Adapting anaerobic consortium to pure and complex lignocellulose substrates at low temperature: kinetics evaluation

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Received: 15 May 2018 / Accepted: 8 December 2018 / Published online: 15 December 2018 © The Author(s) 2018

Abstract

Purposes The purpose of this research was to evaluate the kinetics of anaerobic microbial culture during adaptation to pure and complex lignocellulosic substrates at low temperature.

Methods Six pairs of 1.0 L batch reactors maintained at 20 °C were fed pure (xylan and cellulose) and complex (cow manure and wheat straw) lignocellulosic substrate in three successive cycles of 35 days each. The biogas volume and composition, chemical oxygen demand, and volatile solids were monitored to evaluate the kinetics of the culture during the adaptation.

Results Anaerobic culture adapted to digest the pure and complex lignocellulosic substrates at 20 °C in relatively short period (105 days; 3 successive cycles of 35 days each) in batch reactor studies. The first-order model kinetics revealed that the average increase (day\(^{-1}\) cycle\(^{-1}\)) in the reaction rate constant over the successive cycles was 0.0831 (xylan) > 0.0235 (cow manure) > 0.0207 (wheat straw) > 0.0123 (xylan:cellulose mixture) > 0.0041 (cellulose). The rates of the substrate degradation at 20 °C were: 0.085–0.093 day\(^{-1}\) (cellulose), 0.112–0.278 day\(^{-1}\) (xylan), 0.112–0.137 day\(^{-1}\) (xylan:cellulose mixture), 0.069–0.116 day\(^{-1}\) (cow manure), and 0.057–0.106 day\(^{-1}\) (wheat straw).

Conclusions Anaerobic mixed culture can be adapted to pure and complex lignocellulosic substrates and convert them to methane at low temperature (20 °C) in relatively short time (105 days) using a sequential procedure. The culture adaptation to wheat straw proceeded at a slower rate than that for cow manure.

Keywords Adaptation · Low temperature · Anaerobic digestion · Lignocellulose · Cow manure · Kinetics

Introduction

Lignocellulosic agriculture residue and farm waste are abundant biomass of high potential to serve as renewable feedstock for sustainable biogas production and value-added products; however, they are currently undervalued (Paul and Dutta 2018). Anaerobic digestion is a low-cost microbial conversion process (Lynd et al. 2008) which converts biodegradable organic substrates such as manure and farm lignocellulosic waste biomass into methane (Lynd et al. 2008).

Biogas production through anaerobic digestion of lignocellulose substrates such as cellulose, hemicellulose, cow manure, and wheat straw have been reported in mesophilic and thermophilic processes. However, very little is known about such application at low temperature.

Anaerobic digestion is sensitive to the environmental and operating conditions. Growth conditions such as temperature, pH, type of substrate, reactor design and configuration, mode of operation, and other maintenance procedures impose various selective stresses on microorganisms (Angert et al. 1998; Boivin-Jahns et al. 1995; Ferris et al. 1996). A very small fraction of microorganisms may benefit from these stresses and thus dominate the culture and contribute to its intended functionality. Fastidious microorganisms such as mono-trophic species as well as microorganisms which function in symbiotic interaction with other species will likely be screened out upon the elimination of their substrate or their symbiotic partners, respectively.

Upon exposure to different conditions than the ones prevalent in their indigenous environment, microorganisms
may adapt by changing their physiological state. Introducing 10% sucrose in the growth medium of a mixed anaerobic culture maintained on a mixture of acetate and propionate decreased its methanogenic activity by 30–70% (Dolfing and Bloeman 1985). After 3 months, the methanogenic activity of the culture reflected its growth media composition rather than its original activity before adaptation (Dolfing and Bloeman 1985).

Feeding the culture a single new substrate imposes stress on the various microorganisms and alters their proportions in the reactor. Consequently, the proportion of some trophic groups may increase, decrease, or even disappear. Upon such effects, the culture may transform into a new one. Adapting anaerobic mixed culture on lignocellulosic substrates at low temperature may come at the expenses of fats- and protein-metabolizing microorganisms (Gavala and Lyberatos 2001; Gavala et al. 1999; Guieysse et al. 2001). Precisely, adaptation may induce a population shift from the substrate sensitive to the substrate tolerant microorganisms. Moreover, the kinetics of the culture may change as well. Gavala and Lyberatos (2001) reported that adaptation affected the consumption and kinetics (the maximum specific utilization rate) of various compounds in the mixed anaerobic culture significantly. The adaptation affects the stoichiometry of the reactions which produce the volatile fatty acids during the acidogenesis and acetogenesis (Gavala and Lyberatos 2001).

Temperature is an important factor because it affects both the growth of microbes and the thermodynamics of the bioreactions. The maximum growth rate of microbes increases with temperature. Generally, the growth rates double for each 10 °C increase in temperature over the range of 10–35 °C (Rittmann and McCarty, 2001). Therefore, upon adaptation to low temperature the culture’s growth rate decreases (van Lier et al. 1997). The growth rate of microbes at psychrophilic conditions is less than that at mesophilic conditions.

