

1 **Title: Evaluating the effect of biochar addition on the anaerobic digestion of swine**
2 **manure: Application of Py-GC/MS**

3

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18 This is a pre-print of an article published in Environmental Science and Pollution

19 Research. The final authenticated version is available online at:

20 <https://link.springer.com/article/10.1007/s11356-018-2644-4>

21

22 Cite as:

23 Gómez, X., Meredith, W., Fernández, C. et al. Environ Sci Pollut Res (2018).

24 <https://doi.org/10.1007/s11356-018-2644-4>.

25

26 **Abstract**

27

28 The anaerobic digestion process of swine manure was studied when char was used as
29 supplement for improving performance. The use of pyrolysis-gas chromatography/mass
30 spectrometry (Py-GC/MS) was proposed for assessing the organic matter degradation.
31 The assessment on biogas production was carried out using samples of swine manure
32 (SM) supplemented with char in one case and pre-treated by microwave irradiation in
33 the other. This experimental set-up allows for the comparison of the biological
34 degradation observed under these two different configurations and therefore aids in
35 understanding the effect of char particles on the process. Results showed similar
36 performance for both systems, with an average improvement of 39% being obtained in
37 methane production when compared to the single digestion of SM. The analysis of
38 digestate samples by Fourier Transform Infrared (FTIR) Spectroscopy and Py-GC/MS
39 showed improved degradation of proteins, with the Py-GC/MS technique also capable
40 of identifying an increase in microbial derived material when char was added; therefore
41 highlighting the relevant role of carbon conductive particles on biological systems. Py-
42 GC/MS along with the use of (FTIR) Spectroscopy has proven to be useful tools when
43 evaluating anaerobic digestion.

44

45

46 **Keywords:** *Biochar, PY-GC/MS, FTIR, microwave pre-treatment, anaerobic digestion*

47

48

49 **Introduction**

50

51 The potential use of carbon-based conductive materials in anaerobic digestion processes
52 has gained great interest in recent years due to a great variety of reports claiming the
53 benefits associated with the increase in methane yields and attenuation of inhibitory
54 effects (Cai et al. 2016; Cuertos et al. 2016). Biochar (obtained from the thermochemical
55 conversion of biomass) and its use in land application has been traditionally proposed as
56 a means for increasing the carbon storage capacity of soils, enhancing their properties
57 and influencing the soil bacterial community (Wang et al. 2015) and a suitable soil
58 amendment for minimising the negative effects of heavy metals on soils (Gusiatin et al.
59 2016). In recent years, the addition of biochar to digestion systems has proven to be a
60 good alternative for providing new synergies between two traditional processes, as it is
61 the pyrolysis of lignocellulosic biomass and the digestion of wastes (Song et al. 2017).
62 In addition, the use of microalgae chars (Kim et al. 2016) and lignocellulosic biomass
63 offers a good prospect to escape from the dilemma of biofuel versus food production
64 (Roth and Spiess 2015).

65

66 Recent approaches for increasing the economic feasibility of waste treatment processes
67 involve the conjunction of thermal and biological treatments in an attempt to increase
68 energy efficiency and reduce the volume of side streams which can find limited disposal
69 options. Swine manure is an organic stream that presents a high energy-production
70 potential (Han et al. 2018). However, the use of complex substrates has been associated
71 with inefficiencies in the digestion process due to the slow rate of the first hydrolysis
72 stage. In an attempt to increase biogas yield and improve the degradation of organic
73 materials, the application of different pre-treatments have been studied (e.g. thermal,

74 alkaline, acid, chemical, mechanical, etc.) (Feki et al. 2015; Donoso-Bravo et al. 2016;
75 Martínez et al. 2017), although in some cases, their large scale implementation may be
76 limited due to their low economic feasibility.

77

78 On the other hand, the combination of anaerobic digestion and pyrolysis allows for new
79 synergies, which results in improvements thanks to the addition of biochar to the reactor
80 liquor. Recent studies on the performance of anaerobic digestion supplemented with
81 char report on the enhancement of the process by avoiding inhibitory conditions or
82 promoting the growth of methanogens (Luo et al. 2015; De Vrieze et al. 2016). Direct
83 interspecies electrons transfer (DIET) has also been suggested as one of the reasons for
84 obtaining better degradation rates when carbon-based conductive materials are added
85 (Dang et al. 2016). Although, several reports can be found in literature about the effects
86 of adding conductive carbon materials to biological systems, there is still missing
87 information about the evolution of the organic matter during biological transformations,
88 and to what extent the improvement obtained with the addition of conductive materials
89 may be comparable to those obtained from the application of different pre-treatment
90 technologies.

91

92 Pyrolysis-gas chromatography/mass spectrometry (Py-GC/MS) has been widely applied
93 for the study of complex materials such as soil organic matter and recently for biomass
94 characterisation (Liu et al. 2016; Zhu et al. 2016). The main advantages of the pyrolysis
95 route for analysing complex organic samples are minimal sample preparation, minimum
96 time of analysis and relative low cost (Prasad et al. 2006). There exists a wide
97 experience in the application of spectroscopic techniques for evaluating the
98 transformation of the organic material by biological process such as composting, soil

99 organic matter degradation and anaerobic digestion (Mopper et al. 2007; Kataki et al.
100 2017) allowing to elucidate the main relevant factors involved. In the present research
101 the use of Py-GC/MS, along with spectroscopy techniques as it is Fourier Transform
102 Infrared (FTIR) Spectroscopy has been investigated as tools for evaluating the effects of
103 biochar addition on anaerobic digestion systems.

104

105 The aim of this experimental work was the assessment of the anaerobic digestion of
106 swine manure (SM) following biochar addition as a means for improving system
107 performance. A comparison of performance is carried out to evaluate the effect of
108 supplementing the process with biochar particles. The process was evaluated using as
109 substrates SM and pre-treated SM by microwave irradiation. The pre-treated system
110 was used as reference for evaluating the modifications attained during digestion. Py-
111 GC/MS and FTIR were used for evaluating the biological performance.

112

113 **Materials and methods**

114

115 **Substrates and inoculum**

116

117 SM was collected from a livestock farm located in the surroundings of the City of León
118 (Spain). The manure was stored at 4 °C until required for use. The anaerobic sludge
119 used as inoculum was obtained from the wastewater treatment plant of León where the
120 anaerobic digester operates under mesophilic conditions (32 – 34 °C at an average
121 hydraulic retention time of 30 d). The digested sludge obtained from the wastewater
122 treatment plant was stored in closed vessel with a biogas release device and at room

123 temperature conditions (22 °C) for a month. This was done to allow the release of
124 biogas and avoid interferences during the batch experiments.

