Green hydrogen and platform chemicals production from acidogenic conversion of brewery spent grains co-fermented with cheese whey wastewater: adding value to acidogenic CO₂

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The biotechnological production of fuel and chemicals from renewable, organic carbon-rich substrates offers a sustainable way to meet the increasing demand for energy. This study aimed to generate platform chemicals, which serve as precursors for the synthesis of fuels and various materials, along with green hydrogen (bio-H₂) by co-fermenting two different waste streams: brewery spent grains and cheese whey (CW). Reactors fermenting a fixed quantity of brewery-spent grains were loaded with CW at 20, 30, and 40 g COD per L, and microbial production of short-chain (SCCA) and medium-chain carboxylic acids (MCCA) along with bio-H₂ was assessed. The reactor with the highest organic load (40 g COD per L) produced the highest amount of SCCA (21.67 g L⁻¹) whereas bio-H₂ was with 30 g COD per L (181.35 mL per day). In the next phase, the generated gas (H₂ + CO₂) was continuously recirculated within the reactor to enhance SCCA production by a further 19.9%. In the later stages of fermentation, MCCA production indicated the occurrence of chain elongation from the accumulated lactic acid. Consumption of H₂ and CO₂ during gas recirculation highlighted the role of bio-H₂ as an electron donor and acidogenic CO₂ as a precursor molecule in the chain elongation process. As a result, no external reducing agent was required and only limited CO₂ was released in the atmosphere, making the overall process more sustainable and cost-effective.

Introduction

Fossil-based fuels in the form of coal, oil, and natural gas remain the source of 80% of the world’s energy, but they also strongly contribute to global warming.⁰ Burning fossil-based fuels accounts for 89% of human-derived CO₂ emissions and, according to the Intergovernmental Panel on Climate Change, could cause global mean surface temperatures to rise by 1.5 °C above the pre-industrial mark in as little as a decade. To prevent catastrophic warming, it will be necessary to replace these fossil-based fuels with a sustainable alternative for energy and material synthesis.² Hydrogen is considered one of the most promising energy carriers due to its elevated energy density (141.9 MJ kg⁻¹), clean emissions (H₂O as the only combustible byproduct), and widespread abundance.¹,³⁻⁵ Recently, the hydrogen has been pitted as a significant upcoming energy source in our global landscape and thus both develop and developing nations and are making the low carbon energy source a central part of their strategy to decarbonize.

Currently H₂ production via electrolysis of water, steam reforming of natural gas followed by cracking oil products, coal gasification remains the main route, whereas less energy-demanding biological processes for hydrogen production are considered a promising alternative, which have garnered increasing attention.⁶ Dark fermentation/acidogenic fermentation is a versatile process capable of efficiently converting various organic substrates (waste/wastewater) to bio-hydrogen (bio-H₂) under ambient temperature and pressure.⁶⁻⁸ The added advantage of acidogenic fermentation is co-production of short-chain carboxylic acids (SCCA), including acetic (C₂), propionic (C₃), butyric (C₄), and valeric (C₅) acids, which can serve as platform chemicals for industrial applications.⁹,¹⁰ The demand for volatile fatty acids (VFAs), including the above SCCA, is expected to increase over the coming years due to their numerous applications as fuel precursors, as well as in pharmaceutical, and household chemical formulations.¹⁰ while global production of chemicals doubled over the past two decades, reaching 2.3 billion tons in 2017, only 2% of them were bio-based. Furthermore, fossil-based chemical production consumes 20% of the energy used for industrial purposes. Switching from a fossil-based to a bio-based economy remains a challenge, but it represents also a necessary step to meet the UN Sustainable Development Goals.

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Important, SCCA can be upgraded to value-added caproic acid (C6) and other medium-chain carboxylic acids (MCCA) via reverse β-oxidation in the presence of an external electron donor, such as ethanol, methanol or lactic acid. During reverse β-oxidation, SCCA are elongated via the addition of two carbon atoms per cycle, while the electron donor is converted to acetyl-CoA, acetoacetyl-CoA, and butyryl-CoA. The latter can react with acetate to generate butyrate, while another acetyl-CoA can react with butyryl-CoA to form caproyl-CoA, and thereby lead to caproate.

The conversion of SCCA to MCCA is carried out by microorganisms and a crucial role is played by electron donors. Increasing attention has been garnered by new solutions such as a carbonate platform with mixed culture fermentation for the production of MCCA (6 to 12 carbon atoms) through carboxylic chain elongation. With high caloric value and slightly hydrophobic properties, caproic acid is a suitable intermediate for biofuel (i.e., isobutyl hexanoate as drop-in additive in A-1 jet fuel), pharmaceuticals, and biochemical production. Moreover, upgrading SCCA to MCCA via reverse β-oxidation can help overcome the limitations associated with the high costs of recovery and purification of SCCA. Moreover, MCCA have a higher market value than SCCA, estimated at 2000–3000 $ per t, and a market demand of 25 000 tons per year.

The aim of this study was to evaluate the production of acidic bio-H2 and carboxylic acids from a renewable feedstock such as brewery-spent grains (BSG) co-fermented with cheese whey (CW). These two waste streams complement each other, as BSG is rich in protein and polysaccharides, while CW is rich in lactose. Co-fermentation enhances system stability and the synthesis of microbial metabolites due to synergistic effects that promote a more diverse microbial community, better nutrient balance, and access to trace elements essential for the fermentation process.

