

1 **Modelization of anaerobic processes during co-digestion of**
2 **slowly biodegradable substrates**

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23

24 **Abstract**

25 The influence of the soluble substrates over the anaerobic processes has been extensively
26 investigated, but little is known about the effects of particulate substrate. The biodegradation
27 of these substrates starts with the hydrolytic step, this process is slower than the other ones
28 involved in the biodegradation of particulate substrates and usually becomes the rate-limiting
29 step. This study investigate the effect of the initial total solids (TS) concentration on the
30 anaerobic co-digestion of two slowly biodegradable organic substrates. The wastes mixtures
31 were prepared at different dilutions in the range from 10% to 28% TS. From these experiments
32 it was observed that as TS concentration increased, the methane production decreased. These
33 results were modelled and it was observed that neither hydrolysis nor fermentation stages
34 controlled the methane production rate. Being a substrate inhibition event experienced at the
35 methanogenic stage the responsible of the lower methane production when operating at high
36 TS concentrations.

37

38 **Keywords:** Anaerobic co-digestion; hydrolysis rate; inhibition; particulate substrate.

39

40 1. INTRODUCTION

41

42 The anaerobic co-digestion can be defined as the simultaneous biological treatment of two, or
43 **more biodegradable wastes**. The combination of substrates with different compositions could
44 be used to enhance the biogas production due to the equilibration of the nutrients balance in
45 the mixture, mainly C/N ratio (Bohutskyi et al., 2018; Zahan et al., 2018), at the same time that
46 the particulate substrates as well as toxic **or inhibitory** compounds concentrations could be
47 diluted (Mata-Alvarez et al., 2000; Xie et al., 2016).

48 **In the anaerobic digestion process, the presence of particulate substrates is relevant for the**
49 **extension of the treatment due to the multiphase, multistage and sequential reactions**
50 **required for its final transformation**. The term hydrolysis define the breakdown of particulate
51 **substrates** into biodegradable soluble substrates (Henze et al., 1987). Before its

52 biodegradation, it is necessary to hydrolyse the particulate substrates (Levine et al., 1985). In
53 most of the cases, the rate of hydrolysis is much slower than that of the **consumption** of the
54 soluble substrates that it generates. **Because of that, usually the hydrolysis stage became** the
55 rate limiting step in the biological treatment of the particulate pollutants (Henze et al., 1987).

56 In the literature, it has been described that the hydrolytic processes play a dominant role in
57 the delicate balance of electron donor/electron acceptor ratio of several bio-processes (Gujer
58 and Zehnder, 1983; Levine et al., 1991; Rodríguez Mayor et al., 2004; de los Ángeles Fernandez
59 et al., 2016). In the literature, different experimental approaches have been used to evaluate
60 the anaerobic hydrolysis process. **Some generalised approaches cannot be used for accurate**
61 **estimation of the kinetic and stoichiometry of the anaerobic hydrolysis process. This is because**
62 **the different populations/wastes involved in the study leads to large uncertainty (Morgenroth**
63 **et al., 2002). In other cases, the studies of the hydrolysis involve the measurement of specific**
64 **hydrolytic enzymes, intermediates or end-products (Brethauer et al., 2011). However, caution**
65 **should be employed, when applying these results because slight changes in the wastes could**

66 yield different results. Because of that, methods based on real effluents seems more
67 convenient (Kouas et al., 2018).

68 Nowadays, the IWA Anaerobic Digestion Model No. 1 (ADM1) (Batstone et al., 2002) is one of
69 the most widely used for anaerobic digestion modelling. This model accurately describes the
70 stages taking place in the anaerobic digestion process, but presents a complex structure.
71 Because of its complex structure, this model requires a detailed substrate characterization,
72 which could be difficult to obtain, and the definition of the stoichiometric coefficients and
73 kinetics rates of a wide number of processes (Goel et al., 1998). The large number of
74 parameters make this model difficult to identify and may result in parameter correlation,
75 leading to a significant uncertainties. Currently, several simplified anaerobic digestion models
76 (Giovannini et al., 2018; Kouas et al., 2018; Kouas et al., 2019) adequately predict the process
77 in most of the cases avoiding the complexity of the ADM1. However, when the wastes present
78 components different to the conventional ones (Mata-Alvarez et al., 2000) or when the waste
79 presents high solids contents, these simplified models present limitations in its accuracy
80 (Morgenroth et al., 2002; Vavilin et al., 2008; Mao et al., 2019).

81 In this context, the aim of this work was to develop and validate a, mass balance based,
82 simplified model describing the hydrolysis and subsequent processes in anaerobic co-digestion
83 of two slowly biodegradable organic wastes, the 2POMW and the CM. The work pays special
84 attention to the effects of the initial solids concentration, covering the wet as well as the dry
85 anaerobic digestion, and to the hydrolytic stage on the operational evolution and biogas
86 production.

