comamonas testosteroni*, *Klebsiella planticola*, *Pseudomonas putida*, *Pseudomonas fluorescens*, and *Serratia liquefaciens*. Cadmium-resistant bacterial isolates were able to remove cadmium from solution and the efficiency of cadmium removal correlated with the amount of additionally synthesized proteins in the cell fractions.

Although many researchers have studied the bioremediation of cadmium from industrial waste, none has used mutated cadmium resistant bacteria for this purpose. In this investigation, a cadmium resistant bacteria was isolated from active sludge processing plant of a food factory near the city of Kerman in Iran. By mutational enhancement technique the bacterial strain was employed for removal of Cd⁺⁺ using batch bed reactor.

**MATERIALS AND METHODS**

*Effluent sampling and source*

500g soil from depth of 0.1 meter and 5L of active sludge effluent in depth of 21cm from waste processing plant of a food factory at Kerman city, Iran, were collected in sterile 15L can and transferred to International Center for Science, High-Technology, Kerman, Iran, for further analysis. The pH of active sludge and soil were measured by pH meter (Metrohm- 691).

Concentrations of cadmium in active sludge and biomass were measured by atomic absorption spectrometry (Philips, PU 9100X). Before analysis of cadmium concentration, all samples were filtered through 0.45µm pore size hydrophilic membranes filters (Sartoreious, Millipore, Germany). The cadmium used was in the form of 3CdSO₄·8H₂O with 95% purity, purchased from Merck Co. Ltd. (Germany). Muller-Hinton agar and broth were obtained from Hi-Media (Mumbai, India). Mutagenic agents of Acridine Orange and Acriflavin were purchased from Merck Co. Ltd.

*Preliminary screening of cadmium resistant bacteria*

The preliminary screening of cadmium resistant bacteria was carried out from both active sludge and soil samples by two methods. One method was based on serial dilution technique (Shakibaie *et al.*, 2008) in which 1mL of active sludge was added to a tube containing 9mL of sterile 0.75% normal saline (10⁻¹) and mixed well. 1mL supernatant of this dilution was transferred to another 9mL sterile normal saline (NaOH 0.07%) tube to obtain final volume 10⁻². Dilution was repeated till 10⁻⁸. 200µL of each dilution was inoculated on to sterile Muller-Hinton agar (MHA) plates containing 0.5mM 3CdSO₄·8H₂O solution, spread thoroughly with sterile glass spreader and incubated aerobically at 35°C for 24- 96 hours. For control sample, inoculation was done on Muller-Hinton agar medium without.
cadmium sulphate and incubated along other plates. Similarly, all above processes were done for one gram soil sample as well.

In the second method, 1mL of active sludge effluent was inoculated in 10mL tube with 9mL Muller-Hinton Broth medium containing 0.5mM 3CdSO₄·8H₂O solution and mixed well by agitation and incubated at 35°C for 24 hours. All above processes were repeated for one gram soil sample as well. 200µL of soil and sludge suspensions were inoculated onto series of 20mL sterile Muller-Hinton agar plates and incubated at 35°C for 24-72 hours. After growth, one loopful of each colony was suspended in 1mL sterile Muller-Hinton broth containing 40% glycerol in 1.5mL Eppendorff tubes, mixed well and stored at -70°C (Shakibaie, et al., 1999).

**Determination of cadmium sensitivity**

After the preliminary isolation of the cadmium resistant bacteria, the minimum inhibitory concentration (MIC) of Cd²⁺ was determined by the agar plate dilution method as described by Malik and Jaiswal, (2000). 250mL of 0.05M stock solutions of the metal salt (3CdSO₄·8H₂O) were prepared in sterile DD/W to obtain final and concentrations of 1, 2, 3, 4, 5, 6 and 7 mM Cd²⁺, respectively. The Petri plates were inoculated with 200µL log phase liquid culture of isolated bacteria and incubated at 35°C for 24-96 hours. The MIC was defined as the lowest concentration of the Cd²⁺ that inhibits the visible growth (number of colonies) of the organisms. The Cd²⁺ sensitivity and resistance of the isolates were calculated according to published papers (Devicente et al., 1990; Sarby et al., 1997; El-Helow et al., 2000).