125

126 The chemical characteristics of SM and inoculum are presented in Table 1. Char was
127 produced from residual biomass of almond shell provided by “BIOTERM
128 AGROFORESTAL SL” (Córdoba, Spain). Raw biomass and char samples were
129 characterised as described by Albuquerque et al. (2016). Almond shells had volatile
130 matter (VM) content of 821.4 g/kg and an ash content of 5.5 g/kg. The char was
131 produced in a semi-continuous electrically heated reactor at 550 °C. The char thus
132 produced had a VM content of 88.6 g/kg and an ash content of 31.2 g/kg. Further details
133 of the operating conditions and char chemical and physical characteristics are described
134 by Gómez et al. (2016). The mean particle size was 225 µm for the biochar fraction
135 used for digestion experiments.

136

137 Table 1

138

139 **Batch digestion experiments**

140

141 Digestion experiments were performed using SM and microwave pre-treated SM as the
142 substrate. Batch digestion tests were carried out in Erlenmeyer flasks with a working
143 volume of 250 mL. Flasks were stirred at 125 rpm and maintained at 35 ± 1 °C by
144 means of a water bath. Each reactor contained 1.5 g of volatile solids (VS) from
145 manure. Inoculum was added to attain a VS ratio of 1.0 —inoculum-substrate (I/S)—.
146 Water was added to complete the 250 mL volume. Four reactors were used in each
147 experiment. Two replicates were used for measuring biogas production and

148 composition, while the other two were used for monitoring the liquid phase. A control
149 assay was also run in parallel to measure the background gas production from the
150 inoculum. Reactors were denoted as SM for the one digesting this substrate and
151 SM_MW for the one treating microwave pre-treated SM.

152

153 Char was added to a set of batch digestion experiments to evaluate the effect on gas
154 production and organic matter degradation. 3.0 g of char (dry basis) were added to each
155 batch reactors. The experimental conditions were kept the same way as described above
156 but with the addition of char in this latter case. Therefore, these reactors were denoted
157 as SM_Char and SM_MW_Char.

158

159 Gas and liquid samples were taken twice a week to measure the composition of biogas
160 and the concentration of volatile fatty acids (VFAs). Gas production was measured daily
161 using a liquid displacement bottles filled with an acid and high salinity solution to avoid
162 CO₂ absorption. The gas stored by these devices was sampled using gas syringe and
163 composition was measured by gas chromatography as described in analytical techniques
164 section. The duration of the experiments was based on the observation of total stoppage
165 of gas production.

166

167 Curves of specific methane production were fitted to a modified Gompertz equation (1).
168 This model has been successfully tested for adjusting biogas data obtained from batch
169 digestion assays (Cuetos et al. 2013):

170

$$171 \quad P(t) = P_{\max} \cdot \exp\left[-\exp\left[\frac{R_{\max} \cdot e}{P_{\max}} (\lambda - t) + 1 \right] \right] \quad (1)$$

172

173 Where $P(t)$ is the cumulative values of the specific methane production (mL/kg VS),
174 P_{max} is the maximum value obtained for the specific methane production (mL/kg VS),
175 R_{max} is the maximum production rate (mL/kg VS d), λ is the lag-phase time (d) and e is
176 2.71. The software OriginPro was used for fitting data to the equation and obtaining the
177 model parameters P_{max} , R_{max} , and λ .

178

179 **Microwave pre-treatment**

180

181 SM was pre-treated using a 1500 W/2455 MHz microwave oven MARS (CEM, North
182 Carolina, USA). The pre-treatment assay was performed by fixing power at 600 W
183 (maximum efficiency 80%). The temperature was increased with a ramp of 10 °C/min
184 until reach 80 °C and hold during 15 min. Temperature was controlled by an Infrared
185 (IR) temperature probe. The fresh SM was pre-treated by microwave (solid content of
186 23.1 g/L, see Table 1). 500 g of this substrate were pre-treated by equally distributing
187 the sample in 10 rotating vessels on the carousel.

188

189 **Analytical techniques**

190

191 Kjeldahl nitrogen (KN), total solid (TS), VS, alkalinity, ammonium, and pH were
192 measured in accordance with American Public Health Association (APHA) standard
193 methods (2012). Total phosphorous (TP) was measured by the use of inductively
194 coupled plasma atomic emission spectrometer (ICP-AES) (Perkin Elmer Optima 2000
195 DV). Gas production data were normalised to standard temperature and pressure (0 °C
196 and 1 atm). Biogas composition and VFAs were analysed using a gas chromatograph
197 (Varian CP 3800 GC) as described by Martínez et al. (2017).

198

199 Proximate analysis was performed according to ASTM 3302 for total moisture, UNE
200 3219 for analysing volatile matter and UNE 32004 for ash analysis. Proximate analyses
201 of the oven dried samples were carried out using a LECO MAC-300 thermogravimetric
202 analyzer (TGA). A LECO CHN-600 apparatus was used for the analysis of C, H and N
203 according to ASTM 5373. Sulphur was measured on a LECO SC-132, according to
204 ASTM 4239. The O content was calculated by difference of the sum of C, H, N, and S
205 values from 100% (on a dry ash-free basis). Biochar particle size analysis was
206 performed using a Laser Diffraction particle Size Analyser LS 13 320 Beckmann
207 Coulter. Samples selected for analysis were those comprising the initial state of the
208 experiment (SM and inoculum) and those obtained from the final stage of the batch
209 digestion test (digestates from the different experimental sets).

210

211 The surface of solid samples obtained after dismantling of reactors was analysed by
212 scanning electron microscopy (SEM). Char and digestate samples were obtained after
213 sedimentation of the reactor liquor and drying at 105 °C. Samples were ground using an
214 agate mortar. Samples were sputter-coated with gold in high vacuum (0.05 – 0.07 mbar)
215 with a coater Blazers SCD 004. The samples were examined using a JEOL JSM 6840
216 LV scanning electron microscope.

217

218 Infrared spectra of substrate and digested samples were recorded using an FTIR Thermo
219 Scientific Nicolet iS5 (ID7 ATR accessory, a monolithic diamond ATR crystal with
220 high-efficiency) spectrophotometer over the 4000 – 650 cm^{-1} range at a rate of 0.5 cm/s .
221 Sixteen scans were collected with 0.482 cm^{-1} spacing, averaged for each spectrum and
222 corrected against ambient air as background.

223

224 **Py-GC/MS**

225

226 Py-GC/MS analysis was conducted using a CDS Analytical 5200 pyroprobe, coupled
227 with an Agilent GC-MS (7890B GC; 5977A MSD). The probe has a temperature range
228 of 1–1400 °C, and a heating rate up to 20 °C/ms. A sample of homogeneous powder
229 (approximately 5 mg) was placed in a quartz capillary tube (length 25 mm; diameter 2
230 mm) and secured at each end with quartz wool. Pyrolysis was carried out to a final
231 temperature of 700 °C, with a heating rate of 20 °C/s and the final temperature then held
232 for 60 s. Products were retained on a Tenex trap before desorption at 280 °C (for 4 min)
233 and passage to the GC via a heated transfer line (310 °C). Product separation was
234 performed on an HP-5MS column (30 m x 250 µm x 0.25 µm). The GC oven
235 temperature was initially held at 50 °C for 0.5 min, then heated to 300 °C at a rate of 4
236 °C/min, where it was held for 5 min. The MS (EI of 70 eV) scanned in the mass range
237 of m/z 40 – 400, with an ion source temperature of 200 °C. Individual compounds were
238 identified using a NIST MS library and published data.

