

Enhanced bioconversion of hydrogen and carbon dioxide to methane using a micro-nano sparger system: mass balance and energy consumption

著者別名	清水 和哉, 雷 中方, 小林 幹佳, 張 振亞
journal or publication title	RSC Advances
volume	8
number	47
page range	26488-26496
year	2018-07
権利	(C) The Royal Society of Chemistry 2018 Open Access Article. This article is licensed under a Creative Commons Attribution-NonCommercial 3.0 Unported Licence.
URL	http://hdl.handle.net/2241/00154023

doi: 10.1039/C8RA02924E

Cite this: *RSC Adv.*, 2018, 8, 26488

Enhanced bioconversion of hydrogen and carbon dioxide to methane using a micro-nano sparger system: mass balance and energy consumption†

Ye Liu,^a Ying Wang,^a Xinlei Wen,^a Kazuya Shimizu,^a Zhongfang Lei,^a Motoyoshi Kobayashi,^a Zhenya Zhang,^a ^{*,a} Ikuhiro Sumi,^b Yasuko Yao^b and Yasuhiro Mogi^b

Simultaneous CO₂ removal with renewable biofuel production can be achieved by methanogens through conversion of CO₂ and H₂ into CH₄. However, the low gas–liquid mass transfer ($k_{L}a$) of H₂ limits the commercial application of this bioconversion. This study tested and compared the gas–liquid mass transfer of H₂ by using two stirred tank reactors (STRs) equipped with a micro-nano sparger (MNS) and common micro sparger (CMS), respectively. MNS was found to display superiority to CMS in methane production with the maximum methane evolution rate (MER) of 171.40 mmol/L_R/d and 136.10 mmol/L_R/d, along with a specific biomass growth rate of 0.15 d⁻¹ and 0.09 d⁻¹, respectively. Energy analysis indicated that the energy–productivity ratio for MNS was higher than that for CMS. This work suggests that MNS can be used as an applicable resolution to the limited $k_{L}a$ of H₂ and thus enhance the bioconversion of H₂ and CO₂ to CH₄.

Received 5th April 2018

Accepted 30th May 2018

DOI: 10.1039/c8ra02924e

rsc.li/rsc-advances

Introduction

Carbon dioxide (CO₂) has been regarded as the largest contributor to global warming, accounting for about 60% of the greenhouse gas effects.¹ According to a recent survey,² following the energy industry, the industry and product sector (7%) is the second emission source in Japan, especially the iron and steel industry. CO₂ is usually discharged as a waste product due to its inert, non-reactive, and low Gibbs free energy properties. Various CO₂ removal technologies including absorption,³ adsorption,⁴ cryogenic distillation,⁵ and membranes⁶ have been proposed and investigated to mitigate its emission. Taking into consideration that CO₂ is another important carbon source, the conversion of CO₂ to chemicals and energy products that are currently produced from fossil fuels is promising due to the high potential market and promising benefits.^{7,8} In comparison to chemical-based CO₂ capture, biological conversion of CO₂ together with hydrogen (H₂) to methane (CH₄) attracts particular interest, because the profits generated from CO₂ utilization can offset a portion of the capture cost under mild operational conditions.^{9,10} Bioconversion of CO₂ to CH₄ by hydrogenotrophic methanogens is of considerable interest because

this process realizes energy storage and conversion, as well as using the biological-based CO₂ capture and sequestration technique.^{11,12} The calorific value can be improved by converting nonflammable CO₂ to CH₄ (with a calorific value of 55 kJ kg⁻¹). Moreover, methanogens can produce not only methane but also valuable materials such as enzymes, amino acids, vitamins and so on.¹³

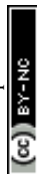
However, there are two key points in the bioconversion procedure. The first one is the source of the energy donor H₂. In the previous study, H₂ can be rapidly produced *via* water electrolysis,¹⁴ or obtained from biological process (bio-hydrogen).^{15,16} While considering H₂ is also a co-product in the steel sector, it is a promising alternative to utilize the waste H₂ to realize the bioconversion. Meanwhile, CH₄ as the product of the bioconversion is also regarded as an energy carrier for electricity storage, which is more easily transported or stored than H₂.¹⁷

The second point, also the big obstacle to the successful development of the technology for scaling up is the poor gas–liquid mass transfer rate ($k_{L}a$) of low soluble H₂ gas.^{18–20} When gas is sparged into the liquid, the $k_{L}a$ principally depends on the size and number of bubbles present,²¹ which are affected by many factors such as agitation speed,²² gas or liquid flow rate,²³ reactor geometry^{11,23} and the nature of the liquid.²⁴ Various methods were employed to improve the $k_{L}a$ value of H₂. Kougias *et al.*²⁵ demonstrated successful *ex situ* biogas upgrading in different systems by the action of hydrogenotrophic methanogens. Compared with the continuous stirred tank reactor (CSTR) and bubble column reactor, the two connected up-flow

^aGraduate School of Life and Environmental Sciences, University of Tsukuba, 1-1-1 Temodai, Tsukuba, Ibaraki 305-8572, Japan. E-mail: zhang.zhenya.fu@u.tsukuba.ac.jp

^bJFE Steel Cooperation, 2 Chome-2-3 Uchisaiwaicho, Chiyoda, Tokyo, 100-0011, Japan

† Electronic supplementary information (ESI) available. See DOI: 10.1039/c8ra02924e



reactor gained the highest efficiency of biogas upgrading with the CH₄ content of 98%. Bassani *et al.*¹⁹ constructed a granular UASB coupled with a separate chamber for H₂ injection. The $k_L a$ value of H₂ was improved by different packing materials (rashing rings and alumina ceramic sponge), which lead to a maximum increase of 24% in CH₄ content. While the $k_L a$ value of hydrogen was not mentioned. Diaz *et al.*²⁰ evaluated the potential of a pilot hollow-fiber membrane bioreactor for the conversion of H₂ and CO₂ to CH₄. The system transformed 95% of H₂ and CO₂ fed at a maximum loading rate of 40.2 m_{H₂}³/m_R³d and produced 0.22 m³ of CH₄ per m³ of H₂ fed at thermophilic conditions. $k_L a$ -H₂ of 430 h⁻¹ were reached in the bioreactor by sparging gas through the membrane module. Luo and Angelidaki²³ used the CSTR equipped with a column diffuser (pore diameters 0.5–1.0 mm) or ceramic diffuser (14–40 μm) to realize *in situ* biogas upgrading by co-digestion of manure and whey with the addition of H₂. The decrease in pore size of diffusers lead to an increase in $k_L a$ value from 6.60 h⁻¹ to 16.05 h⁻¹. While much higher $k_L a$ values of H₂ (105–776 h⁻¹) were reached by up-flow reactors in the research of Bassani *et al.*²⁶ Converse to the result of Luo and Angelidaki,²³ the smaller pore size lead to the lower $k_L a$ value, which was illustrated that the more efficient gas bubble break was caused by the larger pore size devices. Although various enhancement measures have improved the $k_L a$ of H₂ and CH₄ production significantly, the energy consumption and practical viability of them should be carefully evaluated for their applications in large scale fermentation systems. What's more, not only the CH₄ yield and $k_L a$ -H₂, but also the intermediates variation and biomass growth should be taken into deep consideration, to understand the evolution of carbon and hydrogen.

