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Effect of total solids content on biohydrogen production and lactic acid accumulation

2	during dark fermentation of organic waste biomass
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14 15 16 17 18 19 20 21 22 23	*Corresponding author. Present address: Nepal Engineering College, NEC- Center for Postgraduate Studies, G.P.O. Box: 10210, Kathmandu, Nepal. Tel: +977 01 5221006, Fax: +977 01 015221001 <i>E-mail address</i> : <u>anishghimire@gmail.com</u> (Anish Ghimire)

- 27 28 Abstract

28 29	Production of biohydrogen and related metabolic by-products was investigated in Solid		
30	State Dark Fermentation (SSDF) of food waste (FW) and wheat straw (WS). The effect of		
31	the total solids (TS) content and H ₂ partial pressure (pp _{H2}), two of the main operating		
32	factors of SSDF, were investigated. Batch tests with FW at 10, 15, 20, 25 and 30 % TS		
33	showed considerable effects of the TS on metabolites distribution. H_2 production was		
34	strongly inhibited for TS contents higher than 15% with a concomitant accumulation of		
35	lactic acid and a decrease in substrate conversion. Varying the pp_{H2} had no significant		
36	effect on the conversion products and overall degradation of FW and WS, suggesting that		
37	pp_{H2} was not the main limiting factor in SSDF. This study showed that the conversion of		
38	complex substrates by SSDF depends on the substrate type and is limited by the TS		
39	content.		
40			
41	Keywords: Biohydrogen; Lactic acid; Dark fermentation; Total solids; Food waste		
42			
43	Highlights		
44	• H ₂ and by-products production were investigated in solid state DF of food waste.		
45	• Effects of TS content and H ₂ partial pressure on substrate conversion were studied.		
46	• 15% TS was found to be limiting for H ₂ production during solid state dark		
47	fermentation.		
48 49			

50 1. Introduction

51 Interest in Dark Fermentation (DF) bioprocesses has increased over the past decade because 52 it has shown the potential to produce 'green' hydrogen (H₂) from widely available and low-53 cost organic biomass with a low environmental impact (Azwar et al., 2014). Renewable 54 waste biomass such as agricultural residues, organic fraction of municipal solid waste 55 (OFMSW) and agro-industrial waste might give an additional competitive economic 56 advantage to industrial units treating waste feedstocks by anaerobic digestion for local 57 supply of H_2 (De Gioannis et al., 2013; Guo et al., 2010; Urbaniec and Bakker, 2015). 58 OFMSW and lignocellulosic residues such as wheat and rice straw are potential substrates, 59 as they are abundant and do not compete with food crops as for the production of first 60 generation biofuels (Ghimire et al., 2015a; Guo et al., 2010). 61 Based on the total solids (TS) content of the digestion mixture, anaerobic fermentation 62 processes such as anaerobic digestion (AD) processes can be categorized as: i) wet AD, 63 which operates at low solids content (<15% TS) and ii) dry AD at high solids content 64 (>15% TS) (Motte et al., 2013). Alternatively, AD processes are categorized as: (i) wet 65 (<10% TS), (ii) semi-dry (10–20% TS) and (iii) dry (>20% TS) (Karthikeyan and 66 Visvanathan, 2012). Dry anaerobic fermentation processes can be operated at high substrate 67 concentrations with little water addition, which offers the advantages such as of: (i) 68 economic benefits by reducing reactor volumes and energy consumption for heating the 69 digester, (ii) more efficient handling of the digestate by reducing the drying post-treatment 70 operation and (iii) a higher simpler technicity (Karthikeyan and Visvanathan, 2012; Motte 71 et al., 2013). Dry AD processes have been successfully applied to many different 72 agricultural feedstocks (Liotta et al., 2015; Mussoline et al., 2012).

