

Microbiological and Polycyclic Aromatic Hydrocarbon Profile of *Pomadasys commersonnii* Harvested from Crude Oil Impacted Creeks in Borikiri, Port Harcourt, Nigeria

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ABSTRACT

This study evaluated the microbiology and level of PAH contamination of *Pomadasys commersonnii* and surface water along Ikpukulu Creek in Borikiri. Standard microbiological procedures were used to assess and identify total heterotrophic bacteria (THB), total fungi and hydrocarbon utilizing bacteria (HUB) and fungi (HUF). PAHs were analyzed using Gas chromatographic–Flame Ionization Defector. Results revealed that, THB counts of *Pomadasys commersonnii* ranged from 2.4×10^6 to 1.2×10^7 cfu/g, HUB; ranged from 2.0×10^3 to 1.3×10^4 cfu/g, Fungi; ranged from 1.9×10^3 to 2.3×10^5 cfu/g and HUF ranged from 2.2×10^2 to 2.5×10^3 cfu/g. Genera of HUB and HUF identified from fish and water samples were; *Pseudomonas*, *Bacillus*, *Aeromonas*, *Proteus*, *Serratia*, *Citrobacter*, *Staphylococcus*, *Escherichia*, *Corynebacterium*, *Acinetobacter*, *Flavobacterium*, *Micrococcus*, *Enterobacter*, *Penicillium*, *Mucor*, *Rhizopus*, *Fusarium*, *Cladosporium*, *Metshnikowia*, *Candida*, *Cryptococcus*, *Leucosporidium*, *Rhodotorula*, *Aspergillus*, *Alternaria*, *Geotrichium*, *Saccharomyces*, and *Trichoderma*. PAHs level in fish samples ranged from 0.025mgkg^{-1} – 0.035mgkg^{-1} with a mean of 0.031mgkg^{-1} , while PAHs in water ranged from 0.0035 – 0.0045mgL^{-1} with a mean of 0.004mgL^{-1} . Highest PAH recorded was benzo(b)fluoranthene (60%). Results showed that the fish biomagnified pyrenes, benzo(a)anthracene and benzo(b)fluoranthene. Contamination factor (c^1f) was 0.12 ($c^1f < 1$), which indicates, Cancer Risk Index using indeno1,2,3,(cd)pyrene as bioindicator showed no cancer risk in consumption of *Pomadasys commersonnii* harvested from Ikpukulu Creek. There is significant difference at $p < 0.05$ in PAH level between surface water and *Pomadasys commersonnii* indicating the fish has the tendency of biomagnifying some PAHs. Therefore, the inhabitants should be enlightened of the dangers of these pollutants (PAHs) in food chains. Some of the hydrocarbon utilizing microbes in this study could be useful in bioremediation of hydrocarbon and PAH contaminated environments.

Keywords: Crude oil, PAH, Creek, *Pomadasys commersonnii*, biomagnify, carcinogen

Introduction

Contamination of water bodies with hydrocarbons from partial refined crude oil discharged into the ecosystem is a common occurrence in contemporary time in Rivers State (Ekweozor, 1996). Apart from

interfering with the chemistry of both water and sediments, the survival of both micro and macro fauna are threatened by the components of crude oil in the ecosystem (FME, 2006). Polycyclic aromatic hydrocarbons (PAHs) are a group of hydrocarbons, with fused aromatic (benzene) rings between 2 – 6. PAHs occur in the environment naturally or as a result

of man's activities. PAHs arising from fossil fuel are called pyrogenic while those from crude oil are called petrogenic. About thirty parent PAHs are found in crude oil of which sixteen are regarded as important pollutants due to their high carcinogenic, mutagenic and teratogenic activity (Douglas *et al.*, 2020). The 16 listed PAHs as priority pollutants are; naphthalene, acenaphthylene, acenaphthene, flourene, anthracene, phenanthrene, fluoranthene, pyrene, chrysene, benzo (a) anthracene, benzo (b) fluoanthene, benzo (k) flouranthene, benzo (a) pyrene, benzo (g,h,i) pyrene, Indeno (1, 2, 3 – cd) pyrene, and dibenz (a,h) anthracene (Douglas *et al.*, 2020; Ariyo and Obire, 2022). PAHs are the first substance to be associated with carcinogenesis (Eisler, 1987). Among them benzo (a) pyrene is the known human carcinogen and is as an indicator for PAH exposure. PAHs are fractions of the crude oil that enter into the food chain and are bioaccumulated and biomagnified through feeding (Edori and Edori, 2021). Polycyclic aromatic hydrocarbons (PAHs) are a group of hydrocarbons of carbon between 2 - 6 are fractions of the crude oil that enter into the food chain and are bioaccumulated and biomagnified through feeding.

In order to maintain osmotic pressure, fishes always drink water, in this manner, PAHs gain entrance into the fish and alter some vital enzymatic reactions thus leading to cell necrosis (Temitayo *et al.*, 2019). Due to the specific heat of water, water current, wave actions and convection, the marine ecosystem and estuaries are able to stabilize the physicochemistry of the water. By photo-oxidation some PAH component are rid of the ecosystem within a short time, some are biodegraded by hydrocarbonclastic microbes (Vinithin *et al.*, 2015). However, some gets adsorbed onto surfaces in the ecosystem and become persistent (Jih *et al.*, 2018). Research works have been ongoing in various ecosystems due to continuous discharge of petroleum and refined products from artisanal refineries along the coasts of Niger Delta. Some studies have established that there are bacterial strains that have adapted to these discharges and hence have the capacity to degrade the pollutants even at very high concentrations (Obire and Nwaubeta, 2002). A study carried out on Bonny River on hydrocarbon degraders-Drilling fluid utilizing bacteria genera isolated were: *Pseudomonas*, *Bacillus*, *Micrococcus* and *Enterobacter*, with *Pseudomonas* having the highest

frequency of 35.7%, followed by *Bacillus* with the frequency of 30.7%, *Micrococcus* had 15.4% and *Enterobacter* 15.4%.Fungi genera isolated include *Aspergillus*, *Penicillium*, *Rhizopus* and *Mucor* (Nrior *et al.*, 2017).

