

Chapter

**BIOLOGICAL TREATMENT OF LEACHATE
IN A SUBMERGED ANAEROBIC
MEMBRANE BIOREACTOR**

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ABSTRACT

In this chapter, a simulated **Organic Fraction of Municipal Solid Waste** (OFMSW) was treated in an anaerobic two-stage membrane process. The OFMSW feedstock was fed to a ten litre **hydrolytic reactor** (HR) where solid and liquid fractions were separated by a coarse mesh, while the **leachate** was fed to a three litre **submerged anaerobic membrane bioreactor** (SAMBR) with *in situ* **membrane** cleaning by biogas sparging beneath a flat sheet microfiltration module. The aim was to develop and optimize this two-stage process where the use of a membrane in both reactors to uncouple the Solid and Liquid Retention Times (SRT and HRT) would allow us to improve the current performances obtained with single stage designs. The **Denaturing Gradient Gel Electrophoresis** (DGGE) technique was used to monitor the microbial population in the reactors and have a better understanding of the archaeal and bacterial distribution in a two-stage process.

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It was found that **Chemical Oxygen Demand (COD)** removal was greater than 90% at a **Hydraulic Residence Time (HRT)** of 1.6-2.3 days at a maximum **Organic Loading Rate (OLR)** of 20 g COD/L.day. Even though the influent COD of the leachate was constantly changing giving rise to a transitory Food to Microorganisms ratio over time, the permeate COD from the SAMBR was typically between 300 and 500 mg/L_s which can therefore be defined as a stabilised leachate. Because of the fluctuating properties of the leachate produced in the HR_s the process was deemed more representative because the SAMBR treated a leachate with varying organic strength which is what can happen on a full scale.

The COD removal in SAMBR2 was 94.5% on average, and only 1.6% in the subsequent aerobic polishing bioreactor (AMBR-**Aerobic Membrane Bioreactor**), so that a total COD removal of 96.1% was achieved at 0.4 day HRT. On average, 26% of the recalcitrants from SAMBR2 could be degraded aerobically in the AMBR. However, as in SAMBR1, at HRTs lower than 2 days, particulate solids in the leachate built up at the bottom of the SAMBR, eventually leading to the diffuser blocking. At MLTSS beyond 20 g/L, the **transmembrane pressure (TMP)** culminated at 850 mbar and the flux dropped to 0.5 **L.m⁻².hr⁻¹** LMH until the end of the experiment. The permeate of the SAMBR was low in COD and relatively constant which promoted the growth of **autotrophic bacteria** in the subsequent AMBR_s so that 97.7% of the NH₄⁺-N was removed at a maximum nitrogen loading rate of 0.18 kg NH₄⁺-N/m³.day.

Calcium in the leachate was found to precipitate in the AMBR because of the higher pH. A sample taken from the membrane consisted most likely of pure **hydroxylapatite** Ca₅(PO₄)₃(OH) which had a needle shape, whereas the background of the precipitate consisted more of nodules of calcium carbonate with traces of manganese, iron, magnesium, aluminium, sulphur and sodium.

The very high **molecular weight (MW)** aromatic organics in the leachate fed to the SAMBRs were almost fully degraded in the bulk of the SAMBRs. Moreover, their permeate was absolutely free of them which indicates a full rejection of these compounds by the membrane. The medium MW compounds in the range of 395 - 646 kDa were more likely to be rejected by the membrane while the MW ≤ 395 kDa were observed in the permeate. Regarding the evolution of the medium and low MW compounds over the 200 days, it can be stated that overall there was no build-up in the absorbance in any effluent.

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1. INTRODUCTION

The amount of waste produced in England and Wales, annually, is around 434 million tonnes, of which 8% is Municipal Solid Waste (MSW) (Phillips et al., 2005). Moreover, household waste is growing by around 2% each year. If this rate continues, nearly twice as many new waste management facilities will be required by 2020 to reach UK and EU targets. Municipal Solid Waste (MSW) is a growing concern in the world, and European authorities discourage municipalities choosing landfilling as their waste management strategy. The European landfill directive (99/31/EC) requires that biodegradable municipal waste landfilled should be reduced by 2010 to 75% of that produced in 1995. In addition, the Household Waste Recycling Act 2003 in the UK requires that waste collection authorities shall ensure that by the end of 2010 they should collect at least two types of dry recyclable waste separate from the remainder of the waste. The aim of the Act was to increase the recycling rate of household waste by recycling or composting, and implementation of the Act should allow the UK to reach the target of 25% of MSW being recycled or composted by 2005, 30% by 2010 and 33% by 2015. Unfortunately, in 2003/2004, the majority (72%) of municipal waste in England was still disposed of to landfill, 9% was incinerated and only 19% was recycled or composted (DEFRA, 2005). The last figure increased to 36% in 2008 (DEFRA, 2008), but MSW is still a huge problem.

Currently, biological treatment methods such as composting and **Anaerobic Digestion (AD)** offer the only route for recycling organic matter and nutrients from the organic fraction of municipal solid waste (OFMSW) (Braber, 1995). Composting can diminish the organic matter by about 50% through the formation of carbon dioxide and water (Held et al., 2002), but it represents an energy consuming process (around 30-35 kWh is consumed per tonne of waste input) while AD is a net energy producing process (100-150 kWh per tonne of input waste). AD is a process by which bacteria degrade organic matter and convert it to mainly carbon dioxide and methane. The main advantage of this technology is the destruction of organic components without the addition of oxygen, and it produces useful by-products such as a gaseous fuel and stabilized solid residue that can be sold as a **soil fertilizer**. In contrast, aerobic treatment is plagued by the

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production of large amounts of activated sludge (about 10 times more than AD) and consumption of significant amounts of energy due to aeration requirements (Braber, 1995; Mahmoud, 2002; Speece, 1996). Digester technology for solid waste has basically followed the conventional approach used for sewage sludge digestion over the last fifty years. The result is that the market is currently dominated (over 90%) by single tank designs (Vandevivere et al., 2003) aimed at promoting good mixing. In sewage sludge digestion the rate-limiting factor is the slow growth of the methanogenic population; this necessitates long retention times (15-20 days), which in turn leads to low loadings because of the low solids concentration of the feed sludge. This approach is inappropriate for high solids digestion, however, where the rate-limiting step is in the hydrolysis of the substrate, and hence excessively long retention times are required even at moderate loading rates. In Europe, the first full scale AD plant treating OFMSW were commissioned in the last 1980s and are still in operation demonstrating that it can be considered as a mature technology and more than 168 industrial plants have been built since (Monson et al., 2007).

The anaerobic digestion of solid waste and the organic fraction of municipal solid waste (OFMSW) in particular has been reported to be more stable and robust in two stages (Banks and Humphreys, 1998; Vieitez et al., 2000). The first phase includes a hydrolytic reactor (HR) in which **enzymes** transform complex molecules into simple ones that can be converted into fatty acids; these acids are then degraded by **methanogens** in the second reactor. The slow-growing methanogens can therefore operate at an optimum pH independent of the pH in the first reactor. The literature revealed that the hydrolytic stage takes place either in a well-mixed reactor, or in a leach-bed where the leachate is recirculated to the top of the reactor to promote the hydrolysis of solid waste. The loading rate can be as high as 20-25 g VS.L⁻¹.d⁻¹ in a Continuously Stirred Tank Reactor depending on the proportion of easily degradable waste, but the hydraulic retention time (HRT) has been restricted to 2 days due to process limitations. In batch leach-bed processes, the OLR is low and they require a large footprint.

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In an anaerobic system, the slow growth rate of mixed cultures posed a major problem for reactor design as long minimum solid retention times (SRT) are required. Thus, anaerobic bacteria are easily washed out because their doubling times are normally higher (5-6 days) than the HRT. As efficient anaerobic digester performance depends on the hydrolysis rate of soluble solids and the biomass concentration, the challenge was to develop a reactor which enables the HRT to be uncoupled from the SRT. Thus the objective of this work was to investigate the use of membranes in anaerobic bioreactors to achieve this. Possible problems that may arise include the clogging of the membrane due to the wide molecular weight (MW) distribution from the complex substrates. This may significantly lower the flux which will in turn affect the loading rate to the methanogenic reactor.

A common problem associated with the continuous wet anaerobic fermentation process is that inhibition can occur due to the build-up of light metal ions and ammonia (Gallert et al., 2003). Another issue is the generation of approximately 100-320 L wastewater per ton of waste input (Dierick, 2006; Fricke et al., 2007). Therefore, recycling the stabilized leachate to the head of a continuous wet process treating OFMSW could significantly reduce the amount of wastewater produced, and reduce the environmental impact of MSW disposal. This was one of the objectives of the present work.

Mechanical-biological treatment (MBT) of the OFMSW is now the main strategy to reduce biodegradable MSW in waste in the UK and in Europe (Scaglia and Adani, 2008). It consists of mechanical pretreatment followed by an aerobic (composting-like process) or anaerobic process so that waste impacts are reduced. These processes have attracted attention because they produced a stabilized waste, which has a low impact when disposed of in landfill (Adani et al., 2004). However, landfill leachate still represents a serious environmental threat. In the UK, the leachate from the OFMSW disposed in landfills is currently being treated anaerobically in ponds or aerobically by Sequencing Batch Reactor (SBR) (Robinson,

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2005); raw leachate is fed to an aerated reactor. After a certain time, aeration is shut down and the solids are allowed to settle. The supernatant, i.e., the treated leachate, is discharged to sewer or a watercourse. On lab-scale, several researchers have used UASB (Upflow Anaerobic Sludge Blanket), fixed-film and packed bed reactors to treat the leachates from MSW and the COD removal efficiencies were typically higher than 90% (Anderson and Saw, 1992; He et al., 2005). Aerobic Membrane reactors have also been used for biological oxidation and nitrification/denitrification (Ahn et al., 2002; Laitinen et al., 2006), but the use of an anaerobic MBR for the stabilization of the leachate is not widespread. In particular, Submerged Anaerobic Membrane Bioreactors (SAMBRs) have been developed at Imperial College (Hu, 2004), and stable operation was reported for high strength feed (20,000 mg COD/L) at 20 hours HRT and a permeate flux of 1.5-2 LMH (litres m⁻²hour⁻¹) (Akram, 2006), but no information is available about its behaviour for high-strength leachates from the OFMSW.

The aim of this chapter was thus to explore the feasibility of treating the OFMSW in a two-stage process where the OFMSW is first hydrolyzed in a hydrolytic reactor, and leachate is treated in an anaerobic membrane bioreactor so that the more stable and efficient process can achieve better performance than current technologies. The performance of the process was investigated in a semi-continuous mode, i.e., the HR is fed once every day or every two days. AD of the OFMSW can take place either in dry or wet systems depending on the Total Solids (TS) content of the reactor. For wet fermentation, the dry matter content is adjusted to 8-16% by addition of process water, whereas for dry systems no or only little process water is added to moisten the feedstock. An example of a full scale wet two-stage system is the Schwarting-Uhde process which can sustain an OLR of up to 6 kg VS/m³.day, whereas a full scale dry 2-stage process such as the BRV plant can achieve up to 8 kg VS/m³.day (Trosch and Niemann, 1999). When a biomass retention scheme is added such as in the BTA and Biopercolat designs, an OLR up to 15 kg VS/m³.day can be applied successfully (Gallert et al., 2003; Wellinger et al., 1999). The biofilm growth in the second stage of the Biopercolat process allows the system to

run at an overall retention time of 7 days. In the BTA process, the HRT could be reduced to 5.7 days.

For laboratory and pilot scale anaerobic leachate treatment experiences, OLRs from 3 to 22 kg COD/m³.day with COD removal efficiencies of 68 - 97% and HRTs between 1.5 and 2.6 days have been reported previously (Chang, 1989; Henry et al., 1987; Kennedy et al., 1988). On the other hand, aerobic leachate treatments found in the literature have been applied to leachate with COD values between 3,000 and 48,000 mg/L. COD removal efficiencies reported for aerobic systems are higher than 70%, with HRTs ranging from 2.5 to 20 days (Boyle and Ham, 1974; Cook and Foree, 1974; Maris et al., 1984; Robinson and Maris, 1985; Uloth and Mavinic, 1977). However, less sludge is generated and less energy is required if an anaerobic step is followed by an aerobic one. In this process sequence, the final aerobic stage serves as a post-treatment to improve the final effluent quality (Agdag and Sponza, 2005; Hoilijoki et al., 2000). For instance, Borzacconi et al. (1999) loaded a UASB at an OLR of 20 kg COD/m³.day at an HRT of 2 days, and achieved a COD removal greater than 80%; the subsequent aerobic rotating biological contactor achieved 72% COD removal. Xu et al. (2008) removed 7% of the SCOD in an anaerobic filter, while 82% was removed in the subsequent aerobic MBR at 9.5 days HRT. They pointed out that most of the polycyclic aromatic hydrocarbons (PAHs) were biodegraded in the anaerobic step. The final effluent, however, still contained 1,000 mg SCOD/L which suggests that there were some biorefractory organics that were hardly removed by both biological processes and the membrane process. Another process advantage is the possibility of removing ammonia from leachate in the aerobic step, but it is known that high influent COD promotes heterotrophic growth and inhibits ammonium oxidation (Cheng and Chen, 1994; Hanaki et al., 1990). Different process configurations have been reported for the simultaneous removal of COD and ammonia from landfill leachate. Im et al. (2001) used an up-flow anaerobic biofilm reactor (36°C), an aerobic activated sludge reactor (23°C) and a clarifier achieving an organic removal rate of 15.2 kg COD/m³.d in the anaerobic reactor and ammonium removal rate of 0.84 kg

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N/m³.day in the aerobic reactor operating at 4 days HRT. Agdag and Sponza (2005) obtained 98% COD removal of food waste leachate at an OLR of 16 kg COD/m³.d in two UASBs (HRT=1.25 day) and an aerobic CSTR used in sequence. Ninety nine percent of NH₄⁺ was removed at 4.5 days HRT in the aerobic CSTR. Chen et al. (2008) used an anaerobic-aerobic moving-bed biofilm system and achieved a COD removal of 92% at an OLR of 15.7 kg COD/m³.d, while 97% of NH₄⁺-N was removed when the HRT of the aerobic step was more than 1.25 days. Jokela et al. (2002) obtained over 90% nitrification at 0.13 kg N/m³.day at 25°C and 1.4 day HRT in an upflow filter with crushed bricks.

The objectives of this chapter were numerous: the effect of the inoculum on the behaviour of the SAMBR was investigated; the stability of the SAMBRs and the AMBR was tested at different HRTs and OLRs; and an AMBR operating at ambient temperature was set up to determine whether the recalcitrants from the SAMBR could be biodegraded aerobically. After 200 days of operation, another objective was to see if there was a build-up of recalcitrants with time due to the permeate recycle, or if there was slow degradation, and **GC-MS analysis** was performed to determine what if any these recalcitrants were. Another pertinent question related to continuous wet anaerobic fermentation process with effluent recycle is whether light **metals ions** (Na⁺, K⁺, Mg²⁺, Ca²⁺) and Cl⁻, PO₄³⁻, SO₄²⁻ and **ammonia** accumulate to inhibitory levels (Gallert et al., 2003) and these parameters were therefore monitored throughout the 200 days of operation. **Leachate recirculation** over a tank filled with MSW is relatively well documented (Bilgili et al., 2007; Hao et al., 2008), but recirculation of stabilized leachate from membrane bioreactors is not. During anaerobic treatment of organic wastes, approximately 100-170 L wastewater per ton of waste input are generated (Fricke et al., 2007). Therefore, recycling the stabilized leachate to the head of a continuous wet process treating OFMSW could significantly reduce the amount of wastewater produced, and reduce the environmental impact of MSW disposal.

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2. MATERIALS AND METHODS

2.1. Experimental Setup

2.1.1. *Hydrolytic Reactor*

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Figure 1. 3-D view of the hydrolytic reactor (left) and stirrer (right).

The hydrolytic reactor (HR) is a 20 liter cylinder made of acrylic sheet; a stainless steel mesh follows a concentric arrangement inside the cylinder and has a grid of 1mm square holes. A stirrer moves inside the mesh allowing two pieces of rubber to rub against it. Figure 1 shows the reactor and the stirrer. The reactor is wrapped in a copper coil in which hot water is circulated and glass fiber is used to cover the coil so that the reactor internal temperature is maintained at $35 \pm 1^\circ\text{C}$. The stirrer speed is set at 64 rpm.

The HR (10L working volume) was operated intermittently (15 min ON-15 min OFF). The HR was fitted with a 50 micron stainless steel macro filter (Spectrum Laboratories Inc.) on the inside of the stainless steel mesh (1 mm) in order to retain the large partially hydrolyzed particles and

thereby separate the coarse solids from the leachate being fed to the SAMBRs, so that the SRT and the HRT were uncoupled.



Figure 2. Inside view of the Hydrolytic Reactor (HR) without the stirrer (left). Photo of the laboratory HR set-up (Right).

The HR was inoculated with 4L of biomass from a previous batch test in the HR at a residence time of about 50 days. The inoculum was sieved through a 180 micron screen and its TSS and VSS were 2.74 and 2.07 g/L, respectively. The HR was loaded with 400 g OFMSW on a dry matter basis (≈ 340 g VS), and the volume was adjusted to 10L with tap water containing NaHCO_3 so that the HR was started up at 4,000 mg equivalent CaCO_3/L of alkalinity. The HR was then fed semi-continuously with a feedstock of 10% Total Solids that was prepared by adding leachate from the HR to the simulated OFMSW in order to blend the mixture and obtain a homogeneous slurry. Fresh tap water was only added to the HR to keep a constant working volume. Until day 159, the HR was fed once every two days, then from day 160 onwards, it was fed once a day. The digestate from the HR was removed just before feeding, and it was dewatered by hand compression over a 500 microns sieve. The liquid fraction was used to moisten the fresh feedstock, while a fraction of the dewatered digestate was taken, weighed and discarded to set a specific SRT in the HR.

2.1.2. Methanogenic Reactor

A schematic diagram of the experimental set up used is shown in Figure 3. Each submerged anaerobic membrane bioreactor (SAMBR) has a working volume of 3 litres, and is made of acrylic panels. Each reactor contains a standing baffle designed to direct the fluid to the upcomer and downcomer regimes. The reactors were maintained at $35 \pm 1^\circ\text{C}$, and the biomass was continuously mixed using headspace biogas that was pumped (Charles Austen Pumps, Model B100SEC) through a stainless steel tube diffuser to generate coarse bubbles. The bubbles push the sludge flow upward between the membrane module and the reactor wall in the upper section. A flask (500 mL) was placed before the biogas pump in order to collect any sludge going to the gas line due to excessive foaming or excess of liquid in the reactor so that the biogas pump was protected against any liquid that could harm the mechanism. The sparging rate was controlled by a gas flowmeter (2 - 20 LPM, ColeParmer, USA) to minimise membrane fouling. A data logging system (USB-1408FS, Measurement computing) was used to monitor transmembrane pressure (PMP1400, 1 bar A, Druck), and permeate flow rates (13 – 100 mL/min, Flow Sensor Rytan Sensor, McMillan). The membrane module (Kubota) had 0.1 m^2 of total surface area with a pore size of 0.4 micron. The membrane panel was comprised of a solid acrylonitrile butadiene styrene support plate (5 mm thick) with a spacer layer between it, and a welded flat sheet membrane, which was made of polyethylene, on both side. The membrane was coated with a special chemical (10%w ethanol) to give a hydrophilic surface to the membrane.

The HR and SAMBR1 were connected in series: the leachate was fed to SAMBR1 and the permeate from SAMBR1 was recycled to the HR in order to maintain the moisture and alkalinity of the system. On day 45, SAMBR2 was fed on leachate in parallel with SAMBR1 in order to compare the effect of inoculum on the start-up of SAMBR. The HR, SAMBR1 and SAMBR2 were maintained at $35 \pm 1^\circ\text{C}$. The biogas sparging rate was set at 5 L/min (LPM) to minimize cake formation on the

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membrane. On day 130, an **Aerobic Membrane Bioreactor** (AMBR) operating at ambient temperature (21-22°C) was started up to treat the permeate of SAMBR2. The permeate of the AMBR was then returned to the HR. The two SAMBRs and the AMBR were three litre reactors fitted with a **Kubota polyethylene flat sheet membrane** of 0.1 m² of total surface and a pore size of 0.4 microns.

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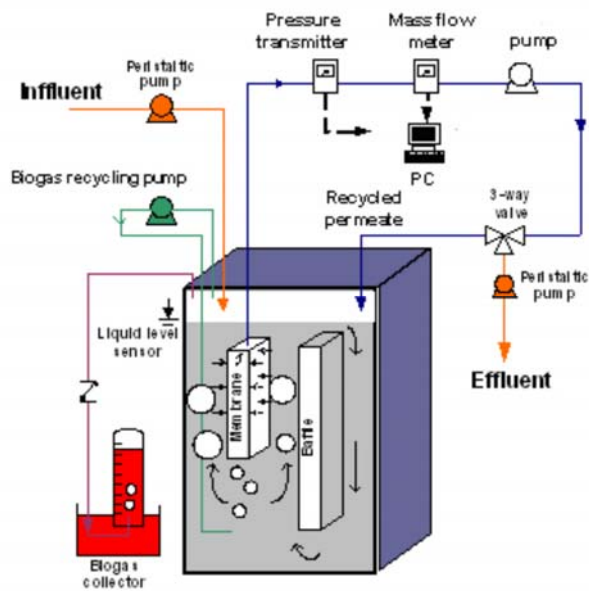


Figure 3. Schematic diagram of the **Submerged Anaerobic Membrane Bioreactor**.

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SAMBR1 was inoculated with 0.5 L of seed from a SAMBR fed on leachate from the same simulated OFMSW at a HRT of 4 days (Figure 4). The volume was adjusted to 3 L with the anaerobic biomedium defined in Owen et al. (1979) so that the initial TSS and VSS were 3.31 and 2.54 g/L, respectively.

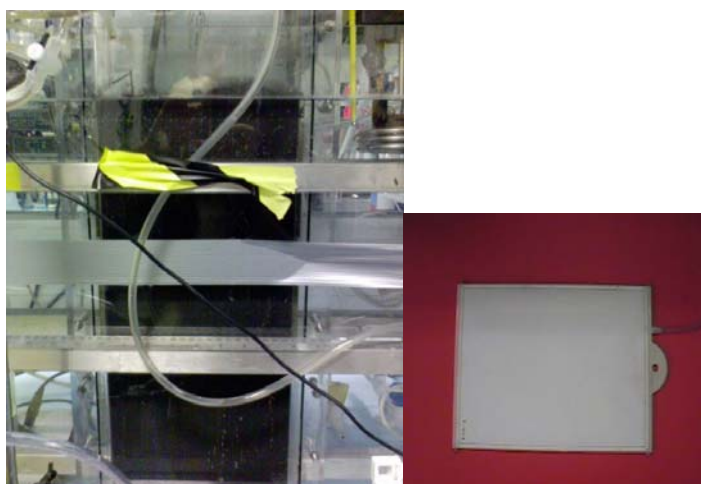


Figure 4. Left: photo of the Submerged Anaerobic Membrane Bioreactor (SAMBR) during start-up. The tip of the submerged membrane module can be seen above the liquid. Right: Kubota membrane module in the SAMBR.

SAMBR2 was inoculated with biomass from a 4 litre chemostat batch-fed (once a week) on a 8 g COD/L feed with a composition given elsewhere (Nachaiyasit and Stuckey, 1995). The feed consisted of peptone and meat extract (25% on a COD basis) and a synthetic VFAs mixture (75% on a COD basis). The ratios of the VFAs compared to acetic acid were 1.2, 0.05, 0.22, 0.08, 0.23 for propionate, iso-butyrate, n-butyrate, iso-valerate, n-valerate, respectively. These ratios were typically observed in the raw leachate obtained in previous tests from the simulated OFMSW. The supernatant of the chemostat was discarded and the settled solids were used to inoculate SAMBR2. The volume was adjusted to 3L with the anaerobic biomedium defined in Owen et al. (1979) so that the initial TSS and VSS were 2.56 and 1.78 g/L, respectively.

The AMBR was inoculated with biomass from a dye wastewater plant at an initial MLTSS and MLVSS of 3 and 2.3 g/l, respectively. Air was used to mix the reactor content at 2 LPM.

2.2. Analytical Methods

2.2.2. pH

pH was measured using a calibrated pH meter (Jenway, Model 3020). Values obtained were accurate to within ± 0.02 units.

2.2.3. Oxido-Reduction Potential

The oxido-reduction potential (ORP) was measured with a platinum-band ORP electrode (Cole-Palmer, USA) connected to a Jenway 3020 pH meter. Values obtained were accurate to within ± 3 mV.

2.2.4. Biogas Composition and Production Rate

The composition of biogas was determined using a Shimadzu GC-TCD fitted with a Porapak N column (1500×6.35 mm). The carrier gas was Helium set at a flow rate of 50 ml/min. The column, detector and injector temperature were 28, 38 and 128°C, respectively. The peak areas were calculated and printed out on a Shimadzu Chromatopac C-R6A integrator. Samples of 1 ml were collected using 1 ml plastic syringes (Terumo). The coefficient of variation for 10 identical samples was $\pm 2\%$.

In the HR, the gas production rate was measured using Tedlar bags connected on the top. To empty the Tedlar bag, it was connected to the top of a column full of water. A valve at the bottom of the column was then opened to let the water drain into a collection vessel as the gas filled the column. The water stopped running when the Tedlar bag was empty which enabled the volume of biogas produced to be measured. The percentage of methane was analyzed by taking a sample at the top of the column. ~~As this method measures a volume of gas that is expanded due to sucking by water flowing down by gravity, the volume must be corrected using the equation provided in Appendix B.~~

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In the SAMBR, the **gas production rate** was measured using the water displacement method with cylinders filled with water acidified with 2% (v/v) H₂SO₄ and 10% (w/v) NaCl according to Agdag and Sponza (2007). The top of the SAMBR was connected to the top of the cylinders, and when gas was produced the water left the cylinder from the bottom and was collected in a vessel placed above it. A one-way valve was placed between the cylinder and the collection vessel so that the water in the collection vessel did not exert any pressure on the gas in the cylinder.

