Influence of *Mucor mucedo* immobilized to corncob in remediation of pyrene contaminated agricultural soil

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Abstract

In recent years, immobilization agents were introduced into organic contaminated soil remediation and more and more materials were screened and used as the immobilizing carrier. However, effect of the decomposition of the immobilizing carrier on the bioremediation was rarely concerned. Therefore, the decomposition experiment of immobilizing carrier - corncob was carried out in the lab with the efficient degradation fungi - *Mucor mucedo* (MU) existing, and PAHs residues E4/E6 of the dissolved organic matter and microbial diversity during the decomposition process were studied. The results showed that: a) During the decomposition, the degradation of pyrene (Pyr) was mainly in the first 28d in which the content of extractable Pyr decreased rapidly and the highest decrease was in the treatment with only *Mucor mucedo* (MU) added. b) Analysis of E4/E6 changes showed that rich microorganisms could promote aromatization and condensation of humus. c) From the diversity index analysis it can also be seen that there is no significant difference in effects of PAHs on the uniformity of microorganisms. These results will not only be useful to have a better understanding of the bioavailability of contaminants adsorbed to biodegradable carriers in PAHs contaminated soil remediation, but also be helpful to perfect the principle of immobilized microbial technique.

Keywords: Immobilizing carrier, Microbial diversity, *Mucor mucedo*, PAHs degradation

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1. Introduction

Polycyclic aromatic hydrocarbons (PAHs) contamination has become a common soil pollution type and has been highly concerned as the contamination degree exacerbating. The magnitude of pollution levels has increased from μg/kg to mg/kg and the detection rate has risen to more than 80% from less than 20%. In addition, approximately 20% of major agricultural products’ content of PAHs exceeds the standard [1]. PAHs contamination will result in degradation of soil function and reduction of agricultural production. More importantly, through food chain or food web contaminants will be taken into and accumulated in human body, which endanger human health potentially. A large number of studies showed that PAHs content taken into human body through soil was greater than that through water and atmosphere [2]. Therefore, remediation of PAHs contaminated soil has become the important research subject. The microbial remediation technique has become a continuous concern for its low cost, high efficiency and environmental responsibility [3-4]. In addition, introduction of immobilization agents to PAHs contaminated soil remediation has made some achievements in immobilizing carrier screening and processing etc., which is gradually shifting from experimental research and development to field application [5-7]. Therefore, how to screen high efficient degrading bacteria is the key to efficient application of microbial remediation technology. According to Wang et al [8], three strains through separate and mixed immobilization, respectively, were used to degrade pyrene and benzo[a]pyrene (B[a]P) in soil. The results showed that the degradation efficiency of immobilized microorganism was much better than that of
free bacteria and compared with separate immobilization, the degradation efficiency was generally higher through mixed immobilization. Li et al [9] studied phenanthrene and pyrene in soil based on the immobilized microbial technique (IMT) and Li believed that IMT was helpful to enhance the competitiveness of introduced microorganisms in soil and was an effective method for the degradation of PAHs in soil. Gentili et al [10] immobilized Rhodococccusorynebacterioides to chitin and chitosan to repair oil polluted seawater and the results showed that compared with separate free bacteria treatment, the removal rate of contaminants by immobilized bacteria was increased by approximately 30%. Researches also showed that *Phanerochaetechrysosporium* immobilized by bagasse could greatly enhance the activity of manganese peroxidase and the removal rate of anthracene in water [11]. By immobilizing Mucor sp. to corn cob in order to repair B[a] P contaminated soil, Su et al [12] found that immobilized Mucor sp. had a better environmental resilience and a faster reaction speed, obviously resulting in a higher removal rate of B[a]P. According to Dzul-Puc et al [13], in order to repair B [a] P contaminated soil *Phanerochaete chrysosporium* was immobilized to pine needle powder and bagasse, respectively, and found that the B [a] P degradation rate was faster when bagasse was taken as immobilizing carrier. Bacteria could also be immobilized to plant residues, which could enhance removal rate of PAHs by immobilized bacteria and indigenous microorganisms to a certain extent [14-16].

Therefore, in this paper the decomposition process of immobilizing carrier (organic material) was simulated in laboratory in order to study the effects of high
efficient degrading bacteria on the decomposition process of immobilizing carrier and microbial diversity changes, so as to provide theoretical support for revealing the influence mechanism of immobilizing carrier in degradation of contaminants in soil.

