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# A Review of the Role of Anthropogenic Effects on Microorganisms in Soil

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#### Authors' contributions

This work was carried out in collaboration between all authors. Author WMP designed the study and wrote the first draft of the manuscript. All authors managed the literature searches. All authors read and approved the final manuscript.

#### Article Information

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**Review Article** 

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# ABSTRACT

Anthropogenic activities are progressively releasing pollutants to the soil, including heavy metals and other pollutants emanating from industries and overpopulated human settlements. Pollution by heavy metals and other physicochemical materials is gradually becoming an area of concern, especially in sub-Saharan Africa. The paper reviews the effects of anthropogenic activities on soil ecosystem integrity including physicochemical parameters as well as biological parameters in the soil. The review focuses on impact of the application of fertilisers in soil and seasonal influence in levels of physicochemical parameters in soil ecosystem as well as microorganism population shifts in soils affected by pollution as a need to alleviate soil contamination. The review established gaps to further understanding of the relationship between soil physicochemical parameters, soil biological parameters and influence of seasonality in the resilience of the soil ecosystem as a way to foster the integrity of soil. Keywords: Soil ecosystem integrity; soil microorganisms; soil physicochemical parameters; anthropogenic activities; seasonal influence.

#### **1. INTRODUCTION**

The concept framework (Fig. 1) outlines a structure for understanding how urbanisation, industrialisation and agricultural activities impact on soil ecosystem. The framework highlights the effects of urbanisation, industrialisation and agricultural activities on soil ecosystem integrity. Urbanisation is characterised with increased human population. industrialisation is characterised by increased emissions and agricultural activities involve overuse or misuse of land. Jointly, urbanisation, industrialisation and agricultural activities produce wastes that end up contaminating the environment, consequently leading to acidification, eutrophication, siltation, accumulation toxic substances. of

loss of biodiversity and loss of ecosystem processes.

Fig. 2 depicts pathways between soils and both abiotic and biotic components in the environment and shows how soil contaminants find their way to human diet. Polluted soils are interlinked in various ways to human diet. Crops that are planted in contaminated soils are likely to be contaminated in turn. Runoff from polluted soils contaminates water bodies including rivers, lakes and the sea where animals and humans abstract water for consumption. These calls for essence to sporadically determine levels of pollution of soils in order to ensure that adequate measures are put in place to ensure that wastes are disposed of in an amicable manner hence ensure

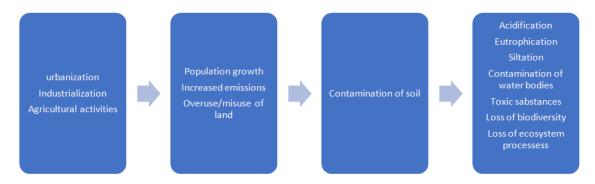


Fig. 1. Showing effects of urbanisation, industrialisation and agricultural activities on soil ecosystem integrity

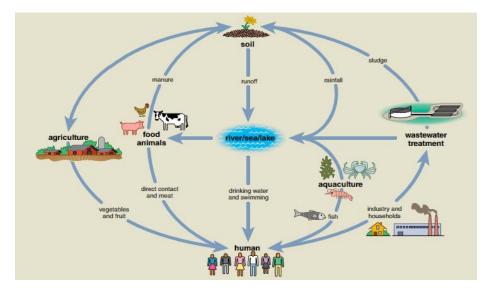


Fig. 2. Showing linkage between polluted soils and human health

soil ecosystem integrity is maintained. This study describes levels of pollution in soils and determines the impact of pollution on biodiversity and distribution of soil microorganisms in the soil.

## 2. BACKGROUND INFORMATION

Pollution is the introduction of foreign material into an environment thereby compromising the integrity of that environment. According to World Health Organization (WHO), mismanagement of chemicals and wastes cause pollution of air, water and soil. Various advanced effects have accompanied oil production activities in Niger Delta region after discovery crude oil in 1950s. Discovery of crude oil was accompanied with increased pollution associated with population and industrialization. Hence the native human inhabitants and other biotic communities have been directly or indirectly affected by pollution from oil production activities [1], water and soil ecosystems [2,3] and air [2,3]. Pollution is a major threat to biodiversity and ecosystem integrity in the environment [4]. In addition, it is a risk factor to human health and can cause heart diseases. stroke. cancer. infections. developmental disabilities and neurological disabilities among others. Polycyclic aromatic hydrocarbons (PAHs) are carcinogenic and can induce malignant tumors [5].

Among economic activities in Rivers State, Niger delta include; exploration, drilling, processing of crude oil and farming. Intensification of economic activities has increased chances of contamination of the environment with petroleum hydrocarbons. A vast number of additives are used in the exploration and processing of oil and can be sources of contamination of the environment. Such petroleum contaminants are classified as priority pollutants [6]. These pollutions can either occur onshore or offshore. Petroleum hydrocarbons have aliphatics. monoaromatics. polyaromatics. resins. (Benzene, Toluene, Ethylbenzene and Xylenes) BTEX, PAHs and many additives used in the process. Most of the additives are teratogenic, carcinogenic, hemotoxic, anoxic, and metabolic retarders which can cause acute or chronic effects [7]. Microorganisms have been reported to evolve and adapt to these environments by development of mechanisms of degradation of these hydrocarbons [8]. Degradation of hydrocarbons is tough, persistent and can cause irreversible changes to the environment.

#### 3. CHEMICAL SOIL HEALTH INDICATORS

Industrialization, urbanization and agricultural activities are progressively releasing pollutants to soil, including heavy metals. Pollution by heavy metals is gradually becoming an area of concern especially in sub-Saharan Africa, including Nigeria [9]. The major heavy metals of concern include; Arsenic, Lead, Zinc, Cadmium, Nickel, Copper and Mercury [10]. Heavy metals occur naturally in undisturbed soils. These occurrence of heavy metals in soils can be attributed to characteristics of the parent rock, environmental management practices, particle size distribution, soil age and influences from the air that comes in contact with soil. Nickel (Ni) and Zinc (Zn) are essential for plant growth, while Pb is considered a stern environmental contaminant [11]. Heavy been used expansively in metals have herbicides, pesticides farm chemical or throughout the world, to protect crops by eliminating pests and thereby increasing agricultural output. This could be source of residual heavy metal content in agricultural soils [12].

The mostly applied pollution indicator is a chemical concentration. Findings by Adamu [13] reveal that soils around areas of Minnna Railway Station, in Niger State of Nigeria, are contaminated with Zn, Cu and Pb. The soils were classified as moderate to highly contaminated [13]. Pollution of soils around the Railway Station demonstrates the effect of industrialization on soil integrity in the industrialized areas. This was attributed to leakages in fuel storage tanks as well as piping in trains or during repairs of machines. Eventually a cumulative effect can be achieved after a period of spillage. A similar study was conducted by Owolabi and Hekeu [14], in Lagos and Ota in Nigeria. Heavy metal resistant bacterial isolates; Aerononas spp. Anthrobacter Corynebacterium spp, spp, Pseudomonus spp and streptococcus spp were isolated and tested in the laboratory and analyzed against levels of tolerance to different concentrations of Cd, Cr and Pb. Bioremediation of environments that are contaminated with aromatic hydrocarbons such as BTEX has gained attention. Pseudomonus spp has been closely associated with Benzene, Toluene, Ethylbenzene and Xylenes (BTEX) contaminated environments and have been said to be useful in decontamination of the environments that are polluted with BTEX [15]. Rate of degradation of BTEX by organisms can be affected by presence of other contaminants present in the soil [16].

Osterreicher-Cunha [16] described ethanol to reduce the rate of degradation of BTEX and bioventing reduced effect of ethanol on reduction of rate of degradation of BTEX. Benzene, Toluene, Ethylbenzene and Xylenes (BTEX) often creates an anoxic environment and consequently anaerobic bioremediation techniques are preferred. Aromatic compounds for example Benzene and Toluene are electron donors for nourishment of bacteria [15].

