Bioremediation of phenanthrene-polluted soil using *Bacillus kochii* AHV-KH14 as a halo-tolerant strain isolated from compost

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**Abstract**

**Background:** Phenanthrene (PHE) is a polycyclic aromatic hydrocarbon (PAH) with crystalline structure of C14H10, which was produced from incomplete combustion of hydrocarbons and fossil fuels and can cause harmful biological effects. Bioremediation using halophilic bacteria is paid attention over chemical methods due to considerable benefits.

**Methods:** In the present study, a halo-tolerant bacterium *Bacillus kochii* strain AHV-KH14 was isolated from municipal compost, and used for the bioremediation of PHE from the contaminated soil. The effects of operational parameters including soil/water ratio, initial inoculum size, PHE concentration, and salinity on the bioremediation performance were investigated.

**Results:** A biodegradation efficiency of about 98% was obtained for PHE concentration of 50 mg/kg and salinity level of 1.5%. By increasing salinity content PHE concentration, PHE biodegradation rate decreased significantly. It was found that the bioremediation efficiency decreased with increasing PHE concentration. It was also revealed that for the unwashed soil sample, cumulative concentrations of different hydrocarbons played an important role in decreasing the efficiency of bioremediation.

**Conclusion:** A natural hydrocarbon-contaminated soil sample with total petroleum hydrocarbon (TPH) concentration of 2350 mg/kg was subjected to bioremediation using the selected conditions of operational parameters, and a biodegradation rate of 17.7% was obtained.

**Keywords:** Biodegradation, Soil pollution, Phenanthrene, *Bacillus kochii* strain AHV-KH14, Compost


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**Introduction**

Polycyclic aromatic hydrocarbons (PAHs), as a group of organic compounds, are among the most common contaminants in the environment. PAHs are released into the environment through bringing fossil fuels, such as coal, oil, coal, and natural gas, as well as anthropogenic activities (1). Emissions from the anthropogenic activities, such as the production of plastics, pesticides, and dyes, coal burning power plants, and shipping activities, are considered as the main sources of these compounds in the environment (2). PAHs are mainly produced as a result of incomplete combustion of wood, tobacco coal, oil, and petrol. PAHs are released into the environment in gas and particle phases. Atmospheric PAHs can be deposited to the earth through wet and dry deposition processes. So that, these compounds can be added to surface soil and cause soil contamination (3). Short-term exposure to these compounds is associated with eye irritation, nausea, vomiting, diarrhea, and confusion (4). But there is a growing concern regarding their long-term effects. Previous studies have shown that long-term exposure to PAHs can cause cancer in some individuals. Moreover, the US Environmental Protection Agency (USEPA) has classified several types of PAHs as probably carcinogenic to humans. The presence of PAHs in soil poses serious health threats to human population (5). Phenanthrene (PHE) is one of the most widely used PAHs that has low biodegradability and high persistence in the environment. The USEPA has classified PHE as a priority pollutant. Therefore, they must be removed before being released into the environment. Until now, many technologies, such as solvent extraction, phytoremediation, photocatalytic remediation, electrochemical remediation, thermal destruction, and bioremediation, have been tested for
the remediation of soils contaminated by hydrocarbons. Biological processes have received many attentions from scientists and researchers due to their less toxicity to the environment, easy maintenance, high-performance, and economic aspects of the contaminated soils remediation. Bioremediation is a biological process used for the treatment of organic pollutants from various media, such as water, soil, air, and surface of materials. In bioremediation, microorganisms or their enzymes are applied to treat and remove organic pollutants from various media (6). Bioremediation, which relies principally on living organisms, especially fungi, bacteria, and plants, is a reliable technique for the degradation of persistent organic pollutants (POPs), such as pesticides, solvents, pharmaceuticals, and PAHs. Bioremediation as a promising technique offers many advantages including low damage to ecosystem, low operating costs, producing few harmful byproducts, and ease of use in different conditions (7). The bioremediation efficiency of microorganisms depends on various parameters, such as pollutant type, microorganism type, substrates, pollutant concentration, and environmental conditions. The selection of suitable species of microorganism with regard to the target contaminant has a dramatic impact on the bioremediation performance (8). So far, various microorganisms like Penicillium chrysogenum, Pseudomonas putida, Coprinellus radians, Tyromyces palustris, Gloeophyllum trabeum, Trametes versicolor, Acinetobacter sp., Pseudomonas sp., Bacillus subtilis, Ralstonia sp., and Microbacterium sp. have been applied for the bioremediation of POPs (9-11). Among them, Bacillus strains have shown high potential to degrade PAHs from surface soil (12). To the best of our knowledge, this is the first report demonstrating the Bacillus subtilis strain AHV-KH14 for the bioremediation of PHE from contaminated soil. Therefore, in the present study, a halotolerant bacterium Bacillus kochii strain AHV-KH14 was isolated from municipal compost, and then, used for the bioremediation of PHE from the contaminated soil.

