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1 **Effect of post digestion temperature on serial CSTR biogas**  
2 **reactor performance**

3

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11

1 **Abstract**

2 The effect of post digestion temperature on a lab-scale serial continuous-flow stirred  
3 tank reactor (CSTR) system performance was investigated. The system consisted of a  
4 main reactor operated at 55 °C with hydraulic retention time (HRT) of 15 days followed  
5 by post digestion reactors with HRT of 5.3 days. Three post digestion temperatures (55  
6 °C, 37 °C and 15 °C) were compared in terms of biogas production, process stability,  
7 microbial community and methanogenic activity. The results showed that the post  
8 digesters operated at 55 °C, 37 °C and 15 °C gave extra biogas production of 11.7%,  
9 8.4% and 1.2%, respectively. The post digester operated at 55 °C had the highest biogas  
10 production and was the most stable in terms of low VFA concentrations. The specific  
11 methanogenic activity tests revealed that the main reactor and the post digester operated  
12 at 55 °C and 37 °C had very active acidogens and methanogens. In contrast, very low  
13 methanogenic activity was observed at 15 °C.

14

15 **Keywords:** biogas; post digestion; temperature

16

## 1 **Introduction**

2 Anaerobic digestion of manure or other slurries is commonly performed in continuous-  
3 flow stirred tank reactors (CSTR). A single CSTR is simple to operate but less efficient  
4 compared to immobilized-cell configurations, such as upflow anaerobic sludge blanket  
5 (UASB). However, manure slurry is not suitable for direct application in the UASB  
6 reactor due to its high content of particulate matter. In CSTR, the slowly degradable  
7 fibers in manure are normally not fully utilized during the digestion (Ahring and  
8 Angelidaki, 2000). It has been reported that only 50-70% of organic matter is converted  
9 to biogas in a manure digester with typical hydraulic retention time (HRT) of 15-30  
10 days (Hartmann et al., 2000). Moreover, a single CSTR digester can “loose” biogas  
11 production due to “short-circuit”, where parts of organic material in the feed remain in  
12 shorter duration in the reactor than the nominal retention time (Angelidaki et al., 2005).  
13 There are several reports on improving biogas production in the CSTR process, for  
14 example, by increasing hydraulic retention time (Hansen et al., 1998), co-digestion with  
15 other organic wastes (Ahring et al., 1992), or by pre-treatment of the substrate to  
16 improve degradation of recalcitrant materials (Ahring and Angelidaki, 2000; Hartmann  
17 et al., 2000). Simulation results using Anaerobic Digestion Model 1 (ADM1) (Boe et  
18 al., 2005) showed that a serial CSTR configuration consisting of one reactor (or so-  
19 called main digester) with long retention time and another reactor (or so-called post  
20 digester) with short retention time could improve biogas production and achieve better  
21 effluent quality in terms of VFA (volatile fatty acids) concentration compared to a  
22 single CSTR reactor. Once the main digester has sufficient volume to maintain stable  
23 operation, extra biogas production could be obtained by post digestion (Angelidaki et  
24 al., 2005). Full-scale biogas reactors are commonly connected to effluent storage tanks

1 to store digested manure before recycling to agricultural farmlands. These effluent  
2 storage tanks (known also as post-digesters) have normally covered their top for  
3 collection of the extra biogas that might be produced in post digesters. However, these  
4 tanks have often no temperature control installed, which can affect the microbial  
5 activity and composition and consequently the biogas production. It is well-known that  
6 one of the main factors affecting anaerobic digestion process is the temperature  
7 (Hashimoto, 1983). Over the last decades, the effect of temperature on biogas  
8 production, effluent quality (in terms of VFA concentrations) and process stability have  
9 been widely investigated (Harris and Dague, 1993; Kugelman and Guida, 1989). Some  
10 studies showed better performances of thermophilic anaerobic digestion compared to  
11 mesophilic one as indicated by the lower VFA concentrations and higher biogas  
12 production (Kim et al., 2002; Mackie and Bryant, 1995). In general, higher process  
13 temperatures enhance microbial activities, resulting in higher biogas production. With  
14 regards to microbial community structure, thermophilic conditions result in lower  
15 microbial diversity compared to mesophilic conditions (Karakashev et al., 2005;  
16 Sekiguchi et al., 1998). Additionally, the thermophilic reactors are more susceptible to  
17 ammonia inhibition compared to the mesophilic reactors. Indeed, high temperatures  
18 increase free ammonia concentrations which negatively affect the anaerobic digester  
19 performances (Angelidaki and Ahring, 1994). Furthermore, increasing temperatures  
20 also increase energy costs for heating the post digesters. Therefore, the benefits of  
21 increasing post digestion temperatures should be thoroughly investigated.

