Sunflower seed husk combined with poultry droppings to degrade petroleum hydrocarbons in crude oil-contaminated soil

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ABSTRACT
To remediate crude oil-contaminated soil, poultry droppings and sunflower seed husk were incorporated at various concentrations. Initially, the pH, moisture content, total organic carbon (TOC), total nitrogen content, and bacterial population of soil, poultry droppings, and sunflower seed husk were determined. The initial bacterial population was $35 \times 10^8$ colony forming units/g soil (dry weight basis). The total petroleum hydrocarbon (TPH) content in crude oil-contaminated soil was 7,048.4 mg/kg, in which 857.9 mg/kg accounted for eicosane. Naphthalene and acenaphthalene were the two polyhydroxy aromatic hydrocarbons present in soil at low concentrations. After the amendment, the pH, moisture content, microbial population, and TPH content of soil were determined on days 20, 40, and 60. The TPH concentration was considerably decreased in the treatment T4 wherein 250 g of soil was mixed with 125 g of poultry droppings and 125 g of sunflower seed husk. The indigenous bacterial population was also increased tremendously. The dehydrogenase enzyme activity was increased in the amended soil (T4: $0.74 \pm 0.06 \mu g TPF/g/h$). After the treatment of soil, the germination percentage and vigour index of maize, lady’s finger and tomato seeds were enhanced.

Keywords: Biodegradation, Petroleum hydrocarbon, Poultry droppings, Sunflower seed husk

1. Introduction

Crude oil is a naturally occurring, unrefined petroleum, and yellow-to-black liquid found within the Earth, and it is composed of hydrocarbons, organic compounds, and small amounts of metal [1]. Most of the components present in crude oil have low water solubility because of their low polarity and high substrate hydrophobicity. Soil contaminated with crude oil causes severe environmental issues because crude oil particles bind to soil particles and the contaminated soil particles become nonbioavailable for microorganisms, resulting in poor contact between food and microorganism and thus limiting the rate of mass transfer for biodegradation [2-4]. Furthermore, the contaminated soil possesses complex compounds, such as alkanes, benzenes, and methylbenzenes, and these compounds cannot be easily removed and thus leach into groundwater systems [5]. In addition, these compounds can penetrate through macro- and micropores in soil, thus limiting water and air transport necessary for organic matter conversion [6]. Furthermore, crude oil present in soil may impair aeration and infiltration of water into soil, as well as inhibit plant growth. Thus, the physical properties of soil, including water holding capacity, total porosity, and macro porosity, would be decreased [7]. Various physical and chemical methods employed for treating crude oil-contaminated soil are depicted in Fig. 1.

Considering the demerits of existing technologies, an alternative cheaper and environmentally friendly approach for enhancing petroleum hydrocarbon degradation is required. Studies have used animal and plant waste as a biostimulating agent to remediate crude oil-contaminated soils. For hydrocarbon removal, contaminated soil has been treated with sewage sludge (mixed in a ratio of 7:3) [8]; inorganic fertilisers (N, P, and K), cow dung, and palm kernel husk ash [9]; aged refuse from landfills [10]; rice husk combined with chicken manure [11]; sawdust [12]; and poultry manure [13]. Agricultural and industrial residues enhanced the colonisation of microorganisms due to favourable sorption properties and high mechanical resistance [14, 15].

The carrier or immobilisation matrix provides a protective microenvironment to microbial inoculants, either physically by providing a protective surface or pore space or nutritionally by providing specific substrates [16]. These materials should be nontoxic both...
to microbial inoculants and soil organisms and also be biodegradable while possessing such chemical characteristics that enable them to stay in soil long enough for the immobilised bacteria to perform their function.

