

This is the submitted version of the article: Saraya, Tahseen, et al. *Anaerobic bioremediation of PAH contaminated soil: assessment of the degradation of contaminants and biogas production under thermophilic and mesophilic conditions* in Environmental engineering and management journal, vol. 14, no. 1 (Jan. 2015), p. 153-165

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34 **1. Introduction**

35

36 Soil contaminated with polycyclic aromatic hydrocarbon (PAHs) is a global
37 environmental problem that is continuing to increase because these compounds are widely
38 present in many substances used as part of modern life (Agarwal, 2009). Warnings about the
39 dangers and potential side effects of PAHs have motivated researchers to investigate their
40 bioremediation and detoxification. It is well known that, for many organic contaminants,
41 aerobic treatments are more efficient and widely used than anaerobic ones because of their
42 broader catabolic range (Cajthaml et al., 2002). However, aerobic processes are not the only
43 options that need to be considered because different types of contaminants can be present for
44 which such processes are difficult to apply. High number of materials contaminated with
45 PAHs could be treated by using anaerobic processes (Popescu et al., 2009; Soceanu et al.,
46 2009), and therefore it is necessary to understand their fate under these conditions, as well as
47 understand if anaerobic treatments could be applied as a biological alternative process.

48 Although anaerobic bioremediation of PAH-contaminated soil has been investigated
49 under a number of different conditions relatively little is still known and more research is
50 necessary (Zhang et al., 2000). Recently, various electron acceptors have been used for the
51 anaerobic bioremediation of different compounds such as hydrocarbons (Callaghan et al.,
52 2006; Dionisi et al., 2006; El-Hadj et al., 2006). Evidence of anaerobic degradation of various
53 PAHs have been reported: phenanthrene and fluorene have been degraded by sulphate-
54 reducing bacteria (Coates et al., 1997) and under nitrate-reducing conditions (Eriksson et al.,
55 2003), and iron-reducing and methanogenic conditions promote the degradation of similar
56 compounds (Chang et al., 2002; Chang et al., 2006; Meckenstock et al., 2004; Sayara et al.,
57 2010). It is worth mentioning that a gradient of contaminant concentrations are found in
58 different contaminated matrices, and these concentrations vary from low, such as wastewater
59 sludge ($\mu\text{g}/\text{kg}$, dry matter (DM)), to high, such as soils or sediments (g/kg , DM) (Chang et al.,
60 2003; Chang et al., 2008; Christensen et al., 2004; Trably et al., 2003).

61 Anaerobic digestion under methanogenic conditions can be carried out under different
62 temperature including mesophilic (37°C) and thermophilic (55°C) regimes (De Baere, 2000).
63 However, mesophilic treatments are considered more economical because less energy is
64 required. As a result, most anaerobic full-scale plants work under mesophilic conditions
65 (Ferrer et al., 2008) even though thermophilic treatments have a higher biogas production rate
66 because of the associated faster reaction rates (Cecchi et al., 1991). Accordingly, the study of

67 these two different treatments to gain a full understanding of the mechanisms and behavior of
68 this process is recognized as an alternative source for renewable and clean energy.

69 In a normal start-up of a batch anaerobic digester a certain amount of inoculum needs
70 to be added together with the substrate to be digested to generate the required conditions and
71 microorganisms needed to start the biological reactions (Liu et al., 2009). Theoretically, the
72 degradation of organic matter including PAHs can be promoted by increasing the density of
73 microbial activity either by favorable conditions or by increasing the amount of
74 microorganisms present. Equally, the adaptability of the introduced microorganisms is also
75 crucial for promoting the digestion process (Fernández et al., 2005). Consequently, the
76 determination of the ratio between the treated material and inoculum is of great importance. In
77 this regard, the minimum amount of inoculum required to provide sufficient activity with the
78 maximum load of organic material is considered to be crucial for process design.

79 The role of anaerobic degradation for PAH-contaminated soil is an area in which little
80 work has been reported. To our knowledge, there is a lack of information about the
81 bioremediation of PAH-contaminated soil under strict methanogenic-anaerobic conditions. To
82 date, there have been no reports of the behavior of such processes under various temperatures,
83 and no information is available about the influence of the main factors that typically affect the
84 process such as contaminant concentration and soil to inocula ratio.

85 Hence, the objective of the present study was to comparatively assess the process
86 performance under different operating conditions by using two different types of inoculum
87 requiring different temperature: one thermophilic and one mesophilic. Moreover, two main
88 controlling factors such as the PAH concentration and the soil to inocula ratio were evaluated
89 to assess their effect on the PAH degradation rate. The biogas production was measured as an
90 indicator of process activity, whereas PAH degradation was evaluated by measuring the
91 concentration decrease and the production of biodegradation metabolites using advanced
92 techniques. Also, phospholipid fatty acid (PLFA) analysis was used to determine the
93 evolution of the microbial communities present in each sample. In both the thermophilic and
94 mesophilic case the experiments were systematically carried out by using an experimental
95 design technique considering the different levels of the studied factors, to allow for statistical
96 validation of the results obtained.

97

98 **2. Experimental**

99

100 *2.1. Soil*

101 The soil used in this study was an uncontaminated soil classified as sandy loam soil. It
102 was collected from the surface horizon (0–30 cm) of an agricultural field located in Prades
103 (Tarragona, Spain). The soil was air-dried and then sieved (2 mm) to remove any debris and
104 kept at 4°C until use. The soil texture consisted of 73.4% sand, 18.6% silt and 8% clay. No
105 PAHs were detected in the soil before being used for experimental purposes. Other properties
106 of the soil are shown in Table 1.

107

108 2.2. Contaminants

109 Fluorene, Phenanthrene, Anthracene, Fluoranthene and Pyrene, all of 98–99% purity,
110 were selected as contaminants in order to evaluate their degradation under anaerobic
111 methanogenic conditions. These compounds were obtained from Sigma-Aldrich (Barcelona,
112 Spain). These five PAHs are included on the list of the 16 USEPA priority pollutants. The
113 weight percentage of each compound as part of the total PAHs concentration (Σ PAHs) was
114 33, 31, 10, 22 and 4%, respectively. These percentages were chosen to simulate a real
115 creosote sample the percentages of which were determined following fractionation of a
116 commercial creosote sample (Creosote lot: 42-13B, Chem Service, Sugelabor, Spain) in our
117 laboratory by using the method 3611B of the USEPA, in which the volatile part was ignored.
118 The PAHs were mixed in the appropriate proportions to create a stock solution. This stock
119 solution was used to spike the soil to obtain the desired concentration according to the
120 experimental design range (0.1–2 g/kg, DM) measured as the total concentration of the five
121 PAHs.

122

123 2.3. Anaerobic inocula

124 In this study, two types of inoculum with different optimal temperatures, thermophilic
125 (55°C) and mesophilic (37°C), were used. The mesophilic-anaerobic inocula were obtained
126 from the digested effluent of an anaerobic digester after the solid-liquid separation in a waste
127 treatment plant (Barcelona, Spain). The inoculum contained 10.02% of total solids, which is
128 composed of the entire microbial consortium involved in the anaerobic digestion process,
129 together with some difficult to degrade or inert solids. The thermophilic inoculum was
130 obtained from a thermophilic treatment plant (Terrassa, Spain). The thermophilic inoculum
131 was filtered through a 4 mm sieve to remove some big parts of the feedstock and 11.8% total
132 solids remained. In both cases the treatment plants are fed with the source-selected organic
133 fraction of municipal solid wastes. The two inocula were kept separately in plastic containers
134 with a gallon capacity under strict anaerobic conditions and incubated in water baths at 37°C

135 and 55°C for about two weeks before use to remove any biodegradable materials. No
136 contamination by PAHs was detected in the inocula prior to the experiments. Other
137 characteristics of the inocula are presented in Table 1.