2. Materials and Methods

2.1. Materials

Corncob was collected from the Shenyang Ecological Experimental Station, Chinese Academy of Sciences, in the Liaoning Province of China (E 123°21′58.6″, N 41°31′6.53″). PAHs-degrading fungus, *Mucor mucedo* (MU) which had been successfully applied to remediation of contaminated soil of farmland in Shenfu sewage irrigated area (SIA, southeast of Shenyang, Liaoning province, China) were screened from heavily contaminated soil of SIA using the method of enrichment and culture in a mineral plate where PAHs were added as sole carbon and energy source and cultured according to the study of Li et al. [17-18]; Pyrene (Pyr) was purchased from sigma cooperation with purity of 98%; Contaminated soil was collected from topsoil (0-20cm) of long-term contaminated farmland in upstream of Shenfu irrigated area and fresh soil was stored at 4°C.

2.2. Experimental Methods

2.2.1. Corncob pretreatment

The dried corncobs were crushed to particles with diameter 0.6~0.8 mm and were pretreated by soaking in calcium hydroxide for 24 hr and then mixed with other accessories in the proportion of the formula [18] And stirred to constitute the
microorganism immobilizing carrier. The water content of the carrier mixture was maintained between 30%–50%, while pH was adjusted to 7.0 with Ca (OH)\(_2\) solution. The carrier mixture was then sterilized by autoclaving twice at 121°C, 1×10\(^5\) pa for 90 min.

2.2.2. Soil solution

100 g pre-activated field soil (wet) was mixed with 500 ml sterilized distilled water. The mixture was shaken on a reciprocal shaker (175 rpm) for 3h then stand 16h to clarify soil solution. The aqueous suspensions (soil solution) were used in the following experiment. The pre treatment of the field soil meant culture the field soil at 28 °C for 3-5 days to activate the microbial populations in soil in which the water content was kept in 20-30% (W/W).

2.2.3. Inoculated Fungi

MU stored in laboratory was inoculated into potato liquid nutrient medium (40.0 g saccharose, 4.0 g (NH\(_4\))HPO\(_4\), 1.0 g KH\(_2\)PO\(_4\), 0.5 g MgSO\(_4\)·7H\(_2\)O and 0.05 g vitamin B1 per L) and then was cultured for 3 days in an incubator at 28 °C.

2.2.4. Experimental design

The target PAHs, pyrene, was dissolved in HPLC grade acetone and kept at -20°C until required. According to the Table 1, the treatments were prepared in the sterile room. The pre-treated corncobs, soil solution, pyrene and MU were added and mixed together. The water content of the mixture was maintained between 60%–70%, and the mixture was cultured at 28 °C for 120 d. All the treatments were triplicates. The samples were collected at 0 d, 7 d, 14 d, 28 d, 42 d, 63 d and 120 d. The collected
samples were stored in 0-4℃. The sterilized water was added each three days based on the weight balance.

**Table 1.** the treatments during the decomposition

<table>
<thead>
<tr>
<th>Treatments</th>
</tr>
</thead>
<tbody>
<tr>
<td>A (control experiment)</td>
</tr>
<tr>
<td>1</td>
</tr>
<tr>
<td>2</td>
</tr>
<tr>
<td>3</td>
</tr>
<tr>
<td>B</td>
</tr>
<tr>
<td>1</td>
</tr>
<tr>
<td>2</td>
</tr>
<tr>
<td>3</td>
</tr>
</tbody>
</table>

Concentration of Pyr is 20mg kg⁻¹ dried corncob; Percentage of soil solution and MU is Volume ratio in which MU means *M. Mucedo*.

2.3. Determination

2.3.1. Pyr extraction and analysis

Extraction of pyrene was performed according to Song et al. [19]: 20ml of dichloromethane and 3.0 g air-dried corncob were added to each corncob sample in the glass centrifuge tube, then, samples were extracted by sonication at 30-35 ℃ for 2h. The extraction was repeated three times and the extracts were combined. The final extract was completely dried by evaporation of the dichloromethane under a stream of nitrogen. The residuals were dissolved in 2ml filtered methanol and filtered through a 0.22-μm hydrophobic syringe filter (Nantong Filterbio Membrane CO., LTD). The concentrations of PYR were determined by high performance liquid chromatography (Agilent 1200s) equipped with a fluorescence detector [20].