87

88 2. MATERIALS AND METHODS

89 2.1 *Experimental design*

90 The experiments were designed to evaluate the hydrolytic as well as the subsequent processes
91 of the anaerobic biodegradation of the solid wastes mixtures. Two organic wastes were used:
92 two phase olive mill waste (2POMW) and Cattle manure (CM). 2POMW was collected from an
93 olive oil mill (Cooperativa Nuestra Señora de los Remedios) located in Olvera, Cádiz (Spain). CM
94 was obtained from a semi-intensive livestock farm located in El Puerto de Santa Maria Cádiz
95 (Spain). Both substrates were homogenized and stored at -4°C to preserve its original
96 characteristics. The main physical–chemical characteristics of 2POMW and CM used in this study
97 can be found in the supplementary material, Table S1.

98 On the one hand, the 2POMW is a by-product from oil olive extraction process in which a
99 horizontal centrifuge is used to separate the oil fraction from this residue. This by-product was
100 a semisolid waste, slightly acidic, presenting a high solid and organic matter content. The
101 2POMW contains compounds as lignin, hemicellulose, cellulose, fats, water-soluble
102 carbohydrates and proteins (Morillo et al., 2009). Additionally, the 2POMW also presented a
103 high C/N ratio (41.23) and a concentration of soluble phenolic compounds of 1,6 g L⁻¹. In the
104 literature, it has been described that the presence of phenolic compounds in 2POMW depends
105 on the fruit (type, maturity, etc.), climatic conditions and processing technique (Alburquerque
106 et al., 2004; Morillo et al., 2009). The CM contains the faeces and urine from the animal, used
107 bedding, sand and sediments (Cong et al., 2018). On the other hand, the CM was also
108 characterized by a high organic nitrogen content, but presented low C/N ratio (18.52) and high
109 pH values. A high proportion of the CM's organic load correspond to cellulose, hemicelluloses
110 and lignin (Bernal et al., 2009).

111 Previous studies demonstrated that the anaerobic digestion of 2POMW and CM yielded a
112 maximum biogas production when mixed in a 75:25 ratio (2POMW:CM) (Pagés Díaz et al.,
113 2011; Giuliano et al., 2013; Rubio et al., 2019). Based on these results, the mixtures used in this

114 study were prepared keeping the ratio 2POMW:CM constant but modifying the total solids
115 percentage by diluting with demineralised water. Working in this way the reactors operated at
116 a 10%, 15%, 20% and 28% of total solids (TS) percentage. The TS percentage were selected in
117 order to cover the study of the performance of wet and dry anaerobic digestion. These tests
118 were named, by their TS percentage, as R10, R15, R20 and R28.

119

120 **2.2 Reactor set-up and operation**

121 The anaerobic reactions were carried out in laboratory-scale batch anaerobic digesters of 2 L
122 working volume and 1 L for the head space volume. A scheme of the reactor can be found in
123 the supplementary material, Figure S1. These reactors were hermetically sealed to ensure
124 anaerobic conditions during the digestion process, the wastes contained in the reactor were
125 continuously mixed by means of a mechanical stirrer. These reactors were operated in the
126 mesophilic range, at 35°C, and at a HRT of 15 d. The reactors were filled with the co-substrates
127 mixture up to 80% of effective volume (1600 mL) and were completed with 400 mL of
128 inoculum. These reactors were inoculated with a mesophilic seed from a laboratory digester
129 acclimatised to the treatment of the 2POMW and CM. The main physical–chemical
130 characteristics of the inoculum used in this study can be found in the supplementary material,
131 Table S1. The digester had two ports for sampling and biogas output. Samples were taken
132 three times a week and subsequently analysed. The biogas produced was collected in a 5 L
133 Tedlar® bags for its subsequent analysis. The volumetric biogas production was quantified
134 using a high precision gas meter (Ritter® Drum-type Gas Meter, 0.1 mbar). All the parameters
135 determined were analysed in triplicate. In order to ensure similar initial pH values, the pH of all
136 the mixtures were adjusted to 8.0 by adding a solution, 2.8 M, of Na₂CO₃.

