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Enrichment and Identification of Askarel oil (PCB blend) degrading bacteria enriched from landfill sites in Edo State, Nigeria.

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ABSTRACT

Microbial degradation following aerobic biodegradation is one of the means used by microorganisms for the removal of persistent organic pollutants from the soil. Bacterial species namely: *Pseudomonas*, *Micrococcus* *Arthrobacter* and *Acinetobacter* capable of utilizing askarel oil (PCBs blend) were isolated from landfill soil samples from Uzogholo, Mgboaku and Auchi in Edo State of Nigeria. The isolation of these bacterial species followed an enrichment in minimal salt media where Askarel oil served as the sole carbon source. These bacterial isolates were characterized based on their cultural, morphological, biochemical characteristics and comparison with standard reference organisms. The potentials of these bacterial species to utilize askarel oil were assessed by measuring changes in the turbidity and pH of the enrichment medium containing varying concentrations of the Askarel oil. From the weekly results obtained, significant increases in the mean turbidity ranged between (0.065- 0.371) and decreases in mean pH ranged between (6.15- 3.55) for the 21 days incubation period. Some of the bacterial organisms exhibited remarkable utilization of askarel oil at different concentrations in 15 μ L and 20 μ L of the askarel minimal salt medium. From these findings, the bacterial species of remarkable potential can be isolated, re-engineered via biotechnology or bioaugmentation for effective remediation of landfill sites polluted with Polychlorinated biphenyls compounds and its derivatives in Nigeria.

Keywords: Aerobic biodegradation, bacterial species, biotechnological tools, Edo State, Enrichment, Minimal salt medium.

INTRODUCTION:

Environmental pollutants of anthropogenic origin are serious concern when introduced into environment at concentrations that impair the biological functioning of the ecosystem or pose risks to plants, animal and human welfare (Scullion, 2006). Polychlorinated biphenyls (PCBs) and its derivatives are among the organic pollutant that have long continued to be a source of concern due to its high persistence, carcinogenic, mutagenic and tetraogenic potential. Polychlorinated biphenyls are organic compounds with several chlorine atoms attached to the biphenyl. Commercial products of Polychlorinated biphenyls can be prepared industrially following catalytic chlorination of biphenyl with anhydrous chlorine in the presence of iron fillings or ferric chloride as a catalyst. Theoretically, about 209 possible polychlorinated biphenyls isomers, differing from each other by the level of chlorination and substitution position exists. In general, the most common isomers are those with an equal number of chlorine atoms on both rings or differ in only one chlorine atom between the rings (Langer, 2005). As a

result of potential accumulation and toxic effects within the environment, the manufacture of PCBs were banned in most countries. However due to their non-flammability, chemical stability, high boiling point and electrical insulating properties, PCBs still have applications in electrical equipment such as heat transfer, and hydraulic equipment; as plasticizers in paints, plastics and rubber products; in pigments, dyes and carbonless copy paper. In transformers, they serve as coolants and insulating fluids, in capacitors especially components of early fluorescent light fittings, locomotive's electrical transformers, stabilizing additives in flexible PVC coatings of electrical wiring and electronic components, pesticide extenders, cutting oils, reactive flame retardants, lubricating oils, hydraulic fluids, sealants for caulking in schools and commercial building (Rudel *et al.*, 2008). According to reports of Faroon *et al.*, (2003) more than 1.5 million tons of PCBs were manufactured worldwide between 1927 and the early 1980s of which a significant amount has been released to the environment. Due to their hydrophobic nature, PCBs tend to be adsorbed strongly to organic matter, thus leading to their

accumulation in soils and in the food chain (Sawhney *et al.*, 1987). In organic solvents, such as lipids PCBs tend to be soluble. This influences their transport and persistence in the environment. PCBs solubility in water is very low and decreases with increasing degree of chlorination. The solubility of the compound plays an essential role in its degradation. Compounds with high aqueous solubility can be easily degraded by microorganisms than those with low solubility (Imamoglu *et al.*, 2002). Over the years, PCBs residues may have entered the environment (atmospheric, aquatic and terrestrial) through inadequate elimination by incineration of PCB items. In the atmosphere, owing to the dynamic fluxes of wind speed, there exists a relatively low percentage of PCBs. Furthermore, this process is one of the most common means of PCB transportation and global dispersion in the environment. (Klecka *et al.*, 2000; Mandalakis *et al.*, 2002). Because PCBs can undergo long range of atmospheric transport ie moving from source regions to more remote locations; PCBs can be detected in the biota from different regions of the world, where they have been neither used nor produced. (Wania, 2003; Braune *et al.*, 2005). According to Offenberg *et al.*, (2005) the emission of PCBs into the coastal atmosphere from industrial areas may lead to increased depositional fluxes. In the reports of Gevao *et al.*, (2006) equipment dumps, landfills, sewage sludge and PCB-coated silos represent large PCB reservoirs which frequently find their way to contaminate the terrestrial area and groundwater. PCBs have been found to cause a variety of adverse health effects in animals which range from immune, reproductive, nervous and endocrine disorders. In human there exist supportive evidence for potential carcinogenic and non-carcinogenic effects of PCBs (USEPA, 1987). Transformation of PCBs and there derivatives is a critical step determining their fate and persistence in the environment. The effectiveness of remediation of contaminated sites depends on degradation rates and the conditions present in the environment. These includes: the degree of biphenyl chlorination; position of chlorine atoms in the biphenyl nucleus; species of microorganisms, their activity and in-between interactions; the structure of a given PCB compound; the presence of substituents and their position in the molecule; the presence of toxic or inhibitory substances; solubility, temperature, pH, light, and concentration of the pollutant (Imamoglu *et al.*, 2002). Biodegradation using microorganisms depends on the array of enzymes produced them; and possible use of the pollutant as source of carbon