73	Dry AD anaerobic fermentation processes, in particularly dry DF, can also be attractive for
74	commercialization of bio-H $_2$ processes for the same reasons. Dry DF which operates at a
75	TS content higher than 15% (sometimes also defined as > 20% TS), is also known as Solid
76	State Dark Fermentation (SSDF). SSDF has recently been reported as suitable for
77	biohydrogen production from wheat straw (WS) (Motte et al., 2015, 2014). In their study,
78	Motte et al. (2015) proposed to combine SSDF with a mechanical pre-treatment process to
79	enhance the overall conversion of lignocellulosic biomass, which makes it more feasible for
80	upscaling in a biorefinery concept.
81	However increasing the TS content during dry anaerobic fermentation can impact both the
01	However, increasing the 15 content during dry anaerobic refinentation can impact both the
82	total substrate degradation and the biogas production (Abbassi-Guendouz et al., 2012;
83	Motte et al., 2013). At high $(19 - 25 \%)$ TS content, mass and energy transfer are lowered
84	by the low water content (Motte et al., 2014; Valdez-Vazquez & Poggi-Varaldo, 2009).
85	These can lead to a local accumulation of by-products up to inhibitory concentrations or
86	result in decreased metabolic conversion rates. Fernández et al. (2008) reported 17%
87	decrease of the degradation rates in dry AD of OFMSW, when the TS content increased
88	from 20 to 30%. Cazier et al. (2015) showed that high hydrogen partial pressures (pp_{H2})
89	inhibited wheat straw WS hydrolysis in dry AD, suggesting that local accumulation of
90	metabolic intermediates adversely impacted the overall conversion of the biological
91	process. More particularly, Motte et al. (2013) reported a significant decrease in H_2
92	production along with a sharp decrease of WS conversion when performing SSDF at 19 $\%$
93	TS. Moreover, a reduction of the H_2 yields and shifts in the microbial community
94	composition towards lactic acid metabolism was observed at high TS content (Motte et al.,

95	2014). In contrast, only few studies have addressed the effect of increasing the TS content
96	on DF of food waste (Valdez-Vazquez and Poggi-Varaldo, 2009).
97	Therefore, this study aims to investigate the effect of the TS content on organic waste
98	conversion in SSDF using food waste (FW) and Wheat Straw as high and low
99	biodegradable substrate, respectively. In addition, the effect of the pp_{H2} on organic waste
100	hydrolysis and metabolic pathways was evaluated using SSDF batch tests.
101	
102	2. Material and Methods
103	2.1 Substrate feedstock
104	Batch SSDF experiments were carried out with synthetic FW (20.6% TS and 19.7% VS),
105	freshly prepared with a well-defined composition as described elsewhere (Ghimire et al.,
106	2015b) and similar to the composition of FW in European countries reported by
107	VALORGAS (2010). WS with a TS and VS content of 95% and 97%, respectively, was
108	used as a model lignocellulosic biomass substrate.
109	
110	2.2 Source of inoculum
111	Waste activated sludge was used as inoculum after heat-shock treatment (90 $^{\circ}$ C, 15 min) to
112	enrich spore forming Clostridium sp. and inhibit methanogenic and homoacetogenic
113	activities that consume H_2 (Ghimire et al., 2015a). It was sampled from the secondary
114	clarification unit of a municipal wastewater treatment plant located in the center of France
115	with a capacity of 285,000 population equivalents. The waste activated sludge was
116	centrifuged at 6,500 rpm for 20 min (4 $^{\circ}$ C) to reach 13.9% TS and 9.7% VS content for
117	achieving a higher TS content during all experiments.
110	