137

138 **2.3 Analytical techniques**

139 The analytical parameters used for the physicochemical characterization of co-substrates and

140 monitoring the batch test were determined according to the Standard Methods (American
141 Public Health Association, 2005). Total solids (TS), total volatile solids (VS), pH and total
142 nitrogen Kjeldahl (TNK) were determined directly from the samples. Soluble chemical oxygen
143 demand (COD_s), dissolved organic carbon (DOC), alkalinity, total phenols and volatile fatty
144 acids (VFA) were measured over samples previously lixiviated. To do that, 10 g of sample were
145 mixed with 100 mL of distilled water during 30 minutes. Then the mixture was filtered through
146 a 0.45 µm glass-fibre filter. The DOC was determined by combustion/non-dispersive infrared
147 gas analysis method using a total organic carbon analyser Shimadzu® TOC-5000 (Fernandez et
148 al., 2008). Total phenols were determined by liquid chromatography according to the
149 procedure described in the literature (Medina et al., 2011). The chromatographic system
150 consisted of a Waters 717 plus autosampler, a Waters 600E pump, and a Waters 996 diode
151 array detector (Waters Inc., Milford, MA). A Spherisorb ODS-2 (5µm, 25 cm X 4.6 mm i.d.,
152 Waters Inc.) column was used. Separation was achieved using an elution gradient with an initial
153 composition of 90% water at pH 2.3 adjusted with phosphoric acid) and 10% methanol. The
154 concentration of the methanol was increased to 30% over 10 min and maintained for 20 min.
155 Subsequently, the methanol percentage was raised to 40% over 10 min, maintained for 5 min,
156 and then increased to 50%. Finally, the methanol percentage was increased to 60%, 70%, and
157 100% in 5 min periods. A flow of 1 mL/min and a temperature of 35°C were used. Phenolic and
158 oleosidic compounds were monitored at 280 and 240 nm, respectively. For determination of
159 VFA, the following procedure was used: samples were lixiviated and then filtered through a
160 0.22 µm Teflon filter, acidified with a solution 1:2 (v/v) of phosphoric acid, spiked with phenol
161 as internal standard and, finally, analysed in a gas chromatograph (Shimadzu® GC-2010)
162 equipped with a flame ionization detector and using a capillary column filled with Nukol
163 (polyethylene glycol modified by nitroterephthalic acid). The temperatures of the injection
164 port and detector were 200 and 250 °C, respectively. Hydrogen and synthetic air were used
165 for the gas chromatograph flame ionization at 40 and 400 mL min⁻¹. Total acidity (TVFA) was

166 calculated by the addition of individual VFA levels, taking into account the molecular weights
167 of the different VFAs in order to express this parameter as acetic acid concentration.

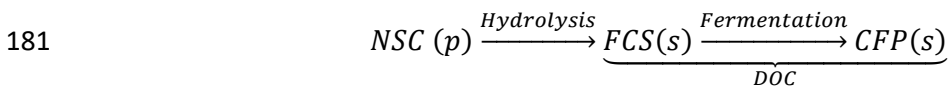
168 The main components of biogas (hydrogen, methane and carbon dioxide) were determined by
169 gas chromatography (Shimadzu® GC-14 B) with a stainless steel column packed with Carbosive
170 SII and a thermal conductivity detector. The injected sample volume was 1 mL and the
171 operational conditions were as follows: 7 min at 55 °C; followed with a ramp of 27°C min⁻¹ until
172 the temperatures reached 150 °C; detector temperature: 255 °C; injector temperature: 100°C.

173 The carrier was helium and the flow rate used was 30 mL min⁻¹ (Fdez-Güelfo et al., 2012).

174

175 **2.4 Determination of non-solubilized carbon (NSC), Fermentable carbonous substrate (FCS)**
176 **carbonous fermentation products (CFP).**

177 To evaluate the performance of the hydrolysis and the subsequent processes, as well as to
178 quantify the biodegradable fractions involved in each process, the trends of non-soluble carbon
179 (NSC), fermentable carbonous substrate (FCS) and carbonous fermentation products (CFP)
180 were determined. A scheme of the transformation is presented below.



182 The NSC is the particulate fraction of the organic carbon to be hydrolysed. The FCS is the
183 fraction of solubilized organic matter that has been transformed into fermentable substrates.

184 CFP represents the fraction of soluble organic carbon in acid form, i.e. the fraction
185 corresponding to VFAs. The sum of FCS and CFP fractions account to the **Dissolved Organic**
186 **Carbon (DOC).**

187 The NSC and FCS were determined according to equations (1) and (3) proposed in literature
188 (Fdez-Güelfo et al., 2012). The CFP was calculated according to equation (3) where A_iH ,
189 represents the concentration of each individual VFA measured by gas chromatography; n_i , is
190 the number of carbon atoms of each A_iH ; MW_i , is the molecular weight of each A_iH . The total
191 organic carbon (TOC) was calculated from equation (2) as suggested by (Navarro et al., 1993).

