

# Bioremediation of heavy metal in crude oil contaminated soil using isolated Indigenous microorganism cultured with E coli DE3 BL21

Oluwamodupe Emmanuel Giwa, Francisca Omolara Ibitoye

**Abstract**— Soil contamination from crude oil have often been observed in recent years to increase the heavy metals and some hydrocarbon level in the environment from the soil to the plant and animals from the soil and hence the risk of bioaccumulation of this toxic compounds in the ecosystems which may threaten the human health in the endemic society. Bioremediation potency of individual indigenous bacteria isolated from soil polluted with crude oil was evaluated. Conventional method of identification was used to isolate and identify the indigenous microbes and the following were identified; *Bacillus* spp, *Staphylococcus aureus*, *Micrococcus* sp and *Pseudomonas aeruginosa*. The microbial accounts of total viable count after bio-augmentation  $4.3 \times 10^8$ ,  $2.7 \times 10^8$ ,  $2 \times 10^7$  and  $1.6 \times 10^7$  CFU g<sup>-1</sup> for *Pseudomonas aeruginosa*, *Bacillus* spp, *Micrococcus* sp and *Staphylococcus aureus* respectively. Each microbe was bio-amplified in an improvised bioreactor containing nutrient broth and re-inoculated into a 20 gram of sterilized polluted soil with crude oil to ensure mono-bioremediation. The heavy metal analyses were carried out using AS machine in the space of 60 days. There was a significant different at a probability level 0.05 in the degree of bioremediation in all the treatment using t-test, comparing the Bio-Augmented Mechanic Site Sample + PET system and Bio-Augmented Mechanic Site Sample. PET System *E. coli* DE3 BL21 aided in a synergistic relationship with each selected bacteria to achieve remediation of the polluted soil which may be associated with natural gene sharing and protein amplification by the PET system. Moreover, the gene in each isolated indigenous bacteria encoding bioremediation should be excised and cultured with PET system (*E. coli* DE3 BL21). The proteins harvested may be used directly to study its bioremediation potentials

**Index Terms**— Bio-augmentation, heavy metal, bioremediation, indigenous microbes, PET System *E. coli* DE3 BL21.

## I. INTRODUCTION

Heavy metals are universally found in the soil, however, heavy metal contamination in a wide sort of source including industrial waste, refuse dump site and land filling, demolition site wastes, excavation of the soil and weathering of parent materials and oil spillage has been commonly dispersed around the ecosystem due to increase in human activities. The ubiquitous of heavy metals in the ecosystem has been supposed to be due to human and natural undertakings, some which are naturally occurring and are available in an insoluble form which is not readily available for plants or microbial

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uptakes however, the anthropogenic sources are present in soluble and mobile reactive forms [1]. Most of these heavy metals are recycled in the environment via transportation either release from gas exhausts due to complete or incomplete combustion of crude oil, leading to air pollution and or by oil spillage common via shipping of crude oil and during milling of the ore from the core leading to water pollution. Another common practise is the uncontrollable discharge of spent oil into the soil at the mechanics site and dung hills. This scenario has been reported to further leads to bioaccumulation in water bodies into the aquatic life which are eventually eaten by Man.

Furthermore the alteration of the heavy metals could affect the ecosystem microbiota due to many factor among which include the change in the pH of the soil. Only the acidophilic microbe survives low pH, neutrophilic in the neutral pH and alkaliphile in the basic pH respectively. It is worthwhile to know that most of the heavy metals are toxic to the microbial metabolism. The specific heavy metal toxicity has been studied and findings reveals that copper disrupt cellular function and inhibit enzyme activities [2], [3], Cadmium damage nucleic acid, denature protein, inhibit cell division and transcription, inhibits carbon and nitrogen mineralization [3], [4], lead denatures nucleic acid and protein, inhibits enzymes activities and transcription [3], [5], [6], zinc death, decrease in biomass, inhibits growth [7].

However, microorganism has also been studied to employ varieties of methods to remediate the soil from these heavy metals polluted environment. These include Biosorption, bioaccumulation, biotransformation, and biomineralization. Bioremediation is an innovative technique for the removal and recovery of heavy metal ions from polluted areas, and involves using living organisms to reduce and/or recover heavy metal pollutants into less hazardous forms, using the activities of algae, bacteria, fungi, or plants. It has been employed for the removal of heavy metals from contaminated wastewaters and soils. These organisms help to detoxify hazardous components in the environment. The process can function naturally or can be improved through the addition of electron acceptors, nutrients, or other factors [1]. Soil polluted with heavy metals has been studies worldwide using consortium and specific microbes, however little or no knowledge has been established about the individual potentials in synergy with PET (*E. coli* DE3 BL21) systems.

