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Long-term acclimation of anaerobic sludges for high-rate methanogenesis from LCFA

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ABSTRACT

Inhibition of methanogens by long chain fatty acids (LCFA) and the low numbers of LCFA-degrading bacteria are limitations to exploit biogas production from fat-rich wastewaters. Generally reactors fail due to excessive LCFA accumulation onto the sludge. Here, long-term acclimation and bioaugmentation with a LCFA-degrading coculture were hypothesized as strategies to enhance methanogenic conversion of these compounds. Anaerobic sludges previously exposed to LCFA for more than 100 days converted a specific biomass-associated substrate of $(3.2 \pm 0.1) \text{ kg} \cdot \text{kg}^{-1}$ with very short lag phases (<1 day), whereas non-acclimated sludges showed lag phases of 11–15 days for metabolizing $(1.6\text{--}1.8) \text{ kg} \cdot \text{kg}^{-1}$. Addition of a coculture of *Syntrophomonas zehnderi* and *Methanobacterium formicicum* to sludges previously loaded with LCFA and containing different amounts of biomass-associated substrate (from $(0.5\text{--}3.2) \text{ kg} \cdot \text{kg}^{-1}$) did not improve methane production neither lag phases were shortened, indicating that the endogenous microbiota are not a limiting factor. Clearly, we show that long-term sludge acclimation to LCFA is essential for high rate methanogenesis from LCFA.

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1. Introduction

Long chain fatty acids (LCFA) are produced through lipids hydrolysis and are often abundant in lipid-rich wastewaters. LCFA have a high energy potential, but their conversion to methane in continuous anaerobic systems is frequently hindered by the poor solubility and association to the sludge [1]. High LCFA concentrations may increase the lag phase period in β -oxidation process [2], which is partially due to transport

limitation [3]. The accumulation of LCFA onto microbial cells creates a physical barrier and holds back the transfer of substrate and products [3]. Another limitation in LCFA conversion is often the low number of syntrophic bacteria in anaerobic sludges [4–6], which can limit the conversion rates of LCFA and/or intermediates. Efficient conversion of high LCFA loads in a continuous anaerobic reactor (organic loading rates (OLR) up to $21 \text{ kg m}^{-3} \text{ d}^{-1}$; 50% of COD as oleate) has been reported after a step-wise prolonged feeding [7]. It was suggested that the observed high rates are associated with the development

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of stable syntrophic community of LCFA-degrading bacteria and methanogenic archaea. Amendment of non-acclimated granular sludge with *Syntrophomonas zehnderi*, a LCFA-degrading bacterium, resulted in a 2-fold increase in the initial methane production rate from oleate batch conversion [8]. However, information on the long-term effect of bio-augmenting syntrophic bacteria in LCFA-fed reactors is lacking.

Here, we explore potential strategies to maintain and/or restore microbial activity after an LCFA shock. We studied long-term acclimation of anaerobic sludges to LCFA and investigated if longer acclimation times are beneficial to shorten the recovery of methanogenic activity after LCFA overload (LCFA overloaded sludge is used to designate sludge that has been withdrawn from a bioreactor continuously loaded with LCFA and containing biomass-associated substrate; accumulation of substrate onto the biomass normally precedes reactor failure episodes). In another experiment, LCFA overloaded sludges were bioaugmented with a coculture of *S. zehnderi* and *Methanobacterium formicium* to evaluate the potential of accelerating the recovery of methanogenic activity.

2. Materials and methods

2.1. Methane production by LCFA-overloaded sludges

2.1.1. Source of LCFA-overloaded sludge

A total of 54 sludge samples were collected from laboratory-scale anaerobic bioreactors and used for studying the effect of acclimation time on the lag phases of biomass-associated substrate conversion to methane, and its influence on the degradation of simple substrates (H_2 - CO_2 , acetate, propionate

and butyrate). Reactors had been operated with LCFA-based wastewaters containing oleate at a temperature of (308.15–310.15) K. Table 1 shows the conditions at the time of sampling and the contact time of the sludges with oleate. All the analyzed samples had biomass-associated substrate due to overload with oleate.