Recently, a special attention has been paid to the application of micro-nano bubbles (MNBs) technology in many fields, including medicine science,²⁷ food science,²⁸ aquaculture,²⁹ and water remediation.³⁰ Several special characteristics of MNBs, such as high specific area (surface area per volume) and high stagnation in liquid phase, increase the gas dissolution. Moreover, it has been reported that the collapse of micro-bubbles, due to the high density of ions in gas-liquid interface just before the collapse, will lead to free radical generation, which might be favorable for microbial metabolism and further stimulate the bioactivity.³¹ Up to now, however, little information can be found on the combination of MNBs with methanogenesis. Except the methane production performance and mass transfer of H₂, it is worthy to explore the interaction of bubbles and microbes.

In this study, MNBs were applied for the bioconversion of H₂ and CO₂ to CH₄, aiming at supplying a sufficient interaction of gaseous substrates and methanogens. The effect of MNBs on methanogens was achieved by operating two bioreactors equipped with micro-nano sparger (MNS) and common micro sparger (CMS), respectively. The gas-liquid mass transfer of hydrogen was also determined in this study. Bubble size distribution of the two spargers were analyzed to calculate the specific surface area. By analyzing the variations of reactants and products in the two bioreactors and conducting the mass balance analysis, deeply understand the evolution procedure of

carbon and hydrogen. Also, the energy consumption analysis was conducted to figure out the commercialization potential of MNS.

Materials and methods

Experimental apparatus and operation conditions

This study was carried out in two identical stirred tank reactors (STRs) equipped with a micro-nano sparger (MNS) (Foamest Column 16-60, Nac sales corporation, Japan) and a common micro sparger (CMS) (HA003, Haohai, China), respectively. The two reactors have a same total volume of 1.1 L (head-space : liquid = 6 : 5, v/v). The H₂ and CO₂ gas mixture (80/20, v/v) was transferred from the head space of the reactor to the liquid phase by a diaphragm pump (GS-6EA, E.M.P.-Japan Ltd, Japan), with a continuous recirculation rate of 40 mL min⁻¹. A gas holder connected to the head space of the reactor was used to allow fermentation to proceed without creating vacuum and maintain a positive and constant pressure inside the reactor during the fermentation. A magnetic stirrer was employed for both reactors to maintain mixing (500 rpm) and temperature maintenance (37 ± 2 °C). All the tube connections, stoppers, and seals were made of butyl rubber and glass. The diagram of the experimental setup in this study is illustrated in Fig. 1.

Inoculum and medium

The acclimated inocula, collected from the pond sediment (Matsumi Ike, Tsukuba campus) which has been well adapted to the H₂/CO₂ gas mixture (80/20, v/v) for 4 months' methane production, were introduced at a ratio of 1 : 4 (v/v) into the medium. The compositions of medium were the same as a previous study.³²

Analytical methods

The contents of H₂, CO₂, and CH₄ in the gas phase were analyzed by gas chromatograph (Shimadzu GC-8A, Japan) equipped with a thermal conductivity detector connected to

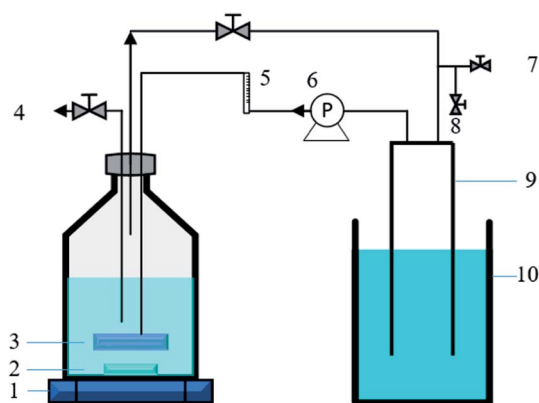


Fig. 1 Schematic diagram of the batch fermentation system. (1) Magnetic stirrer, (2) rotor, (3) micro-nano sparger (or common micro sparger), (4) liquid sample port, (5) flow meter, (6) gas recirculation pump, (7) gas input port, (8) gas sample port, (9) gas holder, (10) saturated sodium bicarbonate solution.



a chromatopac data analyzer (Shimadzu C-R4A, Japan). A stainless steel column packed with Porapak-Q was used for the analysis, with the temperatures of both detector and injector at 60 °C and of the column at 80 °C, respectively. N₂ was used as the carrier gas at an inlet pressure of 199 kPa and an outlet pressure of 150 kPa, respectively. The gas samples were taken at an interval of 24 h.

Total solid (TS), volatile solid (VS), and biomass concentration were analyzed according to the standard methods.³³ Volatile fatty acids (VFAs) were analyzed by gas chromatography (GC-FID, Shimadzu C-R8A, Japan). Soluble carbon was analyzed by a TOC analyzer (Shimadzu TOC-V_{CSN}, Japan). The liquid samples were collected every three days, which were used for VFAs analysis after filtration through 0.22 μm filters. 3% phosphoric acid solution was added to the filtrate to acidify samples at a volume ratio of 1 : 9 for the analysis of volatile fatty acids (VFAs). Gas chromatograph (GC-8A, Shimadzu) equipped with Unisole F-200 30/60 column and flame ionization detector (FID) was used for quantification of VFAs, including acetic acid (HAc), propionic acid (HPr), iso-butyric acid (iso-HBu), *n*-butyric acid (*n*-HBu), iso-valeric acid (iso-HVa) and *n*-valeric acid (*n*-HVa).³⁴

The bubble size distribution from the two sparger systems was analyzed by Nano Sight (nano scale bubbles) and a high-speed camera (micro scale bubbles) with distilled water as the media. The mean bubble size was utilized for the calculation of specific surface area.