For the proper maintenance of the biological equilibrium in soil, the physical, chemical, and biological characteristics of soil are crucial [17, 18]. Any disturbance in soil affects the biological equilibrium because it is susceptible to modifications caused in the ecosystem [18]. However, the stability and fertility of soil are predominantly determined by the activity of enzymes and distribution of microorganisms. Soil harbours numerous enzymes including the representative of all enzyme classes such as oxidoreductases, hydrolases, isomerases, ligases, and transferases. These enzyme groups are involved in the biogeochemical cycling [18] of organic matter for nutrient mineralisation and energy generation [19]. The dehydrogenase enzyme belonging to the oxidoreductase group is considered crucial because it occurs in all living microorganisms [20] and is closely connected with microbiological redox reactions [20]. The dehydrogenase enzyme plays a vital role in indicating pollution in soil, which is known by the absence of the enzyme in the polluted environment. A reduction in enzyme activity and microbial abundance is attributed to the recalcitrant nature and toxicity of compounds [21]. Hence, for remediating such soils, it is essential to increase the microbial profile with necessary enzyme systems capable of degrading pollutants.

North Chennai situated in Tamil Nadu, India, is home to several oil refineries and other heavy industries and thus is one of the most polluted areas in the state capital [22]. A preliminary site analysis showed that alkanes, benzene, toluene, ethylbenzene and xylene (BTEX), and polynuclear hydrocarbons (PAHs) were found in the soil and groundwater of the Manali region [23]. Because of oil spills in the Manali region, groundwater was heavily contaminated, leading to human health issues [24]. Thus, a cost-effective and environmentally friendly method for removing oil pollution from soil is required. Furthermore, microorganisms present in crude oil-contaminated soil should have a protective shield to degrade pollutants. However, for the efficient degradation of pollutants, the microbial population should be increased by supplementing nutrients to soil. Studies have attempted to remediate crude oil-contaminated soil by using residues of crops and animals for enhancing the nutrient content and increasing the microbial population. However, no study has examined the combined effect of poultry droppings and sunflower seed husk on the remediation of crude oil-contaminated soil. Hence, the present study aimed to remediate crude oil-contaminated soil with sunflower seed husk and poultry droppings.

2. Materials and Methods

2.1. Sample Collection

2.1.1. Soil

The soil sample used in this study was collected from Manali, Tamil Nadu, India (13.1636°N, 80.2586°E). The area is highly polluted by petroleum products containing a high amount of hydrocarbons, resulting in contaminated soil, surface, and sub-surface water. The soil was collected from outside the industrialised area and near the Santhankadu Lake. The soil sample was collected at a depth of 15 cm and stored in polythene bags. The soil sample was sieved through a 2-mm mesh to obtain a uniform texture of the soil.

2.1.2. Poultry droppings

Poultry droppings (faeces of chickens) were collected from a nearby poultry farming facility. The droppings were dried and ground into powder and stored in plastic containers at 4°C without freezing until further use.

2.1.3. Sunflower seed husk

The hull of sunflower seeds was bought from Sai Agros Ltd. (Parrys, Chennai, Tamil Nadu, India). The hulls were shredded into smaller pieces, sun dried for a week, ground into powder, and stored in plastic containers at 4°C for further study.

2.2. Characterisation of Soil, Poultry Droppings, and Sunflower Seed Husk

The pH was determined using a pH meter with a soil to water ratio of 1:2.5 and an equilibrium period of 30 min [25, 26]. Other parameters, namely moisture content [27], total nitrogen [28], total organic carbon (TOC) by wet oxidation [29], and phosphorus [30], were determined. The bacterial population was estimated using the soil extract agar medium [31]. The oil content of the polluted and remediated soil samples was determined through gas chromatography (GC) by using the toluene extraction method [32] and sonication water bath method [33]. The hydrocarbon content of the samples was determined using GC-mass spectrometry (MS)-Agilent 6890N GC, Santa Clara, USA, equipped with MS and a GC column (DB-5 MS 30 m × 0.25 mm and 0.25 μm). The heat programme of the column was started at 100°C (hold: 2 min), which was increased at a rate of 15°C/min until it reached 160°C and then at a rate of 5°C/min until it reached 280°C (hold: 10 min). Then, 1 μL of the extract was injected into the splitless mode.
percentage, vigour index (VI), and mean time to germination, were determined. Initially, the seeds were surface sterilised with 0.58% sodium hypochlorite for 1 min and subsequently washed with deionised water and air dried to prevent fungal infection. The seeds were grown in polystyrene Petri plates (100 × 15 mm in size). Each Petri plate was filled with 100 g of soil obtained after the degradation study (amended with poultry droppings and sunflower seed husk). For comparison, one set of experiment was conducted using soil without any crude oil contamination and amendments. Approximately 20 seeds were placed in each Petri plate, and for each treatment, five replications were maintained.