Methane formation reactions are carried out by two groups of microorganisms: bacteria and archaea (methanogens). Temperature affects the growth and activity of both groups. Lowering the temperature decreases the activity of hydrolytic bacteria thus lowers the rate of hydrolysis. Hydrolysis is generally found to be the rate-limiting step in anaerobic degradation (Dhaked et al. 2010). On the other hand, methanogens are also temperature-dependent (Nozhevnikova et al. 2001). Lowering the temperature may change the dominating metabolic pathway in the microbial culture, particularly methanogens.

To maintain the same methane production output and feedstock loading for anaerobic digestion at low temperatures it is possible to increase the digester volume to provide longer solids retention time (SRT). This accounts for the decreased microbial activity and the slow microbial growth rate. Alternatively, the organic loading rate (OLR) should be decreased.

Thermodynamically, the reaction free energy is a function of the temperature. The free energy of a reaction increases with decreasing temperature. A reaction becomes thermodynamically favorable or feasible when its free energy is negative. Anabolic reactions, which form new cell biomass, have positive free energy. Therefore, they have to be coupled with catabolic reactions (substrate degradation) which have negative free energy so that the overall reaction becomes thermodynamically favorable (von Stockar et al. 2006). Methane formation is growth-associated bioreaction. Although methane production occurs at temperatures near zero (Zeeman et al. 1988) the growth rate is very slow and methane yield is low.

Increasingly, psychrophilic anaerobic digestion is shown to be a promising and an economical process for animal manure and farm residue treatment in cold regions (Gunnigle et al. 2015; Massé et al. 2010; Safley and Westerman 1994). However, the biotechnological potential of psychrophilic anaerobic digestion is underutilized (Kasana and Gulati 2011). Psychrophilic culture could evolve from thermophilic and mesophilic cultures through adaptation and enrichment. Analogously, adaptation and enrichment of mixed anaerobic culture could be used to develop and intensify the lignocellulose (cellulose and hemicellulose) degrading activity in the culture. Although some studies reported biomethane production from lignocellulosic substrates at mesophilic and thermophilic conditions (Adl et al. 2015; Janke et al. 2015; Kang et al. 2014), very few studies (Saady and Massé 2013) have been published on psychrophilic anaerobic digestion of cellulose and hemicellulose. Little is known about the culture kinetics when integrating low temperature with feeding lignocellulose substrate.

Currently, lignocellulose substrates and their by-products such as cellulose and xylan (hemicellulose) are investigated extensively to improve the technologies of their microbial conversion to biogas, bioethanol, biobutanol, etc. Dairy manure is composed of heterogenous undigested or partially digested lignocellulosic biomass. For example, the by-mass percentage of cellulose, hemicellulose, and lignin in dairy manure is 25%, 13–20%, and 15%, respectively (Yue et al. 2011). Also, straw such as wheat straw is usually used as a bedding material in livestock farms.

The objective of this study was to assess the adaptation of the anaerobic microbial consortium through the kinetic of biodegradation of lignocellulosic substrates [hemicellulose (xylan), cellulose, xylan:cellulose mixture, cow manure, and wheat straw] and methane production in batch reactors at low temperature (20 °C).
Materials and methods

Experimental setup

The anaerobic digestion experiments were performed in six pairs of (1.0 L; working volume = 0.5 L) bottles maintained at 20 °C for three successive cycles to digest (xylan, cellulose, xylan:cellulose mixture, cow manure, and wheat straw). All reactors received 0.5 L of culture. The reactors were labeled as follows: R0: Control (contained 0.5 L of culture only; no substrate was added), R1: xylan (fed with 1.24 g of xylan), R2: cellulose (fed with 0.69 g cellulose), R3: xylan:cellulose mixture (fed with 0.62 g xylan and 0.34 g cellulose; 2:1 COD ratio), R4: cow manure (fed with 10.26 g cow manure), and R5: wheat straw (fed with 1.37 g of shredded wheat straw). The reactors were kept static and were mixed for 1 min once a week just before taking liquid samples. After preparing the reactors and feeding them with the substrates, they were purged with nitrogen (N₂) gas to establish anaerobic condition. The reactors were then closed with rubber stoppers and sealed with aluminum crimps.

Inoculum and substrates

The seed inoculum was made up as follows: 2.0 kg of the culture (30% total solids) obtained from dry anaerobic digestion reactors (40 L) was diluted with deionized water, homogenized, and sieved through 250 µm sieve to remove large fibers and any floating matter. The mixed liquor passed through the sieve was used as inoculum (TS 0.95%). For cycle 2 onwards the same culture at the end of the previous digestion cycle was re-used as inoculum for the next cycle. Table 1 gives the total solids (TS), volatile solids (VS), and chemical oxygen demand (COD) of the inoculum. The alkalinity of the dilute culture was 1850 mg L⁻¹ as CaCO₃ and was adjusted by adding 5.0 g of CaCO₃ into 8 L of dilute culture. The final measured alkalinity was 2600 mg L⁻¹ as CaCO₃. The dilute culture was incubated for 2 weeks to get rid of any residual substrate. The experiments started when the dilute culture biogas production reached zero.

Pure substrates [cellulose (insoluble fibers of 40 µm in particle size) and xylan (insoluble powder from birchwood) ] [Sigma Aldrich (Canada)] were used in this study. Bedding-uncontaminated fresh manure from dairy cows was collected, transferred into a plastic container, and stored at 4 °C until it was fed to the bioreactors. The wheat straw was shredded (<0.25 mm) and stored at 20 °C in plastic drum until its use. The substrate’s TS, VS, and COD are given in Table 1. The inoculum-to-substrate ratio ranged from 3.1 to 5.6 (Table 1) to ensure fast start-up of the anaerobic digestion depending on results published previously (Chynoweth et al. 1993; Hansen et al. 2004; Hashimoto 1989; Møller et al. 2004).