The mass transfer coefficient ($k_L a$) of H₂ was determined before inoculation. H₂ was supplied to the reactors continuously. H₂ gas samples were collected from a three-way gas sampling port at an interval of 5 min. The first gas sample was collected at 2 min after the introduction of H₂ gas into the reactor. Once being collected, the gas sample was injected into a 20 mL sealed vial which contained some water at a same gas phase to liquid phase ratio as the reactor. Then the gas and the liquid was well mixed using a vortex mixer for 1 min and allowed 1 h to equilibrate the gas and the liquid phases. A gas sample from the head space was then taken and analyzed for gas composition in the gaseous phase using GC-TCD (Shimadzu GC-8A, Japan). The gas content in the head space was then converted to the aqueous phase concentration according to Henry's law (eqn (1)).

$$K_H = \frac{P}{X} \quad (1)$$

where, K_H is the Henry's law constant (atm), P is the partial pressure of gas above the aqueous phase (atm) and X is the mole fraction of gas in the solution (unitless). The Henry's law constant used for H₂ in this analysis was 7.52×10^4 atm (at 35 °C and 1 atm).

SEM images were taken by Field Emission Scanning Electron Microscope (FE-SEM, S-4800; Hitachi Hitec Corp, Japan) using the samples during the stable operation of the two reactors (on day 38). The samples were diluted by ten times and dried for one night in the oven (40 ± 1 °C) before being used for analysis.

Calculation

Determination of $k_L a$ of hydrogen. Assuming that the concentration in the liquid phase at the gas-liquid interface is in equilibrium with the gas concentration in the gaseous phase, the volumetric mass transfer coefficient ($k_L a$) in the absence of microorganisms was determined using the following equation (eqn (2)).

$$\frac{dc}{dt} = k_L a (C_i - C) \quad (2)$$

where, C is the gas concentration in the liquid phase (mg L^{-1}) at any given time t (min), and C_i is the saturated gas concentration (mg L^{-1}). eqn (2) can be further simplified to eqn (3),

$$\ln\left(\frac{C_i - C_0}{C_i - C}\right) = (k_L a)t \quad (3)$$

where, C_0 is the initial gas concentration in the liquid phase (mg L^{-1}).

Hydrogen conversion efficiency. Hydrogen was provided as the sole electron donor for the batch experiments. As the total mass of H₂ supplied into the systems ($m_{G_{H_2, in}}$) is the sum of the mass utilized by the microbes and the mass left (effluent) in the system, the H₂ conversion efficiency (%) was calculated according to eqn (4),

$$\eta_{H_2} = 100 \times \left(\frac{m_{G_{H_2, in}} - m_{G_{H_2, eff}}}{m_{G_{H_2, in}}} \right) \quad (4)$$

where $m_{G_{H_2, in}}$ is the mass flow rate of H₂ fed into the reactor per day and $m_{G_{H_2, eff}}$ is the mass flow rate of H₂ in the effluent gas. The utilized H₂ ($m_{H_2, utl}$) was the difference between $m_{G_{H_2, in}}$ and $m_{G_{H_2, eff}}$, as shown in eqn (5):

$$m_{H_2, utl} = m_{G_{H_2, in}} - m_{G_{H_2, eff}} \quad (5)$$

$m_{H_2, utl}$ can be classified into two parts, H₂ employed for microorganisms growth (anabolism) and consumed to produce energy (catabolism). Since the H₂ as an energy source can be transferred to VFAs as the intermediates and CH₄ as the final product, the utilized H₂ can be quantified according to eqn (6).

$$m_{H_2, utl} = m_{CH_4, H_2} + m_{VFAs, H_2} + m_{growth, H_2} \quad (6)$$

where m_{CH_4, H_2} is the mass of CH₄ as equivalent H₂, m_{VFAs, H_2} is that of H₂ transferred into VFAs, and m_{growth, H_2} is that of H₂ employed for microbial growth.

Carbon balance analysis. The carbon was supplied in three phases in this study: CO₂ (gas phase), Na₂CO₃ (liquid phase), and inoculum or biomass (solid phase), which can be expressed as follows:

$$C_{Total, in} = C_{in, G, CO_2} + C_{L, medium} + C_{S, biomass} \quad (7)$$

After the reaction, CO₂ can be transferred into CH₄, VFAs, and utilized for microbial growth, and other metabolites. Finally, the unreacted CO₂ was left in the system.

$$C_{Total, in} = C_{G, CH_4} + C_{G, CO_2, eff} + C_{L, VFAs} + C_{L, other} + C_{S, biomass} \quad (8)$$



Estimation of Monod kinetic parameters for the batch experiments. In order to estimate the maximum specific growth rate (μ_{\max}), the nutrients were supplied sufficiently to create the condition of nutrients concentration (C) $\gg K_s$ (i.e. $C/(K_s + C) \approx 1$). During the exponential growth phase, the Monod equation can be simplified to eqn (9).

$$\left(\frac{dX}{dt}\right)_{\text{growth}} = \mu_{\max} X \quad (9)$$

where μ_{\max} is the maximum specific growth rate (d^{-1}). Then, μ_{\max} can be obtained by eqn (10):

$$\mu_{\max} = \frac{1}{t} \ln \frac{X_t}{X_0} \quad (10)$$

where X_0 is the initial biomass concentration (g-biomass per L) and X_t is the biomass concentration at time t during exponential growth with no limitation of nutrients (g-biomass per L).

Energy consumption analysis. The electricity consumed by the systems was estimated by monitoring the energy consumption of the related devices. The energy input was the sum of electricity consumed by pumps and stirrers, as shown in eqn (11):

$$E_{\text{consumed}} = E_{\text{pump}} + E_{\text{stirrer}} \quad (11)$$

where E_{consumed} , E_{pump} and E_{stirrer} are the total energy consumed by the system, the energy consumed by pump and stirrer, respectively.

The energy-product ratio (R) was calculated by eqn (12).

$$R = \frac{Y_{\text{CH}_4, \text{total}}}{E_{\text{consumed}}} \quad (12)$$

where R is the energy-product ratio (L kW^{-1}), $Y_{\text{CH}_4, \text{total}}$ is the accumulated methane production during the whole experimental period (L). The energy consumption is mainly attributable to the heat loss during the operation with a small amount utilized by the growth of microbes.