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2.2.5. **Volatile Fatty Acids (VFAs)**

For **acetic, propionic, n-butyric, iso-butyric, n-valeric, iso-valeric and n-caproic acids** analysis, the sample was filtered through a 0.2 micron filter, and then acidified by adding one drop of concentrated sulphuric acid, and eventually analyzed using a Shimadzu Gas Chromatograph with a flame-ionised detector and a SGE capillary column (12m x 0.53mm ID-BP21 0.5micron). The mobile phase was helium and the injector, column and detector temperature were 200, 80 and 250°C, respectively. The relative standard deviation was ± 3% for ten identical samples.

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2.2.6. **Gas Chromatography - Mass Spectrometry (GC-MS)**

For the GC-MS analysis, the analytes of interest (low molecular weight hydrophobic compounds) were extracted using a solid phase extraction (SPE) procedure. The Oasis HLB cartridge (Waters Corporation) was first conditioned with 3 mL methyl tertiary-butyl ether (MTBE), 3mL methanol and 3 mL deionized water (DW). A sample (500mL) at pH 2 was then loaded onto the cartridge and filtered dropwise. The cartridge was then washed with 3mL of 40% methanol in DW to remove organic interferences, re-equilibrated with 3mL DW, washed with 3mL 10% methanol/2% NH₄OH to remove humic interferences and finally 6mL 10% methanol/90% MTBE. The final matrix was then evaporated to 200 microliter.

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The samples were then analyzed using a 5890 Series gas chromatograph equipped with an autosampler and a 5970 mass spectrometry detector (Hewlett-Packard, USA). Analytes were separated

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using a SGE HT5 column of 25m x 0.22mm with a film thickness of 0.1 micron. The temperature program was: 50°C, hold 2 min, rate 8°C min⁻¹ to 350°C, hold 30 seconds. Helium was used as a carrier gas at a flowrate of 2 ml/min. The injector temperature was set at 270°C. The MS was operated in the electron impact ionisation mode (70eV). The transfer line and ion source temperatures were 290°C and 220°C, respectively, and the quadrupole was not heated. Scan runs were made with a range from m/z 33 to 500. The chromatograms were analysed using the NIST05 library and the compound was deemed identified if the match percentage was higher than 70%. A retention index (RI) was attributed to the unidentified peaks according to Van den Dool and Kratz (Van Den Doole and Kratz, 1963):

$$RI = 100 \cdot \left[z + \frac{t_i - t_z}{t_{z+1} - t_z} \right]$$

where t_z and t_{z+1} are retention times of the reference n-alkanes hydrocarbons eluting immediately before and after the chemical compound i . Table 1 lists the retention times of the n-alkane hydrocarbon standard used in the Van den Doole-Kartz retention index equation. Regarding the concentrations of the analytes, two sets of standards were run with o-hydroxybiphenyl and bis(2-ethylhexyl)phthalate which are components detected in this study. Standards of 20, 2 and 0.2 mg/L were used to obtain the calibration with a coefficient of correlation R^2 greater than 0.99 in both cases. The limit of detection of the GC-MS used in this study was 50 ng/L.

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2.2.7. Particle Size Distribution

Particle size measurements were made using a Malvern Instruments Particle Size Analyser Model 2600C with a helium neon laser. The standard deviation was within $\pm 5\%$. The value reported for this parameter is the $d(4,3)$ value, which is the volume mean diameter of particles, which takes into account the number and volume of particles in a particular medium. This is different from the Sauter mean $d(3,2)$ which can be defined as the diameter of a sphere that has the same volume/surface area ratio as a particle of interest.

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2.2.8. **Heavy Metals**

All the glassware was washed with Aqua Regia (1 volume of concentrated HNO₃ with 3 volumes of concentrated HCl) prior to the experiment. For the determination of heavy metals in a solid sample, Aqua Regia was used to digest a pre-weighted sample. The solution was heated for 1 hour with a watch glass and was not allowed to boil. After cooling, the watch glass and the beaker were rinsed with distilled water and the final solution was filtered through a 0.45 micron filter. A blank with deionized water was run following the same procedure. The sample and blank were analyzed with PerkinElmer Optima 2000 DV Inductively Coupled Plasma-Optical Emission Spectrometer (ICP-OES) in triplicate. The coefficient of variation is different for each element and is given along with the results. The detection limit is given in Table 2 for each element.

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Table 1. Retention times of standard hydrocarbons detected by the GC-MS used in this study

Retention time (min)	Hydrocarbons	Formula
6.568	decane	C ₁₀ H ₂₂
7.256	undecane	C ₁₁ H ₂₄
9.219	dodecane	C ₁₂ H ₂₆
11.083	tridecane	C ₁₃ H ₂₈
12.866	tetradecane	C ₁₄ H ₃₀
14.532	pentadecane	C ₁₅ H ₃₂
16.109	hexadecane	C ₁₆ H ₃₄
17.614	heptadecane	C ₁₇ H ₃₆
19.014	octadecane	C ₁₈ H ₃₈
20.44	nonadecane	C ₁₉ H ₄₀
21.737	eicosane	C ₂₀ H ₄₂
23.028	heneicosane	C ₂₁ H ₄₄
24.16	docosane	C ₂₂ H ₄₆
25.194	tricosane	C ₂₃ H ₄₈
26.338	tetracosane	C ₂₄ H ₅₀
27.337	pentacosane	C ₂₅ H ₅₂
28.355	hexacosane	C ₂₆ H ₅₄
29.33	heptacosane	C ₂₇ H ₅₆

Retention time (min)	Hydrocarbons	Formula
30.267	octacosane	C ₂₈ H ₅₈
31.174	nonacosane	C ₂₉ H ₆₀
32.074	triacontane	C ₃₀ H ₆₂

Table 2. Detection limits of heavy metals analyzed by Inductively Coupled Plasma-Optical Emission Spectrometer

Element	Detection Limit (ppb)
Al	10
B	10
Cd	1
Co	2
Cr	2
Cu	4
Fe	1
Mn	1
Mo	5
Ni	5
Pb	10
V	5
W	10
Zn	2
Na	5
K	5
Ca	0.5
Mg	1

2.2.9. Ion Chromatography

The ions Na⁺, K⁺, Mg⁺⁺, Ca²⁺, Cl⁻, NO₂⁻, NO₃⁻, PO₄³⁻, SO₄²⁻ were analysed using the DIONEX Ion Chromatograph as described by the American Public Health Association (APHA, 1999). The coefficient of variation for the ions listed above are respectively 2, 0.2, 0.5, 0.6, 1.1, 1.8, 1.8, 3.4 and 5.1 for 5 identical samples. The detection limit was between 50 and 100 ppb.

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2.2.10. *Size Exclusion Chromatography (SEC)*

For size exclusion chromatography (SEC), an Aquagel OH-40 column (Polymer Labs) was used with deionised water as eluent delivered at a flowrate of 0.5 mL/min. The sample volume was 50 microliter, and the column was maintained at ambient temperature with a Shimadzu UV detector set at 254 nm. Unbranched standards of polyethylene oxide (PEO) and polyethylene glycol (PEG) and glucose were used to calibrate the system, hence the results obtained are quoted relative to these linear standards detected by a Shimadzu refractive index detector.

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2.2.11. *Scanning Electron Microscope - Energy Dispersive X-ray (SEM-EDX)*

Fouled membranes and sludge samples were analyzed by scanning electron microscopy (SEM) and energy dispersive X-ray (EDX) spectroscopy. SEM samples were fixed overnight at 4°C in 3% glutaraldehyde and kept at pH 7.2 by a 0.1M phosphate buffer. Samples were then dehydrated in a graded ethanol/water series (10-30-50-70-90-100%) for 20 minutes at each concentration, and then dried for a day at 30°C. Samples were sputtered-coated with gold or carbon (30mA for 2.5 minutes, vacuum 0.2 Torr) prior to SEM/EDX analyses. Specimens were examined and photographed under a scanning electron microscopy (JEOL JSM-5610LV). Energy Dispersive X-Ray analysis was conducted with an EDX-60 (Oxford instrument-incax-sight). The EDX analyser was connected to a scanning electron microscope (model JSM-840A). More than one point on the sample was analyzed by EDX. The exact number of replicates is given along with the results.

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2.3. Physico-Chemical Methods

2.3.1. *Alkalinity*

The alkalinity of wastewater is based on its acid-neutralizing capacity, and is the sum of all the titratable bases. The measured value may vary significantly with the end-point pH used. The method used in this chapter

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followed the procedure described in Standard Methods (APHA, 1999). Alkalinity of samples were titrated potentiometrically to an end-point of pH 4.5 with 0.1 N H₂SO₄. The sulphuric acid solution was prepared by dissolving 2.8 mL of concentrated sulphuric acid in 1L of deionised water. The coefficient of variation for 10 identical samples was within 2.7%. The calculation of alkalinity is shown below:

$$\text{Alkalinity (as mg CaCO}_3\text{/L)} = \frac{A \times N \times 50,000}{\text{mL sample}} \quad (1)$$

where: A= mL standard acid used, N = normality of standard acid.

2.3.2. **Total Solids, Volatile Solids and Fixed Solids**

The measurement of Total Solids (TS) and Volatile Solids (VS) was applied as described in Standard Methods (APHA, 1999). Empty aluminium dishes were placed in a furnace at 550°C for 1 hour. After cooling in a dessicator, the sample was injected into the pre-weighted dish, and the liquid fraction evaporated overnight in an oven at 103-105°C until the weight stabilised. The resulting weight was recorded for the measurement of TS and then the dish was placed in a furnace at 550°C for 1 hour. For VS measurement, the final weight was then recorded on removal from the furnace. Calculations were then performed as given in Standard Methods (APHA, 1999). Fixed solids (FS) or ash percentage was simply obtained by difference. The coefficient of variation for 10 identical samples was within 4%.

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2.3.3. **Total Suspended Solids and Volatile Suspended Solids**

The measurement of Total Suspended Solids (TSS) and Volatile Suspended Solids (VSS) was carried out according to Sections 2540-B and 2540-E of Standard Methods (APHA, 1999). Firstly, three glass fiber filters (GFC, Whatman) were washed by filtering 20 mL deionized 3 times. The filters were placed on aluminium dishes and placed in a furnace at 550°C for 1 hour. They were then placed in a dessicator until needed. To measure TSS, a known volume of sludge was filtered through the filter, and the liquid fraction was evaporated overnight in an oven at 103-105°C.

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They were then allowed to cool down in a desiccator. The resulting weight was recorded for the measurement of TSS and then the tray was placed in a furnace at 550°C for 1 hour. For VSS measurement, the final weight was then recorded on removal from the furnace. Fixed Suspended Solids was obtained by subtracting VSS from TSS. Calculations were then performed as given in Standard Methods (APHA, 1999). The coefficient of variation (COV) for ten identical samples was 4%, 3.1% and 7.1% for TSS, VSS and FSS, respectively.

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2.3.4. COD Measurement

The measurement of Chemical Oxygen Demand (COD) was based on the Standard Closed Reflux Colorimetric Method described in Section 5220-D of Standard Methods (APHA, 1999). Digestion solution was first prepared by adding 10.216 g of $K_2Cr_2O_7$ (Merck), previously dried overnight at 103°C, 167 ml of concentrated H_2SO_4 (Merck, UK) and 33.3 g of $HgSO_4$ (Merck, UK) into 500 ml of distilled water. The mixture was then left to cool at room temperature before diluting to 1000 ml. Samples of 1 ml were added to a Hach reflux tube, followed by 0.6 ml of digestion solution. Then 1.4 ml of sulphuric acid reagent (2.5% w/w silver sulphate in sulphuric acid) was carefully run down the inside of the tube so that an acid layer was formed under the sample/digestion solution layer. The tubes were tightly sealed and inverted three times to mix properly. The mixtures were then refluxed in a Hach COD reflux reactor (Model 45600) at 150°C for 2 hours. After cooling, the samples were analysed on a Shimadzu UV/VIS scanning spectrophotometer (Model UV-2101/3101 PC) at a wavelength of 600 nm. Potassium hydrogen phthalate (KHP) (Merck, UK) was used to prepare standard solutions in the range of 20-900 mg/l. KHP has a theoretical COD of 1.176 mg O_2 /mg.

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Soluble COD (SCOD) was obtained by centrifuging the samples at 10,000 rpm for 5 minutes (Biofuge Stratos, Heraeus Instruments), then at 17,000 rpm for 20 minutes to ensure an easy filtration through a 0.2 micron filter to remove fine suspended material and any residual biomass. The coefficient of variance for ten identical samples was $\pm 2.6\%$.

Total COD (TCOD) was obtained by taking 1ml of sample and diluting it to 100 ml with deionised water. Although the sample was blended for 1 minute in a 1L Waring blender, the presence of particles in suspension made it difficult to take representative samples. The coefficient of variance for ten identical samples was 9.9%.

2.3.5. **Ammonia** Measurement

The measurement of ammonia was carried out according to the Nesslerization method in Standard Methods (APHA, 1999). A sample of 0.25 ml was diluted with deionized water to 25 mL in a volumetric flask. Three drops of mineral stabilizer (Hach Lange GmbH) were added followed by three drops of polyvinyl alcohol dispersing agent (Hach Lange GmbH). The mixture was then mixed by inverting the flask. One milliliter of Nessler reagent (VWR) was then added and the yellow colour was then analyzed after ten minutes on the spectrophotometer at a wavelength of 495nm. The coefficient of variance for ten identical samples was $\pm 6.6\%$.

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2.3.6. **Total Nitrogen** Measurement

The measurement of total Nitrogen was carried out according to the TN kit manual from Hach Lange GmbH. An alkaline persulfate digestion converted all forms of nitrogen to nitrate. Sodium metabisulfite was added after the digestion to eliminate halogen oxide interferences. Nitrate then reacts with chromotropic acid under strongly acidic conditions to form a yellow complex with an absorbance maximum at 410 nm. The coefficient of variance for ten identical samples was $\pm 5.5\%$.

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2.3.7. **Total Phosphorus** Measurement

For the measurement of Total Phosphorus (TP) a known mass of sample was digested in a mixture of hydrochloric and nitric acids (3:1) at 150°C until no solid particles were visible. The digested acid mixture was then transferred to a 50 mL flask, 1 drop of phenolphthalein indicator solution was added and sodium hydroxyde was added until a faint pink color appeared. Deionized water was added to reach the 50 ml mark. The

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measurement of TP was then carried out according to the vanadomolybdophosphoric acid colorimetric method described in Standard Methods (APHA, 1999): in a dilute orthophosphate solution, ammonium molybdate reacts under acid conditions to form a heteropoly acid, molybdophosphoric acid. In the presence of vanadium, yellow vanadomolybdophosphoric acid is formed. The intensity of the yellow colour is proportional to phosphate concentration. ~~the~~ The absorbance was read on a spectrophotometer at 470 nm. The coefficient of variance for ten identical samples was $\pm 0.6\%$.

2.3.8. **Carbohydrates** Measurement

Carbohydrates are hydrolysed by sulphuric acid; dextrose and phenol, in the presence of sulphuric acid, react further to produce a colored complex that is measured spectrophotometrically. Samples of 0.4 ml were added to each test tube and then mixed well with 0.4 ml of 5% (w/v) phenol (Sigma, UK). Two ml of H_2SO_4 was added to each test tube and left at room temperature (18-26°C) for 20-30 minutes to allow color development. The content of each tube was transferred to cuvettes and the samples were analysed on a Shimadzu UV/VIS scanning spectrophotometer (Model UV-2101/3101 PC) against the blank at a wavelength of 485 nm (Dubois et al., 1956). Dextrose was used to prepare standard solutions in the range of 0 - 200 mg/l, but the method gives a linear relationship for some five carbon sugars such as D-xylose and D-arabinose. To convert into COD, 1g carbohydrates assumed as $C_6H_{12}O_6$ is equivalent 1.07 g COD (Sanders, 2001). The coefficient of variance was within 2.5% for ten identical samples.

2.3.9. **Protein** Measurement

A Protein kit (Stephenson's modification) was purchased from Sigma, England (Kit no. P5656). Samples of 1 ml were added to each tube with 1 ml of Lowry Reagent Solution. The samples were mixed well and kept at room temperature for 20 minutes. With rapid and immediate mixing, 0.5 ml of Folin and Ciocalteu's Phenol reagent was added to each tube and mixed well after addition. These samples were left at room temperature for

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30 minutes. The content of each tube was transferred to cuvettes and the samples were analysed on a Shimadzu UV/VIS scanning spectrophotometer (Model UV-2101/3101 PC) against the blank at a wavelength of 725 nm. The detection limit was 5 mg/L, and the coefficient of variance within 3.6% for 10 identical samples. As the precise chemical formula of the proteins detected was not determined, the percentage of soluble COD represented by protein had to be estimated by assuming a stoichiometric conversion factor of 1.5 which is derived from the typical formula of proteins ($C_{16}H_{24}O_5N_4$) presented in Rittmann and McCarty (Rittman and McCarty, 2001).

2.3.10. *Biochemical Methane Potential Test*

The assay was conducted using the media and serum bottle techniques developed by Owen et al. (Owen et al., 1979). The gas used to purge the assay bottles during preparation of the assays was a mixture of 70% N_2 and 30% CO_2 at a flow rate of approximately 0.5 L.min⁻¹. Two mL sample, 14 mL biomedica and 4 mL biomass were transferred into 20-ml serum bottles under anaerobic conditions by continuously flushing the bottles with a mixture of N_2 and CO_2 . Blanks were run with no carbon source and the methane obtained in these control bottles was due to the autolysis of the inoculum. Each serum bottle was then capped with leak proof Teflon seals. Ultimate Methane potential was measured by the amount of methane produced (converted to standard temperature and pressure) minus the methane produced by the blanks and divided by the mass of volatile solids added. Gas production was measured using a glass syringe and the gas was then wasted, whereas gas composition was measured with the GC described above. Duplicate samples were run and the coefficient of variation in the worst case was $\pm 6\%$.

2.4. MOLECULAR BIOLOGY METHODS: **DENATURING GEL GRADIENT ELECTROPHORESIS (DGGE)**

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The DNA from a mixed culture was extracted using the FastDNA SPin for soil kit from MP Biomedicals. The PCR mixture (50 μ L) contained 2 μ L of each primer, 0.2 μ L of the Taq Polymerase (Promega), 2 μ L of dNTPs stock solution (10 μ M), 5 μ L of Taq buffer solution containing $MgCl_2$ and between 1 and 20 μ L of DNA template. The final volume was adjusted to 50 μ L using DNA-free water. Archaeal DNA was amplified using a nested PCR reaction in a G-Storm thermocycler. First, the DNA template was amplified using the primers 46F and 1017R in 25 μ L PCR volume. The temperature program was: initial denaturation 95°C for 3 minutes, then 35 cycles of denaturation at 95°C for 1 minute, annealing at 40°C for 1 minute, elongation at 72°C for 1 minute and the final elongation took place at 72°C for 7 minutes (Akarsubasi et al., 2005; Gray et al., 2002). Then, 1 microliter of the PCR product was used for the second PCR reaction in 50 μ L using the primers 344F-GC and Univ522R. The temperature program was: initial denaturation 95°C for 3 minutes, then 35 cycles of 95°C for 1 minute, 53°C for 1 minute, 72°C for 1 minute and the final elongation took place at 72°C for 7 minutes (Akarsubasi et al., 2005; Gray et al., 2002).

Bacterial DNA was amplified using a single PCR reaction in the same thermocycler. The bacterial DNA was amplified using the primers 341F-GC and 907R (Fernandez et al., 2008; Liu et al., 2008) in 50 μ L PCR volume. The temperature program was: initial denaturation 94°C for 5 minutes, then 30 cycles of 94°C for 1 minute, 52°C for 1 minute, 72°C for 1 minute and the final elongation took place at 72°C for 10 minutes. The DNA specific to ammonia-oxidisers was amplified using the primers CTO189FGC and CTO654R in 50 μ L PCR volume. The temperature program was: initial denaturation 94°C for 1 minutes, then 35 cycles of 92°C for 30 seconds, 57°C for 1 minute, 68°C for 45 seconds and the final elongation took place at 68°C for 5 minutes (Kowalchuk et al., 1998; Stephen et al., 1998).

The DNA specific to fungi was amplified using the primers ITS4 and ITS5 in 50 μ L PCR volume. The temperature program was: initial denaturation 95°C for 1 minutes, then 30 cycles of 95°C for 1 minute, 58°C for 30 seconds, 72°C for 1 minute and the final elongation took place

at 72°C for 10 minutes (White et al., 1990). Presence of PCR products was confirmed by electrophoresis on 1% agarose gels stained with ethidium bromide. DGGE of the 16Sr DNA PCR products was carried out using the DCODETM system (Bio-Rad Laboratories Ltd) according to the manufacturer's instructions and protocols. The percentage of polyacrylamide varied according to the range of base pairs (Table 3) as recommended in the manufacturer's manual:

Table 3. Polyacrylamide concentration in the DGGE vs. the base pair of the PCR products

Gel %	base pair
6%	300-1000 bp
8%	200-400 bp
10%	100-300 bp

This implied that a 6% gel was made up if bacterial primers were used, while a 8% or even a 10% gel was required for archaeal primers. The PCR products were electrophorised at 60V for 16 hours at 65°C with a denaturing gradient ranging from 40 to 60% for bacterial primers (Liu et al., 2008), 45-65% for archeal primers and 38-50% for ammonia-oxidising bacteria (Kowalchuk et al., 1998; Stephen et al., 1998). A 100% solution contains 7 mol/L urea and 40% formamide. The gel was stained in 1 X TAE buffer containing SYBR green before visualizing on a UV transilluminator and photographed. The brightest DGGE bands were cut out, eluted in DNA-free water overnight at 4°C and then re-amplified. The purified PCR product was then cloned with the pCR 2.1-TOPO cloning kit (Invitrogen) according to the manual instructions. For the cloning reaction, 1 µL of vector was mixed carefully with 1 µL of salt solution (Invitrogen) and 4 µL of purified PCR product. Meanwhile chemically competent *E. coli* were placed on ice and 2 µL of the cloning reaction was mixed with the competent cells. The cells were transformed using a heat shock protocol (30 seconds at 42°C), then incubated for 1 hour at 37°C in S.O.C. medium and eventually spread onto Luria-Bertani (LB) agar plates containing 50 µg/mL of Kanamycin to incubate overnight at 37°C. The LB medium consisted of tryptone (10g/L), yeast extract (5g/L), NaCl (10g/L)

and 15g/L of agar in 1L deionized water. The pH was adjusted to 7 and the medium was autoclaved (120°C for 20 min.). Kanamycin was added in a sterile environment. Three well-separated colonies were picked from the plate using a pipette tip and were grown between 12 and 16 hours in LB medium (without agar) at 37°C. The plasmid DNA was then purified using the purification kit from Qiagen (QIAprep Spin Miniprep Kit) and the resulting DNA was then kept frozen until it was sent off for sequencing (Cogenics, UK). DNA sequences analyses were performed using the BLAST server of the National Center for Biotechnology Information (<http://www.ncbi.nlm.nih.gov>).

The Expect value (E) is a parameter that describes the number of hits one can "expect" to see by chance when searching a database of a particular size. It decreases exponentially as the percentage of the match increases. Essentially, the E value describes the random background noise. For example, an E value of 1 assigned to a hit can be interpreted as meaning that in a database of the current size one might expect to see 1 match with a similar score simply by chance. The lower the E-value, or the closer it is to zero, the more "significant" the match is.

2.5. Feedstock Composition and Properties

2.5.1. Composition

The simulated OFMSW used for this study was made up of paper, food wastes and garden wastes. Kitchen wastes (KW) came from a canteen in Southampton University; the left-overs were passed through a kitchen grinder and mixed in a large tank with a drill mixer and then frozen until the experiment. The composition of the simulated paper waste used for the study is listed in Table 4. Paper wastes were kept at room temperature. Garden waste was collected from the Downend Quarry centralised composting site near Fareham (Hampshire, UK) and were kept at 4°C until the experiment.

The OFMSW feedstock used in this study consisted of 41.3% Kitchen Wastes (KW), 10.8% Garden Wastes (GW) and 47.9% Paper Wastes (PW)

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on a wet basis. This composition comes from an assessment scheme for kerbside collection of dry recyclables undertaken by the district of Eastleigh in the UK. It was deemed a representative paper waste composition because Eastleigh has a good mix of socio-economic groups. Moreover, data of KW, GW and PW are given for summer and winter months, allowing us to see the variation (mainly in garden waste) over the seasons and to calculate an annual average. Finally, Eastleigh has a well-established dry recyclables collection of card, newspaper, cans, bottles etc and the data is set out so that actual refuse (Total MSW minus recyclables) is available. This is desirable as legislation says that all city councils in the UK now have to pick up at least two dry recyclables in their waste collections, meaning that Eastleigh refuse MSW is representative of future MSW in the UK. The TS content of the mixture of waste was adjusted with tap water to obtain the OFMSW feedstock at 10% TS. In other words, to prepare 1kg of feedstock at 10%TS, about 830 g of tap water was blended with 18.6g of GW, 71g of KW and 82g of PW.

Table 4. Composition of the simulated paper waste used for the study

Type of paper	%
Newspaper	21.2
Magazine	12
Office paper	7.9
Card and paper packaging	10.5
Cardboard	1.2
Card non packaging	0.6
Liquid carton	1.4
Tissue paper	15.06
Paper plate	15.06
Toilet paper	15.06
TOTAL	100

2.5.2. Properties

2.5.2.1. Physico-Chemical Analysis

The SCOD for the KW, GW and PW was obtained by drying the waste (105°C) overnight, grinding to obtain a powder and placing it in a dessicator for one day. A suspension was thereafter prepared by mixing 2 g of waste per liter in a blender. The suspension was then mixed with a magnetic stirrer for one day at ambient temperature and then filtered through a paper filter (Fisherbrand QT280). The low SCOD/COD ratios in Table 5 show the poor tendency to dissolve and this tendency is even worse (8.4%) for the OFMSW. It is hypothesized that most of the soluble material originates from the KW which has the highest SCOD/COD ratio. Nopharatana et al. (2006) found a SCOD/TCOD ratio equal to 7% in a similar experiment. Moreover, Table 5 reveals that the COD/VS ratio for the feed is about 1.3 g O₂/g VS, which is close to values found in the literature for OFMSW: 1.4 (Hartmann and Ahring, 2005; Nopharatana et al., 2006), 1.284 (Anderson and Saw, 1992). The high ash content of garden waste could be explained by the presence of sand and stones among the organic material.