The wavelength for the fluorescence detector was 240 nm. Separation was performed using a ZORBAX Eclipse C18 analytical column (250×4.6 mm id, 5μm)
with an operating temperature of 35℃. The isocratic immobile-phase consisted of methanol and water (90:10 v/v) at a flow rate of 0.7 µl min⁻¹. The injection volume was fixed at 10µl. The moisture was determined [by ISO 11465:1993] to allow data presented on a dry matter basis.

2.3.2. E4 / E6 ratio in dissolved organic matter (DOM) of corncob

The E4/E6 ratio is extensively used in soil science to indicate the humification degree (decomposition of organic matter); the progressive humification, which had the strong adsorption for pyrene due to the strengthened hydrophobic property, is indicated by a decreasing E4/E6 quotient [21]. The E4/E6 ratio was determined by dividing the absorbance at 465nm (Abs465) by that at 665nm (Abs665) for the individual samples [22]. DOM was obtained according to Long et al.[23]. 1.0 g sample was put into a centrifuge tube and 20 ml of ultrapure water was added and shaken for 4 h at 200 rad·min⁻¹. The supernatant was obtained after the mixture was centrifuged at 4000 rpm for 10min4 h shock at 200 rad·min⁻¹. Then, the supernatant was filtrated through the 0.45um water membrane and stored in polypropylene bottles at 4℃ prior to analysis (). The obtained DOM was measured at the wavelength of 465 nm and 665 nm by UV-765 spectrophotometer. If the color of leach liquor was too deep and was beyond reading range of the spectrophotometer, before determination, the leach liquor needed to be diluted with ultrapure water several times before determinationat first.

2.3.3. Population structure changes of soil microorganisms

Population structure of soil microorganismswas analyzed by polymerase chain reaction - denaturing gradient gel electrophoresis (PCR-DGGE) method. PCR-DGGE
technique is a common method for the analysis of microbial population structure in microbial ecology, which has been widely used at present. Given laboratory conditions, PCR-DGGE technique was used to determine changes of microbial community structure and microbial diversity.

PCR-DGGE in present study was carried out according to the study of Su et al. [24]. The Corncob samples (0.1 g wet weight) were used for DNA extraction with the TaKaRa MiniBEST Bacteria Genomic DNA Extraction Kit Ver. 3.0. The primer pair GC-954F / 1369R was used to obtain DNA fragments 496 bp in length. All PCR was performed with a PTC-200 thermal cycler (MJ Research, Waltham MA). The PCR products then were verified by running a 1.2 % (w/v) agarose gel electrophoresis. DGGE was carried out using a DCode Universal Mutation Detection System (Bio-Rad, Richmond, CA). The PCR-DGGE banding pattern was analyzed by Quantity-One image analysis software (Bio-Rad) and Shannon index (H), Shannon evenness (E) and Simpson dominance (D) were calculated according to the following equations.

\[ H = \sum_{i=1}^{S} P_i \ln(P_i) \]  

(1)

\[ E = H / \ln S \]  

(2)

\[ 1 / D = 1 - H = \sum_{i=1}^{S} P_i^2 \]  

(3)

Where: \( P_i \) is the ratio of the gray scale of DGGE band \( P_i \) to that of all bands in each lane; \( S \) is the total number of DGGE bands.
3. Results and Discussion

3.1. Pyr Degradation

In group B where Pyr was added, the contents of extractable Pyr all decreased rapidly in the first 28 days and reached to a relatively low value in day 28 since the immobilizing carrier began to decompose (Fig. 1), after which they undergone rises and falls until the presence of the minimum in day 120. In initial decomposition stage, the decline rate of extractable Pyr in B2 was greater than that in B3 and B1. Content of extractable Pyr in B3 changed little during day 14 to day 63, but after day 63 it decreased dramatically. After the troughs in day 28 in B1 and B2, the content of extractable Pyr in B1 and B2 rose first and then fell with the peak values in day 42 and day 63, respectively. The content of extractable Pyr in B2 where only *Mucormucedo* was added was far lower than that in B1 and B3 (except in day 63). The content of extractable Pyr in B3 and in B1 had significant difference from 14d to 42d. After 120 days’ decomposition, there was no significant difference in contents of extractable Pyr in B1, B2, and B3. As the decomposition deepening, the content of extractable Pyr in undecomposed parts decreased gradually. In addition, Pyr usually served as a carbon source for microorganisms, leading to the decomposition of Pyr and the decrease of extractable Pyr. However, the increase of pyrene concentration during the degradation process was probably because of the release rate of Pyr sequestrated by corncob during the degradation of Pyr [17].
Fig. 1. The exactable Pyr in corncob during the decomposition process