22         The objective of this study was to investigate the effect of temperatures on the  
23 post digester performances. Three post digesters operated at temperatures of 55 °C, 37  
24 °C and 15 °C were compared in terms of biogas production, effluent quality and process

1 stability. The structure of the microbial communities and methanogenic activity were  
2 also determined.

3

## 4 **Materials and Methods**

### 5 **Substrate characteristics**

6 Substrate fed to the main reactor was cow and pig manure mixed with approx. 25 % of  
7 industrial wastes (mainly dairy and slaughterhouse wastes). The substrate was obtained  
8 in one batch from a full-scale biogas plant (Lemvig, Denmark). The substrate was  
9 blended to ensure homogeneity, and then stored at -20 °C for the whole period of  
10 experiment. The frozen substrate was thawed and kept at 4 °C for 2-3 days before use.  
11 Substrate characteristics are shown in Table 1.

12

### 13 **Reactor setup and operations**

14 The experiment was carried out in a serial configuration of CSTR digesters (Figure 1).  
15 The main digester (referred as R0) was made of a 9 L reactor with 7.2 L working  
16 volume and a hydraulic retention time (HRT) of 15 days. Reactor R0 was built from  
17 double glass cylinder fitted with stainless steel plates on top and bottom. The top plate  
18 supported the mixer, mixer motor, feed tube, and effluent tube, temperature measuring  
19 port and sampling port. The bottom plate had one sampling port. The operating  
20 temperature in reactor R0 was maintained at 55 °C by circulating hot water through the  
21 space between the reactor glass walls. Reactor R0 was connected to three post digesters  
22 as shown in Figure 1. The three post digesters were operated in parallel with operating  
23 temperatures of 55 °C (referred as R55), 37 °C (referred as R37), and 15 °C (referred as  
24 R15), respectively. Each post digester was made of a 1 L glass reactor with 800 mL

1 working volume and HRT of 5.3 days, which was in the range of typical HRT of post  
2 digester used in the Danish full-scale biogas plants. The operating temperatures in the  
3 post digesters were controlled by circulating hot water (for R55 and R37) or cooling  
4 liquid (for R15) through a water jacket surrounding the reactors.

5 The amount of 120 mL substrate was fed into the main reactor four times per day using  
6 peristaltic pump with timer control. Each post digester was fed with 150 mL of the  
7 effluent from the main reactor once per day by manually start a peristaltic pump. Rest of  
8 the effluent from the main reactor (less than 30 mL per day) was collected in the  
9 effluent bottle of the main digester. All reactors were continuously stirred throughout  
10 the experiment by motor mixer in the main reactor and by magnetic stirrers in the post  
11 digesters. Operating pressure of all reactors was around 1 atm. The slight overpressure  
12 from the produced biogas and the influent inside the reactors pushed out the effluents  
13 from liquid surface through the effluent tubes on top of the reactors. The effluents were  
14 collected in the effluent bottles. The effluent bottles were connected with gas meters  
15 that registered the produced biogas production.

16

### 17 **Analytical methods**

18 The biogas production was measured by an automated displacement gas metering  
19 system with a 100 mL reversible cycle and registration (Angelidaki et al., 1992). The  
20 water used in gas meter was acidified to pH 3 by HCl and NaCl was added to prevent  
21 gas dissolution. Methane content of the biogas was measured by a gas chromatograph  
22 (Mikrolab, Århus) equipped with thermal conductivity detector and a glass column, 6ft  
23 x 3mm I.D., packed with Poropack Q (10/80). The injector, detector and oven  
24 temperature were isothermal at 55 °C. The carrier gas was Helium with flow rate 40

