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#### CONTENTS

#### Aquatic environment

Immunotoxic potential of aeration lagoon effluents for the treatment of domestic and Hospital wastewaters in the freshwater mussel Elliptio complanata	
Francçis Gagné, Chantale André, Marlène Fortier, Michel Fournier	· 781
Spatial distribution of archaeal and bacterial ammonia oxidizers in the littoral buffer zone of a nitrogen-rich lake	
Yu Wang, Guibing Zhu, Lei Ye, Xiaojuan Feng, Huub J. M. Op den Camp, Chengqing Yin	· 790
Accelerated biodegradation of nitrophenols in the rhizosphere of Spirodela polyrrhiza	
Risky Ayu Kristanti, Masahiro Kanbe, Tadashi Toyama, Yasuhiro Tanaka, Yueqin Tang, Xiaolei Wu, Kazuhiro Mori	· 800
Sorption of 2,4-dinitroanisole (DNAN) on lignin	
Rabih Saad, Zorana Radovic-Hrapovic, Behzad Ahvazi, Sonia Thiboutot, Guy Ampleman, Jalal Hawari	· 808
Sewage sludge disintegration by high-pressure homogenization: A sludge disintegration model	
Yuxuan Zhang, Panyue Zhang, Boqiang Ma, Hao Wu, Sheng Zhang, Xin Xu	· 814
Degradation kinetics and mechanism of aniline by heat-assisted persulfate oxidation	921
Xiaofang Xie, Yongqing Zhang, Weilin Huang, Shaobing Huang	· 821
Degradation of some typical pharmaceuticals and personal care products with copper-plating iron doped Cu <sub>2</sub> O under visible light irradiation Jing An, Qixing Zhou	۶ <u>۰</u> ۶
	. 821
Preparation of high concentration polyaluminum chloride by chemical synthesis-membrane distillation method with self-made hollow fiber membrane Changwei Zhao, Yong Yan, Deyin Hou, Zhaokun Luan, Zhiping Jia	824
Characteristics of gas-liquid pulsed discharge plasma reactor and dye decoloration efficiency	. 834
Bing Sun, Nyein Nyein Aye, Zhiying Gao, Dan Lv, Xiaomei Zhu, Masayuki Sato	. 840
Photolysis kinetics and influencing factors of bisphenol S in aqueous solutions	040
Guiping Cao, Jilai Lu, Gongying Wang	. 846
Comparative study of leaching of silver nanoparticles from fabric and effective effluent treatment	010
Aneesh Pasricha, Sant Lal Jangra, Nahar Singh, Neeraj Dilbaghi, K. N. Sood, Kanupriya Arora, Renu Pasricha	. 852
Atmospheric environment	052
Size distribution and chemical composition of secondary organic aerosol formed from Cl-initiated oxidation of toluene	
Mingqiang Huang, Weijun Zhang, Xuejun Gu, Changjin Hu, Weixiong Zhao, Zhenya Wang, Li Fang	· 860
Real-world fuel efficiency and exhaust emissions of light-duty diesel vehicles and their correlation with road conditions	
Jingnan Hu, Ye Wu, Zhishi Wang, Zhenhua Li, Yu Zhou, Haitao Wang, Xiaofeng Bao, Jiming Hao	· 865
Operating condition influences on PCDD/Fs emissions from sinter pot tests with hot flue gas recycling	
Yongmei Yu, Minghui Zheng, Xianwei Li, Xiaolei He ·····	· 875
Size distribution of chemical elements and their source apportionment in ambient coarse, fine, and ultrafine particles in Shanghai urban summer atmosphere	
Senlin Lü, Rui Zhang, Zhenkun Yao, Fei Yi, Jingjing Ren, Minghong Wu, Man Feng, Qingyue Wang	· 882
Synergistic effects of non-thermal plasma-assisted catalyst and ultrasound on toluene removal	
Yongli Sun, Libo Zhou, Luhong Zhang, Hong Sui	· 891
Absorption characteristics of new solvent based on a blend of AMP and 1,8-diamino-p-menthane for CO <sub>2</sub> absorption	
Sang-Sup Lee, Seong-Man Mun, Won-Joon Choi, Byoung-Moo Min, Sang-Won Cho, Kwang-Joong Oh	· 897
Terrestrial environment	
Toxicity and subcellular distribution of cadmium in wheat as affected by dissolved organic acids	
Dandan Li, Dongmei Zhou ·····	· 903
Changes in the sorption, desorption, distribution, and availability of copper, induced by application of sewage sludge	
on Chilean soils contaminated by mine tailings	
Tatiana Garrido, Jorge Mendoza, Francisco Arriagada ·····	· 912
Mechanism of lead immobilization by oxalic acid-activated phosphate rocks	
Guanjie Jiang, Yonghong Liu, Li Huang, Qingling Fu, Youjun Deng, Hongqing Hu	· 919
Methyl-β-cyclodextrin enhanced biodegradation of polycyclic aromatic hydrocarbons and associated microbial activity in contaminated soil	
Mingming Sun, Yongming Luo, Peter Christie, Zhongjun Jia, Zhengao Li, Ying Teng	· 926
Inhibitory effect of nitrobenzene on oxygen demand in lake sediments	
Xiaohong Zhou, Xuying Wang, Hanchang Shi	· 934
Environmental health and toxicology	
Endogenous nitric oxide mediates alleviation of cadmium toxicity induced by calcium in rice seedlings	
Long Zhang, Zhen Chen, Cheng Zhu ·····	· 940
Species-dependent effects of the phenolic herbicide ioxynil with potential thyroid hormone disrupting activity: modulation of its cellular	
uptake and activity by interaction with serum thyroid hormone-binding proteins	
Sakura Akiyoshi, Gobun Sai, Kiyoshi Yamauchi	· 949
Environmental catalysis and materials	
A screen-printed, amperometric biosensor for the determination of organophosphorus pesticides in water samples	0.57
Junfeng Dou, Fuqiang Fan, Aizhong Ding, Lirong Cheng, Raju Sekar, Hongting Wang, Shuairan Li	
A GFP-based bacterial biosensor with chromosomally integrated sensing cassette for quantitative detection of Hg(II) in environment	Cille
Hımanshu Prıyadarshi, Absar Alam, Gireesh-Babu P, Rekha Das, Pankaj Kishore, Shivendra Kumar, Aparna Chaudhari	. 963
Serial parameter: CN 11-2629/X*1989*m*188*en*P*26*2012-5	and and a
A GFP-based bacterial biosensor with chromosomally integrated sensing cassette for quantitative detection of Hg(II) in environment Himanshu Priyadarshi, Absar Alam, Gireesh-Babu P, Rekha Das, Pankaj Kishore, Shivendra Kumar, Aparna Chaudhari Serial parameter: CN 11-2629/X*1989*m*188*en*P*26*2012-5	
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## Methyl-β-cyclodextrin enhanced biodegradation of polycyclic aromatic hydrocarbons and associated microbial activity in contaminated soil