Statistic analysis. All the data were expressed as mean value \pm standard deviation in this study. The significance of difference in the quantitative variables (e.g. CH_4 content in the output gas) between the two reactor systems was analyzed by one-way analysis of variance (ANOVA) using Origin 9.0 (Originlab, USA), and significance was assumed at $p < 0.05$. Moreover, regarding the microbial community, statistical analysis was carried out as previously described by Tsapekos *et al.*³⁵ to identify the significant abundance difference in microorganisms among the samples.

Results and discussion

Bioconversion performance

Methane production. To compare the effect of MNS and CMS on CH_4 production, the MNS reactor (MNSR) and CMS reactor (CMSR) systems were established to produce CH_4 from H_2 and CO_2 . A start-up period was noticed in both reactors with a gradually increased methane production rate (Fig. 2), due to the adaption of methanogens to the new conditions. While the

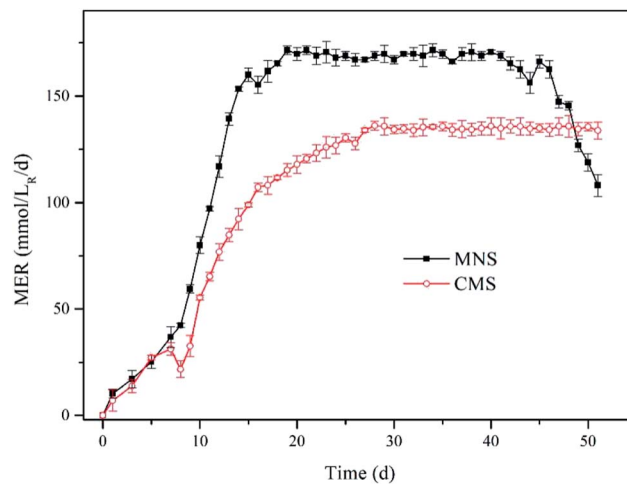


Fig. 2 Methane evolution rate (MER) for the two reactors. Solid squares denote MNSR and open circles denote CMSR.

methane production trended to be stable from day 14 to day 45 with the maximum methane evaluation rate (MER) of 171.40 $\text{mmol/L}_R/\text{d}$ in the MNSR. Similarly, the stable phase in the CMSR was detected from day 27 to day 50, achieving the maximum MER of 136.10 $\text{mmol/L}_R/\text{d}$. The enhancement of methane yield in the MNSR is significant with the p value lower than 0.05. The delayed achievement on the maximum MER in the CMSR was most probably attributed to its lower biomass growth rate. At the very beginning, both the aqueous substrates and the gaseous substrates were sufficient for the microbes, so that the microbial growth exhibited a rapid increase which agrees with the increase in methane production, the major product in this kind of systems. However, when the biomass increased to some extent, the dissolved hydrogen concentration (DHC) became the limiting factor for the microbes, leading to the maximum methane production maintained at a stable period. From day 46 to day 51, the MER of MNSR began to decrease gradually, as a result of the exhaustion of some nutrients in the medium. The results from this study indicate that MNSR can produce more CH_4 than CMSR.

The finding of this work is to some extent similar to that of Weimer & Zeikus³⁶ who used *Methanosarcina barkeri* MS as the inoculum, achieving the MER of 5.539 $\text{mmol L}^{-1} \text{h}$ in a 2 L reactor under fed-batch mode conditions. Their result is comparable to the MER of CMSR, while lower than that of MNSR. The current work is also comparable to Roennow & Gunnarsson³⁷ who carried out the fed-batch experiments in a 1 L reactor. For recent studies, the methane production has been improved greatly by various methods. Lee *et al.*³⁸ converted H_2 and CO_2 to methane by a fixed bed reactor with the mixed culture, and got the conclusion that the maximum methane production of 143.97 $\text{mol L}^{-1} \text{d}^{-1}$, which is much higher than this study. The low methane yield in this study was closely related to the low feeding rate of gaseous substrates.

H_2 utilization efficiency. In order to confirm that the enhancement on CH_4 production by MNSR was brought about by improvement of $k_L a$ using MNBs, H_2 balance analysis was used to investigate whether the bioconversion efficiency of H_2



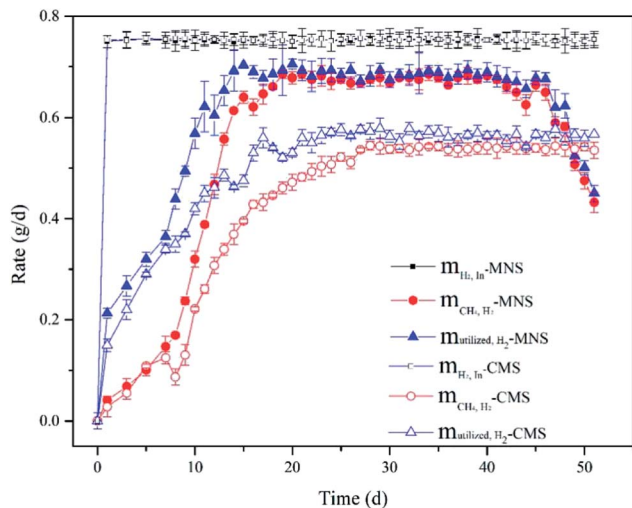


Fig. 3 H₂ balance analysis for the two reactor systems.