1 mL/min. Theoretical biogas production was calculated according to Angelidaki and  
2 Sanders (2004). Ammonium and total nitrogen, pH, total solids (TS), and volatile solids  
3 (VS) were determined according to standard methods (Greenberg et al., 1992). Free  
4 ammonia was estimated as described in Angelidaki et al. (1999). VFA was measured  
5 using a gas chromatograph (GC) HP 5890 Series II equipped with flame ionization  
6 detector and a FFAP fused-silica capillary column, 30m x 0.53mm I.D., film thickness  
7 1.5 µm. Nitrogen was used as a carrier gas. The oven temperature was initially set at 70  
8 °C and then increasing 10 degrees/min. to 190 °C and kept at final temperature for 3  
9 min. The injection port and detector temperature were 150 °C and 200 °C, respectively.

10

#### 11 **Pulse load tests**

12 At day 94, a lipid pulse load was introduced to study the effect on the process stability  
13 of the different setups. The pulse load was added by directly injecting 219 g of rapeseed  
14 oil, which corresponding to 14.6 times of the normal VS load, into the main reactor R0.

15

#### 16 **Specific methanogenic activity (SMA) tests**

17 At day 62, batch experiments for measuring specific methanogenic activity (SMA) on a  
18 specific substrate were carried out (Angelidaki and Schmidt, 2002). 40 mL basal  
19 anaerobic (BA) medium prepared as described in Karakashev et al., (2005) was  
20 dispensed anaerobically in 100 mL serum bottles. The media was supplemented with  
21 different substrates- acetate (20 mM), propionate (10 mM), butyrate (10 mM),  
22 hydrogen/carbon dioxide (50/50) under 1 atm, and glucose (10 mM). After addition of  
23 vitamin solution and Na<sub>2</sub>S.9H<sub>2</sub>O as a reducing agent the medium was inoculated with 10  
24 mL fresh samples from each reactor and incubated in respective temperature of



1 inoculums. Bottles with BA medium and inocula only but without substrates were used  
2 as controls (blanks). All the tests were prepared in duplicates. Methane production and  
3 VFA levels were monitored closely for the first 10 days and thereafter for 50 days.

4

### 5 **Microbial community composition**

6 Fluorescent *in situ* hybridization (FISH) method of Hugenholtz et al. (2001) was used to  
7 assess main phylogenetic groups of methanogens, i.e. *Methanomicrobiales*,  
8 *Methanococcales*, *Methanobacteriales* *Methanosaetaceae* and *Methanosarcinaceae*, as  
9 previously described by Karakashev et al. (2005).

10 Polymerase chain reaction-temperature gradient gel electrophoresis (PCR-TGGE) was  
11 used to determine microbial community profiles of the reactors according to Muyzer  
12 and Smalla (1998).

13

### 14 **Results**

15 Biogas production, VFA concentrations and pH from all reactors are presented in Figure  
16 2. The experiment was carried out for 111 days and steady state was obtained after day  
17 45 when VFA and biogas production had reached relatively stable values (daily  
18 variations lower than 14% in the main reactor). The process parameters during steady  
19 state are summarized in Table 1. Methane content was found similar in all reactors  
20 around 70%. During steady state average biogas production in reactors R0, R55, R37  
21 and R15 were of 24.6, 2.9, 2.1 and 0.3 mL biogas/mL feed, respectively (Figure 2a). In  
22 comparison, the theoretical biogas production resulting from full decomposition of the  
23 organic matter in raw manure would have been approximately 40 mL-biogas/mL-feed

1 (0.5 L-CH<sub>4</sub>/gVS). The additional biogas obtained from the post digesters were 11.7%,  
2 8.4% and 1.2% for R55, R37 and for R15 respectively.

3 During steady state, total VFA concentrations in reactor R0, R37 and R15 were  
4 similar (20 mM in average), while reactor R55 had significantly lower VFA  
5 concentration (4 mM). This corresponded to the low %TS and %VS in the reactor R55  
6 compared to other reactors. Reactor R37 and R15 seemed to have biomass accumulation  
7 due to lower removal of organic matters. Acetate and propionate were predominant  
8 VFA in all reactors. Total VFA concentration in reactor R0 followed the trend of total  
9 VFA in the feed, and total VFA concentrations in reactor R37 and R15 also followed  
10 the trend of VFA in reactor R0 (Figure 2b). In contrast, the VFA concentrations in  
11 reactor R55 were constantly low, almost independent from the VFA concentration in  
12 R0. The pH values in all reactors were relatively stable in the range of 7.7-8.1. Levels of  
13 ammonium nitrogen in all reactors were similar within a range of 3.3-3.8 g-N/L (Table  
14 1). The corresponding calculated free ammonia was ranged from 0.13 to 1.1 g-N/L.