16S rRNA

In this study, 16S ribosomal DNA (rDNA) sequencing using universal 27F and 1525R primers of the genomic DNA was used for the genetic identification of the isolated strain (14). The genomic DNA was extracted by boiling (15). Polymerase chain reaction (PCR) was carried out using Biometra thermocycler (Whatman, Biometra, Göttingen, Germany). A 50-μL solution containing Lyophilized 1-Taq Maxime PCR Premix (iNIRON Biotechnology, Korea), 41 μL double distilled H2O, 5 μL extracted DNA, and 2 μL of each primer (10 pmol) was applied to perform the reaction of amplification. The PCR conditions were as follows: initial denaturation step (95°C, 5 minutes) followed by 35 cycles of denaturation (94°C, 20 seconds), annealing (56°C, 30 seconds), polymerization (72°C, 90 seconds), and a final extension step at 72°C for 15 min. The purification of the PCR products was carried out using QIAquick Gel Extraction kit (Qiagen, Germany), and direct sequencing in both strands via an ABI PRISM 3730xl DNA Analyzer and the BigDye Terminator v3.1 Cycle Sequencing Kit chemistry (Applied Biosystems, USA) under contract by Bioneer Inc. (South Korea). The edition and assembling of Sanger sequence reads were carried out using DNA Sequence Assembler v4 (2013). The analysis of sequence data was performed using BLASTn from NCBI (http://www.ncbi.nlm.nih.gov), and classified using the EzTaxon server http://ezbiocloud.net/eztaxon. The phylogenetic analysis was performed in MEGA 6.0 by neighbor-joining algorithm using Kimura-2-parameter model and 1000-bootstrap replication.
Soil preparation
The soil samples required for bioremediation experiments were derived from a contaminated area in the Southwest of Iran, using soil cores from depth of 0-30 cm. Afterwards, the samples were air-dried, sieved (2 mm), shaken, and rinsed by acetone for the removal of organics, then, sterilized and finally, stored in plastic ice boxes. For synthetically contamination of the soil, the PHE was first solubilized in n-hexane, and then, added at desired levels of 50, 100, and 200 mg/kg to the samples. The solvent was evaporated under hood during 24 hours. The oxides present in the soil sample was determined using X-ray fluorescence analysis. According to results presented in Table 1, the soil sample was a sandy-clay type based on the texture. It should be noted that a portion of soil sample was kept without any processing as a naturally-contaminated soil sample.