was stimulated by the MNBs or not. The utilized H₂ concentration was calculated by eqn (5). As shown in Fig. 3, H₂ was not exhausted in both MNSR and CMSR, due to the decreased dissolved H₂ concentration along with operation time, resulting in weakened driving force for gas-liquid mass transfer. As the same trend with CH₄ production, H₂ consumption increased gradually, and then became stable for some period. The maximum H₂ utilization in the MNSR was detected on day 20 with the maximum η_{H_2} of 95%. In comparison to MNSR, $m_{\text{H}_2,\text{util}}$ in the CSR was lower, correspondingly yielding to a lower maximum η_{H_2} of 80% (day 25). Albeit the methane yield in this study was low, the H₂ utilization efficiency of MNS is high and comparable with other studies,^{20,38} indicating the smaller bubbles can be utilized effectively by the methanogens. This observation implies that MNS can transfer more H₂ from gas phase into liquid phase during the same operation duration compared with CMS. An ideal condition for the bioconversion is that H₂ could be transferred at a high rate without any accumulation. The maximum dissolved hydrogen concentration in the water is about 1.6 mg L⁻¹ at normal pressure.³⁹ While the utilized hydrogen concentration in the two reactors is much higher than 1.6 mg L⁻¹, illustrating that H₂ was consumed by the microbes. The H₂ converted to CH₄ ($m_{\text{CH}_4,\text{H}_2}$) was lower than the utilized H₂ concentration ($m_{\text{H}_2,\text{util}}$), indicating that there could be a prior in biomass growth than CH₄ production. From day 20 on, the $m_{\text{CH}_4,\text{H}_2}$ in the MNSR contributed over 95% to the total utilization of H₂, and maintained at this high proportion for about 20 days. The CH₄ production in both reactors did not show a continuous increase, demonstrating the lack of dissolved hydrogen for methanogens. Specifically, the contribution of $m_{\text{CH}_4,\text{H}_2}$ in the CMSR to $m_{\text{H}_2,\text{util}}$ was slightly lower (about 90%), indicating H₂ was also engaged in VFAs production and biomass growth.

Biomass enrichment

To determine the effect of MNBs on biomass growth, biomass concentration was quantified in time course (Fig. 4). The superiority of the MNSR was not obvious till day 5. After day 5

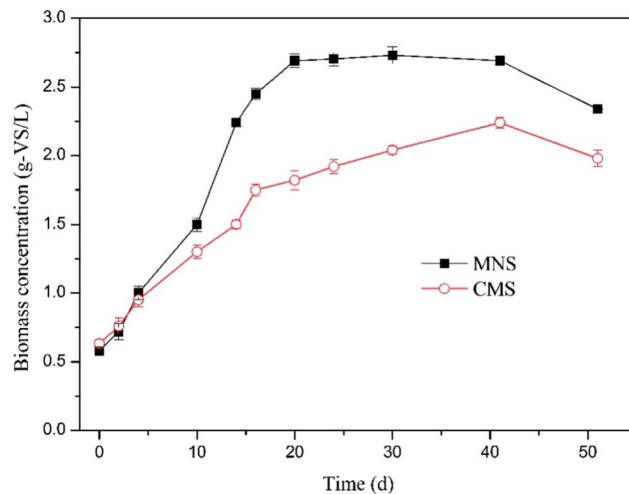


Fig. 4 Variation in biomass concentration in the two reactors. Solid squares denote MNSR and open circles denote CMSR.

the MNSR experienced a rapid increase in biomass and earned a higher maximum specific growth rate of 0.15 d⁻¹ in comparison to 0.09 d⁻¹ in the CMSR. The methanogens in the two reactors were shown in Fig. S1.† Results suggest that methanogens use more dissolved H₂ in MNSR due to its higher k_1a . However, the biomass concentration showed a stable trend following the rapid growth period, demonstrating that the dissolved H₂ concentration was still the limiting factor which dominates H₂ utilization for catabolism reactions (CH₄ production). The inocula used in this study was the acclimated anaerobic sludge, a mixed culture. Compared to the pure culture, generally the mixed culture is more advantageous regarding availability and cost, if not thinking about the relatively lower conversion rate due to the competitive communities. Moreover, the MNSR can be applied in large-scale tests according to the results from the acclimated anaerobic sludge.

Variation in volatile fatty acids (VFAs)

Anaerobic conversion of CO₂ can support a variety of microorganisms from different trophic groups within a microbial community. Therefore, the pathways involved in CH₄ production from CO₂ become more complex when taking the mixed anaerobic consortium into consideration. For a mixed methanogenic culture, *i.e.* the acclimated anaerobic sludge used in this study, it is essential to consider all possible reactions that are involved in the conversion of CO₂ to CH₄. The possible pathways indicate that hydrogenotrophic methanogens can directly convert H₂ and CO₂ to CH₄ through the pathway (a) (Fig. S2†). On the other hand, homoacetogenic bacteria can participate in the conversion of the H₂ and CO₂ to acetate, a thermodynamically favorable reaction (pathway (b)). Then the acetoclastic methanogenesis will occur according to the pathway (c). Conversely, syntrophic acetate-oxidizing (SAO) bacteria can convert acetate to H₂ and CO₂ (pathway (d)) when acetoclastic methanogenesis is deficient.⁴⁰ The SAO reaction becomes thermodynamically favorable at low H₂ partial pressure (<10⁻⁴ atm at 35 °C).^{41,42}



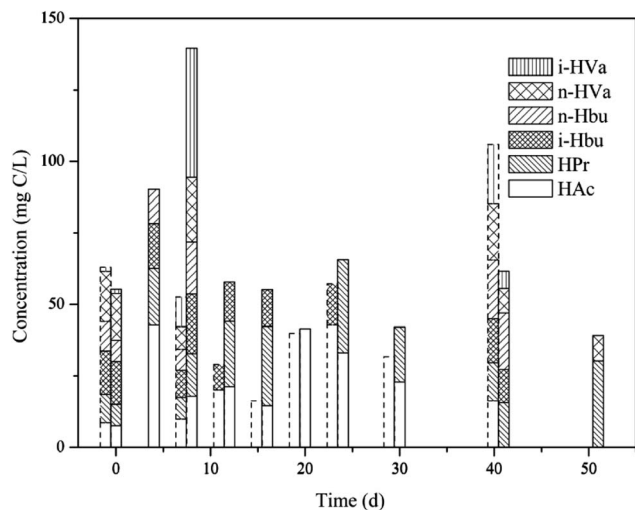


Fig. 5 VFAs variations in the two reactors (dashed line: MNSR; solid line: CMSR).

As shown in Fig. 5, VFAs accumulation was not detectable in the MNSR during the first week. Then VFAs concentration increased with the increase in CH_4 production. However, at the end of the experiments VFAs was not detectable in the MNSR, indicating that VFAs in this reactor experienced a production followed by consumption progress. It has been pointed out that a high dissolved H_2 concentration inhibits propionate and butyrate conversion to acetate or H_2 and CO_2 during anaerobic digestion, yielding lower conversion rate or the whole process breakdown.²⁰ Thus, a higher acetic acid concentration in the MNSR can be used for more CH_4 production when compared to the CMSR, possibly contributed by the higher dissolved H_2 concentration in the reactor.