and energy. Following the isolation of two bacterial strain (*Achromobacter* species) by Ahmed and Focht in 1973, the overall perspective of PCBs as immutable chemicals were changed. From then onwards, quite a number of bacterial strains capable of utilization of PCBs have been noted. (Furukawa *et al.*, 1979; Bopp 1986; Bedard *et al.*, 1987, Kim and Picardal 2000 and Adebusoeye *et al.*, 2008). However these notable achievements, in most developing countries such as Nigeria most PCB items and PCB waste oils are discharged into the environment without proper monitoring and determination of the fate of such compounds (Adebusoeye *et al.*, 2008). Furthermore from research studies, the isolation and studies on degradation of PCB item and its derivatives (transformer oil) in Nigeria is still at its infancy. However this, there are few notable scientists in Nigeria that have provided groundbreaking research for continuous search for more bacterial species capable of utilizing PCB and its derivatives as carbon sources. To contribute in the quest for more bacterial species capable of growth and utilization of PCB items /(transformer oil-askarel oil),author seek to isolate, characterize and determine the potentials of bacterial species from landfill sites in Edo State ,Nigeria capable of utilization of askarel oil (a PCB blend) as carbon source.

MATERIALS AND METHODS

Chemical and Reagents: All chemicals and reagents were of analytical grade. The (PCB blend) Askarel was generously provided by Power Holding Company of Nigeria. All other chemicals and reagents were obtained from Sigma- Aldrich Chemicals Co Ltd England.

Sample Collection: The soil samples were collected from landfill sites containing organic wastes, sewage sludge, old electrical equipment and waste oil in Uzoghola, Mgboaku and Auchu in Edo State, Nigeria. Geographical coordinates are 7° 4' 0" North, 6° 16' 0" East. The site had been contaminated with wastes from organic and inorganic pollutants for several years. The soil samples were collected randomly at sites with indications of low to high level wastes. The soil was collected from a depth of about 2-7cm deep using a sterile hand trowel. The soil samples were placed in separate sterile jars and transported back at ambient temperatures for further studies. The trowel surface was sterilized with 70% ethanol prior to collection of each sample.

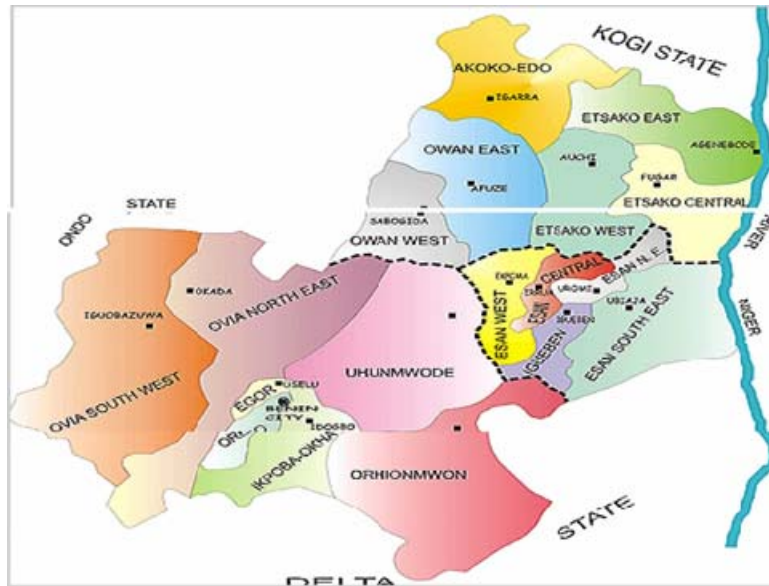


Fig : Map showing the Edo state

Source : Google.com

Isolation and Enrichment of Aerobic Indigenous Bacterial Strains: The enrichment and degradation potential of askarel oil were conducted in Minimal salt medium containing in CaCO_3 (2mg/L), $\text{MgSO}_4 \cdot 7\text{H}_2\text{O}$ (0.1g/L), $\text{ZnSO}_4 \cdot 7\text{H}_2\text{O}$ (1.44mg/L), $\text{CuSO}_4 \cdot 5\text{H}_2\text{O}$ (0.25mg), H_3PO_4 (0.06mg/L), HCL (51.3 μL) and Askarel oil (PCB blend) about 20ppm as carbon source. The pH was adjusted to 7.0. Cultures were incubated in 160mL serum bottles containing a liquid volume of 30mL with mouth plugged with sterile cotton wool and incubated at room temperature (25°C) for a period of three weeks. Incubation at 37°C is lethal to many soil bacteria. For the bacterial isolation from enrichment culture, transfers to fresh Askarel minimal salt medium using about 10% of inoculums from the previous enrichment was done weekly and incubated at 25°C. This procedure was repeated for four successive transfers. Pure cultures were isolated from the enrichment media by plating out on nutrient agar sprayed with Askarel. Discrete single colonies were selected and inoculated on Minimal agar medium sprayed with Askarel. The process was repeated severally to obtain pure cultures capable of growth on Askarel oil (Liu *et al.*, 2002, Nwinyi *et al.*, 2008; Nwinyi,2010).