192
$$NSC = TOC - DOC \quad (1)$$

193
$$TOC = VS \cdot 0.51 \quad (2)$$

194
$$CFP = \sum_{i=2}^{i=7} AiH / MW_i \quad (3)$$

195
$$FCS = DOC - CFP \quad (4)$$

196

197 **2.5 Model structure**

198 The hydrolysis process sum up several steps such as lysis, non-enzymatic decay, phase
 199 separation, diffusion, adsorption, reaction, physical breakdown, etc. of particulate substrate
 200 (Vavilin et al., 2008). Because of that, the first order kinetics appears to be not applicable
 201 under all circumstances and therefore it is needed a model that accurately describe the
 202 disintegration and hydrolysis steps. In the literature, the Contois model has been
 203 demonstrated to adequately describe experimental data sets from a wide range of organic
 204 wastes (Sötemann et al., 2006; Nopharatana et al., 2007; Vavilin et al., 2008). The Contois
 205 model can be written as presented in equation (5):

206
$$\rho_{process} = k_{m,process} \cdot X \cdot \frac{S}{K_{S,process} \cdot X + S} = k_{m,process} \cdot X \cdot \frac{S/X}{K_{S,process} + S/X} \quad (5)$$

207 where $\rho_{process}$ is the process rate (g C kg⁻¹ fresh weight d⁻¹); $k_{m,process}$ is the maximum specific
 208 uptake rate of the process (d⁻¹); $K_{S,process}$ is the half-saturation coefficient for the ratio S/X (g C
 209 kg⁻¹ fresh weight); X is the hydrolytic (disintegration) biomass concentration (g C kg⁻¹ fresh
 210 weight) and S is the particulate compound concentration (g C kg⁻¹ fresh weight).

211 The fermentative and the methanogenic stages can be described by Monod Kinetics (Jeong et
 212 al., 2005; Fernandez-Morales et al., 2010; García-Gen et al., 2013). In addition, the anaerobic
 213 biodegradation of 2POMW produces a large quantity of polyphenols which could cause
 214 inhibition (Rubio et al., 2019). This inhibition event is usually observed by a decrease of the
 215 methane production and an accumulation of VFAs (Chen et al., 2008; Zhang et al., 2019). This
 216 phenomenon can be explained by the relationship between the polypohenols and the

217 propionic generation (Morillo et al., 2009) which leads to an inhibition in the methanogenic
218 stage. Because of that, a non-competitive function was included for modelling the inhibition in
219 the methanogenic stage according to the Hill function (Hill and Barth, 1977) this function is
220 presented in equation 6:

$$221 \quad I_P = b \cdot \left[1 - \frac{S_P}{K \cdot S_P + S_{P,lim}} \right] \quad (6)$$

222 where I_P is propionic inhibition factor of acetoclastic methanogens, S_P is the propionic
223 concentration (g C kg⁻¹ fresh weight), $S_{P,lim}$ is the mean propionic threshold concentration (g C
224 kg⁻¹ fresh weight), K is the Hill coefficient which defines the slope of the drop in the inhibition
225 function.

226 The processes rates and stoichiometry of the developed model is presented in Table 1 as a
227 Petersen matrix.

228
229

Table 1. Petersen matrix of the main anaerobic degradation processes taking place during the co-digestion of POMW and CM.

Component									
Process	NSC	FCS	CFP	CH ₄	CO ₂	Xh	Xf	X _{CH4}	Process rate
Hydrolysis	-1	(1-Yh)				Yh			$kh \cdot \frac{NSC/Xh}{K_{s,h} + NSC/Xh} \cdot Xh$
Fermentation		-1	(1-Yf)				Yf		$kf \cdot \frac{FCS}{K_{s,f} + FCS} \cdot Xf$
Methanogenesis			-1	(1-Y _{CH4})·f _{CH4}	(1-Y _{CH4})·f _{CO2}			Y _{CH4}	$k_m \cdot \frac{CFP}{K_{s,m} + CFP} \cdot X_m \cdot \left[1 - \frac{S_p}{K \cdot S_p + S_{p,lim}} \right]$
Decay Xh	1					-1			$k_{dec,Xh} \cdot Xh$
Decay Xf	1						-1		$k_{dec,Xf} \cdot Xf$
Decay X _m	1							-1	$k_{dec,Xm} \cdot X_m$
Nomenclature	<i>Yh: Yield of biomass on the hydrolysis process</i> <i>Yf: Yield of biomass on the fermentation process</i> <i>Y_{CH4}: Yield of biomass on the methanogenic process</i> <i>f_{CH4}: Yield, catabolism only, of methane</i> <i>f_{CO2}: Yield, catabolism only, of carbon dioxide</i> <i>kh: Hydrolysis rate</i> <i>kf: Fermentation rate</i> <i>km: Methanogenesis rate</i>					<i>ks,h: Half saturation constant of the substrate in the hydrolytic process</i> <i>ks,f: Half saturation constant of the substrate in the fermentation process</i> <i>ks,m: Half saturation constant of the substrate in the methanogenic process</i> <i>Xh: Hydrolytic biomass</i> <i>Xf: Fermentative biomass</i> <i>Xm: Methanogenic biomass</i> <i>k_{dec,Xh}: Hydrolytic biomass decay rate</i> <i>k_{dec,Xf}: Fermentative biomass decay rate</i> <i>k_{dec,Xm}: Methanogenic biomass decay rate</i>			

