

## RESEARCH ARTICLE

# Biogasification of methanol extract of lignite and its residue: A case study of Yima coalfield, China

Jianmin Liu<sup>1,2</sup>, Hengxing Ren<sup>1,3</sup>, Yi Jin<sup>1,2\*</sup>, Huan He<sup>4</sup>, Linyong Chen<sup>1,3</sup>, Guofu Li<sup>3</sup>, Baoyu Wang<sup>1,3</sup>

**1** School of Resources & Environment, Henan Polytechnic University, Jiaozuo, P R China, **2** Collaborative Innovation Center of Coalbed Methane and Shale Gas for Central Plains Economic Region, Jiaozuo, P R China, **3** State Key Laboratory of Coal and CBM Co-mining, Jincheng, P R China, **4** School of Chemical Engineering and Technology, China University of Mining and Technology, Xuzhou, China

\* [jinyi2005@hpu.edu.cn](mailto:jinyi2005@hpu.edu.cn)



## OPEN ACCESS

**Citation:** Liu J, Ren H, Jin Y, He H, Chen L, Li G, et al. (2022) Biogasification of methanol extract of lignite and its residue: A case study of Yima coalfield, China. *PLoS ONE* 17(10): e0275842. <https://doi.org/10.1371/journal.pone.0275842>

**Editor:** Bawadi Abdullah, Universiti Teknologi Petronas, MALAYSIA

**Received:** April 1, 2022

**Accepted:** September 24, 2022

**Published:** October 12, 2022

**Copyright:** © 2022 Liu et al. This is an open access article distributed under the terms of the [Creative Commons Attribution License](https://creativecommons.org/licenses/by/4.0/), which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited.

**Data Availability Statement:** All relevant data are within the manuscript.

**Funding:** This work was supported by the National Natural Science Foundation of China (No. 41972175), the Science and Technology Major Project of Shanxi Province, China (No. 20181101013-1), Henan Science and Technology Research Project (No. 182102310901), Shanxi Province Science Foundation for Youths (No. 201801D221354), and the Fundamental Research Funds for the Universities of Henan Province (No. NSFRF170803).

## Abstract

To investigate the biogas generation characteristics of the organic matter in lignite, methanol extraction was conducted to obtain the soluble fraction and the residual of lignite, which were subsequently taken as the sole carbon source for biogas production by a methanogenic consortium. Afterward, the composition of compounds before and after the fermentation was characterized by UV-Vis, GC-MS, and HPLC-MS analysis. The results indicated that the methanogenic microorganisms could produce H<sub>2</sub> and CO<sub>2</sub> without accumulating CH<sub>4</sub> by utilizing the extract, and the methane production of the residue was 18% larger than that of raw lignite, reaching 1.03 mmol/g. Moreover, the organic compounds in the methanol extract were degraded and their molecular weight was reduced. Compounds such as 1, 6-dimethyl-4-(2-methylethyl) naphthalene, 7-butyl-1-hexylnaphthalene, simonellite, and retene were completely degraded by microorganisms. In addition, both aromatic and non-aromatic metabolites produced in the biodegradation were detected, some of which may have a negative effect on the methanogenesis process. These results revealed the complexity of the interaction between coal and organism from another point of view.

## Introduction

Coal-bed methane (CBM), a new energy currently being promoted and developed in China, is a clean energy of high quality. Many reports have indicated that nearly 20% of the methane gas in the developed CBM resources is produced by microorganisms [1]. This has drawn the attention of many researchers worldwide to the promotion of CBM production by microorganisms and proposed a mechanism for the bioconversion of coal to methane [2]. Biogas obtained from in-lab gas production simulation experiments typically consists of methane and unconverted hydrogen and carbon dioxide [3–5]. Although the methane concentration in the biogas is lower than that in the original coal seam, the presence of carbon dioxide allows the syngas to

**Competing interests:** The authors have declared that no competing interests exist.

be further produced by other means such as methane dry reforming, providing a new way for the clean and efficient utilization of coal [6, 7].

Biogasification of coal is a process in which methane is produced by anaerobic degradation of organic matter in coal by methanogenic bacteria. During this process, soluble organic matter is released from coal and continuously degraded by microorganisms to form precursor substances such as acetic acid, carbon dioxide, hydrogen, and C1 compounds such as methanol, which are finally converted into methane [8–10]. Heteroatoms such as nitrogen, oxygen, and sulfur in coal macromolecules are considered to be the active sites of biodegradation [11–15]. The soluble oxygen-containing organics are readily released from coal and come into contact with microorganisms, however, their effects on each stage of the biogasification process are unknown.

Methanol, a relatively highly polar, readily available organic solvent, has enriched oxygen-containing compounds in coal [16]. In addition, the methanol extract of coal usually contains alkanes, aromatics and other heteroatomic compounds, which are involved in the biodegradation of coal [17–21]. Although aromatic compounds seem more resistant, they can still be degraded under anoxic conditions through the cleavage of aromatic rings [22, 23]. Solvent extraction also causes changes in coal organic composition and pore structure, which also have implications for microbial-coal interactions [24, 25]. Therefore, the research on the biodegradation of methanol extracts of coal and its residue is beneficial for exploring the interaction mechanism between microorganisms and coal.

In this research, methanol was employed as an organic solvent for the Soxhlet extraction to obtain the organic matter of Yima lignite. And microorganisms with good anaerobic gas production effect on lignite, pre-stored in the laboratory, were used as bacterial sources. Afterward, extracts and residue were utilized as substrates to conduct gas production simulation experiments. Finally, various analytical methods were combined to analyze the gas production of extracts and residue, as well as the composition and content of organic matter in the gas production process. This research provided an experimental basis for the subsequent analysis and degradation mechanism of biogas-generating active organic components in coal.

## Materials and methods

### Lignite methanol extraction and GC-MS analysis

Lignite was collected from the No. 2–3 coal seams of Qianqiu Mine in Yima field in Henan province, with a buried depth of 798.5 m and a coal thickness of 6.82 m. The sedimentary age of the lignite sample with  $R_o = 0.48\%$  was the Middle Jurassic. After sampling on site, coal was stored in a nitrogen-filled sealed tank. Before conducting the experiments, the oxide layer was removed and pulverized to below 120 meshes. After being dried at  $70^\circ\text{C}$  to a constant weight, it was stored in a sample bag and named YM.