During the experimental period, the moisture content of the soil was maintained at 50% by spraying sterile water. The germination of seeds was monitored for 20 days, and for every two days, the number of seedlings was counted and removed from the Petri plate. Seed germination was determined based on the final germination percentage, number of days to first germination (delay of germination), and mean time to germination.

The mean time to germination was determined using the following formula [35].

\[ \text{Mean time to germinate} = \frac{\sum (n_i \times d_i)}{N}, \]

where 
\( n_i \) = Number of germinated seeds at day \( i \)
\( d_i \) = Incubation period in days
\( N \) = Total number of germinated seeds in the treatment

**Vigour Index**

VI was calculated on the 7th day [36] using the following formula:

\[ \text{VI} = \frac{\text{Total length of the seedling} \times \text{germination} \%}{\text{Total length of the seedling}} \]

The total length of the seedling was measured using a wooden scale, and the values were expressed in cm/plant.

### 2.5. Scanning Electron Microscope analysis

Bacterial species adsorbed on the amendments were examined using a scanning electron microscope (SEM; Quanta 250 FEG, Thermo Fischer Scientific) [37]. Crude oil-contaminated soil amended with poultry droppings and sunflower seed husk was removed from pots and fixed overnight with 5 mM phosphate buffer (pH 7.0) with 2.5% glutaraldehyde for an hour, followed by repeated washing with ethanol at different concentrations (10%, 20%, 40%, 60%, 80%, and 100%). The samples were dried and prepared using gold sputtering before imaging.

### 2.6. Statistical Analysis

Data were subjected to analysis of variance [38], and differences in significance among the means were compared using Duncan’s new multiple range test at a significance level of 5%.

### 3. Results and Discussion

#### 3.1. Characteristics of Soil, Poultry Droppings, and Sunflower Seed Husk

The samples of soil, poultry droppings, and sunflower seed husk were analysed for pH, moisture content, TOC, total nitrogen, total phosphorus, and bacterial population, and the results are listed in Table 1.

The pH of soil, poultry droppings, and sunflower seed husk
was 8.5, 7.2, and 5.8, respectively. The moisture content of poultry droppings was higher (18%) than that of soil (11.4%) and sunflower seed husk (6.5%). The TOC content of poultry droppings was higher (58.1%) than that of sunflower seed husk (52.1%) and soil (1.47%). The nitrogen content of soil, poultry droppings, and sunflower seed husk was 0.09%, 3.6%, and 1.5%, respectively. The phosphorus content of soil, chicken manure, and sunflower seed husk was 0.02%, 2.8%, and 0.14%, respectively. The initial bacterial population of soil was approximately $35 \times 10^8$ colony forming units (CFUs)/g soil (dry weight basis). Soil characteristics are considered crucial parameters for microbial growth and hydrocarbon degradation. The optimum pH for microbial growth and organic matter decomposition is 6.5-7.5 [39]. Moreover, soil microbes thrive better in neutral than in acidic soil [39]. An increase in soil pH towards the neutral condition indicates the presence of more favourable conditions for soil microbes [40]. Soil with a pH of 5.5-8.8 and a moisture content of 25%-80% enhance microbial growth and biodegradation [41]. However, sunflower seed husk with a moisture content of < 9.0% [42] can be a stable medium to protect microorganisms in crude oil-contaminated soil [43].