Some PAHs e.g. Benz(a) pyrene are well reported carcinogens by the IARC (International Agency for Research on Cancer). Standards are therefore set by EU NO 835/2011 establishing permissible levels of PAH markers in seafoods. Such include 0.006mgkg⁻¹ of Benzo(a) pyrene or 0.01mgkg⁻¹ of the total level of Benzo(a) pyrene, chrysene, Benzo(a) anthracene and Benzo(b) fluranthene (PAHs 4) or 0.005mgkg⁻¹ or indeno 1,2,3, (cd) pyrene. PAHs are lipophilic and easily bioaccumulate in the tissues of fish. The consumption of seafood forms part of the culture of the people of Rivers State. Fish contains the heart friendly omega-3-protein without cholesterol (EUEPA, 2009).

Pomadasys commersonnii (commonly known as spotted grunter) is found in coastal water and estuaries. It can tolerate fresh water. It feeds mostly on crustaceans, worms and bivalves. It is ray finned and a native of brackish and marine waters of West Indian Ocean (Ibim and Bongili, 2018). It has a maximum length of 80cm, found within a depth of 0 – 49m in the water body. It belongs to the class Actinoptcroii, order pereformes (Ibim and Bongili, 2018). It is also known as *Pomadasys perculare*, *Pristopoma operculare*, *Labrus commersonnii*, *Pristipoma opercularis* (Ibim and Bongili, 2018). It is one of the choice fish though expensive in Rivers State. Apart from tilapia, mullet and sardine, *Pomadasys commersonnii* is one of the abundant fishes harvested along the Ikpukulu Creek (Vincent-Akpu and Okosiemiema, 2019).

Materials and Methods

Study Location

The study area are creeks in Borikiri Wetlands connecting the Bie-Ama and Pereama, Borikiri Sandfill area of the Port Harcourt, Rivers State Nigeria. It lies approximately between Longitude 7^o 1.0" E to 7^o 2'27.33"E and Latitude 4^o 43'32.2" N to 4^o 45' 0"N (Vincent-Akpu and Okosiemiema, 2019).

These are marketing points of kerosene, gasoline, lubricating oils and diesel from artisanal refineries conveyed from “kpo fire” camps along the coastal areas of Rivers State.

This study area is an Estuary with brackish water salinity ranging from 10% - 30%. The climate is tropical wet with lengthy rainy season and very short dry season. The vegetation is mangrove with subsoil characterized by typical fibrous clayey mud that shows a large value of compressibility (Ideriah *et al.*, 2012). Several anthropogenic activities take place along the creek, such as disposal of refuse and excreta, frequent blasting of crude oil conveying boats as well as recreation.

Sampling

Fish (*Pomadasys commersonnii*) were harvested by net throwing in the intertidal zones and open sea. Twenty fishes were randomly collected from each of the four sampling sites (Ikpukulu jetty, MTN Mast, Island and Okilo Polo) every month from November 2019 – October 2020. Samples were washed thoroughly with sea water and placed in foils and labeled and taken to the Institute of Pollution Studies (IPS) Rivers State University Nkpolu-Oruworukwu, Port Harcourt in an ice-chest as described by Isaac *et al.* (2018)

Sampling for surface water was done from November, 2019 to October, 2020. A total of five water samples were collected each month from each of the four sample stations; making 20 water samples per month. Sample bottles were carefully opened under the water within a depth of about 10cm from the surface. Bottles were rinsed with surface water samples about 3 times before sample collection.

The five composites of each sample stations were mixed together and homogenized to obtain a bulk composite (APHA, 1995). Water samples for microbiological analysis were properly labeled and put in ice box. Water samples were filtered and preserved on site with hydrochloric acid for PAH analysis and transported in an ice chest to the laboratory after proper labeling (APHA, 1995).

Samples of collected *Pomadasys commersonnii* were allowed to thaw and cool to room temperature. Fish samples were homogenized using clean sterile ceramic mortar and pestle. One gram (1g) of each part of sample was added to 9ml of normal saline for the serial dilution.

Microbiological Analysis

Total Heterotrophic bacterial Counts (THBC)

Total heterotrophic bacteria from the samples were enumerated using the spread plate technique as described by Prescott *et al.* (2005). An aliquot (0.1 ml) of the dilutions 10^{-4} to 10^{-6} were aseptically transferred into properly dried nutrient agar plates in duplicate, spread evenly using a bent glass rod and incubated at 37°C for 24 hours. After incubation, the bacterial colonies that grew on the plates were counted and sub-cultured into fresh nutrient agar plates using the streak plate technique.

Discrete colonies on the plates were aseptically transferred into 10% (v/v) glycerol suspension, well label and stored as stock cultures for preservation and identification (APHA, 1995). Total Heterotrophic Bacterial Counts (THBC) were calculated from the mean value of colonies counted from the duplicate plates using the below formula: THBC (CFU/g) = Number of Colonies/ (Dilution (10^{-7}) x Volume plated (0.1 ml)).

Total Heterotrophic Fungal Counts

The total Heterotrophic fungi were enumerated using the spread plate method as described by Prescott *et al.* (2005). An aliquot (0.1 ml) of the dilutions 10^{-2} to 10^{-3} was aseptically transferred into properly dried Sabouraud Dextrose Agar plates containing antibiotic (tetracycline and penicillin) to inhibit bacterial growth in duplicate (Cheesebrough, 2006).

Plates were spread evenly using a bent glass rod and incubated at 35°C for 3 to 5 days. After taking counts, the fungal colonies were sub-cultured onto Sabouraud Dextrose Agar slant in bijoux bottles for preservation (APHA, 1995).

Isolation and Enumeration of Hydrocarbon Utilizing Bacteria (HUB)

Collected water samples from the ice box were cooled to room temperature and diluted in tenfold serial dilution with sterile physiological saline to give an initial 1:10 dilution. 0.1 ml of prepared dilutions were pipetted out and placed on mineral salts medium supplemental with 50 ugml⁻¹fungizol miconazole nitrates to prevent fungal contamination.

Isolation and enumeration were done using spread plate techniques (Okpokwasili and Amanchukwu, 1988; Chikere *et al.*, 2009) using vapour phase transfer technique on mineral salts agar for HUB. The plates were incubated at 30°C for 7 days for HUB, while for THB, incubation was at 28°C for 24hours. Enumeration of isolates was done and expressed as CFU/ml for surface water and CFU/g for fish gills and tissues.

Hydrocarbon Utilizing Fungal Counts

The population of hydrocarbon utilizing fungi was determined by inoculating 0.1 ml aliquot of the serially diluted samples onto mineral salt agar media using vapour phase transfer method according to Nrior *et al.* (2017). The mineral salt medium used was amended with 250mg of tetracycline to inhibit the growth of hydrocarbon utilizing bacteria.