15 After adding a pulse load of 219 g rapeseed oil into reactor R0 on day 94, a  
16 double increase in biogas production and subsequent increased VFA in reactor R0 were  
17 observed. Similar responses were noticed in all post digesters. Nevertheless, reactor  
18 R55 showed smaller increase in VFA concentration compared to the increase observed  
19 in R37 and R15. Indeed, the effect of pulse load was more pronounced in R15 and R37.  
20 In all reactors, pH did not change significantly after the lipid pulse load (Figure 2).

21 In regards to microbial activities, specific methanogenic activity (SMA) tests  
22 revealed that the degradation rate of the tested substrates increased with the increase of  
23 temperatures from 15 °C to 55 °C (Table 2). Higher degradation rates of acetate,  
24 H<sub>2</sub>/CO<sub>2</sub>, butyrate and propionate were observed in R0 and R55 compared to R37 and

1 R15. A higher methanogenic activity from acetate was noticed in R37 than R55. No or  
2 very low degradation was noticed for all the tested substrates in R15. These results were  
3 supported by FISH and PCR-TGGE analysis of microbial community composition in all  
4 reactors (pictures not shown). According to the FISH results, both *Bacteria* and  
5 *Archaea* (especially *Methanosaetaceae*) were more abundant in R0, R55 and R37  
6 compared to R15, where the relative abundance was in order of R0>R55>R37.  
7 Regarding the PCR-TGGE results, the microbial community profiles at day 62 and day  
8 93 were different for both abundance and community structure. From the bacterial and  
9 archaeal community profiles, the relative abundances of microorganisms were  
10 compared based on the intensity of DNA bands, and summarized in Table 3.

11 Profiles of bacterial community showed four well separated bands representing  
12 four different phylogenetic groups of dominant *Bacteria* in the samples. At day 62, band  
13 1 existed only in reactor R0 and R15. Band 3 was more abundant in reactor R37 and  
14 R55, while band 4 was more abundant in R0 and R15. Bacterial diversity in reactors  
15 R37 and R55 was significantly lower compared to R0, while the bacterial profile in  
16 reactor R15 was similar compared to R0. At day 93, band 1 disappeared from reactor  
17 R0 and R15, and band 3 disappeared from all reactors. Comparing day 93 to day 62,  
18 relative abundance of band 2 in reactors R0, R55 and R15 increased with 27 %, 32 %  
19 and 35 %, respectively. For reactor R37, band 4 abundance increased with 38 %  
20 compared to day 62 but the abundance of band 2 was relatively constant with time.

21 *Archaea* community profile showed four well separated bands representing four  
22 different phylogenetic groups of dominant *Archaea*. At day 62, band 2 was lacked from  
23 R37. The diversity of *Archaea* in reactor R0 and R55 did not change significantly with  
24 time. However, a change in abundance was observed. Band 2 in reactor R0 and R55 was

1 enriched at day 93 compared to day 62. In contrast, at day 93 band 2 disappeared from  
2 reactor R15 and appeared in R37.