For a large-scale application, fermentation of this waste/wastewater can be conjugated with the existing anaerobic digestion/sewage treatment plants allowing to settle overnight. The supernatant (mostly water) was removed and the thickened sludge with a volatile solids (VS) content of 0.56 g g⁻¹ was used as biocatalyst. Prior to use, 2-bromoethanesulfonic acid (4 g L⁻¹) was added to the sludge to suppress methanogens. To promote an active bacterial population, the sludge was incubated at ambient temperature for 72 h with a nutrient solution containing 5 g L⁻¹ glucose, 0.5 g L⁻¹ NH₄Cl, 0.25 g L⁻¹ KH₂PO₄, 0.25 g L⁻¹ K₂HPO₄, 0.3 g L⁻¹ MgCl₂, 25 mg L⁻¹ CoCl₂, 11.5 mg L⁻¹ ZnCl₂, 10.5 mg L⁻¹ CuCl₂, 5 mg L⁻¹ CaCl₂, 15 mg L⁻¹ MnCl₂, 16 mg L⁻¹ NiSO₄, and 25 mg L⁻¹ FeCl₃, before inoculation in the reactor system.

**Experimental**

**Inoculum**

Anaerobic sludge was collected from a biogas plant in Luleå, Sweden. The sludge was filtered using a stainless steel mesh to remove grit and other solid particles (e.g., hair and paper) and allowed to settle overnight. The supernatant (mostly water) was removed and the thickened sludge with a volatile solids (VS) content of 50% H₂ and 50–60% CO₂, in which the cogenerated CO₂ limits the use of bio-H₂ as a fuel. Many studies have demonstrated the utilization/removal of aceticogenic CO₂ by adopting different strategies such as chemical/physical adsorption, membrane/vacuum separation, electrochemical processes etc. Despite these processes significantly upgrade the bio-H₂, poses a few limitations to these processes as the captured CO₂ is released back into the atmosphere and amplifies greenhouse heating. CO₂ storage and capturing technologies on the other hand are expensive, at the same time developing a process towards sequestration of CO₂ for the sustainable production of fuels and chemicals become a current research hotspot and has important strategic and real economic significance.

Currently numerous studies has been carried out finding a potential mitigation options such as utilization of low carbon dependent fuels as chemicals/feedstock’s and fuels including biomass as well as CO₂ capture and storage (CCS) in order to reduce greenhouse gas (GHG) emissions. Moreover, the separation processes for CO₂ demands retrofitting to the traditional processes, which directly influence the overall investment. On the other side, the biological routes offers an attractive approach for the utilization/conversion of CO₂ as the operating principle towards CO₂ reactions occurs naturally in microbes (eqn (1)–(3)).

\[2\text{CO}_2 + 4\text{H}_2 \rightarrow \text{CH}_3\text{COO}^- + \text{H}^+ + 2\text{H}_2\text{O} \quad (1)\]

\[2\text{CH}_3\text{COO}^- + \text{H}^+ + 2\text{H}_2 \rightarrow 2\text{CH}_3(\text{CH}_2)_2\text{COO}^- + 2\text{H}_2\text{O} \quad (2)\]

\[4\text{CO}_2 +10\text{H}_2 \rightarrow \text{CH}_3(\text{CH}_2)_2\text{COO}^- + \text{H}^+ + 6\text{H}_2\text{O} \quad (3)\]

"CO₂-reducing acetogen" or a “homoacetogen” takes acetyl-CoA biochemical pathway for the formation of acetic acid as fermentation product from CO₂. Thus, here we studied the utilization of the in situ CO₂ produced during acidicogenic co-fermentation of cheese whey and BSG for an enhanced biosynthesis of microbial metabolites, at the same time to limit the release of CO₂ into the environment.

**Feedstock preparation and characterisation**

BSG used in this study was provided by Skellefteå Bryggeri (Skellefteå, Sweden). Prior to use, BSG was oven-dried at 65 °C for 12 h and stored in a sealed bag. Highly heterogeneous BSG was homogenised using a kitchen mixer (SM-1FP; Wilfa), which delivered particles of 0.5–1 cm. The homogenised BSG contained 96.2% ± 0.02% w/w total solids, of which 94.2% ± 0.03% w/w were VS. Cellulose, hemicellulose, and lignin accounted for 29.35%, 16.64%, and 13.33% w/w of BSG, respectively.
The organic content of CW was determined as 75.6 g chemical oxygen demand (COD) per L while pH was 5.7. A major fraction of CW was represented by lactose (48 g L\(^{-1}\)), along with traces of lactic acid (0.04 g L\(^{-1}\)), acetic acid (0.08 g L\(^{-1}\)), propionic acid (0.01 g L\(^{-1}\)), and butyric acid (0.07 g L\(^{-1}\)). Based on the required organic load (20, 30, and 40 g COD per L), CW was diluted with tap water.

**Experimental procedure**

The experiments were conducted in 18 identical 2000 mL glass bottle reactors (triplicates of six experiments) in two phases (P-I and P-II) using the AMPTS-II automated analytic system (Bio-process Control). During P-I, the effect of varying the organic load of CW on carboxylic acids and bio-H\(_2\) recovery was evaluated. The biogas (H\(_2\) + CO\(_2\)) produced during acidogenic fermentation was recirculated in P-II to provide a source of inorganic carbon (CO\(_2\)) and an electron donor (H\(_2\)) for the homoacetogens in the mixed culture. Based on the experimental design and conditions, reactors were labelled as R\(_{20}\) (20 g COD per L), R\(_{30}\) (30 g COD per L), and R\(_{40}\) (40 g COD per L) when operated in P-I, and GC-R\(_{20}\) (20 g COD per L), GC-R\(_{30}\) (30 g COD per L), and GC-R\(_{40}\) (40 g COD per L) in P-II, with GC corresponding to gas recirculation (Table 1 and Fig. 1). All reactors were operated for 56 days in batch mode under mesophilic conditions (35 °C). No further nutrients were added during fermentation because BSG was sufficiently rich. Prior to start up, the pH in the reactors was adjusted using 2 M HCl/NaOH, after which it was set manually to 6.0–6.5. Nitrogen gas was sparged into the reactor for 30 min to maintain anaerobic conditions. The reactors were kept in suspension mode during the reaction phase by continuous mixing with a stirrer fixed to the cap. All fermentation tests and measurements were conducted in triplicate, and the average values and standard deviation were reported.