The plates were inverted and incubated at 28°C for 5 days. Fungal counts were expressed and recorded as Colony Forming unit per gram (CFU/g) for fish gills and tissues and CFU/ml for surface water.

Identification of Bacterial and Fungal Isolates

Bacterial isolates were identified based on cultural, morphological and biochemical characteristics using Holt *et al.* (1994).

On the other hand, the fungal spores /isolates that utilized carbohydrate as their sole carbon energy source were viewed macroscopically and microscopically using Lactophenol Cotton Blue Stain and the slide culture technique. This was carried out using the steps as described in Cheesbrough (2006).

Polycyclic Aromatic Hydrocarbon (PAH) Analysis

PAH Analysis of Surface Water

Two hundred and fifty milliliters (250ml) of water samples were measured into a separating funnel rinsed with dichloromethane. To the 250ml water sample 25ml dichloromethane was added. The mixture was shaken vigorously to extract all organic materials. The organic extract was passed through a column containing cotton wool, silica gel and anhydrous sulphate for cleaning and dehydration. The organic extract obtained was injected into gas chromatographic column. A µl of the concentrated sample was injected by means of hypodermic syringe through a rubber septum into the column, the vapour fraction of the PAHs were automatically detected as it emerges from the column by the flame ionization Detector (FID). The results were expressed in mg l⁻¹.

PAHs Analysis of *Pomadasys commersonnii*

Samples of collected *Pomadasys commersonnii* were allowed to thaw and cool to room temperature. Fish samples were homogenized using clean sterile ceramic mortar and pestle. Two grams (2g) of sample were weighed into a clean extractor container. 20ml of extraction solvent (hexane) was added into sample and mixed thoroughly and allowed to settle. The mixture was carefully filtered into Buchner funnels. The extracts were concentrated to 2ml and transferred for clean-up. For clean-up, 1cm of moderately packaged glass wool was placed at the bottom of 10mm internal diameter x 250mm long chromatographic column. Slurry of 2g activated silica in 10ml dichloromethane was prepared and placed into the chromatographic column. To the top of the column, 0.5cm of sodium sulphate was added.

The column was rinsed with additional 10ml of dichloromethane. The column was pre-eluted with 20ml of hexane. This was allowed through the column at the rate of about 2 minutes until the liquid in the column was just above the sodium sulphate layer. Immediately, 1ml of the extracted sample was transferred into the column. The extraction bottle was rinsed with 1ml of hexane and the dissolved extract was added to the column as well. The stop cork of the column was opened and the eluent was collected in a

10ml graduated cylinder. Hexane was added to the column in 1-2ml increment just before the exposure of the sodium sulphate layer to air. Accurately measured volume of 8-10ml of the effluent was collected and labeled-polyaromatic. The concentrated polyaromatic fractions were transferred into labeled glass vials with Tefion and rubber crimp caps for gas chromatographic analysis.

One microliter (1µl) of the concentrated sample was injected into the column by means of hypodermic syringe through a rubber septum into the column. Separation occurs as the vapour constituents' partition between the gas and liquid phases. The sample was automatically detected as it emerged from the column by the Flame Indicator Detector (FID) (RPI / NNPC, 1985a; RPI / NNPC, 1985b).

Results

Results of the total heterotrophic bacterial and total fungal counts in the surface water and in *Pomadasy commersonnii* from November 2019 to October, 2020 are as shown in Figure 1 and Figure 2 respectively. The THB counts ranged from 2.4×10^6 to 1.2×10^7 cfu/g while total fungi ranged from 1.9×10^3 to 2.3×10^5 cfu/g.

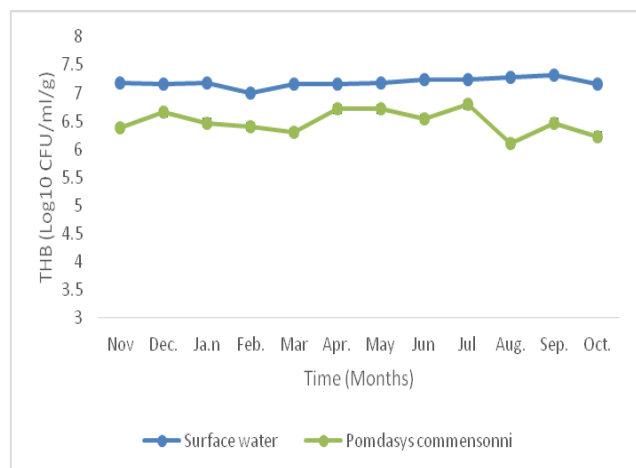


Fig. 1: Total Heterotrophic Bacterial Counts in Surface Water and *Pomadasy commersonnii* (Nov. 2019- Oct. 2020)

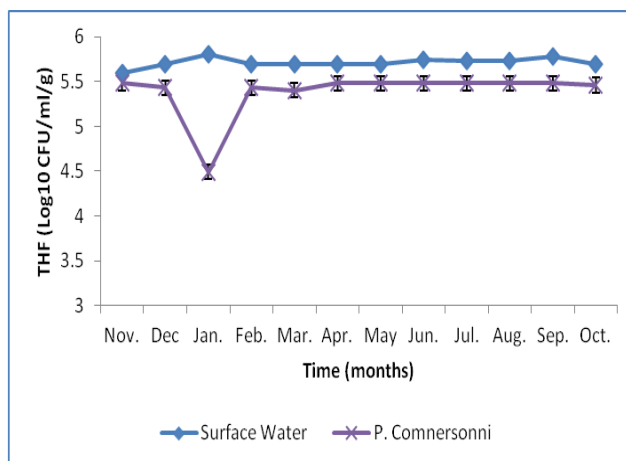


Fig. 2: Total Heterotrophic Fungi (THF) Count in Surface Water and *Pomadasy commersonnii* (Nov. 2019- Oct. 2020)

The results of the counts of hydrocarbon utilizing bacteria (HUB) and hydrocarbon utilizing fungi (HUF) are as shown in Figures 3 and 4 respectively. The counts of hydrocarbon utilizing bacteria ranged from 2.0×10^3 to 1.3×10^4 cfu/g, while counts of Hydrocarbon utilizing fungi ranged from 2.2×10^2 to 2.5×10^3 cfu/g.