## II. METHOD

### COLLECTION OF MATERIALS

The crude oil contaminated with soil was collected from the soil at Igbokoda, Ilaje local government area Ondo State which has a characteristic of black colours due to oil spillage

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and the soil surface was hardened. Loamy soil was also collected from Rufus Giwa Polytechnic Farm site. The sample was packaged into a sterile polytene bag and was brought to the Science Laboratory Technology Department, at Rufus Giwa Polytechnic Owo for evaluation. The sample was stored at adequate temperature before experimental work.

### PREPARATION OF SAMPLES

#### Preparation of soil sample

200g of the selected soil samples (crude oil contaminated soil, crude oil contaminated sample mixed with cowdung at 50:50, and loamy/black soil) was weighed using analytical balance into three different containers. 500ml each of distilled water was measured and added into the different container containing the polluted soil and it was mixed vigorously.

#### ISOLATION OF BACTERIA FROM SOIL SAMPLE

A mixture of 10g of the polluted soil samples were measured into 90ml of sterile water aseptically to make slurry, a drop of 0.1ml of each soil sample slurry were aseptically injected into separate sterile Petri dishes. Sterilized Nutrient agar of about 5ml was poured on the top of the sample to isolate total *Pseudomonas* count, total micrococcus count, Tryptosol agar for total *Bacillus* count and Manitol salt agar for total *Staphylococcus* count. The plates were incubated for 24hours at 37°C in an inverted position.

#### IDENTIFICATION OF BACTERIAL ISOLATES

The bacterial isolates were identified and classified using a combination of the methods as recommended by [8], [9], [10], [11], [12]. The identification of bacterial was based on cultural, morphological characteristics, staining properties, sugar fermentation and biochemical characteristics.

#### MAINTENANCE OF STOCK CULTURE

The pure culture were inoculated into sterile agar slant in McCartney bottles and incubated at 37°C for 48hours in other to maintain growth of microorganism, the stock cultures were transferred aseptically by sub-culturing from old slants to freshly prepared plates and incubated at 37°C for 24hours.

#### BIOSTIMULATION OF ISOLATED MICROBE

Two litres nutrient broth was used to stimulate the growth of each microorganism isolated from the soil polluted with crude oil. The soil samples polluted with crude oil was sterilized to kill all the living soil microorganisms. 20g of the sterilize soil sample was then mixed with each isolated microorganism respectively in 2 litres nutrient broth to form a slurry in a constructed bioreactor [13]. Each soil microorganism was allowed to colonize the soil polluted with crude oil and the heavy metals were monitored. Each isolated microbe was inoculated with PET system *E.coli* DE3BL21 strains in a closed system. The soil sample slurry containing broth was digested with acid H<sub>2</sub>SO<sub>4</sub> for heavy metal quantification analysis using A.A.S (Atomic Absorption Spectrophotometer) machine before, during and after the treatment. Treatment of metal by bacterial cells was calculated as the ratio of ions removal.

$$\% R (\%) = (A-B)/A \times 100$$

Where is the R=Removal ratio (%); A=Concentration of heavy metal ions in the primary solution ( $\mu\text{g mL}^{-1}$ ) and

B=Concentration of heavy metal ions in the treated solution ( $\mu\text{g mL}^{-1}$ ) [14].

### III. RESULTS

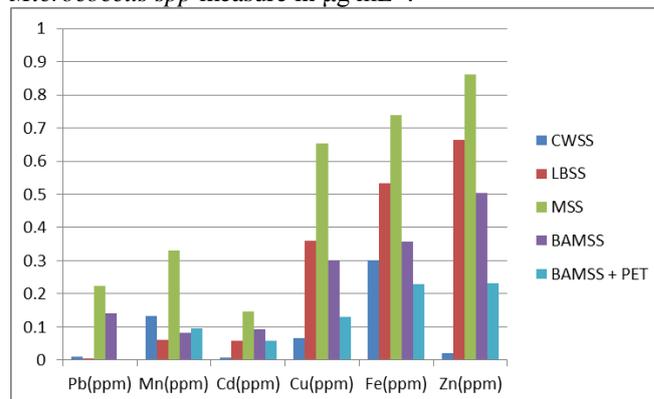
**Table 1: Morphology, staining, biochemical and sugar fermentation properties of bacteria isolate from crude oil polluted soil on various selective and general purpose agars**