The methanol extraction process of coal was as follows: (1) 50 g of lignite was weighed, and 250 mL methanol was employed as the extraction solvent to perform Soxhlet extraction at  $68^\circ\text{C}$  for 80 h [17]; (2) after the extraction was completed, the extract was concentrated at  $45^\circ\text{C}$  using a rotary evaporator. Then, the concentrated extract was made up to 100 mL with methanol and recorded as M1; (3) the residual coal was dried to a constant weight at  $70^\circ\text{C}$  and stored in the sample bag, recorded as M2. The calculation formula for methanol extraction rate was as follows:

$$P = (m_0 - m_1)/m_0 \times 100\% \quad (1)$$

where  $P$  is the extraction rate;  $m_0$  is the mass of raw coal, and  $m_1$  is the mass of residual coal. The methanol extraction rate of Yima lignite was 3.12%.

After diluting M1 10 times with methanol, the organic components of the extracts were analyzed by gas chromatography-mass spectrometry (GC-MS, Agilent 7890A-5795C). The column was VF-WAXms (30 m×250 μm×0.25 μm), the post-operation temperature was 280°C which was maintained for 5 min, and no split injection was applied. The inlet temperature was 250°C, the injection volume was 0.8 μL, the purging rate was 15 mL/min, and the purging time was 0.2 min. Also, the carrier gas was helium with high purity, the column flow rate was 1.0 mL/min, and the initial temperature was 60°C which was kept for 2 min. Then the temperature was raised to 250°C at a rate of 10°C/min and was maintained for 20 min. The MS was operated in the electron impact mode, with an ionization energy of 70 eV. The mass spectrometric identification was performed using the mass spectral database NIST2008.

### Biogas production experiment of methanol extract and residue from lignite

The biogas production experiment consisted of four experimental groups: M1, M2, YM, and CK. All groups were set up in triplicate. The substrate of each experimental group was M1 2 mL, M2 2 g, YM 2 g, and methanol 2 mL, respectively. The components of the medium used in the experiment were as follows: K<sub>2</sub>HPO<sub>4</sub> (2.9 g), KH<sub>2</sub>PO<sub>4</sub> (1.5 g), NH<sub>4</sub>Cl (1.8 g), MgCl<sub>2</sub> (0.4 g), yeast extract (0.2 g), L-cysteine hydrochloride (0.5 g), deionized water (1000 mL).

Microorganisms with a good anaerobic gas production effect on lignite, pre-stored in the laboratory, were employed as bacterial sources. The bacterial group composition at the level of phylum predominantly included: *Firmicutes* (27%); *WWE1* (25%); *Bacteroidetes* (21%), *Synergistetes* (13%), *Proteobacteria* (5%), and *Chloroflexi* (1%), and *Archaea* which mainly belonged to the *Euryarchaeota*. The total number of bacteria per ml was  $1.3 \times 10^7$ , and the inoculation amount was 4%. The experimental period was 180 d.

Gas chromatography (GC, Agilent 7890) was employed to analyze the gas composition in the process of biogas production, with a Carbonplot column (60 m×320 μm×1.5 μm), TCD detector, and gas-tight injection needle. The injection volume was 0.5 mL. The inlet temperature, column temperature, and the detector temperature were 150°C, 30°C, and 200°C, respectively.

### Organic composition analysis of extract and residue gas production system

The fermentation broth was initially aspirated from the anaerobic bottle with a sterile sampler. Then it was collected through a 0.22 μm microporous filter membrane. A dual-beam ultraviolet-visible light spectrophotometer (UV-Vis, Unico, UV4802) was employed to perform spectral scanning. The scanning range was 190–400 nm with an interval of 1 nm. The deionized water was used as a blank to qualitatively analyze the composition of organic matter in the fermentation broth.

Quantitative analysis of polar organic components of the fermentation broth was also carried out by liquid chromatography-mass spectrometry (HPLC-MS, Agilent 1290 6530 QTOF equipped with electrospray ionization source) equipped with an Agilent Zorbax C8 (1.8 μm×4.6 mm×50 mm). The mobile phase was methanol and 0.1% formic acid with a 0.5 mL/min flow rate. The column temperature was 25°C, and the injection volume was 10 μL. The mass spectrometry acquisition mode was positive ion mode, and the fragmentor voltage and the capillary voltage were 130 V and 3500 V, respectively. N<sub>2</sub> was used as collision gas and dryer gas. The mass-to-charge ratio scanning range was 50–450 m/z.

The non-polar organic matter in the fermentation broth was initially enriched with a solid-phase extraction column (Agilent Bond Elut C18, 500 mg, 120 μm, 6 mL) through the following steps: (1) 6 mL of methanol and 6 mL of deionized water were added to the extraction column sequentially to activate it. (2) 10 mL of the sample filtered with a 0.22 μm

microporous membrane was added and passed the column at a flow rate of about 2 mL/min. (3) The column was rinsed with 10 mL of deionized water initially, and then the column was blow-dried with nitrogen for 10 min. (4) The organic matter was eluted with 2 mL of methanol. Then the eluent was concentrated to a volume of 1 mL with nitrogen at 45°C. After the enrichment, the sample was analyzed using the GC-MS method described in section 2.1.

## Results and discussion

### The organic composition of methanol extract of lignite

The GC-MS total ion current chromatogram (TIC) of the extract is shown in Fig 1. The library search was performed on the chromatographic peaks. The 20 compounds with a matching degree greater than 60 are shown in Table 1.

Alcohols, esters, carboxylic acids, amides, phenols, aromatic hydrocarbons, and heterocyclic compounds were mostly detected in the methanol extract. More than half of these compounds contained oxygen atoms, indicating that methanol extraction had an enrichment effect on oxygen-containing compounds in coal [16]. The oxygen element in the extract primarily existed in the form of hydroxyl, carbonyl, ester and amide groups, and the nitrogen element chiefly existed in the form of the heterocycle. The presence of nitrogen and oxygen heteroatoms provided potential sites for biodegradation. The aromatic hydrocarbons in the extract were principally alkyl substituents of naphthalene and phenanthrene, of which 7-butyl-1-hexylnaphthalene showed the highest abundance.