2.1.2. Biodegradation of the biomass-associated substrate

Sludge samples described in Table 1 were washed and centrifuged twice with basal medium prepared as described by Alves et al. [9]. Vials (25 cm³) containing 12.5 cm³ of basal medium were inoculated with the anaerobic sludges to a final concentration of VS between (3–5) kg·m⁻³ and sealed with butyl rubber stoppers and aluminum caps. The headspace of the bottles was then washed with a mixture of 80% N₂ and 20% CO₂ (total gas pressure in bottles' headspace was set at 100 kPa). Sludges were incubated without the addition of any substrate besides the substrate (mainly LCFA) associated to the sludge. Assays were performed in triplicate. Incubation was performed at 310.15 K while shaking (2.5 Hz). Biogas production was monitored with a pressure transducer and methane was quantified by gas chromatography (GC) [9]. Cumulative methane production data were fitted to the modified Gompertz equation (Section 2.4) to estimate lag phase time and maximum methane production in the assays. The maximum amount of methane produced (corresponding to the plateau of the methane production curves) was converted to its COD equivalent (0.064 kg mol⁻¹) and was considered as an indirect measurement of the amount of biomass-associated substrate (as previously described by Pereira et al. [12]). Specific biomass-associated substrate was calculated by dividing the determined COD equivalent by the VS content in the vials at the end of the assays.

Table 1 – Conditions prevailing and exposure time to oleate of sludge sources used in this work.

Reactor	Operation mode	Substrate	Co-substrate:Oleate (%)	Time in contact with oleate (d)	References
Fixed bed	Continuous	Skim milk + oleate	50:50	103	[9]
Fixed bed	Continuous	Oleate	0:100	214	[9]
Fixed bed	Continuous	Skim milk + oleate	50:50	103	[9]
Fixed bed	Continuous	Oleate	0:100	214	[9]
Fixed bed	Continuous	Skim milk + oleate	50:50	20–55 ^a	[10]
Fixed bed	Continuous	Skim milk + oleate	50:50	114–147 ^b	[10]
EGSB	Continuous	Skim milk + oleate	50:50	70	[11]
EGSB	Continuous	Skim milk + oleate	50:50	70 ^c	[11]
EGSB	Continuous	Oleate	0:100	70–219 ^d	[11]
EGSB	Continuous	Oleate	0:100	70–219 ^e	[11]
Column	Fed-batch	Skim Milk + oleate	50:50	103	[3]
EGSB	Continuous	oleate	0:100	178	[3]
UASB	Cycles	Skim Milk + oleate	50:50	17–213 ^f	[7]
UASB	Continuous	Skim Milk + oleate	50:50	473–665 ^g	[7]

EGSB – Expanded granular sludge bed; UASB – Up-flow anaerobic sludge blanket.

^a 6 samples were collected in this period.

^b 3 samples.

^c 2 samples, one at the top and other at bottom of the reactor.

^d 8 samples.

^e 7 samples.

^f 9 samples.

^g 12 samples.

2.1.3. Biodegradation of direct and indirect methanogenic substrates by LCFA overloaded sludges

The degradation of H₂–CO₂, acetate, propionate and butyrate by LCFA-overloaded samples was studied in batch assays according to the procedure described by Pereira et al. [12]. Briefly, sets of serum bottles were prepared as detailed in Section 2.1.2 and were individually supplemented with acetate (30 mol m⁻³), propionate (30 mol m⁻³), butyrate (15 mol m⁻³) or a mixture of H₂ (80%) and CO₂ (20%). Cumulative methane production was monitored, and the specific methane production values were calculated by dividing the initial slope of the cumulative methane production curves by the VS content of each vial at the end of the experiment. Background methane production due to residual substrate was measured in blank assays and discounted.