3

#### 4 **Discussion**

5 Many researchers have reported strong effect of temperature on the microorganisms  
6 where lowering the operational temperature led to a decrease in the maximum specific  
7 growth and substrate utilization rates. Under psychrophilic conditions (10-15 °C),  
8 chemical and biological reactions proceed much slower than under mesophilic  
9 conditions (30-40 °C) (Lettinga et al., 2001). The results from the present study also  
10 showed that operating temperature in the post digesters had significant effect on biogas  
11 production, methanogenic activities and microbial community. The higher additional  
12 biogas production obtained from R55 and R37 compared to R15 (Figure 2a) suggested  
13 that operating temperature in the post digesters should be maintained at temperature  
14 higher than 15 °C. Additional biogas production in post digesters came from residual  
15 VFA and residual organic matter present in the effluent of the main reactor (R0  
16 effluent). If the biogas production in post digesters only came from residual VFA  
17 decomposition, then the amount of biogas produced should be close to the value of  
18 biogas equivalent from VFA in R0 effluent, which was 1.1 mL-biogas/mL-feed (Figure  
19 3a). However, it was observed that the amount of biogas obtained in reactors R55 and  
20 R37 were 2.9 and 2.1 mL-biogas/mL-feed, respectively, which were higher than the  
21 biogas equivalent from VFA in R0 effluent. This implies that VFA were also produced  
22 in reactor R55 and R37 from acidogenesis of residual organic matters. Comparatively,  
23 reactor R15 had very low biogas production and VFA production. From Figure 3b, the  
24 combination of R0 with R55 could obtain biogas production of 27.5 mL-biogas/mL-

1 feed, which corresponds to around 70% of the theoretical biogas potential in the feed  
2 (40 mL-biogas/mL-feed), i.e. to approx. 70% utilization of the organic matter in  
3 manure. The combination of R0 with R37 and R15 could obtain 67% and 62% of the  
4 theoretical biogas potential in the feed, respectively. This showed that maintaining high  
5 temperature in the post digester could increase biomass utilization in the system.  
6 Moreover, R55 and R37 were more efficient than R15 to recover the biogas yield from  
7 VFA in R0 effluent during overload. As it was seen that adding lipid pulse at day 94  
8 increased biogas production and VFA concentration in reactor R0 with subsequently  
9 increased biogas production in R55 and R37 compared to R15.

10 The low methane yield in R15 revealed that methanogenesis was inactive, and  
11 the similar VFA concentrations between R0 and R15 meant that acidogenesis occurred  
12 at very low extent. Moreover, compared to R55 and R37, reactor R15 was more  
13 dependent on R0 as its variation in acetate concentration strongly followed the trend of  
14 R0 (Figure 2b). The constant low VFA concentrations and high biogas production in  
15 R55 confirmed its stability and independence from the variations in R0, although the  
16 high free ammonia level in R55 (1.1 g-N/L) could be inhibitory according to the  
17 threshold limit reported from Hansen et al. (1998). The high concentrations of free  
18 ammonia in reactors R0 and R55 were mainly due to high operating temperature  
19 (Angelidaki and Ahring, 1994). Moreover, the total ammonia concentration in all  
20 reactors were in the range of 3-4 g-N/L. It was previously shown that the total ammonia  
21 concentration up to 4 g-N/L did not result in process inhibition both at thermophilic and  
22 mesophilic conditions (Angelidaki et al., 2005).

23 The results from SMA tests showed a good correlation to the results from  
24 reactor operational parameters measured at steady state. The high initial degradation

1 rates (Table 2) showed for all tested substrates in reactor R0, R55 and R37 suggested a  
2 high activity of all microorganisms (both acidogens and methanogens). Moreover,  
3 glucose degradation in reactor R37 was as efficient as that in R55, indicating that  
4 acidogenesis were equally active in these two reactors. The higher aceticlastic  
5 methanogenic activity from acetate in R37 compared to R55 suggested that aceticlastic  
6 methanogens were more active at mesophilic conditions than at thermophilic. This  
7 could be due to the higher methanogenic diversity in mesophilic compared to  
8 thermophilic processes (Sekiguchi et al, 1998). Some aceticlastic methanogenic species  
9 present in R0 could grow better in R37 than in R55 under the same post digester HRT.  
10 The low activity on glucose in R15 suggested that acidogenesis from hexoses was low.  
11 Very low methane production from acetate and H<sub>2</sub>/CO<sub>2</sub> and no consumption of butyrate,  
12 propionate in R15 suggested that both acetogens and hydrogenotrophic methanogens  
13 were inactive.

14 The FISH results showing high abundance of different phylogenetic groups of  
15 *Bacteria* and *Archaea* in R0, R37 and R55 indicated that acidogens and methanogens  
16 were active in these reactors. This is in good agreement with the results from reactor  
17 operation and SMA tests. Moreover, the FISH results showing low microbial abundance  
18 in R15 suggested that the microorganisms might be metabolically inactive, resulting in  
19 low content of ribosomal RNA which could be detected by FISH technique. This  
20 observation correlated to almost no biogas production and very low activity in the SMA  
21 test at low temperature. Nevertheless, we believe that the microorganisms were not  
22 washed out from the reactor R15 although they were inactive, since R15 was running at  
23 very short retention time (HRT 5.3 days) and always received the fresh effluent from R0  
24 which contained large amount of active microorganisms.

