

Assessment of Biodegradation Potential on Different Soil Particles

Yahaya, S. and *Bappa, A.M.

Department of Microbiology, Faculty of Life Sciences, Bayero University, Kano

Abstract: An explosion in the world's population has led to an increase in the demand for crude oil and its products resulting in an increased environmental pollution and thus leading to loss in biodiversity. Environmental reclamation by natural technique is believed to be eco-friendly and cost effective. This research investigated the biodegradation of crude oil supported on different soil particle sizes. Soil samples were randomly collected from different locations within Botanical garden of BUK and sieved into four (4) different particle sizes (0.6mm, 0.3mm, 0.15mm and 0.075mm) 200g of each of the sieved soil particle sizes were weighed and introduced into masonjar bottle. These were then uniformly contaminated with 60ml of crude oil. The set up were replicated in triplicate with each having two (2) controls. They were incubated aerobically at 37°C for 56 days. Enumeration of total aerobic heterotrophic bacteria (TAHB) was done on Nutrient agar (NA) and hydrocarbon utilizing bacteria (HUB) on Bushnell'Haas medium (BHM) supplemented with crude oil. The bacterial isolates were identified based on Grams reaction and biochemical tests. The degradation efficiency was confirmed by GC-MS analysis, which indicated that the microbial isolates utilized most of the crude oil components. The result shows that the mean microbial counts for both TAHB decrease from 5.93 ± 0.1 to $5.38 \pm 0.08 \times 10^5$ cfu/g during 56 days period of the study. The result also shows an increase in the mean counts of HUB from $0.00 \pm 0.0 \times 10^3$ to $3.74 \pm 0.03 \times 10^3$ cfu/g. Particle size A has the highest increase. The HUB identified were *Bacillus* spp., *Pseudomonas* spp., *Staphylococcus* spp., *Escherichia coli* and *Proteus* spp. The results indicate that larger particle size favors faster hydrocarbon biodegradation.

Keywords: Reclamation, Biodegradation, Hydrocarbon

INTRODUCTION

An increase in the world's population has led to an upsurge in the demand for petroleum and petroleum products, which has apparently become a source of pollution to the environment (Akoachere *et al.*, 2008). The wide spread contamination of most of arable lands, creeks, swamps and natural source of water with petroleum and petrochemical products particularly in the Niger Delta region of Nigeria, is due to an increasing petroleum exploration, refining and other related industrial activities (Okpokwasili and Odokuma, 1990; Odokuma and Okpokwasili, 1993). The contamination of these habitats poses major public health and socio-economic hazard which most often has developed into impetuous protestation between some of the oil companies and the surrounding communities (Akpe *et al.*, 2015).

Soil texture and soil structure are both unique properties of the soil that have a profound effect on the behavior of soils, such as water holding capacity, nutrient retention and supply, drainage, and nutrient leaching (Silva *et al.*, 2000). The texture of soil is determined primarily by the size of the mineral particles that make up the soil (Oades, 1993). The mineral particles of a soil are present in a wide range of size and in association with organic matter (Tisdall, 1994). Fine earth fraction includes all soil particles that are less than 2 mm. Soil particles within this fraction are further divided into three (3) separate size classes, which includes sand, silt, and clay. The size of sand particles range between 2.0 and 0.05 mm; silt, 0.05 mm and 0.002 mm; and clay, less than 0.002 mm (Silva *et al.*, 2000).

Soil is a rich source of microorganisms capable of degrading hydrocarbons and residual oil (Atlas and Bartha, 1999). The rate of crude oil biodegradation in the soil seems to be rapid (Allamin, 2014). This may be due to the fact that the microorganisms in the soil have efficient ability in utilizing the residual crude oil as a source of carbon and energy. Indigenous and adapted microorganisms are more efficient for biodegradation of oil pollutant Akoachere *et al.* (2008).

¹*Correspondence Author:

Email: muhammadalkalibappa@gmail.com

Phone: 07033175716

Consequently, there is an increased interest in promoting environmental methods in the process of cleaning oil polluted sites. These methods are less expensive and do not introduce additional chemicals to the environment compared to the physicochemical methods (Akpe *et al.*, 2015).

