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Evaluation of the effect of glycerol supplementation on the anaerobic digestion of real municipal solid waste in batch mode



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ABSTRACT

The effect of low glycerol supplementation (<1% v/v) in the thermophilic anaerobic digestion (AD) of real municipal solid waste (RMSW) on methane production (MP) is not known. This study explores the effect of five different glycerol supplementations (0%, 0.1%, 0.25%, 0.5% and 1%) on effluent characteristics, anaerobic consortia and MP. Specifically, by adding 0.25% v/v crude glycerol to the feed, the methane production rate increased by 48% (from 7.40 ± 1.17 l CH₄/l to 11.01 ± 1.66 l CH₄/l), in line with the increase in total volatile solids (TVS) removal (from 65 ± 7% to 81 ± 7) and methanogenic activity (from $110 \times 10^{-12} \pm 17 \times 10^{-12} l$ CH₄/cell to 156 \times 10^{-12} \pm 24 \times 10^{-12} l CH₄/cell). Extra glycerol added to the feed (0.5% or 1%) was shown to be non-feasible for thermophilic AD of RMSW, as inhibition of methanogenesis was observed. Fluorescent in situ hybridization (FISH) studies showed that the percentages of *Eubacteria*, *Archaea*, H₂-utilising methanogens (HUM) and acetate-utilising methanogens (AUM) were stable within the 94.0–97.0%, 6.0–3.0%, 1.8–3.6% and 0.6–2.8% ranges, respectively.

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1. Introduction

Municipal solid waste (MSW) generation is increasing significantly in current society. Almost 1.9 billion tons of MSW are generated worldwide every year, which means about 218 kg/person/ year [1]. Anaerobic digestion (AD) is an attractive treatment strategy for generating energy from putrescible MSW and for mitigating the problem of disposal [1–3]. It also has a reduced environmental impact, especially with respect to the greenhouse effect. Due to the advantages of AD, many research studies have sought to optimize the AD of MSW, including the interesting option of the co-digestion process, which increases the load of biodegradable organic matter and produces a higher biogas yield [4-8]. Studies on the AD of MSW have shown that the C/N ratio of this waste presents average values of 10:1, below the optimum for anaerobic digestion (25:1) [9], while methane production (MP) is reduced due to the washing up of microorganisms and not to overloading [10,11]. Therefore, an increase in the loading rate of the AD process via the addition of readily biodegradable organic substances, such as glycerol, a major by-product of biodiesel production, could be an ideal strategy [5]. Biodiesel constitutes a promising alternative as a renewable fuel and its production capacity has been well developed in recent years [12,13]. The increased production of biodiesel supposes an increase in crude glycerol. The global biodiesel market is expected to reach 37 billion gallons by 2016, with an average annual growth rate of 42%. This means about 4 billion gallons of crude glycerol will be produced each year. A recent study [7] has demonstrated the effectiveness of glycerol supplementation (1% v/v) on the hydrogen production steps in thermophilic-dry dark fermentation of real MSW (RMSW) in batch mode. However, no previous studies have been published on the effect of glycerol addition on the biodegradation of RMSW and subsequent methane generation or on the effect on the microbial groups involved in the digestion process, which have varying sensitivities to environmental changes, especially acetogenic bacteria and methanogenic *Archaea*.

The aim of this study was to investigate the effect of different glycerol supplementations (0%, 0.1%, 0.25%, 0.5% and 1%) on the AD of RMSW. This residue is a known substrate for methane production (MP) [9,10,14,15]. For this purpose, anaerobic batch reactors were used to determine the anaerobic biodegradation and methane generation potential.

The effect of operational parameter modifications on dissolved chemical oxygen demand (COD_D), volatiles fatty acids (VFA), total volatile solids (TVS), ammonia, alkalinity, pH, microbiological population and MP were considered to study the effect of glycerol on



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Nomenclature					
AD AUM COD FISH GP HUM RMSW MP	anaerobic digestion acetate-utilising methanogens chemical oxygen demand fluorescence in situ hybridization gas production H ₂ -utilising methanogens real municipal solid waste methane production	MSW OFMSW OLR SMP TS TVS VFA	municipal solid waste organic fraction municipal solid waste organic loading rate specific methane production total solids total volatile solids volatile fatty acids		

the anaerobic batch processing of RMSW. The anaerobic consortia was analysed by fluorescent in situ hybridization (FISH), employing different oligonucleotide probes.