Biogas measurement

A calibrated digital gas pressure gauge (Dwyer Model: DPQA-08, Dwyer Instruments Inc., Michigan City, IN, USA) was used to measure the pressure of the biogas produced in the batch bioreactor. The volume of methane production was calculated based on standard temperature and pressure (STP) (273 K; 1 atm); i.e., normalized liter (L CH₄) using Eq. 1.

\[
V_{\text{CH}_4}^{\text{STP}} = \frac{V_m}{\alpha} \cdot \frac{T_s}{T_m} \cdot \frac{P_m}{P_s}
\]

where \(V_m\) is the measured volume of biogas, \(\alpha\) is the percentage of CH₄ in biogas, \(T\) and \(P\) which are the actual temperature and atmospheric pressure at the time of measurement, and \(T_s\) and \(P_s\) are the standard temperature and pressure. \(V_{\text{CH}_4}^{\text{STP}}\) is the volume of methane at the standard temperature and atmospheric pressure. The volume of CH₄ produced was corrected to that produced in the control reactors. The reported total specific methane yield (SMY) is the cumulative specific CH₄ yield which was calculated as the ratio of CH₄ produced (g L) over the mass of the total VS fed to the bioreactor at the beginning of the experiment. The COD recovery in biogas was calculated as the percentage of CH₄ produced relative to the theoretical biochemical methane potential (350 L CH₄ kg⁻¹ COD).

Table 1 Physicochemical characteristics of the inoculum and substrates

|                     | Inoculum (control) | Xylan | Cellulose | 2:1 xylan:cellulose mixture | Cow manure* | Wheat straw |
|---------------------|--------------------|-------|-----------|-----------------------------|-------------|-------------|
| Total solids (%)    | 0.97               | 93    | 96        | 95                          | 13          | 89          |
| Volatile solids (%) | 0.73               | 87    | 96        | 90                          | 11          | 85          |
| COD (g g⁻¹)         | 0.01               | 1.11  | 1.27      | 1.21                        | 1.15        | 1.23        |
| Organic loading rate (g COD L⁻¹) | NA     | 2.75  | 1.73      | 2.24                        | 3.04        | 3.36        |
| Inoculum-to-substrate ratio (g VS g⁻¹ VS) | NA   | 3.4   | 5.6       | 4.2                         | 3.1         | 3.1         |

*Alkalinity (5750 mg L⁻¹ as CaCO₃), pH (6.2), acetic acid (3370 mg L⁻¹), propionic acid (880 mg L⁻¹), butyric acid (1390 mg L⁻¹)
Analytical methods

Solids and chemical oxygen demand

Analytical methods used for the physicochemical [total chemical oxygen demand (TCOD), pH, alkalinity, and total and volatile solids (TS and VS, respectively)], biogas composition, and volatile fatty acids (VFAs), elemental analysis, and fiber analysis were according to the Standard Methods (APHA 1992).

Alkalinity was determined according to the method number 2320B Potentiometric titration to preselected pH 4.38. The TCOD was determined by the closed reflux colorimetric method (APHA 1992). The total solids (TS) and volatile solids (VS) content were determined using method number 2540B for TS and method number 2540E for VS. Briefly, the TS content was determined by drying a 10 mL sample for 24 h at 105 ± 3 °C. The VS content was determined by incinerating the dried solids for 3 h at 550 °C. The VS content was taken as the weight lost during the incineration; i.e., the VS content was calculated as the difference between the dried solids and fixed solids (ash remained after incineration).

Volatile fatty acids

The concentrations of individual volatile fatty acids (VFAs) including acetic, propionic, butyric, isobutyric, butyric, valeric and isovaleric acids have been measured according to the method number 5560D. The analysis used a Perkin Elmer gas chromatograph (GC) model 8310 (Perkin Elmer, Waltham, MA, USA) equipped with an autosampler to facilitate the analysis, fitted with FID, and equipped with a J&W Scientific DB-FFAP high resolution column (30 m × 0.53 mm × 1.00 μm; Chromatographic Specialties Inc., Ontario) (Saady and Massé 2013, 2015).

Helium, flowing at 9.5 mL min⁻¹, was employed as the carrier. The injector temperature was maintained at 200 °C, while the detector temperature was set at 250 °C. A 10 g of sample was diluted in 20 mL of deionized water, mixed thoroughly and the liquid part was separated and transferred into a centrifuge tube. After centrifugation for 30 min at 46,300g, 2 mL of the supernatant was transferred into 3 mL centrifuge tube, acidified with phosphoric acid, and centrifuged for 15 min at 16,300g. A 0.5 mL of the supernatant was filtered through a 0.45 μm nylon syringe filter into a 1.5 mL GC tube. The detection limit of acetic acid was 3.5 mg L⁻¹ while the detection limit of all other acids was 1.5 mg L⁻¹. The instrument performance was checked by throwing several standard samples systematically every five unknown samples injection until the end of the analysis of the batch of samples.

