

Enhancement of biogenic methane production by co-degradation of coal and straw: microbial and organic analysis

Hongguang Guo¹, Yatong Cheng¹, Kaixin Duan¹, Weiguo Liang¹, Zaixing Huang², Michael Urynowicz³, and Muhammad Ali⁴

¹Taiyuan University of Technology

²China University of Mining and Technology

³University of Wyoming

⁴Quaid-i-Azam University

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Abstract

Co-degradation of coal and straw could produce significantly higher methane which was potential to increase biogenic CBM. In this study, the success of microflora and organic compounds during co-degradation was determined by MiSeq and GC-MS, and compared with cultivations with only coal (C) and with only straw (RS). The results showed that the methane production in co-degradation was 12 times higher than that in cultivation C. A shift of dominant methanogen was caused by the addition of straw from acetoclastic *Methanosaeta* in inoculum to methylotrophic *Methanomethylovorans* in 7 days, then hydrogenotrophic *Methanobacterium*. The bacteria and fungi with ability to degrade macromolecules in coal and metabolize VFAs were enriched which would facilitate methanogenesis. VFAs, especially butanoic acid, were dominant in intermediates of co-degradation which contributed to methane production as their content were negatively correlated with methane production. The different component of intermediates and microbial communities among co-degradation, cultivations C and RS suggested that the metabolic pathway in co-degradation was distinctive and the fracture of coal molecules was almost completed in the first 7 days of cultivation. Coal might also serve as the suitable microhabitat for microorganisms to avoid the threat from environment in addition to function as methanogenic substrates.

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