2.2. Bioaugmentation of LCFA-overloaded sludge with a coculture of *S. zehnderi* and *Methanobacterium formicicum*

2.2.1. Bioaugmenting culture

A coculture of *S. zehnderi* DSM 17840^T and *M. formicicum* DSM 1535^T was grown in bicarbonate-buffer mineral salt medium [13]. The methanogen was pre-grown in sealed bottles pressurized with a gas mixture (80% H₂ and 20% CO₂, 170 kPa). After *M. formicicum* growth, the headspace of the bottles was flushed and pressurized under sterile conditions with 80% N₂ and 20% CO₂, 170 kPa and further inoculated with a coculture of *S. zehnderi* and *M. formicicum*. The cultures were incubated at 310.15 K, statically and in the dark and two successive additions of 0.5 mol m⁻³ sodium oleate (≥99%, Fluka) were made. Then, the coculture was anaerobically centrifuged (15 691 m s⁻², 15 min, 277.15 K) and distributed in two bottles under N₂ atmosphere; one of these bottles was heat treated two times (394.15 K, 40 min) to inactivate the culture.

2.2.2. Source of LCFA-overloaded sludge

To obtain LCFA-overloaded sludge, a 2 dm³ (working volume) plexiglas up-flow anaerobic column reactor was operated at (308.15–310.15) K in continuous mode for 82 days. Anaerobic granular sludge from a brewery wastewater treatment plant (SCC – Sociedade Central de Cervejas e Bebidas, S.A., Portugal, 38° 52' 58" N, 9° 3' 25" W) was used as inoculum with a VS content of 16 kg m⁻³ and OLR between 2 and 6 kg m⁻³ d⁻¹ were applied. Reactor operational parameters are shown in Table 2. Biogas flow rate and composition, total and soluble COD, volatile fatty acids (VFA) and VS were monitored during the experiment.

COD removal efficiency, methane production rate and methane yields decreased during the operation, and LCFA tended to accumulate in the bioreactor (Table 2, Supplementary Figure S1). Mixed liquor samples (sludge and liquid medium) with increasing LCFA concentrations were withdrawn from the reactor at three different operation times (days 26, 55 and 82) and used to perform batch bioaugmentation experiments (Section 2.2.3). These sludge samples were characterized in terms of LCFA, VFA and VS. The specific biomass-associated LCFA content was calculated as previously described in Section 2.1.2.

2.2.3. Batch bioaugmentation experiments

Mixed liquor samples (35 cm³) withdrawn from the reactor at days 26, 55 and 82 were transferred to 70 cm³ bottles under 80% N₂ and 20% CO₂. *S. zehnderi* coculture was added to a final concentration of 0.24 kg m⁻³. Inactivated coculture was used in non-bioaugmented controls. To guarantee uniform hydrogenotrophic activity in bioaugmented and non-bioaugmented assays, *M. formicicum* active culture was added to all the vials, including controls (0.05 m³ m⁻³). Vials were incubated at 310.15 K statically and in the dark. Methane concentration in the vials headspace was monitored throughout the experiments. Methane production data were converted to its COD equivalent (kg) and divided by the vials biomass content (kg). Specific cumulative methane production data were then fitted to the modified Gompertz equation (Section 2.4). VFA and LCFA were quantified at the beginning and at the end of the assays and expressed as equivalent COD.

2.3. Analytical methods

In the batch biodegradation assays methane was quantified by gas chromatography (Chrompack CP 9000) using a Porapak Q column and a flame ionization detector. N₂ was used as carrier gas. Injection port, column and detector temperatures were 373.15 K, 308.15 K and 493.15 K, respectively. In the continuous reactor (Section 2.2.2), biogas flow rate was measured using a Ritter MilliGascounter (Dr. Ing. Ritter Apparatebau GmbH, Bochum, Germany), and the methane content was analyzed using a Micro-GC CP-4900 (Varian Inc.) with a 10 m PPU column heated at 353.15 K. A thermal conductivity detector (328.15 K) was used and helium (at 150 kPa) was the carrier gas. Injection port temperature was 383.15 K. Total and volatile solids were determined according to Standard methods [14]. Total and soluble COD was determined using COD-cuvette test kits (Hach-Lange GmbH, Düsseldorf, Germany) and a DR 2800 spectrophotometer (Hach-Lange GmbH). VFA were

Table 2 – Reactor operational parameters and performance data during continuous operation.

