



Screening of diesel oil degrading bacteria from petroleum hydrocarbon contaminated soil

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Abstract

Four bacterial strains were isolated from diesel contaminated soil in TVS oil depot, Trichy, Tamil Nadu. From the soil sample four morphologically different bacterial species were isolated and the active strains were *Pseudomonas aeruginosa*, *P. putida*, *Arthobacter* sp and *Bacillus* sp. The diesel degrading efficiency of isolated organisms was tested in BH medium supplemented with diesel and DCPIP for 14 days. Among the four isolated species incubated and monitored in diesel supplemented medium for 14 days, the maximum diesel biodegradability was noticed in *P. putida* sp at fourth day itself and a TPC 64×10^5 CFU at a temperature of 30°C and pH 5. This study revealed a qualitative evaluation of potentials of hydrocarbonoclastic bacteria degradation of hydrocarbon. Thereby giving a measureable ability of these groups of bacterial possible use in hydrocarbon impacted soil remediation.

Keywords: *Pseudomonas putida*, petroleum hydrocarbon, DCPIP, Bioremediation.

Introduction

Petroleum based fuels are one of the most prevalent pollutants, particularly in developing countries (Joshi and Pandey, 2011). The widespread nature of petroleum products and their use is strongly associated with anthropogenic discharge of hydrocarbons into the environment (Bidoia *et al.*, 2010). Environmental pollution arising from petroleum leakages in storage tanks, spillage during transportation of petroleum products, deliberate discharge of petroleum products and various industrial processes is hazardous to soil and water ecosystems (Geetha *et al.*, 2013). This also results in huge disturbances of the abiotic and biotic components of the ecosystem (Okoh, 2006).

A major concern for petroleum hydrocarbon pollution is the presence of heavy compounds such as polycyclic aromatic hydrocarbons (PAHs), asphaltenes

and many branched compounds with twenty or more carbon atoms (Bidoia *et al.*, 2010).

Biological and non-biological approaches are being used for remediation of oil pollution. Bioremediation is one of the principle strategies for remediation, wherein the pollution can be removed by use of microorganism or by any biological process that uses microorganisms or their enzymes to return the environment altered by contaminants to its original condition (Olu- Arotiowa *et al.*, 2007).

Many species of microorganisms including bacteria, yeasts and fungi obtain both energy and tissue-building material from hydrocarbons. A wide range of studies have dealt with biotransformation, biodegradation and bioremediation of petroleum

hydrocarbons and interested in exploiting crude oil-degrading organisms for environmental cleanup has become central to petroleum microbiology. In the course of biological restoration of hydrocarbon contaminated soil, the main factors that affect the effect of remediation include the pH value, soil moisture, oxygen supply, the nutrient level, bacterial diversity and the temperature, among which the impact of petroleum hydrocarbon degrading bacteria on the effect is critical (Venosa and Xueqing, 2003; Chaillan *et al.*, 2006). The ability of many microorganisms in order to biodegradation of hydrocarbons has been studied by Liangli and Hungchen, (2009); Sarikhani *et al.* (2011); Ebrahimi *et al.* (2012) and Geetha *et al.* (2013). These methods are less expensive and do not introduce additional chemicals to the environment.

Hence in this investigation indigenous bacteria which degrade diesel and hydrocarbons are isolated from diesel contaminated site and screened for their hydrocarbon degradation efficiency. They were further characterized by morphological, cultural and biochemical techniques.

Materials and Methods

The diesel contaminated soil samples were aseptically collected from TVS oil depot, Trichy, stored in a sterilized container. The soil samples were placed in 4°C and transported to the laboratory immediately and maintained at 4°C until microbial analysis.

Diesel sample was collected from petrol bunk and the indicator 2, 6-dichlorophenol indophenol (DCPIP) used for the study was obtained from Lobachemie.