Biogas composition

The content of CH₄ and CO₂ in biogas were measured weekly using a Hach Carle 400 AGC gas chromatograph (Model 04131-C, Chandler Engineering, Houston, TX, USA) configured for the application 131-C. The application uses a column (1/8 inches) composed of 1.8 m (805 porapak N + 205 Porapak Q), 2.1 m (80% molecular sieve 13X + 20% molecular Sieve 5A), and 1.8 m (80% OV-101 on chromosorb WHP). The column and thermal conductivity detector were operated at 85 °C with a helium gas flow rate of 30 mL min⁻¹. The GC Calibration was performed weekly with a certified gas standard (27.3% CO₂, 1.01% N₂, 71.69% CH₄, 0.53% H₂S). The quantification regression equation has been generated using the same certified gas standards. The linear quantification regression equation covered the range from 0.001 to 100% (pure gas injection) with an R² value of 0.9991, 0.9988, and 0.9968 for CH₄, CO₂, and H₂S. Additionally and routinely, the GC performance checkup during any gas test was conducted at the beginning, every five injections, and at the end of the analysis of each batch of gas samples. This checkup was conducted by injecting triplicate samples of the certified gas standard at the beginning and injecting one standard gas sample after every five injections of unknown samples until the end of the analysis.

Elemental analysis

Elemental analysis was conducted on xylan, cellulose, and glucose; the glucose was used as a control to the elemental analysis. Elemental analyses for CHN and O were carried out with the Carlo-Erba elemental analyzer, model 1106, according to standard procedure and the results are accurate to ± 0.3%. The analysis for carbon (C), hydrogen (H), and nitrogen (N) has been performed based on a modification of the classical (Pella and Colombo 1972). Samples have been weighed accurately to 0.0001 mg, tightly enclosed in tin capsules, and analyzed. The analysis for oxygen (O) has been performed based on the Unterzaucher modified method (Pella and Colombo 1973). Samples have been weighed accurately to 0.0001 mg, tightly enclosed in silver capsules, and analyzed.

Results and discussion

Methane production

The SMY (mL CH₄ kg⁻¹ VS fed) obtained after 35 days of incubation (corrected to that produced in the control cultures) during the successive cycles are given in Table 2. The
elemental analysis revealed that the molecular formulae for cellulose and xylan are \( \text{C}_{5.928}\text{H}_{10.049}\text{O}_{4.965} \) and \( \text{C}_{5.148}\text{H}_{8.418}\text{O}_{4.418} \), respectively.

The theoretical methane yields for cellulose and xylan were calculated using the results of the elemental analysis and Buswell and Mueller’s Eqs. (2, 3) (Buswell and Mueller 1952).

Table 3 gives the percentage of the theoretical methane yield recovered in the successive cycles in the cultures fed pure and complex substrates. The cultures fed with the pure substrate (xylan, cellulose, xylan:cellulose mixture) recovered between 82 and 96% of the theoretical methane yield in cycle 3 compared to between 76 and 82% in cycle 1.

The highest percentage of theoretical methane yield recovery (96.0%) was in the culture fed a mixture of xylan:cellulose mixture whereas the lowest percentage (48.7%) was in the culture fed with cow manure:

\[
\text{C}_5\text{H}_9\text{O}_5\text{N}_n\text{S}_s + \left(\frac{c - \frac{h}{4} - \frac{a}{2} + \frac{3n}{8} + \frac{s}{2}}{2}\right)\text{H}_2\text{O} \rightarrow \left(\frac{c}{2} - \frac{h}{8} + \frac{a}{4} + \frac{3n}{8} + \frac{s}{4}\right)\text{CH}_4 + \left(\frac{c}{2} - \frac{h}{8} + \frac{a}{4} + \frac{3n}{8} + \frac{s}{4}\right)\text{CO}_2 + n\text{NH}_3 + s\text{H}_2\text{S}
\]

The rate kinetic and the extent of substrate biodegradability can be obtained from methane production curve pattern during the batch anaerobic digestion. The SMY profiles in the successive cycles of anaerobic digestion of pure and complex substrates in R1–R5 at low temperature (20 °C) are given in Fig. 1a–e. Generally, for all substrates and during all successive cycles the anaerobic digestion proceeded well without any observable inhibition. The \( \text{CH}_4 \) production profiles are composed of two phases: phase I which exhibited linear kinetics where the \( \text{CH}_4 \) production increases linearly with time and phase II in which the \( \text{CH}_4 \) production decreased towards a plateau. Nevertheless, some variations in the \( \text{CH}_4 \) profiles during the successive cycles can be noticed as well.

The rate of substrate degradation can be correlated to the \( \text{CH}_4 \) production rate since the latter can be calculated from the \( \text{CH}_4 \) production profiles. During the first cycle, the culture started degrading all substrates-fed immediately at a relatively high rate. Notice that the duration of this phase varied for the different substrates particularly during cycle 1; it was longer for the cultures fed pure substrate or their mixture (> 10 days) than that for the cultures fed complex substrates (cow manure and wheat straw) (6–7 days). In addition, the rates of \( \text{CH}_4 \) production in cultures fed with the pure substrates were faster than those of complex substrates (Table 4).