138

139 *2.4. Experimental system*

140 Experiments with both inocula were performed in triplicate by using sealed 1-L bottles
141 (Traveller SIGG[®], Spain) as reactors. They were tightly sealed and equipped with a ball valve
142 that can be connected to a digital manometer to determine the pressure of biogas produced.
143 The contaminants were prepared in a stock solution (5 g total PAHs in 500 mL
144 dichloromethane) and used to spike the soil at different concentrations (0.1–2 g/kg total
145 PAHs; approximately 300–6000 mg chemical oxygen demand (COD)/L) determined
146 according to the experimental design technique. The solvent was allowed to evaporate at
147 room temperature. Next, the soil and inoculum were mixed together based on the dry weight
148 fraction at different ratios ranging from 0.2:1 to 5:1 (w/w, soil/inoculum). As inoculum and
149 soil are different in their total solids content, distilled water was added to the samples to
150 correct for this and ensure the same total solids content for both. Thus, all samples had the
151 same amount of total solids equal to that corresponding in each inoculum (11.8 and 10.02%).
152 All the bottles were then tightly sealed and purged several times with compressed nitrogen
153 gas to ensure anaerobic conditions (Ponsá et al., 2011) before being incubated under strict
154 anaerobic static conditions. The mesophilic samples were incubated in a temperature-
155 controlled chamber at 37°C, and the thermophilic samples were incubated in an oven adjusted
156 to 55°C. The total experimental time was set to 50 days in line with previous experiments
157 (Sayara et al., 2010). After 30 days one sample from each set of experimental conditions was
158 removed from the incubator to evaluate the process behavior. The remaining samples (in
159 duplicate) were left for the full 50 days. A blank experiment (in triplicate) with only inoculum
160 was used to evaluate the baseline of biogas production by an endogenous matter. These
161 samples did not contain any additional nutrients or organic matter. All the results are
162 expressed as the average measured after 30 and 50 days.

163

164 *2.5. Analytical methods*

165

166 *2.5.1. Determination of biogas and analysis*

167 The production of biogas was followed and determined quantitatively by measuring
168 the pressure increase in the headspace of the sample bottle by means of a SMC (ISE30)

169 Pressure Switch manometer (1 MPa, 5% accuracy). The quantity of the produced biogas in
170 each sample was determined by subtracting the biogas produced by the inoculum (blank
171 experiment) from that produced in the sample. Because it is crucial to be able to accurately
172 determine the differences between the production of biogas from the inoculum and the PAH
173 experiments, the inoculum was maintained under anaerobic conditions for 15 days prior to use
174 to remove the biodegradable organic matter and to reduce biogas production owing to
175 endogenous activity (Ponsá et al., 2011). The net value of biogas produced was calculated and
176 expressed under normal conditions (0°C and 1 atm). Biogas analysis, particularly the CH₄ and
177 CO₂ contents were determined by using a gas chromatograph (GC 5890 Capillary Hewlett
178 Packard) as described by Sayara et al. (2010).

179

180 2.5.2. PAHs analysis

181 To determine the concentration of PAHs after 30 and 50 days of incubation, the
182 contents of the sample bottles were dried by using a lyophilizer (Benchtop 5L, Virtis Sentry,
183 NY). Lyophilized samples were extracted by using an ASE 200 System (Dionex, Voisins-le-
184 Bretonneux, France). The extraction cell (11 mL) was loaded in an oven and extracted with
185 hexane-acetone (3:1, v/v). The cell was heated (150 °C, 7 min) and subsequently extraction
186 took place at 150 °C under 103.4 bar for 5 min. The cell was then flushed with fresh solvent
187 (60% of total cell volume) and finally the solvent was purged from the cell with nitrogen gas
188 for 60 seconds. For each sample the extraction cycle was performed three times. The resulting
189 organic extracts were collected in 40 mL vials, dried under vacuum at room temperature and
190 finally dissolved in acetonitrile for subsequent analysis. Reverse phase high-performance
191 liquid chromatography analyses were performed by using a system consisting of a 2695
192 Separations Module (Waters, Milford, MA) equipped with a LiChroCart column filled with
193 LiChrospher[®] PAH (250 x 4 mm; particle size 5 µm; pore size 150 Å; Merck, Darmstadt,
194 Germany), a 2996 diode-array detector and 2475 fluorescent detector (Waters, Spain).
195 Separation of the PAHs was achieved with a gradient program, by using a mixture of
196 methanol:acetonitrile (1:1 v/v, solution A) and Milli-Q water. After 5 min of isocratic elution
197 with 70% of solution A, the eluent was changed slowly to be 100% solution A over 15 min
198 and maintained constant for the following 20 min. PAHs were identified by using UV
199 spectrophotometry and comparison of retention times to commercially available standards
200 (Dr. Ehrenstorfer, Augsburg, Germany). The concentrations of the five PAHs were calculated
201 according to the USEPA method 610. The compounds were quantified with the fluorescent
202 detector under excitation/emission conditions: phenanthrene and anthracene at -250/390 nm;

203 fluorene, fluoranthene and pyrene at –280/340 nm. Calibration curves with the standards were
204 prepared over a linear range (0.1–10 µg/ml) for each compound, and the recovery rate was
205 approximately 95%.

206

207 2.5.3. Metabolite identification

208 Samples that had been previously extracted (See section 2.5.2) were analyzed by using
209 gas chromatography (Agilent HP 6890 Series II) coupled to a mass selective detector under
210 electronic impact ionization (Agilent HP 5973) with a HP5-MS (30 m x 0.25 mm x 0.25 µm;
211 Agilent, Spain). The operating conditions of the chromatograph were as follows: injector
212 (splitless 1 min) 320°C, injection volume 1–3 µl (depending on the sample), oven
213 temperature: 50°C (1 min), rate 7°C/min, final temperature 320°C with helium as carrier gas
214 at 0.7 ml/min. The detector remained in solvent delay mode for 3.2 min and the mass range
215 measured was 40–400 (m/z). The products detected were identified by comparing the mass
216 spectral data to the Wiley 7® library.

217

218 2.5.4. Volatile fatty acids

219 The total volatile fatty acid composition (acetate, propionate, *iso*-butyrate, *n*-butyrate,
220 *iso*-valerate, *n*-valerate; (g/L)) was determined by using gas chromatography (GC). Samples
221 were centrifuged for 10 minutes (10⁴ rpm) then filtered through to remove any solid particle
222 (0.25 µm). Known volumes of the filtrates were mixed equally (1:1, v/v) with pivalic acid as
223 standard, before analysis by GC. Samples (1 µl) were injected into the GC (GC 5890
224 Capillary Hewlett Packard) equipped with a flame ionization detector (280°C) and a splitless
225 injector (260°C). A HP-Innowax column (Crosslined polyethylene Glycol; 30 m x 0.53 mm
226 x1 µm; Agilent) was used. The temperature was maintained at 80°C for 1 min before
227 increasing at a rate of 5°C/min until 150°C and then at 20°C/min until 230°C was attained.

228

229 2.5.5. Evaluation of the bioavailability of PAHs

230 The bioavailable fractions of PAHs were estimated by using sequential supercritical
231 fluid extraction (SFE) (Cajthaml and Šašek, 2005). The extraction was performed with a
232 PrepMaster extractor (Suprex, Pittsburgh, USA) equipped with a VaryFlow restrictor
233 operating at 40°C with a downward stream of carbon dioxide (5.5 SFE/SFC, Supercritical
234 fluid chromatography and extraction system, Messer Technogas, Prague, Czech Republic).
235 The samples (1 g, dry basis) were extracted at 50 °C and 200 bar. Each extraction was carried
236 out in duplicate and the compounds were collected after 5, 10, 20, 40, 60, 80, 120, 160, and

237 200 min. SFE represents a desorption model that generally assumes that the extraction is
238 controlled by two rate constants differing in orders of magnitude (Williamson et al., 1998).
239 The “F fraction”, which represents the rapidly desorbed fraction of the target chemical from
240 soil, is usually assumed to be representative of equilibrium release conditions, and the next,
241 more slowly released, portion is considered to be kinetically rate-limited. The chemical
242 release data is then modeled by an empirical two-site model, consisting of two first-order
243 equations:

$$244 S_t = F S_0 \exp(-k_1 t) + (1 - F) S_0 \exp(-k_2 t) \quad (1)$$

245 Where S_t is the pollutant concentration remaining in the soil after time t ; F is the fraction of
246 chemicals that is rapidly released; S_0 is the original concentration of the pollutant in soil; k_1
247 and k_2 are the first order rate constants. Prism version 5.0 (GraphPad, USA) was used to
248 calculate the F values according to Equation 1.