Table 5. Physical and chemical properties of the simulated OFMSW and its components used in this study. (Standard deviation is indicated in brackets for sample in triplicate). dm = dry matter

	Kitchen Wastes	Garden Wastes	Paper Wastes	Simulated OFMSW
Gravimetric tests				
Total Solids (%)	22.9	33.7	94.4	53.8 (1.3)
Volatile Solids (% of TS)	95.4	66.5	85.5	84.4 (1)
Ash (% of TS)	4.6	33.5	14.5	14.7 (1.2)
COD tests				
COD (mg O ₂ /gdm)	1,369.7 (135)	870.1 (45)	1,330 (61)	1,089.4 (120)
SCOD (mg O ₂ /gdm)	471.2 (5)	71.7 (4)	15.9 (2)	91.1 (7)
COD (mg O ₂ /g VS)	1,462.8 (279)	1,041.5 (88)	1,555.7 (72)	1,331.2 (155)
SCOD (mg O ₂ /g VS)	380.5 (17)	136 (5)	18.5 (2)	111 (10)
SCOD/COD Ratio (%)	34.3 (3)	8.2 (0.6)	1.2 (0.2)	8.4 (1.1)

In addition to COD tests, carbohydrate and protein concentrations were measured on filtered (0.45 µm filter) and unfiltered samples of the simulated OFMSW feedstock. For the unfiltered samples, carbohydrates, on a COD basis, represent 78%, whereas proteins represent only 7%, as shown in Table 6, while the remaining fraction may be aromatic compounds, lipids, humic and fulvic acids, etc. For the filtered samples (≤ 0.45 micron), the proportion of carbohydrates is only 1.7% which implies that carbohydrates from the OFMSW are huge and complex molecules, of which a tiny fraction is composed of small molecules that are solubilised. This is in line with Nopharatana et al. (2006) who found that only 2% of the SCOD was made up of the simple sugars, glucose and maltose, and VFAs like lactic and acetic acids. Table 6 also implies that about 95% of the soluble fraction are not carbohydrates, nor proteins which means that most of the soluble fraction consists of complex molecules which must be broken down into simpler molecules before they can be taken up by microorganisms.

Table 6. Concentration and percentage of carbohydrates and proteins in total and soluble fractions of OFMSW (standard deviation is indicated in brackets)

Unfiltered sample	Carbohydrates	Proteins	COD
mg/g VS	976.9 (59)	62.3 (11)	1,331.2 (155)
% of TCOD	78.5 (8)	7 (1)	100
Filtered sample ($\leq 0.45\mu\text{m}$)			
mg/g VS	1.8 (0.1)	2.7 (0.2)	111.3 (10)
% of TCOD	1.7 (0.1)	3.7 (0.3)	100

Eventually, other relevant properties of the feedstock such as pH, total nitrogen (TN) and total phosphorus (TP) were also measured in order to characterize the simulated feedstock of the OFMSW. The pH was found to be in the range 6.3 - 6.9. TN and TP were found to be 13.1 and 1.3 mg/g TS, respectively, whereas the respective standard deviations were 0.4 and 0.1.

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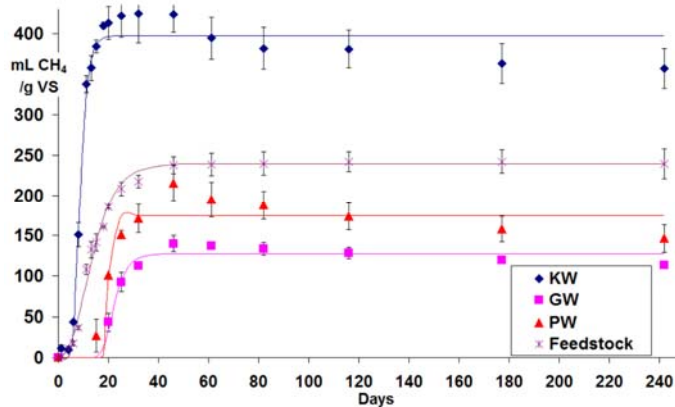
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2.5.2.2. Biochemical Methane Potential Test

The test was carried out with fresh screened ($\leq 500 \mu\text{m}$) anaerobic sludge from a conventional sewage sludge anaerobic digester (Mogden, UK). The biomass TSS concentration was $34.4 \pm 0.6 \text{ g/L}$ and VSS was $21 \pm 0.5 \text{ g/L}$. The BMP test was determined in duplicates in 38 mL serum bottles, with 20 mL of media at an initial I/S ratio of 2.1, except for the feedstock for which it was 0.7. Some authors (Lay et al., 1997; Nopharatana et al., 2006) used the Gompertz equation that describes the cumulative methane production in batch assays.

$$M = P \cdot \exp \left\{ -\exp \left[\frac{R_m \cdot \exp(1)}{P} \cdot (\lambda - t) + 1 \right] \right\} \quad (2)$$

where: M is the cumulative methane production (mL), P is the methane production potential (mL), R_m the maximum methane production rate ($\text{mL} \cdot \text{d}^{-1}$), λ is the duration of lag phase (d) and t is the duration of the assay at which cumulative methane production M is calculated (d). The parameters P, λ and R_m were estimated by applying a least squares fit of the above equation to the experimental data set using the Solver in Excel.



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Figure 5. Biochemical Methane Potential on the components and the feedstock of OFMSW used in this study. markers: average of measurements; line: fitted using Gompertz Equation; error bars: standard deviation of triplicates.

Figure 5 shows that all wastes display a lag phase due to the non-acclimated inoculum used for the experiment. The biostabilization of the organic substrate in anaerobic batch reactors is known to display a slow phase which is the necessary period for adaptation and beginning of bacterial mass multiplication, and a rapid decomposition phase when the bacterial growth is the highest (Lopes et al., 2004). The period of the lag phase is mainly influenced by the initial concentration of microorganisms and substances stimulating the growth of bacteria (Lay et al., 1997). The lag phase, λ , however, is shorter for KW (5.8 days) and Feedstock (4.5 days) than for PW (18.4 days) and GW (17.5 days). This lag phase may be explained by the close association between cellulose and lignin in PW and GW, which impedes access to cellulose. This association might prevent bacteria having direct access to hydrolyzable material which could explain why the blank bottles produced more gas at the very beginning of the test. This translated to negative values for PW and GW during the first twenty days. Interestingly, the lag phase for the Feedstock was less than the one for KW. This is possibly due to the fact that the Feedstock was blended to reduce the particle size and to take a homogeneous sample, and not the KW, or due to a better C/N ratio for the combined waste. The hydrolysis might have thus been enhanced for the feedstock, and gas production might have started earlier.

Kitchen Wastes are readily hydrolysed and converted to methane. The ultimate methane potential was 357 mL CH₄/g VS, which corresponds to 66% (± 7) of the theoretical methane production based on its COD content. After 50 days, the bottle containing KW stopped producing methane while blank bottles were still producing a little bit which explains why the curve decreased. This biodegradability is consistent with Shin et al. (1993), Lee et al. (1997) and Zhang et al. (2007) who found 356, 343 and 348 mL CH₄/g VS, respectively.

Garden wastes are slowly biodegraded due to their high lignin content which is not degradable under anaerobic conditions. The ultimate methane

potential was estimated to be 113 mL CH₄/g VS, which is 33.1% (± 3) of the theoretical methane production based on the COD content of that waste. This value is consistent with the 124 mL CH₄/g VS found by Owens and Chynoweth (1993).

Paper waste is heterogeneous because it is made up of easily degradable paper such as office paper, but also less degradable paper such as newspaper. It is, however, degraded faster than garden wastes and eventually reached 146.6 mL CH₄/g VS, which corresponds to 27.9% (± 2) of the theoretical methane production on a COD basis. The value of ultimate methane production was close to the value reported in the literature (Owens and Chynoweth, 1993).

The feedstock had an intermediate behaviour, as expected because it consisted of a mixture of the three wastes; the ultimate methane potential was estimated to be 239.1 mL CH₄/g VS, which is 45.5% (± 2) of the theoretical methane production based on the COD content of the feedstock. The methane yield was consistent with values reported in the literature. For instance, Nopharatana et al. (2006) found 0.24 $\frac{\text{mL CH}_4}{\text{g VS}}$. As a conclusion to this test, the BMP revealed that 46% of COD of the feedstock was converted to methane, which is close to the value found by Nopharatana et al. (2006) who carried out a BMP test on the OFMSW (48%). Moreover, 87% of the total methane production was obtained in 32 days. Better performance could probably be obtained with an acclimatised biomass. After 177 days, all the bottles contained a gas in the head space with a methane percentage comprised between 50 and 60%. All the experimental results and the Gompertz coefficients are gathered in Table 7.

Table 7. Gompertz coefficients and experimental ultimate biodegradabilities

Waste	Gompertz coefficients			BMP mL CH ₄ /g VS	biodegradability %
	P (mL CH ₄)	R _m (ml CH ₄ /day)	λ (day)		
KW	397.1	72.6	5.8	357.1	66
GW	127.4	13.6	17.5	113.7	33.1
PW	175.2	63.1	18.4	146.6	27.9
Feedstock	237	13.8	4.5	243.8	46

2.5.2.3. Effect of Biomass **Acclimatisation** on the Yield

Another BMP test was carried out on the OFMSW to determine the influence of acclimatisation on the methane production rate. The acclimatised biomass was taken from a 4L chemostat batch-fed with leachate from OFMSW and operating at 20 days HRT; the mixing was stopped 2 days prior to effluent removal in order to let the biomass settle and to keep it within the chemostat. The leachate was produced in another 4L chemostat batch fed with the OFMSW at an OLR of 1 g VS/L.day. The unacclimatised biomass was taken from a drum containing anaerobic sludge from a conventional sewage sludge anaerobic digester (Mogden, UK). In order to compare the two inocula, their VSS content was adjusted to 0.5 g VSS/L and the BMP test was conducted in duplicates at an initial I/S ratio of 0.25.

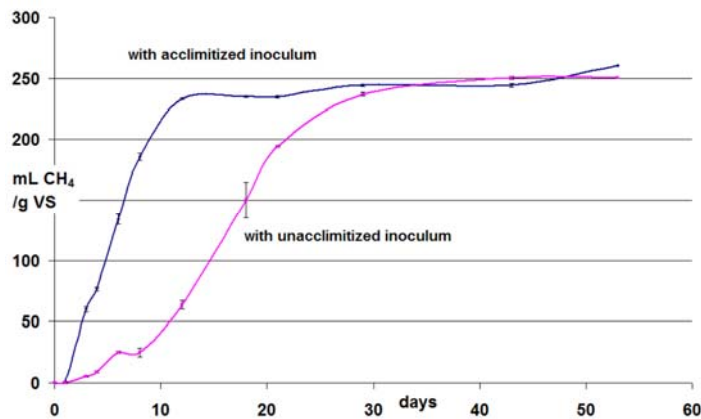


Figure 6. Effect of the acclimatisation of the biomass on the Biochemical Methane Potential test of OFMSW. The error bars show the standard deviation.

Figure 6 shows that an acclimatised biomass resulted in a similar methane potential. However, the lag phase (obtained by ~~the Gompertz~~ equation) was found to be significantly shorter for the acclimatised

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biomass: 1.4 days versus 7 days. It is suggested that the acclimatised biomass includes microorganisms producing more enzymes compared to the unacclimatised ones. It can be seen that the unacclimatised inoculum gives a sluggish methane production rate but eventually both curves reached similar ultimate biodegradability circa 250 ml CH₄/g VS. Based on the COD content, the biodegradability was 49.6 and 47.7% for the acclimatised and the unacclimatised inocula, respectively. The biochemical methane potential assay can be used to determine the hydrolysis constant if the biodegradation kinetics is assumed to follow a first-order reaction according to (Lee et al., 2009):

$$r = -k \cdot S \quad (3)$$

where k is the reaction rate constant (d⁻¹), S is the substrate concentration (g VS/L) and t is the time (days). Integrating this equation with initial and final boundary conditions,

$$\ln\left(\frac{S}{S_0}\right) = -k \cdot t \quad (4)$$

The production of methane is related to the substrate degradation by

$$\frac{S}{S_0} = \frac{Y_{max} - Y}{Y_{max}} \quad (5)$$

where Y is the cumulative methane yield at time t and Y_{max} is the ultimate methane yield. From the two equations above:

$$\ln\left(\frac{Y_{max} - Y}{Y_{max}}\right) = k \cdot t \quad (6)$$

Using the data from the BMP using acclimatised inoculum and by plotting the equation above, the slope gave the value of the hydrolysis constant as 0.21 d⁻¹ ($R^2 = 0.976$). Thus the kinetics of gas production is satisfactorily predicted with the first-order model.

2.5.2.4. Heavy Metals

The **heavy metals** content of the OFMSW used in this study are listed in Table 8. The high content of calcium, iron, sodium and potassium can be pointed out in this table. These are presumably due to KW. Zhang et al. (2007) carried out an elemental analysis of food waste and found concentrations of 21,600 ppm calcium, 766 ppm iron, 900 ppm potassium and 1,400 ppm magnesium.

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Table 8. Heavy metal concentrations in the simulated MSW (standard deviation is indicated in brackets for sample in triplicate)

Element	ppm of TS
Ca	38,698.66 (4733.6)
Fe	2,325.39 (266.3)
Na	2,256.59 (265.5)
K	1,848.02 (208.5)
Mg	821.34 (93.9)
Zn	588.17 (66.3)
Al	551.94 (66.9)
Mn	88.75 (10)
Cu	47.79 (5.4)
Co	45.16 (5.1)
B	25.73 (4.4)
Pb	8.93 (3.6)
W	6.83 (0.9)
Ni	4.73 (0.9)
Cr	4.2 (0.5)
Cd	2.1 (0.2)
V	< 2.6
Mo	< 2.6

3. RESULTS AND DISCUSSION

3.1. Performance of the **Hydrolytic Reactor**

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3.1.1. Effect of the OLR on VS Removal

The TCOD in the leachate varied over a wide range, between 4000 and 26,000 mg/l, possibly due to the HR being fed every two days until day 159, intermittent mixing, and due to occasional stirring difficulties. It can be seen from Figure 5-7 that the TCOD did not change with changes in OLRs from 0.5 to 16 g VS/L.day. However, the value of TCOD did depend upon the occasional presence of solid particles in the sampling line at the time of sampling. Similarly, the SCOD did not vary significantly when a step increase in OLR was effected in the HR, and was always in the range 530 – 2,900 mg/L. The low SCOD observed in the leachate is thought to be due to poor hydrolysis because of the ligno-cellulosic type of feedstock, and an inadequate amount of inoculum used to seed the HR.

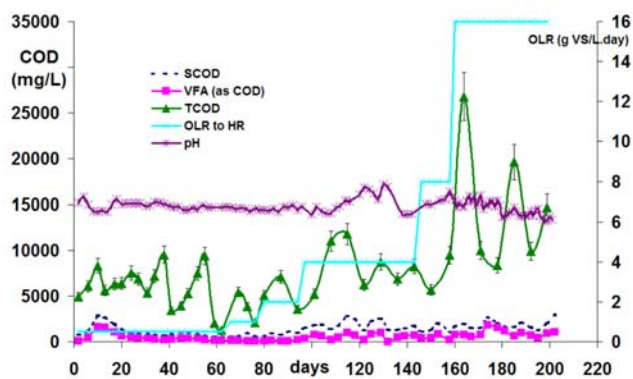


Figure 7. Evolution with time of TCOD, SCOD and VFAs in the effluent of the HR (left axis), and OLR and pH (right axis). Reprinted from (Trzcinski and Stuckey, 2009b).

The initial inoculum to substrate ratio (I/S) may also play a role during start-up of anaerobic digesters. The initial I/S ratio was 0.02 based on the initial load of 340 g volatile solids fed during start up. Then the HR was fed continuously at an OLR of 0.5 g VS/L.day but with intermittent mixing as well as occasional stirring difficulties at TS above 5%. Table 9 presents

the VS removal percentages at the various OLRs and HRTs tested. The VS removal percentage was calculated as follows:

$$\text{VS removal\%} = 100\% \left(1 - \frac{\text{mass VS removed} + \text{mass VS accumulated in HR}}{\text{mass VS fed in HR}} \right) \quad (7)$$

Where the masses were considered over a period longer than 15 days so that steady-state can be assumed and the mass of VS accumulated in the HR is the difference between the mass of VS in the HR at the beginning and the end of the period considered. The VS removal percentages shown in Table 9 are 65.4, 43.8, 35.5, 22 and 13.8% VS destruction at 0.5, 2, 4, 8 and 16 g VS/L.day, respectively, assuming that the volatile solids production due to bacterial growth and the transfer of volatile solids to the SAMBR were negligible.

The transfer of volatile solids to the SAMBR was very limited thanks to the separation between coarse solids and leachate by the perforated stainless steel mesh within the HR. Nevertheless, a small fraction of solids could still pass through and be pumped to the SAMBRs. At the end of the experiment large holes and rust were observed in the 50 micron stainless steel presumably due to a low quality steel or due to pieces of wood in the GW that may have pierced the mesh. This transferred VS fraction over 200 days was estimated as 37.8 and 69.3 g VS for SAMBR1 and SAMBR2, respectively, which can be considered as negligible. For instance, during the period at 16g VS/L.day (day 159 to day 199) the total VS mass transferred to SAMBR1 and SAMBR2 together equalled 91 g changing the VS removal % in the HR to 12.4 instead of 13.8. The former is the actual VS removal in the HR, while the later could be named the "apparent VS removal" and in this study they were similar and thus the difference was neglected. The low VS removal percentages were also due the low volatile solids retention times calculated as the ratio of mass of volatile solids in the HR that is equal to $X \cdot V$, where X is the VS concentration in g/L in the HR and V is the HR working volume in L, and the mass of volatile solids removed from the HR per day (W in g VS/day):

$$\text{VS Retention Time (days)} = \frac{x.V}{w} \quad (8)$$

Consequently, the anaerobic biodegradability of the digestate that was taken out of the HR was consistent with the lower VS removal observed as the OLR was increased. The BMP of the digestate was carried out at an I/S of 0.1-0.2 using the acclimatised sludge from SAMBR2 as inoculum, and the ultimate biodegradability was found to be 167.7, 229.7 and 296.6 mL CH₄/g VS fed at OLRs of 0.5, 8 and 16 g VS/L.day, respectively.

Table 9. Comparison of volatile solids retention times, volatile solids removal percentages, fresh water consumption, hydraulic retention times, percentages of fresh water addition compared to recycled process water and digestate, methane potential at different organic loading rates in the hydrolytic reactor. n.a. = not applicable

OLR (g VS/L.day)	0.5	2	4	8	16
Duration (days)	63	17	47	14	40
VS RT (days)	67.8	49	16.6	6.4	3.3
VS removal (%)	65.4	43.8	35.5	22	13.8
Average fresh water consumption (mL/day)	3.7	n.a.	68	202	652
HRT (days)	15	9	7.8	4	2.2
% of fresh water added compared to recycled process water	0.6	n.a.	5	8	14
Digestate methane Potential (mL CH ₄ /g VS)	167±6.2	n.a.	n.a.	229.7±6.9	296.6±24

Table 9 also contains the HRT of the HR, i.e., the **hydraulic retention time** or leachate retention time, which is the average retention time of a unit volume of liquid in the reactor and is calculated as the ratio of the reactor volume and the leachate flowrate to the SAMBRs. Obviously, the lower the HRT, the higher the flowrate of leachate to the SAMBRs and the more likely the **acidification** of the HR can take place because methanogens do not have time to grow in sufficient amount to degrade the VFAs in the HR. However, it is possible that the mesh inside the HR acts as a bacterial support where **methanogens** can attach and form colonies with acidogens so that they can survive on VFAs produced in the close

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environment. Figure 6-8 shows the methane content of the biogas produced in each reactor. It can be seen that methane started to be produced very slowly compared to the SAMBR. The methane percentage increased gradually to circa 40% around day 100, but on day 112 the HR was opened and the 50 microns macrofilter was removed and replaced by a new one which let some air into in the headspace. As a result, the methane content dropped and remained low as the reactor had to be opened more frequently due to technical issues such as mixing problems or solids that could not be removed from the bottom pipe due to the high TS content.

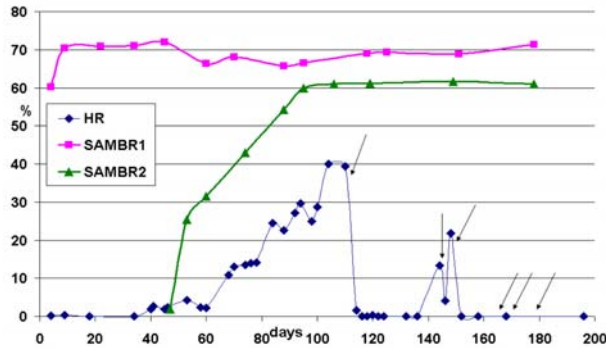


Figure 8. Evolution of the methane content in the HR, SAMBR1 and SAMBR2. The arrows show when the HR was opened for maintenance work on the macrofilter.

3.1.2. Effect of the OLR on VFAs Production and Composition

The evolution and composition of VFAs over time in Figure 7-9 shows that acetate was the main VFA at steady-state, but propionate became the main acid after the shock at 4, 8 and 16 g VS/L.day on days 101, 146 and 164, respectively, which is a few days after the organic shocks took place (Trzcinski and Stuckey, 2009b). The HRT also affects the VFAs distribution because it was reduced to 4 and 2 days on day 146 and 164, respectively. From day 160 onwards, the HR was fed every day at 16 g VS/L.day at HRTs ranging from 4 to 1 days, and propionate remained the main acid until the end of the run.

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Gallert et al. (2003) also observed a higher and longer-lasting propionate accumulation when the HRT was reduced from 7.1 days to 5.7 days at an OLR of 15 kg COD/m³.day; they correlated this with 1% hydrogen in the off-gases. Propionate oxidation is known to be the bottle neck reaction during the **methanogenesis** of complex substrates because their growth rate is only 0.13 d⁻¹ (Wallrabenstein et al., 1995), and they can be washed out at a HRT below 8 days (Gallert et al., 2003). The pH dropped between 6 and 6.5 due to the OLR of 16 gVS/L.day applied after day 159, but with the accumulated **alkalinity** (5,000 mg equivalent CaCO₃/L on day 199) and the recycling of the SAMBR1 and AMBR permeates, the pH did not drop any further, which highlights the advantage of recycling the SAMBR **permeate**. A BMP test (I/S = 1.3) on the leachate on day 193 revealed that 57% (± 10%) of the leachate was converted to methane. The rest of the COD was either not hydrolysable or converted to biomass.

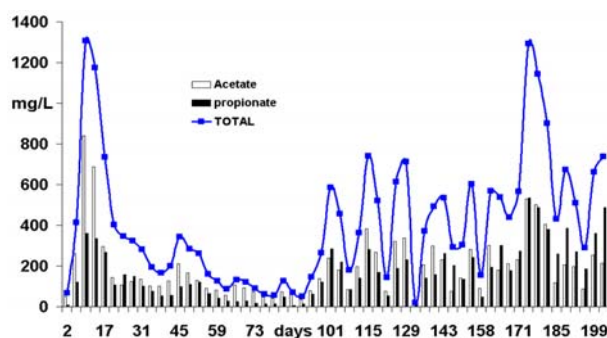


Figure 9. VFA distribution in the effluent of the HR. Reprinted from (Trzcinski and Stuckey, 2009b).

3.1.3. **Scanning Electron Microscopy** of the Fibers in the HR

In order to gain more understanding of the hydrolysis process, a Scanning Electron Microscope was used to visualize the fibres of the OFMSW. Since the KW is easily degraded, and since GW represented a small percentage of the feedstock, it can be assumed that the fibres

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originated from the PW. Different magnifications were taken of both the virgin fibres taken from the feedstock, and the fibres from the HR after anaerobic digestion. All the pictures are shown in Figure 810. The virgin fibres are on the left-hand side (Pictures A, C, E and G), and the treated fibres are on the right-hand side (Pictures B, D, F and H). The pictures show evidence that the fibres are being peeled off and small nodules looking like bacteria could be seen on the surface (Figure 911). The surface is much rougher than virgin fibres which is due to bacterial hydrolysis that targets the cellulose and hemicellulose from the fibrils, but it could also be due to the shear of the stirrer in the HR. Figures B and D show that pieces up to 10 microns can be peeled from the main fibre, but still remain attached to it. This could be the result of a partial hydrolysis of cellulose that was consumed until only lignin remained accessible to enzymes. Some holes were observed in the virgin fibres of picture E. This could be specific to a certain type of fibre (cardboard, newspaper, office paper, etc.) or it could be the result of solubilisation of hemicellulose during the autoclaving of the virgin fibres. Solubilisation of hemicellulose has indeed been reported possible in hot water (Hendriks and Zeeman, 2009). Liao et al. (2004) also observed holes in manure fibres before acid treatment and they attributed this to a partial degradation of cellulose.

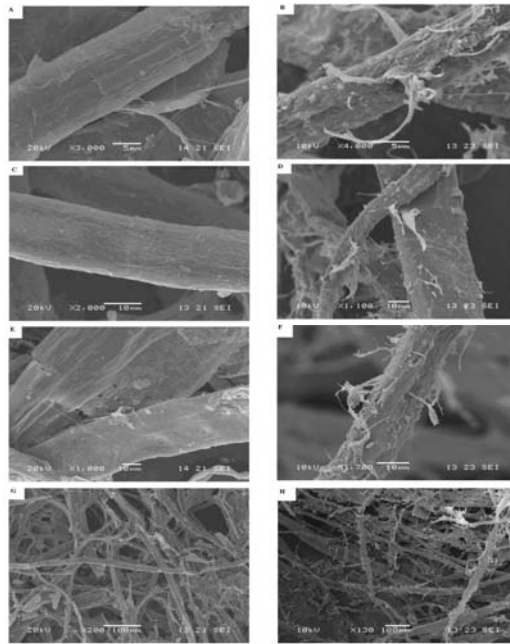


Figure 10. Scanning Electron Microscopy of the raw fibres from the OFMSW feedstock (left) and treated fibres from the HR (right).