For comparison and taking R5 (wheat straw) as a reference, the rate of methane production for R1 (xylan) was 1.3–1.6 times that of R5, and for R2 (cellulose) it was 1.5–1.6 times that of R5. Similarly, R3 (fed mixture of the pure substrates) produced methane at a rate of 1.3–1.6 of that
produced by R5. Notice that the rate of methane production in R4 (cow manure) was almost 90% of that calculated for R5.

Generally, the specific methane yield (SMY) and its rate of production increased successively from cycle 1 to 2 and then to cycle 3 in all reactors except for R3 (fed with pure substrate mixture) which exhibited marginal decrease (0.2%) in cycle 3. The SMY increased by 7.6% from cycle 2 to 3 for R1, and increased by 4.5% and 9.7% for cycle 2 and 3, respectively, in R2 (fed with cellulose). Although the SMY increased by 17.9% from cycle 1 to 2 in R3 no further increase was measured in cycle 3. The cultures fed complex substrate exhibited a successive increase in the SMY. The increase was 8.5% and 17.2% in cycle 2 and cycle 3, respectively, in R4 (fed with cow manure). Similarly, the increase in specific methane yield was 9.75 and 16.7%, respectively, in R5. It is worthwhile to notice that the percentage of increase in the SMY in pure substrates-fed culture was less than the corresponding increase in SMY in the culture fed with a complex substrate. This is likely due to the nature of the substrate. Cow feces contains carbohydrates, proteins, and fats which require varying degradation times; therefore, they could have contributed to the increase in the SMY observed during cycle 2 and 3. Wheat straw, on the other hand, is slowly degraded substrate (Pohl et al. 2013); thus, the observed increase in SMY could have been from the residual of the substrate from the previous two cycles. However, the CH4 production profiles changed significantly in cycle 2 and 3. In R1, the rate of methane production increased during the exponential phase in the order cycle 3 > cycle 2 > cycle 1 and this phase lasted longer compared to that in cycle 1 (Fig. 1a). Notice that 90% of the maximum SMY obtained in R1 has been achieved in 12 days during cycle 3 compared to 20 days in cycle 1 and 14 days in cycle 2. This indicates an increase in the activity of the xylan degrading microorganisms from cycle 1 to 3.

Notice that 90% of the maximum SMY in R2 has been achieved in 17 days during cycle 3 compared to 22 days in cycle 1 and 19 days in cycle 2. The R2 experienced a lag phase in methane production of 2 days in cycle 1

| Reactor | Maximum CH4 production rate (L CH4 kg⁻¹ VS day⁻¹) |
|---------|--------------------------------------------------|
| Cycle 1 | Cycle 2 | Cycle 3 |
| R1      | 8.549 ± 0.254 | 8.563 ± 0.137 | 9.214 ± 0.014 |
| R2      | 9.011 ± 0.443 | 9.763 ± 0.323 | 10.88 ± 0.191 |
| R3      | 8.083 ± 0.283 | 9.531 ± 0.369 | 9.503 ± 0.169 |
| R4      | 5.131 ± 0.506 | 5.569 ± 0.251 | 6.529 ± 0.180 |
| R5      | 5.506 ± 0.386 | 6.037 ± 0.294 | 7.043 ± 0.474 |

Fig. 1 Profile of specific methane yield (SMY) during the first 35 days of the anaerobic digestion at 20 °C in successive cycles (values shown are average of duplicate bioreactors)
and 4 days in cycle 2 and 3. The lag phase is the time needed by the microorganisms to acclimatize to a new environment (Echiegu 2015) and substrates. Generally, the physiological processes by which microorganisms adapt to the new environment or substrates dictates the length of the lag phase (Rolfe et al. 2012). Non-fiber carbohydrates have been found easily accessible to microbial degradation (Janke et al. 2015) with short lag phase whereas substrates with increased fiber content such as the cellulose fed to R2 were reported to exhibits increased lag phase (Nielfa et al. 2015). Moreover, the duration of the lag phase is also related to the initial buffer capacity of the fermentation liquor (Mao et al. 2015). The alkalinity of the mixed liquor was not adjusted at the beginning of the cycle 2 and 3. Additionally, by the end of cycle 1 the content of the undegraded more recalcitrant organics which remained in the reactor after fermentation increased and might have lengthened the lag phases to 4 days in cycle 2 and 3 compared to 2 days in cycle 1; similar observation has been reported previously by Wilkie et al. (2000).

In the culture fed mixture of pure substrates (R3), the exponential phase of methane production lasted longer in cycle 3 (14 days) and cycle 2 (8 days) compared to cycle 1 (3 days). Notice that 90% of the maximum SMY in R3 has been achieved in 14 days during cycle 3 compared to 20 and 17 days in cycle 1 and 2, respectively. This represents 70–82% increase in the activity of the microorganisms degrading the xylan:cellulose mixture in cycle 2 and 3, respectively.