From the whole process, the concentration of total VFAs in the MNSR was lower than that in the CMSR, indicating more substrates were converted into CH_4 . Meanwhile in the CMSR, VFAs accumulation was detected at the beginning of experiment. From day 12, the total VFAs concentration decreased while maintained at a relatively stable level, reflecting that VFAs consumption in the CMSR was not so efficient.

Carbon mass balance analysis

To evaluate carbon conversion efficiency in the reactors, carbon elements in the liquid, solid and gaseous phases were quantified as illustrated in Fig. 6. The C content in biomass was estimated according to a theoretical formula ($\text{C}_{60}\text{H}_{87}\text{N}_{12}\text{O}_{23}\text{P}$) for microbes. In addition, the calculation of $C_{L,\text{TVFAs}}$ was based on the individual VFA. During the whole experiments, except for the initial carbon in the biomass and medium, the carbon input is only from CO_2 in the gas phase. The carbon balance was analyzed according to eqn (8). Being consistent with the results from above two sections, a relatively more carbon conversion into the solid phase ($C_{S,\text{Biomass}}$) was achieved in the MNSR in comparison to the CMSR. And the final biomass yield in the MNSR was 2.34 g-biomass per L, which was 1.98 g-biomass per L in the CMSR. Anyhow, the liquid phase carbon percentage gradually was found to decrease to a low level in both reactors,

which might be utilized as a source for biomass growth. Considering the gaseous phase carbon fractions, C_{G,CH_4} in the two reactors showed a remarkable difference. Still, a considerable part of CO_2 ($C_{G,\text{CO}_2,\text{Rest}}$) was remained in both systems. Seen from the whole conversion process, the input CO_2 tended to be employed for microbial growth first, and then the C_{G,CH_4} increased with the increase in biomass growth.

Energy consumption

The economic and practical feasibility of this enhancement approach should be evaluated for its applications in large scale fermentation systems. MNS clearly has a significant stimulation potential for hydrogenotrophic methanogenesis. Enrichment of methanogens in the MNSR provides a greater H_2 -bioconversion potential than in the CMSR, suggesting that the gas-liquid mass transfer limitation is minimized. However, the advantage could be mitigated by high energy consumption. In this context, energy consumption analysis is essential for both the MNSR and CMSR. Table 1 summarizes the results relating to the energy consumptions by the MNSR and CMSR according to eqn (11) and (12).

As shown in Table 1, both the work of pump (E_{pump}) and stirrer (E_{stirrer}) in the MNSR were higher than those in the CMSR, leading to the higher energy consumption in the MNSR system. However, when evaluating the technology, the product value (CH_4 in this work) also should be taken into account. In this work, the energy-product ratio (R) was employed to represent the potential for practical application. As for the MNSR, although a higher CH_4 yield is corresponding to a higher energy consumption, its R value is still higher than the that for the CMSR, demonstrating its great potential for scaling-up application. In fact, continuous-type reactors dominate the industrial scale fermentation systems, and the methane yield can also be improved by increasing the gas recirculation rate. As proposed by Szuhaj *et al.*,⁴³ the energy for H_2 production could be supplied by the renewable resources such as the wind or solar energy. In this case, the application of MNSR for bioconversion of H_2 and CO_2 is more meaningful.

Gas-liquid mass transfer evaluation

In order to find out the reason for the enhancement effect by MNS, the H_2 gas-liquid mass transfer was evaluated based on the dissolved H_2 concentration. Gas mass transfer in the two reactors occurs in two zones. Gas transfer in the headspace happens through a very thin liquid boundary layer between the bulk gas and the culture cells. The mass transfer in the liquid phase is characterized through quantification of the volumetric coefficient ($k_L a$). Vega *et al.*⁴³ have described the multiple steps when mass transfer occurs from gas to liquid phases, which involve (1) the absorption of a gaseous substrate across the gas-liquid interface, (2) the transfer of the dissolved gas to the fermentation media, and (3) diffusion through the culture media to the cell surface. And the most sparingly soluble gases utilized in the biochemical reactions trigger the major resistance in the liquid film around the gas-liquid interface.^{44,45}



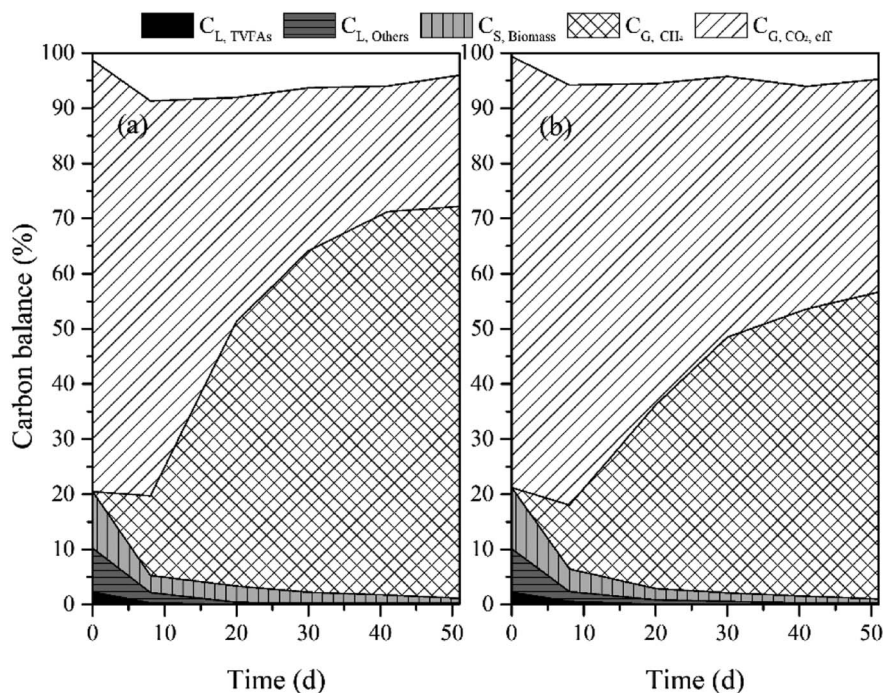


Fig. 6 Carbon balance analysis for MNSR (a) and CMSR (b).