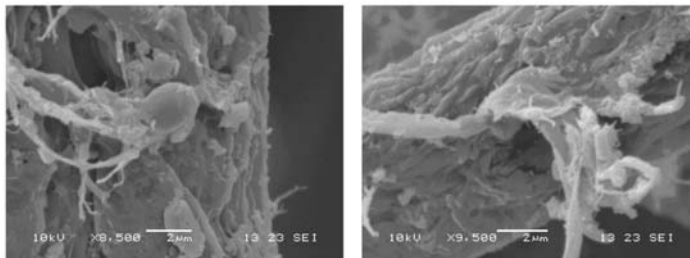


Figure 11. Close zoom on the fibres treated in the HR. This is probably bacterial aggregates attached to the fibre.

3.2. Performance of SAMBR1

3.2.1. Effect of the HRT on COD Removal

The OLR to the SAMBR was not constant because of the fluctuations in the TCOD of the leachate from the HR (Figure 57), and as a result the SCOD in SAMBR1 (Figure 4012) sometimes increased sharply over time. For instance, an OLR to the SAMBR of 8 g COD/L.day was observed temporarily on day 164, and a decrease of the HRT to 2.1 days led to a sharp peak of SCOD in the reactor but not due to VFAs build-up, indicating that the hydrolysis was rate limiting. On day 185, a temporary OLR of 19.8 g COD/L.day was also observed, but the stability of SAMBR1 at such a high OLR could not be assessed properly due to the transient character of the OLR. Despite the varying OLR, the permeate SCOD (effluent SCOD in Figure 4012) remained typically in the range 300-500 mg /L. A build-up of the SCOD was not observed and the SCOD was even found to decrease slightly on few occasions. This can be partly attributed to the greater consumption of fresh water towards the end of the run to keep up the volume in the HR (see Table 9), but the decline of SCOD was also due to the very high MLTSS (28.7 g/L) at the end of the run, and was not due to the enhanced rejection by the membrane because the SCOD in the bulk liquid was also found to decrease slowly.

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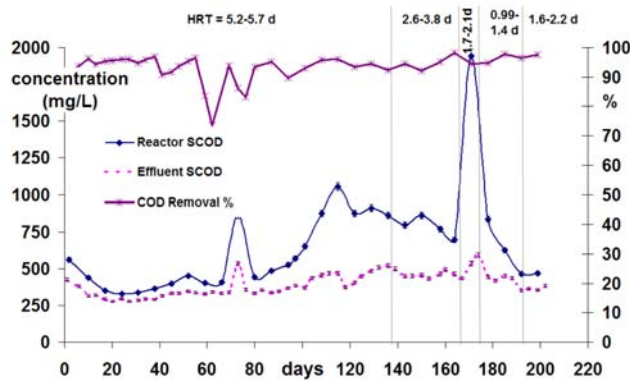


Figure 12. SCOD and VFAs inside SAMBR1 and in its permeate (left axis). COD removal in SAMBR1 (right axis). Reprinted from (Trzcinski and Stuckey, 2009b).

3.2.2. Effect of the TSS on the Flux

The COD removal remained circa 95% throughout the run, while the VFA concentration was virtually zero, suggesting that the methanogenic population could cope with an HRT as low as 1 day. However, SAMBR1 could not be operated in a sustainable way at a HRT below 1.6-2.3 days due to a membrane flux limitation at 0.54-0.78 LMH. At an HRT below 2 days, the rate of particulate COD destruction became less than the feeding rate, resulting in the build-up of solids at the bottom of the reactor which eventually blocked the diffuser and, on day 182 no more bubbles were scouring the membrane. At the same time, the MLTSS increased to 28.7 g/L (Figure 4+13) which also badly affected the flux. This indicates that the performance of the SAMBR treating leachate containing particles was limited to 1.6-2.3 days HRT by the hydrolysis and not the VFA degradation. Interestingly, as the HRT dropped the inorganics in suspension followed the same trend as TSS and VSS with a sharp increase to 13.6 g/L showing that at HRTs lower than 1 day the FSS build up in SAMBR. Again because the hydrolysis of solids is too slow compared to the feeding rate, the inorganics bound to the ligno-cellulosic matrix are not solubilised fast enough and build up. As a result the VSS/TSS ratio

plummeted to circa 50%. The transmembrane pressure (TMP) in SAMBR1 was virtually zero until day 60.

SAMBR1 was treating leachate from day 0 to day 60 at a HRT of 5.5 day, at a stable flux of 8.5 LMH and a MLTSS around 5 g/l with virtually zero TMP. The MLTSS could remain that low because the HRT was 5.5 days which provide enough time for the hydrolysis to proceed without noticeable build up in MLTSS. On day 24, it was attempted to increase the flux by increasing the speed of the recycling pump but that led to a sharp increase in TMP to 450 mbar which indicates that operation over the critical flux was taking place and that a cake layer was forming. Based on the step increase in the recycling pump setting, it could be approximated that the flux was near 11 LMH which was thus deemed over the critical flux. Between day 76 and day 105, the effect of the sparging rate on the COD rejection was investigated using TMP data which were analysed in the corresponding section. After day 105, the flux in SAMBR1 remained relatively stable at 3.5 LMH although the MLTSS varied over a wide range (4 - 19 g/L). This can be explained by the increase in sparging rate to 10 LPM that results in the resuspension of solids that settled down in the reactor at low sparging rate.

On a few occasions, however, the sparging rate had to be reduced due to foaming problems which caused the solids in suspension to settle down and this resulted in a lower apparent MLTSS. With this in mind, the great variation in MLTSS should not be seen as a result of changing performance in the reactor but rather as non-homogeneous sampling for MLTSS measurement. The flux of 3.5 LMH was accompanied by TMP values of up to 500 mbar whereas the operation after day 170 at a MLTSS of 28.7 was characterized by very low flux circa 0.5 LMH and very high TMPs in the range 550 - 800 mbar (data not shown). Normal operation of the Kubota membrane is associated with fluxes in the range 20-27 LMH at a maximum of 15 mbar TMP and 20 g TSS/L (Ramphao et al., 2004), but Van Zyl et al. (2007) reported a flux of 2-3 LMH at 36 g TSS/L when treating synthetic petrochemical effluent which is similar to what was observed in this study although their TMP was much lower (50 mbar) probably due to the feed containing only soluble organics. At such high

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MLTSS, it is thought that the diffuser was blocked due to the solids settling at the bottom of the reactor. This led to a much reduced scouring of the membrane. In order to lower the MLTSS and improve the flux, sludge was removed from SAMBR1 at a mean rate of 90 mL/day after day 180 (SRT=33 days), but this did not help in reducing the MLTSS.

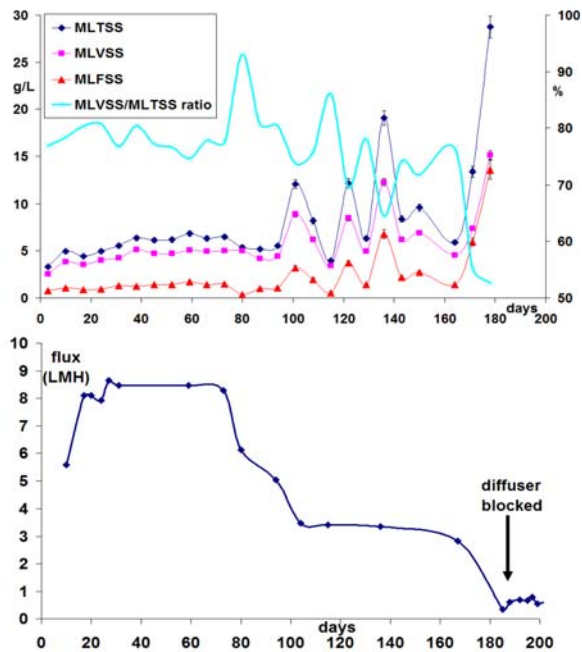


Figure 13. Evolution with time of the TSS, VSS, FSS and VSS/TSS ratio (top), and evolution of the flux in SAMBR1 (bottom).

3.2.3. Effect of the Bulk COD on Membrane Rejection

The SCOD inside the reactor remained higher than the effluent values throughout the experiment, which demonstrates that the presence of a cake/gel layer on the membrane surface improves the effluent quality: this is in line with previous work on the SAMBR (Akram, 2006; Choi and Ng,

2008). Nevertheless, membrane rejection did not increase with time but varied according to the bulk SCOD. The rejection of solute can be due to the dynamic layer formed by the foulant, and the attached biomass acts as a secondary filter. Choi and Ng (2008) observed that solute rejection rate increased with more severe fouling. During fouling, pore blocking and pore narrowing can take place which would enhance solute rejection. They also observed lower TOC in the permeate compared to the supernatant, which they attributed to a combination of biodegradation by the biofilm (cake layer) developed on the membrane surface and further filtration by cake layer and narrowed pores. Hence, lower permeate concentration and faster permeability decline are observed as membranes are fouled. Membrane rejection was expressed as a percentage:

$$\text{Rejection} = 100\% \cdot \frac{SCOD_{bulk} - SCOD_{permeate}}{SCOD_{bulk}} \quad (9)$$

In this study it was observed that the higher the bulk SCOD, the higher the rejection (Figure 14), which suggests that the high molecular weight COD is degraded in the bulk liquid until it becomes small enough to pass through the membrane or the biofilm pores.

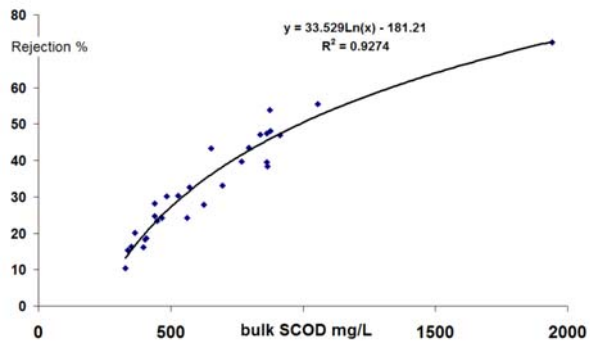


Figure 14. Correlation between the bulk SCOD in SAMBR1 and the membrane rejection. Reprinted from (Trzcinski and Stuckey, 2009b).

3.3. Performance of SAMBR2

3.3.1. Effect of **Inoculum** on Start-Up

Previous studies (Akram, 2006) have shown that a shorter start-up period and higher COD removal in SAMBRs can be obtained by increasing organic load at a lower constant HRT rather than gradually decreasing the HRT at constant high feed strength. This approach was followed to start up a SAMBR, although Akram (2006) used a sucrose-based wastewater that is easily degradable, while the leachate used in this study was lignocellulose-based. For an easily degradable substrate, VFA accumulation can occur in the SAMBR due to the overloading of the methanogens and possibly the lack of syntrophic associations necessary to degrade reduced intermediates. For this reason, prior inoculation into a CSTR is helpful for the development of an active inoculum enriched in methanogens (Akram, 2006). With this in mind, an inoculum was fed on synthetic VFAs as the main carbon source (75% on a COD basis) in a 4 litre chemostat prior to inoculating SAMBR2. Prior to inoculating SAMBR2, a specific acidogenic activity test was conducted on the two different inocula, the one from SAMBR1 and the one from the chemostat batch fed with meat extract, peptone and synthetic VFAs. The test was determined in triplicate in 38 mL serum bottles, with 20 mL of media. The same amount of glucose was fed to both sets of bottles to result in 2 g COD/L for the test at an I/S ratio of 1, and Figure 43-15 reveals that indeed the acidogenic and methanogenic biomass of the inoculum fed with synthetic VFAs was more active than the inoculum taken from SAMBR1 on a same MLVSS basis. Akram (2006) also reported a higher activity from an inoculum taken from a CSTR compared to the SAMBR. The reason for this is the bigger flocks formed in a gently mixed CSTR compared to the SAMBR where the shear rate is more important. Several authors have identified strong mixing conditions and high shear forces as a detrimental factor for VFA degradation since it supposedly affects microbial proximity, and therefore interspecies hydrogen and/or formate transfer (Brockmann and Seyfried, 1997; Kim et al., 2002).

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As a result the smaller flocks in the SAMBR can release more hydrogen due to inefficient interspecies hydrogen transfer, and this results in reduced methane production through the hydrogenotrophic reaction. This can also be due to the non-living fraction of MLVSS in the inoculum from SAMBR1 as the MLVSS measurement does not distinguish the non-living VSS from the living fraction. As a result, the inoculum from SAMBR1 was likely to contain less acidogens and methanogens explaining the slower methane production rate during the first two days. It is interesting to note that eventually the inoculum from SAMBR1 ended up producing significantly more methane than the inoculum from the chemostat. With the inoculum from SAMBR1, 79% (± 0.6) of the glucose was converted to methane, while that percentage reached 72.3% (± 0.9) with the inoculum from the chemostat. This again can be explained by the amount of VSS introduced into the BMP bottles. As the inoculum from SAMBR1 contained non-living VSS consisting of lignocellulosic fibres, the actual living VSS was less than in the inoculum from the chemostat. Therefore less COD was used for biomass growth and the spare COD could then be metabolized to produce more methane. The extra methane produced could also come from the lignocellulosic fibres that were present in the inoculum from SAMBR1, and that were further hydrolyzed in the serum bottles. As the acidogenic test showed that the inoculum from a chemostat fed on meat extract, peptone and synthetic VFAs was more active on a MLVSS basis, it was hypothesized that it would be beneficial for the start-up of the SAMBR in order to avoid VFA accumulation. Also the smaller inoculum required to seed the SAMBR due to a higher methanogenic activity would result in lower MLTSS concentrations for a successful start-up and maybe a better flux. VFA concentrations in SAMBRs 1 and 2 were both virtually zero. This indicates that an inoculum acclimated to VFAs does not bring further advantages regarding VFAs degradation because both SAMBRs could start-up at a HRT of 5.2-5.7 days with no VFA accumulation. Previous work on leachate from MSW has also shown that microorganisms grown in another medium are unable to out-compete native solid waste microorganisms for the cellulose in a foreign (leachate based) medium (O'Sullivan and Burrell, 2007). In this

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study, ~~the~~ bacteria fed on peptone and meat extract may have been inhibited when suddenly fed with leachate, or merely may have not been able to metabolize lignocellulosic compounds causing a delay in VFAs production; this would explain why the methane content of SAMBR2 displayed such a long lag phase before reaching a normal value of 60% methane in the biogas (Figure 68). Moreover, the methane content of the biogas in SAMBR2 gradually increased to a maximum of 61% after 50 days, whereas in SAMBR1 it reached 60% after four days of operation and then slowly stabilized at values between 69 and 71% (Figure 68), which suggests that the inoculum fed on synthetic VFA was not suitable for start-up because it did not contain hydrolytic and acidogenic bacteria for a leachate medium. As a result, the slow production of VFAs was translated into a slow increase in methane production.

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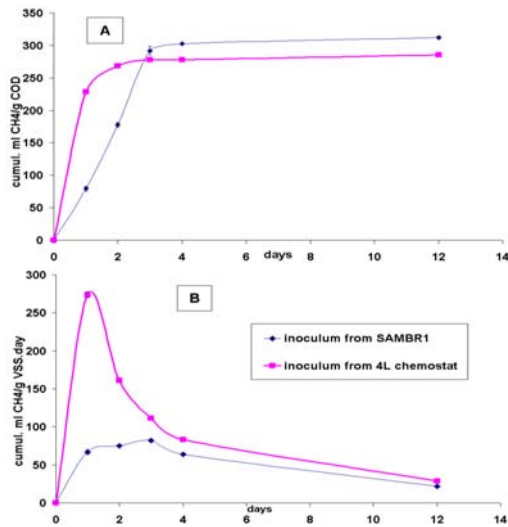


Figure 15. Specific acidogenic activity test on the inoculum from SAMBR1 acclimatised to the leachate medium and the inoculum from a 4 litre chemostat enriched with methanogens in a synthetic medium of peptone, meat extract and VFAs. Panel A: cumulative methane production per g COD fed. Panel B: cumulative methane production per g MLVSS per day. Reprinted from (Trzcinski and Stuckey, 2009b).

However, it could be argued that methane percentage in the headspace is not a good indication of the methane production because it depends on the carbon dioxide solubility, which in turn depends on the pH, alkalinity and buffer capacity. In this case, the pH and the alkalinity (Figure 2023) in SAMBR1 and SAMBR2 were very similar which provides a sound prerequisite to compare headspace methane content. Furthermore, the bulk SCOD in SAMBR2 kept increasing during start-up from 100 mg/L on day 46 to 800 mg/L on day 73 after which it began to decrease. This strongly suggests that, for a lignocellulosic-based feed, it is paramount to start up the SAMBR with a hydrolytic population acclimatised to the leachate medium to avoid SCOD build up in the bulk. Eventually, no biogas was produced in SAMBR2 until day 106, whereas the specific methane production in SAMBR1 was in the range 0.11 - 0.18 L CH₄/g COD fed during the first 30 days.

3.3.2. Effect of the HRT on COD Removal

The HRT of the AMBR was equal to the HRT of SAMBR2 because the two reactors were connected in series. The COD removal in SAMBR2 was 94.5% on average, and only 1.6% in the AMBR so that a total COD removal of 96.1% was achieved. The VFA concentration was virtually zero inside SAMBR2 and the permeate, and thus were omitted from Figure 1416. No significant change in total COD removal efficiency of both reactors was observed when the HRT was decreased from 5.2-5.7 to 0.37 days. In a moving-bed biofilm reactor system with an anaerobic-aerobic arrangement, Chen et al. (2008a) observed that at 1.5 days HRT the COD removal of the anaerobic reactor dropped to 81%, whereas the aerobic COD removal increased to 11%, but nonetheless the total COD of the system remained stable. Although the contribution of the aerobic step to the total COD removal of the system was low in this study because of the membrane rejection in SAMBR2 (1.6% on average), it should be emphasized that on average 26% of the recalcitrants from SAMBR2 could be degraded aerobically in the AMBR. Bohdziewicz et al. (2001) reported less than 10% aerobic biodegradation of anaerobically treated landfill leachate. The COD in the permeate of the AMBR was approximately 300

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mg/L at the end of the experiment, which is close to the 390 mg/L reported by Agdag and Sponza (2005). The SCOD in the bulk of the reactors as well as in their permeate decreased slightly towards the end of the operating period.

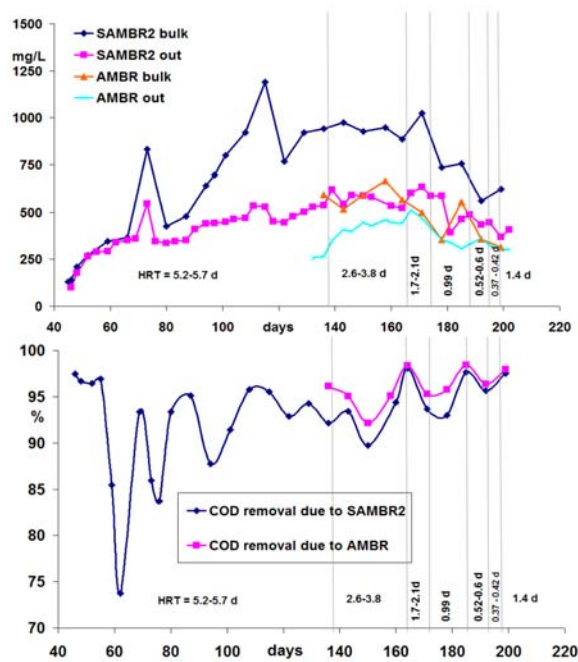


Figure 16. SCOD inside and in the permeate of SAMBR2 (out) and AMBR at different HRTs (top). Reprinted from (Trzcinski and Stuckey, 2009b). COD removal in SAMBR2 and AMBR (bottom).

3.3.3. Effect of the TSS on Flux

Figure 14-17 shows the evolution with time of the TSS, VSS and the FSS. Interestingly, the initial flux in SAMBR2 was much higher than in SAMBR1; the initial flux in SAMBR1 and 2 was 8.6 and 21.6 LMH, respectively although both reactors were started up at similar TSS of 3.3

and 2.6 g/L for SAMBR1 and SAMBR2, respectively. The only difference was in the different inoculum used to seed the SAMBRs. SAMBR2 was inoculated with an inoculum almost free of colloids, whereas SAMBR1 was inoculated with bacteria acclimated to the leachate medium but containing colloids that are known to be detrimental for the flux. Nonetheless, as SAMBR2 was fed on leachate containing colloids its flux also started to decrease over time. As a result, the TMP started to rise so rapidly that the flux had to be reduced manually. This was the case on day 50 where the TMP rose to 200 mbar, on day 53 where it reached 250 mbar and the flux was reduced to 15.3 LMH. Again, on day 59 the TMP rose to 350 mbar and the flux was reduced to 12 LMH, which stabilized the TMP at 200 mbar. With time the TMP rose again to 300 mbar on day 70 and the flux was reduced to 10.1 LMH to stabilize the TMP between 150 and 200 mbar. This approach was also followed by other researchers (Howell et al., 2004) in order to avoid serious fouling of the membrane.

For relatively ideal wastewater such as synthetic or municipal wastewater, the permeate rate is usually reduced if the TMP exceeds 10kPa (= 100 mbar) because this is equivalent to a 1m liquid head corresponding to the typical liquid depth above the membranes in full size systems. In only 30 days, its flux decreased to the same flux as in SAMBR1 meaning that the advantage of a “colloid free” inoculum was effective only for a short period. As in SAMBR1, the low HRT resulted in the build-up of MLTSS, MLVSS and FSS as well as the drop in the VSS/TSS ratio over time. The operation after day 100 was characterized by an increase in TMP to reach 650 mbar on day 120. The TMP remained constant as well as the flux (around 5 LMH), although the MLTSS increased gradually from 1 to 19 g/L on day 178, which emphasized the independence of the MLTSS on the flux in a certain range of values as reported by other researchers (Hong et al., 2002; Ross et al., 1990; Yamamoto et al., 1989). Then after day 180, the MLTSS went over 20 g/L, and even reached 46 g/L on day 195, which eventually blocked the diffuser and no more bubbles could be seen scouring the membrane, which was correlated with the TMP culminating at 850 mbar and the flux dropping to 0.5 LMH until the end of the experiment. Thus, as in SAMBR1, at HRTs lower than 2 days, particulate

solids in the leachate built up at the bottom of the SAMBR eventually leading to the diffuser blocking (see Figure 18). During the last 20 days of the run, sludge from the bulk was removed in an attempt to decrease the MLTSS in SAMBR2. On average, 130 ml were removed per day (SRT = 23 days), but that was not enough to decrease the MLTSS as the feeding rate of particles was too high compared to its removal rate.

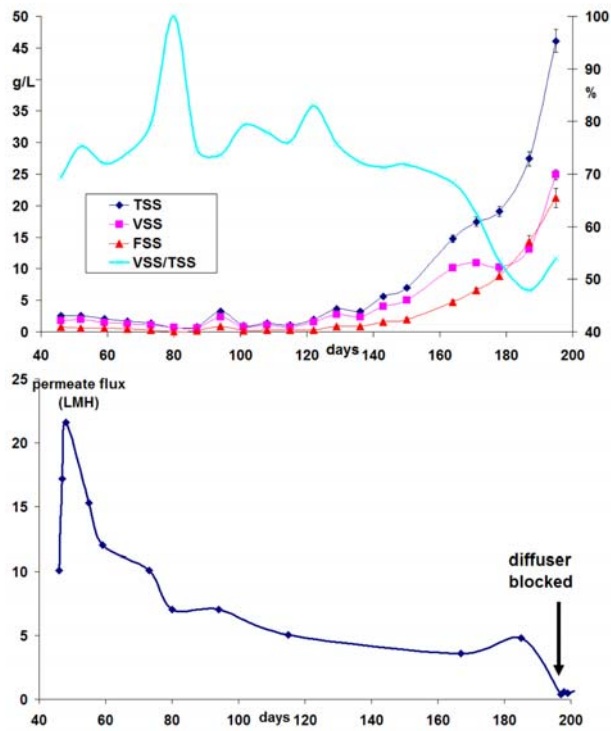


Figure 17. Evolution with time of the TSS, VSS, FSS and VSS/TSS ratio (top), and evolution of the flux in SAMBR2 (bottom).



Figure 18. Left: Kubota membrane module partially scoured by biogas bubbles. Right: blocked diffuser located at the bottom of the SAMBR with only the far left hole remaining open for gas bubbles formation. Reproduced from (Trzcinski and Stuckey, 2009a).

3.3.4. *Effect of the Biogas Sparging Rate on COD Rejection*

Previous work on the SAMBR has shown that the presence of a biofilm is responsible for a cleaner effluent (Akram, 2006; Choo and Lee, 1996; Harada et al., 1995). Both SAMBRs were fed on leachate at the same HRT of 5.2-5.7 days while the sparging was set at a low value in order to investigate the effect of sparging rate on the rejection of COD by the membrane. Another objective was to see how much time it takes to develop a biofilm and whether this biofilm could result in an enhanced rejection and a lower permeate COD. This could have a significant impact because it would mean that the permeate COD could be controlled by changing the sparging rate and thereby obtain cleaner effluent. On day 76, the two SAMBRs were deemed to be at steady state as they recovered from the sudden shock in OLR that made their COD rise to about 500 mg/L. The sparging rate was then set at 2LPM in both SAMBR. The MLTSS were relatively low and constant around 5 and 1 g/L in SAMBR1 and 2, respectively. It was observed that no significant change occurred and that the permeate SCOD remained stable at around 350-360 mg/L in both reactors. In contrast, the TMP skyrocketed to circa 580 and 380 mbar, in

SAMBR1 and SAMBR2, respectively (Figure 19). This indicated that the flux became greater than the critical flux as soon as the sparging rate was reduced and that a cake layer can form very quickly above the critical flux. The flux was manually reduced to 6.1 and 7 LMH in SAMBR1 and SAMBR2, respectively, to avoid operation at high TMP. The TMP went back to 150 and 100 mbar in SAMBR1 and SAMBR2, respectively, and this TMP was probably not high enough to build a biofilm that would have led to enhanced rejection, and as a result the permeate SCOD did not decrease. Although the flux had been manually reduced to 6.1 LMH in SAMBR1, the TMP slowly increased to 650 mbar at which point the flux was only 5 LMH (measured on day 94). Thus a cake layer had been formed at 6.1 LMH but at a much lower rate than at 8.3 LMH. The rate of increase in TMP indicates the degree of compaction of the biofilm which can translate to denser cake layer and COD rejection. The higher the rate of increase in TMP, the more compact the biofilm will be, and the thicker the biofilm will be. Also the degree of compaction will determine the easiness and the rate at which the biofilm can be removed if the sparging rate is increased.