2. Materials and methods

2.1. Inoculum and substrate

Anaerobic digester effluent from single-phase dry-thermophilic AD of RMSW was used as inoculum. The RMSW and glycerol used to feed the reactors came from an industrial trommel (30 mm) located at the Las Calandrias MSW treatment plant (Jerez de la

Table 1

Physical-chemical and microbiological characterisation of the RMSW used in the trials.

Parameter	Average	Range	
COD _D (g/l)	170(10)	160-180	
TVFA (g acetic/l)	5(1)	4-6	
TVS (g/kg)	225(10)	215-235	
C:N (Organic matter)	13(3)	10-16	
Total population (10 ⁷ cells/ml)	36(7)	29-43	
EU (%)	88(2)	86-90	
Archaea (%)	12(2)	10-14	
AUM (%)	7(1)	6-8	
HUM (%)	5(1)	4-6	

Average values are shown with standard deviations in parentheses.

Table 2

Characterisation of the glycerol used in the trials.

Parameter	Average	Range
$COD_D(g/l)$	1400(100)	1300-1500
TVFA (g acetic/l)	<0.1	<0.1
TVS (g/kg)	820(10)	810-830
H ₂ O	12(1)	11-13
К	<0.1	-
Na	5.5(0.5)	5.0-6.0

Average values are shown with standard deviations in parentheses.

Table	3

Oligonucleotide probes used in the study.

Frontera, Cádiz-Spain) and the Abengoa Bionergy biofuel company (San Roque, Cádiz-Spain), respectively.

The RMSW was stored in 25 kg drums at 4 °C to avoid anaerobic degradation by the microorganisms present in the solid waste itself. The characterisation of the substrates used in the tests is shown in Tables 1 and 2.

2.2. Experimental reactors

To investigate the effect of different glycerol supplementations on the AD of RMSW, five different amounts of glycerol were tested (0%, 0.1%, 0.25%, 0.5% and 1%). The batch system used was made up of twelve reactors (2 for each condition + 2 blanks (inoculum without waste)), each with a total volume of 2 l and a useful volume of 1.7 l. Each reactor was equipped with an independent agitation system capable of maintaining an uniform moisture content and of redistributing the soluble substrate and microorganisms, as well as electric control of temperature. Ten digesters were initially loaded with a mixture of inoculum (700 g) and substrate (351 g RMSW in 11 of tap water), resulting in a final concentration of 40% v/v inoculum, which is considered optimum for biogas production and substrate acclimatization [16]. Different amounts of glycerol were then added to the reactors to give glycerol percentages in the 0% v/v (Test 1), 0.1% v/v (Test 2), 0.25% v/v (Test 3), 0.5% v/v (Test 4) and 1% v/v (Test 5) range (two for each test). The two blank experiments were conducted in order to determine the amount of endogenous methane production of the inoculum. This amount was accordingly subtracted from all the experimental results. The reactors were monitored until reaching a MP plateau, which represented a monitoring period of at least 25 days.

2.3. Analytical methods

The analytical methods employed in this study can be grouped into two categories: the physical-chemical parameters controlling the process of degradability, and methods for quantifying the microbial population in the reactors.

2.3.1. Physical-chemical analysis

For process monitoring and control of the reactors, the following analytical determinations were performed: COD_D, pH, TVS, VFA, ammonia and alkalinity. All analyses were carried out in

Probes	Probe sequences (from5' a 3')	Target	Formamide (%)	Time (h)	T (°C)	Reference
EUB338 S-D-Bact-0338-a-S-18 ARC915 MB1174 MX825	GCTGCCTCCCGTAGGAGT ACTCCTACGGGAGGCAGC GTGCTCCCCCGCCAATTCCT TACCGTCGTCCACTCCTTCCTC TCCCACCCTCGCCCAACCTAGC	Eubacteria None (negative control) Archaebacteria HUM AIIM	20 20 35 35 20	1.5 1.5 1.5 1.5 1.5	46 46 46 46	Zahedi et al. [10,11] Zahedi et al. [10,11] Zahedi et al. [10,11] Zahedi et al. [10,11] Zahedi et al. [10,11]

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