## 4. MICROORGANISMS AS SOIL HEALTH INDICATORS

Typically, soil contains 10<sup>8</sup> to 10<sup>10</sup> bacteria per gram of soil [17]. Evolution of complex microbial communities in soil is driven by both abiotic and biotic pressures. Microorganisms continue to adapt their ever-changing environment by acquiring genes that previously were unavailable through either point mutations that alter regulation of enzyme kinetics or genes acquired through horizontal or lateral gene transfer [18]. Microorganisms change rapidly and unlike macroscopic organisms. However. microorganisms cannot be observed directly, but can be cultured and their morphological and selected physiological characteristics used to classify them.

Bacteria are least structurally complex microorganisms but offer greatest diversity and metabolic flexibility. Bacteria are microscopic single celled organisms that can inhabit varied ecosystems in the environment. Bacteria can be arouped into decomposers or heterotrophs. mutualists. pathogenic, lithotrophs or chemoautotrophs [19]. Decomposers rely on carbon materials; pesticides, hydrocarbons and pollutants. Mutualists form close other relationship with plants for example nitrogen fixing bacteria. Pathogenic bacteria include; and Xymomonas, Erwinia spp, other agrobacterium that cause gall formation in plants. Lithotrophs or chemoautotrophs, obtain their energy from compounds of nitrogen, sulfur, iron or hydrogen but not carbon compounds [20]. Bacteria from the four groups are very important integral component of the ecosystem. Bacteria are part of nutrient cycling and produce substances that bind soils together into small aggregates. Bacteria competes with disease causing organisms, nitrogen fixing bacteria form symbiotic relationship with roots of legumes, nitrifying bacteria transform NH<sub>4</sub><sup>+</sup> to NO<sub>2</sub><sup>-</sup> then to NO3<sup>-</sup>, while denitrifying bacteria convert nitrate to nitrogen (N<sub>2</sub>) or nitrous oxide (N<sub>2</sub>O) gas.

Actinomycetes decompose a wide range of compounds that are rigid to decompose, for instance, cellulose and chitin, as they are active at high pH [21,22].

Soil microorganisms play a vital role in nutrient cycling [23]. They are actively involved in cycling of Nitrogen, Sulphur and Phosphorus. The symbiotic relationship between microorganisms and plants contributes above 20% of all plant nitrogen requirements [24]. Microorganisms degrade organic materials into mineralized carbon dioxide, water, mineral nitrogen and phosphorous among others. Thus. microorganisms play a vital role in soil formation and maintenance of soil ecosystem integrity by production of secretions and their cellular debris upon their death. Biological adaptation and responses to environmental changes are among factors that drive need to assess soil integrity and evolution of microorganisms to suit their evolving habitats. Evaluation of diversity of microorganisms in soil is a reliable indicator of soil integrity and changes in soil integrity [25].

Microorganisms can easily be used as soil health indicators. Fecal indicators indicate the presence of fecal contamination for example Escherichia coli [26]. In addition, microorganisms can also be used as process indicators, for instance total heterotrophic bacteria or total coliforms can indicate disinfection [27]. The total coliform includes; Escherichia, Enterobacter, Klebsiella and Citrobacter species. Total coliforms are easy to detect and specifically include all aerobic, gram-negative, non-spore forming and rodshaped bacteria that produce gas upon fermentation of sugars in culture media within 48 hours at 35°C. Bacteria have been proven to grow on natural surfaces [28]. Microbes that inhabit harsh areas like polluted areas are prone to develop abilities that eventually enable them to survive, hence microorganisms are very resilient. Bacteria play a role in degradation of hydrocarbons in polluted areas [8]. These include bacterial species of genera Azotobacter, Pseudonomus, Micrococcus, Cellulomonas among others. Thus, microorganisms have been employed in degradation of toxins in the environment and have also been used in microbial enhanced oil recovery [29].

#### 4.1 Total Heterotrophic Bacteria

The total heterotrophic bacteria include all bacteria that utilize organic nutrients as a source of their energy [30]. Heterotrophic bacteria are

found all over the world in water, soil, air, food, and on vegetation. Heterotrophic microorganisms grouped photoheterotrophs, can be as organotrophs chemoheterotrophs. and lithotrophs [31]. Examples of heterotrophic bacteria include Bacillus spp, Criptobacter spp, Escherichia Coli. Micrococcus spp, Pseudomonus Serratia spp, spp and Staphylococcus spp among others [32]. Heterotrophic plate count refers to particular microbes isolated by a particular standard method of isolation by use of sufficient media time and conditions. Other terminologies have been used to refer to heterotrophic bacteria including; standard plate count, total viable count, total count, plate count, total bacterial count, water plate count, aerobic mesophilic count, colony count and autochthonous flora [30]. Heterotrophic bacteria refer to all bacteria requiring organic nutrients for growth, however, diverse methods target only particular subpopulation of total heterotrophs in a sample. It is difficult to differentiate which subpopulation includes pathogenic or nonpathogenic organisms [30].

# 4.2 Total Fungi

Fungi are chemoheterotrophs and mostly obligatory, which means they derive their energies from chemicals, and need to devour other organisms in order for them to survive [33]. Fungi are physically larger organisms as compared to bacteria which are most abundant. Similar to protozoans; 18S rRNA gene is being used for fungal identification, by use of internal subscribed spacer (ITS) [34]. Fungi are primarily found in soil but can be described as ubiquitous. Fungi are important in industrial processes of biotechnology and fermentation. Anaerobic fungi ferment sugars to produce ethanol, wine, lactic acid and acetic acid. They also produce secondary metabolites which have revolutionized agriculture and medicine through production of penicillin by Penicilliun notatun, griseofluvin produced by *Penicillium griseofulvum* and biotechnology.

It is hypothesized that fungi help reduce competition from other microorganisms for nutrients [35]. Saprophytic fungi are known to secrete extracellular enzymes that breakdown complex polymers; cellulose, lignin and chitin to simple forms [35]. Fungi has been found to degrade a variety of complex environmental contaminants; *Aureobasidium pullulans*  decgrades polyvinyl chloride [36], Penicillium, Stachbotrys, Allescheriella, and Phlebia degrade petroleum hydrocarbons aromatic and agrochemicals [37,38]. Mycorrhizae form symbiotic important relationship the in environment. They colonize plant roots in exchange for CO<sub>2</sub> from the plant and the mycorrhizal fungi in turn solubilize phosphorous and bring soil nutrients; nitrogen, phosphorous and water. For example, the ectomycorrhizae that grows on root surfaces and the endomycorrhizae that grows within the root cells of plants. Parasitic and pathogenic fungi are a major problem in farming as it has been associated with reduction of yields and economic losses. Pythium, Rhizoctonia and Verticillium are pathogenic to roots of plants [35]. Fungal genera that has been isolated and identified in different levels of industrialization (highly, moderate, low and very low) in Port Harcourt, where Mucor, Aspergillus, Candida, Penicillium, Rhizopus, Trichosporon and Sacharomyces spp have been isolated and identified [32]. Industrialization influences distribution and abundance of microorganisms in soil [32].

# 4.3 Distribution of Microorganism Communities in the Environment

Microorganism communities are complex environmental components that are necessary for maintenance of soil ecosystem integrity [39]. Soil aggregates and pores within and around them create microhabitats that support different microorganism communities [40]. Within soil habitats, there is difference in quantity and chemistry of organic substances that are likely to be drivers of the microbial community difference [40]. Environmental conditions within and between soil aggregates such as physicochemical parameters are varied and thus resulting in diverse microorganism community among the different soil microhabitats [40]. Soil microorganisms plays vital roles in the soil ecosystem; regulates release of nutrients and maintenance of soil structure and texture [41]. When microorganisms are well established within soil aggregate, microbial structure and microbial community plays an important role in processing and protecting soil organic matter [40]. Soil microorganisms are associated with various biochemical processes and are important in maintenance of soil fertility [42], and also maintenance of ecosystem functions [43]. Microorganisms play a very important role in regulation of carbon dynamics in soil ecosystems [44,45]. Further, soil function, processes and integrity are influenced by soil microorganisms [46-48].