**Biochemical and gas analyses**

COD of CW was analyzed using the Spectroquant NOVA 60A COD cell test kit (Merck Millipore). Changes in pH were measured with a pH meter (pHenomenal-pH1100L; VWR). Total solids (TS) and volatile solids (VS) were estimated as described by Matsakas et al. (2020). Microbial metabolites, including lactic (H\(_{1}\)lac), acetic (H\(_{1}\)ac), propionic (H\(_{3}\)pr), butyric (H\(_{4}\)bu), valeric (H\(_{5}\)va), and caproic (H\(_{6}\)ca) acids, were analysed by high-performance liquid chromatography (HPLC) and quantified with calibration curves generated.
from commercially available standards (10 mM, Volatile Free Acid Mix; Sigma). The HPLC apparatus (PerkinElmer) was equipped with a Flexar LC pump, Bio-Rad Aminex HPX-87H column (300 m × 7.8 mm), and PerkinElmer-200 refractive index detector. Column temperature was maintained at 65 °C. The mobile phase consisted of 5 mM H₂SO₄ and was eluted at 0.6 mL min⁻¹. Biogas production and composition was analyzed using a mass spectrometer (GAM 400; InProcess Instrument).

Results and discussion

Total carboxylic acids production

Co-fermentation of protein and carbohydrate-rich BSG and CW enhanced carboxylic acid production beyond what had been observed previously when using BSG as sole carbon source.³⁶ Production performance was investigated by varying the COD of CW (20, 30, and 40 g COD per L) while maintaining a fixed BSG content (35 g VS). Total carboxylic acid production increased with fermentation time in all reactors (Fig. 2), with variations based on the initial COD. Specifically, production was more or less similar until day 8, when it ranged around 3.45–4.66 g L⁻¹ by day 16 (Fig. 2a). On day 24, the three reactors displayed diverging patterns, with R₄₀ attaining a production of 19.63 g L⁻¹, followed by R₃₀ (12.89 g L⁻¹) and R₂₀ (11.58 g L⁻¹). By day 40, reactor R₄₀ reached 24.53 g L⁻¹, while R₃₀ followed with 22.06 g L⁻¹ and R₂₀ with 13.21 g L⁻¹. By the end of the experiment (day 56), reactor R₃₀ decreased slightly to 20.91 g L⁻¹, whereas R₄₀ gradually increased production to 26.35 g L⁻¹ and R₂₀ to 16.14 g L⁻¹. Overall, reactor R₄₀ achieved 1.6-times and 1.2-times greater production than R₃₀ and R₂₀, respectively.

Previously Teixeira et al., found a good carboxylic acids production of 43.8 g COD as a major microbial metabolite from an untreated BSG during a long-term (HRT 41 days) fed-batch acidogenic fermentation.²⁷ Liang and Wan demonstrated a mixture of carboxylic production from BSG at alkaline pH 10 with higher fraction of acetic acid (6.3 g L⁻¹).³⁸ In a two-step conversion of acid pretreated hydrolysate of BSG, Guarda et al. (2021) found a mixture of carboxylic acids (9 ± 1.59 g COD per L) from a continuously operated reactor at an organic load rate (OLR) of 8.11 ± 0.87 g COD per L per day.²⁹ Fermenting the hydrolysate derived from BSG through acid pretreatment, Mussatto et al. (2007) showed a lactic acid production of 5.4 g L⁻¹ by Lactobacillus delbrueckii.⁴⁰

Composition of carboxylic acids (P-I)

Total carboxylic acids included H₅Lac, H₅Ac, H₅Pr, H₅Bu, H₅Val, and H₅Ca. Their individual concentration varied with respect to fermentation time, VS, and COD load (Fig. 3). The reactor with the highest COD achieved also the highest H₅Lac output (Fig. 3a). The concentration of H₅Lac was maximal between 8 and 24 days, with peak production of 9.7 g L⁻¹ (10.38 g COD per L, R₄₀; day 24), followed by 6.8 g L⁻¹ (R₃₀; day 24) and 3.6 g L⁻¹ (R₂₀; day 16). These amounts corresponded to 41.12%, 37.6%, and 31% of the total carboxylic acids accumulated in the reactor, and reflected the influence of a higher COD, and hence lactose content, in CW. Mixed culture fermentation converts monosaccharides and disaccharides to H₅Lac.⁴¹ Lactose-rich CW is initially broken down to glucose and galactose, after which it is converted to H₅Lac via the glycolytic pathway.²³,⁴² Similarly,