The culture in R4 (fed cow manure) exhibited a steady increase in the SMY of 8.5% and 17.2% in cycle 2 and cycle 3, respectively. The specific methane production rate was generally lower than that measured in cultures fed with other substrates. Interestingly, the exponential phase of the methane production profile lasted longer in cycle 3 (9 days) compared to cycle 1 and 2 (6 days). Around 90% of the maximum SMY in R4 has been achieved in 22 days during cycle 3 compared to 27 and 25 days in cycle 1 and 2, respectively. The increase in the duration of the exponential phase of methane production and the faster recovery of the 90% of the maximum SMY with the increase in cycle number indicates an enhancement in the activity of the microorganisms. This could be due to increase in the amount of the viable/active cells or the production of more active cold-adapted enzymes with high specific activities at low temperatures (high turn over rates) (Casanueva et al. 2010). For R5 (fed wheat straw) no change could be observed in the methane production rate and the duration of the exponential phase of methane production among the four successive cycles. About 90% of the maximum SMY from wheat straw has been achieved in 27, 22, and 22 days in cycles 1, 2, and 3, respectively. It seems that the culture adapted relatively at the same pace to both complex substrates.

In terms of COD recovery in methane, Table 5 gives the calculated percentages for the successive cycles in the cultures fed with the various substrate examined.

The difference between the percentage of COD recovery in methane and the percentage of the theoretical methane yield recovery is likely because the microorganisms used some of the substrate for growth and maintenance. The theoretical methane yield calculated from Eq. 3 is not corrected for biomass growth since Eq. 2 assumes that all the substrate will be converted into gaseous by-products. The microorganisms utilize about 5–10% of the substrate for growth (Rittmann and McCarty 2001). Generally, the COD recovery in all the cultures fed pure or complex substrates increased in cycle 2 whereas in cycle 3 the increase was marginal (1–2%). For example, in R2, the COD recovered as methane by day 15 was higher (72.4%) for cycle 2 and (73.3%) for cycle 3 compared to 55.9% in cycle 1. This might indicate that the culture needed a relatively short time (two successive cycles, i.e., 70 days) to adapt to the new substrates.

**Modeling kinetics of methane production**

The activity of the microorganisms catalyzing the CH₄ production during anaerobic digestion depends on the substrate, environmental conditions (temperature and pH), and physiological characteristics of the cell (Rolfe et al. 2012). The fiber content and its quality vary among the various pure and mixed substrates used in this study (Table 1). Kinetic model can elucidate the changes in the kinetics of the microbial cultures due to adaptation. The model would describe the CH₄ production curves obtained from the anaerobic digestion of the pure and complex fibrous substrates during successive cycles conducted under the same condition. Moreover, the model can provide information useful for predicting the digestion of fiber in animal manure and lignocellulosic biomass based on pure substrates studies. Fitting CH₄ production data to a mathematical model to describe the progress of methane production and estimate the kinetics parameters allows comparing the effects of various substrates and detect if any adaptation took place.

| Reactor | Percent COD recovery in CH₄ at day 15 |
|---------|--------------------------------------|
|         | Cycle 1          | Cycle 2          | Cycle 3          |
| R1      | 55.8 ± 0.7       | 67.6 ± 0.5       | 69.1 ± 0.8       |
| R2      | 55.9 ± 5.3       | 72.4 ± 0.5       | 73.3 ± 1.4       |
| R3      | 51.8 ± 5.9       | 66.9 ± 11        | 67.2 ± 1.3       |
| R4      | 29.3 ± 2.3       | 49.0 ± 1.0       | 52.9 ± 3.9       |
| R5      | 31.1 ± 3.0       | 45.5 ± 2.1       | 46.5 ± 2.0       |
The model was developed to describe the methane generation in the batch bioreactors of volume (V) using mass balance approach (Eq. 4) (Rittmann and McCarty 2001) and through monitoring COD concentration (C) through \( \text{CH}_4 \) production:

\[
\frac{dC}{dt} = Q_o C_o - Q_o C + VrC, \tag{4}
\]

where \( Q_o \) is the input flow to the bioreactor; \( C_o \) is the concentration of COD fed to the reactor, and \( C \) is the concentration of COD at any time, and \( r \) is the reaction rate of methane. Since for batch bioreactor \( Q_o=0 \); the Eq. 4 becomes:

\[
\frac{dC}{dt} = VrC. \tag{5}
\]

For the first-order rate kinetic model, the time rate of change in the concentration of COD in a bioreactor is a function of the concentration (C) and the rate constant (k):

\[
\frac{dC}{dt} = -kC. \tag{6}
\]

Solving this differential equation by integration yields:

\[
\ln \frac{C_o}{C} = kt. \tag{7}
\]

Although Eq. 7 uses the substrate’s concentration (COD) it could be modified to give the methane yield through developing a correlation between a substrate (COD) biodegradability and methane yield at any time (Y). To achieve this, it is assumed that all substrate consumed is to be converted into biogas; this introduces a marginal error because methane production is a growth-associated reaction; therefore, some substrate (COD) will be consumed for growth and production of new cell biomass. Equation 7 can now be written in terms of methane yield as follows:

\[
Y = Y_{\text{max}} (1 - \text{Exp}^{-kt}). \tag{8}
\]

where \( Y \) is methane volume \((\text{L})\) produced at time \( t \) (day), \( Y_{\text{max}} \) is maximum methane volume \((\text{L})\) produced, \( k \) is the reaction rate constant \((\text{day}^{-1})\), and \( t \) is incubation time (day). Now it is possible to estimate the reaction rate constant \((k)\) of the \( \text{CH}_4 \) production reaction by fitting the data of \( \text{CH}_4 \) production with time to the kinetic model (Eq. 7).

