

1 **Improvement of waste-fed bioelectrochemical system performance by**
2 **selected electro-active microbes: Process evaluation and a kinetic study**

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15

16 **Abstract**

17 In this work, bioaugmentation strategy was tested to enhance electricity
18 production efficiency from municipal waste liquor feedstock in microbial fuel
19 cells (MFC). During the experiments, MFCs inoculated with a mixed anaerobic
20 consortium were enriched by several pure, electro-active bacterial cultures
21 (such as *Propionibacterium freudenreichii*, *Cupriavidus basilensis* and
22 *Lactococcus lactis*) and behaviours were assessed kinetically. It turned out
23 that energy yield could be enhanced mainly at high substrate loadings.
24 Furthermore, energy production and COD removal rate showed an optimum
25 and could be characterized by a saturation range within the applied COD
26 loadings, which could be elucidated applying the Monod-model for describing
27 intracellular losses. Polarization measurements showed the positive effect of
28 bioaugmentation also on extracellular losses. The data indicated a successful
29 augmentation process for enhancing MFC efficiency, which was utmost in
30 case of augmentation strain of *Propionibacterium freudenreichii*.

31

32 Keywords: bioaugmentation, microbial fuel cell, *Propionibacterium*
33 *freudenreichii*, *Cupriavidus basilensis*, *Lactococcus lactis*

34

35 1. Introduction

36

37 Microbial fuel cell (MFC) technology can be considered as a rapidly
38 developing alternative for generating electricity using electro-active
39 microorganisms from the chemical energy stored in organic substrates [1 – 3].
40 As various research works demonstrated, besides easy-degradable materials,
41 waste streams may also be utilized in MFCs as feedstock for electricity
42 production [4, 5] e. g. synthetic human blackwater [6], industrial wastewaters
43 [7 – 9], landfill leachate [10] or municipal solid waste [11]. Although in practice
44 MFCs are typically operated with a mixed consortium in the anode chamber, a
45 considerable number of pure cultures have been also tested including different
46 Gram-negative/Gram-positive bacteria, yeasts and algae [12, 13]. In general,
47 such single-strain MFCs are suitable for fundamental research and have
48 limitations for real-field applications due to strict sterility requirements.
49 Nevertheless, they can be viewed as potential candidates for the
50 augmentation of mixed culture MFCs.

51 Bioaugmentation is a well-known strategy for process enhancement (i.e.
52 aiming at the efficient removal of specific components) and relies on the
53 addition of selected microbial species to an initial – mostly natural – microbial
54 consortia/environment [14, 15]. The target compounds to be converted vary
55 widely and can include oil-based contaminations, polycyclic aromatic
56 hydrocarbons (PAHs), phenol, etc. according to the scientific literature [16 –
57 18]. Moreover, microbial augmentation can be advantageous not only in terms
58 of specific substrate degradation but also to improve biofuel (e. g. biogas or
59 biohydrogen) formation as well as integrated applications designed by
60 coupling fermentation and bioelectrochemical treatment [19, 20]. The
61 bioaugmentation in microbiologically-assisted electrochemical systems has
62 been demonstrated with success (i.e. to utilize corn stover [21] or synthetic
63 wastewater [22]) by exploiting specific syntrophic processes and hierarchical
64 structures present in such systems in order to boost electricity generation [23].
65 So far, electro-kinetic analysis of MFCs augmented with *Shewanella haliotis*

66 [22] showed the positive effect of this technique on the grounds of power
67 output and substrate biodegradation. The observed benefits could be mainly
68 attributed to lower activation losses and enhanced shuttling between redox
69 intermediates [22]. In another paper applying electro-active *Pseudomonas*
70 *aeruginosa* and non-electro-active *Escherichia coli* strains for bioaugmentation
71 in MFCs, it could be concluded that the bioelectrochemical cells had taken
72 advantage of synergistic species interactions in the mixed consortia, leading to
73 lower polarization resistance and increased power generation capacity [24].

74 In this work, bioaugmentation of MFCs was carried out by employing
75 pure isolates of electro-active bacteria, namely *Propionibacterium*
76 *freudenreichii*, *Cupriavidus basilensis* and *Lactococcus lactis*, which to our
77 knowledge, have not been used for this this purpose. *P. freudenreichii* is a
78 Gram-positive obligate anaerobic bacteria belonging to the phylum
79 *Actinobacteria* and known as an endogenous mediator-producing strain.
80 Actually, 1,4-dihydroxy-2-naphthoic acid (DHNA) and 2-amino-3-dicarboxy-
81 1,4-naphthoquinone (ACNQ) are reported as electron shuttle molecules,
82 secreted by *P. freudenreichii* [25, 26] which allow its application in mediator-
83 less MFC systems [27]. *C. basilensis* is a flagellated Gram-negative,
84 facultative aerobic β -*proteobacteria* [28] and able to the utilize substances e.g.
85 phenol or aliphatic alcohols as substrates [29, 30]. The members of this genus
86 are described to be capable of producing endogenous mediators for
87 extracellular electron transfer [30, 31]. Since *C. basilensis* is metal-resistant
88 and able to degrade a wide range of materials, its use seems to be promising
89 in wastewater treatment as well as in bioelectrochemical technologies. *L.*
90 *lactis*, a member of phylum *Firmicutes*, is a Gram-positive, facultative
91 anaerobic bacterium with a potential as a biocatalyst in microbial
92 electrochemical cells because of its self-secreted electron accepting and
93 shuttling agent, ACNQ [32, 33]. Furthermore, its important trait is the capability
94 of pursuing electrochemically-modified metabolic pathway besides homolactic
95 fermentation, which leads to the formation of acetate (as by-product) to be

96 consumed by other i.e. exoelectrogenic microorganisms present in an
97 augmented bioelectrochemical reactor [33].