Isolates/ Morphology	Isolate 1	Isolate 2	Isolate 3	Isolate 4
<b>Agar</b>	Nutrient Agar	Tryptose Soy Agar	Nutrient Agar	Mannitol Salt Agar
<b>Shape</b>	Rod	Rod	Cocci	Cocci
<b>Colour</b>	Green	Cream	yellow	White
<b>Margin</b>	Entire	Entire	Entire	Flat
<b>Consistency</b>	Butyrous	Dry	Slimy	Dry
<b>Opacity</b>	Opaque	Opaque	Transparent	Opaque
<b>Gram staining</b>	-ve	+ve	+ve	+ve
<b>Spore staining</b>	-ve	+ve	-ve	-ve
<b>Capsule staining</b>	-ve	-ve	-ve	+ve
<b>Catalase</b>	+ve	+ve	+ve	+ve
<b>Oxidase</b>	+ve	+ve	+ve	-ve
<b>Glucose</b>	+ve	+ve	+ve	+ve
<b>Lactose</b>	-ve	-ve	-ve	+ve
<b>Maltose</b>	-ve	+ve	+ve	+ve
<b>Fructose</b>	-ve	+ve	+ve	+ve
<b>Sucrose</b>	-ve	+ve	-ve	+ve
<b>Probable organisms</b>	<i>Pseudomonas spp.</i>	<i>Bacillus spp.</i>	<i>Micrococcus spp.</i>	<i>Staphylococcus spp.</i>

**Table 2: Total viable count of bio-augmented indigenous bacteria isolated from soil polluted with crude oil**

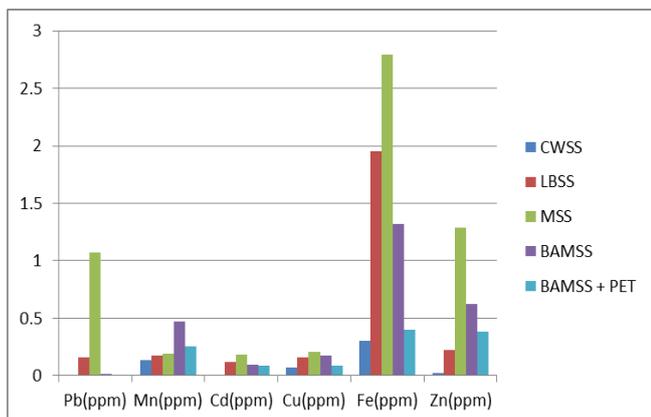
Isolates	TVC CFU/g
<i>Pseudomonas spp.</i>	$4.3 \times 10^8$
<i>Bacillus spp.</i>	$2.7 \times 10^8$
<i>Micrococcus spp.</i>	$2 \times 10^7$
<i>Staphylococcus spp.</i>	$1.6 \times 10^7$

**Figure 1: Heavy metal quantity of three selected soil treatment (loamy soil, mechanic site soil and bio-augmented mechanic site) that is inoculated/bio-stimulated with *Micrococcus spp* measure in  $\mu\text{g mL}^{-1}$ .**



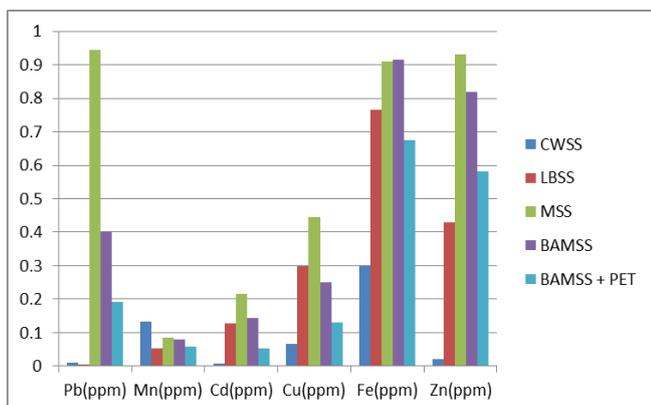
CWSS: Control Without Soil Sample, BAMSS: Bio-Augmented Mechanic Site Sample, MSS: Mechanic Site Sample, LBSS: Loamy/Black Soil Sample, BAMSS+PET: Bio-Augmented Mechanic Site Sample + PET SYSTEM