### 3.2. Total Petroleum Hydrocarbon Content in Soil

The TPH content in crude oil-contaminated soil collected from Manali was determined, and the results are shown in Fig. 2. The TPH content of crude oil-contaminated soil was 7,048.4 mg/kg. Among various petroleum hydrocarbons, the concentration of eicosane was the highest (857.9 mg/kg). The concentration of other petroleum hydrocarbon compounds, namely dotriacontane and hexacosane, was 751.2 and 604.9 mg/kg, respectively. The concentration of PAHs, namely naphthalene, phenanthrene, and acenaphthalene, was 0.6, 0.3, and 0.2 mg/kg, respectively. Crude oil contains more than 17000 compounds [44]. Crude oil is a mixture of aliphatic and aromatic hydrocarbons and is complex in nature [45]. However, the composition depends on the source and refining methods [45]. Soil collected from Anatolia had a TPH concentration of 48,000 ppm [46]. In the present study, TPH and PAH concentrations determined in crude oil-contaminated soil was in the range of milligram per kilogram. This result is in agreement with that of a previous study [47]. Furthermore, a study [47] reported that naphthalene, which is one of the 16 Environmental Protection Agency (EPA) PAHs, is present in the highest concentration in crude oil. However, in the present study, soil collected from contaminated sites contained naphthalene (0.20 mg/kg), phenanthrene (below detection level [BDL]), and acenaphthalene (BDL). Likewise, the naphthalene concentration in soil collected from Manali was higher; this finding is in agreement with those of previous studies [48]. The variation in the TPH concentration in crude oil-contaminated soil collected from different regions might be due to the fact that the extraction efficiency is affected by the ageing and clay content of soil as well as the high molecular weight of PAHs [48].

### 3.3. Effect of Amendments on Soil pH, Moisture Content, and Bacterial Population

The pH of crude oil-contaminated soil without amendment ranged
from 8.0 ± 0.11 to 8.5 ± 0.13 (T1) for various periods. However, the pH of soil after the addition of poultry droppings and sunflower seed husk ranged from 6.2 ± 0.08 to 6.5 ± 0.06 (T2), 6.4 ± 0.04 to 6.6 ± 0.08 (T3), and 6.3 ± 0.04 to 6.6 ± 0.06 (T4) for various periods (Fig. 3).

The moisture content of crude oil-contaminated soil after amendment with poultry droppings and sunflower seed husk was determined for regular time intervals, namely on days 0, 20, 40, and 60, and the results are illustrated in Fig. 3. The moisture content was higher in soil with amendment than in soil without amendment. Among different time intervals, the highest moisture content was recorded on day 60, and the moisture content of soil amended with sunflower seed husk and poultry droppings (T4) was higher (45% ± 0.16%) than that of soil without amendment (16% ± 0.25%).

Soil amended with poultry droppings and sunflower seed husk had a higher bacterial population than did soil without amendment (Fig. 3). The bacterial population was higher on day 60 of treatment (T4: 184 ± 4.0 × 10⁹ CFU/g soil) than for T1 (soil without amendment). The higher bacterial population found in soil amended with sunflower seed husk and poultry droppings might be attributed to nutrients provided by these amendments for the growth of microbial species. Organic amendments have been reported to enhance the multiplication of microorganisms that are capable of degrading and utilising crude oil as the source of carbon and energy [49, 50].

3.4. Degradation of TPH in Crude Oil-Contaminated Soil

The TPH content in crude oil-contaminated soil was determined on day 0, 20, 40, and 60, and the results are represented in Fig. 4. In all the treatments (with and without amendments), the degradation of TPH compounds was observed, and the maximum degradation was observed at day 60. The decane concentration was 763 and 674.5 mg/kg on days 0 and 60, respectively. The docosane concentration was 786.7 and 350.8 mg/kg on days 0 and 60, respectively. Likewise, the concentration of all the TPH compounds was significantly decreased in crude oil-contaminated soil. In all the treatments, a reduction in the concentrations of decane, docosane, dodecane, dotriacontane, eicosane, hexacosane, hexadecane, hexatriacontane, octacosane, octadecane, octane, octatriacontane, and tetracontane was observed on days 20, 40, and 60. The concentrations of tetracosane, tetradecane, tetracontane, and triacontane were BDL from day 20 for the treatments T1, T2, T3, and T4.