### Analysis of biogas production composition of lignite methanol extract and residual coal

The results of biogas production in each experimental group are illustrated in Fig 2. Among the experimental groups, the M1 group had the largest H<sub>2</sub> production (0.72%, the cumulative H<sub>2</sub> production was 0.05 mmol/g). Also, all groups produced CH<sub>4</sub> (methane content of CK>M2>YM), except the M1 group. The CK group had a large amount of CH<sub>4</sub>, CO<sub>2</sub>, and a small amount of H<sub>2</sub>, which indicated that the bacteria employed could directly use methanol as a substrate to produce methane. Furmann et al. [14] found that a small amount of CH<sub>4</sub>

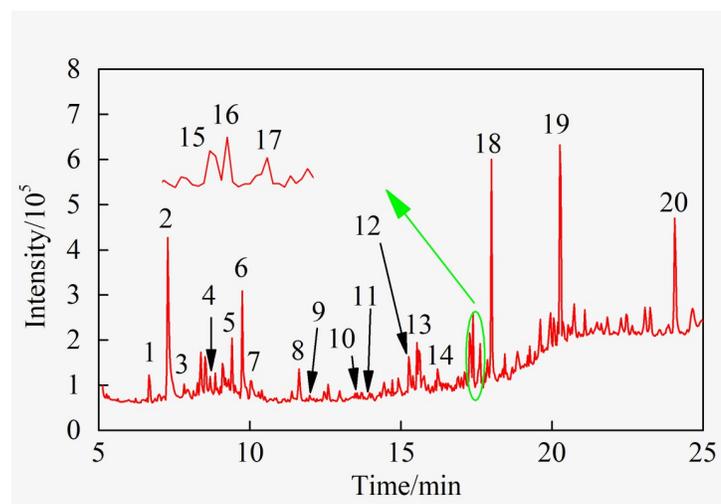


Fig 1. TIC of GC-MS of methanol extract.

<https://doi.org/10.1371/journal.pone.0275842.g001>

Table 1. Identified molecular compounds in methanol extract.

NO.	Retention time/min	Composition	MW	Name	Prob.
1	6.697	C <sub>4</sub> H <sub>6</sub> O <sub>4</sub>	118	Dimethyl oxalate	80
2	7.314	C <sub>2</sub> H <sub>4</sub> O <sub>2</sub>	60	Acetic acid	90
3	7.828	C <sub>2</sub> H <sub>5</sub> BrO	124	2-bromoethanol	77
4	8.684	C <sub>7</sub> H <sub>14</sub> O <sub>3</sub>	146	Pentanoic acid, 2-hydroxy-4-methyl-, methyl ester	73
5	9.438	C <sub>11</sub> H <sub>18</sub>	150	Trans-1,2,3,4,4a,5,8,8a-octahydro-4a-methylnaphthalene	80
6	9.78	C <sub>14</sub> H <sub>22</sub> O	206	3,5-bis(1,1-dimethylethyl)-phenol	87
7	10.054	C <sub>8</sub> H <sub>14</sub>	110	1,2-dimethylcyclohexene	84
8	11.63	C <sub>15</sub> H <sub>22</sub>	202	1-methyl-4-(1,2,2-trimethylcyclopentyl)benzene	98
9	12.007	C <sub>4</sub> H <sub>8</sub> O	72	Trans-2,3-dimethylethylene oxide	79
10	13.72	C <sub>4</sub> H <sub>8</sub> O	72	Cis-2,3-dimethylethylene oxide	73
11	13.96	C <sub>12</sub> H <sub>13</sub> NO	187	1-(1,3-dimethyl-1H-indol-2-yl) ethanone	81
12	15.296	C <sub>13</sub> H <sub>9</sub> F <sub>2</sub> NO	233	N-(4-fluorophenyl)-3-fluorobenzamide	73
13	15.57	C <sub>17</sub> H <sub>34</sub> O <sub>2</sub>	270	Methyl hexadecanoate	95
14	16.255	C <sub>14</sub> H <sub>13</sub> N <sub>3</sub> O <sub>2</sub>	255	2-(3,4-dimethoxyphenyl)-1H-imidazo[4,5-c]pyridine	76
15	17.146	C <sub>10</sub> H <sub>13</sub> NO <sub>3</sub>	227	2-[2-(4-nitrophenoxy)ethoxy]ethanol	72
16	17.317	C <sub>14</sub> H <sub>18</sub>	186	(2-ethyl-3,3-dimethyl-cycloprop-1-allyl)benzene	79
17	17.626	C <sub>15</sub> H <sub>18</sub>	198	1,6-dimethyl-4-(2-methylethyl)naphthalene	77
18	18.002	C <sub>20</sub> H <sub>30</sub>	270	Dehydroabietane	74
19	18.893	C <sub>20</sub> H <sub>28</sub>	268	7-butyl-1-hexylnaphthalene	73
20	20.263	C <sub>19</sub> H <sub>24</sub>	252	Simonellite	71
21	24.1	C <sub>18</sub> H <sub>18</sub>	234	Retene	99

<https://doi.org/10.1371/journal.pone.0275842.t001>

could be detected after the anaerobic degradation of the methanol extract of highly volatile bituminous coal. However, only H<sub>2</sub> and CO<sub>2</sub> were detected in the M1 group, which contained organic matter extracted from coal in addition to methanol. It was thought that the extracted organic matter was the substrate of the fermentation bacteria and hydrogen-producing acetogens in the flora. The substrate promoted the production of more H<sub>2</sub>, but it did not play a corresponding role in the methanogenic process in the system.

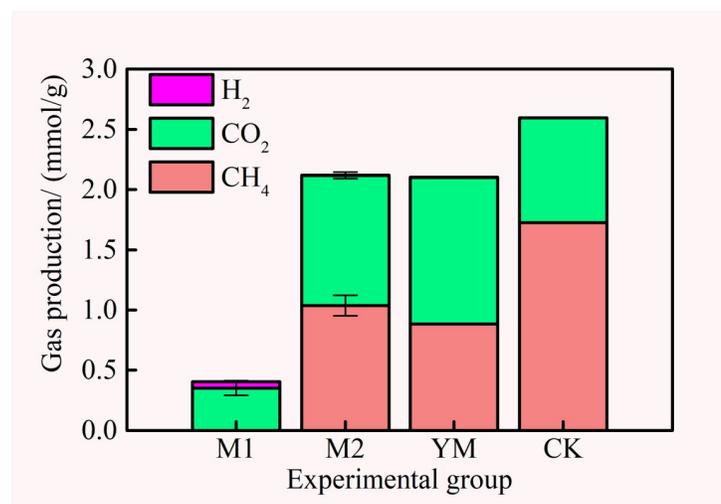


Fig 2. Biogenic methane production of each experimental group.