249

250 2.5.7. *Quantification of microbial biomass*

251 Samples for PLFA analysis were extracted by using a mixture of chloroform,
252 methanol and phosphate buffer (1:2:0.8; v/v/v). LiChrolut Si-60 solid-phase extraction
253 cartridges (Merck, Whitehouse Station, NJ) were used for the separation of the extracts and
254 phospholipid fractions were subjected to mild alkaline methanolysis (Šnajdr et al., 2008). Gas
255 chromatography-mass spectrometry (GC-MS) was used for analysis of the free methyl esters
256 of phospholipid fatty acids (450-GC, 240-MS ion trap detector, Varian, Walnut Creek, USA).
257 The GC instrument was equipped with a split/splitless injector and a DB-5MS column (J&W
258 Scientific, Folstom, 60 m, 0.25 mm, 0.25 μ m film thickness) was used for separation. The
259 temperature program started at 60 °C and was maintained for 1 min in splitless mode. Then
260 the splitter was opened and the oven heated to 160 °C at a rate of 25 °C min⁻¹. The second
261 temperature ramp reached 280 °C at a rate of 2.5 °C min⁻¹ and this temperature was
262 maintained for 10 min. The solvent delay time was set to 8 min. The transfer line temperature
263 was set to 280 °C. Mass spectral data were recorded under electron impact at 70 eV, mass
264 range 50–350 amu. Methylated fatty acids were identified according to their mass spectra and
265 quantified by using their individual chemical standards (Sigma-Aldrich, Prague, Czech
266 Republic and Matreya LLC, Pleasant Gap, USA). Fungal biomass was quantified based on
267 18:2 ω 6,9 content, bacterial biomass was quantified as a sum of i14:0, i15:0, a15:0, 15:0,
268 i16:0, 16:1 ω 7, 16:1 ω 9, 16:1 ω 5, 10Me-16:0, i17:0, a17:0, cy17:0, 17:0, 10Me-17:0, 18:1 ω 7,
269 18:1 ω 9, 10Me-18:0 and cy19:0. Biomass of Gram positive and Gram negative bacteria were
270 estimated by using concentrations of i14:0, i15:0, a15:0, 15:0, i16:0 and 16:1 ω 7, 18:1 ω 7, 16:1

271 $\omega 5$, cy19:0 cy17:0, respectively. To evaluate the anaerobic PLFA markers the quantities of
 272 cy19:0, cy17:0 and 18:1 ω 9 were used (Oravec et al., 2004; Sampedro et al., 2009; Šnajdr et
 273 al., 2008).

274

275 *2.6. Experimental design and statistical analysis*

276 The effect of two factors ($k = 2$): the PAHs concentration (x_1) and the soil to inocula
 277 mixing ratio (x_2), on the bioremediation of PAHs-contaminated soil under anaerobic
 278 conditions were systematically studied by using the central composite design (CCD). More
 279 details about the experimental design technique and its application can be found and reviewed
 280 elsewhere (Deming and Morgan, 1987; Sayara et al., 2010). The design matrix is presented in
 281 Table 2, in which the coded and actual values of the two independent variables (x_1 and x_2) and
 282 the actual response of each combination regarding the biodegradation percentage (Y_D) and the
 283 biogas production (Y_G) are reported. In total, nine experiments (in triplicate) were carried out
 284 according to CCD principles. Control samples, containing only the corresponding inoculum
 285 were also carried out. Statistical analysis was performed for all the variables by using the
 286 Sigmaplot® 8.0 software package (Systat Software Inc, San José, USA) and according to the
 287 statistics recommended for CCD (Deming and Morgan, 1987).

288

289 **3. Results and discussion**

290

291 *3.1. Response surface and statistical analysis*

292 CCD was applied to study the process behavior under different variables. The
 293 degradation (D) percentage (%) and the biogas (G) production ($L\ kg^{-1}\ TS$) after 30 and 50
 294 days were selected as functions to be optimized along with their parameters to represent the
 295 response of the process. Second order models for each variable are illustrated in Equations 2–
 296 9 under thermophilic (T) and mesophilic (M) conditions:

297

298 $Y_{TD30} = 51.9 + 23.28x_1 - 6.69x_1^2 - 0.69x_2^2 - 1.9x_1x_2 \dots\dots\dots(2)$

299 $Y_{TD50} = 32.14 + 25.09x_1 + 6.62x_2 - 5.8x_1^2 - 2.18x_1x_2 \dots\dots\dots(3)$

300 $Y_{TG30} = 3.62x_1 - 7.03x_2 + 3.1x_1^2 + 1.7x_2^2 - 2.4x_1x_2 \dots\dots\dots(4)$

301 $Y_{TG50} = 8.18 + 7.4x_1 - 11.5x_2 - 0.75x_1^2 + 1.9x_2^2 \dots\dots\dots(5)$

302 $Y_{MD30} = 42.7 + 46.18x_1 - 13.23x_1^2 + 0.21x_2^2 - 1.36x_1x_2 \dots\dots\dots(6)$

303 $Y_{MD50} = 20.95 + 68.18x_1 - 22.77x_1^2 + 0.39x_2^2 \dots \dots \dots (7)$

304 $Y_{MG30} = -135.5x_1 + 6.3x_2 + 50.7x_1^2 + 8.7x_1x_2 \dots \dots \dots (8)$

305 $Y_{MG50} = -96.55x_1 + 6.02x_2 + 35.82x_1^2 + 5.6x_1x_2 \dots \dots \dots (9)$

306

307 The correlation coefficients (R) and p values of the functions representing the degradation
308 percentage for both inocula after 50 days are shown in Table 3. The regression model that was
309 selected is considered to be adequate to describe the data and the relationship between the
310 independent and dependent variables. Conversely, the correlation coefficients and p values
311 obtained from the rest of the treatment models designed to represent the biodegradation
312 percentage and the biogas production were not favorable (data not shown). It is proposed that
313 the low organic matter content and the presence of the contaminants influenced the biogas
314 production. However, the values obtained from the statistical analysis of these samples do
315 give a general idea about the process performance when such conditions occur, especially in
316 the case of biogas production/inhibition owing to the presence of PAHs.

317

318 *3.2. Overall degradation of the contaminants*

319 The anaerobic degradation of the PAHs under the different conditions studied was
320 evaluated after 30 and 50 days incubation to monitor the process. Fig. 1 shows the results
321 obtained under thermophilic and mesophilic conditions. After 30 days, all samples showed
322 some degradation and the same trend under the two temperature conditions was observed. The
323 biodegradation observed seems to demonstrate that the native microorganisms of the
324 inoculum (thermophilic or mesophilic) have the catabolic capacity to degrade the PAHs used,
325 and it could be an indication that the methanogenic metabolism becomes coupled to the
326 anaerobic degradation of the PAHs, as observed in other studies (Chang et al., 2006; Yuan
327 and Chang, 2007). However, the samples containing low concentrations of PAHs (0.1g/kg)
328 resulted in the lowest degradation rate, suggesting that microbial activity is governed by the
329 pollutant concentration in which a minimum threshold level is required to initiate degradation
330 and catabolic induction in biodegrading microorganisms (Boethling and Alexander, 1979). It
331 is also worth noting that, at low concentrations, more surface area on the soil particles is
332 available for adsorption of the contaminant relative to high concentrations. Therefore, a
333 stronger interaction between the soil and contaminant could be formed making the
334 accessibility and biodegradation of the contaminant complex (Table 4) in which, for the most
335 part, the “F” fraction at low concentrations is lower than at the corresponding high

336 concentrations. In addition, a part of the biodegradation could be related to the soil's
337 endogenous microorganisms because the used soil was not sterilized.

338 In general, degradation capacities under mesophilic conditions were higher than those
339 obtained under thermophilic conditions. However, thermophilic conditions resulted in better
340 rates of biodegradation when low concentrations of PAHs were used. This phenomenon can
341 be explained by high temperatures, which, in most cases, increase the desorption of PAHs and
342 their solubility, as observed in the determination of the bioavailable portion of the
343 contaminants, relative to the corresponding mesophilic conditions (Table 4). Moreover, under
344 thermophilic temperatures the mass transfer and reaction kinetics are enhanced. Nevertheless,
345 other studies have reported contradictory results. For instance, Chang et al. (2002) found that
346 a temperature of 30°C enhanced PAH degradation better than at 40°C.