119 2.3 Experimental set-up

120 **2.3.1 Effect of TS content on H**₂ production and substrate conversion

- 121 Batch SSDF tests were carried out in triplicates at 10%, 15%, 20%, 25% and 30% TS
- 122 content of FW. In 600 ml flasks, 53.4 g of an inoculated biomass mixture composed of FW
- 123 (20.3 g wet weight with 19.7% VS) and inoculum (4.1 g wet weight with 9.7% VS) with a
- 124 substrate (S) to inoculum (X) ratio (S/X) of 10 g VS substrate/g VS inoculum were mixed
- 125 with 16.0 g of 2-(N-morpholino) ethanesulfonic acid (MES) buffer, 12 ml of 3.2 % NaOH
- and 1 ml of trace metal solution containing (g/L): FeCl₂ 2, CoCl₂ 0.5, MnCl₂ 0.1, NiCl₂
- 127 0.1 , ZnCl₂ 0.05 , H₃BO₃ 0.05 , Na₂SeO₃ 0.05 , CuCl₂ 0.04 and Na₂MoO₄ 0.01 . This

128 mixture had an initial pH of 5.5. The amount of distillated water to be added was calculated

129 with a mass balance on the TS content, including substrate, inoculum, buffer and the other

- 130 solutions added to obtain final TS contents of 10.0 (\pm 0.01), 14.98 (\pm 0.03), 19.89 (\pm 0.04),
- 131 24.92 (± 0.02) and 30.0 (± 0.07) %TS. Batch tests were then incubated at 37 (± 1) °C for 14
 132 days.

133 2.3.2 Effect of partial pressure of H₂ (pp_{H2}) on substrate conversion

134 The study on the effect of the partial pressure of H₂ (pp_{H2}) was carried out in four replicated

batch tests operated with FW or WS at a S/X ratio of 10 and a final TS content of 25 (± 1)

136 %. A final TS of 25% was selected to ensure the effect of a high TS content on substrate

137 conversion. Tests were carried out with a thin layer of digestate (< 1 cm), approximately 22

- 138 (± 2) g, to minimize the effect of gas diffusion (Cazier et al., 2015). H₂ gas was initially
- added in the headspace of 600 ml serum bottles according to the two following sets of tests:
- 140 pp_{H2} equivalent to 542 (\pm 32) mbars (33 \pm 2 % H₂ in the headspace, named as "A") and
- 141 1087 (\pm 29) mbars (66 \pm 1 % H₂ in the headspace, named as "B"). A control was carried

out with only N₂ in the headspace (named as "C"). In all tests, the initial total pressure was set at 1,500 mbars by adding N₂ at the start of the experiments for all conditions. The initial pH was adjusted at 5.5 using the MES buffer. Batch tests were incubated at mesophilic temperature $(37 \pm 1 \text{ °C})$ for two fermentation periods of 14 and 21 days each.

146

147 **2.4 Analytical methods**

148 Prior to liquid metabolite measurements, 5 g of digestate was diluted in 5 g of deionized

149 water, mixed during 30 minutes, centrifuged at 39,121g during 20 min at 4°C and then

150 filtrated at 0.2 µm with a nylon membrane. The remaining liquid phase was then used to

151 quantify organic acids, other metabolic end-products and soluble sugars. Dark fermentation

152 metabolites were measured at the beginning and the end of each experiment.

153 Organic acids were quantified with a gas chromatograph (Perkin Clarus 580) using an Elite

154 - FFAP crossbond® carbowax® 15 m column connected to a flame ionization detector at

155 280°C, as described elsewhere (Cazier et al., 2015). N₂ was used as carrier gas at a flow

156 rate of 6 mL/min. High performance liquid chromatography (HPLC) was used to quantify

157 other metabolites, e.g. lactic acid, alcohols, and soluble sugars. The HPLC chain was

158 composed of an Aminex HPX-87H column (300 mm on 7.8 mm, Bio-rad), a pre-column to

159 filter residues (Micro guard cation H refill cartbridges, Bio-rad) and an automatic sampler

160 (Water 717). A diluted sulfuric acid solution of 0.005 M in milliQ water was used as eluent

161 at a flow rate of 0.4 ml/min.