Experimental procedure
Thirty grams of the synthetically contaminated soil was poured into 500-mL flasks. Operational variables including moisture content (50%, 100%, and slurry), PHE concentrations (50, 100, and 200 mg/kg), salinity (0.5%, 1%, 1.5%, and 2%), and inoculums size (5, 10, 15, and 20 mL), as well as their effects on the biodegradation rate, were investigated using one-factor-at-a-time method (15,16). Experimental samples were shaken at 180 rpm and 37°C and the amount of residual PHE and microbial growth were monitored weekly. Soil sample was extracted using EPA method 3550B for the PHE analysis. The residual was solved in 4 mL n-hexane for the quantitative analysis. The PHE concentration was determined by gas chromatography (GC) system (Chrompack CP 9001) equipped with a flame ionization detector (FID) using (HP-5) capillary column (30 m × 0.25 mm inner diameter, and 0.2 µm film thickness). Nitrogen was used as a carrier gas at a flow rate of 2 mL min⁻¹. The temperature program was as follows: the column temperature was held at 120°C for 1 minute, and then, raised to 240°C at a rate of 20°C min⁻¹, then, held for 1 minute. The injector and detector temperatures were set at 250 and 300°C, respectively. The PHE recovery rate was about 90% immediately after spiking. Considering the value of 10% as non-extracted portion of PHE in the soil, all raw data were multiplied by the factor of 1.1. All experiments were carried out in triplicate. The total petroleum hydrocarbons (TPH) values of the samples were determined with gas chromatography equipped with a flame ionization detector (GC-FID) (Chrompack CP 9001) using (HP-5) capillary column (30 m length, 0.32 mm inner diameter, and 0.2 mm film thickness). Bacterial density was evaluated based on the most probable number (MPN) (16). Finally, a set of bioremediation experiments was conducted on the unwashed soil sample under selected conditions. The naturally contaminated soil was also analyzed to determine the quantity and quality of hydrocarbons using gas chromatography mass spectrometry (GC-MS) analysis (Agilent 7890, USA) with HP-5MS capillary column (30 m × 0.25 mm × 0.25 mm film thickness, 5% phenyl-95% methyl siloxane phase).

Results
It was revealed that PHE biodegradation was inoculum size-dependent. The trend of PHE biodegradation against time indicated that the seed volume favored biodegradation rate until 10 mL, and higher values of seed size did not enhance the PHE removal. The results of this part are presented in Figure 1. Figure 2 shows the results of the effects of water content on the removal efficiency. The PHE removal rate at different moisture levels of 50%, 100%, and 200% (slurry) were 23%, 35.5%, and 62.5%, respectively. Moreover, the effects of soil salinity on the PHE biodegradation rate are shown in Figure 3. As shown in this figure, the removal efficiencies of PHE (initial concentration of 200 mg/kg) at the end of day 56 for salinity levels of 0.5%, 1%, 1.5%, and 2% were 62.5%, 64%, 64.5%, and 32%, respectively. The inhibitory effect of PHE on biomass was not considerable for the PHE concentrations of 50-200 mg/kg. Figure 4A shows the effect of PHE concentration on the process performance. The biodegradation rate for the PHE concentrations of 50, 100, and 200 mg/kg were 98%, 89%, and 64%, at the end of study period. Figure 5 represents the TPH mineralization

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<th>Table 1. Soil characteristics</th>
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Figure 1. Effect of seed size on the PHE biodegradation in the contaminated soil using Bacillus kochii strain AHV-KH14, moisture: 100%, and salinity: 0.5%.
rate during the bioremediation process. As shown in this figure, the TPH mineralization rate was lower than the PHE removal due to the conversion of mother compound (PHE) to biodegradation metabolites. In this regard, the mineralization rate reached 39% at the end of day 56, compared to the removal rate of 64% for the PHE at the same condition.

Discussion

Bacterial strain

Based on the 16S rRNA gene sequences and Neighbor-joining phylogenetic analysis (Figure 6), the bacterial isolate was characterized as Bacillus kochii strain AHV-KH14. This strain is a gram-positive, strictly aerobic, motile, catalase-positive, endospore-forming rods, and halo-tolerant bacterium. Its sequence was registered in Genbank under accession number of MH819661. Neighbor-joining method with MEGA 6.0 was applied for constructing tree. Bootstrap values over 70% (1000 replications) were indicated at each node. Escherichia coli (LN831047) was used as the out-group.