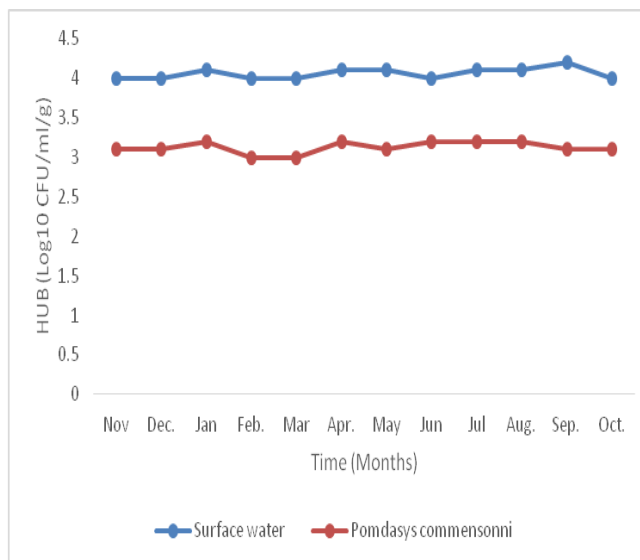
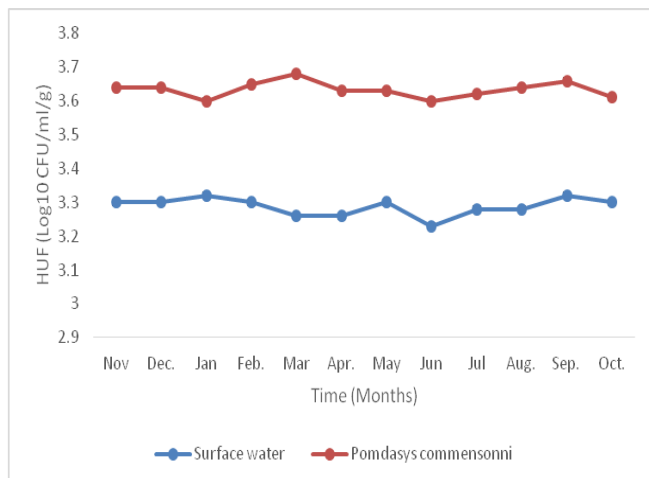


Fig. 3: Hydrocarbon Utilizing Bacteria (HUB) Count in Surface Water and *Pomadasy commersonnii* (Nov. 2019- Oct. 2020)



The occurrences of identified Hydrocarbon utilizing Bacteria (HUB) and the Hydrocarbon Utilizing Fungi (HUF), and their percentage (%) occurrences in the surface water samples and in the fish *Pomadasys commersonni* samples during this study are as shown in Table 1 below.

Fig. 4 Hydrocarbon Utilizing Fungi (HUF) Count in Surface Water and *Pomadasys commersonni* (Nov. 2019- Oct. 2020)

Table 1: Occurrence (%) of Hydrocarbon Utilizing Bacterial (HUB) and Hydrocarbon Utilizing Fungal (HUF) isolates from Surface water and Fish – *Pomadasys commersonni*

Hydrocarbon Utilizing Bacterial (HUB)			Hydrocarbon Utilizing Fungal (HUF)		
HUB isolate	Surface Water Occurrence (%)	<i>Pomadasys commersonni</i> Occurrence (%)	HUF isolate	Surface Water Occurrence (%)	<i>Pomadasys commersonni</i> Occurrence (%)
<i>Proteus</i> sp.	3(4.69)	1(3.03)	<i>Mucor</i> sp.	2(4.76)	4(11.43)
<i>Pseudomonas</i> spp	7(10.94)	3(9.09)	<i>Penicillium</i> sp.	19(45.24)	5(14.29)
<i>Aeromonas</i> sp.	3(4.69)	2(6.06)	<i>Rhizopus</i> sp.	1(2.38)	2(5.71)
<i>Escherichia coli</i>	4(6.25)	1(3.03)	<i>Fusarium</i> sp.	0(0.00)	1(2.86)
<i>Bacillus</i> sp.	6(9.38)	7(21.21)	<i>Cladosporium</i> sp.	5(11.90)	4(11.43)
<i>Serratia</i> sp	4(6.25)	4(12.12)	<i>Metshnikowia</i> sp.	0(0.00)	3(8.57)
<i>Klebsiella</i> sp.	4(6.25)	0(0.00)	<i>Candida</i> sp.	2(4.76)	2(5.71)
<i>Vibrio</i> sp.	3(4.69)	0(0.00)	<i>Cryptococcus</i> sp.	2(4.76)	1(2.86)
<i>Enterococcus</i> sp.	3(4.69)	0(0.00)	<i>Leucosporidium</i> sp.	1(2.38)	2(5.71)
<i>Citrobacter</i> sp.	1(1.56)	1(3.03)	<i>Rhodotorula</i> sp.	1(2.38)	1(2.86)
<i>Staphylococcus</i> sp	4(6.25)	4(12.12)	<i>Aspergillus</i> sp.	4(9.52)	3(8.57)
<i>Streptococcus</i> sp	2(3.13)	0(0.00)	<i>Alternaria</i> sp.	0(0.00)	2(5.71)
<i>Entrobacter</i> sp	3(4.69)	1(3.03)	<i>Geotrichium</i> sp.	0(0.00)	2(5.71)
<i>Shiegella</i> sp	4(6.25)	3(9.09)	<i>Saccharomyces</i> sp.	3(7.14)	2(5.71)
<i>Corynebacterium</i> sp	2(3.13)	2(6.06)	<i>Trichoderma</i> sp.	2(4.76)	1(2.86)
<i>Acinetobacter</i> sp	2(3.13)	1(3.03)			
<i>Chromobacterium</i> sp	2(3.13)	0(0.00)			
<i>Flavobacterium</i> sp	1(1.56)	2(6.06)			
<i>Micrococcus</i> sp	2(3.13)	1(3.03)			
<i>Alcaligenes</i> sp	3(4.69)	0(0.00)			
<i>Norcadia</i> sp	1(1.56)	0(0.00)			
Total	64 (100%)	33 (100%)	Total	42 (100%)	35 (100%)

The mean values or levels of PAHs in surface water and *Pomadasy commersonii* as analyzed are shown Figure 5 and 6 respectively. The PAH levels in surface water ranged from 0.003mg^l⁻¹ to 0.005mg^l⁻¹ with a

mean value of 0.004mg^l⁻¹. The PAH level of *Pomadasy commersonii* ranged from 0.02mg^{kg}⁻¹ – 0.04mg^{kg}⁻¹ with a mean value of 0.031mg^{kg}⁻¹.