230 **3. RESULTS AND DISCUSSION**

231 The increased in the TS concentration leads to an increase in the organic substrates available in
232 the system. This increase in the substrate available for the microbial metabolisms could lead to
233 a higher biogas production or to a lower one because of inhibitory effects in the different
234 stages involved in the anaerobic digestion. In order to study the performance of the anaerobic
235 digestion, several experiments were carried out and modelled. Before the modelling, the mass
236 balance reconciliation was checked, obtaining a reconciliation higher than 90% in all the cases.

237

238 **3.1 Assessment of biomass fractions**

239 The modelization of the experimental results requires the determination of the substrate
240 fractions involved in the processes modelled. In this work, the characterization of the substrate
241 mixtures was carried out following the procedure previously described in the literature
242 (Navarro et al., 1993; Fdez-Güelfo et al., 2012) and the results obtained are presented in Table
243 2.

244 **Table 2.** Carbon fractions in the wastes mixture.

% TS	NSC (g kg ⁻¹ fresh weight)	FCS (g kg ⁻¹ fresh weight)	CFP (g kg ⁻¹ fresh weight)
10	25.6	9.60	0.15
15	34.5	13.62	0.20
20	55.5	17.99	0.33
28.6	73.3	27.00	0.22

245

246 From the characterization of the mixtures, it can be seen the very high NSC fraction, which
247 accounted to about a 73% of the total carbon contained in the mixtures. This particulate
248 fraction is the fraction that could be hydrolysed to form FCS. At the beginning of the tests, the
249 FCS concentration was about 27%. Finally, the smallest fraction was the CFP, which accounted
250 in all the cases percentages lower than 5% of the carbon concentration. The very low CFP

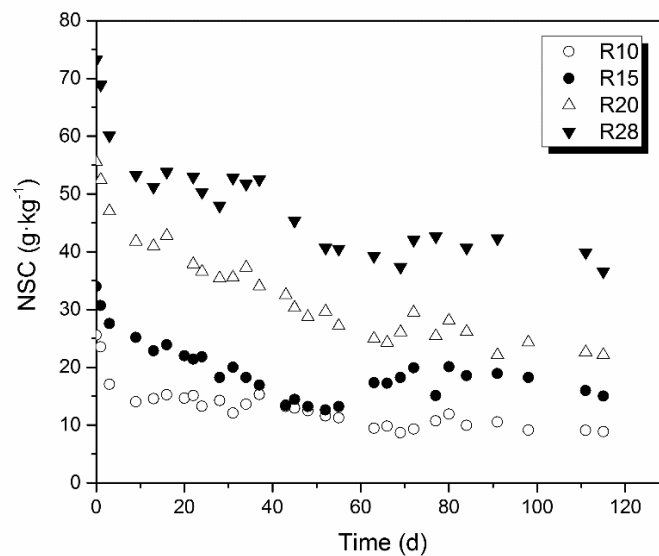
251 concentration could be explained because of its very high biodegradability which leads to a
252 fast consumption when it is generated (de Lucas et al., 2007).

253

254 **3.2. Reactor operation**

255 As stated above, the source of substrate fed to the reactors was the same in all the cases,
256 being the only difference its initial TS concentration which ranged from 10 to 28%. Because of
257 that, the different performances observed could only be explained because of the different
258 initials TS concentrations. In Figure 1 it is presented the evolution of the NSC fraction along the
259 experiments. **As can be seen in this Figure, the NSC fraction reached the steady-state**
260 **conditions after about 80 d.**

261



262

263 **Figure 1.** Evolution of the NSC fraction during the co-digestion of 2POMW and CM.