347 Under the different combinations and conditions, and according to the obtained results
348 regarding the contaminants bioavailability (Table 4), the biodegradation of the contaminants
349 was controlled by their characteristics such as water solubility, number of benzene rings and
350 structural conformation (shape) of the molecules. However, no reliable trend or order in
351 biodegradation is apparent.

352 Analysis of selected components of PLFA demonstrated that anaerobic bacteria were
353 present in all samples, but their abundance in the thermophilic samples was higher than in the
354 mesophilic ones (Table 5). Furthermore, moderate to high PAH concentrations significantly
355 ($p < 0.05$) altered the microbial community structure by changing anaerobic communities in
356 such a way that more anaerobic microorganisms were observed at high PAH concentrations
357 under both conditions, and better degradation rates of PAHs were consequently achieved. The
358 PLFA profiles were in agreement with the rates of PAH degradation (Fig.1) under both
359 conditions with less degradation observed at low PAH concentrations. According to Donald et
360 al. (1998), the microbial community responded to PAHs contamination at both the phenotypic
361 and genotypic levels, which is also in agreement with the obtained results.

362 As expected for biological treatments of PAHs, their concentration decreases with
363 time as a result of the microbial activity, or no change occurs in the case of the process failure
364 or absence of optimal conditions for microbial activity. However, in this study the results
365 were completely contradictory to what was expected. For the majority of samples under both
366 conditions (37 and 55°C), it was found that the PAH concentrations after 50 days of
367 incubation were higher than those obtained after 30 days. We therefore propose the following
368 hypotheses: a) sorption or occlusion of the contaminants (PAHs) in the organic matter during
369 the first 30 days and release after depletion of biodegradable organic matter, b) new PAHs

370 bioformed under these conditions. To examine the potential impact of sorption on the results,
371 representative samples (in duplicate) were extracted twice under extreme conditions: high
372 temperature (200°C), pressure (206.84 bar) and 4 static cycles (10 min). The same samples
373 were also extracted by using a Soxhlet and analyzed as explained elsewhere (Sayara et al.,
374 2010). The extraction and analysis methods resulted in values that corroborated an increase in
375 the PAH concentration. Indeed, Thiele et al. (2002) demonstrated that PAHs were formed in
376 soil under oxygen deficient conditions. In that study, the concentration of high molecular
377 weight PAHs (more than 3 aromatic rings) increased considerably, whereas for compounds
378 with three rings the concentration decreased, probably owing to anaerobic biodegradation.
379 Furthermore, the incubation of edible oil in closed containers at room temperature led to the
380 formation of PAHs (Guillen et al., 2008) indicating that anaerobic conditions are likely
381 responsible for such a phenomenon.

382

383 *3.3. Metabolites*

384 The observed initial decreases in the concentration of PAHs show that the inocula
385 were able to degrade the target contaminants. Anaerobic biodegradation of PAHs was also
386 monitored by analyzing the metabolites produced as a result of microbial activity in the
387 samples (Table 5). In fact, several compounds were identified by GC-MS. These compounds
388 included: phenol, hydroxyfluorene, benzeneacetic acid, cresol, pyridine, fluorenone,
389 fluorenol, carbazole, anthracenone, anthracenedione, biphenyl and anthrone. These
390 compounds are known metabolites generated when PAHs are degraded by microorganisms
391 through several oxidization pathways. Consequently, the metabolites found in this study
392 provide evidence of the microbial capacity to degrade PAHs under the conditions studied.
393 PAH degradation pathways under anaerobic conditions are still not clear, and various
394 hypothesis or arguments are found in the literature. Meckenstock et al. (2004 and Zhang and
395 Young (1997) argued that carboxylation is the initial step in PAH biotransformation under
396 sulphate reducing conditions. On the contrary, Bedessem et al. (1997) proposed hydroxylation
397 as the initial step in PAH biotransformation under the same conditions. Vogel and Grbic-
398 Galic (1986) argued that the anaerobic transformation of benzene and toluene to CO₂ and CH₄
399 occurred through hydroxylation, when phenol and cresol were identified as intermediates of
400 benzene and toluene, respectively. Because these compounds have been identified in the
401 present study, it would be possible to assume that hydroxylation was the initial step occurring
402 during the biotransformation of PAHs in this case. Most likely, PAHs undergo initial ring
403 reduction followed by hydrolytic ring cleavage to yield aliphatic acids for cell growth.

404 Moreover, naphthalenol was detected as metabolite from naphthalene biotransformation in
405 sulfidogenic sediments (Bedessem et al., 1997). All of these observations support the
406 hypothesis that PAHs degrade under methanogenic conditions and hydroxylation is the initial
407 step in their biotransformation. Unfortunately, there are no studies that investigate PAH
408 biodegradation under anaerobic conditions, at thermophilic temperatures.

409

410 *3.4. Biogas production*

411 The cumulative biogas production under thermophilic and mesophilic conditions is
412 shown in Fig. 2. It is clear that the methanogenic inocula under both temperature conditions
413 were exposed to inhibition effects because the PAHs were detrimental for the
414 microorganisms. Usually, under similar conditions, an adapted ecosystem is used to avoid this
415 inhibition. Consequently, as shown in Fig 2, an adaptation period of at least 20 days was
416 needed in most cases except for Runs 2 and 8 under thermophilic conditions. The inhibition
417 rate (negative values indicate that biogas production in the control samples is higher than in
418 the samples containing PAHs) gradually increased with time indicating an increase in toxicity
419 of the samples. However, in most samples, after 20 days' incubation, biogas production
420 changed and inhibition decreased over time, although not completely eliminated in the
421 mesophilic samples. In addition, process recovery was noted with prolonged incubation times
422 and methane production was observed and increased significantly following the inhibition
423 stage (data not shown). This observation demonstrates that both of the inocula were not
424 previously exposed to similar contaminants and agree with the preliminary analysis that
425 showed that they were free from PAHs. Interestingly, the study clearly indicated that although
426 the presence of PAHs considerably inhibits biogas production, PAH degradation is not
427 affected. Indeed, Fuchedzhieva et al. (2008) reported similar behavior for fluoranthene
428 biodegradation, which continued despite cell growth inhibition in the presence of a
429 biosurfactant complex. In general, better results were obtained for the samples treated under
430 thermophilic conditions relative to mesophilic ones, because the higher temperatures facilitate
431 the reaction rates and enhance enzymatic activity. However, microbial activity can be
432 hampered by the presence of toxic metabolites resulting from degradation of the parent
433 compounds. Indeed, analysis showed high rates of toxicity in all samples relative to a non-
434 toxic control (data not shown). This toxicity is likely owing to the PAHs themselves and their
435 metabolites as a result of microbial biodegradation. Additionally, the ratios of saturated to
436 monounsaturated PLFAs (S/M) and the ratios of cyclopropyl PLFAs to their monoenoic
437 precursors (cy/pre) (Table 5) were significantly higher in the mesophilic samples relative to

438 the thermophilic ones, indicating microbial community stress under such conditions (Moore-
439 Kucera and Dick, 2008). However, analysis of volatile fatty acids showed that there is no
440 inhibition caused by such acids because they were not detected in any sample after 30 or 50
441 days. The pH values of the sample mixtures after 30 and 50 days were within 7.3–7.9, which
442 is around the optimal values for anaerobic digestion. Therefore, the fluctuation of biogas
443 production observed was more likely owing to some PAH metabolites that were formed
444 transiently during their anaerobic biodegradation. The oscillation in the biogas production
445 (Fig. 1, thermophilic run 8) clearly shows this effect.

446

447 *3.5. Process response for PAH degradation*

448 The variations in PAH degradation (%) under the conditions studied factors are shown
449 in Fig. 3 and Fig. 4 for thermophilic and mesophilic inoculum, respectively. Under mesophilic
450 conditions, it is clear that the degradation capacity is always proportional to PAH
451 concentration, whereas the soil to inocula mixing ratio was found to influence PAH
452 degradation at low concentrations (less than 1 g/kg). For instance, low (0.2:1) or high (5:1)
453 mixing ratios enhanced the degradation rate, but intermediate ratios negatively affected the
454 process. The same behavior was observed under thermophilic conditions during the first 30
455 days, although PAH concentration influenced the degradation capacities to a smaller extent
456 (Fig. 3A). Nevertheless, after 50 days the degradation capacity response demonstrated that the
457 process is proportional to both factors, with the highest degradation (75%) being obtained at
458 the highest concentration and mixing ratio. The increased toxic potential of higher PAH
459 concentrations on the inocula can explain why degradation decreases as inoculum ratio
460 increases. Similar results were seen with the increase of the inoculum ratio under anaerobic
461 conditions (Chang et al., 2002).