1 Results obtained by PCR - TGGE analysis also showed that the reactor  
2 temperature affected microbial community structures. At day 62, the community profile  
3 of both *Bacteria* and *Archaea* in reactor R15 was very similar to R0. This was due to  
4 the fact that lower temperature resulted in lower specific growth rate (Kevbrina et al.,  
5 2001), thus, the microbial population remained unchanged. At 55 °C and 37 °C,  
6 the microorganisms were more active so the enrichment of some methanogenic group in  
7 the post digester compared to reactor R0 could be observed.

8 Although the degradation rate is very slow at psychrophilic temperature, it has  
9 been reported that stable methanogenesis can still be achieved if the microorganisms has  
10 been long-term adapted to the low temperature (> 4 months) (Kettunen and Rintala,  
11 1997; Lettinga et al., 1999). If this is the case, the post digestion at 15 °C might be  
12 possible if R15 is large enough to build up its own adapted culture. However, in this  
13 experiment the post digesters strongly relied on the microorganisms in R0 effluent  
14 which were thermophiles. The microorganisms coming into R15 were, in some degrees,  
15 exposed to temperature shock and became inactive.

16 Results obtained in this study showed that the post digestion at 37 °C and 55 °C  
17 provided extra biogas yield which could give reasonable compromise between the cost  
18 of heating and the benefit from additional biogas. However, the cost of heating is  
19 strongly depending on the quality of reactor insulation. Before applying this idea in the  
20 full-scale application, an economic evaluation should be done comparing the investment  
21 costs for insulation of the existing post digesters, the operational costs for heating, and  
22 the amount of additional biogas energy expected from the post digesters.

23 Another benefit of using high temperature in post digestion is that the effluent  
24 from the biogas plants would contain less residual organics which will minimize the

1 methane emission from degradation of the residual organics during transportation back  
2 to the farmers and during storage period before manure spreading on agricultural land.

3

#### 4 **Conclusion**

5 Results obtained in this study showed that the temperature could affect biogas  
6 production and microbial ecology in the post digesters. When the main biogas digester  
7 is running at thermophilic temperature, the operating temperature of 15 °C in the post  
8 digesters is very inefficient as the bacteria become inactive as they were unadapted to  
9 low temperature. The post digester operated at 37 °C could obtain similar biogas  
10 production as at 55 °C. The specific methanogenic activity tests showed that the main  
11 reactor had highest microbial activities, followed by the post digester with 55 °C and 37  
12 °C. Moreover, post digestion temperature of 55 °C could also provide the effluent with  
13 low VFA, which means lower potential of smell when recycling on farmland. Thus, it is  
14 recommended that the temperature in the post digester should be kept as close as  
15 possible to the temperature of the main digester to maintain the bacterial activities.

16

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23



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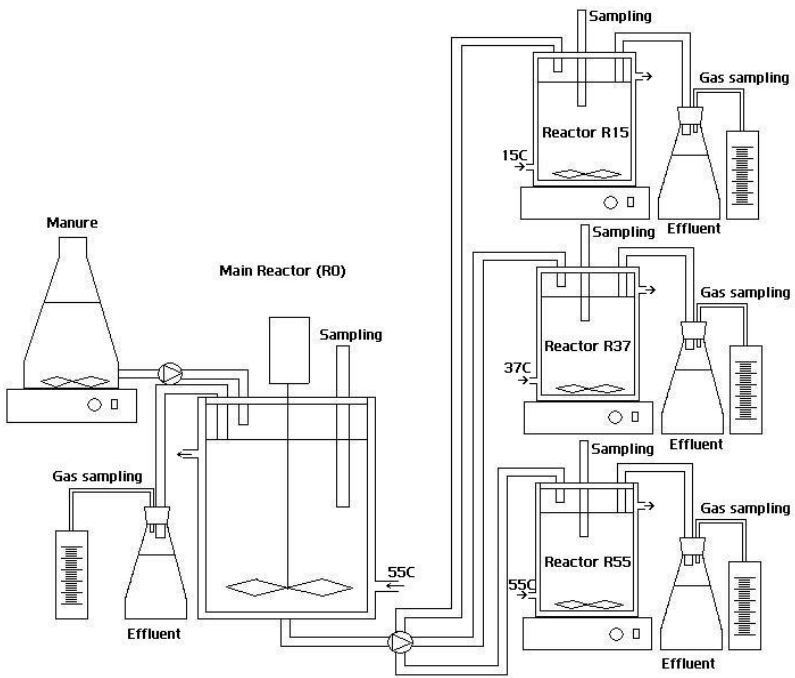
**Table 1.** Average reactor parameters over steady state period (day 45-70) and main substrate characteristics.