Changes in environmental conditions promote complex changes of soil microorganism composition. Effects on major communities are more detectable as compared to minor communities and can be realized through enrichment cultures including sulfate reduction, methane oxidation, halophilic and alkaliphilic and thermophilic responses among others [39]. Complex organic matter including phenols and alkyls has been proven as strong selective force on soil bacterial communities [40]. There is need to advance understanding influence of soil pollution and seasonality on soil microorganism community changes and cycling organic matter in soil [40]. Slight changes in environmental parameters cascades change in microorganism community in same environment and therefore determines the soil ecosystem integrity [39]. Shifts in composition of microbial community is followed up by changes in extracellular enzyme activities in environment which affects soil ecosystem integrity [49]. Specific enzyme activities can have a positive or a negative correlation with soil pH [49]. This suggests that microorganisms in environment adapt to environmental stresses with time through enhancing availability of C and other nutrients [49]. Variation in microorganism diversity in response to increasing pollutant deposition in environment remains uncertain [50], and calls for further studies. Addition of pollutants like nitrogen (N) and change in levels of pH have been described to influence ecosystem functions and can compromise the soil ecosystem integrity [50]. For instance, microorganism biomass is influenced by addition of N, [51-53,50]. Effects of pollutants on microorganism distribution varies among different ecosystems [54,50]. Addition of N to soil decreases soil pH [55], and consequently reduced pH enhances leaching of magnesium and calcium and mobilization of aluminum [50]. Aluminum poisoning and calcium and magnesium limitation to microorganism growth under N condition may affect diversity of soil microorganisms [56,50]. Increased levels of N influence fungal and bacterial diversities in soil ecosystems [50]. Negative effect of addition of N has been reported, which suggested an overall decline in microorganism biodiversity [50].

Fungi are considered to have higher carbon use efficiency as compared to bacteria [57,48]. Composition of soil microorganism community; fungi and bacteria can potentially help describe the unique variations in soil ecosystem integrity [48]. Treatment of soils with fertilizers influences richness and distribution of microorganisms in different soil ecosystems [48]. Chemical NPK fertilizer can influence distribution of soil microorganisms [48]. It is important to conduct studies in soils of cropping system because healthy soils foster food security [42].

Studies reveal that distribution of microorganisms in environment are influenced by level of heavy metal pollution, while other studies reveal that biomass of microorganisms in environment is not influenced by heavy metal pollution levels in environment [41]. Other studies reveal that microorganism biomass in environment can be influenced by other physicochemical parameters rather than heavy metals. Frequent quantification of heavy metals and other physicochemical parameters on their influence on distribution of microorganisms will give а clear soil representation of influence of heavy metals on distribution of microorganisms in natural ecosystems [41].

## 5. EFFECT OF PHYSICOCHEMICAL PROPERTIES ON DISTRIBUTION OF MICROORGANISMS IN SOIL

Physicochemical and biological properties of soil are affected by soil minerals. organic components and microorganisms [58]. Minerals, organic matter and microorganisms in soil ecosystems should rather be handled as dependent rather than independent entities [58]. Soil properties are influenced by cultivation, management, urbanization and industrialization and are responsible for declined and contaminated vields [42]. Activities of microorganisms soil depend in on physicochemical properties of that particular soil [59]. Land use types can affect soil physicochemical parameters and consequently affect soil microbial properties of soil [59]. Microorganism activities in soil are directly or indirectly influenced bv physicochemical parameters in soil [60,59] or weather conditions like precipitation and temperatures. Some of soil physicochemical parameters that are influenced by land-use practices include; TOC, Total Nitrogen and pH [59]. Excess N can acidify soils, deplete soil nutrients, alter amount of organic matter and affect biodiversity [49]. Intensified anthropogenic activities; use of fertilizers, combustion of fuels, cultivation of leguminous plants, among others, can increase N deposition in environment and it is therefore important to know how N levels in environment can influence distribution of microorganisms [49]. The pH and N levels can influence shifts in microorganism community composition in environment [49]. Long term exposure of microorganisms to N in environment can decrease microbial biomass in environment [49]. Fungi are sensitive to elevated N inputs and high levels of N can decrease abundance of fungi and thus a lower fungi/bacteria ratio. This is in contrast to N-rich but phosphorus and other nutrient limited environment [49]. Soils acidification can inhibit growth of microorganisms but fungi are more tolerant to acidic environments as compared to bacteria [61,49] which may reduce soil microbial biomass [62,63,49]. Low pH, elevated electrical conductivity and ammonium concentration mainly inhibits bacteria rather than fungi in environments of high N content [49]. Fungi are able to adapt to acidic environment and bacteria prefer neutral conditions [49]. Shifts in abundance and composition of microorganism community in soils are cascaded with changes in functional activities of microorganisms in various environments and affect soil ecosystem integrity [64,65,49]. Differential effects of activities on different landaffects distribution use types of soil microorganisms in different soil ecosystems found in urbanized, industrialized and agricultural areas [59].

# 5.1 Effect of Fertilizers on Changes of Microorganism Composition in Soil

Fertilizers are used to enhance harvest of crop in agriculture, but if misused can affect soil ecosystem integrity [66]. The expanse/capacity of different types of soil to lose nutrients and pollutants is an important factor affecting ability of soil to lose fertilizers [67]. This to a greater extent can be related to Al, Ca and Fe in a situation of inorganic and organic P and to cation exchange capacity in case of potassium and ammonium. This may consequently decrease loss of nutrients to water and air hence reduces availability of nutrients to crops. Further, value for fertilizers in soil, depend on several reasons; the ratio and nature of organic and mineral components of soil [68], length of growing season climate that determine crop uptake, weather conditions and history of application [69,45].

Distribution of soil microorganisms is influenced by concentration of pollutants. A study conducted at Ebony State, Nigeria, revealed that inorganic fertilizers contribute to spatial distribution of soil

microorganisms [70]. Microbial population decreases with increase in levels of soil contaminants [71]. Levels of cations in soil have also been attributed to influence development in bacteria. A study by Hai-Hang Li [72], used Rhodotorula glutinins and pseudomonas spp to demonstrate that Al <sup>2+</sup>, Ca<sup>2+</sup>, CU<sup>2+</sup>, Zn <sup>2+</sup>, Mn<sup>2+</sup>, Mg<sup>2+</sup>influence bacterial distribution in the environment. Levels of organic matter, soil moisture, soil structure and soil texture affect hydrology and also influence mineralization of soil through influence on soil microorganism activities [73]. Clay content in soil plays an important role in protection of organic matter from microbial degradation [74,75].

Unfavorable pН values can impede decomposition of organic materials by soil microorganisms hence reduced nutrients as there is reduced nutrient cycling. Consequently, there may arise competition for limited nutrients may arise among microorganisms and between microorganisms and plants. Specific microbes are responsible for decomposition of different organic matter hence there is specialization in areas they source for energy, therefore recovery of some nutrients in soil can be increased by suppressing rather that stimulating particular groups of soil microorganisms, for example urea hydrolyzers or ammonium nitrifiers [76]. It is rather necessary to quantify levels of chemicals in soil and their association with soil organisms.