239

240 Two multivariate statistical methods, Hierarchical Cluster Analysis (HCA) and
241 Principal Component Analysis (PCA), were used for the evaluation of the samples
242 based chromatograms obtained from Py-GC/MS. HCA was used to classify the studied
243 samples into different groups. Grouping in clusters was carried out using Ward's
244 algorithm. Results are presented in the form of dendrograms.

245

246 PCA can be applied by creating a discrete data set of signals and representing these as
247 linear combinations of a set of factors. The major advantage of this statistical method is

248 its ability to reduce the dimensionality of the data set by eliminating redundant
249 dimensions and identifying meaningful underlying variables that better represent the
250 differences and similarities in a specific data set (Abdulla et al. 2013). Usually, only the
251 first few principal components in a descending order explain the maximum of the total
252 variance of all original variables. The score plot of the first principal components was
253 used to investigate the inter-relationships between the samples, as it allowed the
254 observation of clusters of samples (Martínez et al. 2016). Both HCA and PCA were
255 performed using OriginPro software.

256

257 **3. Results and discussion**

258

259 *3.1. Digestion experiments*

260

261 Results from proximate and ultimate analysis are shown in Table 2. The inoculum
262 sample presents high ash content associated with the degree of mineralisation of this
263 sample. The digested sludge presented a low volatile content. This sludge was
264 previously stored at room temperature to avoid interferences with the biogas production
265 of the samples during the batch experiments. During this storage period, microbial
266 activity led to the reduction in volatiles and readily degradable material resulting in a
267 decrease in total solid content and enrichment in complex compounds and minerals.

268

269 Table 2

270

271 Digestate obtained from the experiment treating SM as sole substrate, presented also a
272 marked increase in the ash content. This is explained by the high value of the substrate,

273 with an ash content of 27.2%. The mixture of this material with the inoculum which was
274 also characterised by high inorganic concentration leads to an initial ash content of
275 33.6%. Starting the digestion with such high mineral content results in a digestate with
276 remarkably high mineral components. The increase in ash content was of 38.8%, which
277 was also similar to the cases of the pre-treated SM batch experiment.

278

279 The pre-treatment of the substrate by microwave (MW) irradiation did not have any
280 effect on results obtained from proximate analysis, with similar values in volatiles and
281 ash content being reported. The addition of char to the mixture highly affected results
282 from proximate and ultimate analysis due to the high organic carbon content of this
283 material, but no effect was observed on elements measured.

284

285 Differences in methane production and its rate can be observed in Fig. 1a. The digestion
286 of SM was characterised by a delay in biogas production (See λ value in Table 3)
287 probably caused by the change in environmental conditions. Previous to the
288 experiments, the inoculum was stored at room temperature conditions for a month
289 period without any kind of substrate being supplied. The change from the storage to the
290 conditions of the experiments may explain the delay observed in the production of
291 biogas for all assays studied. In addition, the ammonium content of this inoculum was
292 much lower than that of the substrate (Table 1); therefore the initial value of the batch
293 reactors was in the range of 1300 to 1400 mg/L (See Fig. 1b). This change in
294 ammonium concentration and the high mineralisation of the inoculum used may be the
295 main reason for explaining the null biogas production during the initial stage.

296

297 Fig. 1

298

299 Table 3

300

301 The addition of char resulted in a decrease in the lag phase and also affected in a
302 positive way the methane production rate and the final cumulative value. Positive
303 effects have also been observed by different authors when evaluating the use of char and
304 conductive carbon materials in anaerobic digestion (Liu et al. 2012; Cuetos et al. 2016;
305 Dang et al. 2016). The aggregation of cells is considered a key factor for efficient
306 methanisation as a direct result of better electron transfer between obligate H₂-
307 producing acetogens and methanogens. Direct interspecies electrons transfer (DIET) is a
308 syntrophic metabolism in which free electrons flow from one cell to another without
309 being shuttled by reduced molecules such as molecular hydrogen or formate (Dubé and
310 Guiot 2015). It is believed that an improvement in this mechanism by the presence of
311 conductive materials directly results in higher gas yields. It is considered that the
312 presence of conductive carbon materials allows for an effective connection between
313 cells that are not strictly in physical contact with each other. It is suggested that electron
314 transport carriers associated with the outer surface of the cell are able to make the
315 required electrical contacts with the conductive material (Chen et al., 2014; Lovley
316 2017) resulting in an enhancement of methane production (Wang et al., 2017)

317

318 The irradiation of the substrate by MW resulted in the loss of ammonium. The pre-
319 treated sample of SM presented an ammonium concentration of 2775 mg/L which
320 resulted in slightly lower values of ammonium content for batch reactors using this
321 substrate (see Fig. 1b). This change may not seem relevant enough to be considered as
322 the responsible for the difference observed in the lag phase values. The addition of char

323 and the application of a pre-treatment may have facilitated the degradation of the
324 organic material by the anaerobic microflora causing as a result a reduction in the initial
325 lag-phase observed for these systems. The final ammonium concentration measured at
326 the end of the digestion tests present similar values for the four assays evaluated (see
327 Table 3).

328
329 MW irradiation also exerted a positive effect on methane production. Digestion systems
330 fed with the pre-treated substrate showed higher methane content in biogas probably as
331 a result of a better degradation of nitrogen containing material. Higher release of
332 ammonium is observed in Fig. 1b for pre-treated systems, although at the end of the
333 digestion these reactors presented lower ammonium content which may be explained by
334 the loss of this compound by volatilisation. However, the addition of char in
335 experiments using pre-treated SM did not report additional significant changes in the
336 performance of the digestion process. Methane curves obtained from these experiments
337 presented a quite similar gas evolution profile to those from the pre-treated system
338 without biochar.

339
340 The pre-treatment may have facilitated the degradation of complex material like
341 proteins and the lipid fraction of the SM. This effect is observed by the slope of the
342 cumulative methane production curve, which represents the methane production rate.
343 SM digestion systems present a change in the slope of the curve associated with the
344 complexity of the material being degraded. Right after the lag phase, the slope of the
345 curve obtained for the SM digestion system was 25 mL CH₄/kg VS d (days 10–19) and
346 this value was reduced to 6.4 mL CH₄/kg VS d on the following stage (days 21–39).
347 The SM_Char system presented a similar behaviour, but with higher rates being

348 obtained in each stage (29.4 and 9.0 mL CH₄/kg VS d for the two different stages). The
349 application of MW irradiation resulted in a homogenous degradation profile, where
350 differences in the methane production rate were eliminated once the lag phase had
351 elapsed. The improvement of MW pre-treatment on anaerobic digestion has been
352 demonstrated by several authors (Tyagi et al. 2014; Yeneneh et al. 2017). Changes
353 experienced by the organic material and modifications in the specific surface area of the
354 organic particles have been proposed as the reasons given for obtaining an enhancement
355 in digestion when MW pre-treatment is applied (Martínez et al. 2017). In the present
356 study the methane yield from pre-treated systems and biochar aided systems was about
357 420 mL CH₄/g VS, while in the SM reactor this value was 28% lower.