Bushnell-Hass (BH) medium (Atlas, 1995) was used as the enrichment media with 1% (v/v) diesel as the sole carbon source. 10 gm of the contaminated soil was added to 100 ml of BH medium and incubated at 30°C at 170 rpm. After 5 days of incubation, loop full of inoculum from Bushnell-Hass medium was streaked onto the nutrient agar plate and incubated at 30°C for 72 hours. Colonies with different morphological appearance were selected and purified in nutrient agar medium and were transferred to nutrient agar slants and stored for identification and further experimental studies. The cultures obtained were sub cultured monthly in nutrient agar medium.

The bacteria isolated were identified based on physical characterization and the biochemical tests outlined in Bergey's manual of determination bacteriology

(Buchanan and Warwick, 1974). Morphologically distinct colonies were isolated and purified by replicating on the same solid medium to obtain pure cultures.

The identified ten bacterial isolates were tested for their ability of diesel degradation and to determine the most efficient degrader. Each of the four bacterial cultures was inoculated to 100 ml of nutrient broth medium and incubated to 37°C for 24 hours at 150 rpm. The cultures were used as inoculum for the hydrocarbon degradation. 100ml of Bushnell-Hass (BH) medium was prepared with 1% diesel as sole carbon source. DCPIP at a concentration of 20 mg/l of the medium was used. The medium was sterilized at 121°C for 15 minutes. 1 ml each of the inoculum was added to each set of experiment and incubated at 150 rpm for 14 days. Control was also maintained.

The initial and final growth was determined by pour plate method using nutrient agar. The plates were incubated at 37°C for 48 hours. After the incubation period, colonies were counted and total microbial count/ml was calculated.

Diesel degradation was determined by the ability of microorganisms to utilize the diesel as sole carbon source which indicate the change in colour of DCPIP from blue to colourless. The indicator, when oxidized was blue and reduced was transparent. The reaction was observed visually till the end of incubation and also spectrophotometrically (600nm) at an interval of 4 days using Shimadzu model UV-3600 visible spectrophotometer according to Bharathi and Vasudevan (2001).

Results and Discussion

Microorganisms are diverse and are capable of utilizing contaminants as energy and carbon source to survive in natural environment (Singh *et al.*, 2010). Elimination of wide ranges of pollutants from the natural environment is required to enhance a sustainable development of the ecosystem with low ecological impacts (Selvakumar *et al.*, 2014). Microorganisms play a major role in the removal of contaminants taking advantage of their versatile catabolic activity to degrade or convert such compound to harmless substances.

In the present study four diesel degrading bacterial species *Pseudomonas aeruginosa*, *P. putida*, *Arthobacter* sp and *Bacillus* sp were identified in the diesel contaminated sites (Table 1). Similarly Bhasheer *et al.* (2014) reported the hydrocarbon

degrading efficiency of some isolated bacteria from oil polluted sites, such as *Acinetobacter* sp, *Moraxella* sp, *Bacillus* sp, *Vibrio* sp and *Alcaligenes* sp. This agrees with the work of Youssef *et al.* (2010) who reported

that the levels of hydrocarbon present in a contaminated site represent a nutrient rich environment where less-recalcitrant organic carbon may be limiting.

Table 1: Biochemical identification of bacterial isolates

S.no	Test	Strain-1	Strain -2	Strain -3	Strain -4
1	Oxidase	+	--	+	+
2	Catalase	+	+	+	+
3	Carbohydrate	+	+	--	+
4	DNA	--	--	--	--
5	Gelatin	--	--	+/--	+
6	Urea	--	--	+	--
7	Citrate	+	+	+	+
8	MrVp	+	--	+/--	+/--
9	Nitrate	--	+	--	+
10	Starch	+	+	--	+
11	Indole	--	--	--	+
12	Grams stain	+	+	--	--
13	Motility	+	+	+	+
14	Spore staining	+	NA	NA	NA
15	Strains identified	<i>Bacillus</i> sp.	<i>Arthobacter</i> sp.	<i>Pseudomonas putida</i>	<i>Pseudomonas aeruginosa</i>