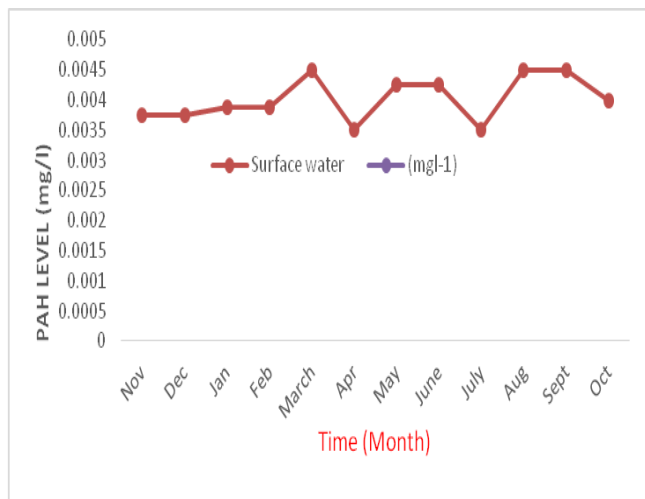


Fig. 5: Changes in PAH Levels in Surface Water (mg/l) in Crude Oil Polluted Stations in Borikiri Wetlands

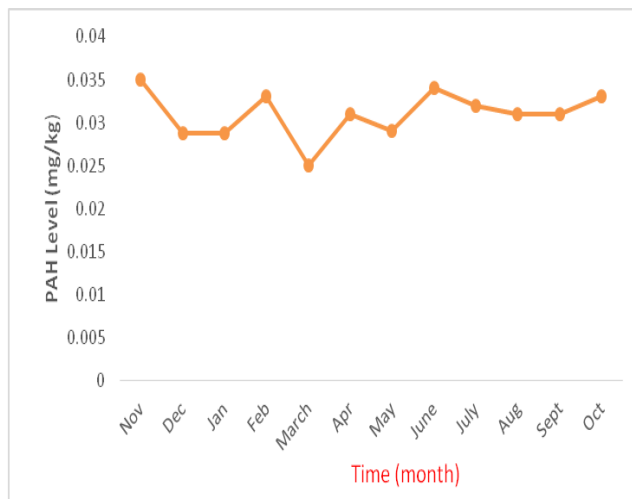


Fig. 6: Changes in PAH Level in Pomadasy commersonii (mg/kg) in the sample Stations in Borikiri Wetlands

The PAHs components are shown in Figures 7 and 8 and in Table 2 below. Specifically, Table 2 shows that out of the 14 PAHs components analyzed; pyrene,

Benzo (a) anthracene and Benzo (b) fluoranthene are not biomagnified by *Pomadasy commersonii*.