Fig. 2 Total carboxylic acids production from a fixed quantity of BSG (35 g VS) co-fermented with a varied organic load of CW (20, 30 and 40 g COD per L) in phase-I (P-I) and phase-II (P-II) where the acidogenic CO₂ was recirculated within the reactor in phase-II.
Pyruvate can be converted through acetyl-CoA to ethanol, HAc, HBu or even other metabolites depending on the microbial microenvironment.\textsuperscript{23} Yu et al. (2004) detected a mixture of carboxylic acids and solvents during acidogenic fermentation of lactose-rich wastewater at pH 5.5.\textsuperscript{43} Fermenting the high strength CW in a anaerobic sequencing batch reactor, Costa et al., reported carboxylic acids production of 10.23 g COD per L accounting for 4.54 g COD per L per day,\textsuperscript{44} Atasoy et al., reported the production of 0.97 g COD per g SCOD with major fraction of butyric acid at alkaline condition in the batch reactor.\textsuperscript{45} HAc and HBu were the two major carboxylic acid fractions, and their concentration gradually increased with fermentation time; whereas HPr remained low throughout the experiment (Fig. 3b–d). HAc biosynthesis was greater in reactor R30 (8.38 g L\textsuperscript{-1}), followed closely by R20 (7.9 g L\textsuperscript{-1}) and R40 (7.6 g L\textsuperscript{-1}) (Fig. 3b). In contrast, total HBu production was significantly higher (9.89 g L\textsuperscript{-1}) under high COD load in CW (R40) compared to R30 (6.84 g L\textsuperscript{-1}) and R20 (5.11 g L\textsuperscript{-1}) (Fig. 3d). HBu biosynthesis occurs via (i) phosphotransbutyrylase and butyrate kinase, or (ii) butyryl CoA:acetate CoA transferase metabolic
Some members of *Clostridium* are capable of elongating H$_{\text{Lac}}$ to H$_{\text{Bu}}$ without the involvement of H$_{\text{Ca}}$ as they can convert butyryl-CoA to H$_{\text{Bu}}$ through phosphorylation instead of via cyclical reverse β-oxidation.\textsuperscript{20} Biosynthesis and concentration of H$_{\text{Pr}}$ were relatively low compared to those of other carboxylic acids and were detected only after day 16 (Fig. 3c). In the present set-up, H$_{\text{Pr}}$ production can be attributed to the degradation of proteins and carbohydrates present in BSG and CW through amino acid catabolic and biosynthetic pathways.\textsuperscript{46} The stronger production of H$_{\text{Pr}}$ in reactor R$_{40}$ (2.25 g L$^{-1}$/C0$_{1}$; day 48) compared to R$_{30}$ (1.53 g L$^{-1}$/C0$_{1}$; day 40) and R$_{20}$ (1.2 g L$^{-1}$/C0$_{1}$; day 40) is related to a greater availability of amino acids. A decline in H$_{\text{Pr}}$ was noticed by the end of the experiment, indicating its biotransformation, together with H$_{\text{Ac}}$ and H$_{\text{Bu}}$, to H$_{\text{Va}}$ and H$_{\text{Ca}}$. An increasing concentration of these carboxylic acids in the reactor, along with their simultaneous transformation to longer-chain molecules during fermentation indicated an efficient and continuous hydrolysis of BSG by a mixed microbial culture, which delivered a continuous stream of sugars for fermentation.

Unlike other carboxylic acids, H$_{\text{Va}}$ and H$_{\text{Ca}}$ were not detected during the initial stages of fermentation (Fig. 3e and f). H$_{\text{Va}}$ production started on day 32 with relatively low levels (0.54 g L$^{-1}$/C0$_{40}$, 0.24 g L$^{-1}$/C0$_{30}$, and 0.12 g L$^{-1}$/C0$_{20}$) (Fig. 3e). These values gradually increased with time and reached a maximum on day 56, with 1.64 g L$^{-1}$/C0$_{40}$, 0.94 g L$^{-1}$/C0$_{30}$, and 0.87 g L$^{-1}$/C0$_{20}$.

**Production and consumption rate of carboxylic acids (P-I)**

The production ($P_{\text{rate}}$) and consumption rate ($C_{\text{rate}}$) of carboxylic acids showed a distinct trend with respect to the initial load of CW in the reactor and fermentation time (Fig. 4). In P-I, $P_{\text{rate}}$ of H$_{\text{Bu}}$ was significantly higher compared to other carboxylic acids. Initially until day 16, the $P_{\text{rate}}$ of H$_{\text{Bu}}$ was ranged between 0.08–0.31 g COD per L per day in all the reactors which later increased on day 24 specifically with R$_{40}$ exhibiting the highest $P_{\text{rate}}$ of 1.15 g COD per L per day. Further from day 48, its consumption was observed with R$_{20}$ (−0.05 g COD per L per day) which later increased to −0.17 g COD per L per day during day 56 and stabilized thereafter. H$_{\text{Bu}}$ consumption was not seen with R$_{30}$ and R$_{40}$ indication its acidogenic production and continuous accumulation in the reactor with a $P_{\text{rate}}$ of 0.11–0.23 g COD per L per day despite its transformation to chain elongated H$_{\text{Ca}}$. $P_{\text{rate}}$ of H$_{\text{Ca}}$ was maximum with R$_{30}$ (0.36 g COD per L per day) followed by R$_{40}$ (0.29 g COD per L per day) and R$_{20}$ (0.24 g COD per L per day) on different time interval of fermentation. H$_{\text{Pr}}$ production was started after day 8, by day 16, $P_{\text{rate}}$ of H$_{\text{Pr}}$ reached to 0.15 g COD per L per day (R$_{30}$), however its maximum $P_{\text{rate}}$ was noticed on day 40 with R$_{40}$ (0.17 g COD per L per day). Afterwards its consumption from day 48, particularly with R$_{30}$ and R$_{40}$ reduced its concentration to less than 0.91 g COD per L and 1.72 g COD per L respectively. On the contrary, although the production declined ($P_{\text{rate}}$ of 0.02 g COD per L per day), its consumption was not documented until the end of the cycle, indicating its continuous biosynthesis from the fermentable sugars by mixed culture. Initially with a $P_{\text{rate}}$ ranging between 0.03–0.12 g COD per L per day, later no great improvement was observed indicating a stabilized production.