<b>Parameters</b>	<b>Substrate</b>	<b>R0</b>	<b>R55</b>	<b>R37</b>	<b>R15</b>
TS (%)	8.21	5.50	5.23	5.88	6.28
VS (%)	4.78	2.52	2.39	2.74	2.92
Total-N (g-N/L)	4.62	4.7	4.7	4.8	4.8
Ammonium-N (g-N/L)	3.26	3.3	3.6	3.8	3.3
Free ammonia (g-N/L)	0.01	0.85	1.10	0.22	0.13
Lipids (g/L)	5	<0.01	<0.01	<0.01	<0.01
pH	6.87	7.89	7.87	7.72	8.09
Total VFA (mM)	259.8	22.8	4.0	19.5	22.7
Acetate (mM)	185.1	15.8	3.4	12.3	15.3
Propionate (mM)	41.2	5.8	0.4	6.1	5.5
Iso-butyrate (mM)	6.3	0.5	0.04	0.5	0.7
Butyrate (mM)	15.9	0.2	0.02	0.1	0.3
Iso-valerate (mM)	6.2	0.3	0.05	0.3	0.7
Valerate (mM)	5.2	0.1	0.02	0.1	0.1
Biogas Production (mL-biogas/mL-feed)		24.6	2.9	2.1	0.3
Extra biogas production (% ) compared to R0		-	11.7	8.4	1.2
Methane (%) in biogas		69.9	68.8	77.9	70.1
Extra methane production (% ) compared to R0		-	11.6	9.3	1.2

**Table 2.** Microbial activity on different substrates.

<b>Substrate</b>	<b>Activity unit</b>	<b>R0</b>	<b>R55</b>	<b>R37</b>	<b>R15</b>
H <sub>2</sub> /CO <sub>2</sub>	mL CH <sub>4</sub> /(gVS.hr)	0.27	0.24	0.16	0.008
Acetate	mL CH <sub>4</sub> /(gVS.hr)	0.50	0.46	0.48	0.005
Propionate	mM Pr/day	0.90	0.70	0.50	-
Butyrate	mM But/day	3.10	1.70	1.30	-
Glucose	mmole CH <sub>4</sub> -equivalent/day	0.47	0.40	0.40	0.043