358

359 The addition of char did not represent any significant changes in the evolution of NH₄⁺
360 content in the different experiments. Values of this parameter were quite similar for the
361 different experiments indicating that the adsorption of this cation on the char surface
362 can be disregarded as the cause for improving the performance of the digestion process
363 when SM was treated.

364

365 Fig. 1 also presents the evolution of VFA in batch tests. The anaerobic degradation of
366 SM was characterised by high VFA production (Fig. 1c). These high values along with
367 the negative effects associated with the ammonium levels in the reactor may have been
368 responsible for the delay observed at the initial stage of the trial. Ammonium content in
369 the reactor was below the threshold value considered as inhibitory (3.0 - 3.7 g/L for
370 ammonium values and 1.0 -1.1 g/L for free ammonia values (Gallert and Winter 1997;
371 Moestedt et al. 2016)). However, the ammonium concentration may have been high
372 enough to affect the performance of the microflora since the anaerobic consortium had

373 not been previously adapted to this type of substrate and was stored at room temperature
374 to avoid interference with the background biogas production.

375

376 Acetic acid reached values around 3 500 mg/L at the initial stage of the fermentation,
377 indicating that the acidification stage and the subsequent consumption of this acid may
378 be suffering imbalances. The addition of char did not change this trend, but the carbon
379 particles may have offered a protective site to the anaerobic microflora explaining
380 therefore the lower values of λ obtained for the SM_Char system. On the other hand, the
381 application of the MW pre-treatment favoured the consumption of this acid due to the
382 volatilisation of ammonium experienced by the substrate, lowering its level in the
383 reactor and therefore enhancing the performance of acetotrophic methanoges. Acetic
384 acid concentration in pre-treated reactors was below 1 500 mg/L. Values of λ were
385 similar for both systems and were close to that derived from the SM_Char system,
386 corroborating the hypothesis of available protective sites for anaerobic microflora.

387

388 **Scanning electron microscopy**

389

390 Images of the char surface and dried digestate particles are shown in Fig. 2, with the
391 porous structure of char being easily observed. Digestate material, obtained from the
392 SM batch digestion experiment, presents an irregular surface which is very different
393 from the surface observed from the solid particles obtained from the experiment of SM
394 digestion supplemented with char. The surface of particles presents a more corrugated
395 structure with the porous associated to the char particles being observed at the centre of
396 the photograph. Digestate solid particles seem to be fully covering the carbon
397 conductive material. Surface analysis performed by the same microscope, reported a

398 carbon content of $87.1 \pm 1.1\%$ for the char particles, this value was just $41.7 \pm 1.7\%$ for
399 the SM digestate and was increased to $65.9 \pm 1.6\%$ as a result of the presence of char in
400 the sample.

401

402 Fig. 2

403

404 **Results from FTIR analysis**

405

406 The degradation of the organic material was studied by means of FTIR spectroscopy.

407 Assignment of main absorption bands was based on Cuetos et al. (2009) and Fierro et

408 al. (2016). Initial samples and digestates were evaluated. Fig. 3 shows the different

409 spectra obtained.

410

411 Fig. 3

412

413 The spectrum of inoculum and substrate shows similar bands with the latter presenting

414 higher intensities in some particular regions. A broad absorption band at $2\,500\text{--}3\,800$

415 cm^{-1} is attributed to O–H vibration and it was common to all samples analysed, with the

416 exception made for the sample of char. The peak centred at $3\,400\text{ cm}^{-1}$ is ascribed to

417 hydrogen vibrations of the OH groups of alcohols, phenols and organic acids, but also

418 to amide hydrogen vibrations. The presence of aliphatic compounds is observed in the

419 range $2\,800\text{--}3\,000\text{ cm}^{-1}$, along with the signals at $1\,450\text{--}1\,410\text{ cm}^{-1}$ associated this latter

420 with the CH_2 scissor deformation vibration. The SM sample presents a higher intensity

421 band in this region which is significantly reduced after the anaerobic treatment (see

422 spectra from digested samples). The protein region associated with nitrogen containing

423 compounds (absorption bands at about 1 650 and 1 540 cm^{-1}) also presents high
424 intensity in this same sample while it was highly attenuated in the inoculum sample.
425 This is also the case for the region at 1 300–1 230 cm^{-1} , which was ascribed to
426 phospholipids (PO_2) asymmetric stretching, C–O in carboxylic acids and protein amide
427 III band (C–H and N–H) (Provenzano et al 2015).

428

429 The inoculum sample also presents an important band at 1 630 cm^{-1} ascribed to
430 aromatic structures, which also remains clearly identifiable in digestate samples. The
431 high intensity band at 1 030 cm^{-1} was also recurrent in the different samples obtained
432 from the digestion experiments and was also reported as characteristic of sludge and
433 digestates (Fierro et al. 2016).

434

435 The degradation of SM under anaerobic conditions leads to the consumption of readily
436 degradable organic matter, therefore the decrease in signals ascribed to aliphatic and
437 carbohydrate regions are in consonance with this statement. The digestate samples
438 present a relative increase of the aromatic signal (1 630 cm^{-1}) with regard to that of
439 proteins at 1 550 cm^{-1} . In addition, the application of microwave pre-treatment allows
440 for an improvement in the degradation of these compounds, an effect which was
441 observed from the change in the relative intensity of the protein peak, with the SM_MW
442 sample presenting a decrease in the intensity of this peak when compared to the one at
443 1 630 cm^{-1} . This effect was also observable when Char was added to the digestion
444 process. However, the combination of pre-treatment and char addition did not lead to
445 additional modifications in the organic matter. Digestate sample obtained in this latter
446 case (char addition and MW pre-treatment), presented a similar FTIR spectra to that of

447 the char supplemented digestate. That is, with similar relative intensities when
448 comparing signals at 1 630, 1 550 and 1 410 cm^{-1} .