**Figure 2: Heavy metal quantity of three selected soil treatment (loamy soil, mechanic site soil and bio-augmented mechanic site) that is inoculated/bio-stimulated with *Bacillus spp* measure in  $\mu\text{g mL}^{-1}$ .**



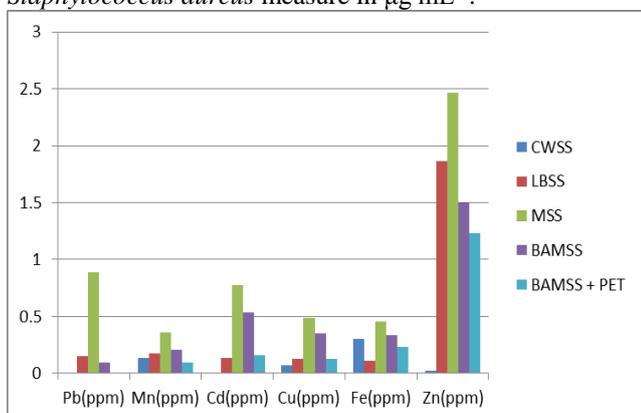
**CWSS:** Control Without Soil Sample, **BAMSS:** Bio-Augmented Mechanic Site Sample, **MSS:** Mechanic Site Sample, **LBSS:** Loamy/Black Soil Sample, **BAMSS+PET:** Bio-Augmented Mechanic Site Sample + PET SYSTEM

**Figure 3:** Heavy metal quantity of three selected soil treatment (loamy soil, mechanic site soil and bio-augmented mechanic site) that is inoculated/bio-stimulated with *Pseudomonas aeruginosa* measure in  $\mu\text{g mL}^{-1}$ .



**CWSS:** Control Without Soil Sample, **BAMSS:** Bio-Augmented Mechanic Site Sample, **MSS:** Mechanic Site Sample, **LBSS:** Loamy/Black Soil Sample, **BAMSS+PET:** Bio-Augmented Mechanic Site Sample + PET SYSTEM

**Figure 4:** Heavy metal quantity of three selected soil treatment (loamy soil, mechanic site soil and bio-augmented mechanic site) that is inoculated/bio-stimulated with *Staphylococcus aureus* measure in  $\mu\text{g mL}^{-1}$ .



**CWSS:** Control Without Soil Sample, **BAMSS:** Bio-Augmented Mechanic Site Sample, **MSS:** Mechanic Site Sample, **LBSS:** Loamy/Black Soil Sample, **BAMSS+PET:** Bio-Augmented Mechanic Site Sample + PET SYSTEM

#### IV. DISCUSSION

The figure 1 above shows the heavy metal quantity of three selected soil treatment (loamy soil, mechanic site soil and bio-augmented mechanic site) The loamy soil as a positive control shows the least contamination with Lead, Manganese, Cadmium, and Copper, however iron and zinc were a bit high in there composition. In the three experimental soil treatment, iron, zinc, copper, manganese were very low while cadmium and lead where under revealing limit. This finding is in line with the findings of [15], which used worms to bio-remediate soil polluted with diesel. Bacteria has proven to have a great biosorbents potentials due to their ubiquity, scope, skill to propagate in an optimize environment [16].

After 60 days of bio-augmentation of indigenous bacteria isolated from each experimental soil sample, most of the heavy metal recorded drastic reduction in the quantity.

*Micrococcus spp* shows a significant reduction in all the selected heavy metals in figure 1. *Micrococcus* has been shown to have a high sorption capacity on heavy metals such as lead and copper [17], *Bacillus* was very effective against lead, iron and zinc in figure 2, while *Pseudomonas aeruginosa* was more effective in the bioremediation of lead, copper, zinc and cadmium in figure 3. The ability of *Bacillus* and *Pseudomonas aeruginosa* to bioremediate soil polluted with hydrocarbon could be associated with the presence of gene responsible for alkane monooxygenase enzymes which is involve in the n alkane oxidation [18], [19]. *Pseudomonas sp* has also been studied to comprises of various plasmid capable of bioremediation among which include pDTG1 and pND15 [20], [21]. *Pseudomonas*, *Bacillus*, and *Micrococcus* species has been studied to show excellent sorption capacity and has been associated to their high surface-to-volume ratios and their numerous potential active chemisorption sites, such as the teichoic acid on the cell wall [22].