Mingming Sun<sup>1</sup>, Yongming Luo<sup>1</sup>, Peter Christie<sup>2</sup>, Zhongjun Jia<sup>1</sup>, Zhengao Li<sup>1</sup>, Ying Teng<sup>1,\*</sup>

1. Key Laboratory of Soil Environment and Pollution Remediation, Institute of Soil Science, Chinese Academy of Sciences, Nanjing 210008, China. E-mail: mmsun@issas.ac.cn

2. Agri-Environment Branch, Agri-Food and Biosciences Institute, Newforge Lane, Belfast BT9 5PX, United Kingdom

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#### Abstract

The contamination of soils by polycyclic aromatic hydrocarbons (PAHs) is a widespread environmental problem and the remediation of PAHs from these areas has been a major concern. The effectiveness of many *in situ* bioremediation systems may be constrained by low contaminant bioavailability due to limited aqueous solubility or a large magnitude of sorption. The objective of this research was to evaluate the effect of methyl- $\beta$ -cyclodextrin (MCD) on bioaugmentation by *Paracoccus* sp. strain HPD-2 of an aged PAH-contaminated soil. When 10% (*W*/*W*) MCD amendment was combined with bioaugmentation by the PAH-degrading bacterium *Paracoccus* sp. strain HPD-2, the percentage degradation of total PAHs was significantly enhanced up to 34.8%. Higher counts of culturable PAHdegrading bacteria and higher soil dehydrogenase and soil polyphenol oxidase activities were observed in 10% (*W*/*W*) MCD-assisted bioaugmentation soil. This MCD-assisted bioaugmentation strategy showed significant increases (*p* < 0.05) in the average well color development (AWCD) obtained by the BIOLOG Eco plate assay, Shannon-Weaver index (*H*) and Simpson index ( $\lambda$ ) compared with the controls, implying that this strategy at least partially restored the microbiological functioning of the PAH-contaminated soil. The results suggest that MCD-aided bioaugmentation by *Paracoccus* sp. strain HPD-2 may be a promising practical bioremediation strategy for aged PAH-contaminated soils.