Application of fertilizers in soil during crop production affects soil microorganisms which are a cognitive indicator of soil health [77]. Long-term trials reveal that mineral fertilizer increase microorganism population compared to unfertilized soils which is in contrast to grassland ecosystems where microorganism population is reduced by increased N input [52,54,77]. Biomass of bacteria, actinomycetes and fungi increases with increased fertilization using mineral fertilizers [77]. Fertilizers affect ratio of bacteria to fungi and also increases ratio of Gram-positive to Gram-negative bacteria [77]. Studies on effect of fertilizers on changes of microorganism populations and communities remain inconsistent as some studies report increasing soil microorganisms with increased fertilizer levels while others report decreasing soil microorganism populations with increased fertilizers concentrations in growth media [78] [79-81,77], which are attributed to site specific factors including soil properties and climate [77]. Fertilization affects microorganism community composition through changes in soil properties which depends on initial characteristics of soil. Effect of fertilizer application strongly depends on environmental factors which is considerably variable among different locations [77]. It is therefore necessary to conduct research on effect of fertilizer application on microorganism population and diversity shifts in polluted environments, in order to have a clear understanding of effects of fertilizers on distribution of microorganisms in soil.

# 6. EFFECT OF SEASONALITY ON DISTRIBUTION OF SOIL MICRO-ORGANISMS

Moisture content in soil limits oxygen availability and influences solubility, mobility and bioavailability of soil pollutants [82]. Wetting and drying cycles can alter bioavailability of organic contaminants in soil [82]. Heavy precipitation episodes and floods lead to increased water erosion and consequently disperse soil pollutants away from polluted sites [82]. Fluctuations in quantities and frequencies of precipitation affect the activities of soil microorganism and ultimately soil ecosystem integrity [82]. In wet seasons, the pH of soils is altered by frequent flooding and drainage [83]. During rainy season pH of acid soils increase while pH of alkaline soils decrease [84,83]. Rain water has high pH which favors ion exchange between OH and phosphate in soils which release P into flood water [83]. Higher concentration of  $Fe^{3+}$  in rainwater causes  $Fe^{3+}$ and  $Fe^{2+}$  form mixed  $Fe^{2+}$  - $Fe^{3+}$  hydroxide with a larger specific surface area, which increases adsorption of soluble P [85,83]. Calcium (Ca) compounds in alkaline soils play an important role in retention of P as most inorganic P is bound to Ca compounds [86,83].

Reduction in levels of rainfall can have major effect in microorganism biomass but minor effect on community compositions. Increase or decrease in quantity of rainfall can cause alteration of carbon cycle in terrestrial ecosystems and hence such ecosystems depend on low-ground microorganism community for resilience of biological processes in soil. However, there is diverse and uncertain findings surrounding seasonal effect on microbial diversity in soil ecosystems [87,88] as there is more than only rainfall as an influencing factor, complex interactions between but and microorganisms abiotic biotic and components. Rainfall reduction can cause change biomass community and of microorganism either directly or indirectly.

Reduced moisture levels in soil decreases solute mobility and constrains the supply of substrate to microorganisms, which is a direct effect that inhibits growth of microorganisms which are decomposers, and consequently cause reduced carbon and nitrogen from detritus entering the soil ecosystem [89,89]. On contrary, some microorganisms, including, Gram-positive bacteria, Actinobacteria and Arbuscular mycorrhizal fungi are tolerant to water logged areas [90]. This indicates that fluctuation in seasonality and rainfall patterns have impact on composition of soil microorganism communities [91,92].

Available moisture or water content controls collisions of soil particles and microorganisms [91,93,45]. Reduced rainfall affects soil total microorganism population [45]. Rainfall reductions affect ability of microorganisms to obtain required nutrients from soil [94,45]. However, not all areas show relationship between rain reductions and microorganism populations [45]. This can be attributed to evolutionary adaptation of soil microorganism population to harsh conditions; temperature and water stress [95,93,96,45]. Fungal populations can withstand reduced rainfall more than bacteria can do [97,93,45]. This can be attributed to possession of filamentous structure that allows fungi to reach and exploit substrates that are available in soil at very low moisture levels [45]. Further, the adaptive traits; thicker cell walls, ability of hyphae to obtain available nutrients in fungi affects responses to rainfall reduction [45]. Thick cell walls assist fungi to withstand high osmotic pressure from environment [98,45]. This can also apply to bacteria, where Gram-positive have advanced osmoregulatory bacteria mechanisms and stronger cell walls as compared to Gram-negative and consequently can manifest better in varied seasonality. For example, Acidobacteria and Actinobacteria are drought tolerant and are hardly affected by rainfall levels Total decrease in microorganism [98,45]. population can also be attributed to warming effects experienced over the dry period [99,100] [45], suggesting that soil microorganisms can be varied according to different ecosystems, for example, rainfall reductions can affect soil microorganisms in a forested area and at same time fail to affect microorganism population in a close by grassland [92,45]. Factors driving microorganism populations among different ecosystems maybe varied with difference in structures of ecosystems [45]. Therefore, reduced microorganism populations in soil can

negatively affect soil carbon dynamics by causing a decline in carbon levels [101-104,45]. Hence seasonal fluctuations may affect microorganism population shifts in polluted areas.

Influence of rainfall on soil microorganisms remains uncertain [105]. Rainfall reductions decrease soil microorganism biomass but with less effect on fungi as compared to bacteria Rainfall has minor influence [105]. on microorganism community composition but can majorly influence microorganism biomass [105]. Increased rainfall declines soil organic carbon and soil respiration which has a positive relationship with changes in microorganism biomass [105]. Microorganism communities respond differently to different levels of precipitations in specified ecosystems [91,92] [106,107,105]. There is still much uncertainty surrounding description of patterns of distribution among microorganism communities to change in rainfall patterns [87,88,105]. Since distribution and abundance of soil microorganisms is dependent on both biotic and abiotic factors, it is difficult to predict microorganism related environment dvnamics in [108,109,105]. Decrease in solute mobility constrains substrates that are necessary for growth of microorganisms and therefore inhibits their growth [93,89,105]. Seasonal influence in availability of oxygen in soil has a cascaded effect to microorganism community population and mainly decomposers [77]. Under anaerobic conditions decomposition of cell wall components is reduced and degradation of leachable and easily hydrolysable compounds is similar under both aerobic and anaerobic conditions [77].

# 7. EFFECT OF URBAN SETTLEMENTS ON DISTRIBUTION OF SOIL MICROORGANISMS

It is vital to assess effects of urbanized settlements on soil ecosystem integrity in order to form a baseline for reference on positive and negative implications that may occur in soils out of the settlement. In order to realize long-term agricultural productivity, soil fertility must be monitored and maintained at its optimal level. Knowledge on soil quality is important for effective management of farms as it forms baseline data for strategies to maintain and improve soil fertility [42].

Metabolic activities of soil microorganisms are mainly influenced by temperature and

physicochemical parameters [110]. In a study by Eze [111], microorganisms were isolated from soils contaminated with cassava effluent. Soils were described to have Escherichia coli, Pseudomonus spp, Bacillus spp, Proteus spp and Penicillin spp. The study revealed presence of microorganisms in soils contaminated with cyanide at a concentration of 3.0 mg/kg. These was an indication that waste material should be well treated before being discharged to the environment. Since poor waste disposal methods leads to contamination of soils [111]. Soil pollution by heavy metal influences population and diversity of soil microorganisms. Microbial functional diversity decreases with increase in level of contamination with cadmium (Cd) [71]. Presence of low levels of microorganisms is correlated to increase cadmium intoxication in soils [71]. Exposure of microorganisms to concentration of pollutants in soil is thus causative in development of adaptive characteristics among different species found in contaminated soils.