Fig. 4. Total petroleum hydrocarbon (TPH) content in crude oil contaminated soil amended with poultry droppings and sunflower seed husk. (a) 20 days after amendment (b) 40 days after amendment (c) 60 days after amendment.
Among PAHs present in crude oil-contaminated soil, naphthalene, acenaphthylene, phenanthrene, fluoranthene and pyrene were detected on day 0, and their concentrations were 0.6, 0.2, 0.3, 0.1, and 0.1 mg/kg, respectively. Other PAH molecules, namely 2-bromo-naphthalene, acenaphthene, fluorene, anthracene, chrysene, benzo[a]anthracene, benzo[a]pyrene, benzo[a]fluoranthene, benzo[ghi]perylene, benzo[a]h[1]anthracene, and indeno [1, 2, 3-cd] pyrene were at BDL. All the PHA molecules were found to be at BDL on days 20, 40, and 60 for all the treatments T1, T2, T3 and T4.

Sunflower seed husk is used to immobilise microbial culture for degrading crude oil-contaminated soil, and it provides a favourable environment for the survival of microorganisms in soil and improves bioremediation [51]. Poultry droppings, which have a high nitrogen content, can serve as soil amendments by adding organic matter as well as by improving the physical properties of soil and soil fertility by adding essential nutrients that improve moisture and nutrient retention. Furthermore, poultry dropping provides essential nutrients for the habitation of microbes, which can produce beneficial metabolites that indirectly reduce the amount of pollutants in soil [52]. Thus, essential nutrients from poultry droppings are crucial for necessary colonisation and survival in order to achieve better activity of microorganisms to mitigate recalcitrant hydrocarbon contaminants from soil. However, the potential of the bacterial community to remediate crude oil determines the treatment efficiency. An increase in the number of water-holding pores and a decrease in the number of water-transmitting pores [53] may be responsible for the increased moisture content in amended soil. Microorganisms present in soil require water for their metabolic activities. Soil water helps in chemical and biological activities of soil [54]. The organic matter present in soil improves binding and water retention [55] as well as promotes aggregate formation [56]. The presence of low organic matter in soil can be a cause of low water accumulation in soil without amendment [57].

Soil amended with 50% amendment material exhibited a higher hydrocarbon-degrading bacterial count compared with control. The high concentration of the amended material supported the growth of crude oil degrading-bacteria in soil; thus, significant stimulation of such bacterial populations was observed [58]. The extent of degradation of hydrocarbons was concurrently improved, indicating that microbial degraders were dependent on amendment supply that acts as a nutrient material. Among different amendment concentrations tested, 50% of the amendment material improved degradation of both aliphatic and aromatic hydrocarbons.

In the present study, higher degradation was observed in soil amended with sunflower seed husk and poultry droppings, and the highest degradation efficiency was shown by the treatment T4 (soil amended with 125 g of poultry droppings and 125 g of sunflower seed husk). Nutrient addition can stimulate the growth of petroleum-degrading microorganisms [59]. Indigenous microbial species were biostimulated by the amendments, and poultry litter is very high in nutrients, such as nitrogen, phosphorus, and potassium, and is used as a high-quality soil conditioner [60, 61]. Sunflower seed husk is composed of lipids, cellulose, and the reduced sugar pentose, and it provides a microenvironment for microorganisms present in soil to multiply and degrade pollutants [62]. In this study, PAH molecules were degraded in all the treatments, and PAH-degrading organisms, such as Sordariomycetes, Actinobacteria, Betaproteobacteria, and Gammaproteobacteria, are capable of degrading TPH molecules in soil [63]. The PAH benzopyrene could be degraded within 18 days by a novel strain that utilises this compound as the sole carbon source [64].

### 3.4.1. Dehydrogenase enzyme activity of crude oil-contaminated soil after treatment

The dehydrogenase enzyme activity of crude oil-contaminated soil treated with poultry droppings and sunflower seed husk was determined after 60 days, and the results are shown in Table 2. For comparison, garden soil (with no prior history of crude oil contamination) was also tested for the dehydrogenase enzyme activity.