**Key words**: polycyclic aromatic hydrocarbons; methyl-β-cyclodextrin; biodegradation; *Paracoccus* sp. strain HPD-2; microbial activity **DOI**: 10.1016/S1001-0742(11)60865-6

#### Introduction

Polycyclic aromatic hydrocarbons (PAHs) are ubiquitous environmental pollutants. Excessive inputs from anthropogenic activities have caused serious contamination and threaten to adversely affect human health (Wilcke, 2007; Gong et al., 2008; Sun et al., 2011). Proper remedial action to remove PAHs from contaminated soils is essential (Ding et al., 2008). Due to their high hydrophobicity and solid-water distribution ratio, PAHs in soil tend to interact with the non-aqueous phase and organic matter, and consequently become less available for further microbial degradation (Johnsen and Karlson, 2005). The utilization of PAHs in soil may result from the combined metabolism of microbial communities or from catabolism by individual strains (Juhasz et al., 1997).

Biodegradation is one of many techniques available for *in* or *ex situ* remediation of PAH-contaminated soils. However, the extent and rate of bioremediation of contaminated soils that can be achieved is often limited by PAH solubility and generally low aqueous PAH concentrations unable to sustain biodegradation (Talley et al., 2002; Yang et al., 2009, 2011). Biodegradation is therefore dependent on the ability of microorganisms to transform and degrade contaminants as well as the accessible concentrations of compounds capable of inducing or sustaining degradation (Allan et al., 2006).

While bioaugmentation with catabolically active microorganisms may be used to minimize microbial constraints, solubility-enhancing agents such as cyclodextrins have been the focus of numerous recent studies regarding their capacity to promote the release and microbial accessibility of contaminants in soils and solutions (Semple et al., 2007; Zhou and Zhu, 2007; Stroud et al., 2009). MCD is a cyclic, bucket-shaped macromolecule presenting a hydrophilic exterior rendering it water-soluble and a hydrophobic cavity enabling the formation of complexes with hydrophobic organic molecules, thereby increasing their aqueous solubility. It is also known to have low toxicity to bacterial cells and a very high solubility (solubility in water > 200 g in 100 mL at  $25^{\circ}$ C) (Petitgirard et al., 2009). Although the ability of cyclodextrins to enhance the biodegradation of pyrene in soil has been demonstrated, most studies of the impact of cyclodextrins on contaminant disappearance have been restricted to single and low molecular weight PAHs (Badr et al., 2004)

<sup>\*</sup> Corresponding author. E-mail: yteng@issas.ac.cn

Stroud et al., 2009) and the effects of MCD addition on the bioremediation of aged PAH-contaminated soils remain to be investigated.

The ultimate goals of any remediation approach are to remove the contaminants from the soil and to restore the capacity of the soil to function according to its potential (Epelde et al., 2009). Soil microorganisms are very sensitive to any shifts in ecosystem function because their activity and diversity are rapidly altered by perturbation (Margesin et al., 2000). Microbiological parameters such as soil enzyme activities and the functional diversity of soil microbial communities may serve as important indices of the impact of pollution on soil health (Shen et al., 2005; Epelde et al., 2008; Teng et al., 2010b).

In the present study several factors were investigated in order to accelerate and improve the bioremediation of an aged PAH-polluted soil, namely the degradative activity of indigenous soil microbial populations compared to that of an added PAH-degrading bacterial strain (*Paracoccus* sp. strain HPD-2), the effect of MCD addition on PAH degradation and on heterothropic microbial population dynamics, and the interaction between MCD addition and bioaugmentation by inoculation with *Paracoccus* sp. strain HPD-2. In addition, soil microcosms were set up to study MCD-assisted bioaugmentation by inoculation with *Paracoccus* sp. strain HPD-2 to evaluate the bioremediation potential of this strategy and examine associated changes in microbial activities in aged PAH-contaminated soil.

#### 1 Materials and methods

#### 1.1 Soil

The soil used for the experiment was collected from the top 15 cm of the soil profile of PAH-contaminated farmland in the Yangtze Delta, East China. The soil was contaminated approximately 30 years previously and the contaminants have therefore undergone a relatively long weathering process. Gravel and plant root residues in the sampled soil were discarded and the soil was air-dried, sieved through a 2-mm mesh, and stored at 4°C in the dark. Some physico-chemical characteristics of the experimental soil (Lu, 2000) show that the soil is a silt loam with 11.1 g/kg total organic carbon, a pH (in water) of 6.4, 1.0 g/kg total N, 14.7 g/kg total K, and 78.4 mg/kg hydrolysable N on a dry weight basis. The concentration of 16 individual PAHs was  $(9942 \pm 91) \mu g/kg dry soil, with concentrations$ of the 3-, 4- and 5(+6)-ring PAHs of (823  $\pm$  30), (5614  $\pm$ 119) and  $(3505 \pm 181) \,\mu$ g/kg, respectively. According to the Canadian Environmental Quality Guidelines released by the Canadian Council of Ministers of the Environment (CCME, 2004), this soil would not be suitable for agricultural use because of the high concentration of PAHs, and especially of 4- and 5(+6)-ring PAHs.