Bioremediation using microorganisms capable of detoxify oil contaminated sites offers a very feasible alternative for an oil spill response due to the fact that majority of the molecule in the crude oil are biodegradable.

Many research activities have been conducted on bio-stimulation and bio-augmentation techniques of bioremediation, but not much has been done on natural attenuation. Therefore, this research which employed natural attenuation technique was aimed at studying the performance of biodegradation on different soil particle sizes which could entails decision when planning for hydrocarbon cleanup.

MATERIALS AND METHODS

Sampling Site and Sample Collection

Soil sample were randomly collected from different locations within the botanical garden of Bayero University Kano, using soil auger at a depth of 10cm (Onifade and Abubakar, 2007). The sample were transported to the microbiology laboratory in a black polythene bags which were mixed together and allowed to air dried.

Bonny light crude oil was collected from Kaduna Refinery and Petrochemical Company (KRPC), Kaduna, Nigeria.

Sieve Analysis of the Soil Sample

The air-dried soil samples were sieved into four (4) different particle sizes using sieve number; 0.6mm, 0.3mm, 0.15mm and 0.075mm respectively. The sieves were stacked with the largest opening at the top and smallest opening at the bottom. The soil was poured into the top of the sieve and covered. The set of sieves were put into a mechanical shaker for 15 minutes. When the shaker stopped, the stacks of sieves were removed. The retained soil samples on each

sieve were poured into a sterile plastic container for further analysis (Sandra, 2011).

Biodegradation Experimental Setup

Exactly two hundred (200) grams each of the sieved soil particle sizes; (0.6mm, 0.3mm, 0.15mm and 0.075mm), were weighed and introduced into a masonjar bottles. Sixty mills (60ml) of crude oil was also added (to serve as the sole carbon and energy source), and allow to diffused through for each of the four different soil particle sizes. It was then shaken gently to ensure homogeneity. These setups were replicated in triplicated for each of the (4) different soil particle sizes, with each having two (2) different controls. These were incubated aerobically with perforated aluminum foil on top of the bottles at room temperature for fifty-six (56) days. Finally, a total of twenty (20) sets were generated as presented below;

A₌ 200g (0.6mm) soil particle size + 60ml crude oil (3×test)

B₌ 200g (0.3mm) soil particle size + 60ml crude oil (3×test)

C₌ 200g (0.15mm) soil particle size + 60ml crude oil (3×test)

D₌ 200g (0.75mm) soil particle size + 60ml crude oil (3×test)

1st CONTROL = 200g particle size + 60ml crude oil + Sterilization (1x for each particle size)

2nd CONTROL = 200g particle size + 60ml crude oil + 4% formaldehyde (1x for each particle size)

Microbial counts and pH of the samples were monitored four times at 14 days interval. Also, at 56days of the experiment, the residual crude oil was extracted from the soil using Soxhlet method. The level of microbial degradation of the crude oil was determined using Gas chromatography/mass spectrophometric analysis.

Soxhlet Extraction

Fifty grammme (50g) of the contaminated soil sample was weighed and wrapped in 11mm Whiteman filter paper. It was gently inserted into the timble.120mills of petroleum ether and 60 mills of diclromethane were placed into the distillation flask.

The temperature was adjusted to 60°C and setup was allowed to run until the solvent through distillation process washes the soil completely. The extract was allowed to air dry for 24 hours before taken for GC-MS analysis.