264

265 The increasing remanent NSC with the increasing initial concentration, could be related to the
266 accumulation of hardly hydrolysable compounds in the reactors quantified within the NSC

267 fraction or due to inhibitory events. In this work, the almost constant percentage of the NSC
 268 removal in all the cases, about 55%, ratified the accumulation of hardly hydrolysable
 269 compounds. Moreover, the final concentration of the NSC fraction accounted about 75% of
 270 total organic carbon (TOC) in the reactors. According to the literature, this value corresponds
 271 to the insoluble lignocellulosic fractions of the wastes used (Albuquerque et al., 2004). These
 272 saccharide chains are connected by hydrogen bonds and aggregated to form a three
 273 dimensional structure of fibrils, which are characterised by its toughness and water insolubility
 274 (Wang et al., 2020). A similar behaviour was observed when removing the FCS fraction, results
 275 not shown. In this case the non-fermentable fraction was also in all the cases almost the same,
 276 about a 50%.

277 Regarding to the biogas production, it must be highlighted the existence of a lag phase when
 278 dealing with high TS concentrations. Table 3 shows the most relevant information related to
 279 the biogas production in the different reactors.

280

281 **Table 3.** Main parameters in the methanogenic stage.

	Units	R10	R15	R20	R28
Lag phase length	d	10	16	22	34
Methane yield	g C kg ⁻¹ fresh weight	5.3	4.8	1.2	1.0
Methane composition	CH ₄ :CO ₂	80:20	75:25	70:30	65:35
Methane selectivity	g C g ⁻¹ C consumed	0.80	0.76	0.71	0.64

282

283

284 As can be seen in Table 3, the length of the lag phase linearly increased when the TS
 285 concentration increases. In the literature, the length of the lag phase has been related to
 286 different operational parameters. Such as inoculum size, physical conditions, inhibitors
 287 presence, etc. (Tsao, 1976; Baranyi and Roberts, 1994). Taking into account that in this work

288 the inoculum was the same in all the cases, the variation in the lag phase length only can be
289 caused by the different initial TS concentrations experienced.

290 When comparing the methane yield, it was observed that it decreased as the initial NSC
291 concentrations increases. This event only could be explained by an inhibition effect when
292 operating at higher TS percentages. It is also remarkable that the higher the TS percentages
293 the lower the CH₄:CO₂ in the biogas obtained, presenting a linear trend with an intercept of 5
294 and a slope of -0.1 (R²= 0.93). The explanation can be found in the two pathways of methane
295 generation, the hydrogenotrophic and the acetoclastic. At the beginning of the processes, the
296 methane was generated mainly by hydrogenotrophic activity, which is characterised by ratios
297 CH₄:CO₂ lower than 2 (Montero et al., 2008). This point was confirmed by the negligible
298 concentrations of hydrogen and the absence of VFAs degradation observed at the beginning of
299 the processes in spite of the initial TS concentration, see Figure S2. After that, the acetoclastic
300 culture could have been developed increasing the methane percentage in the biogas.

301 However, the development of the acetoclastic culture could have been not significant when
302 operating at high TS concentrations. In the literature, it has been described a higher proportion
303 of hydrogenotrophic methanogens in reactors operating with high TS content (Montero et al.,
304 2008). Then, the operation with high TS content in the anaerobic reactors leads to a longer lag
305 phase and to a prevalence of the hydrogenotrophic methanogenic culture which generated
306 biogas with lower methane percentages.

307 With the aim to deep into the mechanisms of the anaerobic transformations taking place
308 during the anaerobic digestion of the wastes studied, the methane selectivity was calculated as
309 the ratio of the methane-carbon generated to the total amount of carbon consumed in the
310 process, the obtained results are presented in Table 3. As can be seen in this Table, the higher
311 the TS percentage, the lower the methane yield and selectivity. These results indicates that a

312 controlling stage or an inhibition event affected the methanogenic reaction (Li et al., 2019; Shi
313 et al., 2019). In order to identify in which stage the inhibition took place, modelling works of
314 the hydrolysis, fermentation and the methanogenic stage were performed by using the model
315 previously described.

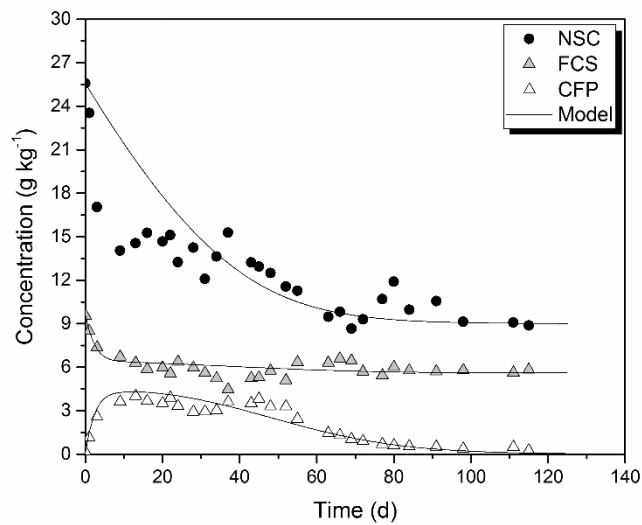
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317 **3.3. Model calibration and validation**

318 The model previously described was calibrated to fit the experimental data set obtained in the
319 co-digestion experiments with 10%, 15% and 28% of TS. As example, the calibration when
320 treating a 10% mixture is presented in Figure 2.