Based on the degradation dynamics of the above five materials, the relation among the microbial population structure and the characteristics of the corncob during the decomposition of the corncob were studied. For instance, by analyzing the changes of microbial components of and humic substances during the decomposition of cow dung + straw, Chen [25] found that inoculation with microorganism resulted in faster decomposition of organic fractions and more formation of humic substances which might affect the degradation of the PAHs [26]. Tang found that DOM from the rice straw enhanced the solubility of pyrene gradually in soil during the decomposition process because more and more hydrophobic sub-fractions were produced in SOM which could bind pyrene strongly [27]. The content and chemical composition of DOM undergone dynamic changes during the decomposition of rice straw [28]. Cheng found that DOM had an obvious solubilizing effect for PAH [29]. According to researches of Xiao, effects of DOM composition on Pyr adsorption were related to concentration of each component of DOM and in addition, the content of Pyr adsorbed by soil decreased with the increase of concentration of DOM.
components [30].

3.2. E4 / E6 Changes

E4 / E6 can describe the type and nature of Humic acid and reflect the condensation degree of aromatic ring in humic acid molecules, the aromatization degree and the molecular weight which are negatively correlated to the value of E4 / E6. The values of E4 / E6 are different for different sources (Fig. 2).

It can be seen from Fig. 2 that E4 / E6 values changed greatly in 0d~14d and 63d~120d, while in 14d~63d the changes were relatively small. E4 / E6 values in all groups declined obviously before day 14 and then generally leveled off, however they increased to some extent after day 63. The large gap among E4 / E6 values of the treatments from 0 to 14d was probably due to the high microbial activity and pyrene degradation during this period, in which the great amount of the secretions from microorganisms and various intermediate products of pyrene were produced being consistent with the degradation process of pyrene. Rising trend during 63d~120d was probably because of the emergence of new active microorganisms. E4 / E6 values in A3 and B3 where soil solution was added were relatively low, indicating that rich microorganisms could promote aromatization and condensation of humus.

Fig. 2 Humic acid E4/E6 changes during decomposition process
3.3. Microbial Diversity Changes during Decomposition Process

Microbial diversity not only represents stability of microbial community, but also reflects the effects of soil ecological mechanism and soil stress on community stability etc. from the other side. Researches and analysis of richness, evenness and other indexes of microorganisms were carried out to provide theoretical support and necessary experimental data. Diversity indexes such as Shannon index (H), Shannon evenness (E) and Simpson dominance (D) can not only reflect the characteristics of soil microbial community structure from different aspects, but also can be used to analyze differences in community types and structures and dynamic changes of community succession.

**Table 2.** Shannon diversity index (H) of soil solution of different treatment at different time

<table>
<thead>
<tr>
<th>Treatments/Days</th>
<th>3d</th>
<th>7d</th>
<th>14d</th>
<th>28d</th>
<th>42d</th>
</tr>
</thead>
<tbody>
<tr>
<td>A1</td>
<td>1.51</td>
<td>1.41</td>
<td>1.34</td>
<td>1.22</td>
<td>1.12</td>
</tr>
<tr>
<td>A3</td>
<td>1.62</td>
<td>1.82</td>
<td>1.26</td>
<td>1.38</td>
<td>1.42</td>
</tr>
<tr>
<td>B1</td>
<td>1.40</td>
<td>1.61</td>
<td>1.25</td>
<td>1.02</td>
<td>1.84</td>
</tr>
<tr>
<td>B3</td>
<td>1.38</td>
<td>1.31</td>
<td>1.20</td>
<td>1.10</td>
<td>1.68</td>
</tr>
</tbody>
</table>