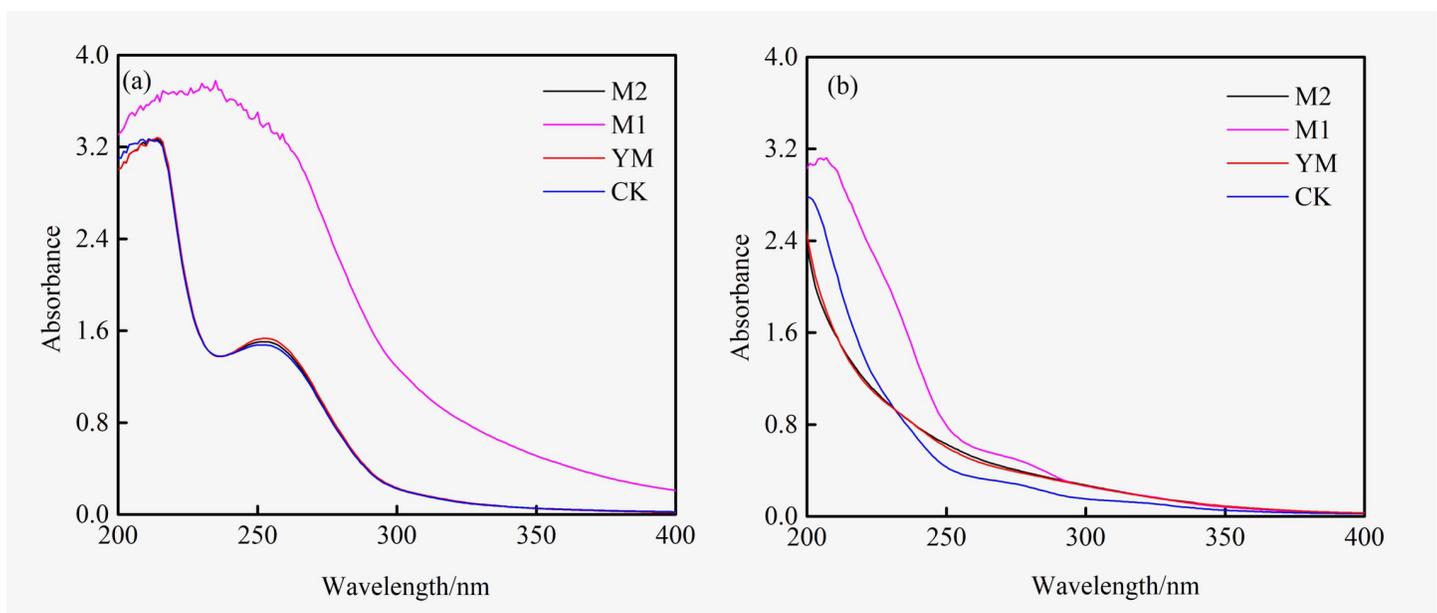
<https://doi.org/10.1371/journal.pone.0275842.g002>

Methane was produced by acetoclastic, methylotrophic, and hydrogenotrophic methanogens [2]. The largest abundance of methanogens in the original flora in this study was *Methanoculleus* [3], the most reported methanogenic microorganism in the literature. It can produce methane using small molecules such as  $H_2/CO_2$  and formate. Theoretically, the M1 group should be able to produce methane. However, based on the current results, it was likely that the organic matter in the extract or the intermediate metabolites produced by the organic matter had a negative impact on the metabolic process of the methanogens. The specific reason still needed to be further elaborated on.

In contrast, after methanol extraction, the methane production of the M2 group reached 1.03 mmol/g and increased 18% compared with the YM group. It was speculated that methanol extraction increased the contact area between coal and microorganisms [26, 27].

### UV-vis analysis of organic matter in the biogas production system of lignite methanol extract and residual coal

The results of the UV-vis analysis of fermentation broth before and after biogas production are exhibited in Fig 3. Before the gas production, the M1 group fermentation broth showed a strong continuous absorption (peak at 225 nm–275 nm), which was generated by the conjugation of the chromophore's double bond with the benzene ring in the system [28]. Combined with the GC-MS analysis results of the extract, the absorption peak primarily originated from aromatic ketones and esters. The M2, YM, and CK groups had shoulder peaks at 220 nm and 254 nm, predominantly generated by the absorption of organic matter in the medium. After biogas generation, the absorption of the M1 group at 225–275 nm was significantly weakened. This result suggested that organic compounds such as aromatic ketones and esters in the extract were degraded. The M1 group had shoulder peaks near 220 nm and 280 nm, primarily from aromatic compounds [29]. However, the M2 group and YM group had no shoulder peaks in this wavelength range, which indicated that the products of the M1 group contained more aromatic compounds.



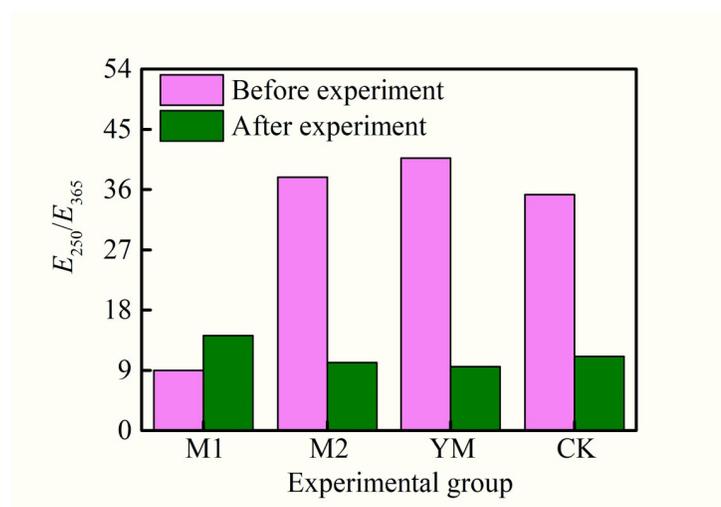
**Fig 3. UV-vis analysis of fermentation liquid before and after biogenic methane production.**

<https://doi.org/10.1371/journal.pone.0275842.g003>

The ratio of the absorbance of the fermentation broth at 250 nm and 365 nm ( $E_{250}/E_{365}$ ) was applied to characterize the molecular weight of soluble organic matter. The larger the ratio, the smaller the molecular weight of the organic matter is [29, 30]. The changes of  $E_{250}/E_{365}$  before and after gas production in each experimental group are displayed in Fig 4. It can be observed that the fermentation broth of the M1 group contained a large amount of macromolecular organic matter before biogas production, and its  $E_{250}/E_{365}$  was substantially smaller than that of other experimental groups.  $E_{250}/E_{365}$  of the M1 group increased slightly after gas production, demonstrating that some organic substances were biodegraded and the molecular weight became smaller. While,  $E_{250}/E_{365}$  of the M2 and YM groups decreased after gas production, which may be due to the release of macromolecular substances in the coal under the biological action [29].