Identification and characterization of the Isolated Bacterial Strains: The pure bacterial strains were identified on the basis of their morphological and biochemical tests. The pure cultures of the bacterial isolates were subjected to various morphological and biochemical characterization tests such as color, shape, elevation, consistency, margin, Catalase test, MRVP (methyl red-voges proskauer test), fermentation of sugars, kovacs citrate, indole, hydrolysis of starch, and

sensitivity tests. In order to determine the identity of bacteria isolates, results were compared with standard references of Bergey's Manual of Determinative Bacteriology 2nd edition (Buchanan and Gibbon, 1974; Olutiola *et al.*, 1991).

Determination of Growth Profile in Different Concentration of Askarel Blend: The isolates were inoculated into different concentrations in 5 μL , 10 μL , 15 μL and 20 μL askarel oil minimal salt medium. This was done to determine the tolerance level, degradation/transformation of Askarel oil through the utilization of Askarel oil as carbon source. The cultures were then incubated at room temperature 25°C for a period of three weeks. The Growth was monitored by measuring weekly the turbidity at 540nm using standardized Hanna H198703 Turbidimeter and pH by Hanna microprocessor pH meter.

RESULTS

Characterization of the isolates: A total of 5 different microbial colonies were selected upon screening individual isolates for growth on Askarel oil minimal media. The isolates were characterized using the morphological, cultural and biochemical characteristics with results shown in table 1.

Determination of growth in different concentration of Askarel minimal salts: The growth (utilization of askarel) by these isolates in different concentration of Askarel minimal salt medium are summarized in the Table 2. All the strains showed visible signs of growth on different compounds. Interestingly growth with production of blue coloration was seen on 15 μL ,20 μL concentrations of Askarel minimal salt containing the *Pseudomonas* spp.

Table: 1. Shows the cultural, morphological and biochemical characteristics of the bacterial isolates

| | A1 | A2 | B1 | B2 | C1 |
|--------------------------------|-----------------------|-----------------------|-----------------------|-------------------|-----------------------|
| | AI | A2 | B1 | B2 | C1 |
| Gram's reaction | - | + | + | + | - |
| Shape | Rod | Cocci | Cocci | Rod | Coccobaccilli |
| Motility | Motile | Non motile | Non motile | Non motile | Non motile |
| Spore formation | - | - | - | - | - |
| Acid fast | - | - | - | - | - |
| Form on Nutrient agar slant | Echinulate | Echinulate | Echinulate | Echinulate | Filiform |
| Growth on Nutrient agar slant. | Large | Moderate | Moderate | Large | Moderate |
| Optical features | Translucent | Transparent | Transparent | Translucent | Transparent |
| Growth on Nutrient Broth | Turbid with pellicles | Turbid with pellicles | Turbid with pellicles | Uniform turbidity | Turbid with pellicles |

| | | | | | |
|----------------------|------------------------|------------------------|------------------------|-------------------------|--------------------------|
| Catalase | + | + | + | + | + |
| Urease | - | + | + | + | + |
| Starch hydrolysis | - | - | - | + | + |
| Growth in 5% NaCL | + | + | + | + | + |
| Growth in pH 6.0 | + | + | + | + | + |
| Growth at 35°C | + | + | + | + | + |
| Lactose fermentation | - | - | - | - | - |
| Sucrose fermentation | - | - | - | - | - |
| Maltose fermentation | - | - | - | - | - |
| Glucose fermentation | - | - | - | - | - |
| Indole | - | - | - | + | - |
| Methyl Red | - | - | - | - | - |
| Citrate | + | - | - | + | + |
| Probable Organism | <i>Pseudomonas spp</i> | <i>Micrococcus spp</i> | <i>Micrococcus spp</i> | <i>Arthrobacter spp</i> | <i>Acinetobacter spp</i> |

Key: + = Growth, - = No growth A=Isolates from Uzogholo, B=Isolates from Mgboaku, C=Isolates from Auch

Table 2.0 Shows the different pH and Optical density readings at (540nm) recorded for the 21 days incubation period at different concentrations (5µL, 10 µL 15 µL, 20 µL)

| MICROORGANISM | pH at 5µL after 1week | pH at 5µL after 2 weeks | pH at 5µL after 3weeks | Mean pH at 5µL |
|--------------------------|-----------------------|-------------------------|------------------------|----------------|
| <i>Pseudomonas spp</i> | 5.57 | 5.92 | 5.85 | 5.78 |
| <i>Micrococcus spp</i> | 5.60 | 5.91 | 5.97 | 5.83 |
| <i>Micrococcus spp</i> | 5.59 | 5.96 | 5.98 | 5.84 |
| <i>Arthrobacter spp</i> | 5.89 | 6.22 | 6.23 | 6.11 |
| <i>Acinetobacter spp</i> | 5.90 | 6.30 | 6.27 | 6.15 |

| MICROORGANISM | pH at 10 µL after 1week | pH at 10 µL after 2weeks | pH at 10 µL after 3weeks | Mean pH at 10µL |
|--------------------------|-------------------------|--------------------------|--------------------------|-----------------|
| <i>Pseudomonas spp</i> | 5.49 | 5.99 | 6.05 | 5.81 |
| <i>Micrococcus spp</i> | 5.57 | 5.91 | 5.95 | 5.80 |
| <i>Micrococcus spp</i> | 5.68 | 6.07 | 6.13 | 5.96 |
| <i>Arthrobacter spp</i> | 5.89 | 6.22 | 6.23 | 6.11 |
| <i>Acinetobacter spp</i> | 5.90 | 6.30 | 6.27 | 6.15 |