321

322 a)



323

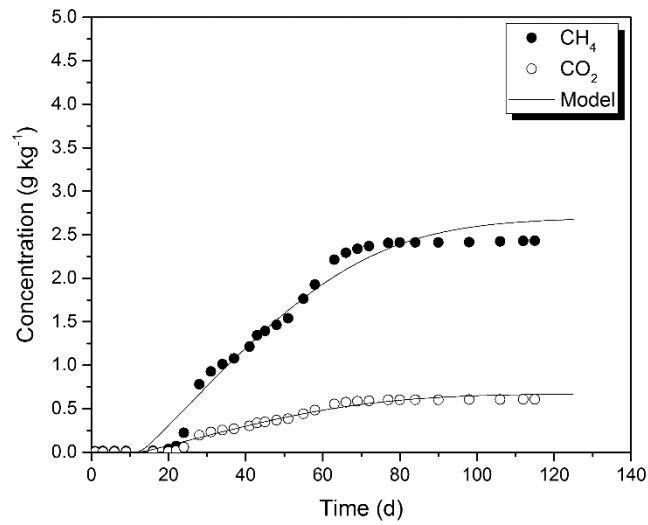
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326

327

328 b)



329

330 **Figure 2.** a) Results of the model calibration of carbon fractions and b) biogas evolution with
331 experimental results of the co-digestion at 10% TS.

332

333 As can be seen in Figure 2, the calibrated model accurately described the performance of the
334 system when dealing with a TS concentrations of 10%. A similar accuracy was obtained in the
335 other concentrations studied, 15 and 28%, in spite of the different initial TS concentrations.

336 The values of the main kinetic and stoichiometric parameters obtained after the calibration of
337 the model are presented in Table 4. The parameter not presented in Table 4 were fitted with
338 the typical values indicated in the literature (Batstone et al., 2002).

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343

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Table 4. Main kinetic and stoichiometric parameters of the model.

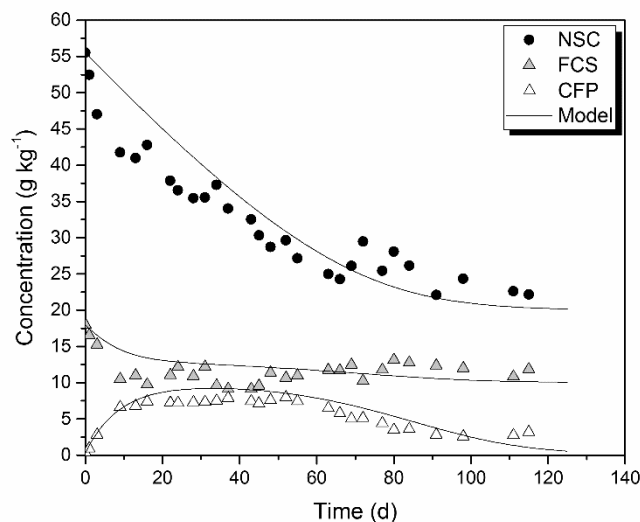
% ST	NSC		FCS		CFP			
	<i>kh</i> (d ⁻¹)	<i>Ks</i> (g C kg ⁻¹)	<i>kf</i> (d ⁻¹)	<i>Ks</i> (g C kg ⁻¹)	<i>km</i> (d ⁻¹)	<i>Ks</i> (g C kg ⁻¹)	<i>K</i>	<i>C_{Lim}</i> (g C kg ⁻¹)
10.0	0.70	10	2.50	4.0	0.75	4.0	0.95	2.7
15.0	0.69	10	2.47	4.0	0.76	4.0	0.90	2.6
28.6	0.70	10	2.50	4.0	0.75	4.0	0.99	2.7
Calibration value	0.70	10	2.49	4.0	0.75	4.0	0.95	2.7
Standard deviation	0.01	0	0.02	0	0.01	0.0	0.05	0.06

345

346 The maximum specific rates were so strongly associated with the biomass concentration that
347 the values could not be estimated individually. As can be seen in Table 4, the fitting values of
348 the *kh* and *kf* were the same in all the cases. The consistent values of the *kh* and *kf* parameters
349 indicates the absence of any inhibition or limitation in both, the hydrolytic and the
350 fermentative processes. Theoretically, the mixture R28 could experience mass transfer
351 limitations due to its very high TS concentration, within the range of the dry anaerobic
352 digestion (Ten Brummeler et al., 1991). However, this limitations do not occur, which could be
353 explained because it was easily hydrolysable and because of the particle size. It is known that
354 small particle size presents high surface to volume ratio, making easier the hydrolysis and
355 subsequent transformations.