**Table 3.** Shannon evenness index (E) of soil solution of different treatment at different time

<table>
<thead>
<tr>
<th>Treatments/Days</th>
<th>3d</th>
<th>7d</th>
<th>14d</th>
<th>28d</th>
<th>42d</th>
</tr>
</thead>
<tbody>
<tr>
<td>A1</td>
<td>0.77</td>
<td>0.80</td>
<td>0.81</td>
<td>0.80</td>
<td>0.79</td>
</tr>
<tr>
<td>A3</td>
<td>0.76</td>
<td>0.78</td>
<td>0.79</td>
<td>0.78</td>
<td>0.80</td>
</tr>
<tr>
<td>B1</td>
<td>0.78</td>
<td>0.79</td>
<td>0.77</td>
<td>0.78</td>
<td>0.80</td>
</tr>
<tr>
<td>B3</td>
<td>0.82</td>
<td>0.81</td>
<td>0.78</td>
<td>0.79</td>
<td>0.82</td>
</tr>
</tbody>
</table>
As can be seen from Table 2, there was no Pyr added in soil solution in A1 and A3 treatments. The bacterial diversity index (H) decreased, which showed that bacterial population number decreased continuously with the continuous consumption of nutrients. Pyr was added in soil solution in B1 and B3 treatments. The bacterial diversity index (H) experienced a rising trend after the first reduction, which showed some vulnerable bacteria was inhibited in early decomposition, while the population number increased, because Pyr provided carbon source for the later decomposition. Comparison of B1 and B3 adding MU, bacterial diversity index (H) was lower than that without added. The reason was competition of carbon source, which was similar to the result that *Mucor* separate treatment of the decomposition was better than the other treatment. As can be seen from Table 3, Shannon evenness index (E) of all treatments varied little, its E value was between 0.77-0.82. As can be seen from Table 4, Simpson dominance (D) of all treatments increased continuously, and reached a maximum value in the 28d. So Pyr adding MU decomposition had greater impact to microbial species in soil, while had little impact to Shannon evenness (E) and Simpson dominance (D) of bacteria.

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**Table 4.** Simpson dominance (D) of soil solution of different treatment at different time

<table>
<thead>
<tr>
<th>Treatments/Days</th>
<th>3d</th>
<th>7d</th>
<th>14d</th>
<th>28d</th>
<th>42d</th>
</tr>
</thead>
<tbody>
<tr>
<td>A1</td>
<td>1.41</td>
<td>1.48</td>
<td>1.50</td>
<td>3.06</td>
<td>1.76</td>
</tr>
<tr>
<td>A3</td>
<td>1.46</td>
<td>1.38</td>
<td>1.36</td>
<td>2.78</td>
<td>1.51</td>
</tr>
<tr>
<td>B1</td>
<td>1.58</td>
<td>1.42</td>
<td>1.61</td>
<td>1.86</td>
<td>1.48</td>
</tr>
<tr>
<td>B3</td>
<td>1.52</td>
<td>1.51</td>
<td>1.53</td>
<td>1.68</td>
<td>1.64</td>
</tr>
</tbody>
</table>

*There was no soil solution in T2 and T5, so fungi DNA was not found.*
4. Conclusions

1. As the decomposition proceeded, the content of extractable Pyr decreased rapidly and then undergone rises and falls until the presence of the minimum in day 120. However, the increase in pollutant concentration during the decomposition process was probably because the release rate of Pyr adsorbed by corncob was faster than the degradation rate of Pyr.

2. Treatment where only MU was added was more effective than other treatments in PAHs degradation especially in earlier period. In addition, Pyr- degrading bacteria also existed in indigenous microorganisms, which significantly reduced the content of Pyr in corncob. Adding both MU and soil microorganisms didn’t significantly improve the efficiency of Pyr degradation, probably due to the competition between MU and soil microorganisms for carbon and nitrogen and the low concentration of MM (5%) compared to the only MU treatment.

3. Microbial life activities and superposition of various intermediate products in decomposition process probably resulted in the large gap among E4 / E6 values in different treatments. E4 / E6 values in two treatments where soil solution was added were relatively low, indicating that rich microorganisms could promote aromatization and condensation of humus.

4. Pyr adding MU decomposition had greater impact to microbial species in soil, while had little impact to Shannon evenness (E) and Simpson dominance (D) of bacteria.
Acknowledgments

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