Period	Time (d)	OLR (kg m ⁻³ d ⁻¹)	HRT (d)	Milk:oleate (%)	COD removal efficiency (%)	Methane yield (%) ^a
I	0–9	2.3 ± 1.1	0.97 ± 0.07	50:50	98 ± 1	55 ± 5
	9–16	4.1 ± 0.5	0.97 ± 0.07	50:50	99 ± 0	69 ± 6
II	16–37	4.1 ± 0.5	0.97 ± 0.07	25:75	97 ± 1	68 ± 15
	37–55	4.1 ± 0.5	0.97 ± 0.07	0:100	65 ± 8	19 ± 4
	55–82	5.7 ± 0.9	0.71 ± 0.01	0:100	64 ± 7	5 ± 2

^a Calculated as the ratio between the methane produced (converted to its COD equivalent) and the COD removed.

analyzed by high-performance liquid chromatography (HPLC; Jasco, Japan) using a Chrompack organic analysis column (30×6.5 mm) with a mobile phase of $5 \text{ mol m}^{-3} \text{ H}_2\text{SO}_4$ at a flow rate of $0.01 \text{ cm}^3 \text{ s}^{-1}$. The column temperature was set at 333.15 K and spectrophotometric detection was performed at 210 nm. LCFA were analyzed as previously described by Neves et al. [15]. Esterification of free fatty acids was performed with propanol, in acid medium (3.5 h at 373.15 K). Propyl esters were further extracted with dichloromethane and analyzed in a gas chromatograph (Varian 3800) equipped with a flame ionization detector and a eq.CP-Sil 52 CB $30 \text{ m} \times 0.32 \text{ mm} \times 0.25 \mu\text{m}$ capillary column (Teknokroma, TR-WAX). Helium was used as gas carrier at a flow rate of $0.02 \text{ cm}^3 \text{ s}^{-1}$. Initial oven temperature was set at 323.15 K for 2 min and final temperature of 498.15 K was attained with a ramp rate 4.72 K s^{-1} . Injector and detector temperatures were 493.15 K and 523.15 K, respectively.

2.4. Data analysis

Experimental methane production data (expressed in COD) from batch incubations were fitted by the modified Gompertz equation (equation (1)) for estimation of the maximum methane production and lag phase time. For methane production curves and fitting results please check Figure S2 in the supplementary material.

$$M(t) = P \cdot \exp \left[- \exp \left[\frac{R_m \cdot e}{P} (\lambda - t) + 1 \right] \right] \quad (1)$$

Where $M(t)$ = specific cumulative methane production ($\text{kg} \cdot \text{kg}^{-1}$), P = maximum specific methane production ($\text{kg} \cdot \text{kg}^{-1}$), R_m = specific methane production rate ($\text{kg} \cdot \text{kg}^{-1} \text{ d}^{-1}$), $e = 2.7182818$, λ = lag phase time (days). R^2 values and the standard error for each variables obtained were calculated.

Statistical significance of the differences observed in the maximum cumulative methane production achieved in bio-augmentation experiments was evaluated using single factor analysis of variances (ANOVA). Statistical significance was established at the $p < 0.05$ level.

3. Results and discussion

3.1. Effect of acclimation on methane production by LCFA-overloaded sludges

LCFA-overloaded sludge samples containing different amounts of biomass-associated substrate were studied. Biomass-associated substrate was mainly composed of LCFA, but small amounts of other compounds were also present when a co-substrate (milk) was included in the feed [7,12]. It has been previously shown that specific amounts of biomass-associated LCFA higher than 1 kg kg^{-1} have a negative effect on the specific methane production rate (estimated by the initial slope of cumulative methane production curves) [12]. Here, we show that the biomass-associated LCFA also has an important effect on the lag phase preceding the onset of methane production, and interestingly this effect is dependent on the previous exposure time of the sludge to LCFA (Fig. 1).