| MICROORGANISM | pH at 15 µL after 1week | pH at 15 µL after 2weeks | pH at 15 µL after 3weeks | Mean pH at 15 µL |
|--------------------------|-------------------------|--------------------------|--------------------------|------------------|
| <i>Pseudomonas spp</i> | 3.89 | 3.46 | 3.30 | 3.55 |
| <i>Micrococcus spp</i> | 3.66 | 4.18 | 4.65 | 4.16 |
| <i>Micrococcus spp</i> | 4.54 | 5.20 | 5.50 | 5.08 |
| <i>Arthrobacter spp</i> | 4.63 | 5.16 | 5.40 | 5.06 |
| <i>Acinetobacter spp</i> | 4.92 | 5.53 | 5.78 | 5.41 |

| MICROORGANISM | pH at 20 µL after 1week | pH at 20 µL after 2weeks | pH at 20 µL after 3 weeks | Mean pH at 20µL |
|--------------------------|-------------------------|--------------------------|---------------------------|-----------------|
| <i>Pseudomonas spp</i> | 3.59 | 3.62 | 3.83 | 3.68 |
| <i>Micrococcus spp</i> | 5.59 | 5.78 | 5.90 | 5.76 |
| <i>Micrococcus spp</i> | 5.55 | 5.42 | 5.62 | 5.53 |
| <i>Arthrobacter spp</i> | 4.18 | 4.30 | 4.62 | 4.37 |
| <i>Acinetobacter spp</i> | 4.84 | 5.47 | 5.69 | 5.33 |

| MICROORGANISM | Optical density at 5µL after 1week | Optical density at 5µL after 2weeks | Optical density at 5µL after 3weeks | Mean Optical density at 5µL |
|--------------------------|------------------------------------|-------------------------------------|-------------------------------------|-----------------------------|
| <i>Pseudomonas spp</i> | 0.053 | 0.069 | 0.110 | 0.077 |
| <i>Micrococcus spp</i> | 0.051 | 0.068 | 0.087 | 0.069 |
| <i>Micrococcus spp</i> | 0.038 | 0.068 | 0.095 | 0.067 |
| <i>Arthrobacter spp</i> | 0.046 | 0.069 | 0.121 | 0.079 |
| <i>Acinetobacter spp</i> | 0.063 | 0.091 | 0.114 | 0.089 |

| MICROORGANISM | Optical density at 10 µL after 1week | Optical density at 10 µL after 2weeks | Optical density at 10 µL after 3weeks | Mean Optical density at 10µL |
|--------------------------|--------------------------------------|---------------------------------------|---------------------------------------|------------------------------|
| <i>Pseudomonas spp</i> | 0.073 | 0.100 | 0.144 | 0.106 |
| <i>Micrococcus spp</i> | 0.043 | 0.085 | 0.124 | 0.084 |
| <i>Micrococcus spp</i> | 0.066 | 0.083 | 0.091 | 0.080 |
| <i>Arthrobacter spp</i> | 0.063 | 0.061 | 0.072 | 0.065 |
| <i>Acinetobacter spp</i> | 0.060 | 0.060 | 0.077 | 0.066 |

| MICROORGANISM | Optical density at 15µL after 1week | Optical density at 15µL after 2weeks | Optical density at 15µL after 3weeks | Mean Optical density at 15µL |
|--------------------------|-------------------------------------|--------------------------------------|--------------------------------------|------------------------------|
| <i>Pseudomonas spp</i> | 0.139 | 0.327 | 0.350 | 0.272 |
| <i>Micrococcus spp</i> | 0.329 | 0.316 | 0.359 | 0.334 |
| <i>Micrococcus spp</i> | 0.244 | 0.216 | 0.232 | 0.231 |
| <i>Arthrobacter spp</i> | 0.196 | 0.203 | 0.220 | 0.206 |
| <i>Acinetobacter spp</i> | 0.175 | 0.166 | 0.182 | 0.174 |

| MICROORGANISM | Optical density at 20 µL after 1week | Optical density at 20 µL after 2weeks | Optical density at 20 µL after 3weeks | Mean Optical density at 20µL |
|--------------------------|--------------------------------------|---------------------------------------|---------------------------------------|------------------------------|
| <i>Pseudomonas spp</i> | 0.305 | 0.396 | 0.412 | 0.371 |
| <i>Micrococcus spp</i> | 0.260 | 0.108 | 0.270 | 0.213 |
| <i>Micrococcus spp</i> | 0.086 | 0.199 | 0.268 | 0.184 |
| <i>Arthrobacter spp</i> | 0.228 | 0.262 | 0.303 | 0.264 |
| <i>Acinetobacter spp</i> | 0.162 | 0.162 | 0.169 | 0.164 |