Bioremediation of PHE-contaminated soil

Effect of seed size

According to the results, inoculum size was effective on the biodegradation rate. At the initial steps and due to low bacterial density in the bioreactor, the startup of biological reaction was slow and vice versa. The PHE removal enhanced with increase of seed size till the value of 10 mL, and there was no significant increase in the biodegradation rate at higher levels (Figure 1). Adaptation period of biomass to PHE as well as low bacterial density in lag phase of growth leads to low biodegradation rate in the initial steps. By increasing the bacterial density at startup, the biodegradation rate could be enhanced somewhat, but from a break point which was 10 mL in the present study,
increasing the seed size will not affect the biodegradation rate due to lack of substrate and intensive competition, which in turn, leads to a sufficient metabolism (17,18). According to literature, a successful strategy to enhance the performance of slurry bioreactors treating hydrocarbons is bioaugmentation (19). Therefore, B. kochii strain AHV-KH14 was efficient in the biodegradation of PHE with seed size of 10 mL (OD_{600} = 1), corresponding to log MPN value of 5.43. The PHE biodegradation efficiencies (200 mg/kg) at the end of day 56 were 30, 35, 35.5, and 36%, for the inoculum sizes of 5, 10, 15, and 20 mL, respectively. As there was no significant difference between 10, 15, 20 mL of seed size (P > 0.05), the value of 10 mL was chosen for the remaining experiments, which is consistent with the results of the study of Zhao et al (20). Qin et al also demonstrated higher biodegradation rates of hydrocarbons for high levels of microbial inoculation (21).

Effect of moisture content

The soil moisture, which is affected by soil texture and organic matter contents, influences the biodegradation rate significantly, since the uniform distribution of substrate and nutrients and even biomass is controlled by water in soil (22). Bioavailability of substrate and its solubility is a major issue for the biodegradation of recalcitrant hydrocarbons by bacterial consortiums and as mentioned before, a uniform environment, in terms of oxygen, bacterial mass, macro and micro nutrients, and target hydrocarbon in a high-gradient medium, such as soil, is provided by moisture (23). Sorption and transportation of metabolic by-products into or out of the bacterial cell is also affected by water in the reaction medium. Consequently, a successful biodegradation is strongly related to water content of the soil (24). Since the soil/water ratio affects the mass transfer in the slurry phase bioreactor, therefore, a desired removal rate can be attributed to water content (25). The PHE biodegradation efficiencies at different moisture levels of 50%, 100%, and 200% (slurry) were 23%, 35.5%, and 62.5%, respectively (Figure 2). According to the results, the PHE removal was improved in the slurry phase, which facilitates the metabolism function of B. kochii strain AHV-KH14 through enhancing the bioavailability of PHE in soil, which is consistent with the results of some studies (26,27). Previous studies on the effect of water content on biodegradation rate indicated that high moisture content improved the hydrocarbon removal in a bioremediation experiment (28,29).

Effect of salinity

Effect of soil salinity on the PHE biodegradation rate was also studied. High salinity levels can adversely affect the biodegradation of organics in soil. Reports indicate that salinity, such as NaCl, can disturb both metabolism and growth simultaneously (30). The PHE removals (concentration of 200 mg/kg) at the end of day 56 were 62.5%, 64%, 64.5%, and 32% for salinity levels of 0.5%, 1%, 1.5%, and 2%, respectively (Figure 3). The results indicated the adverse effect of salinity level of 2% on the PHE biodegradation. Since in this study, bioremediation in saline environment was preferred due to similarity to real conditions and observation of no significant difference between removal data of other salinity levels (P > 0.05), remaining experiments were conducted in salinity level of 1.5%. Findings of the present study indicated that the biodegradation of PHE in the contaminated soil was correlated with salinity up to 1.5%, and in higher salinity values, the metabolic function of B. kochii strain AHV-KH14 was destroyed (31), which is consistent with the results of similar studies about effect of salinity on the bioremediation of hydrocarbon-contaminated soils (32).

