

**Keywords** Biodegradation; Bioremediation; Polycyclic aromatic hydrocarbons; Phenanthrene; Anthracene; Flouranthene; Pyrene

## Introduction

Biodegradation is the chemical dissolving of organic and non-organic pollutants by use of micro-organisms or other biological agents [1]. In recent years, biodegradation of pollutants by microbes has received significant interest as mankind attempts to reduce orgagradiiion; Bioremr

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7KH (IIHFW RI 6RLO S+ RQ %LRUHPHGLDWLRQ RI 3RO\F\F3@`0

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in line with the molecular weight and number of rings in the structure increased population count by at-least one log (from  $10^6$  cells per Degradation was most rapid at pH 7.5 followed by pH 7, pH 8, pH 6.5) in soil sample with microbial inocula. A total number of fungal cfu pH 8, pH 6, pH 8.5, pH 5.5 and pH 5 respectively. In general, soil pH counts measured per gram of experimental soil in each individual week 7.5 exhibited greatest and fastest rates of biodegradation for all the found for each soil pH are exhibited in Figure 6. PAH's over 45 days.

Total bacterial and fungal populations over 70 days in PAH contaminated soil

During biodegradation studies, bacterial populations at day 7, 14, 21, (Figure 5) increased whilst the highest population was evident at day 35. All the time points (from 0 to 70 days) had the greatest bacterial population exhibited at soil pH 7.5 (Figure 5b). The bacterial population dropped from its peak at day 35, dropping each week until the end of the experiment at day 70. The bacterial populations measured in soil samples without microbial inocula were  $(\log 2 \times 10^2)$  below the detection limits at day 0 and after 7 days (Figure 5a). Bacteria were evident in soil without microbial inocula Figure 5a indicating recolonisation of the soil from 14 to 70 day of incubation. The total fungal population was evaluated. The fungal cfu counts per gram of soil exhibited higher difference in soil samples with and without microbial inocula. *Aspergillus* and *Penicillium* strains exhibited

The greatest fungal populations were found in day 29 followed by time point of days 21, 14 and 42 (Figure 6). The highest population was found in soil pH 5 after 21 and at 29 day of incubation. In Figure 6b the greater fungal populations were found at low soil pH acidic conditions. However, alkaline soil pH 8 and 8.5 had higher populations compared to neutral soil pH. Soil pH 7.0 and 7.5 had the lowest fungal populations. Interestingly, *Penicillium* species predominated at acidic soil pH and with lower *Aspergillus* populations whereas at alkaline conditions of (pH 8 and 8.5) *Aspergillus* were predominant and *Penicillium* was not detected. The bacterial populations in microbial inoculated soil were greatest at soil pH 7.5 which also resulted in the greatest degradation rates suggesting that bacteria were involved in the biodegradation of polycyclic aromatic hydrocarbons. Therefore, it may be that microbial community particularly bacteria was more prevalent and active in degradation at pH 7.5. Conversely, fungal populations were greatest at acidic soil pH and with some evident at alkaline soil suggests degradation at lower pH might be initiated by fungal populations.

soilat varying pH:

soil pH. Soil inoculated with microorganisms resulted in differences in ammonification activities were measured at buffer pH 5.5 (Figure 14), in phosphatase activities over time and pH at acidic soil pH when pH 7 (Figure 15) and pH 8.5 (Figure 16) exhibited higher activities compared to neutral and alkaline soil pH. Phosphatase activity measured at buffer pH 8.5 continued to increase over time especially at alkaline and neutral pH reaching a peak of 1-1.2 mg pNP<sub>405</sub> d.w. h<sup>-1</sup> at pH 8.5 at 70 days whilst at the same time point at soil pH of 5.5 the activity was between 0.6-0.8 mg pNP<sub>405</sub> soil d.w. h<sup>-1</sup> (Figure 10b) compared to (Figures 8b and 9b).

#### -Glucosidase soil activity

-glucosidase activities were measured at buffer pH 5.5 (Figure 11), pH 7 (Figure 12) and pH 8.5 (Figure 13) and these showed significantly greater activity with microbial inocula with greater activity measured in buffer pH 8.5 (approximately 0.06 mg pNP<sub>405</sub> soil d.w. h<sup>-1</sup>) at alkaline soil pH and the lower activity rates obtained in buffer pH 7 (approximately 0.04 mg pNP<sub>405</sub> soil d.w. h<sup>-1</sup>) is potentially associated with bacteria respectively.

#### L-Arginine ammonification soil activity

L-Arginine ammonification activity was measured in PAH contaminated J. Arthur Bower's topsoil at varying soil pH. L-arginine





Figure 8: 3KRVS KDWDVH DFWLYLW\ RI - \$UWKXU %RZHU¶V WRSVRLO DWYDLOLQJ RYHU DFWLYLW\ LQ FRQWURO VDPSOHV ZLWKRXW PLFURELDO LQRFXOD E 3KRVS KDWDVH DFWL strains.