![Fig. 4](https://example.com/f4.png) Distribution of carboxylic acids production ($P_{\text{rate}}$) and consumption rate ($C_{\text{rate}}$) pattern in P-I (a–f) with respect to fermentation time.
Biogas production (P-I)

At the end of the experiment (day 56), the composition of the gas in the bags being connected to the headspace of the reactors was analysed. The highest total biogas level was recorded with reactor R30 (16.13 L), followed by R40 (15.30 L) and R20 (14.98 L) (Fig. 5). Most gas (>90%) was produced between 8 and 16 days, thereafter, biogas production declined. Composition analysis of the total biogas revealed that the overall volumetric bio-H$_2$ production amounted to >8 L in all reactors, with the highest accumulation in reactor R20 (67%; day 4), followed by R30 (62%; day 8) and R40 (55%; day 8). The highest cumulative bio-H$_2$ production was recorded with R30 (9.31 L), followed by R20 (9.25 L) and R40 (8 L) (Fig. 5b). The comparatively lower production of bio-H$_2$ at a higher COD might be due to an overload of organic substrate in the system, which slowed down microbial metabolism, as lactose-rich CW could not be metabolised by all microbes. Assessment of the production profile with respect to fermentation time revealed that the largest volume of bio-H$_2$ was generated within 16 days (>95%), declining thereafter. Specifically, reactors R40, R20, and R30 accounted for 83%, 82%, and 79.61% of total volumetric bio-H$_2$ production, respectively. Bio-H$_2$ is generated preferentially during short retention times due to the rapid accumulation of microbial metabolites (mostly carboxylic acids) via acidogenesis. Despite low microbial metabolism after day 12, the reactors were operated for 56 days to further enhance conversion of substrate to carboxylic acids.

Bio-H$_2$ production depends on several operating conditions, such as substrate type/concentration, redox microenvironment, and nature of inoculum. Indeed, a pH of 5.7–6.0 favours acidogenic fermentation and, consequently, both bio-H$_2$ release and chain elongation. Even though the generation of bio-H$_2$ was much lower from day 13 to 28, it nevertheless amounted to almost 1.89 L (R30), 1.62 L (R20), and 1.33 L (R40) (Fig. 5a). As shown here one of the best ways to positively exploit the carbohydrate and protein content of CW and BSG as waste feedstocks, is through generation of bio-H$_2$ and soluble metabolites, such as SCCA and MCCA. When looked into the yields from its initial load of carbohydrate in the reactor, the highest value was found with R20 (216.46 mL H$_2$/gcarbohydrate) followed by R30 (194.66 mL H$_2$/gcarbohydrate) and R40 (146.70 mL H$_2$/gcarbohydrate). BSG co-fermented with CW was found to be an ideal feedstock for acidogenic bio-H$_2$ production due to its high organic load in the form of soluble carbohydrates.
Biogas production (gas recirculation: P-II)

Along with carboxylic acids, acidogenic fermentation cogeners CO$_2$ and H$_2$. In particular, almost 30% of the substrate is broken down to CO$_2$, this impacts carboxylic acid production during fermentation. The utilisation of this CO$_2$ could benefit the production of carboxylic acids. In this study, the biogas released during acidogenic fermentation was recirculated continuously at a flow rate of 80 mL h$^{-1}$. Gas consumption and composition were analysed periodically. Up until day 16, bio-H$_2$ content was 65–67% (GC-R$_{20}$), 57–58% (GC-R$_{30}$), and 52–53% (GC-R$_{40}$), corresponding to a volumetric biogas accumulation of 9.26 L, 9.05 L, and 7.90 L, respectively; whereas CO$_2$ accounted for 30–31% (GC-R$_{20}$), 38–39% (GC-R$_{30}$), and 40–44% (GC-R$_{40}$) of generated biogas. Consumption of both bio-H$_2$ and CO$_2$ was observed after day 16 in all reactors and, by day 24, the proportion of bio-H$_2$ declined slightly to 62% (GC-R$_{30}$), 54% (GC-R$_{30}$), and 46% (GC-R$_{40}$). After day 32, both bio-H$_2$ and CO$_2$ exhibited a significant decline. Eventually, by day 56, bio-H$_2$ content was only 44% (GC-R$_{30}$), 39% (GC-R$_{30}$), and 27% (GC-R$_{40}$); while CO$_2$ amounted to 53% (GC-R$_{30}$), 29% (GC-R$_{30}$), and 33% (GC-R$_{40}$) of biogas. These values indicated significant consumption of H$_2$ and CO$_2$ during recirculation in the reactor.

The availability of H$_2$ and CO$_2$ in the reactor favours homoacetogens, which can grow both autotrophically on H$_2$ and CO$_2$ as the terminal electron acceptor as well as carbon source for biomass production and cell maintenance and, therefore, does not impose a loss of carbon in acetogens. Arslan et al. (2012) reported increased carboxylic acids production by a mixed culture when the reactor headspace was supplemented with H$_2$ and CO$_2$ at a pressure of 2 bar. CO$_2$ released during fermentation is consumed again and converted to acetic acid during acetogenic fermentation. Because 5% to 10% of the reducing equivalents required for fixing the evolved CO$_2$ are used to sustain microbial growth, complete CO$_2$ recycling is not energetically possible without external energy supplementation. During recirculation, bio-H$_2$ serves as electron donor to provide the energy required for biomass production and cell maintenance and, therefore, does not impose a loss of carbon in acetogens. Upon consumption, the yields observed here by the end of the experiment from the initial load of carbohydrate was 101.36 mL H$_2$/g carbohydrate followed by 73.90 mL H$_2$/g carbohydrate and 37.5 mL H$_2$/g carbohydrate with GC-R$_{10}$, GC-R$_{40}$, and GC-R$_{40}$ respectively from its initial yield of 202.46 mL H$_2$/g carbohydrate, 173.47 mL H$_2$/g carbohydrate and 135 mL H$_2$/g carbohydrate respectively noticed during initial phases (8–16 days) of fermentation. With recirculation, the H$_2$ and CO$_2$ in the total biogas was consumed gradually resulted with its decreased volume with time. By the end of the experiment (day 56), with consumption of 3.71 L and 3.4 L of H$_2$ and CO$_2$ respectively, GC-R$_{40}$ was found to be more efficient suggesting its utilization towards formation of microbial metabolites. On the other side, its consumption with GC-R$_{30}$ (5.19 L: H$_2$: 3.33 L: CO$_2$) and GC-R$_{40}$ (4.63 L: H$_2$: 2.21 L: CO$_2$) was slightly less compared to GC-R$_{40}$ (Fig. 5d). No CH$_4$ was detected throughout the process due to suppression of methanogens following addition of 2-bromoethanesulfonic acid.