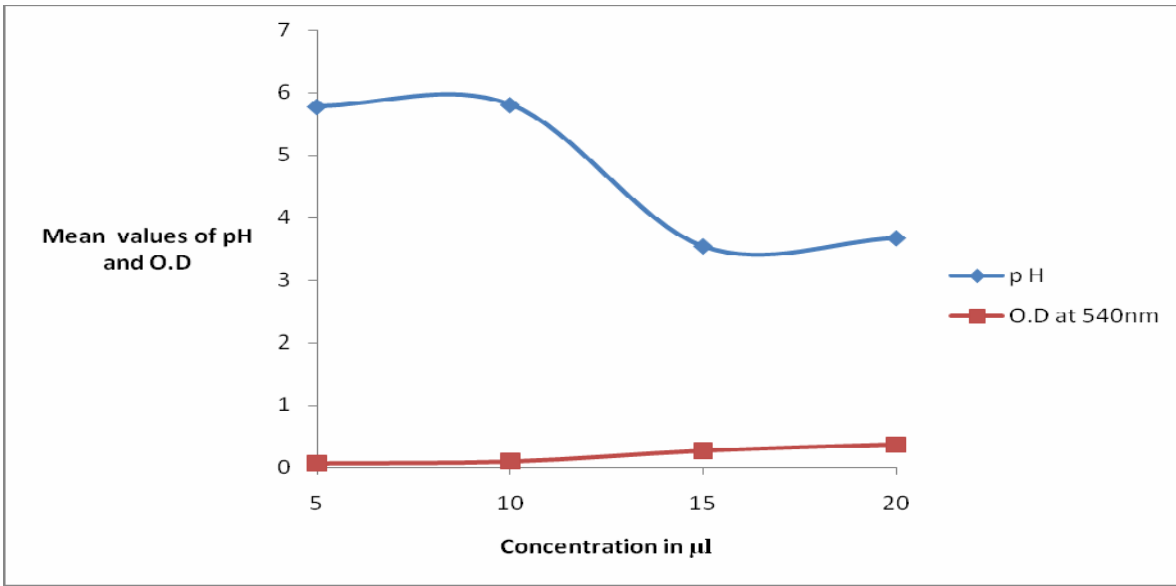


Fig. 1 shows graph for mean values of pH and optical density against different concentration levels of Askarel oil in minimal medium for 21 days by *Pseudomonas* spp isolated from Uzogholo.

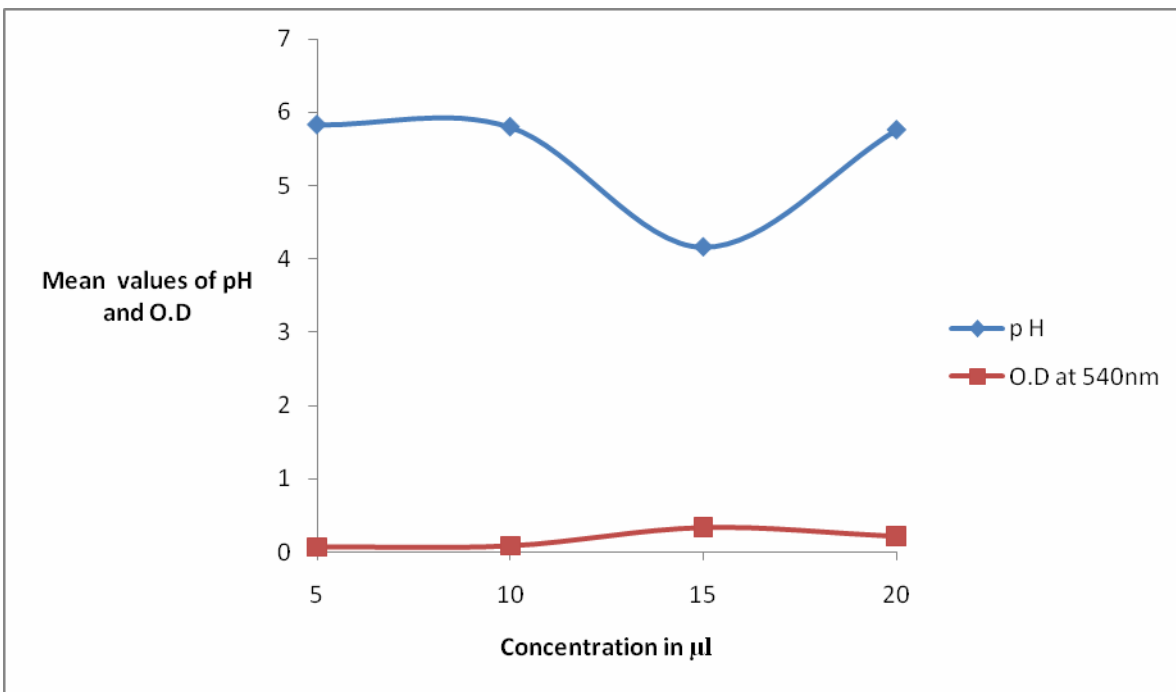


Fig. 2 shows graph for mean values of pH and optical density against different concentration levels(5 μL ,10 μL ,15 μL ,20 μL) of PCB blend in minimal medium for 21 days by *Micrococcus* spp isolated from Uzogholo.