#### 1.2 Bacterial strain and culture conditions

*Paracoccus* sp. strain HPD-2 was isolated from a historically PAH-contaminated soil collected from Wuxi, Jiangsu Province, East China (Teng et al., 2010b). The strain was screened for its ability to degrade the highmolecular-weight (HMW) PAHs fluoranthene, pyrene and benzo[a]pyrene. After inoculation into MS medium containing B[a]P at 3.0 mg/L for 5 days, 89.7% of the B[a]P was degraded by the bacterium. When this strain was grown with pyrene and fluoranthene at 50 mg/L for 7 days, 47.2% and 84.5% of these was degraded, respectively. This strain may therefore have potential in improving HMW-PAH biodegradation. Strain HPD-2 was transferred onto a slant of nutrient agar medium. After 3 days of incubation at 28°C the slant was used to inoculate two 500-mL Erlenmeyer flasks each containing 100 mL of liquid medium from 3 g beef extract and 5 g peptone per liter of deionized water. The flasks were incubated for 48 hr at 28°C on a rotary shaker at 200 r/min and produced cell suspensions of  $2.0 \times 10^8$  CFU/L. Cells were harvested by centrifugation at 5000  $\times g$  for 20 min. The supernatant was discarded and the cells were re-suspended in fresh autoclaved distilled water (repeated twice) for bioaugmentation studies.

#### 1.3 Soil microcosms

The treatments comprised of: control (CK), with no added MCD, no bacterial inoculation, and soil moisture maintained at 60% of water holding capacity (WHC); MCD amendment M1, M2, M3, M4 at rates of 5%, 10%, 15%, and 20% (W/W), respectively, with the MCD added as solid particles mixed thoroughly into the soil; bioaugmentation (CKB), inoculation with 4% (V/W) Paracoccus sp. strain HPD-2 cell suspension; and MCD-assisted bioaugmentation (MB1, MB2, MB3, MB4), combining the same four rates of MCD amendment as above plus inoculation with 4% (V/W) Paracoccus sp. strain HPD-2 cell suspension. The experiment thus consisted of a total of 10 treatments in triplicate. Soil microcosms were set up using a series of beakers (upper diameter 11.5 cm, lower diameter 9 cm) covered with aluminum foil, each containing 1000 g of air-dried contaminated soil. After adjustment of the soil moisture, the microcosms were incubated in a controlled climate chamber for 16 weeks at  $(25 \pm 0.5)^{\circ}$ C. During incubation the soil moisture contents were adjusted by weighing the microcosms. After 1, 2, 4, 8, 12 and 16 weeks soil samples were collected from each soil microcosm using a mini stainless steel soil drill. Each soil sample was divided into two parts, one of which was placed in a small plastic bag at 4°C for subsequent analysis of microbial activity and the other freeze-dried and passed through a 60-mesh sieve prior to analysis for PAHs.

#### 1.4 Extraction and analysis of soil PAHs

PAHs in bulk soil samples were extracted using Soxhlet extraction. In brief, 5 g of freeze-dried sample with filter paper was placed in a porous cellulose thimble (25 mm  $\times$  70 mm) and placed in a Soxhlet extractor. The extractor was then fitted to a 100 mL round bottom flask containing 60 mL dichloromethane and the extraction was performed for 24 hr. All the extracts in the round bottom flasks were dried by rotary evaporation. The residues were dissolved in 2 mL cyclohexane and 0.5 mL of the solute

was transferred, purified with a silica gel column (8 mm  $\times$  220 mm) and washed with a mixture of hexane and dichloromethane (1:1, V/V). The first 1-mL of eluate was discarded because it contained non-polar saturated hydrocarbons and was less retained than PAHs by silica gel. The second 2-mL aliquot of eluate was collected, dried by sparging with N<sub>2</sub> and then re-dissolved in 1 mL acetonitrile for HPLC determination. Determination of 16 EPA PAHs was carried out according to the method of Ni et al. (2008). Briefly, analysis was conducted on a Shimadzu Class-VP HPLC system (Shimadzu, Japan), with a fluorescence detector (RF-10AXL, Shimadzu, Japan). A reversed phase column C18 (VPODS 150-4.6 mm I.D., particle size 5 mm, Shimadzu, Japan), using a mobile phase of water and acetonitrile mixture (1:9, V/V) at a constant solvent flow rate of 0.5 mL/min, was used to separate the 16 PAHs. The excitation and emission wavelengths for individual PAHs were set separately.