162 The gas composition was measured with a gas chromatograph (Perkin Clarus 580)

163 equipped with a thermal conductivity detector at 150°C and an injector heated at 250°C and

164 two capillary columns heated at 60°C. The first column corresponded to a RtU bond for

165 CO₂ analysis, while the second column was a RtMolsieve used for the determination of the

166	O_2 , H_2 , N_2 and CH_4 concentration as described elsewhere (Cazier et al., 2015). Argon at a
167	pressure of 350 kPa and a flow rate of 31.8 ml/min was used as carrier gas. Gas production
168	was monitored through the increase in gas pressure, which was periodically measured with
169	a digital manometer (2000, Leo2 Keller, Winterthur, Switzerland).
170	
171	2.5 Microbial diversity analysis
172	For microbial analysis, $0.2 - 0.3$ g of solid samples were collected at the end of the
173	fermentation period of 14 days from the SSDF batch tests operated at different pp_{H2} . PCR
174	based pyrosequencing, described elsewhere by Carrillo-Reyes et al. (2016), was carried out
175	in order to elucidate the classification of the microorganisms present in the SSDF. The
176	samples for microbial analysis were taken at different pp_{H2} during SSDF of FW and WS.
177	
178	2.6 Data analysis
179	Substrate degradation was estimated by calculating a theoretical chemical oxygen demand
180	(COD) mass balance. For this, the difference in metabolic end-products (accumulated in
181	both the gaseous and liquid phase) at the initial and final state were compared. Direct
182	measurement of COD was not considered in this study, as suggested by Cazier et al.
183	(2015), since the COD measurements of the complex organic residues, such as
184	lignocellulosic biomass and FW, may vary more than 10 %, which may be higher than the
185	overall substrate degradation during the SSDF process.
186	Total substrate degradation was estimated from the amount of COD produced during DF of
187	the substrate estimated per kg of TS added initially, and corresponded to:
188	Total Substrate Degradation = COD of Final State – COD of Initial State

189
$$Total Substrate Degradation = \frac{A_{H_2,f} + A_{met,f} + A_{GC}}{kg TS} - \frac{A_{H_2,i} + A_{met,i}}{kg TS}$$

Where: $A_{H_2,f}$ is the amount of H₂ remaining at the end in the headspace, $A_{met,f}$ the final 190 amount of metabolites accumulated, A_{GC} the total amount of gas (H₂) sampled for analyses, 191 192 A_{H2,i} the initial amount of H2 added and A_{met.i} the initial amount of metabolites in the 193 medium. 194 R (OSX version 3.1.3) with the package Rcmdr (OSX version 2.1.7) was used for statistical 195 analysis of the experimental data. The P value was set at 0.05 and the significance of the results was tested with P values of: * < 0.05; ** < 0.01; *** < 0.001; while results were not 196 197 significantly different when P > 0.05. A multivariate analysis by Principal Components was 198 carried out using FactoMineR (version 1.24, more on http://factominer.free.fr/), an 199 extension of the R software, to correlate the H_2 production with the substrate degradation 200 and formation of other metabolic by-products (Lê et al., 2008).

201

202 **3. Results and discussion**

203 **3.1 Effect of TS content on H**₂ production and substrate conversion

204 SSDF tests were carried out with FW at different TS contents (Fig. 1). Overall, the H₂

205 production was strongly impacted by increasing the initial TS content, which is in

accordance with earlier studies (Motte et al., 2014; Valdez-Vazquez & Poggi-Varaldo,

207 2009). Fig. 1a shows the FW degradation after 14 days of incubation as a function of the

- 208 TS content. Fig. 1b presents the molar yield (mmol/kg TS) of the major metabolic end
- 209 products. Substrate conversion decreased and biohydrogen production was significantly
- 210 lowered when the initial TS content increased (Fig.1b). A simultaneous shift of the
- 211 metabolic pathways was observed at a high TS content. The maximum and minimum