Soils in market places (Uyo, Umuabia, Sokoto and Oka) in Nigeria are polluted and contaminated with heavy metals which influenced distribution of soil microorganisms [112-115]. Growth of microalgae (*Microcystis flos-aquae*) in crude oil contaminated media revealed an exponential growth and reduction of crude oil in media. This is an indication of potential of microalgae in degradation of oil in polluted environments [116] and adaptation by shift in microbial populations, species richness and diversity, hence the role played by microorganisms is diverse.

In Calabar Cross River State of Nigeria, used oil spills has been found to influence distribution of microorganisms in soil [117]. Some of the heterotrophic bacteria isolated from these soils included; Pseudomonus spp, Bacillus spp, Klebsiella spp, Proteus spp, Enterococcus faecalis and Flavobacterium spp [117]. The total hydrocarbon oxidizing bacteria isolated include; Bacillus spp, Pseudomonas spp and Micrococcus spp [117]. The highly prevalent genera were Pseudomonas spp and Bacillus spp an indication that oil degrading microbes are more abundant in areas with oil contamination [117].

## 8. CONCLUSION

Industrialization, urbanization and agricultural activities remain doubtfully associated with

contamination of environment. Sporadic determination of soil physiochemistry and biological parameters is thus of essence in resilience of soil ecosystem integrity. Microorganisms are integral part of ecosystem where they continue to adopt to new niches in the environment hence need for continuous research in order to isolate groups of microorganisms that are resistant to harsh conditions encountered in polluted soils in order to enhance bioremediation of contaminated soils. It is necessary to ascertain the compounded impact of anthropogenic activity on soil ecosystem integrity including both biological and physicochemical parameters as opposed to isolated cases of either biological or physicochemical parameters, in order to develop compounded indication of impact magnitude of anthropogenic activities on soil ecosystem integrity. There is need to advocate for deeper understanding on influence of soil pollution and seasonal variations in soils on microorganism changes and levels of community soil contaminants. Variation in microorganism diversity remains uncertain in unique niches, and is unclear if pollutant influence or does not influence distribution of microorganisms in soil. Influence of fertilizer on bacteria to fungi ratio as well as microorganism population shifts are unique to different habitats and it is therefore important to check how the ratio of bacteria and fungi are affected in different human settlements; urbanized, industrialized and agricultural areas. Individual parameters of soil may seem to carry minimal effect when analyzed in isolation. However, compounded impact may be more descriptive of the soil ecosystem integrity, where the positive and negative impact are assessed simultaneously. This will foster for food security and hence a healthy population as enshrined in the Sustainable Development Goals (SDGs).

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# **COMPETING INTERESTS**

Authors have declared that no competing interests exist.

#### REFERENCES

- Okonkwo CNP, Kumar L, Taylor S. The Niger Delta Wetland ecosystem: What threatens it and why should we protect it? African Journal of Environmental Science and Technology. 2015;9(5):451-463. DOI: 10.5897/AJEST2014.1841
- 2. Adedeji OB, Adetunji VE. Aquatic pollution in Nigeria: The way forward. Advances in Environmental Biology. 2011;5(8):2024-2032.

ISSN 1995-0756

- Ekanem, Jemimah, Nwachukwu Ike. Sustainable agricultural production in degraded oil producing and conflict prone communities of Niger Delta, Nigeria; Journal of Agriculture and Sustainability. 2015;8(1):2015:14-28. ISSN 2201-4357.
- 4 Olalekan, Adekola, Gordon Mitchell. The Delta Niger wetlands: Threats to ecosystem services, their importance to dependent communities and possible management measures. International Journal of Biodiversity Science, Ecosystem Services & Management. 2011;7:1:50-68. DOI: 10.1080/21513732.2011.603138 To link to this article: Available:https://doi.org/10.1080/21513732 .2011.603138

 Desforges JW, Sonne C, Levin M, Siebert U, Guise SD, Dietz R. Immunotoxic effects of environmental pollutants in marine mammals. Environ. Int. 2016; 86:126–139.

 Costa AS, Romao LP, Araujo BR, Lucas SC, Maciel ST, Wisniewski Jr, A, Alexandre MR. Environmental strategies to remove volatile aromatic fractions (BTEX) from petroleum industry wastewater using biomass. Bioresour. Technol. 2012;105: 31–39.

 Meckenstock RU, Boll M, Mouttaki H, Koelschbach JS, Tarouco PC, Weyrauch P, Dong X, Himmelberg AM. Anaerobic degradation of benzene and polycyclic aromatic hydrocarbons. J. Mol. Microbiol. Biotechnol. 2016;26:92–118.

 Wilkes H, Buckel W, Golding BT, Rabus R. Metabolism of hydrocarbons in n-Alkane utilizing anaerobic bacteria. J. Mol. Microbiol. Biotechnol. 2016;26:138–151.

9. Fasinu PS, Orisakwe OE. Heavy metal pollution in sub-saharan Africa and possible implications in cancer epidemiology. Asian Pacific Journal of

Cancer Prevention. 2013;14(6):3393–3402.

Available:https://doi.org/10.7314/APJCP.2 013.14.6.3393

- 10. World Health Organization [WHO]. Training Package for the Health Sector: Adverse Health Effects of Heavy Metals in Children. World Health Organization; 2011.
- 11. Kabata-Pendias A. Trace elements in soils and plants. 3rd Edition, CRC Press, Boca Raton, FL, USA; 2007.
- 12. Defarge N, Spiroux de Vendômois J, Séralini GE. Toxicity of formulants and heavy metals in glyphosate-based herbicides and other pesticides. Toxicology Reports. 2018; 5(October 2017):156-163. Available:https://doi.org/10.1016/j.toxrep.2 017.12.025
- Adamu A, Iyaka YA, Mathew JT, Inobeme A, Egharevba HO. Assessment of some heavy metal contamination and analysis of physicochemical parameters of surface soil within the vicinity of Minna Railway Station, Niger State, Nigeria Journal of Applied Life Sciences International. 2017;10(1):1-9. 2017;Article no. JALSI.28671 ISSN: 2394-1103
- Owolabi JB, Hekeu MM. Heavy metal resistance and antibiotic susceptibility pattern of bacteria isolated from selected polluted soils in Lagos and Ota, Nigeria. International Journal of Basic & Applied Sciences IJBAS-IJENS. 2014; 14(06).
- Sander AB, Weelink Miriam HA. van Eekert Alfons, Stams JM. Degradation of BTEX by anaerobic bacteria: Physiology and application. Rev Environ Sci Biotechnol. 2010;9:359–385. DOI: 10.1007/s11157-010-9219-2
- Osterreicher-Cunha P, Davée Guimarães JR, E. do Amaral Vargas Jr. MI, Pais da Silva. Study of biodegradation processes of BTEX-ethanol mixture in tropical soil. Water Air Soil Pollut. 2007;181:303–317. DOI: 10.1007/s11270-006-9303-y
- 17. Sikorski J. The prokaryotic biology of soil. Soil Organisms. 2015;87(1):1–28.
- Francino MP. The ecology of bacterial genes and the survival of the new. International Journal of Evolutionary Biology. 2012;1–14. Available:<u>https://doi.org/10.1155/2012/394</u> 026
- 19. Hoorman JJ. The role of soil bacteria.

The Ohio State University Extension. 2011;1–4.