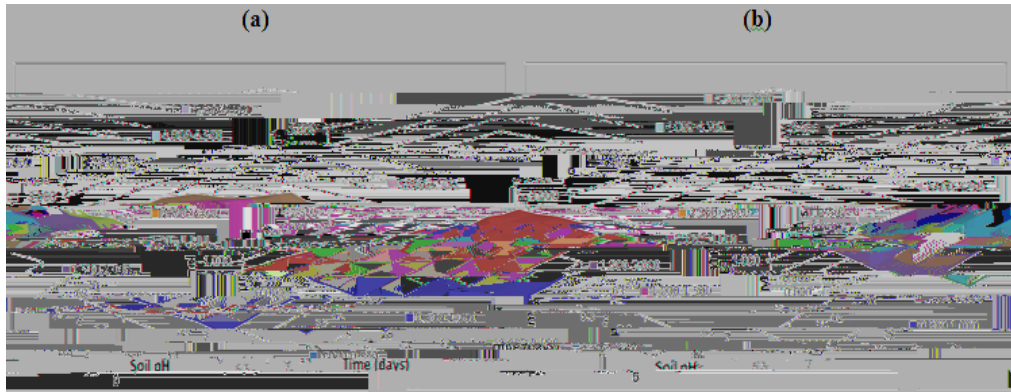


Figure 23: 0DQJDQHVH SHUR[LGDVH DFWLYLW\ RI - \$UWKXU %RZHU¶V WRSVRLO DW YDU\LQJ VRLO S SHUR[LGDVH DFWLYLW\ LQ FRQWURO VDP SOHV ZLWKRXW PLFURELDO LQRFXOD E 0DQJDQHVH microbial strains.

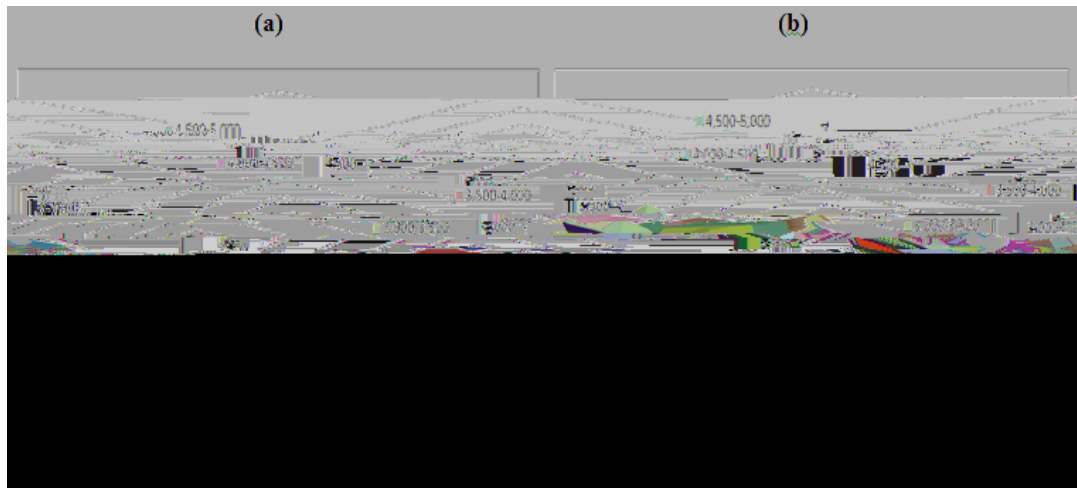


Figure 24: 0DQJDQHVH SHUR[LGDVH DFWLYLW\ LQ - \$UWKXU %RZHU¶V WRSVRLO DW YDU\LQJ VRLO S SHUR[LGDVH DFWLYLW\ LQ FRQWURO VDP SOHV ZLWKRXW PLFURELDO LQRFXOD E 0DQJDQHVH microbial strains.

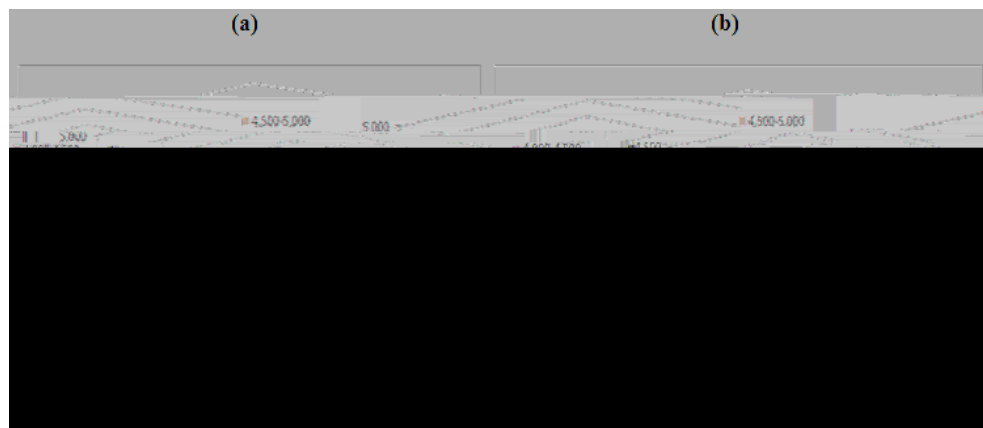


Figure 25: 0DQJDQHVH SHUR[LGDVH DFWLYLW\ LQ - \$UWKXU %RZHU¶V WRSVRLO DW YDU\LQJ VRLO S SHUR[LGDVH DFWLYLW\ LQ FRQWURO VDP SOHV ZLWKRXW PLFURELDO LQRFXOD E 0DQJDQHVH microbial strains.

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