212	substrate degradation of 134 (\pm 22) and 51 (\pm 3) g COD/kg TS was achieved at 10 and 30
213	% TS, respectively. H ₂ accumulated in the headspace only at a TS content of 10%.
214	Hydrogen production drastically decreased and lactic acid production was favored at TS
215	contents higher than 10%. Indeed, from 15% TS onwards, the metabolic pathways shifted
216	towards lactic acid conversion, thus explaining the decrease in H ₂ production (Fig. 1b).
217	The highest substrate conversion of 2,901 (\pm 143) mM metabolites/kg TS was observed at
218	wet conditions of (10% TS) (Fig. 1b), while it was the lowest (1,435 \pm 13 mM
219	metabolites/kg TS) at 30% TS (Fig. 1b). As shown in Fig. 1a and b, such adverse effect of
220	the TS content started between 10 and 15%. Motte et al. (2014) showed similar decrease of
221	the hydrogen yields, related to a metabolic shift with an accumulation of lactic acid, but at
222	higher TS content (19%) for wheat straw only. Further narrowing the study of the TS
223	content within the range $10 - 15$ % TS or considering the soluble carbohydrates available to
224	microbial degradation could further give new insights on the mechanisms behind this
225	limiting effect of the TS content in SSDF.
226	In addition, acetate and butyrate were the two of the major as metabolic by-products in at a
227	TS content of 10%, while lactate dominated at higher TS contents. The PCA correlation
228	plot (Fig. 2a) shows that H ₂ production correlated only with butyrate production,
229	suggesting that acetate resulted from DF and homoacetogenesis pathways. Theoretically,
230	the presence of acetate and butyrate as metabolic by-products is correlated with hydrogen
231	production pathways (Ghimire et al., 2015b). However, a correlation with butyrate
232	production as shown in this study (Fig. 2a), is consistent with previous observations made
233	by Guo et al. (2014) for solid waste where homoacetogenesis can occur. Interestingly, H_2
234	production was also linked to the total substrate degradation, while lactate production did
235	not correlate well with the substrate conversion but more with the initial TS content

236	suggesting an adverse effect of high TS content likely due to the local microbial
237	environment (Fig. 2b). Indeed, the decrease of substrate conversion at increasing TS might
238	be attributed to VFA accumulation, especially lactic acid, decreasing locally the culture pH
239	and inhibiting the hydrolysis process, even though a high amount of MES buffer was added
240	to the medium. Such reaction is very similar to the ensiling process where pH lower than
241	4.5 are favored to stop the microbial process (Kafle & Kim, 2013). Consistently, Lin et al.
242	(2006) reported that xylose removal decreased from 85% to 37% when the culture pH
243	decreased from 8 to 5. Similarly, Fang and Liu (2002) reported a 10% decrease in glucose
244	degradation when the pH decreased from 5.5 to 4. The low culture pH could also have had
245	an adverse effect on hydrolytic enzymes. Indeed, Veeken et al. (2000) reported a decrease
246	in hydrolysis of complex substrates with a decrease in pH, since hydrolytic enzymes are
247	optimally functional at neutral pH (Parawira et al., 2005).
248	Another explanation for the decrease in substrate degradation at higher TS content could be
249	the local accumulation of H ₂ , as reported and hypothesized by Cazier et al. (2015).
250	

3.2 Effect of ppH2 on substrate conversion 251

252 As potential inhibitory factor, the effect of the pp_{H2} on FW and WS degradation at a high

253 TS content (25 % TS) was investigated and different initial pp_{H2} were tested. Fig. 3a shows

254 the substrate degradation (expressed as g COD/kg TS) from FW after 14 and 21 days of DF

255 at different pp_{H2}. The level of inhibition of pp_{H2} on substrate hydrolysis was determined

- 256 based on the difference in total substrate degradation (Fig. 3a). No significant effect of the
- 257 initial pp_{H2} on biomass hydrolysis (ANOVA test, P-values > 0.05) was observed at 25%
- 258 TS. This was further evident from the metabolic products that accumulated at the end of the

259 experimental period, which showed no significant shift when changing pp_{H2} (Fig 3b).

Substrate degradation slightly increased with the fermentation time, and lactic acid and
ethanol accumulated as major metabolites in all tested pp_{H2} with FW.