449

450 *3.4. Results from Py-GC/MS analysis*

451

452 Pyrograms represented in Fig. 4 show a wide diversity of peaks in SM_substrate sample
453 and inoculum. Clear differences are easily observable for some major peaks. The first
454 volatile compounds obtained present as main components butane and acetic acid (see
455 supplementary material SM1 and peak identification table in SM3). These short chain
456 compounds were reduced in digestate samples, with the exception of the SM_Char
457 digested sample which presents a relative intensity similar to that obtained for toluene.
458 Phenol derivatives can arise from the decomposition of proteins, polycarboxylic acids
459 (Bracewell et al. 1980) or polysaccharides (Wilson et al 1983). The different
460 performance in the digestion system when char was added to aid in the mineralisation of
461 the organic material is also observed in the Py-GC/MS profiles. The addition of char
462 allowed for a faster start-up of the batch digestion tests and aided in the degradation of
463 proteinaceous material as it was also observed in FTIR spectra. In the present case, this
464 behaviour is corroborated by the differences observed from the relative intensities of
465 toluene, C₂-benzene, styrene and phenol peaks with regard to those ascribed to short
466 chain molecules. The presence of toluene and styrene has been associated as a lignin
467 pyrolysis product; although styrene arises also very probably from non-hydrolysable
468 proteins, peptides and tannins (Dignac et al. 2006).

469

470 Fig. 4

471

472 Organic compounds of the furan type were identified among the pyrolysis products of
473 the inoculum sample (furfural and 5-methyl furfural, see supplementary material SM1
474 and SM3), but were not discernible in the substrate sample neither in the digested
475 samples. These compounds are considered as the pyrolysis products of polysaccharides
476 (Dignac et al. 2005; El Fels et al. 2014) and furane type compounds have also been
477 identified (furane and benzofurane) in the pyrolysis products of humic acids (Zhao et al.
478 2012) with their presence in the inoculum sample being probably associated with humic
479 and fulvic substances.

480

481 Indole and methyl-indole were identified in the samples studied. These compounds have
482 been recognised as pyrolysis products derived from tryptophan (Kebelmann et al. 2013).
483 Other microbial pyrolysis products selected for tracking microbial signals, this case in
484 sediments, are Benzyl nitrile, 2-furanmethanol, indole, phenol and pyrrole (Zhu et al.
485 2016). In the present study, phenol and Indole were clearly discernible in all
486 chromatograms. Benzeneacetonitrile and benzenepropane nitrile can be also ascribed to
487 microbial signals (See supplementary material).

488

489 Terrestrial lignocellulosic biomass yields a range of products such as acetic acid,
490 furfural, syringol, vanillin, guaiacol, cresol, 2,3-dihydro-benzofuran, coniferyl alcohol
491 and levoglucosan (Wu et al. 2010). Many of these compounds were also identified in
492 the substrate and digestate samples, with particular incidence of cresol which presents a
493 high intensity signal in the chromatograph of inoculum and digested samples.

494

495 SM_substrate and inoculum samples present a large amount of linear fatty acids as well
496 as sterol components (Wang et al. 2015). Lipid fraction of sewage sludge and compost

497 has been analysed by different authors using NMR, pyrolysis–gas chromatography/mass
498 spectrometry with tetramethylammonium hydroxide and spectroscopic techniques
499 (Fierro et al. 2016; Li et al. 2017; Fukushima et al. 2018). In the present study C₁₆ and
500 C₁₈ lipids and their fatty acid forms were identified in the substrate, inoculum and
501 digested sample. These long chain compounds (oleic, stearic, and palmitic acids) have
502 been described as major constituents of wastewaters (Palatsi et al. 2012).

503

504 Sterols presented a great contribution to the chromatogram obtained from the Py-
505 GG/MS analysis of the inoculum sample. Among sterols, cholesterol is the main sterol
506 of animal origin and in particular coprostanol which is used as marker of faeces
507 (Volkman 1986). The recurrent appearance of sterols in digested samples is explained
508 by the accumulation of these components during the digestion process. Due to their
509 complex structure, these molecules were not affected by the microbial degradation.

510

511 Statistical analysis of Py-GC/MS chromatographs allowed the global evaluation of the
512 samples. Results in Fig. 4 show a high correlation between the inoculum sample and the
513 digested swine manure, irrespectively of the application of the MW pre-treatment.

514 Although biogas evolution from the MW pre-treated sample represented an
515 improvement in the biological degradation of the material which was observed by an
516 increase in the biogas production rate (batch tests) and an improvement in the
517 degradation of proteinaceous material (as observed in FTIR analysis), the digested
518 material presents similar pyrolysis products to those obtained from the SM digestate
519 sample, with the exception of the lipidic fraction which present signals with greater
520 intensities in this latter case.

521

522 Statistical analysis also revealed that the presence of char is affecting the distribution of
523 the pyrolysis products with those digestates containing char presenting a closer
524 correlation. Py-GC/MS was capable of discerning differences in the evolution of the
525 organic material when char was added to the digestion process. Pyrograms obtained
526 from the char digestion systems present high intensity signals associated with the
527 presence of microbial biomass and lignocellulosic compounds.

528

529 SM_substrate sample on the contrary, presented a great variety of compounds in the
530 lipidic region (high contribution of -CH₂-) which were greatly reduced in all digested
531 samples evaluated remaining only as major components of this region, those with C₁₆
532 and C₁₈ forms. The degradation of the organic content of SM by anaerobic digestion led
533 to a great decrease of small molecules in pyrograms as evidence of the consumption by
534 microorganisms of less complex compounds leading to the accumulation in digested
535 samples of lignin type material and components derived from microbial biomass. Char
536 addition accentuated the differences related to the distribution of organic compounds
537 and this may be explained by the creation of a more favourable environment leading to
538 an improvement in the development of microbial consortia as corroborated by different
539 authors (Dang et al. 2016; De Vrieze et al. 2016).

540

541

542 **4. Conclusions**

543

544 This study of batch digestion systems of swine manure (SM) showed similar
545 improvements in biogas yields for both the addition of biochar and microwave pre-
546 treatment, demonstrating the great benefits of the addition of conductive carbon
547 materials to biological processes. These results demonstrate the possibility of new
548 synergies between traditional pyrolysis and digestion processes from a perspective of
549 joining these two technologies in future biorefineries.

550

551 Py-GC/MS significantly aided in the interpretation of results regarding the
552 transformation of organic components and was capable of discerning differences in the
553 evolution of the different digestion tests by evaluating the distribution of pyrolysis
554 products obtained from digested samples. A relative increase in microbial biomass
555 pyrolysis products and lignin type material was observed for char containing samples
556 while FTIR spectra although providing valuable information was limited in giving an
557 explanation in the different performance when evaluating char supplemented systems
558 and those of the microwave pre-treatment assay.