Carboxylic acid production during gas recirculation (P-II): adding value to CO$_2$ from acidogenic fermentation

Biogas produced during fermentation was recirculated to evaluate the effect of H$_2$ and CO$_2$ on carboxylic acid production. This strategy significantly enhanced the output of carboxylic acids compared to P-I (Fig. 6). By the end of the cycle (day 56), the reactor with the lowest COD load in CW (GC-R$_{30}$) showed a 21% increment in SCCA + MCCA, followed by 12.29% (GC-R$_{30}$) and 11.75% (GC-R$_{40}$) (Fig. 2b). Compared to P-I, H$_{Ac}$ biosynthesis was 9.55% (GC-R$_{30}$) higher in P-II on day 24, followed by 5.09% (GC-R$_{40}$) and 2.02% (GC-R$_{30}$) on day 32 (Fig. 6a). The accumulated H$_{Ac}$ was completely consumed over time.

In case of H$_{Ac}$, production increased gradually almost from the start (day 4) in all reactors and displayed a significant increment compared to reactors operated in P-I (Fig. 6b). Specifically, an additional production of 2.47 g L$^{-1}$ (GC-R$_{40}$), 1.38 g L$^{-1}$ (GC-R$_{30}$), and 0.89 g L$^{-1}$ (GC-R$_{20}$) in P-II meant that H$_{Ac}$ reached a maximum of 9.87 g L$^{-1}$, 9.76 g L$^{-1}$, and 8.79 g L$^{-1}$, respectively, which was 25%, 14.1%, and 10.12% higher than in P-I. The enhanced H$_{Ac}$ generated in this phase can be attributed to homoacetogens converting CO$_2$ to H$_{Ac}$ in the presence of H$_2$ as electron donor, confirming the impact of gas recirculation on homoacetogens enrichment. Gas recirculation had no major effect on H$_{Pr}$, which remained relatively low and showed a sustained increase only in reactor GC-R$_{40}$ (Fig. 6c).

Chain elongation of H$_{Ac}$ in the presence of an electron donor (H$_2$) led also to enhanced biosynthesis of H$_{Bu}$ (Fig. 6d). H$_{Bu}$ accumulation in P-II was higher with GC-R$_{40}$ (11.22 g L$^{-1}$) compared to GC-R$_{30}$ (7.46 g L$^{-1}$) or GC-R$_{20}$ (6.86 g L$^{-1}$), resulting in 11.2%, 25.9%, and 40% greater H$_{Bu}$ values with respect to P-I. The relatively lower H$_{Bu}$ concentrations recorded with GC-R$_{30}$ and GC-R$_{40}$ over GC-R$_{20}$, despite their higher organic loads, might be explained by H$_{Bu}$ chain elongation to other products. Gas recirculation potentially provides fermenting media with electron donors, which steer the direction and rate of fermentation towards specific products. Greater quantities of H$_{Bu}$ formed in the reactors might accrue from two possible routes: (i) elongation of H$_{Ac}$ to H$_{Bu}$ utilising H$_2$, or (ii) direct reduction of CO$_2$ and H$_2$. Zhou et al. (2017) found a 68.2% increment in carboxylic acids production by sparging H$_2$: CO$_2$ (80 : 20).

H$_{val}$ biosynthesis was also favoured by gas recirculation (Fig. 6e). Indeed, H$_{val}$ production was anticipated from day 32 in P-I to day 24 in P-II, and was accompanied by an overall accumulation of 1.85 g L$^{-1}$ (GC-R$_{40}$), 1.51 g L$^{-1}$ (GC-R$_{30}$), and 0.97 g L$^{-1}$ (GC-R$_{20}$). These values corresponded to an increase of 37.7%, 11.3%, and 9.37%, respectively, compared to P-I.