262 Further tests were carried out with WS under similar experimental conditions with the

263 objective to confirm the results obtained with FW. Although the pH did not significantly

264 decrease at the end of the experimental period, Fig. 3 and the ANOVA analysis (P-values >

265 0.05) show that the pp_{H2} did not affect the DF of the substrate WS. However, the difference

in metabolites accumulated between 14 and 21 d suggests that hydrolysis occurred.

267 Therefore, it seems that some dark fermentative communities present in the SSDF tests

268 investigated with WS were not strongly affected by the presence of H₂. These results

269 contrast with the study of Cazier et al. (2015) who reported an inhibition by high ppH₂ of

WS hydrolysis in dry AD, at pH 7. In their work, inhibition was observed at a $pp_{H2} > 742$

mbars and degradation of wheat straw decreased from 90 (\pm 10) to 20 (\pm 10) g COD/kg TS

in the controls and at a pp_{H2} of 1,555 mbars, respectively. In contrast, in this study, no

273 evidence of the effect of pp_{H2} was observed, suggesting the inhibition of the substrate

274 conversion is mainly due to organic acids (especially lactate) accumulation. Although this

275 difference might be attributed to the lower operational pH in the present study (pH 3.7 -

5.5) compared to the dry AD (pH 7 - 8) process of Cazier et al. (2015), the WS degradation

in the control tests, i.e. 22.4 (± 2) g COD/kg TS (at pH 5.4), was interestingly very similar

to the one reported by Cazier et al. (2015), i.e. $20 (\pm 10)$ g COD/kg TS, when maximum

279 hydrolysis inhibition occurred at a pp_{H2} of 1,555 mbars at pH 7 - 8. Therefore, this

280 difference could be attributed to the fact that substrate degradation in this study was

strongly related to the high content in soluble carbohydrates present in the FW substrate, as

reported by Guo et al. (2013), but not to the substrate to be hydrolyzed such as WS.

- 283 Moreover, the pH might have affected microbial selection and the related metabolic
- 284 pathways. Lactic acid or ethanol were not present as major intermediates in the SSDF of
- 285 WS (Fig. 3c), in contrast to tests with FW (Fig. 3b). This suggests that the microorganisms
- 286 producing these metabolites are dominant in moisture limiting and nutrient rich
- environments such as FW, but not in WS (Sikora et al., 2013).
- 288 This is in good agreement with the characterization of the microbial communities. The
- 289 microbial community characterization (Fig. 4) shows that mainly 3 families represented the
- 290 most abundant OTUs, showing the community present in the SSDF residue after 14 d
- 291 incubation had a low diversity, although the number of distinct operational taxonomic units
- 292 (OTUs) was 1127 and 1877 for the SSDF samples of FW and WS, respectively.. The
- hydrogen (pp_{H2}) favored the development of a *Lactobacillales* population (Fig. 4a). The
- high abundance of *Lactobacillales* (Fig. 4) explains the high lactic acid concentration
- during SSDF of FW (Fig. 3c). Moreover, the high abundance of *Lactobacillales* at T_0
- during SSDF of FW (Fig. 4a) compared to WS (Fig. 4b) implies FW as their major source
- 297 of inoculum. On the other hand, during SSDF of WS, the Clostridiales were present
- abundantly, which agrees with the production of several other metabolites such as acetate,
- propionate and butyrate. This observation is also in good agreement with Motte et al.(2014)
- 300 who previously reported that high TS contents impacted the microbial composition through
- 301 the selection of lactic-acid bacteria from the *Bacilli* class.
- 302
- 303 The production of organic acids by fermentative conversion processes is commonly
- 304 significantly affected by the operating parameters such as culture pH, temperature and
- 305 substrate concentration (Wang et al., 2014, 2015). However, the highest conversion of FW
- 306 $(140.7 \pm 23 \text{ g COD/kg VS}_{added} \text{ at initial pH 5.5, 25 \% TS, 14 days SRT and 37 °C})$ in this