559

560 **Acknowledgments**

561 This research was possible thanks to the financial support of Project CTQ2015-68925-R
562 and UNLE15-EE-3070 funded by Ministry of Economía y Competitividad, Spanish
563 Government

564

565

566 **Conflict of interests**

567 Authors declare no conflict of interest

568

569 **References**

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740

741

742 Figure caption

743

744 **Fig. 1** (a) Specific methane production of SM and (b) ammonium content obtained from
745 batch digestion tests. (c) VFA results obtained from the digestion of SM, (d) pre-treated
746 SM (SM_MW), (e) SM supplemented with char (SM_Char), and (f) and pre-treated SM
747 supplemented with char (SM_MW_Char)

748

749 **Fig. 2** Results from scanning electron microscopy. Surface of char, digestate and
750 digestate containing char particles. Samples were obtained from batch digestion
751 experiments

752

753 **Fig. 3** FTIR spectra of the initial and digestate samples obtained from the batch
754 digestion of swine manure (SM), the pre-treated SM (SM_MW) and systems
755 comprising the addition of Char (SM_Char and SM_MW_Char)

756

757 **Fig. 4** (a) Chromatograms of pyrolysis products released from substrate samples,
758 inoculum and digestates obtained at the end of the experiment. Statistical analysis: (b)
759 HCA and (c) PCA

760

761

762

763 Table 1. Chemical characteristics of inoculum and swine manure (SM).

764

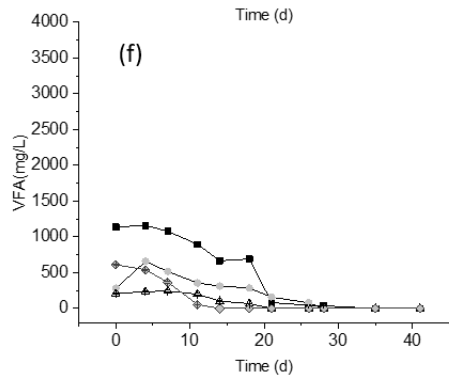
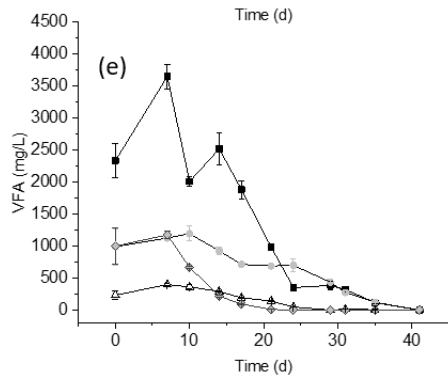
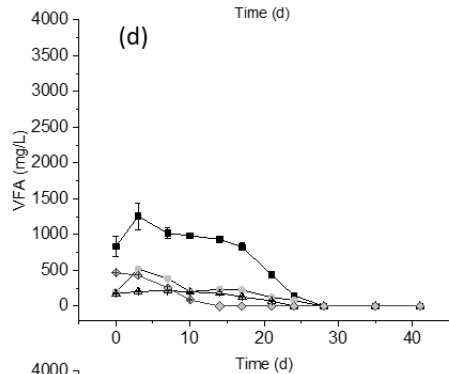
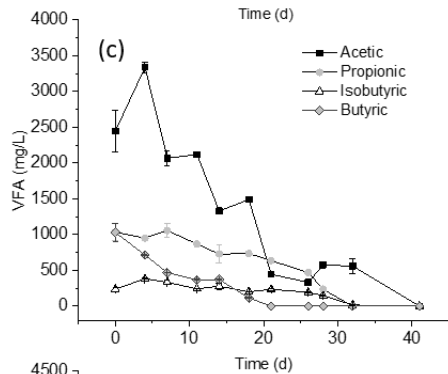
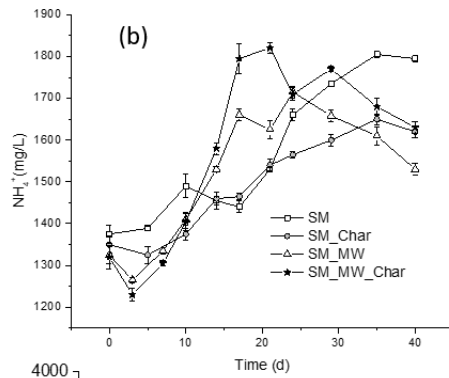
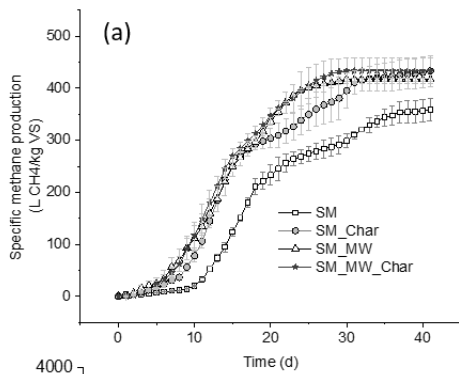
765 Table 2. Results from proximate and ultimate analysis of initial samples and digestates.

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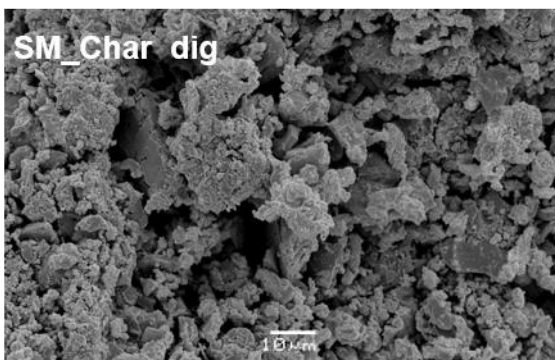
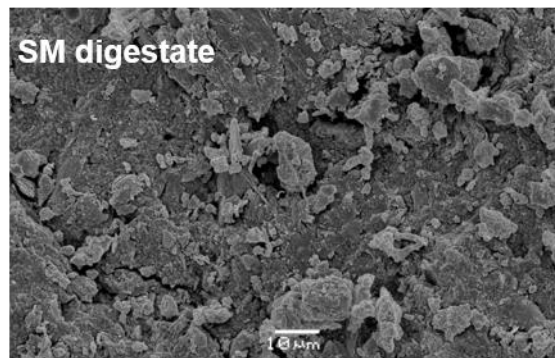
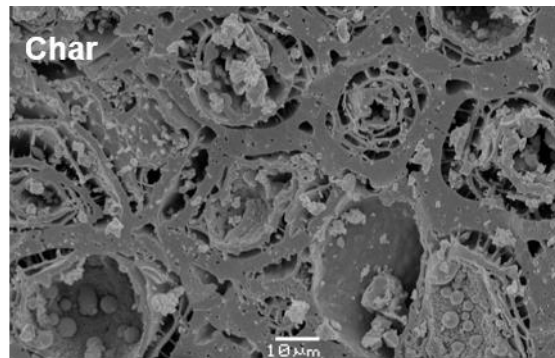
767 Table 3. Results obtained from batch digestion test of swine manure (SM), pre-treated