Enumeration, Isolation and Identification of Microorganisms

Enumeration of Total Aerobic Heterotrophic and Hydrocarbon Utilizing Bacteria

Enumeration of total aerobic heterotrophic bacteria (TAHB) was done on nutrient agar (NA) and Hydrocarbon Utilizing Bacteria (HUB) on Bush Knell Haas agar (BH) supplemented with crude oil. One (1) gram each of the soil samples was weighed and placed into test tubes containing 9ml of distilled water the test tubes were shaken vigorously in order to dislodge the microorganisms that adhered to the soil particles. The content of the tube were serially diluted from 10^1 to 10^4 . $10 \mu\text{L}$ of dilution factor 10^{-4} were plated in triplicates on to sterile Nutrient agar (NA) and (10^{-2}) on Bush knell Haas agar (BH) respectively. The plates were incubated at room temperature ($30 \pm 2^\circ\text{C}$) for 48 hours and 7 days for NA and BH respectively (Allamin, 2014). The colonies, which developed on the plates were counted and recorded as colony forming units per gram (cfu/g) of soil after Bashir (2012). Pure cultures of the isolates were obtained by repeated sub-culturing on media used for primary isolation. The pure isolates were maintained on agar slants for further characterization and identification (Abioye *et al.*, 2012; Allamin, 2014).

Morphological and Biochemical Identification

The phenotypic and biochemical characteristics used in identifying bacteria isolated included Gram staining, colonial appearance, motility, spore forming, urease, catalase, indole, oxidase citrate and methyl red Voges proskauer. The tests were performed using the methods of Cheesbrough (2006).

Measurement of Rates of Crude Oil Biodegradation

Gas chromatographic/ Mass Spectrophotometric Analyses (GC-MS) of Oil Extracts

At fifty- six (56) days of the experiment, the residual crude oil were extracted from the soil using Soxhlet method. The level of microbial degradation of the crude oil was determined using Gas chromatography/ mass spectrophotometric analysis as described by Ijah (1998); Ibrahim *et al.* (2009).

Determination of Physicochemical Properties of the Soil Sample

Total carbon was determined by dichromate wet oxidation method of Walkey and Black as modified by Dyan *et al.* (1999). Nitrogen content was determined using the micro kjeldah digestion method of Brady and Weil (1999). Particle size was determined using method described by Sheldrick and Want (1993). Moisture content was determined using the method of Morris (1999).

Statistical Analysis of Data

The data in this study were subjected to one-way analysis of variance (ANOVA) using instant GraphPad software version 3.05. Relation between variables and comparison of means were tested for level of significance at $p \leq 0.05$.

RESULTS AND DISCUSSION

The soil sample used for the study was composed of 72% sand, 22% silt and 7% clay. The textural class of the soil was sandy, clay and loam. The total nitrogen content of the soil sample was 0.18%, the organic carbon content was 1.5% while the phosphorus content 32.53 mg/kg. The moisture content was 1.02% (Table 1). The presence of these limiting nutrients (C, N and P) in the soil samples analyzed in this study is in consonance with previous reports (Okoh, 2006).

The low level of nutrients in the soil sample could have been caused by leaching or erosion.

This agrees with the findings of Romanus *et al.* (2015) who made similar observation of low percentage of carbon, nitrogen and phosphorus content of a garden soil which was used for bioremediation studies. Soils with appreciable content of these elements have been implicated in bioremediation studies.

The pH of the soil sample monitored during the period of study ranged from 6.55 to 7.66 as shown in (Table 2). The result of the pH observed in this study is within the favorable range for biodegradation of crude oil in polluted soil. Similar observations have been reported by Romanus *et al.* (2013) and Akpe *et al.* (2015).

Table 1: Physicochemical Parameters of the Soil Sample

Sample	Physicochemical Parameters of the soil sample								
	pH	P (mg/kg)	O.C %	N%	MC%	% Sand	% Silt	% CAY	TC
A	7.01	32.53	1.58	0.18	1.02	72.00	22.00	7.00	Sandy, Clay, Loam

Key; PH, O.C=ORGANIC CARBON, N= NITROGEN, MC=MOISTURE CONTENT, TC=TEXTURAL CLASS

Table 2: pH of Crude Oil Polluted Soil Having varying degrees of Particle Size.