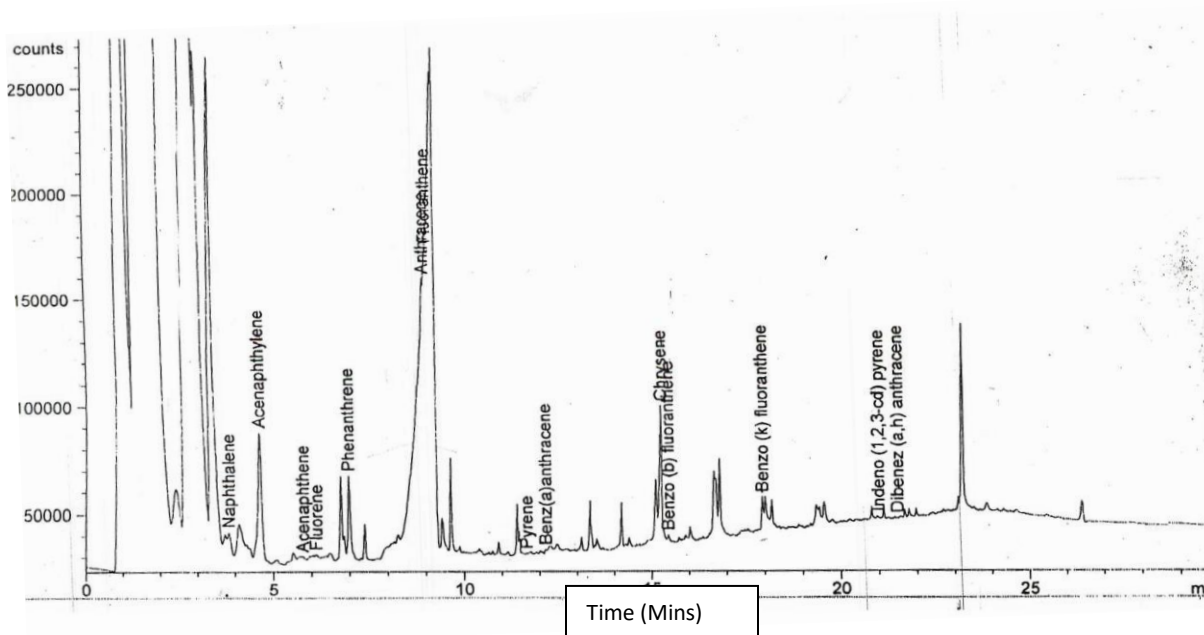


Fig. 7: Chromatogram of PAHs in Pomadasy commersonii from Borikiri Wetlands

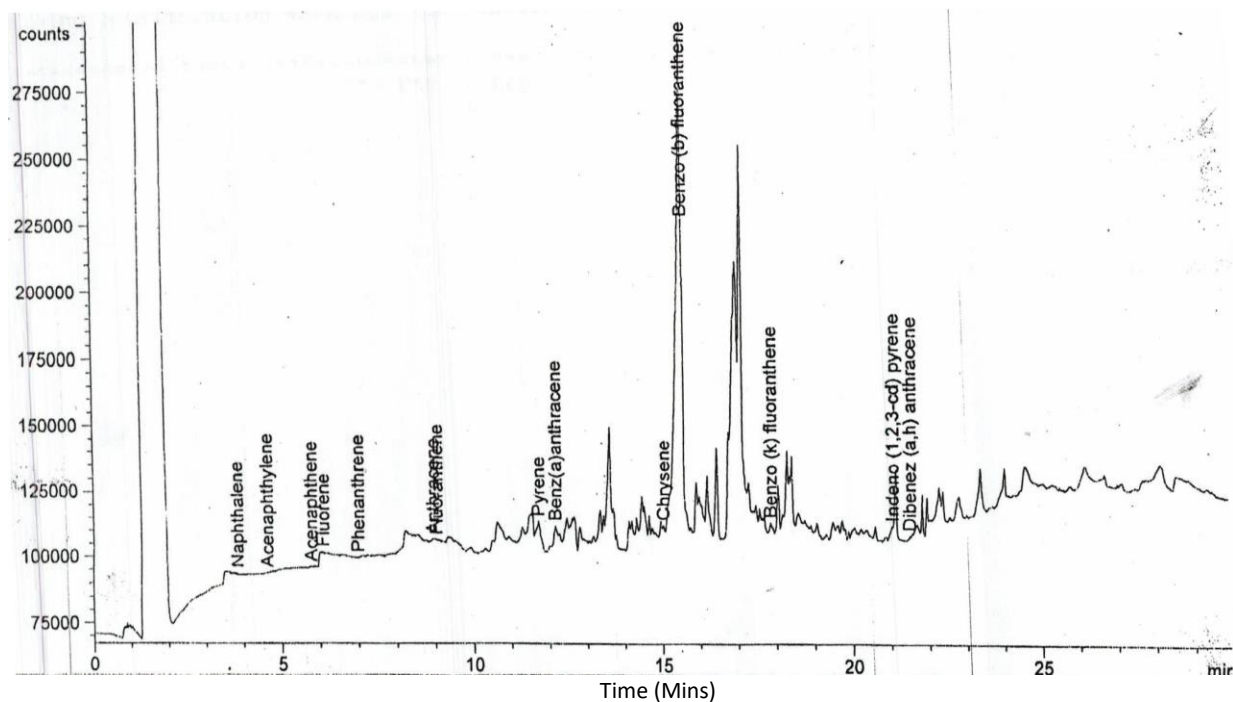


Fig. 8: Chromatogram of Polycyclic Aromatic Hydrocarbons in Surface Water from Borikiri Wetlands

Table 2: Mean values of PAHs Components in Surface Water and *P. commersonnii* in Ikpukulu Creek (Nov. 2019 – Oct. 2020)

PAH Components	Surface Water (mg ^l ⁻¹)	<i>Pomadasy commersonnii</i>	Inference
Naphthalene	0.000016	0.00045	Biomagnified
Acenaphthylene	0.00006	0.0034	Biomagnified
Acenaphthene	0.000075	0.00014	Biomagnified
Fluorene	0.00009	0.0002	Biomagnified
Phenanthrene	0.00003	0.0017	Biomagnified
Anthracene	0.00006	0.014	Biomagnified
Fluoranthene	0.00006	0.0066	Biomagnified
Pyrene	0.0003	0.000026	Biomagnified
Benzo (a) anthracene	0.00019	0.000068	Biomagnified
Chrysene	0.000099	0.0025	Biomagnified
Benzo (b) fluoranthene	0.0024	0.00048	Biomagnified
Benzo (k) Fluorathene	0.00015	0.00063	Biomagnified
Indeno (1,2,3) (d) pyrene	0.0002	0.0006	Biomagnified
Dibenzo (a, k) anthracene	0.00024	0.00064	Biomagnified
Total	0.004	0.031	

Estimated Daily intake of food EDI using indeno 1,2,3, (cd) pyrene

$$= \frac{0.0006\text{mgkg}^{-1} \times 0.1\text{kg}}{70\text{kg}} \dots\dots\dots\text{Equation 1}$$

$$= 0.00000086\text{mg/kg/day}$$

Reference Estimated Daily intake of food:

$$\text{EDI} = \frac{\text{Mass of reference contaminant (mgkg}^{-1}) \times 0.1\text{kg}}{70\text{kg}}$$

.....Equation 2

0.1 kg = 100g = mass of seafood ingested (USEPA, 2011)

70kg = mass of an average man

Reference concentration = 0.005mg/kg

Discussion

This study has revealed the microbiological and polycyclic aromatic hydrocarbon (PAH) profile of *Pomadasy commersonnii* harvested from crude oil impacted creeks and of the surface water in Borikiri, Port Harcourt, Nigeria. The results of Total heterotrophic bacterial count (THBC) from these crude oils impacted surface water falls within 1.2×10^5 Cfu l^{-1} - 5.0×10^7 Cfu l^{-1} (Fig 1) with non-exceeding $A \times 10^7$ Cfu l^{-1} where $1 \geq A \leq 9$. Unimke et al. (2014) recorded bacterial densities of 1.2×10^5 - 2.2×10^6 Cfu l^{-1} for THBC and $1.2 - 1.3 \times 10^5$ Cfu l^{-1} hydrocarbon utilizing bacteria (HUB) for surface water samples in crude oil impacted Imo River estuary and 2.6×10^7 - 2.8×10^9 THBC. Total heterotrophic fungi densities of surface water 5.3×10^5 cfu l^{-1} (Fig. 2); by implication, it simply means that the density of fungi available to metabolize the crude oil in the surface water. These results showed the presence of high active indigenous microbial flora in the fish and surrounding environment.

Results from this research work shows that, the mean bacterial density in surface water from the research sites, 1.6×10^7 Cfu l^{-1} , 1.2×10^4 Cfu l^{-1} (0.075%) are hydrocarbon utilizers (Fig 4). According to Zobell

∴ Reference EDI of *Pomadasy commersonnii*

$$= \frac{0.005\text{mgkg}^{-1} \times 0.1\text{kg}}{70\text{kg}} = 0.000007\text{mg/kg/day}$$

$$\text{Cancer Risk Index} = \frac{0.00000086\text{mg/kg/day}}{0.000007\text{mg/kg/day}}$$

$$= 0.12$$

The Contamination factor (c¹f):

$$C^1f = \frac{\text{Mass of contaminant (mgkg}^{-1})}{\text{Mass of Standard (mgkg}^{-1})} =$$

$$\frac{0.0006\text{mgkg}^{-1} \text{ indeno 1,2,3cd pyrene}}{0.005\text{mgkg}^{-1}} \dots\dots\dots\text{equation 3}$$

Contamination factor of PAH (indeno1,2,3,(cd) pyrene) in *Pomadasy commersonnii* = 0.12

(1971), hydrocarbon utilizing bacteria take advantage of the presence of crude oil in the ecosystem and use them as energy and carbon source since carbon in the marine environment is a limiting factor being in the concentration of 20mg/l. According to this research, one bacterial cell can decompose 5.0×10^{-12} mg crude/hr in a litre of water. Going by this, a bacterial density of 1.2×10^4 cfu l^{-1} which is equivalent to 1.2×10^7 Cfu l^{-1} of the HUB recorded in this research, is a sure way of carrying out a natural attenuation method of bioremediation in the surface water.