462 In both cases the PAH degradation was clearly proportional to the soil to inocula
463 mixing ratio and the best results were obtained when using the highest ratio (5:1). These
464 results are in accordance with those obtained by Hernandez et al. (2008) who reported that
465 increasing the concentration of substituted phenolic compounds enhanced the anaerobic
466 digestion process, but that at higher concentration the total biodegradation decreased. The
467 sequential decrease in biogas production (Fig. 2) observed during the first three weeks
468 demonstrates the difficulties the microflora had to overcome to adapt their biodegradation
469 potential. Here, time was an essential factor to allow for effective microbial adaptation under
470 these conditions. Although PAHs and their metabolites had an inhibitory effect on the
471 methanogenic communities, other anaerobic microbes were able to continue the degradation

472 process. Interestingly, PAH degradation increased when a high soil to inocula ratio was used
473 for both cases. This observation is fundamental when considering an economical evaluation
474 because larger quantities of contaminated soil could be treated with smaller quantities of
475 inoculum. Because no other nutrient sources were available to the microorganisms, it is likely
476 that the organic matter of the soil served as the nutrient source, and that, because the organic
477 matter content was low, this nutrient limitation motivated the microorganisms to use PAHs.
478 However, the addition of an organic matter source, such as compost, could stimulate
479 microbial activity and, consequently, accelerate PAH degradation. This effect was clearly
480 demonstrated in previous studies (Darush et al., 2006; Sayara et al., 2010).

481

482 *3.6. Process response for biogas production*

483 The response of biogas production under thermophilic and mesophilic conditions is
484 shown in Fig. 5. The value for the biogas produced by a sample was calculated by subtracting
485 the value obtained from the control experiment (only inoculum present) from the value
486 obtained from the contaminated soil sample. The negative values indicate that the background
487 biogas production was higher than for the contaminated samples owing to inhibition. The
488 process behavior was completely different under thermophilic and mesophilic conditions,
489 with no biogas being produced under mesophilic conditions, whereas certain amounts were
490 produced under thermophilic conditions in several samples. A soil to inoculum mixing ratio
491 of almost 3:1 was the inflection point in relation to the applied concentrations (0.1–2 g/kg). A
492 soil to inocula ratio higher than 3:1 was gently influenced by the variation in PAH
493 concentration. The biogas production rate slightly increased with increasing PAH
494 concentration under thermophilic conditions, whereas no variation was seen with mesophilic
495 inocula (stagnation state). At mixing ratios below 3:1, different behaviors were observed
496 demonstrating various interactions between the studied variables within the two temperatures
497 domains. Up to a point (almost 1.4g /kg), increasing the concentration of PAHs enhanced the
498 biogas production rate under thermophilic conditions. Thereafter, further increases in PAH
499 concentration negatively affected the microbial activity because of their toxic properties.
500 Moreover, biogas production was enhanced at low soil to inoculum ratios when additional
501 inoculum was introduced. Nevertheless, increasing inoculum quantity negatively affected the
502 biogas production under low PAH conditions (<0.2g/kg). Under mesophilic conditions, the
503 rate of inhibition gradually decreased as the soil to inocula ratio increased until a ratio of 3:1
504 was achieved. Above this ratio, the rate of inhibition plateaued. PAH concentrations below
505 plateau conditions (mixing ratio less than 3:1) influenced the process in two different ways:

506 an increase in PAH concentration was followed by an increase in the inhibition rate until a
507 certain concentration was reached (almost 1g/kg) after which the process changed its behavior
508 and a decrease in inhibition was observed.

509

510 **4. Conclusions**

511

512 A comparative study on the anaerobic bioremediation of PAH-contaminated soil under
513 methanogenic conditions, at thermophilic and mesophilic temperatures, has been conducted.
514 This study indicates that inhibition of biogas production is related to high concentrations of
515 PAHs. Although a high rate of biodegradation was achieved during 30 days, a prolonged
516 incubation time for both thermophilic and mesophilic samples showed an increase in the PAH
517 concentration. This increase was probably caused by the reverse bioformation of these
518 compounds. Further research is necessary to clarify this effect and to determine exactly the
519 optimal times required for biodegradation.

520

521 **Acknowledgments**

522 Financial support was provided by the Spanish Ministerio de Educación y Ciencia (Project
523 CTM2009-14073-C02-01) and by the Czech Science Foundation (Grant No. 525/09/1058).

524

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645

646 **Figure captions**

647

648 **Fig. 1.** Percentage (%) of PAHs remaining after 30 and 50 days incubation. (A) Thermophilic
649 conditions and (B) mesophilic conditions

650 **Fig. 2.** The cumulative biogas production (L/kg TS) after 50 days. (A) Thermophilic
651 conditions and (B) mesophilic conditions

652 **Fig. 3.** Response of PAHs degradation (%) under thermophilic conditions. (A) After 30 days
653 and (B) after 50 days

654 **Fig. 4.** Response of PAHs degradation (%) under mesophilic conditions. (A) After 30 days
655 and (B) after 50 days

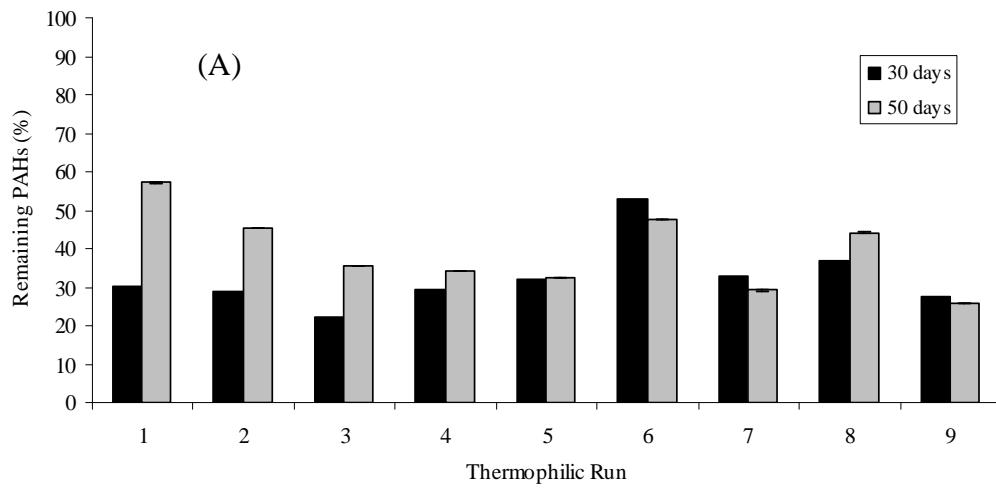
656 **Fig. 5.** Response of the total biogas production (L/kg TS) after 50 days. (A) Thermophilic
657 conditions and (B) mesophilic conditions

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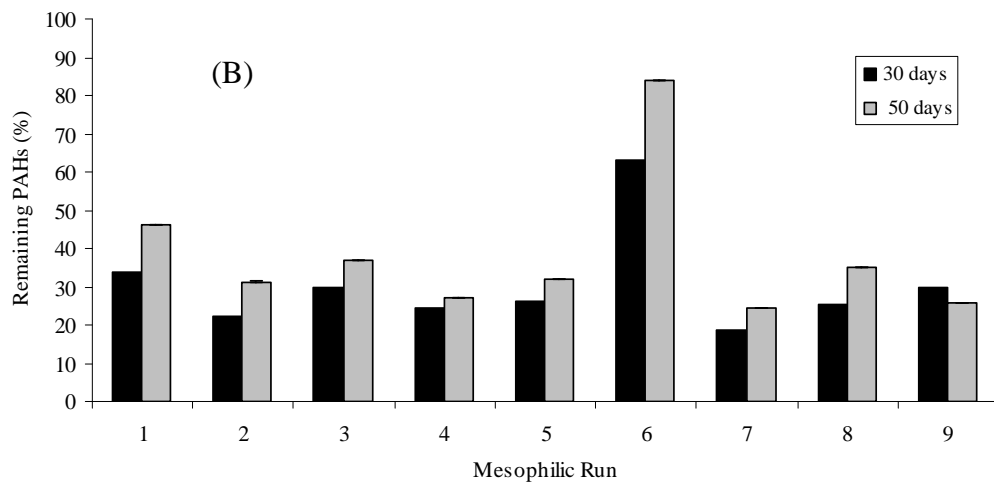
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660 **Fig. 1**