Production and consumption rate of carboxylic acids in P-II

In P-II, both $P_{rate}$ and $C_{rate}$ of carboxylic acids were relatively higher than P-I. $P_{rate}$ of H$_{Ac}$ was 1.12 (GC-R$_{20}$), 1.11 (GC-R$_{30}$) and 2.21 (GC-R$_{40}$) times higher with gas recirculation strategy compared to non-gas circulated reactors (Fig. 7). While the $C_{rate}$ of H$_{Ac}$ was higher with GC-R$_{40}$ (~0.17 g COD per L per day).
(Fig. 7b). In case of HPr, its \( P_{\text{rate}} \) was greater in P-I (0.66–4.75 times higher than P-II), whereas its \( C_{\text{rate}} \) was 1.1–6.27 times higher specifically with GC-R20 and GC-R30 in P-II over P-I indicating its possible conversion to HVal. Previous studies reported the concepts and possible routes involved in elongation of HPr to HVal by chain elongating microbes. On the other side, consumption of HPr was zero in GC-R40 both in P-I and P-II, indicating its continuous production with a \( P_{\text{rate}} \) ranging between 0.04–0.17 g COD per L per day in the reactor (Fig. 7c). While, the \( P_{\text{rate}} \) pattern of HBu between R20 and GC-R20 was more or less similar until day 40. However, its consumption between days 48–56, \( (C_{\text{rate}}: -0.05 \text{ to } -0.17 \text{ g COD per L per day}) \) declined its value to 7.48 g COD per L in R20. While an uninterrupted production in GC-R20 from day 48 to 56 (\( P_{\text{rate}}: 0.14 \text{ to } 0.21 \text{ g COD per L per day}) resulted with net HBu accumulation of 12.49 g COD per L. The reactors loaded with 30 g COD per L, showed a similar trend of HBu \( P_{\text{rate}} \) as observed with 30 g COD per L in P-I. Here the \( P_{\text{rate}} \) was almost similar with R30 and GC-R30 till day 40, which further decreased in R30 from day 48 (\( P_{\text{rate}}: 0.05–0.23 \text{ g COD per L per day}) whereas its consumption was noticed with GC-R30 on day 56. When the CW load was 40 g COD per L, the maximum HBu \( P_{\text{rate}} \) was noticed on day 24 (1.14–1.15 g COD per L per day). Later from day 32 to 48, its \( P_{\text{rate}} \) ranged between 0.22 to 0.25 g COD per L per day, which increased its

**Fig. 6** Influence of biogas recirculation on the composition of carboxylic acids (a–f) observed during P-II operation.
production to 20.42 g COD per L in GC-R40. Whereas the final production was limited to 18 g COD per L due to a lower $P_{rate}$ (0.02 to 0.11 g COD per L per day) during the same course of fermentation time in R40. An enhanced productivity of $H_{Ca}$ with GC-R20, R30, R40 might be due to direct reduction of CO$_2$ and H$_2$ facilitated through gas recirculation (Fig. 7d). Production of $H_{Val}$ was slightly higher in the reactor loaded 40 g COD per L of CW (R40 and GC-R40) with a $P_{rate}$ ranging between 0.04-0.19 g COD per L/day followed by 30 g COD per L (R30 and GC-R30; 0.02-0.16 g COD per L per day) and 20 g COD/L (R20 and GC-R20; 0.03-0.1 g COD per L per day) (Fig. 7e). The $P_{rate}$ of $H_{Ca}$ was varied in all the reactors at different organic load of CW, which influenced its accumulation. With 20 g COD per L load, the maximum $P_{rate}$ was recorded on day 40 (0.29 g COD per L per day) in GC-R30 which was 5.75 times higher over R20, whereas the maximum $P_{rate}$ (0.55 g COD per L per day) with GC-R30 was noticed from day 48-56 which was 1.5 times higher than R30. The $P_{rate}$ with GC-R40 was maximum between day 24–32 (0.63 g COD per L per day) led with accumulation with highest $H_{Ca}$ production (13.02 g COD per L) among all the reactors (Fig. 7f). The $P_{rate}$ was 1.46-2.68 times higher than R40. Chain elongated $H_{Ca}$ production through the reverse β-oxidation pathway was significantly higher with function of gas recirculation. This can be attributed the availability of electron either in the form of lactic acid/bioH$_2$ through continuous recirculation as its formation requires electron donor. Additionally CO$_2$ can be reduced to one mole of $H_{Ca}$ which requires 32 electrons while only 8 electrons are needed to convert CO$_2$ to acetate (eqn (4) and (5)).

$$6\text{HCO}_3^- + 37\text{H}^+ + 32\text{e}^- \rightarrow \text{CH}_3(\text{CH}_2)_4\text{COO}^- + 16\text{H}_2\text{O} \quad (4)$$

Chain elongation in phase-I (P-I)

$H_{Ca}$ was detected first on day 24 at 0.56-0.57 g L$^{-1}$ (Fig. 3f), but was almost double by day 40 in all reactors, with the highest value (2.71 g L$^{-1}$) recorded with R40, followed by 1.49 g L$^{-1}$ (R30) and 0.84 g L$^{-1}$ (R20). $H_{Ca}$ production was accompanied by simultaneous $H_{Lac}$ consumption (Fig. 3a), indicating that the latter was used as an electron donor in the chain elongation process. The highest $H_{Lac}$ production and consumption rates were observed in reactor R40 (Fig. 3a). For all reactors, the highest $H_{Lac}$ production rate was observed on day 16 with R40 (0.68 g L$^{-1}$ per day), followed by R30 (0.33 g L$^{-1}$ per day) and R20 (0.31 g L$^{-1}$ per day). Consumption of accumulated $H_{Lac}$ started on day 24, particularly with R20 (−0.18 g L$^{-1}$ per day), while production was still positive for R30 (0.32 g L$^{-1}$ per day) and R40 (0.22 g L$^{-1}$ per day) (Fig. 3f). This difference in $H_{Lac}$ conversion between individual reactors highlights the significant role played by the initial concentration of lactose in the medium. Several studies have documented $H_{Ca}$ production from ethanol-containing substrate; however, chain elongation using real field waste/wastewater as substrate requires addition of an external electron donor. In this study, $H_{Ca}$ produced from lactose-containing CW by mixed culture fermentation aided in the formation of $H_{Ca}$. From day 32, the consumption of $H_{Lac}$ increased in all reactors at a rate of −0.56 g L$^{-1}$ per day (R40), −0.33 g L$^{-1}$ per day (R30), and −0.2 g L$^{-1}$ per day (R20). Finally, by day 56, $H_{Lac}$ was fully consumed (Fig. 3f). At this point, the highest fraction of $H_{Ca}$ was displayed by R40 (4.68 g L$^{-1}$), followed by R30 (3.61 g L$^{-1}$) and R20 (2.66 g L$^{-1}$) (Fig. 3f).