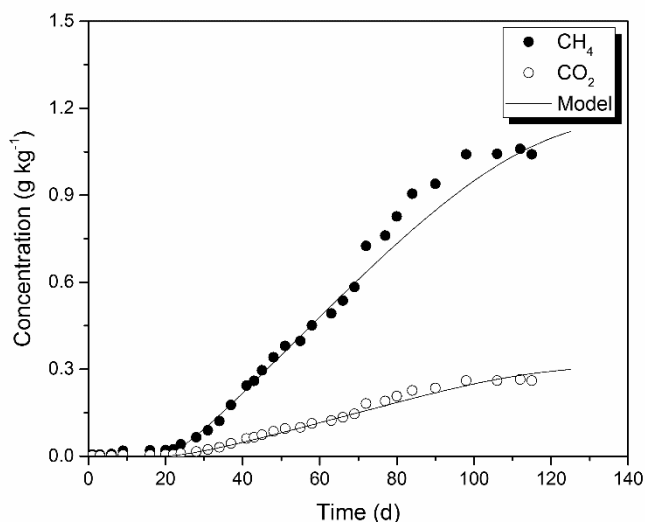
356 With regard to the *km* values, they were the similar in all the cases, indicating that the
357 maximum methanogenic rate is the same in spite of the initial TS concentration. However, this
358 rate was modified by an inhibition expression. The data sets corresponding to the three series
359 of experiments were simultaneously fitted, obtaining different lag phase lengths. The length of
360 these lag phase were linearly proportional to the initial TS concentration. Additionally, the
361 inhibition parameters were also very similar in all the cases, 0.95 for the *K* parameter and 2.7
362 for the threshold concentration, *S_{p,Lim}*. **This inhibitory effect could be caused by the very high**

363 concentration of VFA reached in the liquid bulk, see supplementary material Fig S2. This very
364 high VFA concentration can be explained because of the presence of polyphenols, which leads
365 to a very high propionic acid concentration (Pullammanappallil et al., 2001). When the
366 propionic acid concentration reached values higher than 2.7 g C kg⁻¹ inhibition effects were
367 observed, see figure S2. The explanation this event could be explained because of the
368 dissociated form of the acid can pass across the cellular membrane (Castro-Villalobos et al.,
369 2012). Once inside the cell, a high maintenance energy consumption is required which could
370 inhibit, and even stop, the methanogenic reaction.
371 Once finished the calibration stage, the model was validated using the calibrated parameters
372 previously obtained. The experimental results as well as the predictions of the model obtained
373 during the validation with experimental data corresponding to the 20% TS concentration are
374 presented in Figure 3. As can be observed in these figures an accurate prediction was
375 obtained.
376
377 a)



378

379 b)



380

381 **Figure 3.** a) Validation of COD fractions and b) methane evolution with experimental results of
382 the co-digestion at 20% TS.

383

384 From the results obtained, it can highlighted that the hydrolysis and fermentation rates
385 obtained during the calibration accurately predicted the results obtained when co-digesting
386 the mixture with a 20% TS. This results indicates that, in spite of the very different initial TS
387 concentrations studied in this work, no limitations were observed in the hydrolysis and
388 fermentation rates. However, inhibition events were observed during the methanogenic stage.
389 The inhibition could be caused by the accumulation of fermentation products, mainly
390 propionic acid, which could be caused by the presence of polyphenols in the mixture. The
391 phenolic compounds are characteristic of the by-products from the olive oil extraction and
392 contains a benzene ring conjugated to a propionic acid. In the literature it has been described
393 that propionic acid from phenolic compounds can slow the anaerobic acetoclastic
394 methanogenesis (Palatsi et al., 2011) (Borja et al., 1997).

395

396 **Conclusions**

397 The fractionated disintegration of the substrates provided accurate information for the
398 description of the co-digestion experiments. Additionally, the model developed allowed to
399 accurately predict a wide spectrum of initial TS concentrations. From the modelling results, it
400 was observed that the lower transformation rate was observed in the hydrolysis stage 0.7 d^{-1} .
401 However, when operating at high TS concentrations the inhibition event experienced in the
402 methanogenic stage slow down its rate becoming the controlling stage of the stabilization
403 process. This inhibition event was caused by the propionic acid and described by a Hill inhibition
404 function.

405

406

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