An external standard mixture was used for quantification of the 16 PAHs. The detection limit of the HPLC method for the 16 PAHs was in the range of 0.12-1.57 µg/kg. Method blanks (solvent) and spiked blanks (standards of EPA610 PAH mixture, LA 96245, Supelco, USA, used to spike the soil) were extracted and analyzed by the methods described above. The recoveries and the relative standard deviations of this method for 16 PAHs were in the ranges of 74%-110% and 0.53%-3.57%, respectively. Results of blanks extracted under the same conditions were below detection limits and sample results are presented without recovery ratio correction. The percentage of PAH removal (*R*, %) was given by Eq. (1):

$$R = (1 - C_{\rm s}/C_{\rm i}) \times 100\% \tag{1}$$

where,  $C_s$  (µg/kg) is the concentration of PAHs in each treatment and  $C_i$  (µg/kg) is the initial PAH concentration present in the soil.

#### 1.5 Enumeration of PAH-degrading bacteria in soil

After incubation, PAH-degrading bacteria in soil were counted using a miniaturized most probable number (MPN) method in 96-well microplates with five replicates per dilution (Wrenn and Venosa, 1996). Briefly, phenanthrene, anthracene, fluorene, and dibenzothiophene were added as sole carbon sources to support the proliferation of aromatics-degrading bacteria. Serially diluted samples were inoculated into the wells and the microplates were incubated at room temperature for 3 weeks. Wells turning yellow or brown owing to the accumulation of partial oxidation products of aromatic substrates were treated as positive. Published MPN tables were used to determine the MPN values.

#### 1.6 Soil enzyme activities

Soil dehydrogenase activity (DHA) was assessed by a modification of the method described by Singh and Singh (2005). Weighed 5 g sub-samples of soil were placed in 50 mL polypropylene centrifuge tubes and mixed with 5 mL of 0.5% 1,3,5-triphenyltetrazolium chloride (TTC)

solution. Tubes were incubated for 6 hr at 30°C in the dark. After incubation, triphenylformazan (TPF) formed by reduction of TTC was extracted with three batches of 100 mL methanol. Tubes were shaken in an orbital shaker at 300 r/min for 1 hr, centrifuged (1744  $\times g$ , 5 min), and the supernatant was filtered through filter paper. Blanks without the addition of TTC were carried out in the same manner. The concentration of TPF was determined by spectrophotometry at 485 nm and the results were expressed as g TPF/(g soil·hr).

Polyphenol oxidase activity was performed as described by Clause and Filip (1990) measured by a UV-Vis spectrophotometer. Briefly, 1 g fresh soil was transferred into an 18 mm  $\times$  20 mm glass tube and mixed with 10 mL of 1% pyrogallol solution. After incubation at 30°C for 1 hr, 12.5 mL of 0.5 mol/L HCl was added into the mixture and shaken manually. The gallic acid produced was extracted with 10 mL of ether and measured spectrophotometrically at 430 nm. Activity was quantified by reference to a calibration curve constructed using gallic acid standard and expressed in mg gallnut/(g soil·hr).

#### 1.7 Physiological profiles of the soil microbial community

Soil microbial community physiological profiles were performed as described by Yao et al. (2003). Briefly, 10 g fresh soil was added to 100 mL distilled water in a 250 mL flask and shaken for 10 min. Ten-fold serial dilutions were made and the  $10^{-3}$  dilution was used to inoculate BIOLOG<sup>®</sup> Eco plates (BIOLOG, USA). The plates were incubated at 25°C, and color development in each well was recorded as optical density (OD) at 590 nm with a plate reader at regular 12 hr intervals. Microbial activity in each microplate, expressed as average well-color development (AWCD), was determined as Eq. (2):

$$AWCD = \sum OD_i/31$$
 (2)

where,  $OD_i$  (590 nm) is the optical density value from each well (i). The Shannon-Weaver index (H) was calculated using an OD of 0.25 as threshold for a positive response (Garland, 1996). The H was calculated by  $H = -\sum p_i \ln p_i$ , and Simpson's index ( $\lambda$ ) of diversity (Simpson, 1949) was calculated by  $\lambda = \sum p_i^2$ , where  $p_i$  is the ratio of the activity on each substrate  $(OD_i)$  to the sum of activities on all substrates ( $\sum OD_i$ ).

#### 1.8 Statistical analysis

Statistical analysis was carried out using the SPSS 14.0 for Windows software package. Data were analyzed by twoway analysis of variance. Mean values were compared by least significant difference (LSD) at the 5% level using SPSS software.