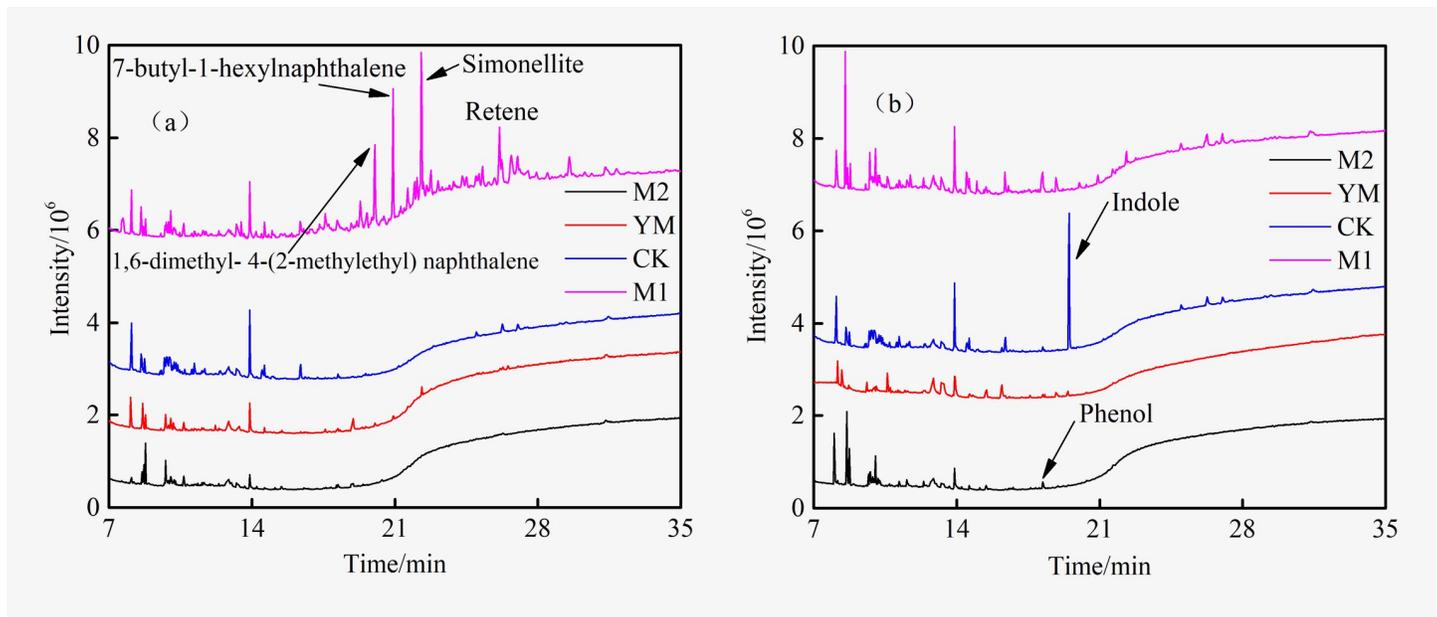
### Composition and change of organic matter in the biogas production system of lignite methanol extract and residual coal

The GC-MS total ion current chromatogram of the fermentation broth of each experimental group before and after gas production is shown in Fig 5. Before gas production, eight compounds were detected in the samples of the M1 group. They included trans-1,2,3,4,4a,5,8,8a-octahydro-4a-methylnaphthalene (No.5 in Table 1), 3,5-bis (1,1-dimethylethyl)-phenol (No. 6 in Table 1), 1-(1,3-dimethyl-1H-indol-2-yl) ethanone (No. 11 in Table 1), methyl hexadecanoate (No. 13 in Table 1), 1,6-dimethyl-4-(2-methylethyl) naphthalene (No. 17 in Table 1), 7-butyl-1-hexyl Naphthalene (No. 19 in Table 1), simonellite (No. 20 in Table 1), and retene (No. 21 in Table 1). No compounds were detected in the remaining experimental groups. After gas generation, the height of the main peaks in the chromatograms of the M1 group samples decreased significantly. The peaks of 1,6-dimethyl- 4-(2-methylethyl) naphthalene (No. 17 in Table 1), 7-butyl-1-hexylnaphthalene (No. 19 in Table 1), simonellite (No. 20 in Table 1), and retene (No. 21 in Table 1) disappeared. This result demonstrated that these compounds could be degraded and utilized by the flora. Different studies have reported that microorganisms of the *Firmicutes* and *Proteobacteria* can degrade polycyclic aromatic hydrocarbons [31, 32], and the microorganisms of the *Bacteroidetes* can degrade phenols [33–35]. These microorganisms in the inoculum might be involved in the degradation of organic matter in the extract.



**Fig 4. The changes of  $E_{250}/E_{365}$  before and after biogenic methane production.**

<https://doi.org/10.1371/journal.pone.0275842.g004>



**Fig 5.** TIC of GC-MS of experimental groups before (a) and after (b) biogenic methane production.

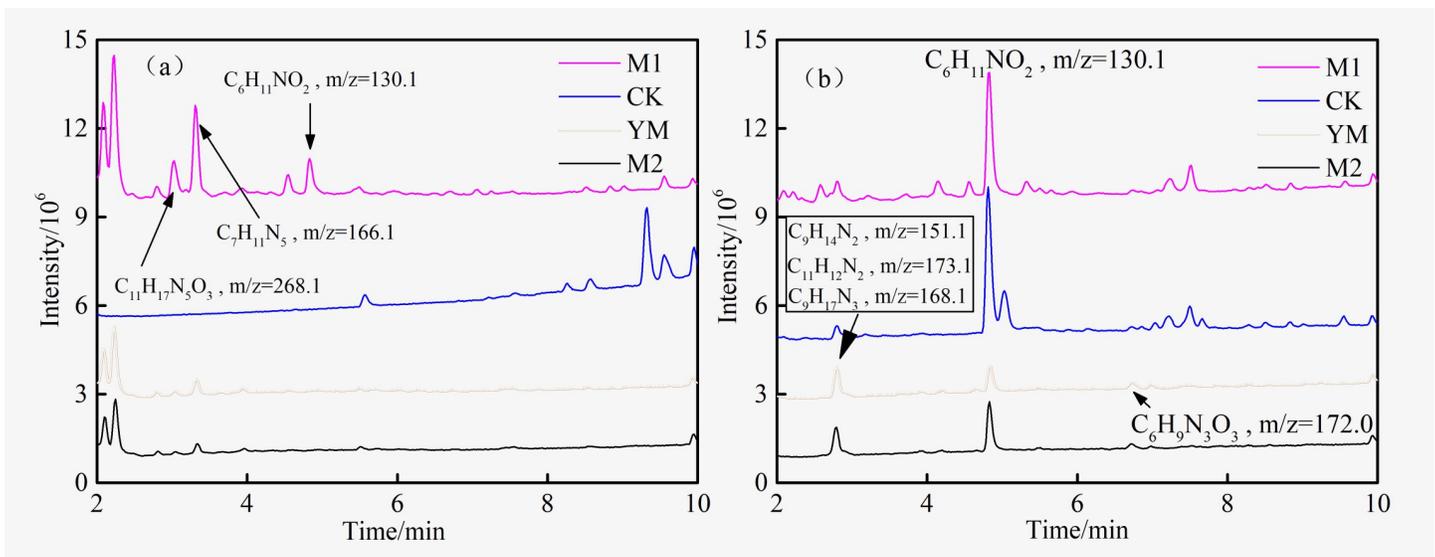
<https://doi.org/10.1371/journal.pone.0275842.g005>

A large amount of indole which was a metabolite produced by the use of methanol by the flora was detected in the fermentation broth after gas production in the CK group (Fig 5B). However, indole was not detected in the M1 group. It was speculated that some compounds might inhibit the production of indole.