Table 1 Energy consumption for the two reactor systems

Reactor	E_{pump} (W)	E_{stirrer} (W)	E_{consumed} (W)	R (L kW ⁻¹)
MNSR	2.80×10^4	1.59×10^4	4.39×10^4	1.80
CMSR	2.69×10^4	1.39×10^4	4.08×10^4	1.45

Because during the batch bioconversion experiments, the $k_L a$ value was continuously varying, the $k_L a$ in this study was evaluated before incubation by analyzing the dissolved H_2 concentration in the liquid. As shown in Fig. 7, the $k_L a$ value in the MNSR (12.95 h^{-1}) is almost twice that in the CMSR (6.6 h^{-1}), most probably due to the increase in specific surface area because of much smaller bubble size.

The specific surface area to liquid volume ratio was calculated by according to both the micro-scale and nano-scale bubble size distribution in the two reactors (Fig. S3†). The average bubble size in the CMSR and MNSR was determined as about $845 \mu\text{m}$ and $220 \mu\text{m}$, respectively, corresponding to the total surface area to liquid volume of $5640 \text{ cm}^2/\text{L}_R$, and $48\,042 \text{ cm}^2/\text{L}_R$. This specific surface area has been increased by one order of magnitude, implying more sufficient contact chance between methanogens and the gaseous substrates.

Results show the significant difference in the $k_L a$ value of MNSR and CMSR. There are several parameters, which have an influence on the $k_L a$ in a reactor. Rittman *et al.*¹⁶ (2015) demonstrated that at a given reactor configuration, agitation, stirrer types, gas flow rate, gas-limitation fundamentals and the sparger type are the most relevant. The $k_L a-H_2$ in different studies were summarized in Table 2. Nielsen *et al.*⁴⁶ proposed that CSTR was the most used reactor configuration for

bioconversion of H_2 and CO_2 to CH_4 by 2011, due to the effective gas-liquid mass transfer in the CSTR. In the study of Bassani *et al.*,²⁶ the effect of sparger type on $k_L a$ value was not very significant, but gas recirculation rate affected the bioconversion significantly. While, in this study, the sparger type is still an important factor. The smaller pore size lead to the higher $k_L a$ value with the assistance by magnetic stirrers to provide sufficient mixing. However, the $k_L a$ obtained from this study is much lower than that from Bassani *et al.*,²⁶ possibly resulted from the lower recirculation rate applied in this study (40 mL min^{-1}). Therefore, the subsequent experiments will be carried out to optimize the $k_L a$ value taking higher recirculation

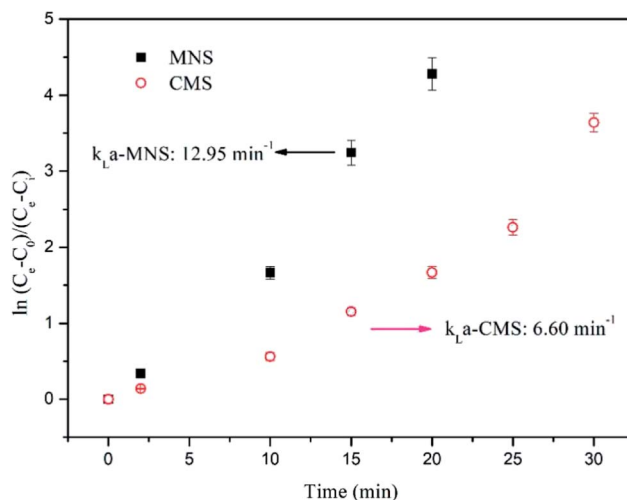


Fig. 7 Determination of $k_L a-H_2$ from dissolved H_2 concentration data. Solid squares denote MNSR and open circles denote CMSR.



Table 2 Comparison in $k_{L,a}$ value among different biogas methanation systems

Bioreactor features	Sparger pore size	Working volume (L)	Cultivation modes	Inoculum	Temp. (°C)	$k_{L,a}$ (h^{-1})	Reference
CSTR	0.5–1.0 mm/14–40 μm	0.6	Chemostat	Digested manure	55	6.6–16.05	Luo and Angelidaki, 2013
UBF	5 μm	19.3	Chemostat	Anaerobic granules	36 ± 2	0.72–20.86	Frigon and Guiot, 1995
CSTR	—	3.07–13.4	Fed-batch	Anaerobic sludge	32–35	0.06–0.16	Pauss <i>et al.</i> , 1990
UASB	0.5 $\mu\text{m}/2 \mu\text{m}$	0.85	Chemostat	Hydrogenotrophic methanogens	55	240–776	Bassani <i>et al.</i> , 2017
CSTR	0.2 μm -MNS/845 μm -CMS	0.5	Fed-batch	Anaerobic sludge	37	12.95-MNS/6.60-CMS	This study

rate and higher temperature into consideration. In a summary, these results demonstrate that $k_{L,a}$ is a key factor for the enhanced CH_4 production from the MNSR.

Conclusions

The enhanced bioconversion of H_2 and CO_2 to CH_4 was realized by using the MNS. The maximum MER of 171.4 mmol/ L_R/d in the MNSR is much higher than that in the CMSR (136.1 mmol/ L_R/d). The MNSR also displayed superior biomass growth with specific growth rate of 0.15 d^{-1} than that of CMSR. The VFAs accumulation was not detectable in MNSR. Higher gas-liquid mass transfer was achieved in the MNSR than that in the CMSR. A higher energy-product ratio and economic analysis indicates that MNS has applicable potential for improvement of CO_2 and H_2 conversion into CH_4 in large-scale plants.

Conflicts of interest

There are no conflicts to declare.

Acknowledgements

This work was financially supported by JFE steel cooperation. The first author acknowledges the support of the China Scholarship Council (No. 201506400049).