307	study is lower compared to the study by Wang et al. (2015) (799 g COD/kg VS _{added} at
308	controlled pH 6.0, 35 °C) and Wang et al. (2014) (918 g COD/kg VSremoved at controlled pH
309	6.0, 30 °C). This could explain the higher conversion of the waste biomass into
310	fermentative products reported in these two studies, compared to this study where the pH
311	was not controlled and the low pH conditions (3.7 - 5.1) finally inhibited the microbial
312	activity. Similarly, a comparison can be made between the study of Motte et al. (2015) who
313	obtained 6 fold higher organic acids production (140 \pm 6 g COD/kg TS $_{added}$ at 23 % TS, 37
314	°C) at an initial pH 8.5, as compared with this study (22 \pm 2 g COD/kg TS _{added} at 25 % TS,
315	37 °C) at initial pH 5.5. Therefore, it can be concluded that under SSDF, the substrate
316	conversion depends on the culture pH, TS content, substrate type and microbial
317	community.
318	The total substrate conversion under SSDF also depends on the substrate type. FW has a
319	high VS content (21 to 27% VS) and is particularly suitable for a concomitant production
320	of bio-hydrogen and platform molecules such as short-chain volatile fatty acids, lactic acid
321	and / or alcohols (Uçkun Kıran et al., 2015; VALORGAS, 2010). Soluble by-products of
322	DF can be further applied in biological processes for (i) wastewater treatment (Elefsiniotis
323	et al., 2004), (ii) production of platform molecules such as biopolymers (Ntaikou et al.,
324	2009), (iii) micro-algal lipids production (Turon et al., 2015), iv) H ₂ production by photo-
325	fermentation, v) feed for microbial electrolysis cells for production of H_2 and other value
326	added chemicals (El Mekawy et al., 2014) or vi) anaerobic digestion for a higher energy
327	recovery, in the form of hythane, i.e. both H ₂ and CH ₄ (Ghimire et al., 2015a).
328	
329	

330	5.	Conclusion

- 331 This study highlighted the effects of limiting parameters on substrate conversion during
- 332 SSDF of FW and WS. The initial TS content significantly affected the substrate
- degradation and metabolic by-products during SSDF of FW. The H₂ production ceased at a
- TS content higher than 15%, accompanied by the accumulation of lactic acid. Moreover,
- the pp_{H2} did not affect the SSDF of food waste due to lactic acid production lowering the
- culture pH, which played a major role in limiting the substrate conversion, as confirmed
- during SSDF of WS.
- 338

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- 343

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477	Figure	Cap	tions

478 Fig. 1. Effect of TS content on substrate degradation (in g COD per kg of initial TS) (a);
479 and accumulation of end metabolites accumulation (mM per kg of initial TS) (b); at
480 different TS content

482 Fig. 2. Principal component analysis correlation circle plots: (a) Hydrogen and major
483 metabolic by-products production and (b) substrate degradation and metabolic products
484

485 **Fig. 3.** Substrate degradation (in g COD per kg of initial TS) at 25% TS using FW and WS

486 (a) as well as end metabolites accumulation (mM per kg of initial TS) in SSDF of FW (b)

487 and WS (c) at different pp_{H2} and at the end of two fermentation times (14 d and 21 d). A =

488 542 (\pm 32) mbars, B= 1087 (\pm 30) mbars, C = 0 mbars, 14 = 14 d and 21 = 21 d

489

490 **Fig. 4.** Microbial community distribution at family level of the sequenced samples during

491 the SSDF using FW (a) and WS (b) at the beginning (To) and end of the 14 d incubation at

different pp_{H2} (0, 542 (± 32) mbars, and 1087 (± 30) mbars) considering most abundant

493 OTUs (only OTUs with a presence of >1.0% were considered)

Fig. 1.



497 Fig. 2.498







Fig. 3.





Fig. 4.