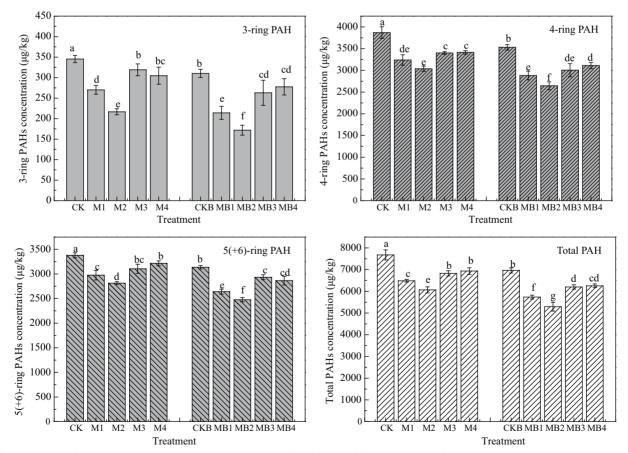
#### 2 Results and discussion

Biodegradation has been proposed as a strategy to enhance the bioremediation of contaminated soil (Hamdian

2007; Teng et al., 2011b). In the present study all treatments showed a significant reduction in PAH concentration following 16 weeks of incubation (p < 0.05) (Fig. 1). However, compared to the MCD amendment and *Paracoccus* sp. strain HPD-2 inoculation combined treatments (MB1, MB2, MB3, and MB4), the PAH removal rate in the bioaugmentation treatment (CKB) was significantly lower (p < 0.05). The present data strongly suggest that microbial factors such as the presence of an adequate consortium of degraders are not the only explanation for the limited extent of biodegradation observed after 16 weeks. It is possible that other factors, such as limited nutrient availability (Teng et al., 2010a) and especially low contaminant bioaccessibility (Allan et al., 2007) may have influenced the degree of degradation.

It is generally accepted that a low level of bioavailability is one of the most important factors involved in the slow biodegradation of hydrophobic organic compounds in soils, as PAHs are very poorly soluble in the aqueous phase where microorganisms are active. It has already been demonstrated that MCD accelerates the degradation kinetics of PAH and PAH bioaccessibility in both the aqueous phase and the solid phase (Bardi et al., 2007). In treatments where MCD was added the extent of PAH degradation was greatly enhanced (Fig. 1) with residual concentrations significantly lower (p < 0.05) than in the control. Soil target PAH concentrations after amendment with MCD and inoculation with HPD-2 after 16 weeks are presented in Fig. 1. For each PAH group there were significant decreases (p < 0.05) in the PAH concentrations for treatments amended with MCD when compared with the initial concentrations. The total PAH concentration in soil with 10% MCD-assisted bioaugmentation (MB2) declined from (7679  $\pm$  230) to (5293  $\pm$  203)  $\mu$ g/kg after 16 weeks. Table 1 shows that MCD amendment, Paracoccus sp. strain HPD-2 inoculation and their combination treatments had significant effects on PAH removal ( $F_{\rm M}$  = 1222.0,  $F_{\rm B} = 802.8$ , p < 0.01). Moreover, the results also showed significant interaction between MCD amendment and Paracoccus sp. strain HPD-2 inoculation ( $F_{M\times B}$  = 3.245, p < 0.05). This suggests that MCD contributes to contaminant desorption by increasing both the mass transfer from the soil particles to the aqueous phase and the solubility of the organic pollutant (Wang et al., 1998; Mayer et al., 2005), and microbial degradation likely acts as a sink for contaminant released into the aqueous/MCD phase by both mass transfer from soil particles to the aqueous phase and transfer from water to microorganisms, or by influencing the hydrophobicity of the surfaces of the microorganisms as shown for some biosurfactants.

Considering the influence that MCD had on specific PAHs, Fig. 2 shows the percentage of compound removal



**Fig. 1** Removal of 3-, 4-, 5-ring and total PAHs in soil before and after 16 weeks of biodegradation. PAH concentration values are means  $\pm$  standard deviations of triplicate determinations. CK: control without MCD amendment or *Paracoccus* sp. strain HPD-2 inoculation; M1, M2, M3, M4: 5%, 10%, 15%, 20% (*W/W*) MCD amendment; CKB: 4% (*V/W*) *Paracoccus* sp. strain HPD-2 inoculation; MB1, MB2, MB3, MB4: 5%, 10%, 15%, 20% (*W/W*) MCD amendment + 4% (*V/W*) *Paracoccus* sp. strain HPD-2 inoculation. Mean values with the same letter are not significantly different among treatments by LSD at the 5% level.