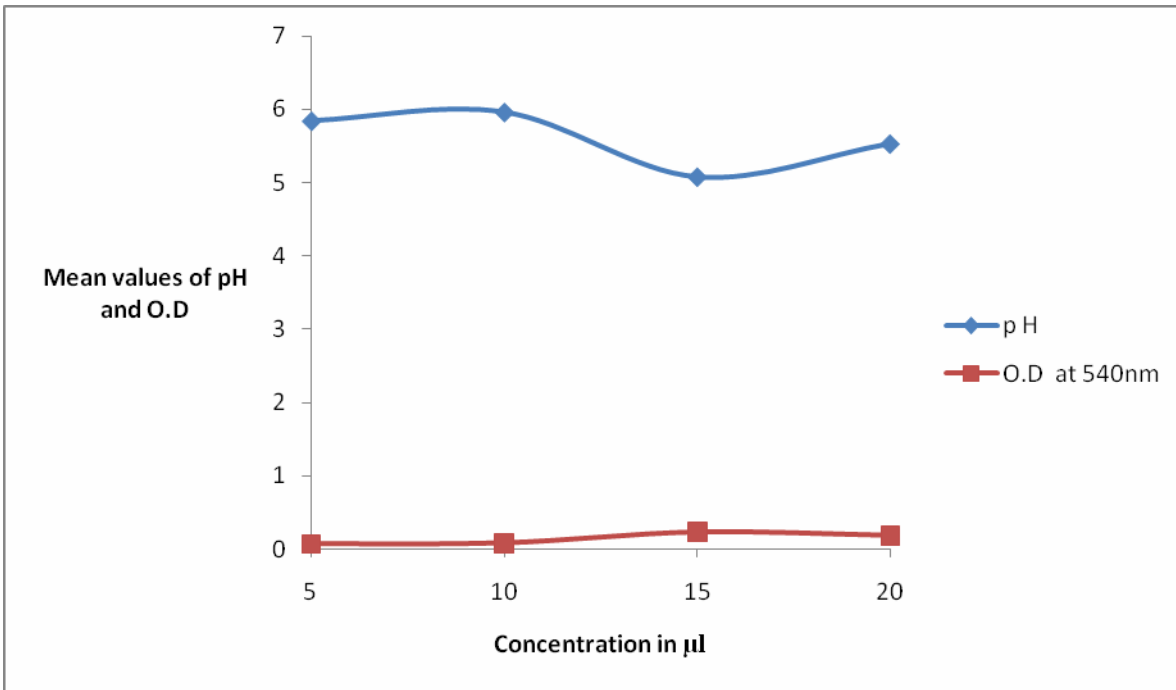


Fig. 3: shows graph for mean values of pH and optical density against different concentration levels of askarel (PCB blend) in minimal medium for 21 days by *Micrococcus* spp isolated from Mgboaku.

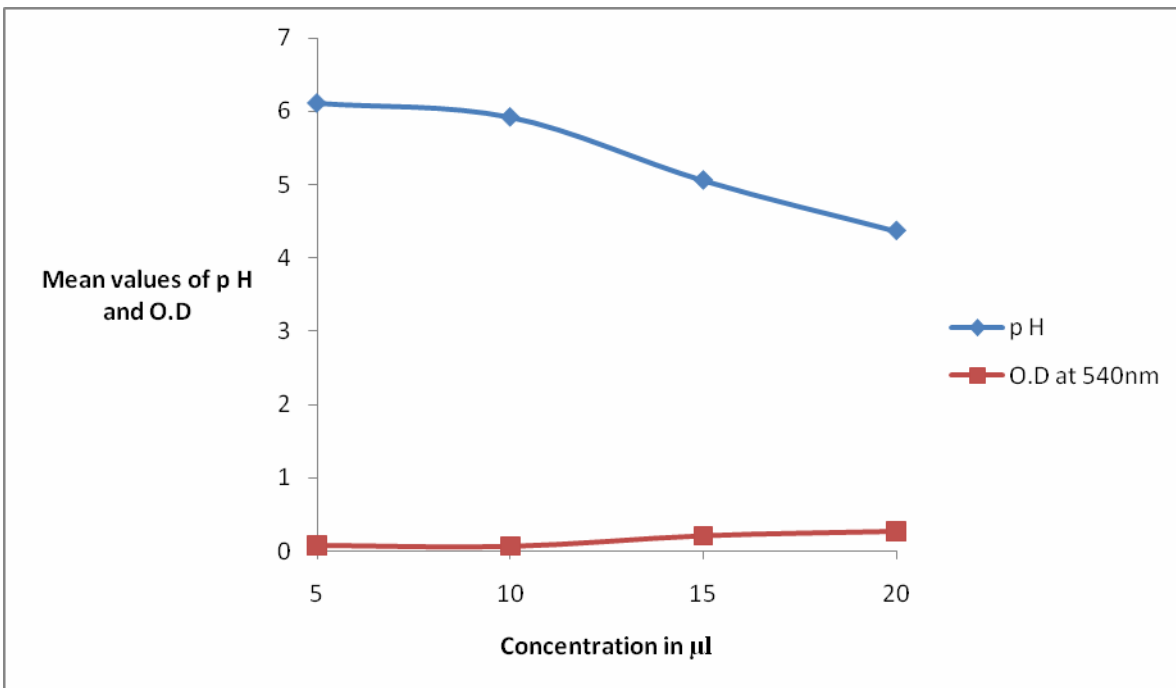


Fig. 4 : Shows graph for mean values of pH and optical density against different concentration levels of askarel (PCB blend) in minimal medium for 21 days by *Arthrobacter* spp.isolated from Mgboaku.