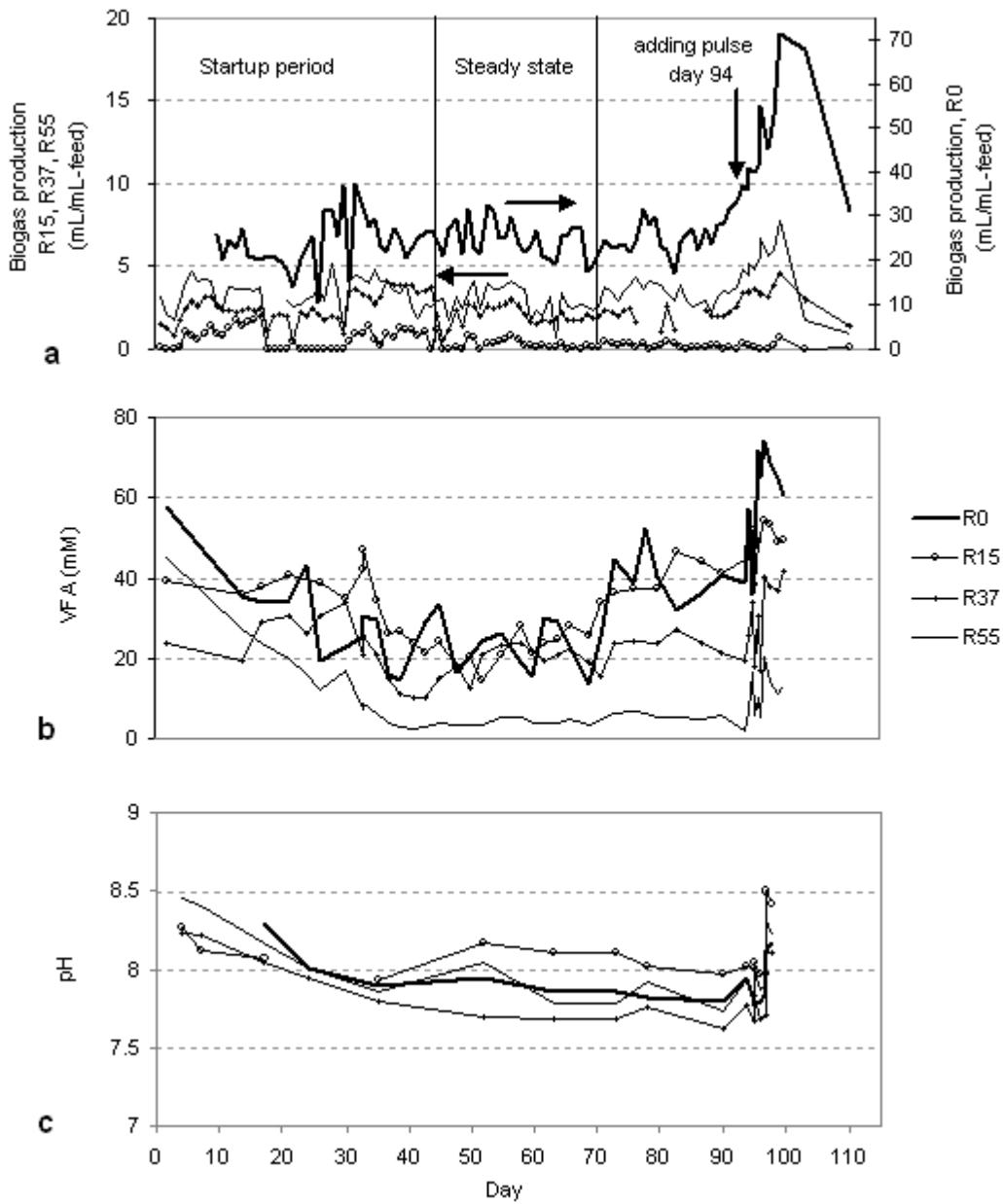




1 Figure 1. Reactor setup for the experiment

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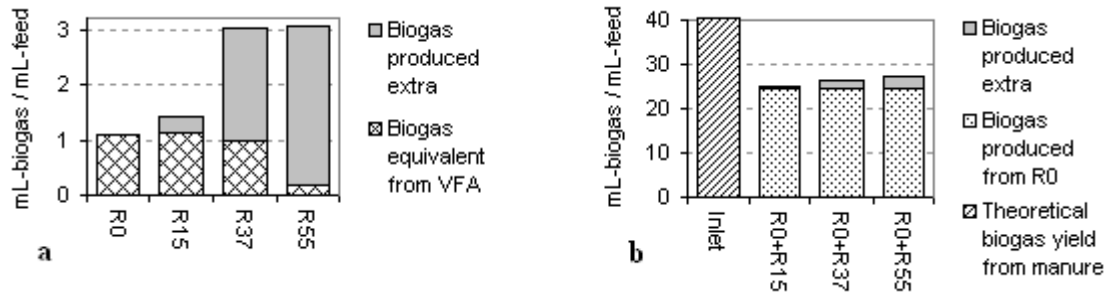


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5 **Figure 2** Experimental results: (a) Biogas production, (b) VFA concentration, (c) pH values

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9 **Figure 3.** Comparison of the actual and theoretical biogas yield; (3a) Actual biogas production  
 10 obtained in post digesters compared to biogas potential from VFA in R0 effluent, (3b) Actual  
 11 biogas production from R0 and post digesters compared to theoretical biogas yield from manure