Time(days)	Sample A	Sample B	Sample C	Sample D
0	6.50	6.53	6.74	6.79
14	6.73	6.54	6.63	6.50
28	7.20	7.52	7.66	7.64
42	6.84	6.85	6.93	6.81
56	6.84	6.85	6.93	6.81

KEY: A=0.66mm particle size B=0.3mm particle size C=0.15mm particle size D= 0.75mm particle size

The results of the Total aerobic heterotrophic bacteria (TAHB) indicated that the mean microbial counts of TAHB reduced from zero to 56 days of the study (Table.3). The highest reduction was observed to be in soil with the largest particle size (0.6mm) which ranged from $5.93 \pm 0.01 \times 10^5$ cfu/g to $5.5 \pm 0.06 \times 10^5$ cfu/g. This could be due to lower water retention capacity of the soil particle size which greatly enhanced hydrocarbon loss. This is in agreement with findings by Sandra (2011), who reported higher loss of hydrocarbons in soil with larger grain size. It also agreed with the study of Kogbara *et al.* (2015) who observed faster rate of hydrocarbon loss in sandy soils. The result also indicated that there were no wide difference in the reduction of mean microbial counts among particle sizes B, C and D. This could be due to them having higher surface area which impede on

the ability to efficiently degrade hydrocarbons. This is in line with the work reported by Cui *et al.* (2011); Sandra (2015) and Romanus *et al.* (2015).

The decrease in the heterotrophic microbial counts could be as a result of adaptation to the polluted environment as well as the toxic effect of crude oil on the microbial population as previously suggested Akoachere *et al.* (2008) and Mbah *et al.* (2009).

Generally, the total heterotrophic bacterial counts were found to be higher than the crude oil utilizing bacterial in all the samples. This could be as a result of nutrient limitation in the enumeration media for crude oil utilizes. This finding is consistent with work Romanus *et al.* (2015), who equally reported higher heterotrophic bacteria counts than hydrocarbon carbon counts in crude oil polluted soils.

Table 3: Total Aerobic Heterotrophic Bacteria Count of the Soil Sample ($\times 10^5$ cfu/g)

Samples	Day 0	Day 14	Day 28	Day 42	Day 52
A	5.93 \pm 0.01 ^{*a}	5.86 \pm 0.02 ^{**}	5.62 \pm 0.04 [*]	5.87 \pm 0.05 [*]	5.55 \pm 0.06 [*]
B	5.83 \pm 0.03 ^{*bc}	5.78 \pm 0.02 ^{*a}	5.65 \pm 0.05 [*]	5.40 \pm 0.06 [*]	5.51 \pm 0.05 [*]
C	5.85 \pm 0.06 ^{*ab}	5.72 \pm 0.04 ^{*a}	5.60 \pm 0.05 [*]	5.53 \pm 0.04 [*]	5.42 \pm 0.05 [*]
D	5.75 \pm 0.04 ^{*c}	5.62 \pm 0.04 [*]	5.60 \pm 0.06 [*]	5.51 \pm 0.03 [*]	5.38 \pm 0.08 [*]
E	0.00 \pm 0.00	0.00 \pm 0.00	0.00 \pm 0.00	0.00 \pm 0.00	0.00 \pm 0.00
F	0.00 \pm 0.00	0.00 \pm 0.00	0.00 \pm 0.00	0.00 \pm 0.00	0.00 \pm 0.00

KEY; A=0.6mm, B=0.3mm, C=0.15mm, D=0.075mm, E = Control, F = Control, Cfu/g=colony forming units per gram, values were Mean \pm SD of triplicates determinations. Values with single * = not significantly different. ** = significantly different, the same alphabets combination along the rows were not significantly different ($p < 0.05$).

Table 4: Hydrocarbon Utilizing Bacterial Counts of the Soil Sample ($\times 10^3$ cfu/g)

Samples	Day 0	Day 14	Day 28	Day 42	Day 56
A	0.00 \pm 0.00 [*]	3.23 \pm 0.06 ^{**}	3.42 \pm 0.09 [*]	3.65 \pm 0.07 ^{*a}	3.74 \pm 0.03 ^{**}
B	0.00 \pm 0.00 [*]	3.01 \pm 0.02 ^{*ab}	3.21 \pm 0.07 [*]	3.56 \pm 0.04 ^{abc}	3.57 \pm 0.03 ^{*ab}
C	0.00 \pm 0.00 [*]	3.03 \pm 0.05 ^{*ac}	3.18 \pm 0.11 [*]	3.58 \pm 0.02 ^{*bd}	3.56 \pm 0.02 ^{*ac}
D	0.00 \pm 0.00 [*]	2.87 \pm 0.11 ^{*bc}	3.18 \pm 0.11 [*]	3.49 \pm 0.02 ^{*cd}	3.57 \pm 0.01 ^{*bc}
E	0.00 \pm 0.00	0.00 \pm 0.00	0.00 \pm 0.00	0.00 \pm 0.00	0.00 \pm 0.00
F	0.00 \pm 0.00	0.00 \pm 0.00	0.00 \pm 0.00	0.00 \pm 0.00	0.00 \pm 0.00