At very high TMPs, Elmaleh and Abdelmoumni (Elmaleh and Abdelmoumni, 1997) also reported a decrease in permeate flux with an increase in TMP which was attributed to the compaction of the foulant layer. Li et al. (Li et al., 2003) explained that cakes formed at higher flux and TMP are much more consolidated than cakes formed marginally above the critical flux. In SAMBR2, although the flux had been manually reduced to 7 LMH, the TMP rose slowly to 400 mbar before day 87. On day 87, SAMBR 2 was set at a high sparging rate (8-10 LPM), while the sparging rate in SAMBR1 remained unmodified. As a result, the TMP in SAMBR2 dropped to virtually zero, while the permeate SCOD in SAMBR2 increased to about 440 mg/L on day 90 and remained stable while the permeate SCOD in SAMBR1 remained below 400 mg/L. It is interesting to note that the increase of the sparging rate did not allow us to re-establish the initial flux in SAMBR2, although the TMP went back to zero.

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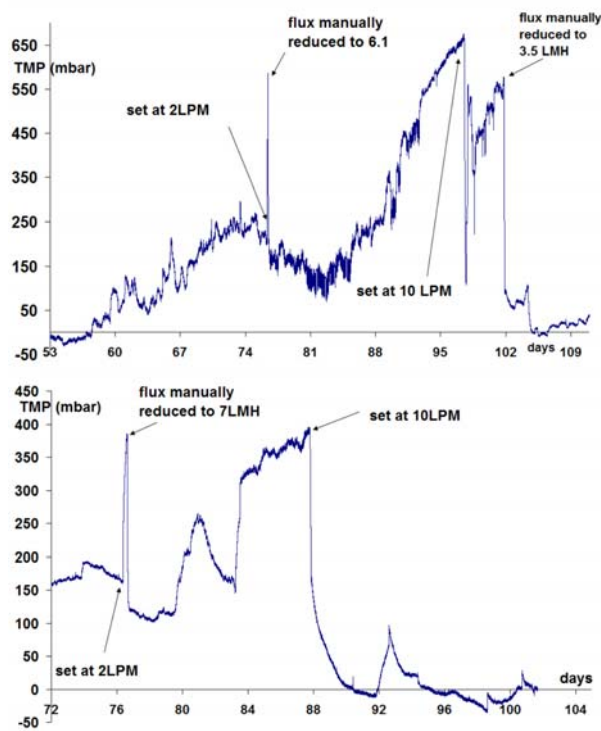


Figure 19. Evolution with time of the TMP in SAMBR1 (top) and SAMBR2 (bottom).

Therefore, it can be stated that in this study low sparging rates even for a short period of time lead to irreversible fouling because of internal fouling of the membrane pores. This is in contradiction with Vyrides and Stuckey (2009) who observed no significant increase in TMP or decrease in flux when operating the biogas pump in an intermittent mode (5 min OFF - 10 min ON) for 120 days. In our study, however, low sparging rates were likely to increase the thickness of the concentration polarization layer because of less turbulence, and this probably led to the aggregation of

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humic substances in the boundary layer. Humic acids can adsorb to particulate matter in the cake layer which results in a more compact and dense cake layer. In addition, the coagulation with calcium may have been enhanced in the concentration polarization layer where the concentration of humic acids is greater, resulting in the formation of aggregates on the membrane. This can lead to a thicker cake layer because calcium facilitates humic acids adsorption to particulate matter and membrane surfaces. Nghiem et al. (2006) also observed a greater TOC rejection associated with an increase in calcium concentration. Moreover, the highest TOC rejection coincided with the highest level of fouling which is consistent with our observations. Eventually on day 97, the sparging rate in SAMBR1 was also set at 8-10 LPM but the effect on the permeate COD was not immediate. In contrast, the effect of increasing sparging rate on the TMP was immediate on day 97: it suddenly dropped to 100 mbar, but unfortunately, the sparging rate could not be maintained at 10LPM due to foaming problems. Consequently, the TMP ended up at 550 mbar and eventually the flux was reduced manually by decreasing the speed of the recycle pump so that the flux was 3.5 LMH. As soon as the flux was lowered, the TMP went back to low values on day 101. It was only on day 104 that the permeate SCOD also rose as in SAMBR2 to values around 440 mg COD/L. Thus seven days (day 104-97) were necessary to see the effect of sparging rate increase in SAMBR1 that kept the low sparging rate mode for 21 days (day 97-76), while it took only 3 (day 90-87) days in SAMBR2 that kept the low sparging rate mode for 11 days (day 87-76).

The effect of the low sparging rate on the flux was immediate and detrimental. As can be seen from Figure 13 for SAMBR1 and from Figure 17 for SAMBR2, the flux drop was 4.8 (8.3 to 3.5LMH) and 3 LMH (10 to 7LMH) for SAMBR 1 and 2, respectively. Thus the effect of the low sparging rate was to block the pores while the partially clogged pores could still let the solutes pass. As a result of the pore blocking mechanisms, fewer molecules could pass which caused a significant increase in the bulk SCOD of both SAMBRs. Furthermore, the increase of the sparging rate to 10LPM did not allow the initial flux to recover showing that irrecoverable fouling occurred in both SAMBRs. The fouling was, however, worse for

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SAMBR1 because of the greater degree of compaction due to the longer period at low sparging rate, and also because its MLTSS was 5 times greater than that of SAMBR2 thus there was more material available in the bulk liquid to build a cake.

Therefore, this experiment performed in duplicate in separate SAMBRs showed similar results; when the sparging was set at a low rate (2 LPM), no significant change was noticed in the permeate COD. This might be due to longer periods required to build up a cake layer under slow TMP increase, and thereby obtain higher COD rejection and lower permeate COD. It is possible, however, that as the TMP increases, the compaction of the cake would be such that the pores of the biofilm will be so narrow that lower permeate COD will be obtained. Although the permeate SCOD did not decrease significantly, the membrane rejection started to increase significantly from day 80 onwards (Figure 20). Thus the effect of lowering the sparging rate was not a decrease of permeate COD, but instead an increase in membrane rejection due to enhanced concentration polarization, keeping at the same time a constant permeate SCOD. Choo and Lee (1996b) also concluded that cake compaction over time had more impact on the SCOD rejection by the membrane. This sounds paradoxical, but can be explained by the very wide range of molecular weight organics present in the bulk; while the lower MW can still pass through the biofilm and the membrane and are responsible for the constant permeate SCOD, the higher MW cannot pass through the pores and remain in the bulk which causes the membrane rejection to increase. These results are consistent with Choo and Lee (1996b) who, after the initial fouling of the membrane, observed a decrease in the SCOD in the bulk and the permeate due to the inoculum acclimatisation, but afterwards the permeate SCOD remained relatively constant for the rest of the experiment and did not decrease any further although the flux was decreasing due to internal fouling and cake layer compaction. This suggests that there is a limit below which the SCOD concentration in the permeate cannot be because the low MW recalcitrants which constitute the SCOD will still be able to pass through the membrane pores.

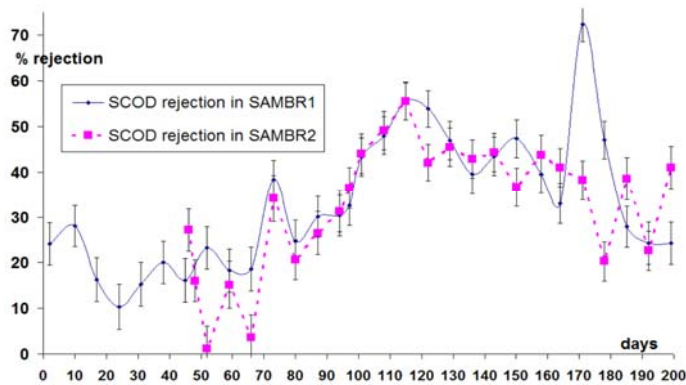


Figure 20. Evolution with time of the SCOD rejection by the membrane in SAMBR1 and SAMBR2.

On the other hand, when the sparging rate was increased to 8-10 LPM, the permeate COD increased typically from 360 to 440 mg/L due to a better scouring of the membrane, which resulted in a thinner concentration polarization layer and a better diffusion through the biofilm. Moreover, this increase in COD was faster for SAMBR2 that kept the low sparging mode for 11 days indicating that part of the cake layer can be easily scoured as indicated by the TMP that went back to zero after the high sparging rate was applied. In contrast, in SAMBR1 the low sparging rate mode was kept for 21 days, and due to the better compaction of the cake, the removal of the reversible fouling was impossible or much slower as indicated by a persistently high TMP, and as a result the effect of a higher sparging rate took longer to be noted. This increase in permeate SCOD was, however, not correlated with a reduction in membrane rejection in any of the SAMBRs shown in Figure 20.

Li et al. (2003) also concluded that the removal of the cake strongly depend on its age; if the cake existed for a short period of time, the lift forces due to surface shear can break some of the enmeshed bonding between the bacteria, allowing for the removal of the cake in the form of

flocs. On the other hand, if the cake has been built up for a longer period, the possibility of breaking the enmeshment is reduced. The aggregation of the cake will also depend on ionic strength: at high ionic strength, which is likely to be the case in landfill leachate, the electrical double layer of bacteria is compressed resulting in more interaction and binding with molecules which allows a dense biofilm. Nevertheless, the change in COD (about 80 mg/L) was relatively small compared to the total COD concentration: this is a 20% increase in permeate SCOD by changing the sparging rate from 2 to 8-10 LPM.

3.4. Performance of the AMBR

3.4.1. Effect of the HRT on Nitrification

The sequential oxidation of NH_4^+ to NO_3^- involves autotrophic NH_3 oxidisers (AOB = ammonia-oxidising bacteria) and autotrophic NO_2^- oxidisers (NOB = nitrite oxidising bacteria). In addition, heterotrophic bacteria can oxidise reduced forms of organic N to NO_3^- (Prosser, 2007). Ammonia-nitrogen in the permeate of SAMBR2 varied between 45 and 215 mg/L. This concentration depends on the hydrolysis efficiency of proteins in the HR which in turn varied according to the VS retention time and the HRT of the HR. Moreover, the protein (only a few measurements were taken) concentration decreased whereas the ammonia concentration increased when passing through the SAMBR, which means that the hydrolysis of proteins was continuing in the SAMBR. Protein determination on days 115, 143, 171 and 199 gave a concentration circa 100 mg/L in the permeate of SAMBR 1 and 2, whereas it was around 200 mg/L in the leachate from the HR. Surprisingly, the concentration in the AMBR permeate was still around 50 mg/L although proteins are known to be degraded readily. It was found in the literature that the determination of proteins using the direct Lowry procedure can suffer from interference due to other chemicals such as tris, ammonium sulfate, EDTA, sucrose, citrate, amino acid and peptide buffers, and phenols. The procedure with protein precipitation, which uses deoxycholate and trichloroacetic acid, eliminates

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all these interferences with the exception of phenols (Bensadoun and Weinstein, 1976). As the procedure with protein precipitation was not used in this work, it is thought that the remaining 100 mg/L in the SAMBR permeate as well as the 50 mg/L in the AMBR permeate were recalcitrant phenolic or proteinaceous compounds interfering with the protein analysis. The 50% degradation of these phenolic compounds in the AMBR shows that further stabilization was taking place in the AMBR which supports the polishing character of the AMBR demonstrated by the COD removal. Figure 21 shows the evolution of inorganic nitrogen in the AMBR. Because the inoculum used in this study came from a dye wastewater plant, it was assumed that it did not contain any nitrifiers. As a result, ammonia-nitrogen was initially not converted to nitrite or nitrate. Ammonia oxidisers may also have been inhibited by undissociated ammonia (NH_3) which was in the range 14 - 23 mg NH_3/L between days 136 and 146. Anthonisen et al. (1976) have observed that free ammonia can inhibit ammonia oxidation to nitrite by *Nitrosomonas* and nitrite oxidation to nitrate by *Nitrobacter* or *Nitrospira* in the range 10-150 and 0.1-1 mg NH_3/L , respectively. The temporary nitrite build-up may be explained by the inhibition of nitrite oxidisers due to the free ammonia ranging from 0.1 to 0.4 between days 146 and 167. Inhibition of nitrifying organisms by free nitrous acid (HNO_2) is unlikely to have occurred as the concentration remained in the range 0.00084-0.0052 mg HNO_2/L , which is far below the inhibitory range of 0.22 to 2.8 mg/L reported by Anthonisen et al. (1976). The growth of *Nitrobacter* (or *Nitrospira*) was confirmed by the slow decrease in nitrite which was correlated with a slow increase in nitrate.

From day 171 onwards, on average 97.7% of the $\text{NH}_4^+\text{-N}$ was converted to nitrite and nitrate in the AMBR at a maximum nitrogen loading rate of 0.18 g $\text{NH}_4^+\text{-N}/\text{L}\cdot\text{day}$ and a HRT of 0.5 day. According to Figure 21, the evolution of nitrogen is the one observed in batch mode (Anthonisen et al., 1976) where the ammonia is sequentially converted to nitrite and then to nitrate. In full scale continuous wastewater treatment plant nitrite is seldom found in the reactors because the conversion of NO_2^- to NO_3^- is much faster than the conversion of NH_4^+ to NO_2^- .

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Therefore, as long as the NOB species are present in the digester running continuously nitrite is almost never found. In this study, nitrite was found because the inoculum did not contain any nitrifiers, and the presence of a membrane kept all the species in the reactor and thereby resulted in a sequential reaction as if the AMBR was batch fed. Regarding the effect of the HRT, it can be said that the growth of AOB and NOB was not impeded by the successive HRT shocks until the shock to 0.99 day where the nitrite concentration did not reach zero which indicates that this HRT could overload the NOB such as *Nitrobacter*. However, previous work has shown that floc size for ammonia degradation is also important (Wittebolle et al., 2008) because a close distance between the nitrite producers and nitrite oxidisers is required to obtain complete nitrification. Wittebolle et al. (2008) have compared nitrification in a sequencing batch reactor and a MBR, and found that residual nitrite in the MBR was correlated with smaller flocs. They measured flocs of 1 to 40 microns in the MBR versus 300 microns in the sequencing batch reactor. Because of the shear effect in the AMBR, it is possible that the residual nitrite was due to a lack of syntrophic association between AOB and NOB. In our study, particle size distribution analysis was carried out in the AMBR, but particles of 1000 microns were detected which could be inorganic precipitates, thus rendering the comparison with data in the literature difficult. In another study (Shin et al., 2005), a temporary nitrite build-up in a MBR was associated with a poor oxygen transfer in the dead space of the MBR. The authors also reported that nitrification was better accomplished with pH control with HCl than without pH control. In this study, the SCOD fed to the AMBR was relatively low (400-600 mg/L) which promoted the growth of autotrophic bacteria. Because of the low organic content, high DO (1.6 mg/L) and good mixing (1.4 LPM) optimal conditions were met for the growth and retention of autotrophic AOB in the AMBR at a HRT as low as 0.37 day. In contrast, Chen et al. (2008a) and Im et al. (2001) could not maintain nitrification at 1.5 and 2.7 days HRT, respectively, because the COD concentration in the feed to the aerobic step sharply increased. Jokela et al. (Jokela et al., 2002) also observed that nitrification efficiency dropped to below 20% when the COD concentration suddenly increased at

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1.4 d HRT. As a result, heterotrophs competed for oxygen with the autotrophs leading to a decrease in nitrification activity. The AMBR permeate containing circa 50 mg/L of nitrite and the same concentration of nitrate was recycled to the HR where no nitrite or nitrate was detected. This was confirmed by the $\text{NH}_4^+\text{-N}$ and TN analysis which gave close concentrations; typically the TN was 10-15 mg/L greater than the $\text{NH}_4^+\text{-N}$ concentration in the HR effluent. The discrepancy was organic nitrogen such as proteins, peptides, nucleic acids or urea; this strongly suggests that denitrification took place in the HR. Another clue is obtained by comparing the TN concentration of the AMBR effluent and the HR effluent; the latter was on average 67% lower than the former which indicates substantial N removal. At this point it is not clear whether Anammox species were responsible for the conversion of nitrite or nitrate to gaseous nitrogen using ammonia that is the end product of protein degradation from the OFMSW feedstock fed to the HR. It is known that NOB can convert nitrite to nitrate if dissolved oxygen is present, and that heterotrophs will consume all the oxygen present in the HR before the NOB could consume the oxygen. However, as the HR was frequently opened for feeding and because of technical problems, it is possible that there was enough DO for NOB to grow and coexist with heterotrophs. Figure 7-8 shows how often the HR was opened for technical problems, and it can be seen that after day 160 there was no methane in the headspace. As the DO was not measured in the HR, it is not clear whether NOB could have grown in the HR, but denitrifiers on the other hand were likely to grow because of nitrate from the AMBR and the carbon sources available. Moreover, the TN analysis also revealed that between 7 and 35% of the TN in the permeate of the AMBR was organic N and that organic N was slowly increasing over time. Hence, heterotrophs could very likely have coexisted in the AMBR using organic N for growth and recalcitrants as a sole carbon source, which may explain why the AMBR acts as a COD-polishing step. On one hand there was organic N as a nitrogen source for the heterotrophs in the AMBR, but on the other hand their increase suggests that these were recalcitrant molecules containing amine groups.

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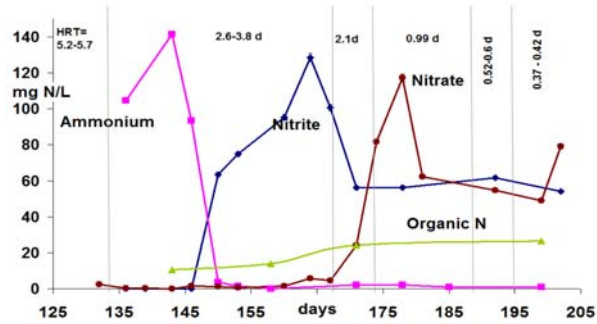


Figure 21. Concentration over time of the organic and inorganic nitrogen in the AMBR permeate. Reprinted from (Trzcinski and Stuckey, 2009b).

3.4.2. Effect of the HRT on the TSS, VSS, FSS, pH and Flux

The AMBR was started up at a TSS, VSS and FSS of 3, 2.3 and 0.7 g/L, respectively, and was fed with the stabilized leachate, i.e., the permeate of SAMBR2. Hence the feed contained no solids in suspension. Figure 22 shows that the VSS/TSS decreased linearly with time because the biomass was either settling out, attaching to the reactor's panels, or was forming a biofilm on the membrane.

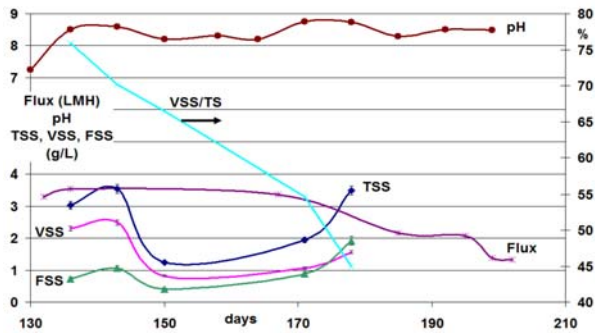


Figure 22. Evolution with time of the pH, flux, TSS, VSS, FSS (left axis) and the VSS/TSS ratio (right axis) in the AMBR.

The pH started at 7.25 but then plateaued at circa 8.5. This is due to the carbon dioxide being stripped out of the liquid phase due to air sparging while the reactor was open to the atmosphere. The feed to the AMBR was saturated in CO₂ because it was used in SAMBR2 to recirculate the biogas at the bottom of SAMBR2 for mixing and scouring purposes. The concentration is CO₂ is set by Henry's law:

$$[CO_{2(aq)}] = \frac{pCO_2}{K_{CO_2}} \quad (10)$$

where pCO_2 is the partial pressure in atmosphere and K_{CO_2} is Henry constant. In SAMBR2, as the biogas contains about 20% CO₂, it can be said that the partial pressure was 0.2 atm and the CO₂ concentration in the liquid is therefore $5.24 \cdot 10^{-3}$ M using Henry's law described in Equation 6.410. In the AMBR which is an open system, the partial pressure of CO_{2(g)} is relatively constant at 0.000355 atm (=355 ppm). Therefore, CO_{2(aq)} is equal to $1.38 \cdot 10^{-5}$ M. This is almost 3 orders of magnitude less than in the SAMBR. In other words, as soon as the feed enters the AMBR the carbon dioxide is stripped out to reach a new equilibrium according to Henry's law. As a result of CO₂ stripping the equilibrium of carbonic acid is changed and there is less dissolved carbon dioxide available for carbonic acid formation:



There is, therefore, less dissociation into protons according to the carbonic acid equilibrium, which caused the pH to rise.



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Regarding the alkalinity, it can be defined as:

$$A = [\text{HCO}_3^-] + 2[\text{CO}_3^{2-}] + [\text{B}(\text{OH})_4^-] + [\text{OH}^-] + 3[\text{PO}_4^{3-}] + [\text{HPO}_4^{2-}] + [\text{SiO}(\text{OH})_3^-] - [\text{H}^+] - [\text{HSO}_4^-] - [\text{HF}] \quad (14)$$

but this can often be approximated by:

$$A = [\text{HCO}_3^-] + 2[\text{CO}_3^{2-}] + [\text{OH}^-] + [\text{H}^+] \quad (15)$$

because phosphates and silicate are negligible, and at pH = 8.5, $[\text{HSO}_4^-]$ and $[\text{HF}]$ are also negligible. As carbon dioxide is stripped out of the system, the pH increases, but the removal of carbon dioxide does not modify the alkalinity of the system. At pH 8.5, the ion HCO_3^- predominates over H_2CO_3 and CO_3^{2-} . Thus when carbon dioxide is removed from the system, the following equilibrium is shifted to the left:



The alkalinity is not changed because the net reaction removes the same number of equivalents of negatively contributing species (H^+) as positively contributing species (HCO_3^-). The precipitation of calcium carbonate, however, has a strong influence on the alkalinity. This is because carbonate is composed of CaCO_3 and its precipitation will remove Ca^{2+} and CO_3^{2-} from the solution. The calcium ion will not influence alkalinity, but CO_3^{2-} will decrease the alkalinity by 2 times the molar concentration of CO_3^{2-} .

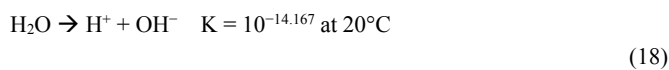
The precipitation of calcium carbonate was observed in our system, and the decrease of alkalinity was confirmed by titrating the effluent from SAMBR2 and AMBR which were found to contain 2250 and 750 mg eq CaCO_3/l , respectively (Figure 23). Borzacconi et al. (1999) have also observed that calcium ions can precipitate as CaCO_3 at a pH around 8. The maximum concentration of calcium before precipitation can be predicted using the solubility product assuming that the interference of other ions can be neglected. The equilibrium is given by the equation:



where the solubility product varies from $K_{sp} = 3.7 \cdot 10^{-9}$ to $8.7 \cdot 10^{-9}$ at 25°C , depending upon the data source (Seely, 2008; Weast, 1963). However, this equation must be taken along with the equilibrium of carbon dioxide with atmospheric carbon dioxide and the equilibrium of bicarbonate and carbonic acid given above.

For the AMBR, it is assumed that the partial pressure of carbon dioxide is $3.5 \cdot 10^{-4}$ atmospheres and that the pH is 8.5. Therefore, according to Henry's equation, $\text{CO}_{2(aq)}$ is equal to $1.38 \cdot 10^{-5}$ M.

This results in a concentration of H_2CO_3 equal to $1.6 \cdot 10^{-8}$ moles per litre taking into account the dissociation constants given in [equations 11, 12 and 13](#). ~~Section 2.3.6 of the literature review which are valid for high ionic strength.~~ When H_2CO_3 is known, the remaining three equations together with (Perrin, 1982):



makes it possible to solve simultaneously for the unknown concentrations. If $4.47 \cdot 10^{-9}$ is taken for K_{sp} , the maximum solubility of Ca^{2+} in the AMBR is only 3.1 mg/L at pH 8.5. Furthermore, this value shows a quadratic dependence with the concentration of H^+ so that the maximum Ca^{2+} concentration is 100 times greater at pH 7 than at pH 8.

In SAMBR2 at 35°C , the pH was 7 and the partial pressure of carbon dioxide was 0.2 atm (maximum 20% CO_2 in the headspace), and the CO_2 concentration in the liquid phase is therefore $5.24 \cdot 10^{-3}$ M. The solubility of calcium at pH 7 is circa 2,850 mg/L, but because of the high partial pressure of carbon dioxide the solubility in SAMBR2 was only 5.1 mg/L. Thus precipitation can occur in the SAMBR as well as in the AMBR if the calcium concentration is greater than the calculated values. If the pH increases to 7.3 due to high ammonia concentrations for instance, the maximum solubility of calcium become 1.3 mg/L, while if acidification to pH 6.8 takes place due to high VFA concentrations, it becomes 12.7 mg/L.