Crude oil-contaminated soil amended with 125 g of poultry droppings and 125 g of sunflower seed husk exhibited the highest dehydrogenase enzyme activity of 0.74 ± 0.06 μg TPF/g/h, followed by control soil (without crude oil contamination and amendments) at 0.66 ± 0.08 μg TPF/g/h. However, the other two treatments (soil with crude oil contamination and amended with 25 g of poultry droppings and 25 g of sunflower seed husk and 75 g of poultry droppings and 75 g of sunflower seed husk, respectively) showed a dehydrogenase enzyme activity of 0.62 ± 0.06 and 0.65 ± 0.08 μg TPF/g/h, respectively. The lowest dehydrogenase enzyme activity of 0.08 ± 0.04 μg TPF/g/h was shown by soil with crude oil contamination and not amended with the substrates.

Compared with soil without amendment, increased dehydrogenase enzyme activity was observed in crude oil-contaminated soil amended with poultry droppings and sunflower seed husk. The addition of crop residues enhanced the soil enzyme activity [65]. Furthermore, organic amendments have been reported to increase the microbial population in soil and induce soil properties to regulate the soil microbial community [66, 67]. The added amendments supplied nutrients for the growth of indigenous oil-degrading microorganisms.

<table>
<thead>
<tr>
<th>Treatments</th>
<th>TPH produced (μg TPF g⁻¹ h⁻¹)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Control ( uncontaminated soil)</td>
<td>0.66 ± (0.08)</td>
</tr>
<tr>
<td>Contaminated soil</td>
<td>0.08 ± (0.04)</td>
</tr>
<tr>
<td>Soil + Poultry droppings (25 g) + Sunflower Seed husk (25 g)</td>
<td>0.62 ± (0.06)</td>
</tr>
<tr>
<td>Soil + Poultry droppings (75g) + Sunflower Seed husk (75g)</td>
<td>0.65 ± (0.08)</td>
</tr>
<tr>
<td>Soil + Poultry droppings (125g) + Sunflower Seed husk (125g)</td>
<td>0.74 ± (0.06)</td>
</tr>
</tbody>
</table>

Values in parenthesis show Standard Error (n = 5)
organisms in soil, which in turn increased the dehydrogenase enzyme activity [15, 68]. In the present study, the highest dehydrogenase enzyme activity was observed in the treatment incorporating sunflower seed husk and poultry droppings. Many studies have reported that organic amendments can increase enzyme synthesis [69, 70] and production by soil microorganisms as well as promote the release of enzymes that are either immobilised by clay and humic particles or entrapped within soil aggregates [71].

3.4.2. SEM analysis
On day 60 after the treatments, bacterial species adhered on the amendments, namely poultry droppings and sunflower seed husk, used for remediating crude oil-contaminated soil were analysed using SEM, and the results are shown in Fig 5. Adherence of bacterial cells was noticed on both the amendments. The adherence of bacterial species on organic amendments improved the tolerance of bacterial cells to toxic pollutants [72]. Many studies have been conducted using different amendments, such as biochar [73], goethite [74], and coconut coirpith [15], to immobilise bacterial species to remediate contaminated sites. However, the adhesion of bacterial species on organic matter could be achieved by polysaccharides produced by bacterial species [75].

3.5. Plant Toxicity Test
The effect of crude oil contamination on the germination of maize, tomato, and lady’s finger seeds was determined. Parameters, namely the percentage of germination, mean time to germinate, delay in germination, and VI, were determined.

In the present study, the percentage of germination for maize varied from 50% to 100% for various treatments. The lowest germination of 50% was recorded for the treatment T1 (crude oil-contaminated soil without amendments), whereas 100% germination was observed in control soil without crude oil contamination (T5). For the treatment T4 (crude oil-contaminated soil amended with 125 g of poultry droppings and 125 g of sunflower seed husk), 95% of germination was observed (Table 3).