KEY; A=0.6mm, B=0.3mm, C=0.15mm, D=0.075mm, E = Control, F = Control, Cfu/g=colony forming units per gram, values were Mean \pm SD of triplicates determinations. Values with single * = not significantly different. ** = significantly different, the same alphabets combination along the rows were not significantly different ($p < 0.05$).

Results of the hydrocarbon utilizing bacteria (HUB) counts indicated that there were no microbial activities at zero day of the study (Table 4). This may suggest period of adjustment and adaptation to the new environment by the indigenous microorganisms. It was observed that HUB counts generally increased from 14 to 56 days of the study. This could have resulted from the adjustment of the indigenous microorganisms to the new environment and their ability to synthesize other metabolites suspected to have been generated during the biodegradation period and conforms with the findings by Akoachere *et al.* (2008). The larger particle size 0.6 mm (A) has the highest HUB counts. This could have resulted from the lower surface area which provided better aeration that greatly

enhanced the ability of the microbes to attack hydrocarbons and utilized them as sole carbon and energy source for their metabolic activities. This is consistent with the study of Kogbara *et al.*, (2015) who reported that sand with larger particle sizes favors faster hydrocarbon degradation due to enhance aeration and thus improved microbial survival and activity during crude oil contamination as well as in the course of remediation. This result also agreed with the existing literature (Romanus *et al.*, 2015; Sandra, 2015).

Analysis of variance for HUB showed significant difference ($P \leq 0.05$) between particle sizes A, B, C and D. However, there were no significant difference among B, C and D. This could be due to their higher specific surface area.

Table 5: Morphological and Biochemical Characteristics of Hydrocarbon Utilizing Bacteria Isolated from the soil sample

Isolate	Biochemical Characteristics																Bacteria
	GR	Shape	Spor e	CAT	OXI	MR	VR	UR	IND	CIT	GLU	LAC	SUC	TSI H ₂ S	GAS	MOT	
1	-	Rods	-	+	-	+	-	+	+	-	+	-	-	+	+	+	<i>Proteus vulgaris</i>
2	+	Rods	+	+	-	-	+	+	+	+	+	-	-	-	-	+	<i>Bacillus cereus</i>
3	-	Rods	-	+	+	+	-	-	-	+	+	-	-	-	-	+	<i>Pseudomonas putida</i>
4	-	Rods	-	+	-	+	-	-	+	-	+	+	+	-	+	+	<i>Escherichia coli</i>
5	+	Cocci	-	+	-	-	-	-	-	-	-	-	-	-	-	-	<i>Staphylococcus aureus</i>

Keys: + =positive, - =negative, CAT= Catalase, OXI= Oxidase, MR= Methyl-red, VR=Voges Proskauer, UR= Urease, IND = Indole, CIT= Citrate, GLU=Glucose, LAC=Lactose, SUC=Sucrose, and MOT=Motility, TSI=Triple sugar ion test.

The bacteria isolated from this study were *Bacillus cereus*, *Proteus vulgaris*, *Pseudomonas putida*, *Escherichia coli* and *Staphylococcus aureus* (Table 5). The predominant form were rod shape bacteria and cocci. The Gram-negative bacteria were mostly isolated. These and other bacterial species isolated from the soil sample are shown in Table 7. The result also correlates the report of previous workers (Esumeh *et al.*, 2009) who isolated more of Gram negative organisms suggesting that they are better degraders of crude oil when compared with their Gram positive counterparts. The higher ability of Gram-negative bacteria to

utilize crude may not be unconnected with the possession of plasmid-borne or chromosomal genes involved in hydrocarbon degradation and porins in their cell wall which helps in the uptake of certain substances by the cell or extrusion of others which may be harmful (Vahaboglon *at al.*, 1996; Jørgensen *et al.*, 2000; Akpe *et al.*, 2013). The crude utilizing bacteria identified in this study have been identified and implicated in crude oil biodegradation by several investigators (Antai, 1990; Allamin 2014; Romanus *et al.*, 2015; Ajayi *et al.* 2008).