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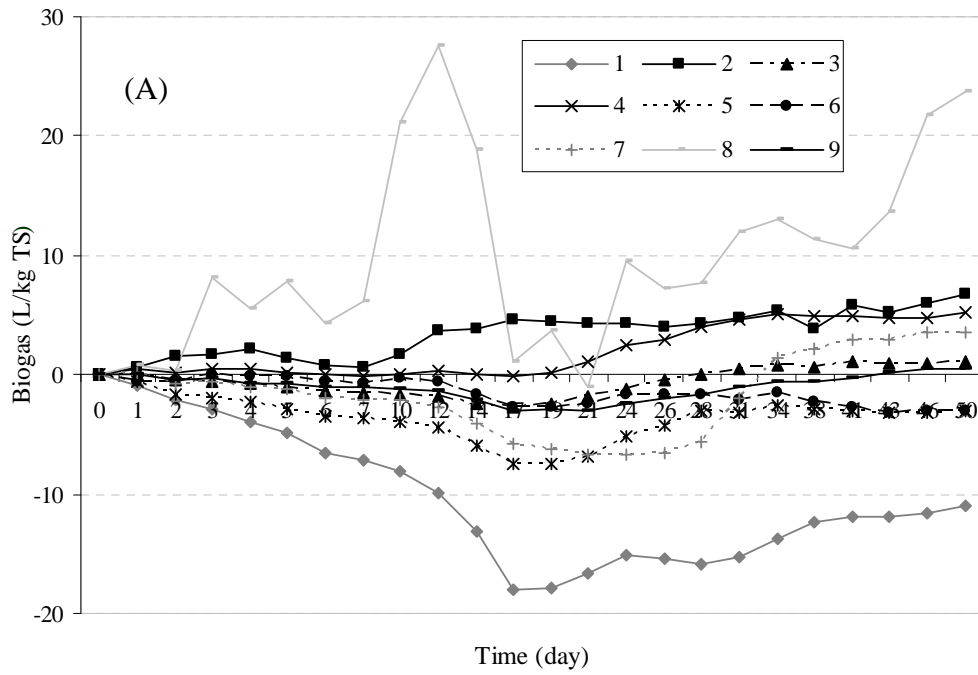
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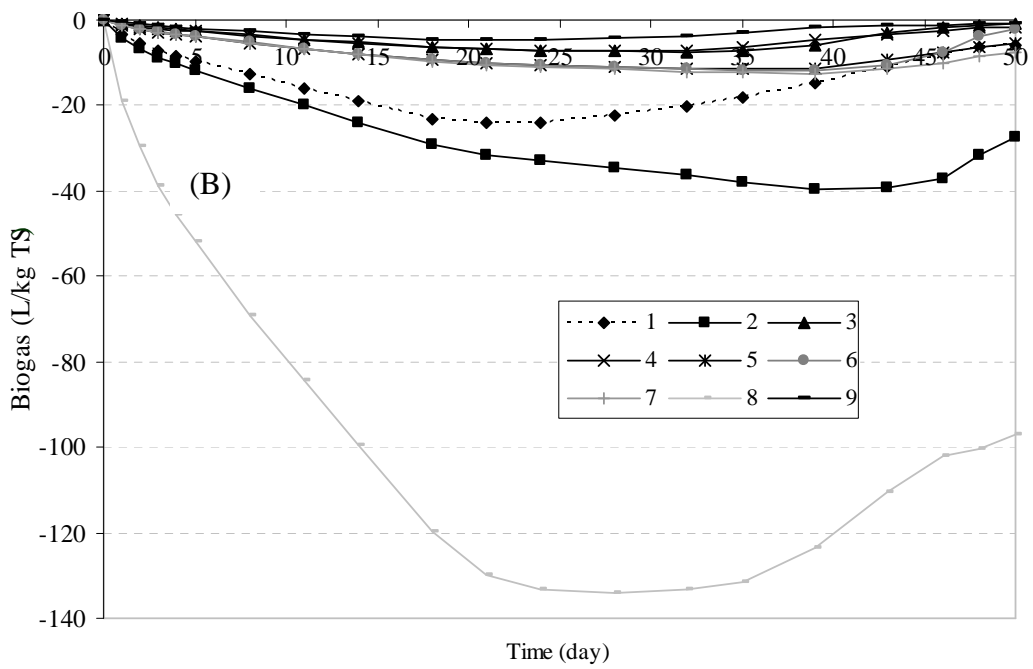
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667 **Fig. 2**

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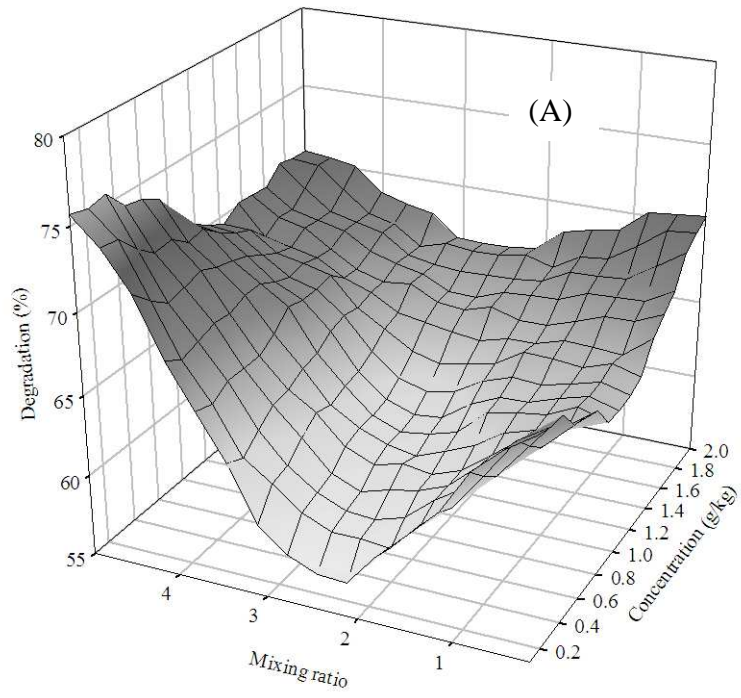
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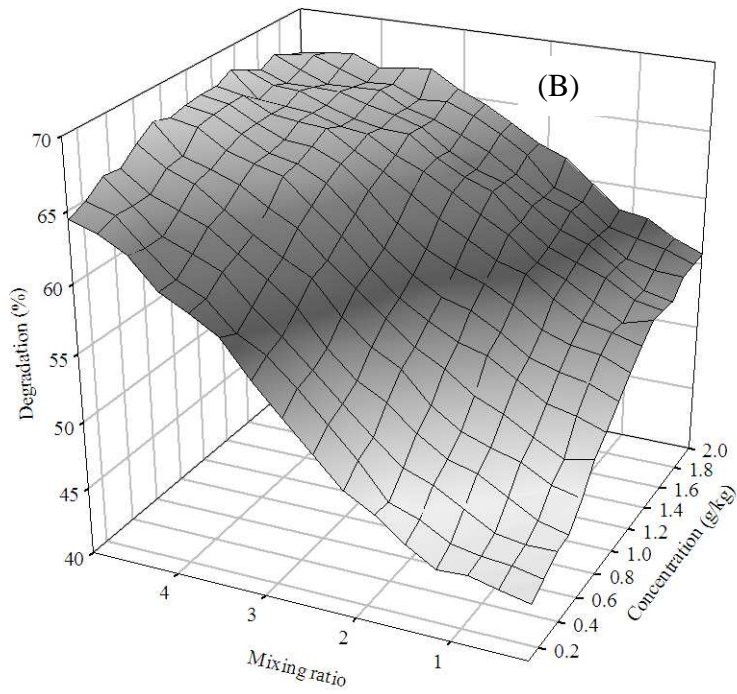
674 **Fig. 3**