768 SM (SM_MW) and char supplemented systems (SM_Char, SM_MW_Char).

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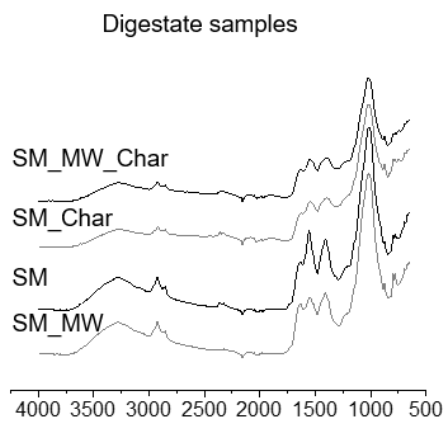
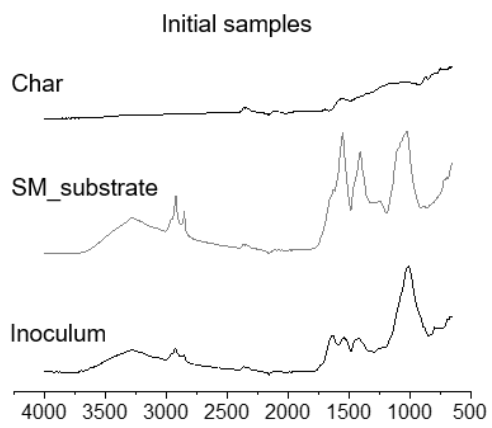


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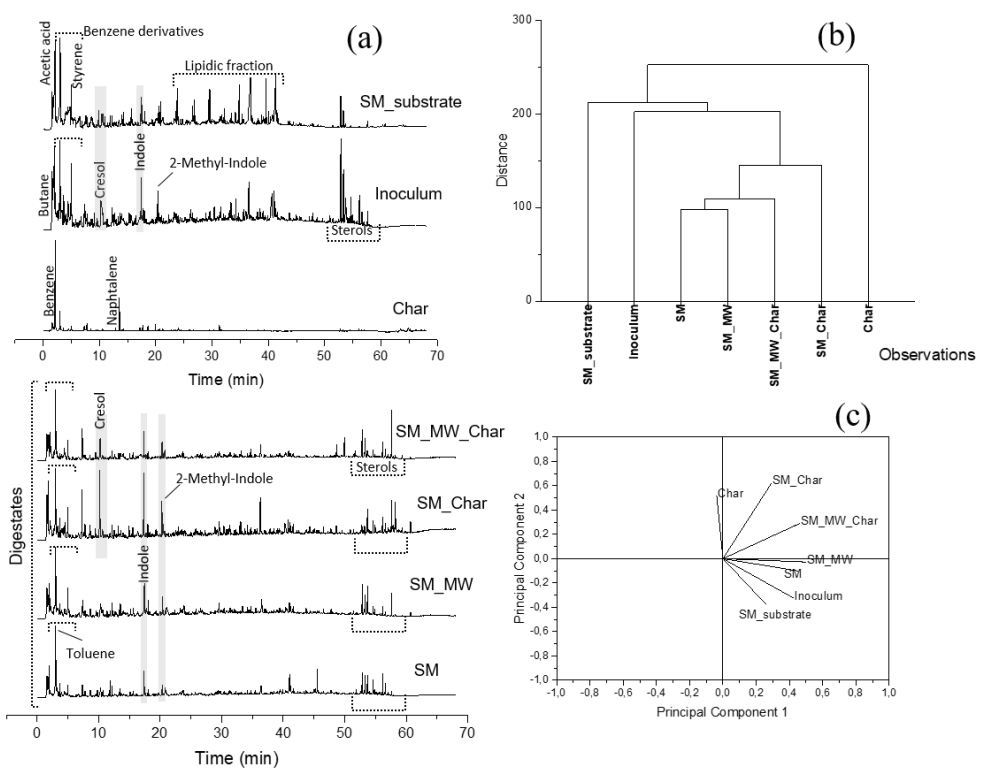


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777

778 Table 1. Chemical characteristics of inoculum and swine manure (SM).
 779

Feedstock	TS (g/L)	VS (g/L)	TN (g/kg) ^a	P (ppm) ^a	pH	C/N	NH ₄ ⁺ (mg/L)
SM	23.1	15.2	3.6	12 207	6.9	10.9	3 085
Inoculum	47.6	27.8	4.2	22 452	7.5	8.5	1 050

780 ^a expressed on a dry basis. TS: Total solids. VS: Volatile solids. TN: Total Kjeldahl
 781 nitrogen.
 782
 783

784 Table 2. Results from proximate and ultimate analysis of initial samples and digestates.
 785

Sample	Proximate analysis			Ultimate analysis				
	Moisture (%)	Volatiles (%)	Ash (%)	C (%)	H (%)	N (%)	S (%)	O (%)
SM_substrate	8.6	61.1	27.2	40.1	4.8	4.3	1.24	49.6
Inoculum	4.8	49.9	40.9	32.0	4.2	4.2	1.5	58.0
Char	3.9	8.9	3.1	90.4	2.1	0.4	0.1	7.1
<i>Digestates</i>								
SM	5.2	44.6	46.7	28.9	3.6	3.2	1.4	63.0
SM_Char	5.3	31.7	28.8	52.8	2.9	2.2	0.9	41.2
SM_MW	5.2	44.0	46.1	28.1	3.4	2.8	1.6	64.1
SM_MW_Char	5.0	30.9	28.1	53.9	2.9	2.1	1.0	40.1

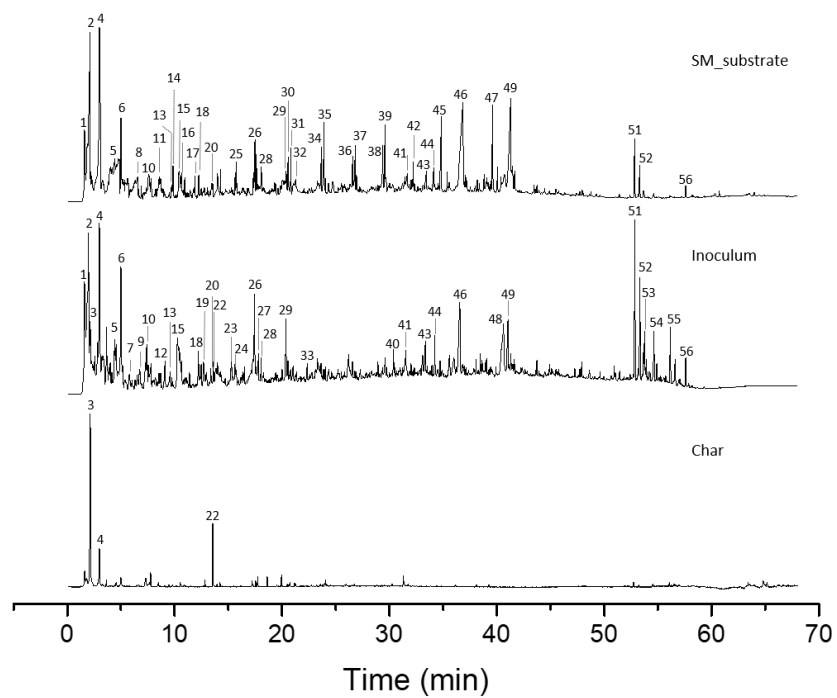
786
 787

788 Table 3. Results obtained from batch digestion test of swine manure (SM), pre-treated
 789 SM (SM_MW) and char supplemented systems (SM_Char, SM_MW_Char).
 790