*Staphylococcus aureus*, though not necessarily a soil flora, showed a notable bioremediation effect on lead, cadmium and zinc (Figure 4). Ahamed [23] has reported isolation of *Staphylococcus aureus* as one of the petroleum hydrocarbon degrading microbes from soil and water samples of ship-breaking yards at Vatiary and Kumira coast in Chittagong. The ability of *Staphylococcus aureus* to grow on culture containing kerosene, diesel and engine oil as carbon source according to the observation of Ahamed [23] shows that the bacteria possess mechanism to utilize and breaks down the compounds. He further noted that the dominant species among the bacteria that grows on culture embedded with kerosene, diesel engine oil belonged to *Bacillus* followed by *Pseudomonas* and *Staphylococcus* species [23], hence our findings is in line with his own.

Furthermore, the introduction of PET system, *E. coli* DE3 BL21 cultured in a LB medium shows a significant rate of enhanced bioremediation in all the selected bacteria. This is not a common practice; however this could be as a result of natural sharing of plasmid and genes, between each of the selected bacteria and the PET system (*E. coli* DE3 BL21). The PET system has been known to be designed to be competent enough to integrate foreign genes in its self and produce up to 80% of the proteins for the incorporated genes. Figure 2 shows great results in the use of bio-augmented *Micrococcus spp* shows great result in lead (Pb), however the value is higher than the acceptable value by [24] of  $0.05 \mu\text{g mL}^{-1}$  and the guidelines for drinking water (WHO, 2004)

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concentration of  $0.01 \mu\text{g mL}^{-1}$ . However the combination with PET System shows a further drastic reduction in the lead concentration below and within the acceptable value. Similar trend was also observed with Cadmium (Cd), since the bio-augmented *Micrococcus spp* reduced the value from 0.146 to 0.093 and with the introduction of the PET system; it was further reduced to 0.057 which is within the acceptable value for USEPA [24] of  $0.05 \mu\text{g mL}^{-1}$ . Bio-augmented *Bacillus sp* in figure 3 shows great results in lead (Pb), with value of 0.013 and in synergy with the PET system, it further reduce to 0.003 both of which are within the acceptable specification by USEPA [24] of  $0.05 \mu\text{g mL}^{-1}$ . The value for cadmium was also recorded to be close within the acceptable specification in both soil treatments with bio-augmentation and introduction of PET system. *Pseudomonas* shows a good trend in reduction both in bio-augmented and the introduction of PET system, however the values obtain in lead (Pb) were both higher than the specification. Nevertheless, with the elongation of period from 60 days to 120 days, reduction to specification range may be achievable. Figure 3 and 4 also follows same trend with *Pseudomonas aeruginosa* showing great impact in the reduction of all the selected heavy metals. The bio-augmented *Staphylococcus aureus* displays high potential of bioremediation, reducing lead from  $0.887 \mu\text{g mL}^{-1}$  to  $0.093 \mu\text{g mL}^{-1}$  as shown in figure 4, a value close to the specification of acceptable range in the soil. The introduction of PET system further led to the reduction to lead. However cadmium was a little bit recalcitrant to remediation. All the soil treatment shows that the soil was not really polluted with copper since the values were below and within the acceptable value by USEPA [24] of  $1 \mu\text{g mL}^{-1}$  and  $2 \mu\text{g mL}^{-1}$  guidelines for drinking water (WHO, 2004).

### V. CONCLUSION AND RECOMMENDATION

Bio-augmentation is a potentially positive method to enhance indigenous microbes of soil polluted with crude oil to remediate the environments. PET System *E. coli* DE3 BL21 also aided in a synergistic relationship with each selected bacteria to achieve remediation of the polluted soil. However, the gene in each isolated indigenous bacteria encoding bioremediation should be excised and cultured with PET system (*E. coli* DE3 BL21). The proteins harvested may be used directly to study its bioremediation potentials.

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