**Induction of mutation**

The mutagenic compounds of Acridine Orange and Acriflavine have the ability to bind and intercalate with the DNA and cause frame shift mutation (Glass, 1982). Two methods for induction of mutation were employed in this study. In the first method, Gradient Plate Technique (GPT), stock solutions of mutagenic agents were prepared by addition of 0.1mg of Acridine Orange and Acriflavine into 100mL sterile double distilled water. Various concentrations of mutagenic agents from stock solution were added to 10mL sterile melted nutrient agar medium, mixed well and kept as sloping condition. After the medium was solidified, 10mL melted soft agar was poured into the sterile Petri plates and kept horizontally till it solidified again.

The cadmium-resistant bacterial strains that were isolated in pervious stages were inoculated throughout gradient plates and incubated at 35°C for 24-48 hours. The colonies that were grown on the highest concentration of the slop, were selected and used for further study. In the second method, Sub-Inhibitory Concentration (SIC) of each mutagenic agent was determined as previously described (Shakibaie et al., 2008). 100µL of log phase bacteria (8 hours grown cell suspension) that was grown on gradient plates of Acridine Orange and Acriflavine were inoculated onto sterile Muller-Hinton Broth (MHB) medium containing different amounts of the above mutagenic agents (400, 800, 1600, 3200 and 6400 µg/mL). All the tubes were incubated at 35°C for 24-48 hours. 100µL from each tube which showed visible growth, was streaked on Muller-Hinton Agar (MHA) medium and incubated at 35°C for 24 hours. Individual colony grown on the highest concentrations of the mutagenic compounds were inoculated on MHA medium containing various concentrations of 3CdSO₄·8H₂O (1-8mM) and incubated for 24-48 hours at 35°C. The bacterial strains that were grown on the highest concentration of 3CdSO₄·8H₂O containing plates, were then isolated and stored for Cd²⁺ removal study.

**Identification of bacterial isolates**

Both, Cd²⁺ resistant gram-negative non-fermentative strains (GNNFR) and gram-positive cocci were isolated. The isolates were tested and characterized by several microbiological key conventional tests for basic differentiation of gram-negative and gram-positive bacteria as previously described in Bergeys Manual of Determinative Bacteriology. Further, the isolates were identified on the basis of biochemical tests of commercial identification systems as shown in Table 1.

**Mass balance experiment**

To determine how much of the Cd²⁺ was
precipitated by the cell in an insoluble form. 1.5mL of the samples were treated with and without cadmium and centrifuged at 10,000rpm for 15 minutes at room temperature. The supernatant and pellet were analysed for Cd\(^{2+}\) content by atomic absorption spectrometry.

For determination of Cd\(^{2+}\) concentrations in the biomasses, bacterial cell residues were dissolved in 1mL 95% nitric acid (Merck Co.), mixed well by vortexing and diluted to 10 mL with sterile DDW. Blanks were treated in the same way and analyzed as described above. Simultaneously, total viable count of the organism was determined each time to see any decrease in the colony count.

**RESULTS**

pH of active sludge effluent was 8.0 and atmospheric temperature was 25\(^\circ\)C, while ambient temperature was 20\(^\circ\)C. Several mesophilic gram-negative and cadmium resistant bacteria were isolated. The enrichment was better as compared to direct culture method for the isolation of Cd resistant bacteria in shorter time. Also, the isolates in primary enrichment method could grow on 6 mM concentration of Cd\(^{2+}\) containing medium. Majority of the bacterial isolates were belonging to gram-negative non-fermentive *Pseudomonas* (4 isolates). One gram-negative coccus was also capable to grow on 2mM concentrations of Cd\(^{2+}\); but on subsequent inoculation, the strain lost its ability to grow on more than 2 mM Cd\(^{2+}\) and was eliminated from the study. Those strains isolated from soil, exhibited MIC=3.5mM and from active sludge as 6mM for 3CdSO\(_4\).8H\(_2\)O, respectively (Table 2).