Parameter	SM	SM_Char	SM_MW	SM_MW_Char
Cumulative CH ₄ production (mL)	448.1 ± 19.7	593.1 ± 50.4	625.5 ± 4.1	649.8 ± 36.9
CH ₄ Yield (mL/g VS)	298.7 ± 23.8	395.4 ± 42.7	416.7 ± 27.9	433.2 ± 37.9
CH ₄ (%)	68.9 ± 1.7	69.3 ± 1.7	73.3 ± 1.6	75.3 ± 1.7
pH _{initial}	7.0 ± 0.1	7.1 ± 0.1	7.7 ± 0.1	7.9 ± 0.1
pH _{final}	8.0 ± 0.1	8.0 ± 0.1	7.8 ± 0.1	8.1 ± 0.1
<i>Modified Gompertz model parameters</i>				
λ (d)	9.19 ± 0.21	6.07 ± 0.22	5.88 ± 0.09	5.83 ± 0.07
<i>P</i> max (mL/kg VS)	349.6 ± 5.1	426.1 ± 6.0	428.8 ± 2.4	441.2 ± 1.9
<i>R</i> max (mL/kg VS d)	21.2 ± 3.9	24.5 ± 4.5	27.5 ± 2.3	28.8 ± 1.8
<i>R</i> ²	0.992	0.990	0.998	0.999

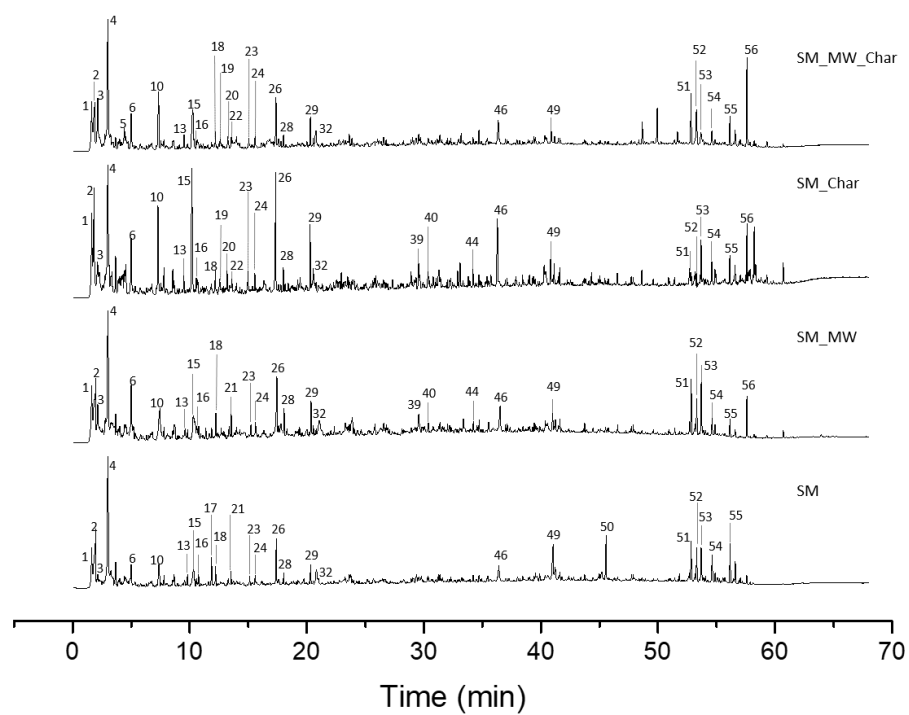
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Fig. SM1. Chromatograms of pyrolysis products released from substrate samples, inoculum and char



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Fig. SM2. Chromatograms of pyrolysis products released from digested samples



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797 Table SM1. Selected pyrolysis fragments identified from substrates and digested
 798 samples.
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No	Compound	m/z
1	Butane	43, 58
2	Acetic acid	43, 60
3	Benzene	78
4	Toluene	91, 92
5	C2-benzene	91, 106
6	Styrene	78, 104
7	Furfural	95, 96
8	Pentanoic acid	60, 73
9	5-Methyl furfural	109, 110
10	Phenol	66, 94
11	2-Propenyl Benzene	91, 103, 117
12	Indene	115, 116
13	o-Cresol, m-Cresol	107, 108
14	acetophenone	77, 105, 120
15	p-Cresol	107, 108
16	2-Methoxy-phenol (guaiacol)	109, 124
17	Benzyl methyl ketone	43, 91, 134
18	Benzeneacetonitrile	90, 117
19	2,5-Dimethyl phenol	107, 122
20	2-Ethyl-phenol, 4-Ethyl-phenol	107, 122
21	Methyl-phenyl-acetate	91, 150
22	Naphtalene	128
23	Coumaran	91, 120
24	Benzenepropane nitrile	91, 131
25	2-Butanone-4-phenyl	91, 105, 148
26	Indole	90, 117
27	Methyl-naphtalene	115, 142
28	Vinylguaiacol	135, 150
29	2-Methyl-indole	130, 131
30	Tetradecene	41, 43, 55, 70, 83, 97
31	Tetradecane	43, 57, 71, 85
32	Ethylguaiacol	137
33	Eugenol	149, 164
34	1-Pentadecene	55, 69, 83
35	Pentadecane	57, 71, 85
36	1-Hexadecene	55, 69, 83
37	Hexadecane	57, 71, 85
38	1-Hexadecanol, 2-methyl	57, 69, 83, 97
39	Tetradecane, 2, 6, 10-trimethyl	57, 71, 85
40	1-Dodecanol 3,7,11-trimethyl	57, 69, 83, 97, 111
41	Tetradecanoic acid	60, 73, 129, 185
42	Phytol	57, 71, 81
43	i-Propyl 12-methyltetradecanoate	
44	Pentadecanoic acid	60, 73, 85, 97
45	2-Heptadecanone	58, 71
46	Hexadecanoic acid	60, 73, 83, 129

47	2-Nonadecanone	58, 71
48	6-Octadecenoic acid	55, 69, 83, 97, 11
49	Octadecanoic acid	60, 73, 129
50	Octadecanamide	59, 72
51	Cholest-3-ene,(5 β)-	135, 147
52	Cholest-2-ene	147, 161
53	Squalene	69,81
54	Cholesta-3,5-diene	147, 159
55	Methyl 3-hydroxycholestan-26-oate	
56	Coprostanol	135, 149, 161

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