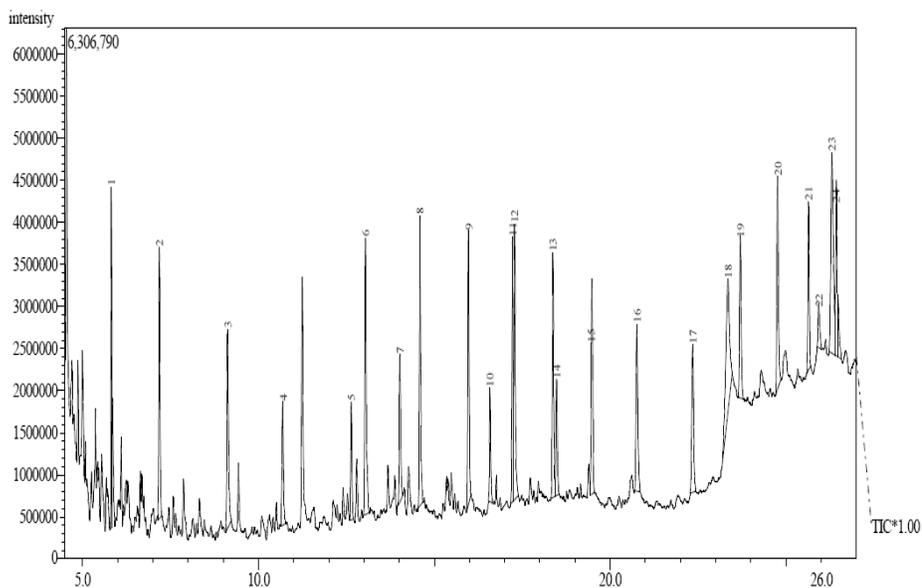


Figure 1: Gas Chromatogram of the Raw Bonny Light Crude Oil (Negative Control).

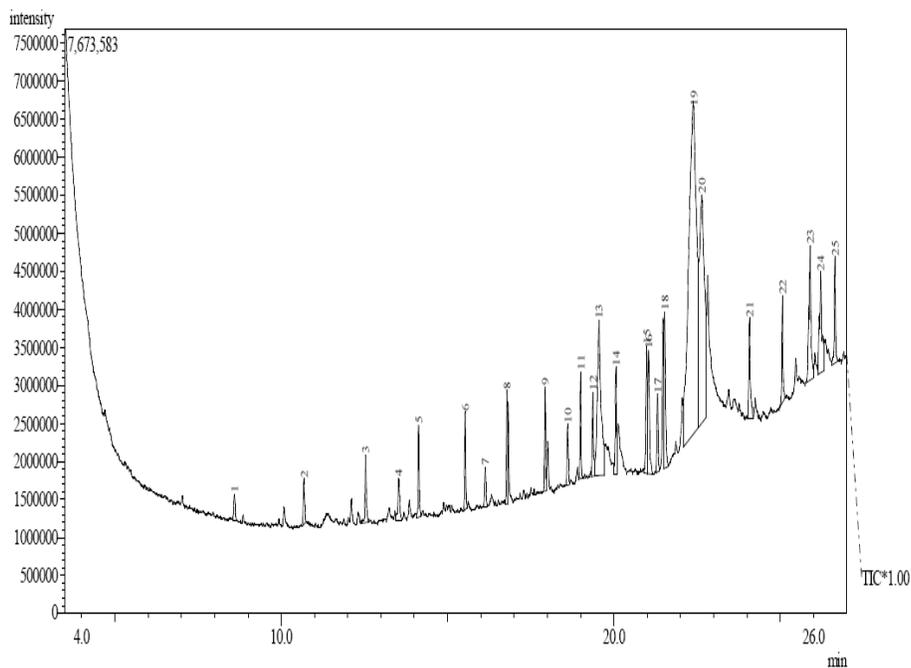


Figure 2: Chromatogram of Soil Particle Size A (0.6mm) Degraded by Indigenous Bacteria after 56 Days of Incubation.