The mean time to germination was 4 and 9 days, for seeds sown in uncontaminated soil (control) (T5) and crude oil-contaminated soil (T1), respectively. The mean time to germination for soil amended with 125 g of poultry droppings and 125 g of sunflower seed husk (T5) was 4 days, which was the same as that for uncontaminated soil (T5) (Table 3). Germination started on day 4 of sowing in all the treatments, except in crude oil-contaminated soil without amendments (T1). In crude oil-contaminated

![Fig. 5. Scanning Electron Micrograph (SEM) of (a) bacterial cells in soil amended with poultry droppings (b) bacterial cells in soil amended with sunflower seed husk.](image)

<table>
<thead>
<tr>
<th>S. No</th>
<th>Treatment</th>
<th>Germination* (%)</th>
<th>Mean Time to Germinate* (days)</th>
<th>Delay in germination* (days)</th>
<th>Vigour Index*</th>
</tr>
</thead>
<tbody>
<tr>
<td>T1</td>
<td>Contaminated soil</td>
<td>50</td>
<td>9.0</td>
<td>8</td>
<td>175.0</td>
</tr>
<tr>
<td>T2</td>
<td>Soil + Poultry droppings (25 g) + Sunflower Seed husk (25 g)</td>
<td>80</td>
<td>4.4</td>
<td>4</td>
<td>544.0</td>
</tr>
<tr>
<td>T3</td>
<td>Soil + Poultry droppings (75g) + Sunflower Seed husk (75g)</td>
<td>88</td>
<td>4.2</td>
<td>4</td>
<td>633.6</td>
</tr>
<tr>
<td>T4</td>
<td>Soil + Poultry droppings (125g) + Sunflower Seed husk (125g)</td>
<td>95</td>
<td>4.0</td>
<td>4</td>
<td>769.5</td>
</tr>
<tr>
<td>T5</td>
<td>Control (uncontaminated soil)</td>
<td>100</td>
<td>4.0</td>
<td>4</td>
<td>800.5</td>
</tr>
</tbody>
</table>

*Values represent mean of five determinations
Furthermore, the germination of the seeds was calculated on day 7. The seeds grown in crude-oil contaminated soil (T1) showed the lowest VI value of 175. The highest VI value of 800.5 was observed in the treatment T4. The dehydrogenase enzyme activity was found to be 0.74 ± 0.06 \( \mu \)g TPF/g soil/h. This could be possible due to the addition of organic amendments, namely poultry droppings and sunflower seed husk. The highest bacterial population of 184 ± 4.0 × 10^9 CFU/g soil was observed in the treatment T4. The dehydrogenase enzyme activity was found to be 0.74 ± 0.06 \( \mu \)g TPF/g soil/h. This could be possible because microorganisms have various enzyme pathways to degrade and utilise different hydrocarbons as the source of carbon and energy. Furthermore, after the treatment with the amendments, the phytotoxicity of soil was reduced. The treated soils enhanced the germination of the seeds within a short period. The germination percentage of plants is damaged by hydrocarbons, which in turn affect the metabolic transport and respiration rate [83]. In the current study, the experiment was conducted only for 60 days. If the experiment would have been performed for additional days, complete degradation of hydrocarbons might have occurred, and soil would have been suitable for growing crops. A field-level study is needed for remediating contaminated soils with locally available agricultural residues.

4. Conclusions

The results of the present study indicated that crude oil-contaminated soil could be remediated using sunflower seed husk and poultry droppings. A considerable reduction in TPH compounds was observed when 250 g of soil was mixed with 125 g of poultry droppings and 125 g of sunflower seed husk. The highest bacterial population of 184 ± 4.0 × 10^9 CFU/g soil was observed in the treatment T4. The dehydrogenase enzyme activity was found to be 0.74 ± 0.06 \( \mu \)g TPF/g soil/h. This could be possible because microorganisms have various enzyme pathways to degrade and utilise different hydrocarbons as the source of carbon and energy. Furthermore, after the treatment with the amendments, the phytotoxicity of soil was reduced. The treated soils enhanced the germination of the seeds within a short period. The germination of the crops tested was 100%, and the highest VI value was obtained. From this study, it can be concluded that hydrocarbon contamination of soil, which is a global issue, can be solved by the addition of organic amendments, namely poultry droppings and sunflower seed husk, which are easily available and inexpensive. Furthermore, remediated soil can be a suitable medium for growing plant species.

Author Contributions

S.B. (M.E. student) conducted the experiment and wrote the original draft. M.S.A. (Assistant Professor) supervised the student and edited the manuscript, and I.R. (Ph.D student) conducted scanning electron microscope studies.
References