One *Xanthomonas oryzae* isolated from soil near factory could grow only on 2mM cadmium containing medium; therefore, it was not used for bioremediation study because of low MIC value. Those *P.aeruginosa* isolates grown on the 3.5-6 mM concentration of Cd\(^{2+}\) containing medium were exposed to 6400mg/L Acriflavine and 9600

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**Table 1:** Biochemical tests for identification of *P.aeruginosa* isolated from active sludge and soil near food factory disposal plant

<table>
<thead>
<tr>
<th>Biochemical tests</th>
<th>Properties</th>
</tr>
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<tbody>
<tr>
<td>Morphology</td>
<td>Short rods</td>
</tr>
<tr>
<td>Gram staining</td>
<td>Gram negative</td>
</tr>
<tr>
<td>Motility</td>
<td>+</td>
</tr>
<tr>
<td>Growth at 42(^\circ)C</td>
<td>+</td>
</tr>
<tr>
<td>Growth at 4(^\circ)C</td>
<td>-</td>
</tr>
<tr>
<td>Fluorescent pigment</td>
<td>+</td>
</tr>
<tr>
<td>Odor</td>
<td>Fruity odor</td>
</tr>
<tr>
<td>H2S production in TSI</td>
<td>-</td>
</tr>
<tr>
<td>Gas production</td>
<td>-</td>
</tr>
<tr>
<td>Growth on Simmon citrate agar</td>
<td>+</td>
</tr>
<tr>
<td>Urea test</td>
<td>+/-</td>
</tr>
<tr>
<td>Oxidase</td>
<td>+</td>
</tr>
<tr>
<td>Arginine dehydrolase</td>
<td>(weak)</td>
</tr>
<tr>
<td>OF</td>
<td>oxidative</td>
</tr>
<tr>
<td>Indole</td>
<td>-</td>
</tr>
<tr>
<td>Gelatinase</td>
<td>+</td>
</tr>
</tbody>
</table>

In case of *X.oryzae*, the yellow pigment was observed.

**Table 2:** Minimum Inhibitory Concentration (MIC) of Cd\(^{2+}\) for three *Pseudomonas* strains isolated from active sludge and soil before and after mutation

<table>
<thead>
<tr>
<th><em>P.aeruginosa</em> isolates</th>
<th>Source</th>
<th>MIC (mM) Before mutation</th>
<th>MIC (mM) After mutation</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Sludge</td>
<td>6</td>
<td>7</td>
</tr>
<tr>
<td>1</td>
<td>Soil</td>
<td>3.5</td>
<td>4</td>
</tr>
<tr>
<td>2</td>
<td>Sludge</td>
<td>2.5</td>
<td>3</td>
</tr>
</tbody>
</table>

The above tests were duplicated and similar observation was made.
mg/L Acridine Orange as shown in Table 3. The isolates could grow well on these concentrations and any changes in the colony characteristics or bacterial morphology was observed after exposure to these mutagenic agents. Table 2 compares MIC values of Cd²⁺ before and after exposure to these two mutagenic agents. The MIC increased to 7 mM for P. aeruginosa isolate 1 after exposure to the above agents.

Fig. 1 (a and b) shows removal of Cd²⁺ by mutated P. aeruginosa isolate 1 and wild type in 30 and 60 mg/L of Cd²⁺ solution, respectively. 94.7% removal was achieved till 60 minutes in 30 mg/L of Cd²⁺ by mutated bacteria. However, in 60 mg/L concentration of cadmium, only 53.58% Cd²⁺ was removed by mutated bacteria and 38.68% by wild type. In all cases the Cd²⁺ concentration rapidly decreased during the first 15 minutes.
the supernatants and remained constant during 3 hours. This indicates that as concentration of Cd$^{2+}$ was increased, the total viable count of the organism decreased and was directly related to the amount of Cd$^{2+}$ removed by the biomass. The Langmuir model was derived based on several assumptions (the surface was homogeneous, adsorption on the surface was localized and each site could only accommodate one molecule or atom).