The nonlinear regression was performed using nonlinear regression function of POLYMATH software version 6.1 (Willimantic, CT, USA) with the objective of minimum sum of squared difference (SSR) between experimentally observed and model-estimated values of methane yield. The substrates examined in this study showed identical patterns of reaction progress with time. The values of the reaction rate constant \((k)\) are given in Table 6.

Since the rate of \( \text{CH}_4 \) production is proportional to the rate of the substrate degradation with time then the greater the value of the reaction rate constant \((k)\) the faster the degradation of the substrates. The ranking of the first-order reaction rate constant \((k)\) in the three successive cycles were as follows: Cycle 1: R3 > R1 > R2 > R4 > R5; Cycle 2: R1 > R3 > R2 > R5 > R4; Cycle 3: R1 > R3 > R2 > R4 > R5. Obviously, the kinetics differentiated between the two categories of substrates (pure and complex). The model provided a numerical indicator \((k)\) value (Table 6) that wheat straw \((k=0.0565 \text{ day}^{-1})\) was the least degradable substrate among those tested in cycle 1 and that cow manure \((k=0.0688 \text{ day}^{-1})\) was the second less degradable substrate while the pure substrates, xylan, xylan:cellulose mixture, and cellulose, showed the same degradability \((k=0.1116, 0.1118, \) and \(0.0932 \text{ day}^{-1}\), respectively). Although the reaction rate constant for R1 degradation increased by 104% (from cycle 1 to cycle 3 upon adaptation) the corresponding reaction rate constant for R2 degradation increased by only 9.5%. Notice that addition of cellulose to xylan at a ratio \((1:2 \text{ by} \) COD) decreased the percentage of increase in the reaction rate constant of this mixture degradation to only 22.1% compared to 104% in R1. This might indicate that the culture did not contain high active cellulolytic microorganisms or their adaptation might need longer time.

Upon adaptation the reaction rate constant calculated for R5 degradation increased by 86.9%, i.e., from 0.0565 day\(^{-1}\) (cycle 1) to 0.1056 day\(^{-1}\) (cycle 3). Similarly, adaptation increased the reaction rate constant in R4 by 68.5%, i.e., from 0.0688 day\(^{-1}\) (cycle 1) to 0.1159 day\(^{-1}\) (cycle 3).

Table 6 The reaction rate constant and the regression coefficient for the successive cycles

| Reactor | Cycle 1 | Cycle 2 | Cycle 3 |
|---------|---------|---------|---------|
| R1      | 0.1116 (0.9935) | 0.1489 (0.9768) | 0.2778 (0.9907) |
| R2      | 0.0851 (0.9440) | 0.0862 (0.9132) | 0.0932 (0.9935) |
| R3      | 0.1118 (0.9671) | 0.1159 (0.9898) | 0.1365 (0.9696) |
| R4      | 0.0688 (0.9900) | 0.0698 (0.9890) | 0.1159 (0.9790) |
| R5      | 0.0565 (0.9800) | 0.0817 (0.9910) | 0.1056 (0.9830) |

\(^a\)The reaction rate constant \((k)\) in \text{day}^{-1}

\(^b\)The regression coefficient \((R^2)\)
readily degradable, but when they are part of a complex structure of lignocellulosic material such as wheat straw they become less degradable. It must be noted that wheat straw contained (7.3%) by-mass lignin whereas xylan and cellulose were pure substrates (lignin free). This explains the difference in the amount of increment in the $k$ value observed in R1 and R3 compared to that observed in R4 and R5. Similarly, cow manure contains fibers composed of cellulose (23.61%), hemicellulose (18.71%) and lignin (11.3%). It must be taken into account that most of the readily digestible cellulose and hemicellulose in the cow diet have been digested in the animal digestive system and what is left of fiber in the cow manure is the part which is protected by the lignin. This explains why the increase in the $k$ value in R4 (0.0235) during the successive cycles of anaerobic digestion at 20 °C was lower than that estimated for R1 (0.0831). However, the increase in $k$ value in R4 (0.0235) over the successive cycles is greater than that in R5 (0.0207) (Fig. 2) might be due to the fats and proteins contained in the cow manure. Wheat straw does not contain any fats, and its content of protein is minimum.

The applicability of a first-order model depends to a large extent on the characteristics and quality of inoculums and that some of the variability in the kinetic constants in models is dictated by the inoculum (Neves et al. 2008). The kinetic of CH$_4$ production from fiber digestion has previously been modeled as a first-order reaction (Kang et al. 2014; Adl et al. 2015; Castano 2012). However, the comparison of the results from this study with values of the reaction rate constant reported in literature should be conducted cautiously because some studies report the first-order kinetic of the hydrolysis step rather than the entire methane production process. The values of $k$ cited from the literature in Table 7 are for the whole anaerobic digestion process unless indicated otherwise. The rate estimated for cellulose (0.085–0.093 day$^{-1}$) in this study is a mid-way between the lower and the upper limits of those reported by other studies (0.05–0.18 day$^{-1}$) (Table 7). This might be due to the effect of the lower temperature (20 °C) compared to that of mesophilic (28–35 °C) for the studies cited. Notice that the rate estimated for cellulose degradation in this study forms about 41–48% of that reported by Desvaux et al. (2001) for cellulose digestion by a pure culture of Clostridium cellulolyticum which always follows first-order kinetics with $k$ value (0.192 day$^{-1}$).

