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Addition of soluble bio-based substances selected for Archaeal communities with higher performances in thermophilic Anaerobic digestion processes.

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In a previous work compost derived- soluble bioorganic substances (CVD-SBO) were successfully used to decrease the ammonia level in the final digestate produced from OFMSW (Francavilla et al. 2016) together with an increase of methane yield.

In this work we evaluated of the Bacteria and Archaea microbial community structure and dynamics in a thermophilic anaerobic process treating OFMSW, according to the addition of a CVD-SBO at different concentration.





Results and Discussion

Materials and Methods

A lab-scale system running in parallel six 6.5L batch bioreactors was used for the investigation. Bioreactors were fed with fermentation slurry sampled from an industrial plant treating OFMSW. Duplicate runs were performed for the control slurry mix containing no added CVD-SBO, and for the same mix added at 0.05% and 0.20% of CVD-SBO, starting the reactors at the same time by heating up to 55°C for 15 days. Microbial analyses were conducted by PCR-DGGE targeting 16SrRNA of Bacterial and Archaeal communities and by sequencing most representative DGGE bands.

Biogas production increased by 5% when 0.2% CVD was added. Moreover, the methane/ CO_2 ratio increased from 2.8 to 3.3. (figures 1-2, table 1). The increase was related to the maximum specific methane production potential (Table 2).

By cluster analysis of PCR-DGGE fingerprinting we found that bacterial communities were not affected, while Archaeal communities clearly clustered according to the addition of CVD to the bioreactor (Figure 3). Generally, under the experimental conditions, archaeal diversity is more affected, with a reduction of abundance and evenness, consistently with increased CVD concentration (table 3).

By analyzing the DNA sequence of the most abundant bacterial and archaeal DGGE bands, we found a general selection of thermophilic bacteria typical of thermophilic AD and a selective effect of CVD towards the increase of acetoclastic *Methanosarcina* and decrease of hydrogenothrophic *Methanoculleus*. (Figure 4)



	Biogas (NI)	CH ₄ (NI)	CH_4/CO_2 ratio	
CVD 0.2	73.20±0.50ª	53.60±0.51ª	3.33±0,086ª	
CVD 0.05	70.50±0.45 ^b	50.20±0.67 ^b	2.90±0.075 ^b	
Control	70.25±0.30 ^b	49.17±0.70 ^b	2.81±0.072 ^b	
Significance	*	**	* *	

Table 1: Analysis of variance (ANOVA) of the main biogas production parameters. Means followed by the same letters in each colon are not significantly different at P \leq 0.05 according to Tukey's test. *, F test significant at P<0.05; **, F test significant at P<0.01.

Н 52.72±2.70^a CVD 0.2 1.43 ± 0.03 2.94±0.60 48.45±2.45^b **CVD 0.05** 1.32 ± 0.21 2.59±0.15 49.10±1.10^b 1.35±0.24 2.47±0.25 Control Significance ns ns

Table 2 - ANOVA results related to coefficients of Gompertz equation for CH_4/CO_2 ratio.

0.00 0.00 CVD_f_0.2_a CVD_f_0.2_b C_b CVD_i_0.2_a CVD_i_0.05_a CVD_i_0.05_b CVD_f_0.05_a CVD_f_0.05_b CVD <u>i</u> 0.05_a CVD <u>i</u> 0.05_b CVD <u>f</u> 0.2_a CVD <u>f</u> 0.2_b C_f_b C_f_a VD_i_0.2_b Cia CVD_i_0.2_b Cia C_i b CVD_i_0.2_a :VD_f_0.05_b VD_f_0.05_a C_f_a C_f_b

Figure 4. Cluster analysis of PCR-DGGE fingerprint of the bacterial and archaeal communities, based on the Bray-Curtis dissimilarity.

DIVERSITY INDICES		Initial (t _o)			Final (t _f)		
		Richness	Shannon	Pielou's	Richness	Shannon	Pielou's
Bacteria	Control	18 (0) ^a	2.73 (0.006)	0.95 (0.002)	13 (0)	2.29 (0.004)	0.89 (0.002)
	CVD 0.05	19 (0)	2.72 (0.048)	0.92 (0.017)	13 (0)	2.30 (0.004)	0.90 (0.001)
	CVD 0.2	17 (0)	2.68 (0.048)	0.95 (0.017)	13 (0)	2.35 (0.053)	0.92 (0.021)
Archaea	Control	28.5 (0.7)	3.07 (0.047)	0.92 (0.007)	17 (0)	2.59 (0.003)	0.92 (0.001)
	CVD 0.05	20 (0)	2.59 (0.016)	0.87 (0.005)	15 (0)	2.54 (0)	0.94 (0)
	CVD 0.2	20.5 (0.7)	2.63 (0.043)	0.87 (0.024)	14 (0)	2.40 (0.023)	0.91 (0.008)

Table 3. Diversity indices estimated by the bacterial and archaeal DGGE profiles. ^aAverage of diversity indexes (standard deviation).

Conclusion

The effect of CVD on the reduction of archaeal diversity might be related to the higher biogas productivity of the reactor with 0.2% CVD added. Possible positive effect of CVD on the selection of the most active methanogenic taxa, triggering the production of methane. Further studies would allow an evaluation on the impact of N-cycle related microorganisms during the DA process.