 Table 1
 Effect of inoculation with Paracoccus sp. strain HPD-2 (B)

 and MCD amendment (M) on the total PAHs removal in the soils after
 16 weeks of treatments

Inoculation $(V/W)$	MCD amendment $(W/W)$	Removal rate of total PAHs
0	0	5.38 ± 1.55 a
0	5%	20.09 ± 1.48 c
0	10%	25.24 ± 1.32 e
0	15%	15.89 ± 1.87 b
0	20%	14.54 ± 1.13 b
4%	0	14.05 ± 1.65 b
4%	5%	$29.30 \pm 1.37$ f
4%	10%	34.78 ± 1.11 g
4%	15%	23.58 ± 1.46 d
4%	20%	$22.93 \pm 1.62$ cd
	$F_{\mathrm{B}}$	802.8**
Two-way ANOVA	$\bar{F_{M}}$	1222.0**
•	$F_{M \times B}$	3.245*

Values are means  $\pm$  standard deviation of triplicate measurements. Mean values with the same letter are not significantly different among treatments by LSD at the 5% level.

\*\* Significance at p < 0.01, \* significance at p < 0.05.

following treatment with 10% MCD amendment and inoculation with HPD-2 (MB2) with respect to the control. MCD amendment enhanced the degradation rates of individual compounds, in the range of 15%–85% enhancement. No significant trends with  $\log K_{ow}$  were observed for degradation enhancement by MCD in soil. However, MCD promoted a maximum biodegradation enhancement for 3-ring PAHs.

#### 2.2 Soil PAH-degrading bacteria and microbial activities

In general, bioremediation of PAH-contaminated soils depends on the presence of degrading microorganisms with the desired catabolic capabilities (Margesin et al., 2000). Bioaugmentation has demonstrated an important role in bioremediation by promoting degrading microorganisms in soil contaminated with such compounds. In the present study a significant increase in counts of PAH-degrading bacteria was observed in bioaugmented microcosms after 16 weeks of incubation compared with

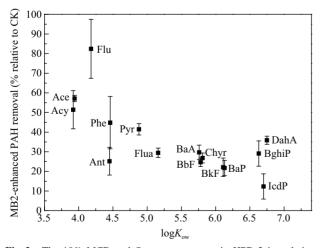


Fig. 2 The 10% MCD and *Paracoccus* sp. strain HPD-2 inoculation enhanced removal of PAHs in soil (relative percentage to the controls without MCD and inoculation) following biodegradation assays. Compounds have been ordered by  $K_{\rm ow}$  value.

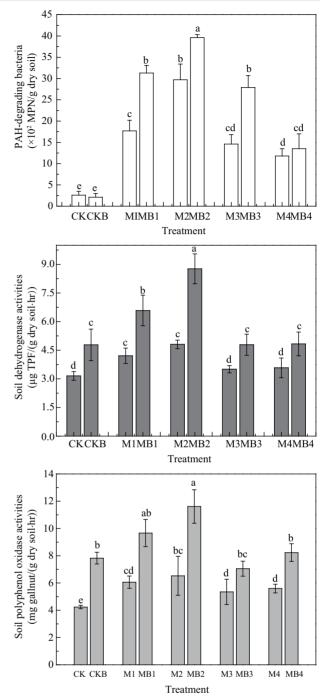


Fig. 3 Soil PAH-degrading bacteria, soil dehydrogenase activities and soil polyphenol oxidase activities under different treatments after 16 weeks of incubation.

the controls (Fig. 3). These results are similar to the results of Teng et al. (2010b) which showed higher proportions of PAH-degrading bacteria in bioaugmented soil treatments. Furthermore, Fig. 3 also shows that PAH-degrading bacteria counts in the MCD-assisted bioaugmented treatments were significantly higher (p < 0.05) than in non-MCD-assisted bioaugmented treatment, and this may have resulted in substantial degradation of 3-, 4- and 5(+6)-ring PAHs in the soil studied.

Soil enzyme activity is one way of describing the general condition of soil microbial populations (Margesin et al., 2000). Soil dehydrogenase activity (DHA), an intra-

cellular process that typically occurs in all intact and viable microbial cells, can be used to determine the presence of viable microorganisms and their oxidative capability (Andreoni et al., 2004). Soil polyphenol oxidase (PPO) is an oxygen-transferring enzyme and can act on specific recalcitrant pollutants by precipitation or transformation to other products, permitting a better final treatment of the waste (Durán and Esposito, 2000). Figure 3 shows that soil DHA and PPO activities increased significantly in the MCD-assisted bioaugmented microcosms. PAHs or their metabolites were likely used as substrates, thus increasing the activity of the enzymes. Changes in enzyme activities in the MCD-assisted bioaugmented microcosms also confirm that MCD-assisted bioaugmentation with Paracoccus sp. strain HPD-2 showed the highest microbiological activity.