Effect of PHE concentration

Concentration of PHE affected the biodegradation ability of B. kochii strain AHV-KH14. According to the obtained results, as the PHE concentration was increased, the removal efficiency by biodegradation was decreased, but the toxic level and inhibitory level were not observed in the PHE concentrations of 50-200 mg/kg (Figure 4A), which indicates the high metabolic function of the isolated strain (B. kochii strain AHV-KH14). Related studies have also verified the adverse effect of high concentrations of organic substrate on the microbial consortium and biodegradation efficiency (33). The removal efficiency of PHE for the initial concentrations of 50, 100, and 200 mg/kg (dry weight) were 98%, 89%, and 64%, respectively at the end of day 56. Researchers have specially reported that the elevated hydrocarbon concentrations are toxic and inhibit the biologic reactions (34). There are some claims for decreasing biodegradation with increasing the initial concentration of pollutant, when it would be adsorbed on the soil particle surface, which hinders the microbial growth and causes toxic effects, which decrease microbial activity, slow metabolism rate, and thus, exhausts the
survival ability of microbial populations (35). Despite of lower removal rate for the initial PHE concentration of 200 mg/kg, around 128 mg/kg substrate was degraded, which is considerable. This acceptable performance of B. kochii strain AHV-KH14 could be attributed to the appearance of a higher bacterial density when a high concentration of carbon source is present. As illustrated in Figure 4B, bacterial density for the initial PHE concentration of 200 mg/kg was significantly higher, compared to the two other levels.

**PHE mineralization**

Mineralization of PHE was studied using a comparative analysis of PHE and TPH. As can be seen in Figure 5, the TPH removal is lower than the PHE removal due to the conversion of mother compound (PHE) to biodegradation metabolites. In this regard, the mineralization rate reached 39% at the end of day 56, compared to the removal rate of 64% for PHE at the same condition, which is consistent with the results of the study of Jorfi et al (36).

**Bioremediation of unwashed soil**

Reliability of the isolated halo-tolerant strain (B. kochii strain AHV-KH14) in the biodegradation of various petroleum hydrocarbons in a real unwashed contaminated soil was also evaluated under selected conditions. As shown in Figure 7, a wide range of different n-alkanes were present in the polluted soil sample. The initial TPH concentration of 2350 mg/kg decreased to 1845 mg/kg (dry weight) (17.7% removal) during 56 days. Unwashed soil was contaminated with 1,2-benzenedicarboxylic acid, octadecane, hexadecane, pentadecane, nonadecane, tetradecane, and cyclotrisiloxane, and a low concentration of PAHs. The variety of contamination and high cumulative concentrations of different hydrocarbons in the unwashed soil were the main reasons for lower removal efficiency of the isolated strain compared to those of synthetically contaminated soil.

**Conclusion**

In this study, a halo-tolerant bacterium, Bacillus kochii strain AHV-KH14, was isolated from municipal compost, and then, used for the bioremediation of PHE from contaminated soil. A biodegradation efficiency up to 98% was observed for the initial PHE concentration of 50 mg/kg and salinity level of 1.5%, and the biodegradation efficiency decreased significantly with increasing salt concentration. The findings showed that the bioremediation efficiency decreased with increasing PHE concentration. The results also indicated that a variety of contaminants and high cumulative concentrations of different hydrocarbons in unwashed soil had a main role in decreasing the efficiency of bioremediation. A natural hydrocarbon-contaminated soil sample with TPH concentration of 2350 mg/kg was subjected to bioremediation using the selected conditions of operational parameters, and a biodegradation rate of 17.7% was obtained. Finally, GC–MS analysis was performed to study changes in PAHs, and it was concluded that the variety of contamination and high cumulative concentrations of different hydrocarbons in unwashed soil are the main reasons for lower removal efficiency of the isolated strain compared to those of synthetically contaminated soil.

**Acknowledgments**

The authors would like to gratitude Ahvaz Jundishapur University of Medical Sciences for its financial support (Grant No: 96s64).

**Ethical issues**

The authors hereby certify that all data collected during the research are as expressed in the manuscript, and no data from the study has been or will be published elsewhere separately.

**Competing interests**

The authors declare that they have no conflict of interests.

**Authors’ contributions**

All authors have contributed in the study design, experiments, data collection and analysis, and manuscript preparation. The final version of this manuscript was reviewed and confirmed by all authors.

**References**