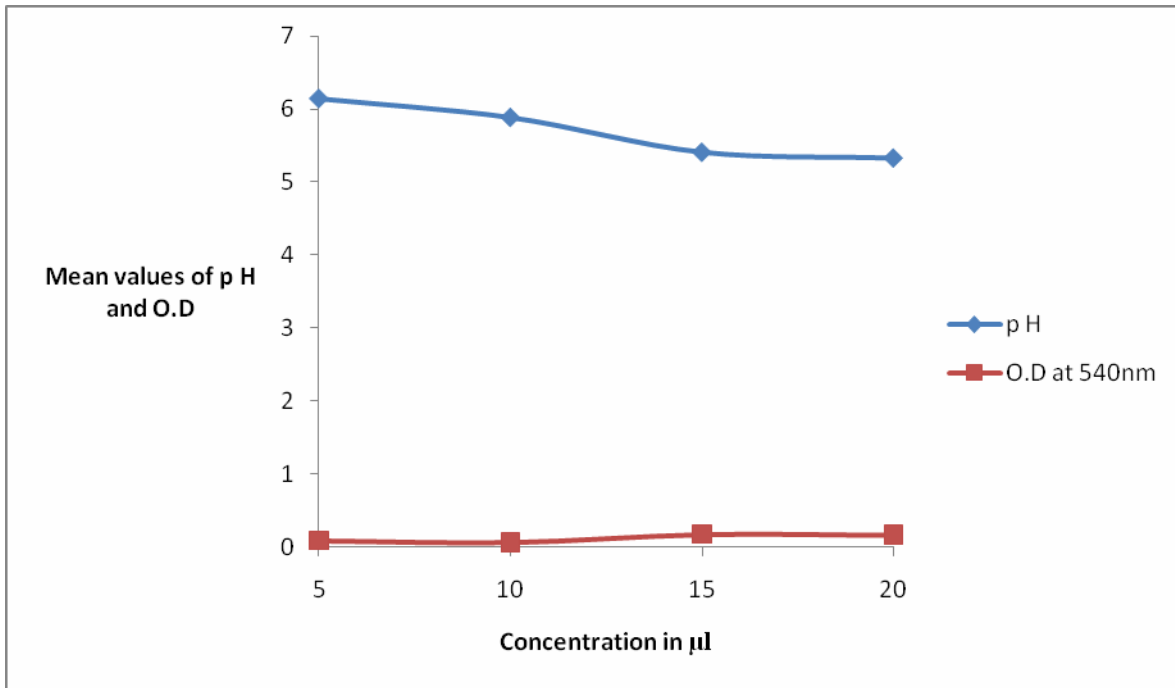


Fig. 5 Shows graph of mean values of p H and optical density(Turbidity) against different concentration levels of askarel (PCB blend) in minimal medium for 21 days by *Acinetobacter* spp isolated from Auchi.

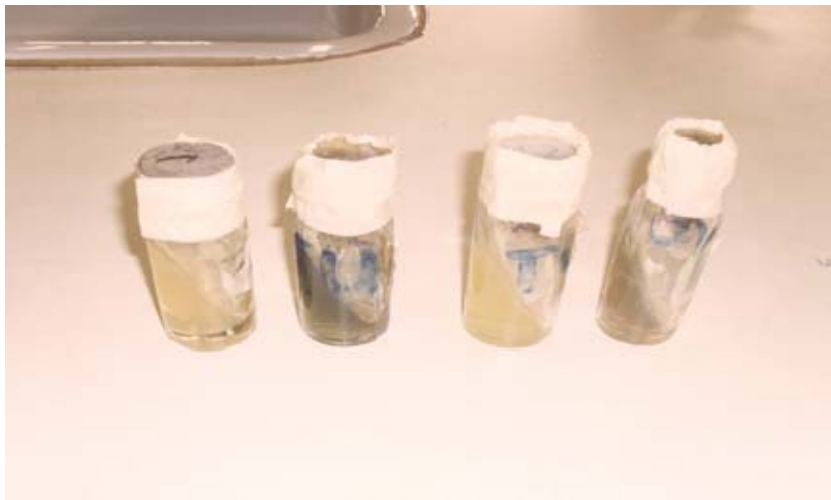


Plate1: shows bacteria isolated from soil samples in Nutrient Agar slant; *Micrococcus* spp, *Pseudomonas* spp, *Arthrobacter* spp, *Acinetobacter* spp respectively.



Plate 2: Shows *Pseudomonas* spp which produces blue coloration at a PCB blend (askarel) concentration of 15µL and 20µL.

DISCUSSION

To make the best of isolation of bacterial species with special abilities from sites of pollution, enrichment method is still an all-important process particularly where the target pollutant serves as the carbon and energy sources. In addition, previous exposures of these bacterial species to the target pollutants often trigger acclimatization and assemblage of important mechanisms that enhance the potential of these microorganisms in degrading the pollutants. In this research investigation, *Pseudomonas* spp, *Arthrobacter* spp, *Micrococcus* spp, and *Acinetobacter* spp were isolated from soils of different landfill sites in Edo State, Nigeria. *Pseudomonas* spp and *Micrococcus* spp were isolated from soil samples from Uzoghlo; *Micrococcus* spp and *Arthrobacter* spp were isolated from soil samples from Mgboaku while *Acinetobacter* spp was isolated from soil samples from Auchi. Polychlorinated biphenyls (PCBs) are biodegraded in two general ways; aerobic metabolism via co-metabolism and anaerobically by reductive dehalogenation (Sierra *et al.*, 2003). In this research biodegradation was carried out aerobically where askarel oil served as principal growth substrate. These isolated organisms were then grown in different concentrations at (5µL, 10µL,