Available:https://doi.org/10.2323/jgam.7.12 8

- 20. Altidor, Martine. Nutritional types of bacteria. Sciencing; 2018. Available:<u>https://sciencing.com/nutritional-types-bacteria-2515.html</u>
- Gleeson C, Gray N. The coliform index and waterborne disease. Problems of microbial drinking water assessment. E & FN Spon, London; 1997.
- Saini A, Aggarwal NK, Sharma A, Yadav A. Actinomycetes: A source of lignocellulolytic enzymes. Enzyme Research; 2015. Available:https://doi.org/10.1155/2015/279 381
- Heritage J, Evans EGV, Killington RA. The Microbiology of soil and nutrient cycling. Microbiology in Action. 1999;2–13.
- 24. Van Der Heijden MGA, Bardgett RD, Van Straalen NM. The unseen majority: Soil microbes as drivers of plant diversity and productivity in terrestrial ecosystems. Ecology Letters. 2008;11(3):296–310. Available:https://doi.org/10.1111/j.1461-0248.2007.01139.x
- Bünemann EK, Mäder P, Wohlfahrt J, Brussaard L, Bongiorno G, Goede R De, Fibl P. Concepts and indicators of soil quality – a review Report type: Report Issue date: October 2016, (October); 2016.
- 26. New Hampshire Department of Environmental Services. Fecal Coliform as an Indicator Organism. Environmental Fact Sheet. 2003;3. Available:https://www.des.nh.gov/organizat ion/commissioner/pip/factsheets/wwt/docu ments/web-18.pdf
- Snel M, Shordt K. The evidence to support hygiene, sanitation and water in schools. Waterlines. 2005;23(3):2–5. Available:https://doi.org/10.4324/97813156 93606
- Møretrø T, Langsrud S. Residential bacteria on surfaces in the food industry and their implications for food safety and quality. Comprehensive Reviews in Food Science and Food Safety. 2017;16(5): 1022–1051. Available:https://doi.org/10.1111/1551-

4337.12283

29. Zhao F, Zhou JD, Ma F, Shi RJ, Han SQ, Zhang J, Zhang Y. Simultaneous inhibition of sulfate-reducing bacteria, removal of H2S and production of rhamnolipid by recombinant Pseudomonas stutzeri Rhl: applications for microbial enhanced oil recovery. Bioresour. Technol. 2016;207: 24–30.

- Allen MJ, Edberg SC, Reasoner DJ. Heterotrophic plate count bacteria - What is their significance in drinking water? International Journal of Food Microbiology. 2004;92(3):265–274. Available:https://doi.org/10.1016/j.ijfoodmic ro.2003.08.017
- James Thomas. Types of heterotrophic bacteria. Sciencing; 2017. Available:https://sciencing.com/typesheterotrophic-bacteria-6884639.html
- 32. Wosu KR, Odokuma LO. Post impact assessment of urbanization on microbial abundance and diversity of soils in Port Harcourt Area. Rep Opinion. 2017;9(4):48-56.
  - ISSN 1553-9873 (print);

ISSN 2375-7205 Available:http://www.sciencepub.net/report .5

DOI: 10.7537/marsroi090417.05

33. Biologydictionary.net Editors. Organ; 2014. Available:<u>https://biologydictionary.net/organ/</u>

Retrieved November 4, 2014

- Lord NS, Kaplan CW, Shank P, Kitts CL, Elrod SL. Assessment of fungal diversiy using terminal restriction fragment (TRF) pattern analysis: Comparision of 18s and ITS ribosomal regions. FEMS Microbiology Ecology. 2002;42:327–337.
- Tugel AJ, Lewandowski AM, eds. (February 2001 -- last update). Soil Biology Primer. Available:www.statlab.iastate.edu/survey/S Ql/soil biology primer.htm [4/06/2018].
- Webb JS, Nixon M, Eastwood IM, Greenhalgh M, Robson GD, Handley PS. Fungal colonization and biodeterioration of plasticized polyvinyl chloride. Appl. Environ. Microbiol. 2000;66:3194–3200.
- Boonchan S, Britz ML, Stanley GA. Degradation and mineralization of highmolecular weight polycyclic aromatic hydrocarbons by defined fungal bacterial cocultures. Appl Environ Microbiol. 2000; 66:107–1019.
- Annibale D, Rosetto A, Leonardi F., Federici V, Petruccioli VFM. Role of autochthonous filamentous fungi in bioremediation of a soil Historically contaminated with aromatic hydrocarbons. Appl. Environ. Microbiol. 2006;72:28–36.

- Imhoff JF. New dimensions in microbial ecology — functional genes in studies to unravel the biodiversity and role of functional microbial groups in the environment. Microorganisms. 2016; 4(May):1–41. Available:https://doi.org/10.3390/microorga nisms4020019
- 40. Bach EM, Williams RJ, Hargreaves SK, Yang F, Hofmockel KS. Greatest soil microbial diversity found in micro-habitats. Soil Biology and Biochemistry. 2018; 118(December 2017):217–226. Available:https://doi.org/10.1016/j.soilbio.2 017.12.018
- Zhang C, Nie S, Liang J, Zeng G, Wu H, Hua S, Xiang H. Effects of heavy metals and soil physicochemical properties on wetland soil microbial biomass and bacterial community structure. Science of the Total Environment. 2016;557–558: 785–790.

Available:https://doi.org/10.1016/j.scitotenv .2016.01.170

- 42. Zhou W, Lv T, Chen Y, Westby AP, Ren W. Soil Physicochemical and Biological Properties of Paddy-Upland Rotation : A Review. Hindawi. 2014;1–8. Available:https://doi.org/doi.org/10.1155/20 14/856352
- Baldrian P, Editor G. Ecology and metagenomics of soil microorganisms. FEMS Microbiol Ecol. 2011;78(August), 1– 2.

Available:https://doi.org/10.1111/j.1574-6941.2011.01184.x

44. Zhou JZ, Xue K, Xie JP, Deng Y, Wu LY, Cheng XH, et al. Microbial mediation of carbon-cycle feedbacks to climate warming. Nat. Clim. 2012;Change 2:106– 110.

DOI: 10.1038/Nclimate1331

- Ren C, Chen J, Lu X, Doughty R, Zhao F, Zhong Z, Ren G. Responses of soil total microbial biomass and community compositions to rainfall reductions. Soil Biology and Biochemistry. 2018a; 116(September 2017):4–10. Available:https://doi.org/10.1016/j.soilbio.2 017.09.028
- Delgado-Baquerizo M, Maestre FT, Reich PB, Trivedi P, Osanai Y, Liu YR, et al. Carbon content and climate variability drive global soil bacterial diversity patterns. Ecol. Monogr; 2016b. DOI: 10.1002/ecm.1216 [Epub ahead of print]

47. Liu, Dong, Huang Yimei, An, Shaoshan, Sun, Hanyin, Bhople, Parag, Chen, Zhiwei. Soil physicochemical and microbial characteristics of contrasting land-use types along soil depth gradients. CATENA; 2017.

DOI: 10.1016/j.catena.2017.10.028

 Liu Y, Delgado-baquerizo M, Wang J, Hu H, Yang Z, He J. New insights into the role of microbial community composition in driving soil respiration rates. Soil Biology and Biochemistry. 2018;118(December 2017):35–41. Available:https://doi.org/10.1016/j.soilbio.2

017.12.003

- Wang C, Liu D, Bai E. Decreasing soil microbial diversity is associated with decreasing microbial biomass under nitrogen addition. Soil Biology and Biochemistry. 2018;120(72):126–133. Available:https://doi.org/10.1016/j.soilbio.2 018.02.003
- Wang C, Lu X, Mori T, Mao Q, Zhou K, Zhou G, Mo J. Responses of soil microbial community to continuous experimental nitrogen additions for 13 years in a nitrogen-rich tropical forest. Soil Biology and Biochemistry. 2018;121(March):103– 112.

Available:https://doi.org/10.1016/j.soilbio.2 018.03.009

- Treseder KK. Nitrogen additions and microbial biomass: A meta-analysis of ecosystem studies. Ecology Letters. 2008;11: 1111–1120.
- 52. Liu L, Greaver TL. A global perspective on belowground carbon dynamics under nitrogen enrichment. Ecology Letters. 2010;13:819–828.
- Zhou Z, Wang C, Zheng M, Jiang L, Luo Y. Soil Biology & Biochemistry Patterns and mechanisms of responses by soil microbial communities to nitrogen addition. Soil Biology and Biochemistry. 2017;115:433– 441.