$$2\text{HCO}_3^- + 9\text{H}^+ + 8\text{e}^- \rightarrow \text{C}_2\text{H}_3\text{O}_2^- + 4\text{H}_2\text{O} \quad (5)$$

![Fig. 7](image-url) Distribution of carboxylic acids production ($P_{rate}$) and consumption rate ($C_{rate}$) pattern in P-II (a–f) with respect to fermentation time.
H_{lac} is thought to act as an electron donor also in the acrylate pathway for the production of H_{pr}, rather than to generate H_{ca} via reverse β-oxidation. Such phenomenon was not observed in the present study, as indicated by a rather stable concentration of H_{pr} throughout the experimental period (Fig. 3c). Besides lactic acid and ethanol, sugars can also donate electrons during microbial chain elongation, leading to H_{ca} production via two-carbon increments. Recent studies suggest that H_{lac} plays an important role in chain elongation of SCCA to MCCA. However, mixed culture fermentation of complex substrates results also in the intermediates such as ethanol, making it difficult to determine the exact role of each molecule in chain elongation. Because in this study ethanol production was only 0.3–0.5 g L^{-1}, it likely played only a minor role in the process. Therefore, we believe that MCCA production was achieved mostly through lactic acid utilisation.

### Chain elongation during gas recirculation in phase-II (P-II)

The availability of H_{lac} along with continuous supplementation of an electron donor through gas recirculation in the fermenting medium not only improved accumulation of the SCCA mixture in the reactor, but also enhanced the biosynthesis of H_{ca}. The concentration of H_{ca} in P-II reached 5.25 g L^{-1} (GC-R_{30}), followed by 4.73 g L^{-1} (GC-R_{30}) and 3.62 g L^{-1} (GC-R_{20}) (Fig. 6f), which accounted for an increment of 10.85%, 23.67%, and 26.51%, respectively, over P-I (Fig. 3f). This observation was in line with the report by Shuai et al. (2019), who observed better chain elongation from mixed acids compared to pure H_{ac} due to the beneficial presence of H_{pr} + H_{lu} in the fermenting medium. An enhanced H_{ca} production could be related also to chain elongation in the presence of both H_{lac} as electron donor and homoacetogenic bacteria as a source of sufficient substrate for the process. Both production and consumption of H_{lac} were influenced by gas recirculation, with the former being slightly higher than in P-I (Fig. 6a). Maximum production rate of H_{lac} was achieved by GC-R_{40} (0.75 g L^{-1} per day), followed by GC-R_{30} (0.43 g L^{-1} per day) and GC-R_{20} (0.35 g L^{-1} per day) on day 16. The consumption rate of H_{lac} was also greater, particularly with GC-R_{30} (~0.35 g L^{-1} per day) on day 32, followed by GC-R_{40} (~0.95 g L^{-1} per day) and GC-R_{20} (~0.66 g L^{-1} per day) on day 40. This result suggested a key role for lactose-derived H_{lac} as an intermediate during reverse β-oxidation and consequent chain elongation. The latter was further promoted by the presence of an additional electron donor in the form of bio-H_{2} during P-II. Indeed, after complete consumption of H_{lac}, the recirculating gas in the reactor played an important role in converting CO_{2} to metabolites. By day 40, as the accumulated H_{lac} in the fermenting media was completely utilized, at this point the production of chain elongated carboxylic acids can be attributed to the availability of H_{2} in the reactor through its recirculation acting as an electron donor, as the H_{2} consumption during this course of time was 2.01 L (GC-R_{30}) followed by 1.9 L (GC-R_{20}) and 1.61 L (GC-R_{20}). At the same time, during reverse β-oxidation, acetate is elongated to butyrate via acetyl-CoA and then butyrate is elongated to caproate via butyl-CoA. Zhang et al. (2013) generated a mixture of carboxylic acids (H_{ac} + H_{bu} + H_{ca}) when fermenting a gas composed of CO_{2} (40%) and H_{2} (60%) in a hollow-fibre membrane biolim reactor containing a mixed culture. The COD equivalent of the carboxylic acids produced from co-fermenting BSG and cheese whey wastewater in two different phases is presented in Table 2.

### Conclusions

The present study shows that complementing two different waste streams, such as CW and BSG as co-fermenting substrates, increases the output of carboxylic acids and bio-H_{2}. Lactic acid biosynthesis is dependent on the initial load of CW. A higher load of CW (40 g COD per L) maximised the production of SCCA to 0.38 g L^{-1} per day, which was about 1.2 to 1.6 times higher than using 20 or 30 g COD per L, respectively. Bio-H_{2} recovery was maximal with 30 g COD per L. Importantly,
gas recirculation allowed acidogenic CO$_2$ to be converted to SCCA at a much higher rate (19.9%) compared to the case without gas recirculation. MCCA production, which requires an electron donor, correlated with the consumption of lactic acid, indicating a lactic acid-based chain elongation process. Finally, the enhanced production of MCCA such as caproic acid, after complete utilisation of lactic acid in the fermenting medium, highlights the benefit of gas recirculation as a source of H$_2$ and its role as a key electron donor.

Author contributions

Omprakash Sarkar: conceptualization, methodology, investigation, writing – original draft; Ulrika Rova: conceptualization, writing – review & editing; Paul Christakopoulos: conceptualization, writing – review & editing; Leonidas Matsakas: conceptualization, methodology, writing – review & editing, supervision.

Conflicts of interest

There are no conflicts to declare.

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