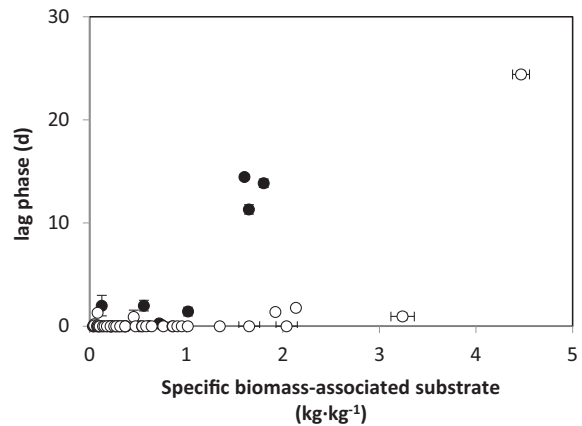


Fig. 1 – Lag phase in methane production from degradation of biomass-associated substrate from different sludge samples. Sludges exposed to LCFA for a period < 100 days (●). Sludges exposed to LCFA for a period > 100 days (○).

For values of specific biomass-associated substrate up to 1 kg kg^{-1} almost no delay in methane production was observed, for all the tested sludges. However, a considerable increase in the lag phases was observed during the incubation of sludges with higher amounts of associated substrate, in particular non-acclimated sludges (exposed to LCFA for less than 100 days). Sludges that had been exposed to extended contact with LCFA (>100 days) performed better during the conversion of the biomass-associated substrate to methane. Sludge with specific substrate content as high as $(3.2 \pm 0.1) \text{ kg} \cdot \text{kg}^{-1}$ showed a lag phase of less than 1 day. Nevertheless, even acclimated biomass showed a high lag phase in methane production (i.e. (24.4 ± 0.3) days) when the specific biomass-associated substrate was about 4.5 kg kg^{-1} . This means that sludges acclimated for long periods (and with progressively increasing loads of oleate) recover their methanogenic activity rapidly when overloaded with LCFA, as long as the LCFA feed is interrupted. On the other hand, non-acclimated sludges need more time to convert the biomass-associated substrate. Silvestre et al. [16] studied biomass adaptation over co-digestion of sewage sludge and trapped grease waste and suggested also that a slow increase in grease waste could favor biomass acclimation, increasing LCFA degradation and reducing their latter inhibitory effect. During activity tests with oleate (approx. 0.25 kg m^{-3}), a lag phase of 15 days was observed when using a sludge inoculum that was not acclimated to fat while no lag phases were observed with the adapted inoculum [16].

Usually, biomass-associated LCFA also have a negative effect on the specific methanogenic activity. Pereira et al. [17] observed a significant decrease of the specific methanogenic activity with $\text{H}_2\text{-CO}_2$, acetate, propionate and butyrate by LCFA overloaded sludges. In our work, specific methanogenic activity tests performed with the sludges exposed to extended contact with LCFA (>100 days), and containing associated substrate up to 5 kg kg^{-1} , showed that biomass acclimation did not avoid the sharp decrease of specific methane production from $\text{H}_2\text{-CO}_2$, acetate, propionate and butyrate (Fig. 2).