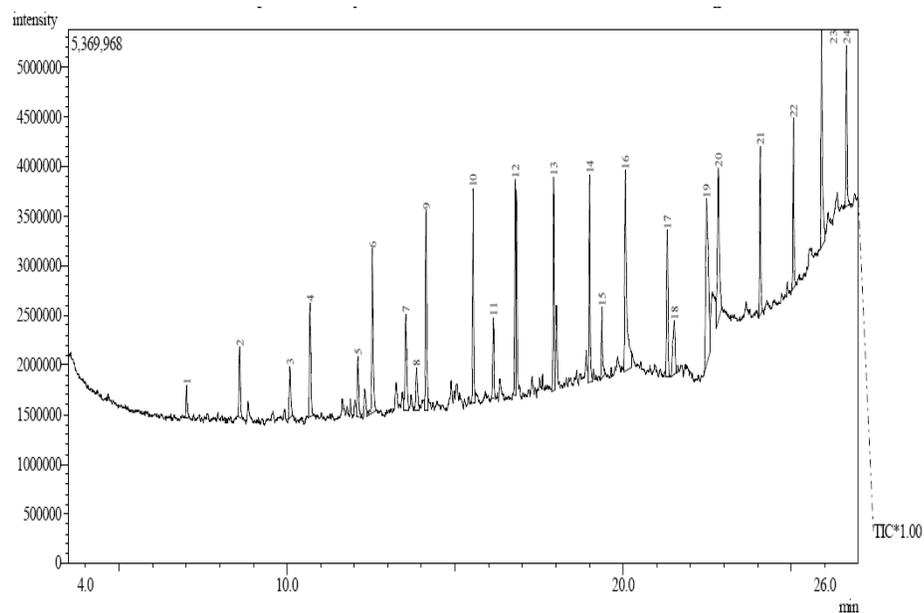


Figure 3: Chromatogram of Soil Particle Size B (0.3mm) Degraded by Indigenous Bacteria after 56 Days of Incubation

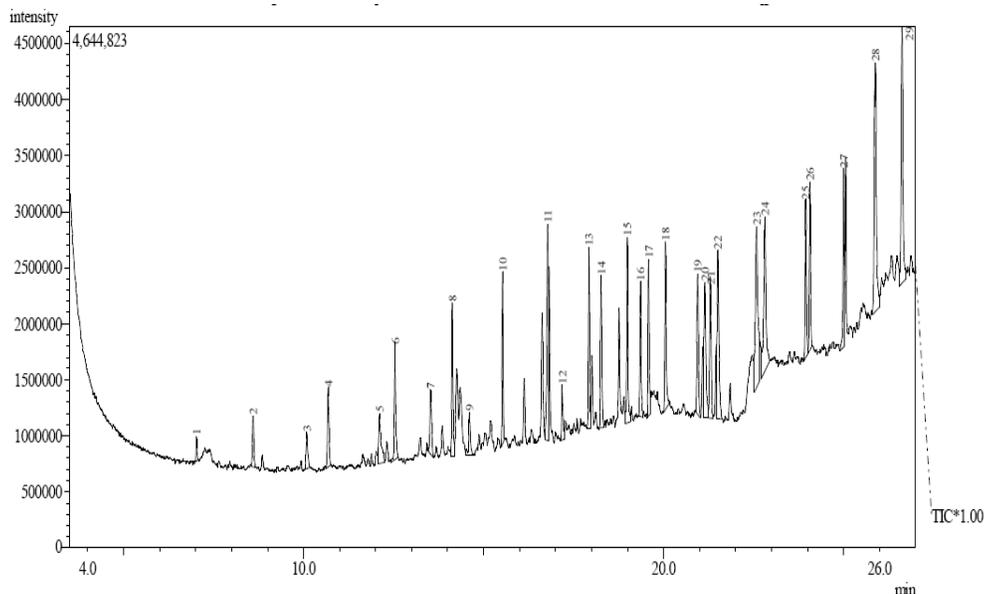


Figure 4: Chromatogram of Soil Particle Size C (0.15mm) Degraded by Indigenous Bacteria after 56 Days of Incubation