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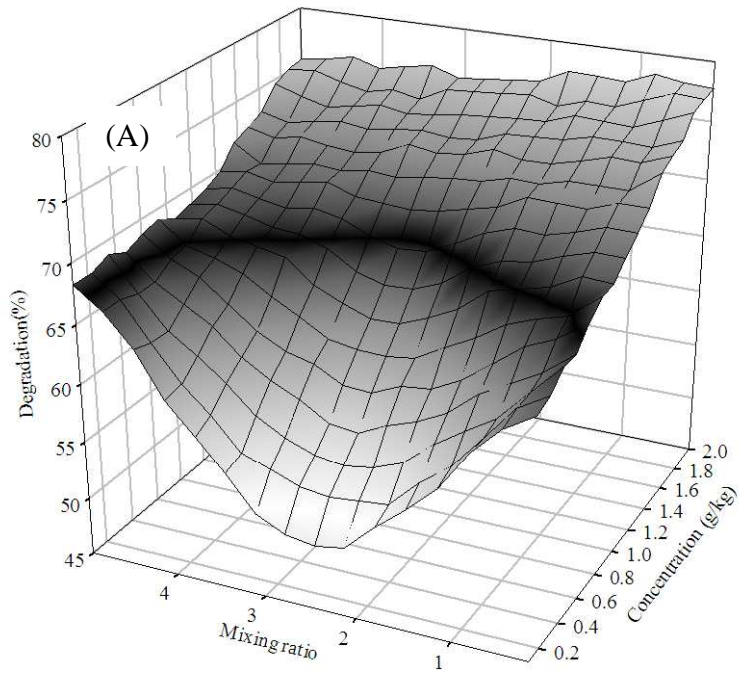
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682 **Fig. 4**

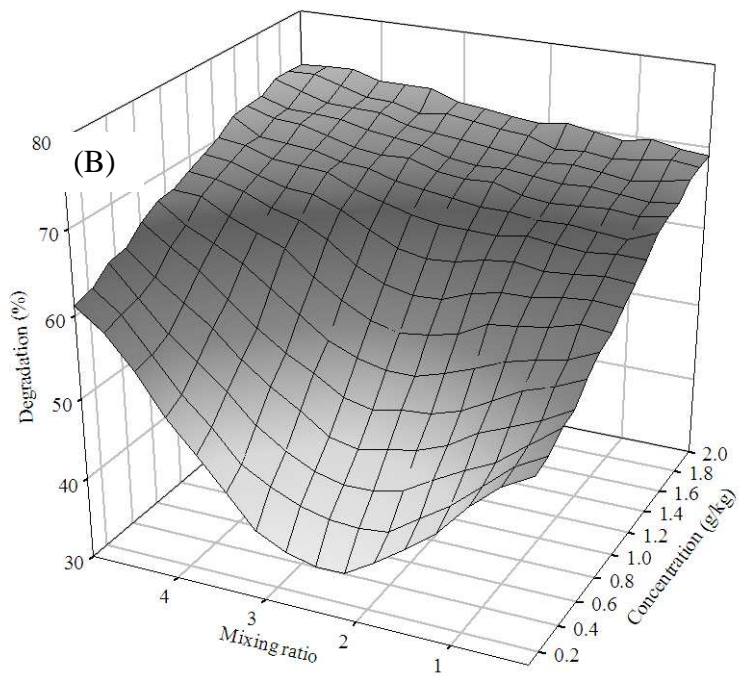
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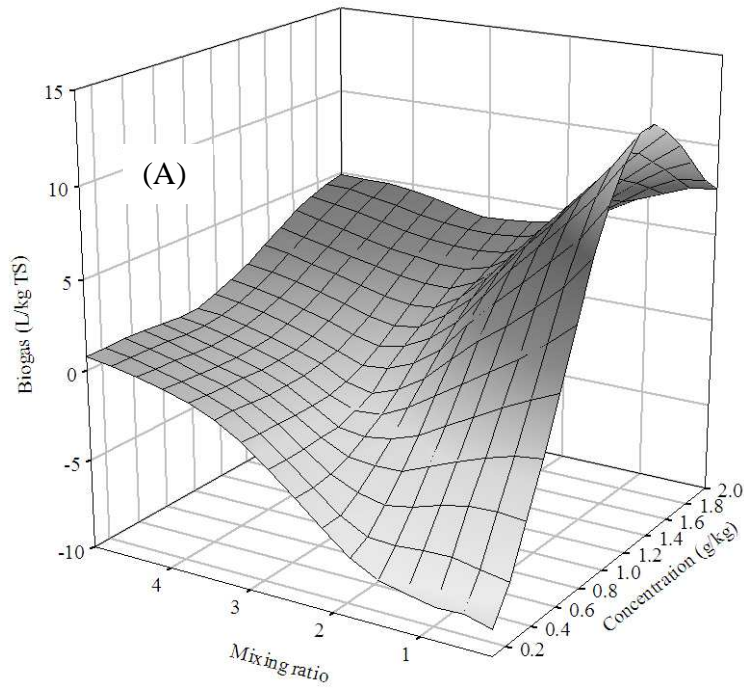
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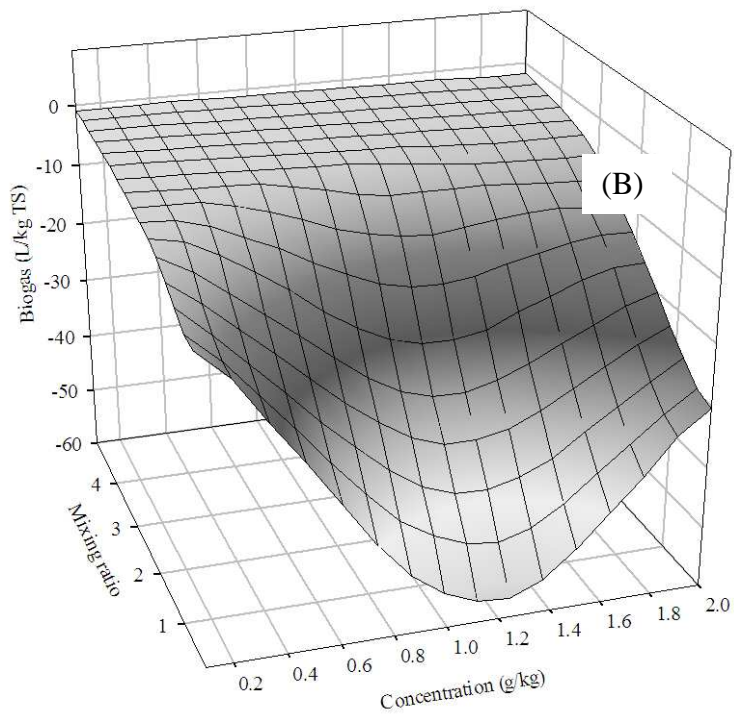
691 **Fig. 5**

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698 **Tables**

699

700

Table 1. Characteristics of inocula and soil used in the experiments

701

| <i>Inoculum/parameter</i> | <i>TS</i> <i>(w.b.)</i> | <i>OM</i> <i>(d.b.)</i> | <i>TOC</i> <i>(d.b.)</i> | <i>TKN</i> <i>(d.b.)</i> | <i>pH</i> | <i>EC</i> | <i>Density</i> <i>(g/L)</i> |
|---------------------------|----------------------------|----------------------------|-----------------------------|-----------------------------|-----------|-----------|--------------------------------|
| Thermophilic | 11.8 | 37.7 | 20.9 | 2.03 | 7.8 | 8.8 | 1113 |
| Mesophilic | 10.02 | 39.1 | 32.8 | 2.5 | 7.9 | 29.9 | 1013 |
| Soil | 93.3 | 3.7 | 1.26 | 0.7 | 6.7 | 0.2 | - |

702 *TS: total solids (%); OM: organic matter (%); TOC: total organic carbon (%); TKN: total Kjeldahl*703 *nitrogen (%); EC: electrical conductivity (mS/cm); w.b.: wet basis; d.b.: dry basis.*