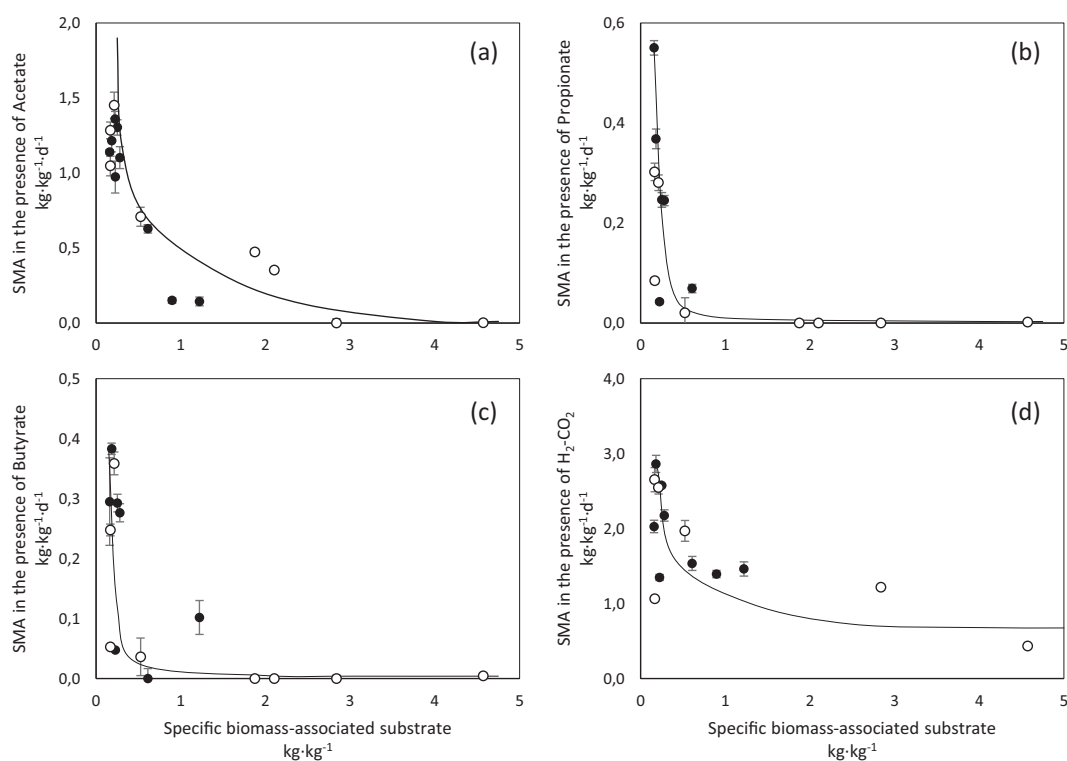


Fig. 2 – Specific methanogenic activity of the different sludge samples, containing different biomass-associated substrate contents, when in the presence of (a) acetate, (b) propionate, (c) butyrate and (d) H_2 - CO_2 as substrates. Sludges exposed to LCFA for a period < 100 days (●). Sludges exposed to LCFA for a period > 100 days (○).

As previously reported by Pereira et al. [3,17], this effect is less severe in the presence of H_2 - CO_2 and likely more related with mass transfer limitations imposed by LCFA accumulation than to metabolic inhibition. A decrease in hydrogenotrophic (with H_2 - CO_2), acetoclastic (with acetate) and acetogenic (with butyrate) activities has also been observed in manure bioreactors after being exposed to oleate pulses [18]. However, activities were not irreversibly damaged and could be recovered after the degradation of the biomass-associated substrate [3,18].

In continuous bioreactors these results have an important application, since the recovery of methanogenic activity, after an episode of LCFA overload, is possible if the feeding is interrupted and sufficient time is provided for degradation of the biomass-associated substrate. This recovery time, corresponding to the lag phases reported in this work, was shown to depend on long-term biomass acclimation, and on the amount of accumulated LCFA (Fig. 1).

3.2. Bioaugmentation of LCFA-overloaded sludge with a coculture of *S. zehnderi* and *M. formicicum*

LCFA-overloaded sludge with increasing LCFA concentrations were used to perform batch bioaugmentation experiments. LCFA and VFA concentrations were assessed at the beginning and at the end of the experiments (Table 3). In all samples withdrawn from the reactor, palmitate accounted for (79–89) % of total LCFA; acetate was the main VFA detected in the samples collected on days 55 and 82. Specific methane

production in bioaugmented vials and non-bioaugmented controls, as well as the different parameters calculated by fitting the modified Gompertz equation are shown in Table 3 (methane production curves in Figure S2).