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Figure 22 also shows that the flux in the AMBR was decreasing over time. Even though the TMP could not be recorded because only 2 transducers were available for the SAMBRs, it is very likely that calcium carbonate precipitating on the membrane was causing the flux to drop. Visual inspection of the membrane indicated a hard, off-white, chalky deposit similar to **limescale**, and contact with a drop of HCl provoked an immediate fizzing. A **precipitate** was also found in the permeate tubing, and a sample was taken and washed with plenty of water and then dissolved in deionized water with HCl. It was then filtered through a 0.2 micron filter and analysed on a Dionex Ion Chromatograph that confirmed that the main element was calcium. In the HR, if a pH of 6 and a CO₂ partial pressure of 0.3 atm are considered, the maximum calcium solubility is 954 mg/L, indicating that **precipitation** is less likely to occur in the HR.

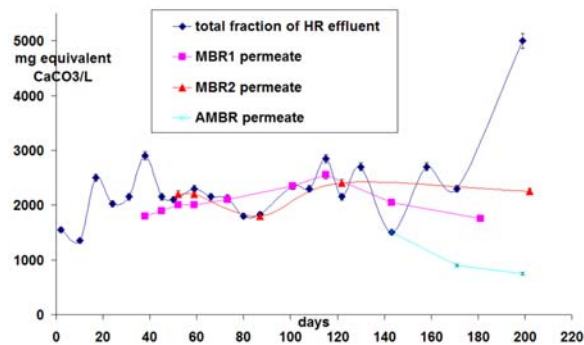


Figure 23. Evolution of the alkalinity as mg equivalent CaCO₃ per litre in the effluent of HR and the permeate of SAMBR 1, SAMBR2 and AMBR.

3.4.3. Scanning Electron Microscope - Energy Dispersive X-Ray (SEM-EDX)

Scanning Electron Microscope (SEM) was used to investigate the morphology and structure of the inorganic precipitate found on the membrane in the AMBR. This technique was coupled to Energy-Dispersive X-rays (EDX) in order to determine the chemical composition

of the sample. The EDX spectra confirmed that the crystals formed contained calcium, oxygen and phosphorus. The carbon peak observed was likely to be due to the sample carbon coating, but it could have come from the membrane material itself (polyethylene). Some oxygen can also be attributed to dissolved organic matter although it is noteworthy that it can also be from the membrane polymer. Several random points on the samples were analyzed by EDX and all the spectra exhibited calcium, oxygen and phosphorus as the main elements. It was found that calcium could precipitate in two different phases; the flat background precipitate (Figure 24), or as long and flat needles (Figure 25) of about 70 microns pointing upwards. The former phase contained small amounts of manganese, iron, magnesium, manganese, silicon, aluminium, sulphur, chloride and sodium (<1% by weight) whereas the needles contained relatively few impurities, which indicates that the needles are a purer form of precipitate than the background (Table 10). Moreover, the needle-shaped precipitate was characterized by significantly higher amounts of phosphorus compared to the background. Unfortunately, as the samples were coated with carbon, the carbon percentage was not available. Thus the comparison of the weight percentages from Table 10 with known minerals is difficult. Nonetheless, it is very likely that the background precipitate was comprised of calcium carbonate containing many impurities.

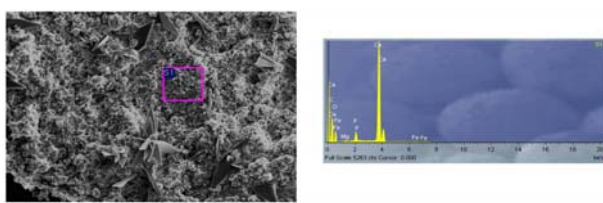


Figure 24. General view of the precipitate on the membrane in the AMBR (left), and elemental chromatogram of the background (right). Reprinted from (Trzcinski and Stuckey, 2016).

Regarding the needles, calcite was not considered due to the high phosphorus content. Several authors have reported that the presence of

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phosphates inhibits calcite growth due to the adsorption of phosphates on the calcite surface, which enables the formation of calcium phosphates (Lin and Singer, 2005; Plant and House, 2002).

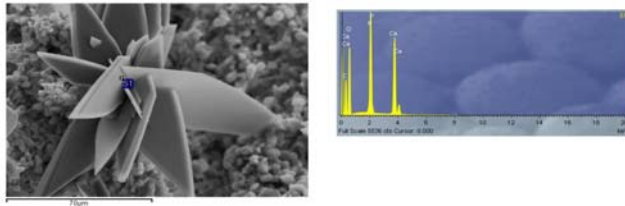


Figure 25. Zoom of the needle-shaped precipitate on the membrane in the AMBR (left), and its elemental chromatogram (right). Reprinted from (Trzcinski and Stuckey, 2016).

Interestingly, it was observed in the SEM picture (Figure 25) that indeed the needles seem to form on the background of calcite, and grow perpendicular to the background. The high amount of phosphorus suggests that it could be a calcium phosphate such as calcium hydrogen phosphate (or brushite $\text{CaHPO}_4 \cdot 2\text{H}_2\text{O}$, $K_{sp} = 10^{-6.69}$ (Montastruc et al., 2003)), dicalcium phosphate anhydrate (CaHPO_4 , $K_{sp} = 10^{-6.9}$ (Montastruc et al., 2003)), amorphous calcium phosphate ($\text{Ca}_3(\text{PO}_4)_2$, $K_{sp} = 10^{-24}$ (Mamais et al., 1994)), octacalcium phosphate ($\text{Ca}_8\text{H}(\text{PO}_4)_3 \cdot 2.5\text{H}_2\text{O}$, $K_{sp} = 10^{-49.6}$ (Montastruc et al., 2003)), hydroxyapatite ($\text{Ca}_5(\text{PO}_4)_3\text{OH}$, $K_{sp} = 10^{-46.8}$ (Barat et al., 2008)) or carbonated hydroxyapatite ($\text{Ca}_{10}(\text{PO}_4)_6(\text{CO}_3)_x(\text{OH})_{2-2x}$, $K_{sp} < 1 \cdot 10^{-100}$ (Ito et al., 1997)). However, calcium phosphate precipitation follows the Ostwald rule, which states that the least thermodynamically stable phase is the first one formed, which works as a precursor of the most stable phase (Montastruc et al., 2003). In this case, carbonated hydroxyapatite would be the least thermodynamically stable phase forming on the calcite background as a precursor for a more stable calcium phosphate. Furthermore, the same authors (Montastruc et al., 2003) found that both dicalcium phosphate dihydrate and amorphous calcium phosphate can crystallize at a pH lower than 7.3, and only amorphous calcium phosphate at a pH higher than 7.3. Therefore, the more

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stable phase in this study would either be dicalcium phosphate dihydrate or amorphous calcium phosphate.

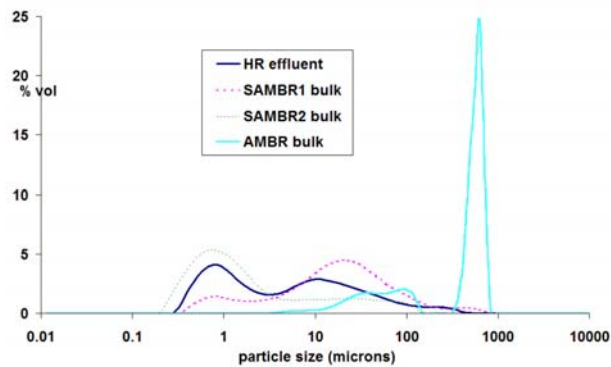
Table 10. Weight percentages of several points of the background and the needle precipitates

Background	Element										
	O	Ca	P	Al	Si	S	Fe	Mg	Cl	Na	Mn
Spectrum 1	46.7	47.98	2.32	0.64	1.09	0.28	0.99	0	0	0	0
Spectrum 2	38.95	55.23	2.08	0.3	1.06	0.28	1.49	0.31	0.29	0	0
Spectrum 3	45.03	50.3	3.43	0	0	0	0.66	0.57	0	0	0
Spectrum 4	55.93	33.55	6.66	0.26	0	0.17	0.82	0.77	0	0.82	1.02
Needles	Element										
	O	Ca	P	Al	Si	S	Fe	Mg	Cl	Na	Mn
Spectrum 5	55.79	26.19	17.22	0.22	0.58	0	0	0	0	0	0
Spectrum 6	30.99	59.04	9.97	0	0	0	0	0	0	0	0
Spectrum 7	64.04	18.78	17.18	0	0	0	0	0	0	0	0

Le Corre et al. (2005) and Doyle and Parsons (2002) observed that calcium ion can affect struvite crystal growth and leads to the formation of an amorphous calcium phosphate. Therefore, it is very likely that struvite precipitation did not occur because the available phosphates precipitated with calcium on the membrane keeping the solution under saturated with respect to struvite. From all these considerations and from the light shed by other authors, it seems very likely that the background was amorphous calcium carbonate and the presence of phosphates inhibited calcite growth due to the adsorption of phosphates on the calcite surface, which enabled the formation of calcium phosphates. The low level of impurities in these needles suggest that they have a more crystalline structure which is why hydroxyapatite is the most likely. Chesters (2009) obtained very similar SEM pictures with flat needles which he assumed to be hydroxyapatite crystals. In the same paper, the authors report the use of a new anti-scalant (Genesys PHO: 2-5 mg/L) to control calcium phosphate scale by inhibition mechanisms so that the solubility of calcium phosphate is raised 150 times.

3.4.4. Particle Size Distribution

It is well known that particles having the same size as the membrane pore are particularly likely to cause fouling. The particle size of the leachate and the contents of the SAMBR and AMBR were measured and compared in Figure 26. It can be observed that the leachate is very heterogeneous as it contains particles ranging from 0.3 microns to 1 mm. As expected, SAMBR1 and SAMBR2 also contains particles with a wide range of sizes including colloids below 0.4 microns which are presumably the particles responsible for pore blocking. The distribution of the leachate and SAMBRs mixed liquor were bimodal with a peak at circa 1 micron and the other at 10 microns, and a high proportion (90%) was below 48, 83 and 30 microns for the HR effluent, SAMBR1 and SAMBR2 mixed liquor, respectively. No straightforward comparison could be made between the leachate and the SAMBR bulks because of the heterogeneous character of the leachate that contained bacterial flocs as well as lignocellulosic fibres. The comparison between the volume percentages are also difficult as the accuracy of the MasterSizer 2000 is within 5% and the peaks on Figure 26 are all smaller than 5%. However, the distribution of the AMBR particles is such that 90% was below 632 microns with a mean diameter $d(4,3)$ equal to 405 microns.



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Figure 26. Particle size distribution of the leachate from the HR and the mixed sludge from SAMBR1, SAMBR2 and the AMBR.

3.5. Salts

3.5.1. Introduction

The salts ions are Na^+ , K^+ , Mg^{2+} , Ca^{2+} , Cl^- , PO_4^{3-} , SO_4^{2-} , and it is well known that high salt levels can cause bacterial cells to dehydrate due to osmotic pressure and that Na^+ is more toxic than any other salt on a molar basis. It is also known that the cation determines the toxicity of salts (McCarty and McKinney, 1961). Sodium, potassium, magnesium, calcium, phosphate and sulphate are present in the influent of anaerobic digesters and they are all required for microbial growth. Moderate concentrations stimulate microbial growth, but excessive amounts can cause inhibition or toxicity (Soto et al., 1993). Figure 27 shows the evolution with time of the ions. Sulfate was omitted because its concentration remained below 10 mg/L in all reactor effluents throughout the 200 days. These results are discussed separately in the following sections.

3.5.2. Sodium

Sodium is essential for methanogens and concentrations in the range 100-200 mg/L were reported beneficial for the growth of mesophilic anaerobes (McCarty, 1964) because of its role in the formation of adenosine triphosphate or in the oxidation of NADH (Feijoo et al., 1995). Sodium concentrations ranging from 3,500 to 5,500 mg/L can be moderately inhibitory and 8,000 mg/L can be strongly inhibitory to methanogens at mesophilic temperature depending on the adaptation period, antagonistic/synergistic effect, substrate and reactor configuration (Kugelman and McCarty, 1964). He et al. (He et al., 2006) report that 3.5-5.5 g/L and 2.5-4 g/L are the concentrations of sodium and potassium ions, respectively, that are optimal for AD. Shin et al. (Shin et al., 1995) showed that 2 to 10 g/L moderately inhibited the methanogenic activity, but

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concentrations above 10 g/L strongly inhibits. Although 8 g/L of Na⁺ and 12 g/L of K⁺ causes severe toxicity in the treatment of wastewaters, methanogens have been reported to acclimate to high salts stress, such as 21.5 g/L Na⁺ and 53 g/L Na⁺. Furthermore, He et al. (2006) observed that 50 g/L Na⁺ inhibited acidogenesis, but surprisingly had the highest amount of substrate hydrolysed because of the highest α -amylase activity at that concentration of Na⁺; the bacteria recovered at a concentration of 25 g/L.

They also observed that denitrification was optimal at pH 7 and that 50 g/L suppressed denitrification. In another study, sodium was found to be more toxic to propionic acid-utilizing microorganisms than acetate-utilizing and lignocellulose-degrading ones (Liu and Boone, 1991; Soto et al., 1993). The literature gives confusing data regarding the optimal and inhibitory levels of salts, and papers have focused more on inhibition studies in batch bottles rather than in continuous systems. Indeed, very little data is available for full-scale plants as ions are not systematically measured to assess the health of anaerobic digesters.

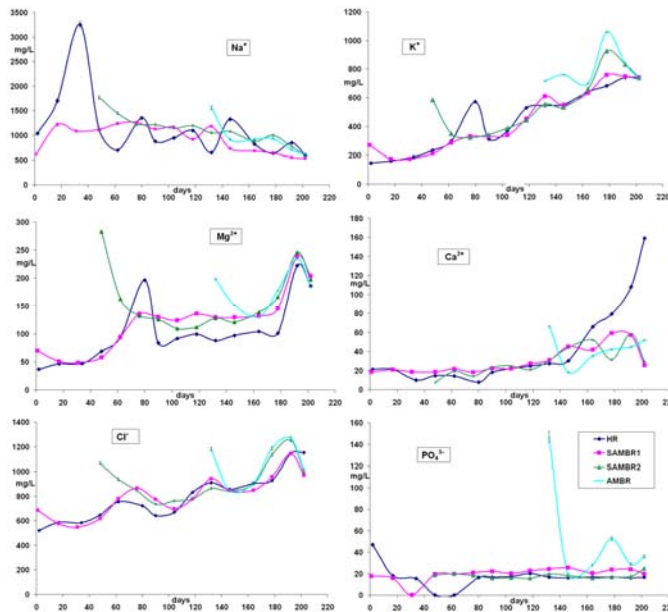


Figure 27. Concentration over time of Na^+ , K^+ , Mg^{2+} , Ca^{2+} , Cl^- , PO_4^{3-} in the HR effluent and the permeate of SAMBR1, SAMBR2 and AMBR. The error bars represent the standard deviation.

In this study, HR and SAMBR1 started with a Na^+ concentration of 1,034 and 612 mg/L, respectively, whereas SAMBR2 and AMBR started with a sodium concentration over 1.5 g/L due to the sodium present in the initial biomedium (Owen et al., 1979) that already contains 1373 mg Na^+ /L. The concentration in the HR effluent and SAMBR1 permeate increased markedly to over 1000 and 3000 mg/L, respectively, in the first 40 days during which time the OLR was low at 0.5 g VS/L.day. There was an initial build-up of sodium due to the high solids destruction, and very little amount of digestate were taken out of the reactor. The levels in both reactors then stabilized around 1 g/L presumably due to sodium uptake for bacterial growth. When SAMBR2 and AMBR started up their levels

levelled off also at circa 1 g/L suggesting a sodium uptake for bacterial growth, and this uptake was not renewed by new sodium ions from the feedstock as the OLR was increased and the solids destruction decreased. Also the sodium content of the OFMSW was low (2.3 ± 0.3 mg Na/g of dry matter). The limited solids destruction at higher OLRs can also explain why more solids were taken out on a daily basis and consequently, the sodium may have been washed out in the final digestate that still contained a bit of process liquid. However, it is thought that this hypothesis was not the main reason because if it was the case the concentration of all other ions would also have decreased over time and that was not the case. It can be seen that at the end of the run, the concentration was below 700 mg/L in all the reactor effluents meaning that the concentration was suboptimal according to He et al. (2006); Owen et al.'s biomedium (1979) contains circa 1,373 mg Na⁺ due to Na₂S₉H₂O and NaHCO₃. This decrease of sodium could have had a severe impact on the process; it could have impeded the growth of new bacteria and decreased the hydrolysis performance because of less hydrolytic enzymes produced as He et al. (2006) stated that high sodium levels promoted α -amylase activity. It could also cause dehydration of cells resulting in their lysis, therefore, it is suggested to maintain the level of sodium between 1 and 2 g/L by addition of more kitchen waste that contains more sodium.

3.5.3. Potassium

Potassium is known to increase cell wall permeability by aiding the cellular transport of nutrients and providing cation balancing (Kayhanian and Rich, 1996). Low concentrations (less than 400 mg/L) can cause an enhancement in performance in both the mesophilic and thermophilic ranges (Chen et al., 2008b) while Kugelman and McCarty (1964) have observed that 0.15 M K⁺ (=5.9 g/L) caused 50% inhibition of acetate-utilising methanogens. Sodium, magnesium and ammonium can also mitigate potassium toxicity, with sodium producing the best results. Figure 27 shows that potassium in all reactors built up steadily over 200 days independently of the OLR. However, the evolution in the HR effluent

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increased suddenly from 297 on day 62 to 572 mg K⁺/L on day 80 when the OLR was set at 2 g VS/L.day. On the other hand, the evolution in SAMBR1 permeate was quite linear ($R^2=0.91$) suggesting that an equal amount of potassium was solubilised independently of the OLR or the HRT applied. This slow increase also indicates that the potassium uptake for bacterial growth is not as important as the sodium uptake. As a result, potassium is likely to build-up in processes and could become inhibitory after longer periods than the one in this study and should therefore be monitored closely. The final concentration in all the reactors effluent was close to the theoretical concentration in Owen et al.'s biomedium (1979) of 680 mg K⁺/L, thus it can be stated the concentration was optimal in the process towards the end of the experiment.

3.5.4. Magnesium

The optimal Mg²⁺ concentration was reported to be 720 mg/L for the anaerobic bacterium *Methanosarcina thermophila* (Schmidt and Ahring, 1993). High concentrations have been shown to stimulate the production of single cells which is detrimental because these are more sensitive to lysis and this is an important factor in the loss of acetoclastic activity (Chen et al., 2008). This study has shown that magnesium does not reach levels higher than 300 mg/L at all the OLRs tested. Although an increase was observed in the HR and SAMBR1 effluent when the OLR was changed to 2 g VS/L.day, the magnesium level remained relatively stable at an OLR of 4 to 16 g VS/L.day between 100 and 150 mg/L, which is relatively close to the level of 215 mg Mg²⁺/L in Owen et al.'s biomedium (1979). This indicates that the OFMSW contains enough magnesium in itself and that no extra source would be required for optimal concentration. The level increased suddenly in all reactor effluents after day 178 due to an artificial spike of trace elements based on Owen et al.'s biomedium (1979) in order to investigate the effect of trace metal addition, but no positive or negative effect were observed.

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3.5.5. Calcium

Calcium is known to be essential for the growth of certain strains of methanogens. It is also important in the formation of microbial aggregates. However, excessive amounts lead to the precipitation of carbonate and phosphate which may result in the scaling of reactors and biomass and reduced specific methanogenic activity, loss of buffer capacity and essential nutrients for anaerobic degradation (Chen et al., 2008b). Regarding the toxicity, the literature provides confusing data. No inhibitory effect was observed at 7000 mg Ca²⁺/L (Jackson-Moss et al., 1989), but Kugelman and McCarty (1964) reported that moderate inhibition can happen at concentration ranging from 2,500 to 4,000 mg/L and that the optimum concentration for methanogens was 200 mg/L. Concentrations greater than 300 mg/L were reported to be detrimental for biofilm formation (Hulshoff Pol et al., 1983). Mathiesen (1989) found that calcium and magnesium supplementation enhanced the methane production and foaming was avoided.

In this study, calcium concentration was low (about 20 mg/L) for the first 100 days until the OLR was set at 4 g VS/L.day on day 97. As the levels were very similar in HR effluent and SAMBR1 permeate, it seems that precipitation did not occur because the calcium concentration was too low. As a reference, Owen et al.'s biomedium (1979) contains 68 mg Ca²⁺/L in the form of CaCl₂·2H₂O. Interestingly, the level started increasing very rapidly to 160 mg/L in the HR effluent but the concentration in the other reactors remained much lower at about 60 mg/l, and even decreased to 20 mg/L at the end of the run indicating precipitation of calcium carbonate in the SAMBR and AMBR. The calcium concentration in the HR seems to be directly proportional to the OLR applied suggesting that calcium salts are readily dissolved in the HR but it could be also due to the lower pH in the HR that dropped below 6.5 after day 180. The concentration reached 80 mg/L on day 178 but the increase after day 178 is also partly due to the artificial spike of trace elements.

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3.5.6. Phosphate

PO_4^{3-} was found at a relatively low concentration in all reactors, except in the AMBR that started up at about 140 mg/L, presumably due to the level in the inoculum. The concentration in the AMBR dropped down to low levels (20 - 40) like in the other reactors. Phosphate is essential for bacterial growth because it is contained in DNA, but high levels are not recommended because it can precipitate as struvite ($\text{NH}_4\text{MgPO}_4 \cdot 6\text{H}_2\text{O}$). Because phosphate concentration is usually lower than ammonia and magnesium in anaerobic digester, it will determine whether struvite precipitates or not.

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3.6. Size Exclusion Chromatography (SEC)

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3.6.1. Introduction

Figure 28 presents the calibration curves used for the SEC. The standards with molecular weights (MW) greater than 219.3 did not appear as a single peak, and could not be resolved using the column Aquagel OH-40. An exponential curve was fitted to the calibration standard and the following equation was obtained ($R^2 = 0.98$):

$$\text{MW (Da)} = 4 \cdot 10^{10} \cdot \exp(-0.9851 \cdot t) \quad (19)$$

where t is the elution time.

Samples of the HR effluent and the SAMBR and AMBR permeates were taken regularly during the continuous experiment in order to investigate the evolution with time of the molecular weight (MW) distribution in the reactor effluents. However, a first objective was to study the difference in MW between the different reactors at a specific moment. Figure 29, Panel A, shows the size exclusion chromatograms of the HR effluent, SAMBR1 bulk and permeate, SAMBR2 bulk and permeate and AMBR bulk and permeate on day 192. This day was deemed representative of the situation in the process because the SCOD in the SAMBR and AMBR was stable. All the chromatograms contain two main

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peaks: one in a region of high MW (circa 11.5 min) and a second sharper peak at low MW (around 17.5 min). A similar observation was made by Ortega-Clemente et al. (2008) and Sierra-Alvarez et al. (1990). The former team was treating pulp mill effluents in an anaerobic fluidized bed reactor, while the latter treated soda pulp liquors of pine, spruce and wheat straw in a UASB. They stated that the lignin fraction removed or bio transformed anaerobically corresponds to low MW lignin derivatives.

The absorbance on the UV detector set at 254 nm was zero before 7.5 minutes and after 20 minutes. Panel B shows a zoom of the very high MW ($t \leq 9$ min) in order to understand the fate of very high MW organics (MW $\gg 200$ kDa). Panel B shows clearly that the very high MW organics were dominant in the leachate fed to the SAMBRs; the bulk of the SAMBRs still contained some of these molecules but considerably less. Interestingly, their permeate was absolutely free of them which indicates a full rejection of these compounds by the membrane. As expected, the AMBR bulk and permeate also did not contain any of these. These molecules eluting before 9 minutes are supposedly huge, but their MW cannot be determined accurately. Nevertheless an extrapolation of the calibration curve would show that organics eluting at 8.4 minutes would have a MW of 10,200 kDa, but this should be interpreted with caution since they are far out of the calibration range (the standard of 219.3 kDa eluted at 12.3 minutes).

The medium MW compounds (11 to 13.2 minutes elution time) were in the range ≈ 800 to ≈ 90 kDa. It can be seen from Figure 29, Panel A that the absorbance of these compounds is greater in the SAMBRs than in the HR as these medium MW compounds are likely to be the result of the hydrolysis of the aforementioned high MW compounds eluting before 9 minutes. Thus the medium MW compounds appear more concentrated than in the leachate due to a further hydrolysis and stabilization of the leachate in the SAMBRs. The stabilization of recalcitrant organics can also result in a higher degree of aromaticity that would also translate to a higher absorbance at 254 nm.

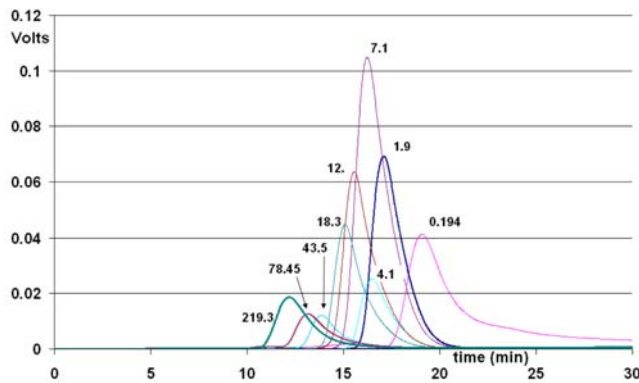


Figure 28. Calibration of the Size Exclusion Chromatography. The Y-axis is the absorbance of the PEG and PEO standards on the refractive index detector.

Interestingly, the absorbance in the SAMBRs permeate is lower which is evidence that these medium MW compounds were rejected by the membrane to some extent. For instance on day 192, based on the relative absorbance, there was a 16 and 18% rejection in SAMBR1 and SAMBR2, respectively. Furthermore, the permeate peaks are sharper than the bulk peaks and the HR effluent peak; it can be seen that SAMBR2 bulk has a wide peak that plateaus between 11.2 and 12.2 minutes while its permeate counterpart is between 11.7 and 12.2, thus showing the rejection of the highest MW compounds amongst the medium MW compounds present in the bulk. The highest standard (219.3 kDa) eluted at 12.3, but nonetheless if the calibration curve is extrapolated beyond 12.3 minutes, it is calculated that elution times of 11.2 and 11.7 would correspond to a MW of 646 and 395 kDa, respectively. As a result, the compounds in that range of 395 - 646 kDa were more likely to be rejected by the membrane while the $MW \leq 395$ kDa were observed in the permeate.