Table 3: Gradient Plate Method (GPM) and Subinhibitory Concentration (SIC) of mutagenic agents Acridine Orange (AO) and Acriflavine (AF) for *P. aeruginosa* strains 1, 2, and 3 isolated from soil and active sludge processing plant effluent

<table>
<thead>
<tr>
<th><em>P. aeruginosa</em> isolates</th>
<th>GMP Concentration (µg/mL)</th>
<th>SIC*</th>
<th>AO 6400</th>
<th>AF 6400</th>
<th>AO 6400</th>
<th>AF 3200</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>9600</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>2</td>
<td>9600</td>
<td></td>
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<td></td>
<td></td>
</tr>
<tr>
<td>3</td>
<td>9600</td>
<td></td>
<td></td>
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</tbody>
</table>

*SIC was defined as maximum concentration of mutagenic agent that bacteria could grow.*

Table 4: Langmuir isotherm coefficients for cadmium

<table>
<thead>
<tr>
<th>Langmuir model for mutated bacteria</th>
<th>Langmuir model for wild type</th>
</tr>
</thead>
<tbody>
<tr>
<td>$q_{\text{max}}$</td>
<td>$k_l$</td>
</tr>
<tr>
<td>4.47</td>
<td>1.58</td>
</tr>
</tbody>
</table>

$q_{\text{max}}$ is the maximum removal capacity of the biomass (mg/g). The above results are average of two independent experiments.

Fig. 2 (a & b) shows the Cd$^{2+}$ removal isotherm for mutated bacteria as well as wild type which follow Langmuir model. Table 4 shows the parameters of Langmuir isotherm for Cd$^{2+}$ derived in this investigation.

**DISCUSSION**

The data obtained in this study clearly shows that with employment of cadmium resistant mutated biomass, bioaccumulation of Cd$^{2+}$ from Cd-containing solution considerably increased. *P. aeruginosa* isolate 1 could efficiently remove 94.7% in 30 mg/L of Cd$^{2+}$ solution within 60 min. The results were consistent with previously report for strain E1 (Zeng *et al.*, 2009). Strain E1 with resistance to 18 mM/L cadmium isolated from Cd-contaminated soil was identified by morphological observation, biochemical and physiological characterization (Zeng *et al.*, 2009). Both living and non-living cells of strain E1 could remove Cd$^{2+}$ from solution, and living cell had better Cd$^{2+}$ removal than non-living cell (Chelliah *et al.*, 2008).

In the other study, Cd-resistant bacteria were isolated from Cd-contaminated soils by Prapagdee and Watcharamusk (2009). One isolate, TAK1, was highly resistant to cadmium toxicity. TAK1 was isolated from soil contaminated with a high Cd concentration (204.1 mg/kg). The removal of the heavy metal ions by some gram-negative bacterial species such as *E. coli* and *P. syringae* were studied by Cohen *et al.*, (1991) and Cabral (1992), respectively. *P. aeruginosa* was found to detoxify Cd$^{2+}$ through production of intracellular cadmium-binding proteins (Hassen *et al.*, 1998). A Cd$^{2+}$-hyperresistant bacterial strain HQ-1 was isolated from a lead–zinc mine by Qing Hu *et al.*, (2007).

Shakibaie *et al.*, (2008) isolated *P. aeruginosa* strains that accumulated high concentrations of copper and zinc after exposure to mutagenic compounds. Similarly, Shakibaie *et al.*, (2003) reported that intracellular accumulation of silver
by Acinetobacter baumannii BL54 was an energy dependent process and occurs through binding to a cysteine rich metall protein. Paul and Jayakumar (2010) applied an analytical study of the cadmium and humic acids contents of two lentic water bodies in Tamil Nadu, India. From present study it can be concluded that the strain 1 can be efficiently used for bioremediation and removal of cadmium containing waste pollutants with minimum cost and high efficiency.

ACKNOWLEDGEMENTS

The authors are grateful of Chairman of International Center for Science, High-Technology and Environmental Science Center, Kerman, Iran, for providing instruments and facilities of this research and Mr. Amini for laboratory assistance.

REFERENCES