Note: NA-Not Applicable; + Positive; - Negative

The degrading efficiency of isolated organisms was studied in BH medium supplemented with diesel and DCPIP. The experimental set up was kept for 14 days of incubation, in order to study the colour change of DCPIP, which is blue in colour and its oxidized form and colourless in its reduced form. This is due to reduction of the indicator by the oxidized product of hydrocarbon degradation which supports the facts that the isolates are potential hydrocarbon oxidizers (Selvakumar *et al.*, 2014). Species of *Pseudomonas*, *Bacillus*, *Micrococcus* and *Proteus* isolated from hydrocarbon contaminated site have been found by several authors to utilize hydrocarbon through oxidation of DCPIP (Roy *et al.*, 2002; Joshi and Pandey, 2011; Patil *et al.*, 2013; Adegbola *et al.*, 2014). The oxidation of DCPIP supports the facts that the isolates were potential hydrocarbon degraders. Absorbance at a wavelength of 600 nm was monitored for the organisms because a peak in absorbance was

observed at 600 nm as reported by Yoshida *et al.* (2001).

Among the four isolated species incubated and monitored in diesel supplemented medium for 14 days, the maximum diesel biodegradability was noticed in *P. putida* sp at fourth day itself and a TPC 64×10^5 CFU at a temperature of 30°C and pH 5. The degradation was also checked spectrophotometrically at an OD of 660nm; where in the OD reading was found to decrease in all the causes till the end of incubation period (Table 2). *Pseudomonas aeruginosa* and *Arthobacter* sp showed the diesel biodegradability at day 7 of incubation with a TPC of 61×10^5 CFU and 55×10^5 CFU, respectively. While *Bacillus* sp showed diesel biodegradability at day 10th of incubation of TPC 54×10^5 CFU. All these were carried at a temperature of 30°C and pH 5 (Table 3).

Table 2: Biodegradation of diesel by isolated organisms

S. No	Day of incubation	OD at 660nm			
		<i>Pseudomonas aeruginosa</i>	<i>P. putida</i>	<i>Arthobacter sp</i>	<i>Bacillus sp</i>
1	4 th day	0.424	0.454	0.395	0.305
2	7 th day	0.402	0.426	0.356	0.294
3	11 th day	0.373	0.408	0.321	0.282
4	14 th day	0.311	0.386	0.295	0.260

Based on oil utilization capacity, *P. putida* is the most active hydrocarbon utilizer in crude oil. Previous observations have identified the *Pseudomonas* genus as the most efficient among hydrocarbon degrading microorganisms (Banat *et al.*, 2000; Saadoun, 2002). Isolation and screening of microorganisms for their

efficiency in utilization of hydrocarbons before field trials is important in bioremediation process and the development of efficient techniques is an important tool in recommending different approaches for bioremediation of hydrocarbon polluted areas (Varjani *et al.*, 2013).

Table 3. Bacterial growth determination during biodegradation of diesel

S. No	Name of the organism	TPC/ml		Final day of incubation
		Initial	Final	
1	<i>Pseudomonas aeruginosa</i>	18 x10 ⁵ CFU	61 X10 ⁵ CFU	7 th day
2	<i>P. putida</i>	20 x10 ⁵ CFU	64 X 10 ⁵ CFU	4 th day
3	<i>Arthobacter sp.</i>	15 x10 ⁵ CFU	55 X 10 ⁵ CFU	7 th day
4	<i>Bacillus sp.</i>	12 x10 ⁵ CFU	54 X10 ⁵ CFU	10 th day

This study revealed a qualitative evaluation of potentials of hydrocarbonoclastic bacteria degradation of hydrocarbon. Thereby giving a measureable ability of these groups of bacterial possible use in hydrocarbon impacted soil remediation

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