The comparison between the "SAMBR2 out" and "AMBR bulk" curves indicates firstly that medium MW compounds are further hydrolysed in the AMBR because they appear at a lower absorbance, therefore at a lower concentration than in SAMBR2 permeate. Secondly,

there is a slight shift of the AMBR curves to the right ($t = 12.7$ min) compared to the SAMBR2 permeate curve that peak at $\approx 11.8 - 12$ min meaning that the medium MW molecules are further broken down in the AMBR.

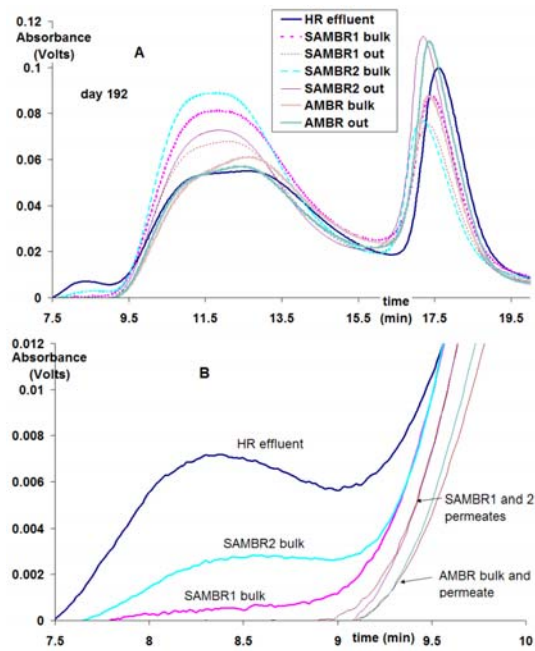


Figure 29. (A) Size exclusion chromatograms of the HR effluent, SAMBR1, SAMBR2 and AMBR bulks permeates on day 192. (B) Zoom of the curves before 10 minutes of elution time.

In terms of MW, it can be stated that medium MW compounds in SAMBR2 permeate are in the range 358 - 294 ($11.8 \leq t \leq 12$ min), while the AMBR bulk contained compounds of MW in the range 198-134 kDa ($12.4 \leq t \leq 12.8$ min). The COD polishing in the AMBR was therefore efficient on compounds of MW in the range 200 - 350 kDa.

Finally, all the samples exhibited a sharp peak with an elution time between 17.1 and 17.8 minutes, which is between the elution times of the standard 1.9 kDa ($t=17$ min) and 0.194 kDa (19.2 min), and thus correspond to low MW compounds or fulvic acids. The difference between these standards is very small in terms of MW and it is assumed that unknown peaks between these two standards are not significantly different.

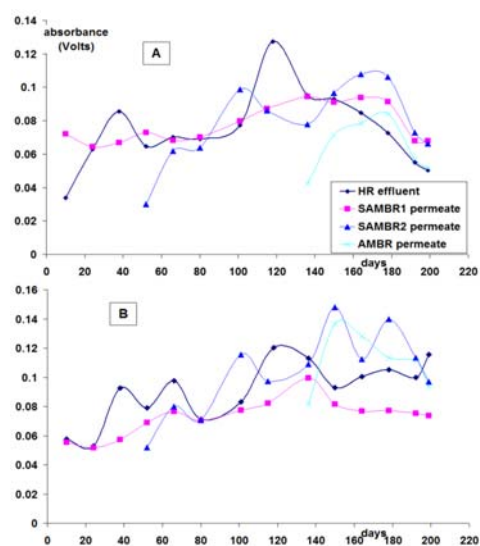


Figure 30. (A) Evolution with time of the absorbance of medium MW compounds (11 to 13.2 elution time, i.e., MW in the range 800 to 90 kDa) in the HR effluent and in the permeates of SAMBR1, SAMBR2 and AMBR. (B) Evolution with time of the absorbance of low MW compounds (17.1 to 17.8 elution time, i.e., MW in the range 1900 to 900 Da) in the HR effluent and in the permeates of SAMBR1, SAMBR2 and AMBR.

The low MW fraction of the HR effluent has a greater absorbance than in SAMBR1 and SAMBR2 bulks. This could be due to the higher concentration of aromatics being absorbed at 254 nm in the leachate compared to the SAMBRs bulks. The low MW fraction in the AMBR has a

greater absorbance than in the SAMBR2 permeate which could be due to the higher aromaticity of the former as fulvic acids are further degraded in the AMBR. In the case of SAMBR2 and AMBR the absorbance was higher in the permeate than in the bulk which could mean that there was an increase in the concentration of these fulvic acids in the permeate. However, it is more likely that the diffusion through the biofilm enables them to be degraded further than in the bulk which provided a greater degree of aromaticity that translated to a higher absorbance at 254 nm and was therefore not due to an increase in concentration. It is noteworthy that this rejection of medium MW and the higher degree of aromaticity of low MW in the permeate of AMBR happened not only on day 192 but also on all the other days sampled (days 136, 150, 164, 178 and 199). Regarding the evolution of the medium and low MW compounds over the 200 days, it can be stated that overall there was no build up in the absorbance in any effluent (HR, SAMBR1, SAMBR2 and AMBR). The panel A of Figure 30 shows the evolution of the medium MW compounds, while panel B shows the low MW compounds. The medium MW compounds in SAMBR1 followed a very similar evolution compared to the SCOD; the absorbance increased slowly until day 136, then plateaued, and eventually, like the SCOD, there was a decrease in absorbance after day 178 probably due to a lower concentration of all organics caused by dilution with more fresh water at high OLRs. The evolution of medium MW compounds in the HR, SAMBR2 and AMBR was more chaotic but eventually their absorbance also decreased.

Interestingly, the absorbance of SAMBR2 was two times lower than that of SAMBR1 during the start-up. This suggests that in SAMBR1 the competent population of hydrolytic bacteria acclimatised to the leachate medium was able to stabilize the organics by increasing their degree of aromaticity, while this was not the case in SAMBR2. At start-up SAMBR2 was virtually free of recalcitrant organics due to its inoculum and because the volume was adjusted with synthetic biomedium (Owen et al., 1979), and hence the concentration of organics with high aromaticity was lower. Similarly, the absorbance of the AMBR at start-up was considerably lower than in the other reactors because of the unadapted bacterial population

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that could not stabilize the refractory organics. In panel B for low MW compounds, the absorbance of SAMBR2 and AMBR was lower than that of SAMBR1, again due to the unadapted inoculum and the organics-free biomedium, but the difference was more significant for the medium MW compounds suggesting that the increase of SCOD in SAMBR2 during start-up was mainly due to the build-up of medium MW compounds that could not be hydrolyzed, rather than the fulvic acids.

3.7. Gas Chromatography - Mass Spectrometry

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3.7.1. Introduction

This technique can provide us with ~~additional-novel~~ insights into the processes taking place in the continuous treatment of OFMSW in two-stage membrane reactors. However, it should be born in mind that library identification should be interpreted with caution as sometimes a high match can be wrong, and this can be detected by comparing the mass spectra of the peak selected with the mass spectra of the proposed compounds. Moreover, a calibration line was drawn for o-hydroxybiphenyl and Bis (2-ethylhexyl)phthalate in order to gain more information on the concentrations of these specific recalcitrants found in the reactors. The raw chromatographs can be seen in Figure 31. A control sample (referred as 'scrap') consisted of 500 mL of DW in which small pieces of the plastic (plastic scrap) used to make the reactor were added and the mixture was shaken for few weeks at 30°C in order to determine which components if any could leach from the reactor's construction material (Table 11). Table 12 to Table 15 gather the peak identification number, the match percentage, the retention time, the area and the name of the components that were detected in the effluent of each reactor, but not in the blank (DW that followed the same SPE protocol) nor the sample with plastic scraps. In addition, the last column comments on the biodegradability by comparing the areas of the respective peaks.

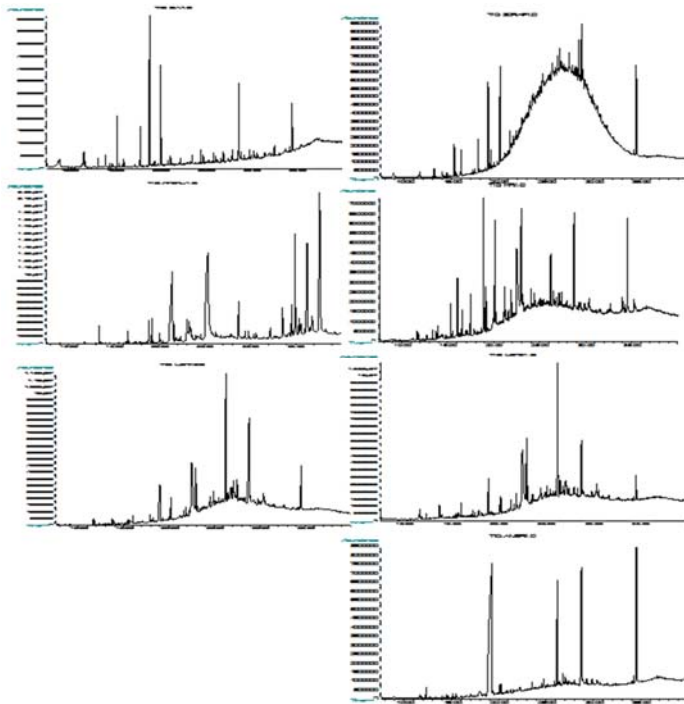


Figure 31. GC-MS chromatographs. From left to right: blank, reactor's plastic scrap, deionized water with anti-foaming agent, effluent of the hydrolytic reactor, SAMBR1 permeate, SAMBR2 permeate, AMBR permeate.

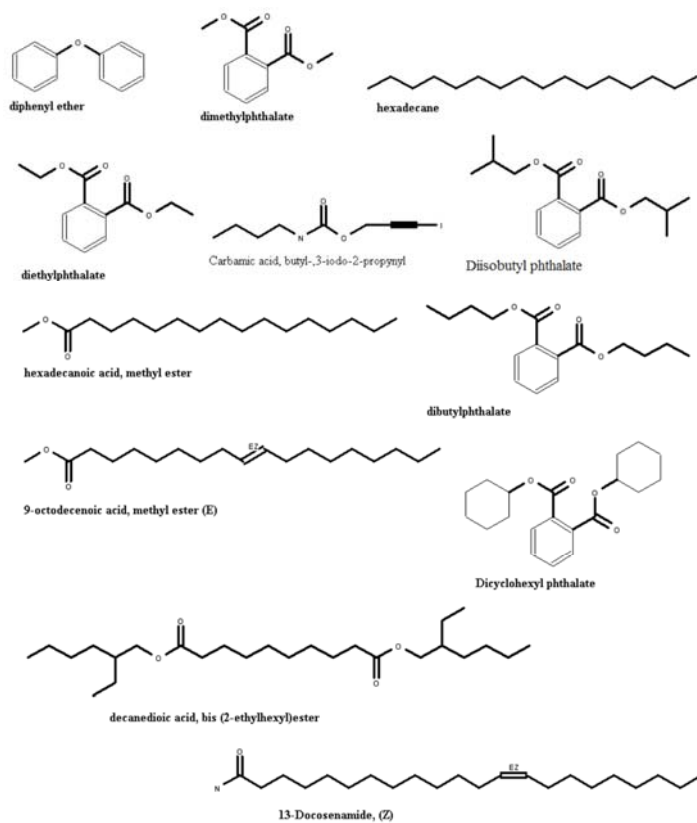


Figure 32. Compounds detected by GC-MS in the reactor's plastic scraps.

Table 11. Compounds detected by GC-MS in the reactor's plastic scraps

Percentage match	Retention time (min)	Compound	Formula
78	11.808	diphenyl ether	C ₁₂ H ₁₀ O
83	13	dimethylphthalate	C ₁₀ H ₁₀ O ₄
97	14.5	hexadecane	C ₁₆ H ₃₄
93	15.304	diethylphthalate	C ₁₂ H ₁₄ O ₄
97	16	Carbamic acid, butyl-,3-iodo-2-propynyl	C ₈ H ₁₂ INO ₂
83	18.901	Diisobutyl phthalate	C ₁₆ H ₂₂ O ₄
99	19.2	hexadecanoic acid, methyl ester	C ₁₇ H ₃₄ O ₂
91	20.107	dibutylphthalate	C ₁₆ H ₂₂ O ₄
92	21.198	9-octadecenoic acid, methyl ester (E)	C ₁₉ H ₃₆ O ₂
78	26.5	Dicyclohexyl phthalate	C ₂₀ H ₂₆ O ₄
91	28.5	decanedioic acid, bis (2-ethylhexyl)ester	C ₂₆ H ₅₀ O ₄
98	28.725	13-Docosamide, (Z)	C ₂₂ H ₄₃ NO

3.7.2. HR Effluent

The analysis revealed that most of the compounds were below 10 µg/L while only two compounds (Bisphenol A and unidentified compounds with RI = 2096) were in the range 14 - 38 µg/L. The chromatograms revealed that butylated hydroxytoluene found in the HR effluent was completely degradable because they were not found in both SAMBRs and the AMBR effluents. However, previous work has shown that butylated hydroxytoluene can leach from plastic and tubings (Shpiner, 2007), but it was not detected in our control sample containing plastic scraps. Similarly, 8,11-octa decadienoic acid, methyl ester (peak ID 8), pentadecanoic acid, 14-methyl, methyl ester (peak ID 7) and tridecanoic acid, 12-methyl, methyl ester (peak ID 5) were three aliphatic molecules that were not detected in the SAMBR permeates due to their complete degradation in this reactor. Surprisingly, padimate O and the phenanthrene carboxylic acid that are aromatics and thus considered as difficult to biodegrade were successfully degraded in the SAMBRs due to the complete retention of

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bacteria. Most of PAHs (polycyclic aromatic hydrocarbons) in leachate are stable and it is difficult to cleave the ring without oxygen. However, recent research has shown that unsubstituted low molecular polycyclic aromatic compounds can be degraded under nitrate-reducing, iron-reducing, sulfate-reducing and methanogenic conditions (MacRae and Hall, 1998; Xu et al., 2008). MacRae and Hall (1998) showed that PAHs could be degraded under denitrifying conditions as long as other nutrients were not limiting. The half-lives of low molecular weight PAH ranged from approximately 33-88 days. Degradation of high molecular weight PAH was slower, or not observed. Half-lives ranged from 143-812 days. Thus, the presence of nitrate from the permeate of the AMBR may have assisted the degradation of the padimate and the phenanthrene under nitrate-reducing conditions in the HR.

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3.7.3. SAMBR1 and 2 Permeates

Table 12 to Table 14 show that o-hydroxybiphenyl and Bisphenol A can be considered as non-biodegradable anaerobically because their areas increased by 9 and 12% in SAMBR1, respectively. In SAMBR2 they increased by 123 and 15%, respectively. The concentration of o-hydroxybiphenyl was found to be 10.2, 11.2 and 22.8 µg/L in the HR, SAMBR1 and SAMBR2, respectively. Bisphenol A (CAS 80057) is a very common compound found in landfill leachate and it is known as a potentially endocrine disrupting industrial chemical (Schwarzbauer et al., 2002; Wintgens et al., 2004). It is found in epoxy resin and is used as a stabilizer for polyvinyl chloride resin. Yasuhara et al. (1999) found that among more than 100 organic compounds, Bisphenol A was at a very high concentration of 61.4 mg/L. Wintgens et al. (2004) and Yamamoto et al. (2001) also found concentrations ranging up to milligrams per litre in raw landfill leachate. Wintgens et al. (2004) stated that the high concentrations can be explained by leaching effects from polycarbonate plastics abundant in the deposited waste. The conditions encountered in the landfill in terms of alkaline pH, mesophilic temperature and high ionic strength might promote the leaching process. They used a pilot plant MBR configuration including denitrification and two-stage nitrification with side-stream

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tubular ultrafiltration that rejected 99% of bisphenol A. They reckon that it seems to be removed most effectively by biodegradation in a system with high sludge age and complete retention of solids.

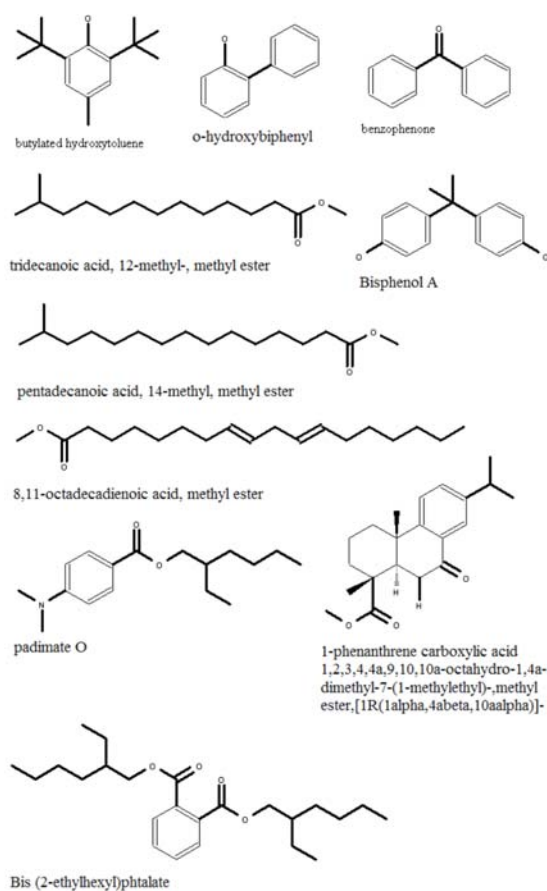


Figure 33. Compounds detected by GC-MS in the HR effluent.

Table 12. Recalcitrant compounds detected by GC-MS in the effluent from the HR. In the case of unidentified compounds, the retention index of the peak is given. Reprinted from (Trzcinski and Stuckey, 2009b)

Peak ID	match %	Retention time (min)	Recalcitrants in HR effluent	Formula	Peak Area
1	93	13.357	butylated hydroxytoluene	C ₁₅ H ₂₄ O	12,281,748
2	94	13.72	o-hydroxybiphenyl	C ₁₂ H ₁₀ O	16,062,321
3		13.91	1462.7		26,872,283
4	93	16.033	benzophenone	C ₁₃ H ₁₀ O	112,030,035
5	93	16.525	tridecanoic acid, 12-methyl, methyl ester	C ₁₅ H ₃₀ O ₂	34,198,411
6		17.486	1691.5		56,558,115
7	99	19.141	pentadecanoic acid, 14-methyl, methyl ester	C ₁₇ H ₃₄ O ₂	62,748,504
8	99	21.113	8,11-octadecadienoic acid, methyl ester	C ₁₉ H ₃₄ O ₂	38,295,039
9		21.271	1964.1		46,078,046
10		21.852	2008.9		95,664,113
11	94	22.481	phenol 4,40-(1-methylethylidene)bis	C ₁₅ H ₁₆ O ₂	570,361,325
12		22.978	2096.1		411,285,760
13	87	24.052	padimate O	C ₁₇ H ₂₇ NO ₂	101,864,975
14	95	24.39	1-phenanthrene carboxylic acid 1,2,3 m,4,4a,9,10,10a-octahydro-1, 4a-dimethyl-7-(1-methylethyl)-,methyl ester,[1R(1alpha,4abeta, 10aalpha)]	C ₂₁ H ₂₇ O ₃	93,127,843
15	72	26.179	Bis (2-ethylhexyl)phthalate	C ₂₄ H ₃₈ O ₄	171,161,815

Several authors have observed that the concentration of Bisphenol A tends to decrease over time in a landfill (Asakura et al., 2004). In our study, bisphenol A was the main peak in the HR effluent at a concentration estimated to be in the range 19-38 µg/L, while in the SAMBR1 and SAMBR2 its concentration increased to 21-44 µg/L. On the other hand, these authors have observed that the concentration of Bis (2-

ethylhexyl)phthalate remained at the same level. In our study it was not detected in the blank and scrap, and its area more than doubled from the HR effluent to SAMBR1 (+283%) and SAMBR2 (+179%) permeates, suggesting that it could be secreted by bacteria themselves, or is the catabolic end product of non-detected compounds. A calibration line was obtained for Bis (2-ethylhexyl)phthalate ($R^2 = 0.99$) and it revealed that the concentration in SAMBR 1 and SAMBR2 was 1.4 and 1 mg/L, respectively. The concentration in the HR and the AMBR was found to be 0.33 and 0.3 mg/l, respectively. Some molecules were found to be slowly biodegradable because their areas decreased when passing through both SAMBRs. These molecules were benzophenone which decreased by 81 and 73% in SAMBR1 and 2, respectively, plus unidentified compounds with retention indices 1462.7 and 2096.1 in SAMBR1 and 2008.9 and 2096.1 in SAMBR2.

3.7.4. SAMBR2 and AMBR Permeates

In comparing the SAMBR2 and AMBR permeates, it can be seen that phenol 2,4-bis(1,1-dimethylethyl) and N-phenethylbenzenesulfonamide were not degraded aerobically because their areas were found to increase when passing through the AMBR: their areas increased by 45 and 786%, respectively. The concentration of the latter reached 44-87 $\mu\text{g/L}$ in the AMBR. Mansouri et al. (2007) also found that phenol 2,4-bis(1,1-dimethylethyl) was among the components that remained after an aerobic fixed bed process treating landfill leachate. As Bisphenol A was not found in the AMBR permeate, it can be stated that it was fully biodegraded aerobically. This is in line with Asakura et al. (2004) who treated raw leachate by aeration and found that bisphenol A decreased from 70.9 - 224 $\mu\text{g/l}$ to 0.11- 0.24 $\mu\text{g/l}$ in the effluent. This is also in line with the Biocatalysis / Biodegradation database from the University of Minnesota (<http://umbdd.msi.umn.edu/index.html>) that states that Bisphenol A is metabolized by a Gram-negative aerobic bacterium (strain MV1) via a novel pathway. The metabolism involves two primary pathways: a minor pathway, in which Bisphenol A is first oxidised to a triol, and a major pathway, in which Bisphenol A is metabolized to form the intermediates 4-

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hydroxybenzoate and 4-hydroxyacetophenone that are converted to carbon dioxide and biomass (Spivack et al., 1994). Other intermediates of the degradation pathway of Bisphenol A include toluene and vanillin. Interestingly, some molecules were found to be non-biodegradable in an anaerobic environment, but could be slowly biodegraded in the AMBR such as 2,5-cyclohexadien-1-one,2,6 bis(1,1-dimethylethyl)-4-ethylidene- and Bis (2-ethylhexyl)phthalate. The former decreased by 68% whereas the latter decreased by 67% in the AMBR. Bis (2-ethylhexyl)phthalate often referred as DEHP is the most common plasticizer (Bauer and Herrmann, 1997).

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Table 13. Recalcitrant compounds detected by GC-MS in the permeate of SAMBR1. In the case of unidentified compounds, the retention index of the peak is given. The last column comments on the anaerobic biodegradability by comparing the peak area of the same compound in the HR effluent. NB = Non-biodegradable, B = Biodegradable

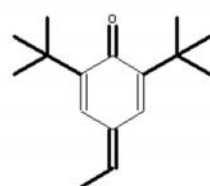
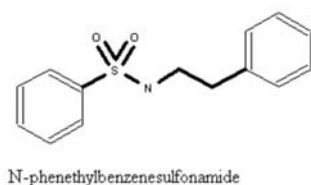
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Peak ID	match %	Ret. time	Recalcitrants in SAMBR1 permeate	Formula	Peak Area	Anaerobic Biodegradability
16		11.68	1333.7		26,996,170	New
2	94	13.72	o-hydroxybiphenyl	C ₁₂ H ₁₀ O	17,502,568	NB (+9%)
3		13.91	1462.7		14,119,684	B (-48%)
4	83	16.03	benzophenone	C ₁₃ H ₁₀ O	21,735,256	B (-81%)
17		18.21	1742.3	C ₁₅ H ₈ O ₂	15,836,896	New
18	78	18.99	n-phenethyl-benzesulfonamide		208,608,266	New
19	78	20.23	2,5-cyclohexadien-1-one,2,6 bis(1,1-dimethylethyl)-4-ethylidene-	C ₁₄ H ₁₅ NO ₂ S	44,712,759	New
11	94	22.48	Phenol 4,40-(1-methylethylidene)bis	C ₁₆ H ₂₄ O	640,395,425	NB (+12%)
12		22.98	2096.1		179,375,192	B (-56%)
15	91	26.18	Bis (2-ethylhexyl)-phthalate	C ₂₄ H ₃₈ O ₄	655,807,845	NB (+283%)
20	95	27.133	Phosphine imide, P,P,P-triphenyl	C ₁₈ H ₁₆ NP	123,496,720	New

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Table 14. Recalcitrant compounds detected by GC-MS in the permeate of SAMBR2. In the case of unidentified compounds, the retention index of the peak is given. The last column comments on the anaerobic biodegradability by comparing the peak area of the same compound in the HR effluent. NB = Non-biodegradable, B = Biodegradable

Peak ID	match %	Ret. time	Recalcitrants in SAMBR2 permeate	Formula	Peak Area	Anaerobic Biodegradability
21	93	12.4	phenol 2,4-bis(1,1-dimethylethyl)	C ₁₄ H ₂₂ O	11,305,194	New
2	94	13.72	o-hydroxybiphenyl	C ₁₂ H ₁₀ O	35,846,178	NB (+123%)
22	94	15.69	benzenemethanol, alpha-phenyl	C ₁₃ H ₁₂ O	36,535,074	New
4	94	16.03	benzophenone	C ₁₃ H ₁₀ O	29,918,768	B (-73%)
18	78	18.99	N-phenethylbenzenesulfonamide	C ₁₄ H ₁₅ NO ₂ S	147,543,256	New
19	78	20.23	2,5-cyclohexadien-1-one,2,6bis(1,1-dimethylethyl)-4-ethylidene-	C ₁₆ H ₂₄ O	44,712,759	New
10		21.85	2008.9		46,417,418	B (-50%)
11	95	22.48	phenol 4,40-(1-methylethylidene)bis	C ₁₅ H ₁₇ O ₂	655,737,185	NB (+15%)
12		22.98	2096.1		262,561,636	B (-36%)
15	91	26.18	Bis (2-ethylhexyl)phthalate	C ₂₄ H ₃₈ O ₄	478,178,027	NB (+179%)



2,5-cyclohexadien-1-one,2,6bis(1,1-dimethylethyl)-4-ethylidene-

Figure 34. New compounds detected by GC-MS in SAMBR1 permeate.