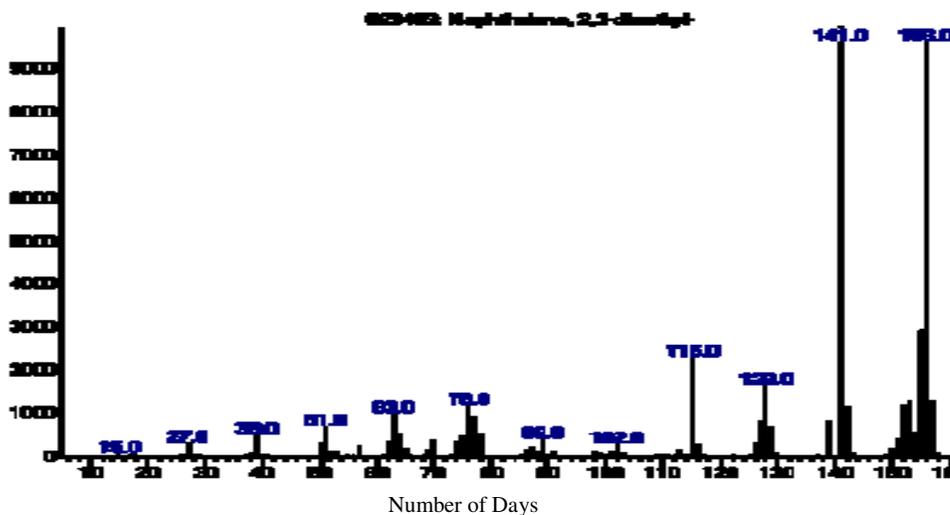


Figure 5: Chromatogram of Soil Particle size D (0.75mm) Degraded by Indigenous Bacteria after 56 Days of Incubation

The result of the GCMS analysis (Figures 1 - 5) further confirmed the biodegradation potentials of the isolates. GCMS analysis was conducted at 14, 28 and 56 days of the study. It was observed that the highest biodegradation occurs at 56 day. This revealed that biodegradation through natural attenuation takes longer time. This finding is consistent with previous work of Victor *et al.*, 2015, who made similar observation that

natural attenuation occurs in his control sample (1kg soil 100ml spent engine oil) at 70 days of the study due to 50% removal of the oil. The result further shows that some components of the oil, particularly the lighter hydrocarbons were completely lost while lower molecular weight were reduced in peak height, indicating that degradation of the component has occurred. Other components of the oil were intact.

This is in agreement with the results of Kawo and Faggo (2017) and Maryam and Ijah (2015) who also reported a decrease in peak height of chromatogram as indication of degradation. It was observed that highest degradation occurred on particle A, followed by B, C and D (Figures 2-5). Generally, the accepted pattern of susceptibility of hydrocarbon components to microbial degradation is n-alkane > branched alkanes > low-molecular weight aromatics > polycyclics. However, system-specific exceptions to this pattern have been found (Makut and Ishaya, 2010).

The findings further suggest that among the four soil particle sizes encountered in this study, the relative ease of oil degradation via natural attenuation is in the order

0.6mm>0.3mm>0.15mm>0.075mm. This study provides information on how different soil particle sizes interact with hydrocarbons that would aid in oil spill bioremediation.

CONCLUSION AND RECOMMENDATION

The study concluded that larger particle size (0.6mm) favors faster hydrocarbon biodegradation. The study also revealed the presence of *Pseudomonas putida*, *Staphylococcus aureus*, *Proteus vulgaris*, *Bacillus cereus*, and *E. coli*. The study also concluded that biodegradation can be influenced by soil particle sizes, concentrations and types of the organisms present in the systems.

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