The $k$ value obtained in this study for xylan (0.112–0.278 day$^{-1}$) is within the values reported previously for carbohydrate (0.120–0.320 day$^{-1}$) (Neves et al. 2008). The $k$ value estimated for cow manure (0.069–0.116 day$^{-1}$) compares well with 0.063 day$^{-1}$ reported by Castano (2012) for anaerobic digestion of dairy manure at 20 °C. Additionally, it compares well with the 0.07–0.08 day$^{-1}$ reported by Kang et al. (2014) for digesting dairy manure at 35 °C. Nevertheless, it is generally within the value (0.08 day$^{-1}$) reported by Hobson (1983) for slowly degradable cow slurry. Although the impact of temperature is obvious the culture adapted well to the new substrate at 20 °C.

Conclusions

Anaerobic culture adapted to digest cellulose, xylan, cellulose and xylan mixture, cow manure, and wheat straw at low temperature (20 °C) in relatively short period (3 successive cycles of 35 days each). First-order model kinetic revealed that the increase in the reaction rate over the successive cycles was 0.0831 (xylan) > 0.0124 (xylan:cellulose) > 0.0113 (wheat straw) > 0.0077 (cow manure) > 0.0041 (cellulose). The rate of the substrate degradation at 20 °C were: 0.085–0.093 day$^{-1}$ (cellulose), 0.112–0.278 day$^{-1}$ (xylan), 0.112–0.137 day$^{-1}$ (xylan:cellulose mixture), 0.069–0.116 day$^{-1}$ (cow manure), and 0.057–0.106 day$^{-1}$ (wheat straw). The physicochemical characteristics of the substrates affect the adaptation of the microbial consortium and lead to differences in the reaction rate constants of the CH$_4$ production. Adding cellulose to hemicellulose decreased the percentage of increase in the reaction rate constant during adaptation compared to hemicellulose only. Unless high active cellulolytic microorganisms present in the culture its adaptation might need longer time.
Acknowledgements This research was partially supported by the Seed, Bridge and Multidisciplinary Fund (2018) from the Memorial University of Newfoundland.

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Table 7 The first-order reaction rate constants of methane production for lignocellulosic substrate

| Substrate                          | Temperature (°C) | The reaction rate constant k (day⁻¹) | References                           |
|------------------------------------|-----------------|-------------------------------------|--------------------------------------|
| Cellulose                          | 20              | 0.085–0.093                         | This study                           |
| Xylan                              | 20              | 0.112–0.278                         | This study                           |
| Cellulose-xylan mixture            | 20              | 0.112–0.137                         | This study                           |
| Cow manure                         | 20              | 0.069–0.116                         | This study                           |
| Wheat straw                        | 20              | 0.057–0.106                         | This study                           |
| Cellulose                          | NR              | 0.180                               | Gunaseelan (2009)                    |
| Cellulose (in cattle manure)       | 30              | 0.05                                | Singh et al. (1982)                  |
| Cellulose                          | 28              | 0.17                                | Heukelekian (1927)                   |
| Cellulose                          | 35              | 0.12                                | Greco et al. (1983)                  |
| Cellulose                          | NR              | 0.192*                              | Desvaux et al. (2001)                |
| Avicel cellulose                   | NR              | 0.14                                | Chynoweth et al. (2001)              |
| Excess cellulose food waste        | 35              | 0.18                                | Qamaruz-Zaman and Milke (2012)       |
| Restaurant waste with excess       |                 | 0.120                               | Neves et al. (2008)                  |
| Carbohydrates                      |                 | 0.320                               | Neves et al. (2008)                  |
| Pig manure and straw               | 35              | 0.02                                | Llabrés-Luengo and Mata-Alvarez (1987) |
| Pig manure                         | 38              |                                     | Adl et al. (2015)                    |
| Hydrolysis                         |                 | 0.105                               |                                     |
| Acidogenesis                       |                 | 0.106                               |                                     |
| Methanogenesis                     |                 | 0.231                               |                                     |
| Cow manure slurry                  | 35              | 0.09                                | Karim et al. (2007)                  |
| Dairy manure                       | 25              | 0.105–0.114                         | Castano (2012)                       |
|                                   | 20              | 0.049–0.063                         |                                     |
|                                   | 15              | 0.044–0.047                         |                                     |
| Dairy manure                       | 35              | 0.07–0.08                           | Kang et al. (2014)                   |
| Cow slurry                         |                 |                                     | Hobson (1983)                        |
| Slowly degradable                 |                 | 0.08                                |                                     |
| Rapidly degradable                |                 | 0.32                                |                                     |
| Dairy manure                       | 38              |                                     | Adl et al. (2015)                    |
| Hydrolysis                         |                 | 0.093                               |                                     |
| Acidogenesis                       |                 | 0.143                               |                                     |
| Methanogenesis                     |                 | 0.146                               |                                     |
| Bedding straw                      | 35              | 0.008–0.010                         | Torres-Castillo et al. (1995)        |
| Rice straw                         | 38              |                                     |                                     |
| Hydrolysis                         |                 | 0.093                               |                                     |
| Acidogenesis                       |                 | 0.920                               |                                     |
| Methanogenesis                     |                 | 1.350                               |                                     |
| Sugarcane straw                    | 38              | 0.075–0.102                         | Janke et al. (2015)                  |

*By pure culture of Clostridium cellulolyticum

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