Available:https://doi.org/10.1016/j.soilbio.2 017.09.015

- Geisseler D, Lazicki PA, Scow KM. Mineral nitrogen input decreases mi- crobial biomass in soils under grasslands but not annual crops. Applied Soil Ecology. 2016; 106:1e10.
- 55. Tian D, Niu S. A global analysis of soil acidification caused by nitrogen addition. Environmental Research Letters. 2015;10: 024019.

- Bowman WD, Cleveland CC, Halada L, Hresko J, Baron JS. Negative impact of nitrogen deposition on soil buffering capacity. Nature Geoscience. 2008;1: 767–770.
- 57. Waring BG, Averill C, Hawkes CV. Differences in fungal and bacterial physiology alter soil carbon and nitrogen cycling: insights from meta-analysis and the- oretical models. Ecology Letters. 2013;16:887–894.
- 58. Huang PM, Bollag JM, SN. Interactions between soil particles and microorganisms: Impact on the terrestrial ecosystem. New Books and Publications. 2002;24(4).
- 59. Liua D, Huanga Y, Ana S, Sunc H, Bhopled P, CZ. Catena soil physicochemical and microbial characteristics of contrasting land-use types along soil depth gradients. Catena. 2017;10(February):0–1. Available:https://doi.org/10.1016/j.catena.2 017.10.028
- 60. Sinsabaugh RS. Enzymic analysis of microbial pattern and process. Biology and Fertility of Soils. 1994;17(1):69–74. Available:https://doi.org/10.1007/BF00418 675
- Rousk J, Baath E, Brookes PC, Lauber CL, Lozupone C, Caporaso JG, Knight R, Fierer N. Soil bacterial and fungal communities across a pH gradient in an arable soil. The ISME Journal. 2010;4: 1340–1351.
- Peng Y, Chen G, Chen G, Li S, Peng T, Qiu X, Luo J, Yang S, Hu T, Hu H, Xu Z, Liu L, Tang Y, Tu L. Soil biochemical responses to nitrogen addition in a secondary evergreen broad-leaved forest ecosystem. Scientific Reports. 2017;7: 2783.61.
- Li Y, Sun J, Tian D, Wang J, Ha D, Qu Y, Jing G, Niu S. Soil acid cations induced reduction in soil respiration under nitrogen enrichment and soil acidifica- tion. The Science of the Total Environment. 2018; 615:1535–1546.
- 64. Kaiser C, Franklin O, Dieckmann U, Richter A. Microbial community dynamics alleviate stoichiometric constraints during litter decay. Ecology Letters. 2014;17: 680–690.
- 65. Ren C, Chen J, Lu X, Doughty R, Zhao F, Zhong Z, Ren G. Responses of soil total microbial biomass and community compositions to rainfall reductions. Soil

Biology and Biochemistry. 2018; 116(September 2017):4–10. Available:https://doi.org/10.1016/j.soilbio.2 017.09.028

- Cassou E, Jaffee SM, Ru J. The challenge of agricultural pollution: Evidence from China, Vietnam, and the Philippines; 2017. Available:https://doi.org/10.1596/978-1-4648-1201-9
- 67. Crouse DA. Soils and plant nutrients, Chpt 1. In: K.A. Moore, and. L.K. Bradley (eds). North Carolina Extension Gardener Handbook. NC State Extension, Raleigh, NC; 2017. Available:https://content.ces.ncsu.edu/exte

nsion-gardener-handbook/1-soils-andplant-nutrients

- Bhogal A, Williams J, Nicholson F, Chadwick D, Chambers K, Chambers BJ. Mineralisation of organic nitrogen from farm manure applications. Soil Use Manage. 2015;32(Suppl 1):32–43.
- Osborne TM, Wheeler TR. Evidence for a climate signal in trends of global crop yield variability over the past 50 years. Environmental Research Letters. 2013;8(2). Available:https://doi.org/10.1088/1748-9326/8/2/024001
- García-Orenes F, Morugán-Coronado A, Zornoza R, Cerdà A, Scow K. Correction: Changes in soil microbial community structure influenced by agricultural management practices in a Mediterranean agro-ecosystem. PLoS ONE. 2016;11(3): 1–9.

Available:https://doi.org/10.1371/journal.po ne.0152958

- Xie Y, Fan J, Zhu W, Amombo E, Lou Y, Chen L, Fu J. Effect of heavy metals pollution on soil microbial diversity and bermudagrass genetic variation. Front. Plant Sci. 2016;7:755. DOI: 10.3389/fpls.2016.00755
- Hai-Hang Li ZY. Zhang, Pan LP. Isolation, identification and characterization of soil microbes which degrade phenolic allelochemicals Journal compilation <sup>a</sup> 2009 The Society for Applied Microbiology, Journal of Applied Microbiology. 2010;108(2010):1839–1849
- 73. Li Y, Liu Y, Niu YWL, Xia X, Tian Y. Interactive effects of soil temperature and moisture on soil N mineralization in a Stipa krylovii grassland in Inner Mongolia, China. Journal of Arid Land. 2014;6:571–580.
- 74. Mc Donald NT, Watson C, Lalor S,

Laughlin SR, Wall DP. Evaluation of soil tests for predicting nitrogen mineralization in temperate grassland soils. Soil Science Society of America Journal. 2014;78: 1051–1064.

- 75. Lehtinen T, Schlatter N, Baumgarten A, Bechini L, Kr€uger J, Grignani C, Zavattaro L, Costamagna C, Spiegel H. Effect of crop residue incorporation on soil organic carbon and greenhouse gas emissions in European agricultural soils. Soil Use Manage. 2014;30:524–538.
- Ruser R, Schulz R. The effect of nitrification inibitors on the nitrous oxide (N2O) release from agricultural soils—a review. Journal of Plant Nutrition and Soil Science. 2015;178:171–188.
- Geisseler D, Linquist BA, Lazicki PA. Soil biology & biochemistry effect of fertilization on soil microorganisms in paddy rice systems e A. Soil Biology and Biochemistry. 2017;115:452–460. Available:https://doi.org/10.1016/j.soilbio.2 017.09.018
- 78. Bhattacharyya P, Chakrabarti K, Chakraborty A. Microbial biomass and enzyme activities in submerged rice soil amended with municipal solid waste compost and decomposed cow manure. Chemosphere. 2005;60:310e318.
- 79. Zhang J, Qin J, Yao W, Bi L, Lai T, Yu X. Effect of long-term application of manure and mineral fertilizers on nitrogen mineralization and microbial biomass in paddy soil during rice growth stages. Plant. Soil & Environment. 2009;55:101e109.
- Datta A, Santra SC, Adhya TK. Effect of inorganic fertilizers (N, P, K) on methane emission from tropical rice field of India. Atmospheric Environment. 2013;66: 123e130.
- 81. Das S, Adhya TK. Effect of combine application of organic manure and inorganic fertilizer on methane and nitrous oxide emissions from a tropical flooded soil planted to rice. Geoderma. 2014;213: 185e192
- Alkorta I, Epelde L, Garbisu C. Environmental parameters altered by climate change affect the activity of soil microorganisms involved in bioremediation. FEMS Microbiology Letters. 2017;364(19):1–9. Available:https://doi.org/10.1093/femsle/fnx 200
- 83. Tian Juan, Dong Guiming, Karthikeyan Raghupathy, Li Lin, HDR. Drained, and