15µL, 20µL) of polychlorinated biphenyls blend (askarel oil) in the minimal salt media. From the observed pH and optical density (O.D) of the growth profile of each bacterial species, obtained results agree with previous reports of organic pollutant utilization. From recent findings, growth of microorganisms in organic pollutants is often indicated by an increase in turbidity and decrease in pH (Nwinyi *et al.*, 2008). In this research growth occurred at different rates among the organisms isolated. *Pseudomonas* spp utilized the askarel oil significantly at concentrations in 15µL and 20µL of askarel minimal media, *Micrococcus* species isolated from Uzoghlo and Mgboaku utilized PCBs blend (Askarel oil) minimal media remarkably at a concentration of 15µL. These findings, showed consistency in behavior between the two *Micrococcus* spp isolated, possibly they could be of the same phylogeny. *Acinetobacter* spp isolated from land fill sites in Auchi showed minimal utilization of PCBs blend (askarel oil). This inability of *Acinetobacter* spp to thrive well in the PCBs blend may be as a result of entrapment of Askarel oil within some part of the biological membrane (Sikkema *et al.*, 1995). Rosenberg *et al.*, (1988) reported that emulsifying and dispersing activities of *Acinetobacter* species appear to be due to different materials, and

is restricted to a relatively small number of the species. Furthermore, PCB congener, 2, 2', 5, 5'-tetrachlorobiphenyl, has been shown to affect the membrane of *Ralstonia eutropha* H850 due to its accumulation in the cytoplasmic membrane (Kim *et al.*, 2001). Although neither the mass flow rate of the cytoplasmic membrane nor that between the cytoplasmic membrane and cytosol is known, it seems likely that the entrapment of PCBs within the membrane reduces both the accessibility of PCBs to the degradation enzymes and the efficiency of PCB degradation. In addition, the accumulation of hydrophobic compounds in the cytoplasmic membrane fluidizes and disrupts its function (Sikkema *et al.*, 1995), thus the accumulation of PCBs in the cytoplasmic membrane may also affect cell viability. Most likely the reduced metabolism rate may be due to high CO₂ concentrations since there is no rapid transfer of oxygen were in most research, additional aeration is provided to assist the organisms to aerobically breakdown the pollutants. From research studies, high CO₂ can inhibit cell growth were the headspace may not have been effectively purged, thus the high CO₂ may have caused the decline in growth rate.

Utilization of Askarel oil (PCBs blend) by *Arthrobacter* spp occurred significantly at 15µl and 20µl. *Pseudomonas* spp showed highest degradation of the askarel oil with pH of 3.55. This is probably because it is not a Gram positive organism. According to (Prescott *et al.*, 2002), Gram positive bacteria are usually more sensitive than Gram negative bacteria towards lipophilic toxic substrates, possible because they lack protection by the outer membrane. Furthermore, *Pseudomonas* spp had been noted as one of the major bacterial species that help degrade most organic pollutant. This is because of their ability to possess varieties of catabolic genes that are resident on the plasmids.

From studies, degradation of PCBs, is initiated by an enzyme called biphenyl 2,3-dioxygenase. Biphenyl 2,3-dioxygenases belong to a large family of Rieske non-heme iron oxygenases (Gibson and Parales, 2000). They comprise a terminal oxygenase composed of a large α- and a small β-subunit, a ferredoxin and a ferredoxin reductase. The ferredoxin and ferredoxin reductase act as an electron transport system to transfer electrons from NADH to the terminal oxygenase. The cis-2, 3-dihydro-2,3-dihydroxybiphenyl formed by this reaction is dehydrogenated by a dehydrogenase to give 2,3-dihydroxybiphenyl, which is subject to extradiol

cleavage. The 2-hydroxy-6-oxo-6-phenylhexa-2,4-dienoate thereby formed undergoes hydrolysis, yielding benzoate and 2-hydroxypenta-2,4-dienoate as reaction products. This sequence of reactions forms the biphenyl upper pathway as described by Ahmed and Focht, in 1973. They were the first to describe that biphenyl-degrading organisms of bacterial species have the capacity to transform several PCB congeners. Since then, a major focus of research on biphenyl-degrading organisms is their capability to transform PCBs, which differs significantly between different isolates (Gibson and Parales, 2000). Following incubation for 7 days in askarel oil (PCB blend) minimal salt containing *Pseudomonas* spp, there was a remarkable change in the medium from colorless to light blue at a concentration of 15µL and 20µL. This change in coloration may be instigated by the catabolic genes. Thus the enzymes produced by the genes may be responsible for the blue colored medium. Also the *Pseudomonas* spp showed highest turbidity (0.727-0.371) and lowest pH values (3.55-3.68). Biodegradation rates are highly variable because they depend on several factors, including the amount and location of chlorination, PCB concentration, type of microbial population, available nutrients, and temperature (Thomas *et al.*, 1992; Robinson and Lenn, 1994). *Pseudomonas* spp, *Arthrobacter* spp, and *Micrococcus* spp utilized the askarel oil (a PCB blend) appreciably at different concentrations in 15µL and 20µL but minimally at 5µL and 10µL. This implies that at higher concentration these organisms degradative enzymes were induced significantly when compared to their behavior at low concentrations of askarel in the minimal medium. Also from bacterial growth kinetics, organisms tend to adapt to the new substrates before actual significant growth occurs and this may provide an explanation to the delay at concentrations in 5µL and 10µL. Thus the organisms were slow in acclimatizing to the Askarel oil (PCBs blend) as evidenced with their performance at an increased concentration of 15µL and 20µL.

In conclusion, *Pseudomonas* spp, *Arthrobacter* spp, *Micrococcus* spp and *Acinetobacter* spp are prevalent microorganisms in soils of tropical ecosystem; therefore further research should be carried out on their degradative competence and also possible expansion of their substrate range using biotechnological tools for effective remediation of landfills polluted with waste polychlorinated biphenyls and their derivatives.

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