It has been reported that, fungi biodegrade PAHs to carbon (IV) oxide and water. A fungal density of 9×10^4 reported in this present study which is included in the range of fungal density is recorded by Zobell (1971), as enough density for microbial crude oil degradation.

The following hydrocarbon utilizing bacteria were identified from the fish and water samples: *Pseudomonas* spp, *Bacillus* spp, *Aeromonas* sp, *Proteus* sp, *Serratia* sp, *Citrobacter* sp, *Staphylococcus* sp, *Escherichia coli*, *Corynebacterium* sp, *Acinetobacter* sp, *Flavobacterium* sp, *Micrococcus* sp, *Enterobacter* sp. While, the following fungal isolates were also identified; *Penicillium* sp, *Mucor* sp, *Rhizopus* sp, *Fusarium* sp, *Cladosporium* sp,

Metshnikowia sp, *Candida* sp, *Cryptococcus* sp, *Leucosporidium* sp., *Rhodotorula* sp., *Aspergillus* sp., *Alternaria* sp., *Geotrichium* sp., *Saccharomyces* sp., *Trichoderma* sp. The bacterial isolates from this research samples like *Micrococcus* sp., *Pseudomonas aeruginosa* and *Bacillus* sp. are reported to possess catabolic genes catechol 2, 3 dioxygenase (nah H), alkane mono-oxygenase (alk B) and catechol 1, 2 dioxygenase (cat A) genes which are PAHs degrading (Oboh, 2006). *Micrococcus*, *Bacillus* and *Citrococcus* are reported by Atlas and Bartha (1992) as alkaliphilic oil utilizing bacteria and are responsible for biodegradation and eventual clean-up of spills in our environment. According to Adekunle and Adebambo (2007), microbial communities exposed to hydrocarbons become adapted, exhibiting selective enrichment and genetic changes. This adaptation makes them extremophiles. These microorganisms are able to withstand this environmental stress due to their ability to release stress mitigating compounds e. g. alkaloids, peptides, and exopolysaccharides which enable them use hydrocarbons as carbon and energy sources. Bacterial densities recorded from other crude oil impacted ecosystems are similar to that reported by Obiukwu and Otokunefor (2014) in these research site viz: Ekerekana creek, Okari Creek, Ogoloma, Edemebikri, Elechi Creek 1.4×10^5 - 5.0×10^7 Cfuml⁻¹ for surface water; 2.5×10^4 - 2.8×10^9 Cfug⁻¹ for sediment. The bacterial isolates were *Pseudomonas* sp., *Bacillus* sp., *Staphylococcus* sp., *Klebsiella* sp., *Vibrio* sp., *E. coli*, *Flavobacterium* sp., *Micrococcus* sp., *Streptococcus* sp. (Obiukwu and Otokunefor, 2014). *Chromobacterium* sp. and *Corynebacterium* sp. were not recorded in these creeks although crude oil impacted. It could thus be said that, some bacteria may be indigenous to some ecosystems just like some macrofauna. Otokunefor and Obiokwu (2014) also recorded a bacterial density of 1.2×10^4 - 1.6×10^9 Cfuml⁻¹ from refinery effluents. It shows that bacterial densities from refinery effluents are in close proximity with crude oil impacted ecosystems. The bacterial isolates have representatives which can biodegrade all the fractions of total petroleum hydrocarbons (TPH) presented in this research; *Pseudomonas* sp., *Bacillus* sp., *Enterobacter* sp., *Staphylococcus* sp., *Citrobacter* sp., *Micrococcus* can biodegrade n – alkanes C₁₄ – C₃₀ as well as resins and asphaltenes being equipped with catabolic genes (nah H, cat A and Alk B).

The Hydrocarbon utilizing Bacteria (HUB) and their percentage (%) occurrences' in the surface water samples and in the fish *Pomadasys commersonni* samples during this study were: *Proteus* sp. 3.03%, *Pseudomonas* spp 9.09%, *Aeromonas* sp. 6.06%, *Escherichia coli* 3.03%, *Bacillus* sp. 21.21%, *Serratia* sp. 12.12%, *Citrobacter* sp. 3.03%, *Staphylococcus* sp. 12.12%, *Enterobacter* sp 3.03, *Shiegella* sp 9.09%, *Corynebacterium* sp 6.06%, *Acinetobacter* sp 3.03%, *Flavobacterium* sp 6.06%, and *Micrococcus* sp 3.03% (Table 1).

The fungi isolated from this research sites and their percentage (%) occurrences were: *Mucor* sp. 11.43, *Penicillium* sp. 14.29, *Rhizopus* sp. 5.71, *Fusarium* sp. 2.86, *Cladosporium* sp. 11.43, *Metshnikowia* sp. 8.57, *Candida* sp. 5.71, *Cryptococcus* sp. 2.86, *Leucosporidium* sp. 5.71, *Rhodotorula* sp. 2.86, *Aspergillus* sp. 8.57, *Alternaria* sp. 5.71, *Geotrichium* sp. 5.71, *Saccharomyces* sp. 5.71, *Trichoderma* sp. 2.86 (Table 1).

Candida sp produces the biosurfactants, sophorolipids and lipomannan that can biodegrade alkanes of carbon chains C₁₀ – C₁₆ the naphthalene, gasoline and kerosene fractions. *Rhodotorula* sp is known as a nuisance in the aviation industry as they contaminate aviation fuel and metabolize it causing adverse effect on the machine. *Rhodotorula* sp is however a potent petroleum degrader and is good news in crude oil impacted ecosystem (Obiukwu and Otokunefor, 2014).

