

Producing methane from coal through biogasification - identifying suitable nutrient solutions for enhancing methane yield

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Abstract

As of 2011, Coalbed Methane (CBM) released through biogasification, contributes ~40% of the total CBM production in the US. This is largely owing to the significant production of biogenic CBM from low rank coal seams in western states of the US. Biogenic CBM in the Illinois basin which has the largest deposit of bituminous coal in the US, however, has not been explored extensively although biogenic methane has been observed at different locations. To enhance methane yield from higher ranked coals, we aimed to stimulate activities of indigenous microorganisms that can convert coal to methane in a collaborative and synergistic way. To achieve this goal, different nutrient solutions have been tested. The top two nutrient media which brought the highest methane production were then investigated on whether they can be simplified for decreasing the cost. In addition, effects from other parameters, such as coal loading, pH, temperature, coal particle size, shaking, inoculum loading, surfactants and solvents have been screened and evaluated. Important parameters were further studied to identify their optimal values. Results from this study can be applied to convert mined out coal and waste coal to methane. Additionally, the developed strategies can be used for enhancing methane production in unminable or abandoned coal seams.

Overall Methodology

Simplify nutrient solution #2 by eliminating expensive components





Two nutrient recipes and optimal conditions for both ex situ and in situ applications

Microbial community analysis



Fig. 4: Yield of gas on bituminous coal under different conditions. A: Methane, B: CO₂.

Screen significant factors for enhancing methane yield using solution #2



Fig. 5: Half normal probability plot indicating positive or negative effects on methane yield.

Analysis of variance table [Part	ial sum of squares - Type]				
	Sum of		Mean	F	p-value	
Source	Squares	df	Square	Value	Prob > F	
Model	70.92	26	2.73	9.46	< 0.0001	signific
A-Particle size	3.52	1	3.52	12.2	0.0013	
B-Temperature	4.47	1	4.47	15.5	0.0004	
C-Inoculum size	0.2	1	0.2	0.7	0.4092	
D-Solid loading	1.05	1	1.05	3.63	0.0646	
E-pH	1.49	1	1.49	5.18	0.0287	
F-Mixing	0.87	1	0.87	3	0.0914	
G-Coenzyme M	0.67	1	0.67	2.32	0.1361	
H-Triton X-100	4.07E-03	1	4.07E-03	0.014	0.906	
J-SDS	7.16	1	7.16	24.84	< 0.0001	
K-Ethanol	22.77	1	22.77	79	< 0.0001	
L-2-Propanol	7.03E-03	1	7.03E-03	0.024	0.8767	
M-Sodium formate	0.07	1	0.07	0.24	0.6261	
AC	0.39	1	0.39	1.34	0.2549	
AD	4.79	1	4.79	16.61	0.0002	
AH	2.11	1	2.11	7.33	0.0102	
AM	0.39	1	0.39	1.35	0.2536	
BF	3.28	1	3.28	11.39	0.0017	
СН	0.27	1	0.27	0.93	0.3409	
DE	2.59	1	2.59	8.99	0.0048	
DJ	3.67	1	3.67	12.73	0.001	
DK	3.52	1	3.52	12.21	0.0013	
EG	2.13	1	2.13	7.4	0.0099	
EK	0.56	1	0.56	1.96	0.17	
FL	1.48	1	1.48	5.13	0.0294	
GM	1.23	1	1.23	4.28	0.0455	
ACH	2.23	1	2.23	7.73	0.0085	
Residual	10.67	37	0.29			
Cor Total	81 59	63				

The Model F-value of 9.46 implies the model is significant. There is only a 0.01% chance that a "Model F-Value" this large could occur due to noise

Replace expensive components in #2 solution with other chemicals





Meniscus spp.	Syntrophus sp.	Ruminococcus spp.	Methanobacteriaceae methanobacterium spgh	Methanobacterium subterraneum
Dasania spp.	Candidatus_solibacter spp.	Crocinitomix spp.	Methanococcus maripaludis	Methanocalculus pumilus
 Desulfofustis spp. Desulfuromusa spp. 	 Thermovirga spp. Maritimimonas spp. 	 Sulfurimonas spp. Alkaliflavus spp. 	Methanobacteriaceae spp.	Methanobacterium spp.
 Desulfuromonas spp. 	 Desulfotignum spp. 	Others	Methanobacterium ferruginis	■ Others

Fig. 2: Distribution of bacteria (A) and archaea (B) in the formation water.

Methane production with or without nutrient solutions







Fig. 6: Effects of corn steep liquor (CSL) and trypticase soy broth (TSB) on methane content (A). Methane yield from bituminous coal incubated with different strength of TSB (B).

Screen significant factors for enhancing methane yield using TSB



Fig. 7. Half normal probability plot demonstrating either positive and negative effect of different parameters (A). Methane yield predicted vs. actual (B).

Table 2: ANOVA results reflecting significant parameters.

	Sum of		Mean	F	p-value
Source	Squares	df	Square	Value	Prob > F
Model	14.97	5	2.99	1059.78	0.0009
significant					
A-temperatu	re 2.72	1	2.72	963.31	0.0010
B-TSB	5.83	1	5.83	2065.49	0.0005
C-Shaking	0.58	1	0.58	206.07	0.0048
D-Coal loadi	ng4.69	1	4.69	1660.54	0.0006
AC	1.14	1	1.14	403.51	0.0025
Residual	5.649E-003	2	2.825E-003		
Cor Total	14 97	7			

146, 145-154.

Table 3: Developed model for predicting methane yield on coal.

Final Equation in Terms of Actual Factors:

Ln(Methane yie	ld) =
+10.28805	
-0.16011	* temperature
+0.037955	* TSB
-0.037384	* Shaking
-0.061257	* Coal loading
+1.25820E-003	* temperature * Shakin

Summary and Conclusions

- Illinois bituminous coal can be converted to methane by a microbial community indigenous to a CBM production site.
- Nutrient addition significantly enhances methane content and yield from coal.
- Two nutrient solutions which have similar costs are developed for biogasifying bituminous coal. 3.

Fig. 3: Gas release from bituminous coal over time. A:methane, B: CO₂

Optimal parameters for improving coal biogasification are identified. 4.

5. The highest methane production rate is between 27.6 and 32.6 ft^3 /ton-day.



Zhang, J.*, Liang, Y.-N., Yau, P.M., Pandey, R.*, Harpalani, S. 2015. A metaproteomic approach for identifying proteins in anaerobic bioreactors converting coal to methane. International Journal of Coal Geology. 146, 91–103. Zhang, J.*, Liang, Y.-N, Pandey, R.*, Harpalani, S. 2015. Characterizing a microbial community dedicated for converting coal to methane in situ and ex situ. International Journal of Coal Geology.