Reflooded Soils. Water. 2017;9:531. Available:https://doi.org/10.3390/w907053 1

- Ponnamperuma FN. The chemistry of submerged soils. Advances in Agronomy. 1972;24:29–96.
- Szilas CP, Borggard OK, Hansedn HCB. Potential iron and phosphate mobilization during flooding of soil material. Water Air Soil Pollut. 1998;106:97–109.
- Hogan DM, Jordan TE, Walbridge MR. Phosphorus retention and soil organic carbon in restored and natural freshwater wetlands. Wetlands. 2004;24:573–585.
- Shen RC, Xu M, Li RQ, Zhao FX, Sheng QK. Spatial variability of soil microbial biomass and its relationships with edaphic, vegetational and climatic fac- tors in the Three-River Headwaters region on Qinghai-Tibetan Plateau. Applied Soil Ecology. 2015;95:191–203.
- Xi NX, Bloor JMG. Interactive effects of precipitation and nitrogen spatial pattern on carbon use and functional diversity in soil microbial communities. Applied Soil Ecology. 2016;100:207–210.
- Manzoni S, Schaeffer S, Katul G, Porporato A, Schimel J. A theoretical analysis of microbial eco-physiological and diffusion limitations to carbon cycling in drying soils. Soil Biology and Biochemistry. 2014;73:69–83.
- Zeglin LH, Bottomley PJ, Jumpponen A, Rice CW, Arango M, Lindsley A, McGowan A, Mfombep P, Myrold DD. Altered precipitation regime affects the function and composition of soil microbial communities on multiple time scales. Ecology. 2013;94:2334–2345.
- 91. Beier C, Beierkuhnlein C, Wohlgemuth T, Penuelas J, Emmett B, Korner C, de Boeck HJ, Christensen JH, Leuzinger S, Janssens IA, Hansen K. Precipitation manipulation experiments - challenges and recommendations for the future. Ecology Letters. 2012;15:899–911.
- 92. Nielsen UN, Ball BA. Impacts of altered precipitation regimes on soil commu- nities and biogeochemistry in arid and semi-arid ecosystems. Global Change Biology. 2015;21:1407–1421.
- Manzoni S, Schimel JP, Porporato A. Responses of soil microbial communities to water stress: Results from a metaanalysis. Ecology. 2012;93:930–938.
- 94. Schimel J, Balser TC, Wallenstein M. Microbial stress-response physiology and

its implications for ecosystem function. Ecology. 2007;88:1386–1394.

- Bradford MA, Davies CA, Frey SD, Maddox TR, Melillo JM, Mohan JE, Reynolds JF, Treseder KK, Wallenstein MD. Thermal adaptation of soil microbial respiration to elevated temperature. Ecology Letters. 2008;11:1316–1327.
- 96. Romero-Olivares AL, Allison SD, Treseder KK. Soil microbes and their re- sponse to experimental warming over time: A metaanalysis of field studies. Soil Biology and Biochemistry. 2017;107:32–40.
- 97. Von Rein I, Gessler A, Premke K, Keitel C, Ulrich A, Kayler ZE. Forest understory plant and soil microbial response to an experimentally induced drought and heatpulse event: the importance of maintaining the continuum. Global Change Biology. 2016;22:2861–2874.
- Evans SE, Wallenstein MD. Climate change alters ecological strategies of soil bacteria. Ecology Letters. 2014;17:155– 164.
- 99. Chen J, Luo YQ, Xia JY, Jiang LF, Zhou XH, Lu M, Liang JY, Shi Z, Shelton S, Cao JJ. Stronger warming effects on microbial abundances in colder re- gions. Scientific Reports. 2015;5:18032.
- Hendershot JN, Read QD, Henning JA, Sanders NJ, Classen AT. Consistently inconsistent drivers of patterns of microbial diversity and abundance at macroecological scales. Ecology. 2017; 98:1757–1763.
- 101. Canarini A, Kiaer LP, Dijkstra FA. Soil carbon loss regulated by drought in- tensity and available substrate: A meta-analysis. Soil Biology and Biochemistry. 2017;112: 90–99.
- 102. Liu LL, Wang X, Lajeunesse MJ, Miao GF, Piao SL, Wan SQ, Wu YX, Wang ZH, Yang S, Li P, Deng MF. A cross-biome synthesis of soil respiration and its determinants under simulated precipitation changes. Global Change Biology. 2016; 22:1394–1405.
- 103. Zhou LY, Zhou XH, Shao JJ, Nie YY, He YH, Jiang LL, Wu ZT, Bai SH. Interactive effects of global change factors on soil respiration and its com- ponents: A metaanalysis. Global Change Biology. 2016a; 22:3157–3169.104
- 104. Zhou X, Zhou L, Nie Y, Fu Y, Du Z, Shao J, Zheng Z, Wang X. Similar responses of soil carbon storage to drought and irrigation in terrestrial ecosystems but with

contrasting mechanisms: A meta-analysis. Agriculture, Ecosystems & Environment. 2016b;228:70–81.

- 105. Ren C, Chen J, Lu X, Doughty R, Zhao F, Zhong Z, Ren G. Responses of soil total microbial biomass and community compositions to rainfall reductions. Soil Biology and Biochemistry. 2018b; 116(September 2017):4–10. Available:https://doi.org/10.1016/j.soilbio.2 017.09.028
- 106. Rousk J, Smith AR, Jones DL. Investigating the long-term legacy of drought and warming on the soil microbial community across five European shrubland ecosystems. Global Change Biology. 2013;19:3872–3884.
- 107. Serna-Chavez HM, Fierer N, Van Bodegom PM. Global drivers and patterns of microbial abundance in soil. Global Ecology and Biogeography. 2013;22: 1162–1172.
- Davidson EA, Janssens IA. Temperature sensitivity of soil carbon decomposition and feedbacks to climate change. Nature. 2006;440:165–173.
- 109. Singh BK, Bardgett RD, Smith P, Reay DS. Microorganisms and climate change: Terrestrial feedbacks and mitigation options. Nature Reviews Microbiology. 2010;8:779–790.
- 110. QL Fu, N, Weng M, Fujii DM, Zhou. Temporal variability in Cu speciation, phytotoxicity, and soil microbial activity of Cu-polluted soils as affected by elevated temperature. Journal Chemosphere. 2018; 194:285-296.
- 111. Eze VC, Onyilide DM. Microbiological and physicochemical characteristics of soil receiving cassava effluent in Elele, Rivers State, Nigeria. Journal of Applied &

Environmental Microbiology. 2015;3(1) (2015):20-24.

DOI: 10.12691/jaem-3-1-4

- 112. Akpoveta OV, Osakwe SA, Okoh BE, Otuya BO. Physicochemical characteristics and levels of some heavy metals in soils around metal scrap dumps in some parts of Delta State, Nigeria Journal of Applied Science and Environmental Management. 2010;14(4):57–60.
- 113. Ogbemudia FO, Mbong EO. Soil reaction (pH) and heavy metal index of dumpsites within Uyo municipality Merit Research Journal of Environmental Science and Toxicology. 2013;1(4):82–85.
- 114. Eze VC, Omeh YN, Ugweje CD. Microbiological and physicochemical assessment of soil contaminated with lairage effluent in Umuahia, Abia State, Nigeria. Journal of Pharmacy and Biological Sciences. 2013;8(2):50–56.
- 115. Imarhiagbe EE. Osarenotor О. Obayaqbona ON. Eghomwanre AF. Nzeadibe BNJ. Evaluation of physicochemical, microbiological and polycyclic aromatic hydrocarbon content of top soils from Oka Market Waste Collection Site, Benin City, Nigeria Appl. Sci. Environ. Manage. 2017;21(1): 112-117.
- Ifeanyi VO, Ogbulie JN. Biodegradation of crude oil by microalgae microcystis flosaquae Nigerian Journal of Microbiology. 2016;30(2):3459-3463.
- 117. Unimke AA, Mmuoegbulam AO, Bassey IU, Obot SE. Assessment of the microbial diversity of spent-oil contaminated soil in Calabar, Nigeria Journal of Advances in Microbiology. 2017;4(4):1-9:2017:Article no. JAMB.34847 ISSN: 2456-7116.

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