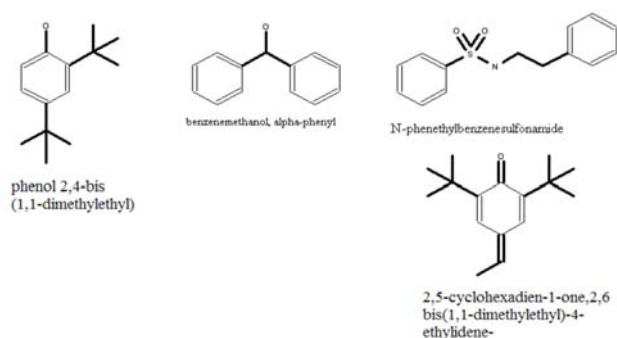


Figure 35. New compounds detected by GC-MS in the permeate of SAMBR2.

Several authors have observed that the concentration of Bis (2-ethylhexyl)phthalate remains constant in a landfill (Asakura et al., 2004; Sakamoto et al., 2000). Because Bis (2-ethylhexyl)phthalate has a high octanol/water partition coefficient, it can be considered as hydrophobic and therefore is expected to be removed when suspended solids are removed. In other words it tends to attach to biomass more than hydrophilic compounds. Asakura et al. (2004) found that it could not be removed by aeration, coagulation/sedimentation nor biological treatment. It has been reported elsewhere that DEHP can be degraded by a common soil bacterium (*Rhodococcus rhodochrous*) with an easily used carbon source, and that the final metabolites are 2-ethylhexanol, 2-ethylhexanoic acid and phthalic acid (Horn et al., 2004; Nalli et al., 2003). Moreover, 2-ethylhexanoic acid was observed to be particularly resistant to further degradation and exhibited acute aquatic toxicity. However, in our study the DEHP concentration increased in the SAMBRs, whereas its concentration was lower in the AMBR, but none of the metabolites given in the paper of Horn et al. (2004) were found. This could be due to a different metabolic pathway in the AMBR or because of their hydrophobicity; these metabolites were preferentially attached to the sludge and hence were not detected. Nevertheless, new molecules appeared in the AMBR permeate such as diisobutyl phthalate and an unidentified peak with a retention index

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of 1510.6. Moreover, the calibration line of Bis (2-ethylhexyl)phthalate was used to calculate the concentration of recalcitrants in the SAMBR2 and AMBR permeate. Then the COD equivalence of each compounds were found to be very low when added together. As a matter of fact, the COD concentrations of recalcitrants in SAMBR2 and AMBR permeate was found to be equal to 17.7 and 20 mg/L, respectively, which contrast with the respective total COD concentrations of about 400 and 300 mg/L. Section [6.3.53.4.1](#) contains details indicating that recalcitrants proteinaceous compounds in the SAMBR2 and AMBR permeates accounted for about 100 and 50 mg/l, which is equal to 150 and 75 mg COD/L. Therefore, about 250 and 225 mg/L of the permeates are still unknown compounds. The use of GC-MS is limited to the identification of non-polar, volatile and thermostable compounds which explains why so little information was found in this study. Techniques such as LC-MS or Matrix Assisted Laser Desorption Ionization-Time of Flight-Mass Spectrometry (MALDI-ToF-MS) would certainly shed more light on the nature of the high MW compounds that were not detected.

3.7.5. *Phthalates and Plasticizers*

Plasticizers are compounds that are added to polymers in order to improve the properties of a plastic such as increasing its flexibility, and several phthalates were detected in this study. It has been demonstrated that plasticizers tend to leach from solid polymer matrices into the environment (Fromme et al., 2002). For instance dimethylphthalate was found in the reactor plastic scrap but was not detected in the reactor indicating that it could be readily biodegraded. Diethylphthalate was found at a similar level in the scrap and in the HR effluent. The fact that it was not detected in the SAMBR permeates indicates that it could be biodegraded completely thanks to the long solid retention times achieved in SAMBRs. Dibutylphthalate was found in the anaerobic reactors but also in the scrap suggesting that it might come from the reactors plastic. Interestingly, its area decreased greatly in the SAMBRs (from 270 million in HR effluent to 25.6 and 19.9 million in SAMBR1 and 2 permeate, respectively), and was absent in the AMBR indicating that a great

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proportion of it can be degraded anaerobically and totally degraded aerobically.

Table 15. Recalcitrant compounds detected by GC-MS in the permeate of the AMBR. In the case of unidentified compounds, the retention index of the peak is given. The last column comments on the aerobic biodegradability by comparing the peak area of the same compound in the SAMBR2 permeate. NB = Non-biodegradable, B = Biodegradable. Reprinted from (Trzcinski and Stuckey, 2009b)

Peak ID	match %	Ret. time	Recalcitrants in AMBR permeate	Formula	Peak Area	Anaerobic Biodegradability
21	91	12.4	phenol 2,4-bis (1,1-dimethylethyl)	C ₁₄ H ₂₂ O	16,343,590	NB (+45%)
23		14.7	1510.7			New
18	78	18.99	N-phenethyl-benzenesulfonamide	C ₁₄ H ₁₅ NO ₂ S	1,307,129,3 72	NB (+786%)
24	81	20.12	1,2-benzenedi carboxylic acid, butyl (2-methylpropyl)ester	C ₁₆ H ₂₂ O ₄	21,287,952	New
19	78	20.23	2,5-cyclohexadien-1-one, 2,6bis (1,1-dimethylethyl)-4-ethylidene-	C ₁₆ H ₂₄ O	14,309,919	B (-68%)
15	91	26.18	Bis (2-ethylhexyl)phthalate	C ₁₆ H ₃₈ O ₄	156,613,458	B (-67%)

3.8. DENATURING GEL GRADIENT ELECTROPHORESIS (DGGE)

3.8.1. Archaea

Archaeal DNA was amplified using the nested **polymerase chain reaction** (PCR) reaction. First, the **DNA** template was amplified using the **primers** 46F and 1017R. Then 1 microliter PCR product was used for the second PCR reaction using the primers 344FGC and Univ522. The objective was to investigate the differences if any between the archaeal

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population in the HR and the SAMBR and the effect of the HRT on their evolution. As Figure 36 shows, there were only three visible bands and with the exception of lane 5 the bands were more dominant in SAMBR1 (Lanes 5 to 10). Previous trials with archaeal primers gave very faint bands as is in Figure 36. Therefore, to be able to cut out some bands and identify any species, it was decided to run three PCR reactions in parallel for the sample SAMBR1 day 137 (lane 8) and to load the 3 times concentrated PCR product on the DGGE gel and thereby obtaining brighter bands. This was indeed a good but time-consuming method to get bright bands as it can be seen that bands in lane 8 (SAMBR1, day 137) are much brighter than in the other lanes. It is believed that the bands in other lanes were very faint because of the small amount of DNA template used in the PCR reaction; as a result all the archaeal species present in the bioreactor did not appear on the DGGE gel. Nevertheless, the presence of bands indicates the predominant species in the reactor. Bands a,b and c were cut from lane 8 but only band b could be identified successfully as an organism from the genus *Methanotherix* named *Methanosaeta* sp. clone DI C03 (Accession number AY454761.1, 100% match, E value = 0.82). Because *Methanosarcina* was also found in the experiment described in the next chapters, it was hypothesized that band c corresponded to *Methanosarcina*. If this is true it would mean that *Methanosarcina* sp. develop preferentially in the HR where the acetate concentration is higher, whereas *Methanosaeta* would be dominant in the SAMBR where acetate concentration is very low. This is in line with Ehlinger et al. (1987) who stated that *Methanosarcina* sp. develop preferentially throughout or in the inlet of reactors, wherever the acetate concentration was equal to or higher than 350 mg L⁻¹. When the acetate concentration falls below 350 mg/L, *Methanotherix* sp. becomes the prevailing species. In the HR (lanes 1 to 5), only band c could be seen except on day 203 (Lane 5) where bands a and b also appear. The appearance of *Methanosaeta* in the HR could be due to the low acetate concentration towards the end (below).

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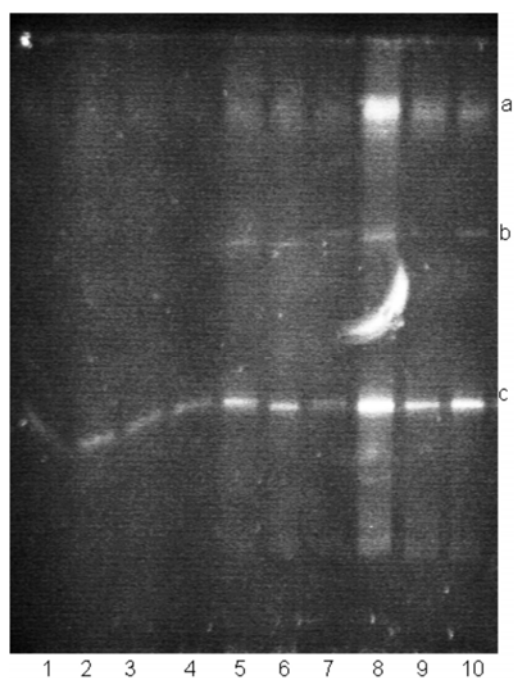


Figure 36. Evolution with time of the archaeal population in the HR (Lane 1 to 5) and SAMBR1 (Lane 6 to 10). Lanes 1 and 6 = day 76; Lanes 2 and 7 = day 119; Lane 3 and 8 = day 137; Lane 4 and 9 = day 178; Lane 5 and 10 = day 203.

3.8.2. *Bacteria*

Bacterial DNA was amplified using a nested PCR reaction. First, the DNA was amplified using the primers 27F and 1492R. Then 1 microliter of PCR product was used as template for the second PCR using the primers 338FGC and 518R. Again the bands were quite faint even with triplicate PCR products with this combination of primers. Nevertheless, it can be seen from Figure 37 that the SAMBR2 started up on day 46 (Lane 5) with no dominant bacteria although that sample contained DNA as it was checked on agarose gel beforehand. On day 119 (Lane 4), three distinct

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dominant species of bacteria (bands a, c and d) could be distinguished, and as the HRT was reduced the number of bands also diminished so that only two bands (c and d) were visible on day 137 (lane 3) and later only 1 band (b) in lanes 1 and 2. Moreover, there was a significant shift in bacterial population as the HRT was decreased because band b appeared only in lane 1 and 2, i.e., when the HRT was less than 4 days. Bands a, b, c and d were cut and after cloning and sequencing as described in Materials and Methods, Table 16 was drawn.

Table 16. Bands cut from the DGGE gel using bacterial primers

Band	Closest Match	Accession Number	Genus	Match %	E value
a	Uncultured bacterium	U81676.2	Uncultured Bacteroidales	100	1.10 ⁻⁹⁵
b	Uncultured bacterium	EF589977.2	Uncultured Bacteroidales	98	2.10 ⁻⁹²
c	Uncultured bacterium	AY693829.1	Uncultured Bacteroidales	100	1.10 ⁻⁹⁵
d	Uncultured bacterium	EF559220.1	Uncultured Bacteroidales	100	1.10 ⁻⁹⁵

As the bands on DGGE gel were quite faint using the nested PCR method mentioned above, another combination of primers was tested: 341FGC and 907R.

Figure 38 shows the same samples amplified with the nested PCR in triplicate on the left hand side and the samples amplified using 341FGC and 907R on the right hand side. It is clear that the latter combination gave rise to much brighter bands on an agarose gel, and the PCR products were therefore loaded on a DGGE gel in Figure 38. It can be seen from Figure 39 that there is no band in lane 11 (day 46) which is when SAMBR2 was started.

The bands could be seen only from lane 12 (day76) onwards and became brighter on lane 13 (day119). This highlights the fact that SAMBR2 did not start with the proper bacterial population and it took at least 30 days to get a similar population as SAMBR1 in lane 6. This lack of

bacteria for start-up was correlated with an increase in bulk SCOD as explained above. Unfortunately, the bright band at the top of the gel which is common to all lanes indicates that some DNA could not migrate through the gel. This is because 40% as a lower gradient was already too high. Another gel with a denaturing gradient of 30-70% was run but the bands were all compressed at the top of the gel. An optimum gradient for these PCR products would have been 20-60%.

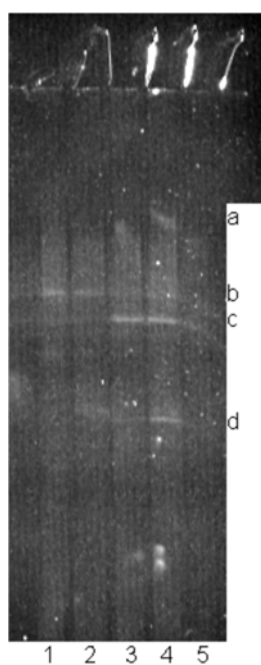


Figure 37. Evolution with time of the bacterial population in SAMBR2 using nested PCR 27F and 1492R then 338FGC and 518R. Lane 1 = day 203; Lane 2 = day 178; Lane 3 = day 137; Lane 4 = day 119; Lane 5 = day 46.

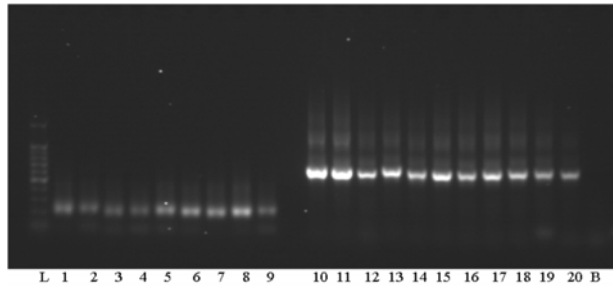


Figure 38. Comparison between the nested PCR products (27F and 1492R then 338FGC and 518R) on the left hand side and the PCR products using primers 341FGC + 907R on the right hand side. L=Ladder; Lane 1=SAMBR1 day119; Lane 2=SAMBR1 day 137; Lane 3=SAMBR1 day 178; Lane 4=SAMBR1 day 203; Lane 5=HR day 76; Lane 6=HR day 119; Lane 7=HR day 137; Lane 8=HR day 178; Lane 9=HR day 203; Lane 10=HR day 137; Lane 11=HR day 178; Lane 12=HR day 203; Lane 13=SAMBR1 day 137; Lane 14=SAMBR1 day 178; Lane 15=SAMBR2 day 137; Lane 16=SAMBR2 day 178; Lane 17=SAMBR2 day 203; Lane 18=AMBR day 137; Lane 19=AMBR day 178; Lane 20=AMBR day 203; B=blank.

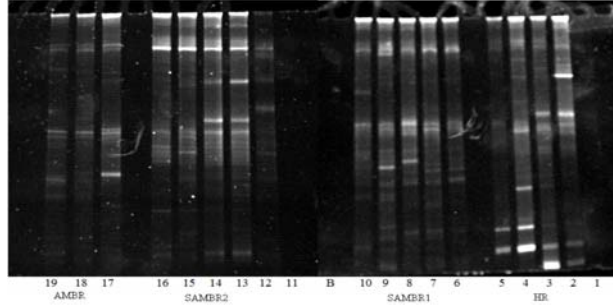


Figure 39. Evolution with time of the bacterial population in the HR (Lanes 1 to 5), SAMBR1 (Lanes 6 to 10), SAMBR2 (Lanes 11 to 16) and AMBR (Lanes 17 to 19) using primers 341FGC and 907R. Lane 1=HR day 76; Lane 2=HR day 119; Lane 3=HR day 137; Lane 4=HR day 178; Lane 5=HR day 203; Lane 6=SAMBR1 day 76; Lane 7=SAMBR1 day 119; Lane 8=SAMBR1 day 137; Lane 9=SAMBR1 day 178; Lane 10=SAMBR1 day 203; B=blank; Lane 11=SAMBR2 day 46; Lane 12=SAMBR2 day 76; Lane 13=SAMBR2 day 119; Lane 14=SAMBR2 day 137; Lane 15=SAMBR2 day 178; Lane 16=SAMBR2 day 203; Lane 17=AMBR day 137; Lane 18=AMBR day 178; Lane 19=AMBR day 203.

3.8.3. Ammonia - Oxidisers

The DNA from ammonia-oxidising bacteria (AOB) was amplified using the primers CTO189F-GC and CTO654R. Bands a, b, c and d were cut from the DGGE gel in Figure 40 and it was found that they all belong to the genus *Nitrosomonas*, except band d which belong to the genus *Nitrospira*. The details of the bands are given in Table 17. It is clear from Figure 40 that there was no AOB species in Lane 4 which corresponds to a few days after the AMBR was started up. This explains why there was no ammonia conversion in the AMBR at the beginning. Then on lane 5 (day 178), two clear bands a (uncultured *Nitrosomonas sp.*) and b (*Nitrosomonas europaea* ATCC 19718) can be seen as well as a faint band c (*Nitrosomonas sp.* NM 41). Eventually on day 203 (lane 6), bands b and c became more dominant as the HRT was below 1 day. It is noteworthy that bands a, b and c are all from the genus *Nitrosomonas* and all are *Nitrosomonas europaea* - like species which are well known AOB. Interestingly, a band was found in the HR but only on day 203, at the end of the run. This band d was found to be *Nitrospira sp.* Nsp40 which is an AOB and was not found in the AMBR. This is, however, controversial as heterotrophs are known to be fast growing microorganisms and AOB would have been unable to compete with them for oxygen present in the feedstock fed once a day in the HR. It is suggested that at low DO levels, AOB used nitrite or nitrate as an artificial electron acceptor and generate nitrous oxide gas (Ritchie and Nicholas, 1972).

The *Nitrospira* and *Nitrobacter* NOB species were not detected because the primers used were specific to AOB. However, these NOB species were expected to be present in the AMBR because nitrite conversion to nitrate was observed from day 167 onwards. To gain more insight into the nitrification process, it would have been useful to use primers specific to NOB such as NSR1113F and NSR1264R for *Nitrospira* (Wang et al., 2007). Alternatively, Wittebolle et al. (2008) have used a nested PCR where the first PCR amplifies specifically AOB whereas the second PCR is performed with total bacterial primers (338FGC and 518R) that will amplify NOB as well. It is possible that some AOB were missed due to the gradient (38-50%) used to make up the gel. Other studies have

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used larger gradients (45-60%) when using primers CTO189FGC and CTO654R (Wittebolle et al., 2008). It can be seen that band a is already at the top of the gel and it is therefore possible that the low gradient (38%) was too high and did not allow some bands to migrate and remained at the top of the gel. Also band d was at the bottom of the gel, and thus it is possible that some bands were missed because they migrate further than band d due to the high gradient (50%) being too low.

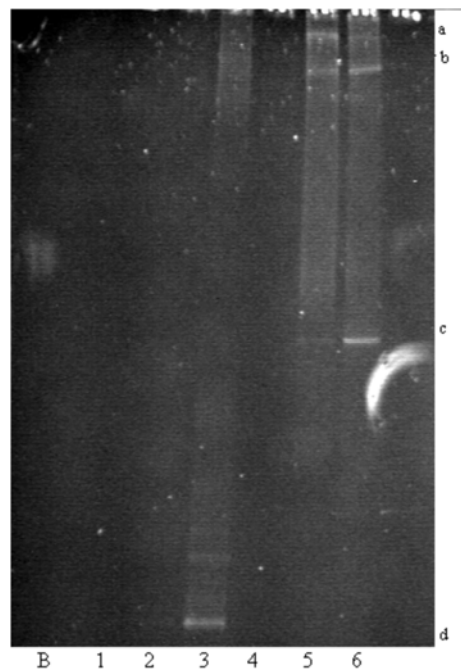


Figure 40. Evolution with time of the population of AOB in the HR and the AMBR using the primers CTO189FGC and 654R. B=Blank; Lane 1=HR day137; Lane 2=HR day178; Lane 3=HR day203; Lane 4=AMBR day137; Lane 5=AMBR day178; Lane 6=AMBR day203.

Table 17. Bands cut from the DGGE gel using AOB primers

Band	Closest Match	Accession Number	Genus	Match %	E value
a	<i>Uncultured Nitrosomonas sp.</i>	AJ307985.1	<i>Nitrosomonas</i>	98	0
b	<i>Nitrosomonas europaea ATCC 19718</i>	AL954747.1	<i>Nitrosomonas</i>	98	0
c	<i>Nitrosomonas sp. NM41</i>	AF272421.1	<i>Nitrosomonas</i>	97	0
d	<i>Nitrospira sp. Nsp40</i>	AY123787.1	<i>Nitrospira</i>	100	0

CONCLUSION

The following conclusions can be drawn from this study:

- The use of a membrane for leachate treatment had several advantages; the complete retention of bacteria allowed for stable operation as no VFAs accumulated even when propionate was the predominant acid in the leachate. The COD removal was greater than 90% at a HRT of 1.6-2.3 days at a maximum OLR of 20 g COD/L.day. Even though the influent COD of the leachate was constantly changing giving rise to a transitory F/M ratio over time, the permeate COD from the SAMBR was typically between 300 and 500 mg/L which can therefore be defined as a stabilised leachate. Because of the fluctuating properties of the leachate produced in the HR the process was deemed more representative because the SAMBR treated a leachate with varying organic strength which is what can happen on a full scale (Kraume et al., 2009).
- An inoculum fed on synthetic VFAs was not suitable for start-up because it did not contain hydrolytic and acidogenic bacteria for a leachate medium. As a result, a slow production of VFAs, and therefore a slow increase in methane production were observed. Furthermore, the bulk SCOD in SAMBR2 kept increasing during

start-up, which strongly suggests that, for a lignocellulosic-based feed, it is paramount to start up the SAMBR with a hydrolytic population acclimatised to the leachate medium to avoid SCOD build up in the bulk. The DGGE picture confirmed this observation.

- The COD removal in SAMBR2 was 94.5% on average, and only 1.6% in the AMBR so that a total COD removal of 96.1% was achieved at 0.4 day HRT. On average 26% of the recalcitrants from SAMBR2 could be degraded aerobically in the AMBR.
- However, as in SAMBR1, at HRTs lower than 2 days, particulate solids in the leachate built up at the bottom of the SAMBR eventually leading to the diffuser blocking. At MLTSS beyond 20 g/L, the TMP culminated at 850 mbar and the flux dropped to 0.5 LMH until the end of the experiment.
- The permeate COD cannot be lowered by decreasing the sparging rate. The main effect of low sparging rate (2LPM) was the enhanced rejection of the membrane and the significant flux drop while the permeate COD remained relatively constant. On the other hand, when the sparging rate was increased to 8-10 LPM, the permeate COD increased typically from 360 to 440 mg/L due to a better scouring of the membrane, which resulted in a thinner concentration polarization layer and a better diffusion through the biofilm. It is likely that the time required to observe the increase in permeate COD depends on the compaction of the cake layer. The increase in sparging rate allowed to recover the operation at 0 mbar TMP but not the initial flux.
- The permeate of the SAMBR was low in COD and relatively constant which promoted the growth of autotrophic bacteria in the subsequent AMBR so that 97.7% of the $\text{NH}_4^+\text{-N}$ was removed at a maximum nitrogen loading rate of 0.18 kg $\text{NH}_4^+\text{-N}/\text{m}^3\cdot\text{day}$. Complete nitrification was not observed in the AMBR as the nitrite concentration did not reach zero. This could be due to the low HRT (1 day) that could have overloaded the NOB such as *Nitrobacter*, or because of the lack of close distance between the

nitrite producers and nitrite oxidisers which is required to obtain complete nitrification.

- Calcium in the leachate was found to precipitate in the AMBR because of the higher pH. A sample taken from the membrane consisted most likely of pure hydroxylapatite $\text{Ca}_5(\text{PO}_4)_3(\text{OH})$ which had a needle shape, whereas the background of the precipitate consisted more of nodules of calcium carbonate with traces of manganese, iron, magnesium, aluminium, sulphur and sodium.
- The very high MW aromatic organics in the leachate fed to the SAMBRs were almost fully degraded in the bulk of the SAMBRs. Moreover, their permeate was absolutely free of them which indicates a full rejection of these compounds by the membrane. The medium MW compounds in the range of 395 - 646 kDa were more likely to be rejected by the membrane while the $\text{MW} \leq 395$ kDa were observed in the permeate. Regarding the evolution of the medium and low MW compounds over the 200 days, it can be stated that overall there was no build-up in the absorbance in any effluent, showing that these aromatics could be slowly degraded.
- Most of the recalcitrant compounds detected by GC-MS in the effluent of the HR were below $10 \mu\text{g/L}$, while only two compounds (Bisphenol A and unidentified compounds with RI = 2096) were in the range $14 - 38 \mu\text{g/L}$. The molecules o-hydroxybiphenyl and Bisphenol A were found to be non-biodegradable anaerobically because their areas increased in SAMBR1 and SAMBR2. The concentration of the former was in the range $11 - 22 \mu\text{g/L}$, while the latter was in the range $21-44 \mu\text{g/L}$. Moreover, as Bisphenol A was not found in the AMBR permeate, it can be stated that it was fully biodegraded aerobically. Some molecules were found to be non-biodegradable in an anaerobic environment, but could be slowly biodegraded in the AMBR such as 2,5-cyclohexadien-1-one, 2,6-bis(1,1-dimethylethyl)-4-ethylidene- and Bis(2-ethylhexyl) phthalate. Nevertheless, the component N-phenethylbenzenesulfonamide

were found to build-up in the AMBR and reached concentrations in the range 44-87 µg/L. Furthermore, new molecules appeared in the AMBR permeate such as Diisobutyl phthalate.

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