Phenolic organic compounds could be connected with the degradation of lignin [36]. Phenol was found in the M1, M2, and YM groups after the experiment (Fig 5B), indicating the decomposition of polymers in lignite [37]. Moreover, these authors identified over 100 compounds from kerogen due to the decomposition of polymers, covering the compounds obtained in this study. Nevertheless, no more metabolites except phenol were detected in all groups. The possible reason was that the produced metabolites were polar and difficult to be detected using the GC-MS method. Therefore, HPLC-MS was employed for further analysis of the fermentation broth.

The HPLC-MS total ion current chromatogram of the fermentation broth of each experimental group was exhibited in Fig 6. For samples of the M1 group after the experiment, the peaks of  $C_{11}H_{17}N_5O_3$  (at 3.0 min),  $C_7H_{11}N_5$  (at 3.3 min), and  $C_{13}H_{11}NO_3$  (at 3.9 min) disappeared (Fig 6A), indicating that these compounds were degraded by the flora. Moreover, the abundances of  $C_9H_{14}N_2$  (at 2.8 min),  $C_{11}H_{12}N_2$  (at 2.8 min),  $C_9H_{17}N_3$  (at 2.8 min),  $C_6H_{11}NO_2$  (at 4.8 min), and  $C_6H_9N_3O_3$  (at 6.7 min) increased significantly after the experiment (Fig 6B), which proved that these compounds were produced during the degradation of the extract. At the same time, YM and M2 groups also showed similar changes. Except for  $C_6H_{11}NO_2$ , the number of rings plus double bonds of other compounds was more than 4. Combined with the results of UV analysis (Fig 3B), these products may be aromatic compounds.

The degradation of the extract might lead to the rapid accumulation of some metabolites, which may have a negative effect on the methane production process. Especially,  $C_6H_{11}NO_2$  existed in the extract and accumulated significantly after biological action in all groups. In another study, the addition of excess pulverized coal resulted in the inhibition of methanogenesis and the accumulation of  $C_6H_{11}NO_2$ , which also showed that  $C_6H_{11}NO_2$  was involved in

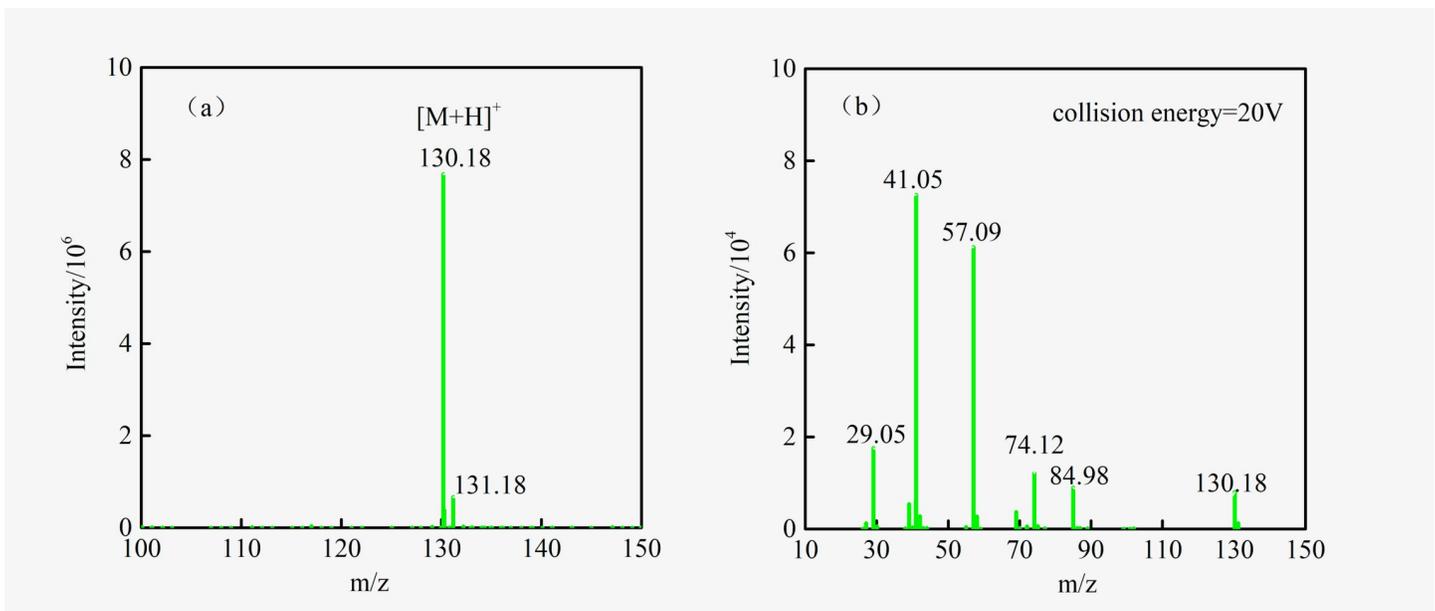


**Fig 6.** TIC of HPLC-MS of experimental groups before (a) and after (b) biogenic methane production.

<https://doi.org/10.1371/journal.pone.0275842.g006>

inhibiting the methanogenesis process [38]. Interestingly, it was not released from coal during the sterilization, which indicated that it was firmly bound to the macromolecular structure. Considering the potential effect of  $C_6H_{11}NO_2$  on the methanogenesis process, its existence in coal also provided insight into the complexity of the process of microbial action.

The number of rings plus double bonds of  $C_6H_{11}NO_2$  is 2, indicating that it is not an aromatic compound. The MS and MS/MS spectra of  $C_6H_{11}NO_2$  are shown in Fig 7, and their specific structure needs to be further identified.