Comparing the initial specific LCFA content and the maximum methane production obtained (P in Table 3) the methane yield accounted only for (35–64) % of the expected values (theoretical values of methane production from oleate and VFA are presented in Supplementary Table S1). The final values of LCFA content were around 2 kg kg^{-1} , representing (59–65) % of the initial LCFA concentration. Palmitate was the main LCFA present at the end of the assays, accounting for (81–94) % of the total LCFA. VFA were also found in concentrations between 0.3 and 0.5 kg kg^{-1} . The non-converted VFA and LCFA accounted for the difference between the initial COD and the COD recovered as methane.

No significant differences were observed in the maximum methane production neither in the lag phases, between the bioaugmented and non-bioaugmented assays (Table 3, Figure S2). Bioaugmentation success relies on the ability of the inoculated microorganisms to continue their activity in the new environment. Palatsi et al. [19] tested the addition of fresh manure to recover LCFA-inhibited bioreactors and observed an increase in maximum methane production rates. However, manure does not function solely as a seed of microorganisms and effects related to the presence of co-substrates or even the dilution of the bioreactor content cannot be disregarded. In LCFA-overloaded sludges, LCFA are largely associated with the biomass that is slowly obtaining the ability to degrade

Table 3 – Specific VFA and LCFA concentrations at the start (t_0) and at the end (t_f) of the bioaugmentation assays and methane production parameters (P , R_m and λ) obtained from methane production curves.

	VFA (kg kg ⁻¹)		LCFA (kg kg ⁻¹)		P (kg kg ⁻¹)	R_m (kg kg ⁻¹ d ⁻¹)	λ (d)
	t_0	t_f	t_0	t_f			
Bioaugmentation 1 (day 26)							
Bioaug.	–	n.d.	0.45 ± 0.07	n.d.	0.27 ± 0.01	0.07 ± 0.01	0
Non-bioaug.	–	n.d.	0.45 ± 0.07	n.d.	0.28 ± 0.01	0.04 ± 0.00	0
Bioaugmentation 2 (day 55)							
Bioaug.	0.086	n.d.	0.69 ± 0.1	n.d.	0.39 ± 0.01	0.01 ± 0.00	1 ± 1
Non-bioaug.	0.086	n.d.	0.69 ± 0.1	n.d.	0.36 ± 0.01	0.01 ± 0.00	3 ± 1
Bioaugmentation 3 (day 82)							
Bioaug.	0.272	1.89	3.2 ± 0.61	0.39	1.25 ± 0.04	0.04 ± 0.00	46 ± 2
Non-bioaug.	0.272	2.08	3.2 ± 0.61	0.41	1.27 ± 0.05	0.04 ± 0.00	38 ± 2

n.d. – not determined; P (maximum specific methane production), R_m (specific methane production rate), and λ (lag phase time) were calculated by fitting the modified Gompertz equation to the experimental data; Person's correlation coefficient (r) ≥ 0.93.

these compounds. Apparently, the bioaugmented methanogenic community used in this study does not have sufficient access to the insoluble substrate to enhance methanogenesis.

4. Conclusions

The amount of biomass-associated LCFA and long-term acclimation (>100 days of exposure) are essential factors for avoiding the delay of methane production imposed by LCFA. Results point to the importance of the physical integrity of the syntrophic communities with the insoluble substrate and *in situ* development of highly-specialized microbial communities. From an application point of view, long-term sludge acclimation and avoidance of excessive LCFA accumulation are essential factors for the efficient conversion of LCFA to methane. Monitoring LCFA accumulation into the sludge is important for preventing serious overloading events. Ideally, specific biomass-associated substrate should be kept below 1 kg kg⁻¹, although well-adapted sludges could still have a good performance with approximately three times this value.

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Appendix A. Supplementary data

Supplementary data related to this article can be found at <http://dx.doi.org/10.1016/j.biombioe.2014.05.012>.

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