References

- 1 G. J. Francisco, A. Chakama and X. Feng, *Sep. Purif. Technol.*, 2010, **75**, 205–213.
- 2 Greenhouse Gas Inventory Office of Japan (GIO), 2017, ch. 2, pp. 3–6.
- 3 U. E. Aronu, H. F. Svendsen and K. A. Hoff, *Int. J. Greenhouse Gas Control*, 2010, **4**, 771–775.
- 4 A. Sayari, Y. Belmabkhout and R. Serna-Guerrero, *Chem. Eng. J.*, 2010, **171**, 760–774.
- 5 G. Xu, F. Liang, Y. Yang, Y. Hu, K. Zhang and W. Liu, *Energies*, 2014, **7**, 3484–3502.
- 6 Y. Zee, L. C. Thiam, W. Z. Peng, R. M. Abdul and S. Chai, *J. Porous Mater.*, 2013, **20**, 1457–1475.
- 7 B. Pau, P. Sebastia, G. Rafael, D. Maria and C. Jesus, *J. Chem. Technol. Biotechnol.*, 2016, **91**, 921–927.
- 8 M. C. A. A. van Eerten-Jansen, N. C. Jansen, C. M. Plugge, V. Wilde, C. J. N. Buisman and A. T. Heijne, *J. Chem. Technol. Biotechnol.*, 2015, **90**, 963–970.
- 9 M. Burkhardt and G. Busch, *Appl. Energy*, 2013, **111**, 74–79.
- 10 M. K. Lam, K. T. Lee and A. R. Mohamed, *Int. J. Greenhouse Gas Control*, 2012, **10**, 456–469.
- 11 H. S. Jee, N. Nishio and S. Nagai, *J. Ferment. Technol.*, 1988, **66**, 235–238.
- 12 D. Ju, J. Shin, H. Lee and S. Kong, *Desalination*, 2008, **234**, 409–415.
- 13 M. Szuhaj, N. Acs, R. Tengolics, A. Bodor, G. Rakhely and K. L. Kovacs, *Biotechnol. Biofuels*, 2016, **9**, 102.
- 14 S. K. Hoekman, A. Broch, C. Robbins and R. Purcell, *Int. J. Greenhouse Gas Control*, 2010, **4**, 44–50.

- 15 M. D. Redwood, R. L. Orozco, A. J. Majewski and L. E. Macaskie, *Bioresour. Technol.*, 2012, **107**, 166–174.
- 16 S. Rittmann, A. Seifert and C. Herwig, *Crit. Rev. Biotechnol.*, 2015, **35**, 141–151.
- 17 N. Nishimura, S. Kitaura, A. Mimura and Y. Takahara, *J. Ferment. Bioeng.*, 1991, **72**, 280–284.
- 18 I. Bassani, P. G. Kougias, L. Treu and I. Angelidaki, *Environ. Sci. Technol.*, 2015, **49**, 12585–12593.
- 19 I. Bassani, P. G. Kougias and I. Angelidaki, *Bioresour. Technol.*, 2016, **221**, 485–491.
- 20 I. Diaz, C. Perez, N. Alfaro and F. Fdz-Polanco, *Bioresour. Technol.*, 2015, **185**, 246–253.
- 21 D. Cheng, J. Cheng, X. Li, X. Wang, C. Yang and Z. Mao, *Chem. Eng. Sci.*, 2012, **75**, 256–266.
- 22 S. R. Guiot, R. Cimpoia and G. Carayon, *Environ. Sci. Technol.*, 2011, **45**, 2006–2012.
- 23 G. Luo and I. Angelidaki, *Appl. Microbiol. Biotechnol.*, 2013, **97**, 1373–1381.
- 24 S. Zhang, D. Wang, F. Bu, X. Zhang and P. Fan, *Surf. Interface Anal.*, 2013, **45**, 1152–1157.
- 25 P. G. Kougias, L. Treu, D. P. Benavente, K. Boe, S. Campanaro and I. Angelidaki, *Bioresour. Technol.*, 2017, **225**, 429–437.
- 26 I. Bassani, P. G. Kougias, L. Treu, H. Porte, S. Campanaro and I. Angelidaki, *Bioresour. Technol.*, 2017, **234**, 310–319.
- 27 A. J. Dixon, A. H. Dhanaliwala, J. L. Chen and J. A. Hossack, *Ultrasound Med. Biol.*, 2013, **39**, 1267–1276.
- 28 F. Kobayashi, H. Ikeura, M. Tamaki and Y. Hayata, *Southeast Asia Symposium on Quality and Safety of Fresh and Fresh-cut Produce*, 2010, 875, pp. 417–424.
- 29 K. Kugino, S. Tamaru, Y. Hisatomi and T. Sakaguchi, *PLoS One*, 2016, **11**, 1–9.
- 30 A. Agarwal, W. J. Ng and Y. Liu, *Chemosphere*, 2011, **84**, 1175–1180.
- 31 F. Ushikubo, T. Furukawa, R. Nakagawa, M. Enari, Y. Makino, Y. Kawagoe, T. Shiina and S. Oshita, *Colloids Surf., A*, 2010, **361**, 31–37.
- 32 Z. Zhang and T. Maekawa, *Biomass Bioenergy*, 1993, **4**, 439–446.
- 33 APHA, *American Public Health Association/American Water Work Association/Water Environment Federation*, Washington, DC, 21st edn, 2005.
- 34 W. Huang, W. Huang, T. Yuan, Z. Zhao, W. Cai, Z. Zhang, Z. Lei and C. Feng, *Water Res.*, 2016, **90**, 344–353.
- 35 P. Tsapekos, P. G. Kougias, L. Treu, S. Campanaro and I. Angelidaki, *Appl. Energy*, 2016, **185**, 126–135.
- 36 P. J. Weimer and J. G. Zeikus, *Arch. Microbiol.*, 1978, **119**, 49–57.
- 37 P. H. Roennow and L. A. H. Gunnarsson, *FEMS Microbiol. Lett.*, 1982, **14**, 311–315.
- 38 J. Lee, J. Kim, W. Chang and D. Pak, *J. Chem. Technol. Biotechnol.*, 2012, **87**, 844–847.
- 39 P. C. Munasinghe and S. K. Khanal, *RSC Adv.*, 2014, **4**, 37575–37581.
- 40 D. Karakashev, D. J. Batstone, E. Trably and I. Angelidaki, *Appl. Environ. Microbiol.*, 2006, **72**, 5138–5141.
- 41 M. J. Lee and S. H. Zinder, *Appl. Environ. Microbiol.*, 1988, **54**, 1457–1461.
- 42 R. Cord-Ruwisch, D. R. Lovley and B. Schink, *Appl. Environ. Microbiol.*, 1998, **64**, 2232–2236.
- 43 J. L. Vega, E. C. Clausen and J. L. Gaddy, *Biotechnol. Bioeng.*, 1989, **34**, 774–784.
- 44 K. T. Klasson, C. M. D. Ackerson, E. C. Clausen and J. L. Gaddy, *Int. J. Hydrogen Energy*, 1992, **17**, 281–288.
- 45 P. C. Munasinghe and S. K. Khanal, *Biotechnol. Prog.*, 2010, **26**, 1616–1621.
- 46 J. Nielsen, J. Villadsen, and G. Liden, *Bioreaction engineering principles*, Springer, New York, 2011, ch. 9, pp. 383–458.