**Fig 7.** The MS(a) and MS/MS(b) spectrum of  $C_6H_{11}NO_2$ .

<https://doi.org/10.1371/journal.pone.0275842.g007>

## Conclusions

Research on bioavailable organic matter in coal and its metabolites is an important part of elucidating the mechanism of coal biogas formation. In this study, many alcohols, esters, carboxylic acids, amides, phenols, aromatic hydrocarbons, and heterocyclic compounds were detected in the methanol extract of Yima lignite, with aromatic compounds showing the most considerable abundance. The methanol extract of Yima lignite was used by the microbial flora to produce H<sub>2</sub> and CO<sub>2</sub> without accumulation of CH<sub>4</sub>. Both Aromatic hydrocarbons and other oxygen-containing compounds in the extract were biodegraded, and their molecular weights decreased. Certain soluble organic compounds in the extract and metabolites of the biodegradation process may negatively affect the methanogenesis process, where the compound C<sub>6</sub>H<sub>11</sub>NO<sub>2</sub> present in both the extract and the product may be the primary inhibitor. The methane production of residual coal increased by 18% compared with the raw coal. In addition, both aromatic and non-aromatic compounds were produced during the biodegradation process, and the appearance of phenol in the product indicated the depolymerization of lignin in coal. The current study demonstrated the complex role of soluble organic matter and its metabolites during the biogasification process.

## Author Contributions

**Conceptualization:** Jianmin Liu, Hengxing Ren.

**Data curation:** Hengxing Ren, Linyong Chen.

**Funding acquisition:** Jianmin Liu, Yi Jin, Guofu Li, Baoyu Wang.

**Methodology:** Jianmin Liu, Hengxing Ren.

**Validation:** Jianmin Liu, Hengxing Ren, Linyong Chen.

**Writing – original draft:** Hengxing Ren.

**Writing – review & editing:** Jianmin Liu, Yi Jin, Huan He.

## References

1. Ritter D, Vinson D, Barnhart E, Akob DM, Fields MW, Cunningham AB, et al. Enhanced microbial coalbed methane generation: A review of research, commercial activity, and remaining challenges. *Int J Coal Geol.* 2015; 146: 28–41. <https://doi.org/10.1016/j.coal.2015.04.013>
2. Zhang D, He H, Ren Y, Haider R, Urynowicz M, Fallgren PH, et al. A mini review on biotransformation of coal to methane by enhancement of chemical pretreatment. *Fuel.* 2022; 308: 121961. <https://doi.org/10.1016/j.fuel.2021.121961>
3. Zhao H, He H, Wang JZ, Tan KL, Zhao N, Ren HX, et al. Variation of microbial community before and after biogas production with Shengli lignite in Inner Mongolia. *J China Coal Soc.* 2019; 44: 1224–1231. <https://doi.org/10.3969/j.issn.1001-1986.2020.06.019>
4. Pytlak A, Szafranek-Nakonieczna A, Sujak A, Grządziel J, Polakowski C, Kuźniar A, et al. Stimulation of methanogenesis in bituminous coal from the upper Silesian coal basin. *Int J Coal Geol.* 2020; 231: 103609. <https://doi.org/10.1016/j.coal.2020.103609>
5. Malik AY, Ali MI, Jamal A, Farooq U, Khatoun N, Orem WH, et al. Coal biomethanation potential of various ranks from Pakistan: A possible alternative energy source. *J Clean Prod.* 2020; 255: 120177. <https://doi.org/10.1016/j.jclepro.2020.120177>
6. Sharifianjazi F, Esmaeilkhani A, Bazli L, Eskandarinezhad S, Khaksar S, Shafiee P, et al. A review on recent advances in dry reforming of methane over Ni- and Co-based nanocatalysts. *Int J Hydrogen Energ.* 2021; In Press. <https://doi.org/10.1016/j.ijhydene.2021.11.172>
7. Yusuf M, Farooqi AS, Keong LK, Hellgardt K, Abdullah B. Contemporary trends in composite Ni-based catalysts for CO<sub>2</sub> reforming of methane. *Chem Eng Sci.* 2021; 229: 116072. <https://doi.org/10.1016/j.ces.2020.116072>

8. Fuertez J, Córdoba G, McLennan JD, Adams DJ, Sparks TD. Potential application of developed methanogenic microbial consortia for coal biogasification. *Int J Coal Geol.* 2018; 188: 165–180. <https://doi.org/10.1016/j.coal.2018.02.013>
9. Wang A, Shao P, Lan F, Jin H. Organic chemicals in coal available to microbes to produce biogenic coalbed methane: A review of current knowledge. *J Nat Gas Sci Eng.* 2018; 60: 40–48. <https://doi.org/10.1016/j.jngse.2018.09.025>
10. Su X, Zhao W, Xia D, Hou S, Fu H, Zhou Y. Experimental study of advantages of coalbed gas bioengineering. *J Nat Gas Sci Eng.* 2022; 102: 104585. <https://doi.org/10.1016/j.jngse.2022.104585>
11. Liu JM, Wang BY, Tai C, Wu L, Zhao H, Guan J, et al. An effective method to detect volatile intermediates generated in the bioconversion of coal to methane by gas chromatography-mass spectrometry after in-situ extraction using headspace solid-phase micro-extraction under strict anaerobic conditions. *Plos One.* 2016; 11: e0163949. <https://doi.org/10.1371/journal.pone.0163949> PMID: 27695055
12. Pinto JSS, Assis LM, Andrade MA, Lancas FM. Pressurized liquid extraction of brazilian coal followed by the extracts characterization by gas chromatography coupled to mass spectrometry. *J Chromatogr Sci.* 2018; 56(8): 761–769. <https://doi.org/10.1093/chromsci/bmy051> PMID: 29931139
13. Wang BY, Tai C, Wu L, Chen LY, Liu JM, Hu B, et al. Methane production from lignite through the combined effects of exogenous aerobic and anaerobic microflora. *Int J Coal Geol.* 2017; 173: 84–93. <https://doi.org/10.1016/j.coal.2017.02.012>
14. Furmann A, Schimmelmann A, Brassell SC, Mastalerz M, Picardal F. Chemical compound classes supporting microbial methanogenesis in coal. *Chem Geol.* 2013; 339: 226–241. <https://doi.org/10.1016/j.chemgeo.2012.08.010>
15. Tao MX, Wang WC, Li ZP, Ma YZ, Li J, Li XB. Comprehensive study on genetic pathways and parent materials of secondary biogenic gas in coalbeds. *Chin Sci Bull.* 2014; 59: 970–978. <https://doi.org/10.1007/s11434-014-0151-7>
16. Zhang XY, Wang F, Li GS, Fan X, Yu YR, Zhao YP, et al. Cluster Analysis of Molecular Characteristics for Soluble Organic Matter in Coals. *Chinese J Anal Chem.* 2019; 47(1): 99–105. <https://doi.org/10.19756/j.issn.0253-3820.181585>
17. Ge XG, Cheng JM, Yang L, Ye YK, Chen LW. An experimental study of *Desulfobrio* sp. anaerobic degradation of >C<sub>12</sub> organic compounds of Huainan coal. *Front Earth Sci-PRC.* 2015; 22(1): 328–334. <https://doi.org/10.13745/j.esf.2015.01.028>
18. Chen LY, Wang BY, Tai C, Guan JD, Zhao H, Wang ML, et al. Composition and conversion of intermediate products in the process of anthracite gasification by microorganism. *J China Coal Soc.* 2016; 41: 2305–2311. <https://doi.org/10.13225/j.cnki.jccs.2016.0318>
19. Haider R, Ghauri MA, SanFilippo JR, Jones EJ, Orem WH, Tatu CA, et al. Fungal degradation of coal as a pretreatment for methane production. *Fuel.* 2013; 104: 717–25. <https://doi.org/10.1016/j.fuel.2012.05.015>
20. Chen T, Rodrigues S, Golding SD, Rudolph V. Improving coal bioavailability for biogenic methane production via hydrogen peroxide oxidation. *Int J Coal Geol.* 2018; 195: 402–414. <https://doi.org/10.1016/j.coal.2018.06.011>
21. Liu F, Guo H, Wang Q, Haider R, Urynowicz MA, Fallgren PH, et al. Characterization of organic compounds from hydrogen peroxide-treated subbituminous coal and their composition changes during microbial methanogenesis. *Fuel.* 2019; 237: 1209–1216. <https://doi.org/10.1016/j.fuel.2018.10.043>
22. Campbell BC, Gong S, Greenfield P, Midgley DJ, Paulsen IT, George SC. Aromatic compound-degrading taxa in an anoxic coal seam microbiome from the Surat Basin, Australia. *FEMS Microbiol Ecol.* 2021; 97(5): fiab053. <https://doi.org/10.1093/femsec/fiab053>
23. Lu Z, Li Q, Ju Y, Gu S, Xia P, Gao W, et al. Biodegradation of coal organic matter associated with the generation of secondary biogenic gas in the Huaibei Coalfield. *Fuel.* 2022; 323: 124281. <https://doi.org/10.1016/j.fuel.2022.124281>
24. Vick SHW, Gong S, Sestak S, Vergara TJ, Pinetown KL, Li Z, et al. Who eats what? Unravelling microbial conversion of coal to methane. *FEMS Microbiol Ecol.* 2019; 95(7): fiz093. <https://doi.org/10.1093/femsec/fiz093>
25. Wang BY, Chen LY, Tai C, Han ZY, Guan JD, Zhao H. A preliminary study of biological coal gasification by exogenous bacteria: Microbiome composition coal type pore and seam fracture. *J China Coal Soc.* 2014; 39(09): 1797–801. <https://doi.org/10.13225/j.cnki.jccs.2014.8009>
26. Hao ZC, Zhang XD, Yang YH, Yang YL, Yu KK. Study on reconstruction mechanism of different solvent to surface and porosity of high rank coal. *Coal Sci Technol.* 2018; 46(3): 108–114. <https://doi.org/10.13199/j.cnki.cst.2018.03.018>
27. Zhang P, Guo H, Duan K, Chen C, Liang W. Effects of microbial anaerobic metabolites on nanoporosity of anthracite. *J China Coal Soc.* 2020; 45(11): 3841–3852. <https://doi.org/10.13225/j.cnki.jccs.2019.1263>