Total heterotrophic fungi densities of surface water 5.3×10^5 cfuml⁻¹; by implication, it simply means that the density of fungi available to metabolize the crude oil in the surface water. Separating the yeast densities from fungi, the yeast density of surface water has a mean of 4.7×10^5 Cfuml⁻¹ of which 09.4% (4.4×10^3 Cfuml⁻¹) is hydrocarbon utilizing Yeast (HUY). The HUY metabolize the petroleum products, they are reduced to the forms other fungi can metabolize. The biota also showed a yeast density ranging from 1.4×10^3 - 4.5×10^3 Cfug⁻¹ of hydrocarbon utilizers. The yeast density of 4.6×10^4 Cfug⁻¹ from *Pomadasys commersoni* (gills), 3.5×10^3 cfug⁻¹ from *Pomadasys commersoni* (tissue). Unimke et al. (2014) records fungi densities of surface water 1.2×10^5 Cfuml⁻¹; Hydrocarbon utilizing fungi of surface water 7.2 - 8.9

$\times 10^4$ Cfuml⁻¹ from Imo River, a crude oil impacted estuary.

The fungi isolated from Ekerekana Creek, Okari, Ogoloma, Edemebikri and Elechi Creek were *Penicillium* sp, *Aspergillus* sp, *Fusarium* sp, *Candida* sp, *Mucor* sp, and *Saccharomyces* sp. The research of Otokunefor and Obiokwu (2014) did not isolate *Metchikowi* sp, *Leucosporidium* sp, *Alternaria* sp, and *Geotrichium* sp. Fungal isolates identified by Douglas and Longjohn (2022) include: *Aspergillus flavus*, *Rhizopus arrhizus*, *Aspergillus niger*, *Rhizopus* sp, *Mucor* sp, *Microsporium* sp, *Cunninghamella* sp and *Candida* sp. from Bonny River.

This research however, similarly isolated *Penicillium* sp as the most dominant fungal isolate.

The PAH level ranged from 0.003 - 0.005 m/l in all the stations with a mean of 0.004 mg/l. There is no significant difference in the PAH levels in the various stations in the different month with $P \leq 0.05$ probability. Daka et al. (2020) recorded 0.009 - 0.0122 mg/l in surface water in Olugbobiri, Bayelsa State. The PAH level in the research site shows a contamination factor (c^1f) of 0.571 which implies low contamination since $c^1f < 1$

Edlund and Jansson (2006) reported *Pseudomonas* and *Flavobacterium* spp as the most dominant species of bacteria in an environment with very high concentrations PAH. *Pseudomonas* spp are reported to reduce PAH components of high molecular weights fractions (Chrysene, Benzo (a) pyrene, indeno 1, 2, 3, (cd) pyrene and benzo (g,h,l) phrylene) (Temitayo et al., 2019). In a consortium of five strains of bacteria, Temitayo et al. (2019) reported complete removal of indeno (1,2,3 cd) pyrene, Benzo (k) flouranthene, Benzo (b) fluoranthene and dibenzo (a,h) anthracene as well as significant reduction of all PAHs. Their presence in this research could be the reason for such small quantity of PAHs.

The levels of PAHs in surface water and *Pomadasy* *commerconnii* are shown in Figures 5 and 6 respectively. The PAH levels in surface water ranged from 0.003mg⁻¹ to 0.005mg⁻¹ with a mean value of 0.004mg⁻¹. The PAH level of *Pomadasy* *commerconnii* ranged from 0.02mkg⁻¹ – 0.04mkg⁻¹ with a mean value of 0.031mkg⁻¹. There is no

significant difference in the monthly PAH levels of *Pomadasy* *commerconnii* with $P < 1$ at 0.05 probability level.

The PAHs components are shown in Table 4 and Figs. 7- 8. Specifically, Table 2 shows that out of the 14 PAHs components analyzed; pyrene, Benzo(a) anthracene and Benzo(b)fluoranthene were not biomagnified by *Pomadasy* *commerconnii*. This implies that, it biominifies Benzo(a)anthracene, pyrene and Benzo(b) fluoranthene.

It thus has a mechanism of utilizing these PAHs as energy sources or it has a device of eliminating these PAHs from its system. Also, anthracene is its highest contaminant (0.14mkg⁻¹ i.e. 45.15%) while its least contaminant is pyrene (0.000026mkg⁻¹ i.e. 0.0084%). PAH is lipophilic and easily accumulate in fatty tissue of fish.

According to EU No 835/2011, benzopyrene is the PAH marker using dietary administration. Initially, PAH 4 (Benzo(a)pyrene, chrysene, benzo(a) anthracene, and benzo(b) fluranthene) were used as a PAH marker. The EU standard stipulates a 0.006mg⁻¹ limit of Benzo(a) pyrene in seafood.

Currently, EU has indicated a 5µgkg⁻¹ (0.005mkg⁻¹) limit of indeno 1,2,3, (cd) pyrene as the most suitable indicator of PAH contamination of sea food. Going by this stipulation, the level of indeno (1,2,3-cd) pyrene in *Pomadasy* *commerconnii* is 0.0006mkg⁻¹ which is lower than the EU standard and giving a c^1f of 0.12 (0.0006mkg⁻¹/ 0.005mkg⁻¹) giving a $c^1f < 1$ implying low contamination.

Research has shown that raw fish from unpolluted water do not contain detectable concentrations of PAHs. PAH concentrations in fish are usually low due to rapid metabolization, while higher molecular weight PAHs, do not seem to accumulate in fish (Eisler, 1987). The tendency of fish to metabolize PAHs extensively and rapidly may explain why benzo(a)pyrene, for example, is frequently undetected, or only detected in low concentrations in livers of fish from environments heavily contaminated with Polycyclic aromatic hydrocarbons (PAHs).

Conclusion

In spite of the ongoing crude oil marketing in the coastal area of Ikpukulu Creek, and the continuous discharge of crude oil into the surface water, *Pomadasy commersonnii* is not affected enough for its consumption to be of health risk by Polycyclic aromatic hydrocarbon contamination. Hence, due to the recalcitrant and carcinogenic nature of PAHs in the environment, it is important to create awareness of the danger of this substance in the environment and how to reduce activities that may lead to increase in its concentration in the environment to the lowest minimum. Therefore, there is need for proper enlightenment campaign to educate the people of the areas on the dangers of the increase in these pollutants in the environment, which may lead to biomagnifications of PAHs in fish and biomagnify in food chains. Results from this study indicate high microbial populations of active hydrocarbon utilizing bacteria and fungi, some of which have been implicated in bioremediation of hydrocarbons and PAHs.

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