#### 2.3 Physiological profiles of the soil bacterial community

Soil is the microbial inhabitancy while soil microorganisms can be sensitive indicators, reflecting subtle changes of soil quality. Organic contaminants are generally considered to have adverse effects on soil microbial functional diversity influencing soil fertility and plant growth, which pose a serious threat to the sustainability of agricultural soils. Therefore, direct measurements of soil microbial community diversity are likely to provide more information for evaluating soil function. The BIOLOG Eco method was used to monitor community-level physiological profiling (CLPP) of soil microorganisms as carbon source utilization patterns and estimate soil bacterial taxonomic diversity (Choi and Dobbs, 1999). The AWCD reflects the sole-carbon-source utilization ability of the soil bacterial community and a measure of soil bacteria activity (Garland, 1996). The Shannon-Weaver index is a measure of the actual richness and evenness of the bacterial population, while the Simpson index is often used to quantify the number of species present, as well as the relative abundance of each species (Garland, 1996; Simpson, 1949). In the present study the community level physiological profiles indicate differences in soil microbial functional diversity among different treatments after 16 weeks of remediation (Fig. 4 and Table 2). The 10% MCD

 
 Table 2
 Sole-carbon-source utilization activity and functional diversity of the soil microbial community after 16 weeks of bioremediation

Treatment	AWCD	Shannon-Weaver index	Simpson index
СК	0.262 ± 0.045 g	2.747 ± 0.434 de	$16.67 \pm 0.01 \text{ f}$
M1	$0.723 \pm 0.084$ f	2.932 ± 0.020 e	$15.68 \pm 0.47$ g
M2	$1.886 \pm 0.029 \text{ b}$	$3.319 \pm 0.027$ bc	26.55 ± 0.76 b
M3	$1.631 \pm 0.135$ cd	3.271 ± 0.022 c	$24.96 \pm 0.80$ c
M4	1.283 ± 0.037 e	$3.123 \pm 0.026 \text{ d}$	$21.01 \pm 0.47 \text{ d}$
CKB	$0.599 \pm 0.169 \text{ f}$	2.749 ± 0.184 e	$12.78 \pm 2.46$ h
MB1	1.227 ± 0.039 e	3.079 ± 0.032 d	19.98 ± 0.79 e
MB2	$2.012 \pm 0.100$ a	3.996 ± 0.010 a	$28.51 \pm 0.53$ a
MB3	1.758 ± 0.063 c	$3.310 \pm 0.012 \text{ b}$	$26.14 \pm 0.39$ bc
MB4	1.314 ± 0.197 d	3.121 ± 0.117 d	$22.02 \pm 1.25 \text{ d}$

Values are means  $\pm$  standard deviation of triplicate measurements. Mean values with the same letter are not significantly different among treatments by LSD at the 5% level.

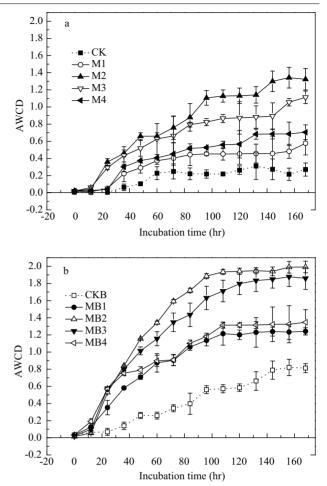


Fig. 4 Variation in average well color development (AWCD) for soil samples from the different treatments after 16 weeks of biodegradation. (a) treatments with only MCD amendment; (b) treatments with MCD amendment and *Paracoccus* sp. strain HPD-2 inoculation. Values are means  $\pm$  standard deviations of triplicate measurements.

treatment showed a higher average well-color development, Shannon-Weaver index and Simpson index. This confirms the previous study of Teng et al. (2010b) which showed marked enhancement of microbial community diversity after 28 days of bioremediation with *Paracoccus* sp. strain HPD-2 inoculation and nutrient addition. It can be concluded that, as a result of bioremediation with *Paracoccus* sp. strain HPD-2 inoculation combined with MCD addition, the microbiological functioning of the PAH-contaminated soil has been at least partially restored.

#### **3** Conclusions

Bioremediation with *Paracoccus* sp. strain HPD-2 inoculation together with 10% MCD addition used in this microcosm study enhanced PAH dissipation in the soil, and increased PAH-degrading bacterial counts and microbial activities in soils, suggesting that this strain of *Paracoccus* and MCD can work together to increase soil microbiological activity, with some restoration of the microbiological functioning of the PAH-contaminated soil. Therefore, MCD-assisted bioaugmentation by *Paracoccus* sp. strain HPD-2 is a promising practical bioremediation strategy for aged PAH-contaminated soil. Elucidation of the potential applicability of this strategy will require further studies involving different soil types and contaminants under field conditions.

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