28. Shi KY, Yin SD, Tao XX, Du Y, He H, Lv ZP, et al. Quantitative measurement of coal bio-solubilization by ultraviolet-visible spectroscopy. *Energ Source Part A*. 2013; 35(15): 1456–1462. <https://doi.org/10.1080/15567036.2010.521800>
29. Zhao TQ, Li P, Tai C, Xiang Y, Li HY, Ma YX, et al. Spectral characteristics of dissolved organic matter in the process of coal gasification by microorganism. *J China Coal Soc*. 2017; 42(Suppl. 2): 525–534. <https://doi.org/10.13225/j.cnki.jccs.2017.1035>
30. Mo L, Wang Q, Bi E. Effects of endogenous and exogenous dissolved organic matter on sorption behaviors of bisphenol a onto soils. *J Environ Manage*. 2021; 287: 112312. <https://doi.org/10.1016/j.jenvman.2021.112312> PMID: 33711663
31. Bacosa HP, Inoue C. Polycyclic aromatic hydrocarbons (PAHs) biodegradation potential and diversity of microbial consortia enriched from tsunami sediments in Miyagi, Japan *J Hazard Mater*. 2015; 283: 689–697. <https://doi.org/10.1016/j.jhazmat.2014.09.068> PMID: 25464311
32. Wu M, Guo X, Wu J, Chen K. Effect of compost amendment and bioaugmentation on PAH degradation and microbial community shifting in petroleum-contaminated soil. *Chemosphere*. 2020; 256: 126998. <https://doi.org/10.1016/j.chemosphere.2020.126998> PMID: 32470727
33. Viggor S, Jöesaar M, Soares-Castro P, Ilmjärvi T, Santos PM, Kapley A, et al. Microbial metabolic potential of phenol degradation in wastewater treatment plant of crude oil refinery: Analysis of metagenomes and characterization of isolates. *Microorganisms*. 2020; 5: 652. <https://doi.org/10.3390/microorganisms8050652> PMID: 32365784
34. Li Y, Li J, Wang D, Wang G, Yue X, Kong X, et al. Denitrifying microbial community structure and bama gene diversity of phenol degraders in soil contaminated from the coking process. *Appl Biochem Biotech*. 2020; 190(3): 966–981. <https://doi.org/10.1007/s12010-019-03144-5> PMID: 31637583
35. Guo H, Zhang Y, Huang Z, Liang W, Urynowicz M, Ali MI. High potential of methane production from coal by fungi and hydrogenotrophic methanogens from produced water. *Energ Fuel*. 2020; 34(9): 10958–10967. <https://doi.org/10.1021/acs.energyfuels.0c02265>
36. Bucha M, Jędrysek M, Kufka D, Pleśniak Ł, Marynowski L, Kubiak K, et al. Methanogenic fermentation of lignite with carbon-bearing additives, inferred from stable carbon and hydrogen isotopes. *Int J Coal Geol*. 2018; 186: 65–79. <https://doi.org/10.1016/j.coal.2017.11.020>
37. Detman A, Bucha M, Simoneit BRT, Mielecki D, Piwowarczyk C, Chojnacka A, et al. Lignite biodegradation under conditions of acidic molasses fermentation. *Int J Coal Geol*. 2018; 196: 274–287. <https://doi.org/10.1016/j.coal.2018.07.015>
38. Chen LY, Li GF, Liu JM, Ren HX, He H, Miao B, et al. Inhibitory effects of excessive pulverized coal on biogenic methane generation from coal. *Coal Conv*. 2021; 44(6): 8–15. <https://doi.org/10.19726/j.cnki.ebcc.202106002>