Table 2. Design matrix including factor levels (coded and actual) and the response values for these two factors

| <i>Ru</i> <i>n</i> | <i>Factor levels</i> | | | | Y_{TD30} | Y_{TD50} | Y_{MD30} | Y_{MD50} | Y_{TG30} | Y_{TG50} | Y_{MG30} | Y_{MG50} |
|-----------------------|--|--|--------------------------------|--|------------|------------|------------|------------|------------|------------|------------|------------|
| | <i>Coded</i> | | <i>actual</i> | | | | | | | | | |
| | <i>Concentration</i> (x_1) (g/kg) | <i>Mixing ratio</i> (x_2) (soil:inoculum) | <i>Concentration</i> (g/kg) | <i>Mixing ratio</i> (soil:inoculum) | | | | | | | | |
| 1 | -1 | -1 | 0.38 | 0.9:1 | 69.96 | 42.89 | 66.36 | 53.82 | -15.2 | -10.9 | -20.2 | -5.9 |
| 2 | +1 | -1 | 1.74 | 0.9:1 | 71.25 | 54.78 | 77.75 | 68.83 | 4.8 | 6.8 | -36.4 | -27.6 |
| 3 | -1 | +1 | 0.38 | 4.3:1 | 77.58 | 64.55 | 70.12 | 63.12 | 0.4 | 1.1 | -6.8 | -0.9 |
| 4 | +1 | +1 | 1.74 | 4.3:1 | 70.71 | 65.87 | 75.43 | 72.92 | 4.5 | 5.3 | -6.3 | -1.7 |
| 5 | 0 | 0 | 1.05 | 2.6:1 | 67.91 | 67.43 | 73.92 | 67.90 | -3.2 | -3.0 | -11.4 | -5.4 |
| 6 | $-\alpha$ | 0 | 0.1 | 2.6:1 | 47.26 | 52.57 | 36.8 | 16.04 | -2.1 | -3.1 | -11.5 | -2.1 |
| 7 | $+\alpha$ | 0 | 2.0 | 2.6:1 | 67.17 | 70.83 | 81.23 | 75.57 | -1.9 | 3.6 | -12.1 | -7.7 |
| 8 | 0 | $-\alpha$ | 1.05 | 0.2:1 | 62.91 | 55.82 | 74.57 | 65.00 | 12 | 23.7 | -133.3 | -97.0 |
| 9 | 0 | $+\alpha$ | 1.05 | 5:1 | 75.59 | 74.21 | 70.00 | 74.17 | -1.0 | 0.4 | -3.8 | -0.70 |

*The response represents the degradation (D) percentage (Y_D) and the biogas (G) production (Y_G) after 30 and 50 days of incubation. Where; T: thermophilic; M: mesophilic and 30 or 50 represents the incubation time (day)

Table 3. Correlation coefficients (R) and p-values of each function (cf. Equations 2–9) representing the treatment process

| | <i>Thermophilic</i> | | | | <i>Mesophilic</i> | | | |
|---|---------------------|------------|------------|------------|-------------------|------------|------------|------------|
| | Y_{TD30} | Y_{TD50} | Y_{TG30} | Y_{TG50} | Y_{MD30} | Y_{MD50} | Y_{MG30} | Y_{MG50} |
| R | 0.65 | 0.93 | 0.65 | 0.70 | 0.86 | 0.89 | 0.78 | 0.79 |
| P | 0.63 | 0.06 | 0.61 | 0.5 | 0.17 | 0.03 | 0.16 | 0.15 |

Table 4. Values of “F” fraction (%) for each component after 50 days for samples 5, 6 and 7 under thermophilic (T) and mesophilic (M) conditions

| | <i>T-5</i> | <i>T-6</i> | <i>T-7</i> | <i>M-5</i> | <i>M-6</i> | <i>M-7</i> |
|--------------|------------|------------|------------|------------|------------|------------|
| Fluorene | 96.0 | 100.0 | 97.15 | 88.94 | 100.0 | 77.75 |
| Phenanthrene | 89.1 | 100.0 | 98.65 | 74.15 | 98.38 | 79.08 |
| Anthracene | 69.7 | 77.09 | 81.38 | 70.89 | 93.33 | 73.58 |
| Fluoranthene | 83.24 | 70.68 | 94.81 | 86.45 | 91.98 | 90.25 |
| Pyrene | 68.17 | ND | 92.94 | 70.23 | 4.31 | 78.75 |

ND: not detected (below the detection level)

Table 5. PLFA structure of the microbial biomass for samples 5, 6 and 7 after 30 and 50 days under thermophilic (T) and mesophilic (M) conditions

| <i>Biomass/Treatment</i> | <i>T-5-30</i> | <i>T-6-30</i> | <i>T-7-30</i> | <i>M-5-30</i> | <i>M-6-30</i> | <i>M-7-30</i> | <i>T-5-50</i> | <i>T-6-50</i> | <i>T-7-50</i> | <i>M-5-50</i> | <i>M-6-50</i> | <i>M-7-50</i> |
|----------------------------------|---------------|---------------|---------------|---------------|---------------|---------------|---------------|---------------|---------------|---------------|---------------|---------------|
| Fungi | 2376.38 | 404.45 | 1031.77 | 1139.98 | 894.50 | 1256.78 | 1233.92 | 1034.08 | 1664.30 | 814.67 | 1139.79 | 804.89 |
| Bacteria | 83862.03 | 1399.42 | 2241.85 | 10669.97 | 11112.80 | 11264.75 | 2751.07 | 3575.27 | 3818.31 | 13201.81 | 13485.32 | 10294.19 |
| Actinobacteria | 455.42 | 134.46 | 272.89 | 665.29 | 632.97 | 669.43 | 290.03 | 380.47 | 329.70 | 544.23 | 712.66 | 600.43 |
| Gram positive | 2521.82 | 916.95 | 1153.36 | 8035.50 | 8616.16 | 8570.36 | 1625.29 | 2253.64 | 2357.06 | 10825.81 | 10476.29 | 8111.61 |
| Gram negative | 1202.15 | 283.71 | 736.54 | 1768.22 | 1648.93 | 1822.75 | 701.02 | 760.37 | 974.36 | 1591.95 | 2021.35 | 1396.72 |
| Anaerobic bacteria | 145317.61 | 24332.03 | 77354.29 | 42321.83 | 35682.23 | 44730.48 | 80674.27 | 83980.56 | 91905.50 | 34616.08 | 46113.00 | 33856.22 |
| Total microbial biomass | 163930.30 | 29044.05 | 87662.33 | 60262.10 | 53775.34 | 63563.36 | 91590.98 | 97311.69 | 105652.35 | 55498.66 | 67997.12 | 50700.74 |
| Sum (cy17:0, cy19:0)= <i>Cy</i> | 145.02 | 59.26 | 116.15 | 475.91 | 439.12 | 485.66 | 106.21 | 136.97 | 111.12 | 422.25 | 525.20 | 349.95 |
| Sum(16:1w7,18:1w7)= <i>Pre</i> | 1028.51 | 214.61 | 605.59 | 1250.36 | 1168.43 | 1287.89 | 580.19 | 602.89 | 843.19 | 1130.94 | 1456.30 | 1012.51 |
| Ratio (<i>Cy/Pre</i>) | 0.15 | 0.27 | 0.20 | 0.38 | 0.38 | 0.38 | 0.18 | 0.23 | 0.13 | 0.37 | 0.36 | 0.34 |
| Saturated PLFA (S) | 12191.59 | 3029.16 | 7229.61 | 6782.45 | 6712.44 | 6976.20 | 7163.70 | 9020.58 | 8519.43 | 7499.67 | 8024.28 | 6256.48 |
| Monosaturated PLFA (M) | 146229.72 | 24497.22 | 77858.54 | 43138.23 | 36452.92 | 45581.91 | 81162.86 | 84467.00 | 92657.62 | 35363.52 | 47083.95 | 34553.03 |
| S/M | 0.08 | 0.12 | 0.10 | 0.16 | 0.19 | 0.15 | 0.09 | 0.11 | 0.09 | 0.21 | 0.17 | 0.18 |
| Anaerobic bacteria/total biomass | 0.89 | 0.84 | 0.88 | 0.70 | 0.66 | 0.70 | 0.88 | 0.86 | 0.87 | 0.62 | 0.68 | 0.67 |