98 To our best knowledge, no comparative study has been done yet with
99 these microbes to investigate bioaugmentation process in MFCs that involves
100 a kinetic approach for the assessment of system behaviour in the course of
101 waste utilization. Therefore, the results demonstrated may have novelty and
102 added-value to support the better understanding of bioaugmentation in MFCs
103 and expand the perspectives of such bioelectrochemical cells.

104

105 **2. Materials and methods**

106

107 **2.1. Seed source and substrates**

108

109 For MFC inoculation, seed source was collected from beet pulp utilizing
110 biogas fermentation unit of Hungarian sugar factory, located at Kaposvár, with
111 an initial microbial community structure demonstrated in our recent work [34].
112 The anaerobic sludge was pretreated (starved) in a laboratory-scale reactor
113 before use for one week at 37 °C. Its main characteristics were the followings:
114 COD content: 12 g L⁻¹, pH = 7.8, Total solids: 6.7 %. As for substrate, pressed
115 fraction of municipal solid waste (LPW) was used. Characteristics of LPW can
116 be found in previous publications [11, 35 – 37]. The most important
117 parameters of the substrate and the flow diagram of its preparation process
118 can be seen in **Fig. 1**.

119

120 **2.2. Preparation of pure cultures of selected electro-active microbes for**
121 **bioaugmentation**

122

123 The pure cultures of selected microbes were purchased from the
124 German Collection of Microorganisms and Cell Cultures (DSMZ). The broth
125 media compositions were the followings: *Lactococcus lactis* (DSMZ-20481)
126 broth – casein peptone (pancreatic digest) 17 g L⁻¹, K₂HPO₄ 2.5 g L⁻¹, glucose
127 2.5 g L⁻¹, NaCl 2.5 g L⁻¹, soy peptone (papaic digest) 3 g L⁻¹, yeast extract 3 g
128 L⁻¹, agar 20 g L⁻¹ (pH = 7); *Cupriavidus basilensis* (DSMZ-11853) broth –
129 peptone 5 g L⁻¹, meat extract 3 g L⁻¹, agar 20 g L⁻¹ (pH = 7); *Propionibacterium*
130 *freudenreichii* (DSMZ-20271) broth – casein peptone (tryptic digest) 10 g L⁻¹,
131 yeast extract 5 g L⁻¹, Na-lactate 10 g L⁻¹, agar 20 g L⁻¹ (pH = 7).

132 The cultures were incubated on agar plates – and in stab agar in case of
133 *P. freudenreichii* – for two days at 37 °C. Thereafter, colonies were harvested
134 and transferred to liquid media (50 mL, without agar) and incubated for two
135 more days under the same conditions. Before use in MFCs, the cell
136 concentration of liquid cultures was determined by Bürker's chamber.

137

138 **2.3. MFC design and setup**

139

140 The design of dual-chamber microbial fuel cells was adopted from our
141 previous work [38]. In this MFC construction, anode and cathode
142 compartments (with 60 mL total volume) were equipped with carbon cloth
143 (Zoltek Corp., USA) and Pt-C (0.3 mg cm² Pt content, FuelCellsEtc, USA)
144 electrodes (64 cm² apparent surface area), respectively. The anode and
145 cathode were connected by Ti wire (Sigma-Aldrich, USA) to the external
146 circuit, containing a 100 Ω resistor. The chambers were separated by Nafion
147 115 proton exchange membrane (Sigma-Aldrich, USA) with diameter of 4.5
148 cm. Before use, the membrane was activated as described elsewhere [38]. In
149 order to maintain aerobic conditions, air was continuously supplemented to the
150 cathode compartment.

151 The anode side of MFCs was filled with 50 mL of mesophilic sludge (pH
152 adjusted to 7) and 5 mL of individual, pure strain liquid culture. Based on cell
153 counting and prior to loading, the liquid cultures were diluted to provide equal
154 cell concentration for each bioaugmented reactors. Thus, initial cell
155 concentration of $3.23 \times 10^7 \pm 2.6 \times 10^6$ cells mL⁻¹ could be reached and
156 maintained in the liquid (5 mL) samples employed for bioaugmentation,
157 irrespective of the strain. The anode chamber was then purged with high purity
158 nitrogen gas to remove dissolved oxygen and ensure anaerobic conditions. In
159 the cathode chamber, 50 mM phosphate buffer (pH = 7) was used as
160 electrolyte. The MFC reactors were running at 37 °C. In addition to the
161 bioaugmented reactors, control MFCs started-up only with (55 mL) inoculum
162 (50 mL slected sludge + 5 mL phosphate buffer) was established, as well.
163 Substrate (LPW, Section 2.1.) additions (0.5, 1, 2 and 4 mL, depending on the
164 aimed COD loading) were carried out by using batch operational mode, after
165 the adaptation phase has been successfully performed (Section 3.1.). During
166 each injection of LPW, the appropriate amount of anolyte was drawn
167 (exchange of volumes) to ensure a consistent working volume. Once the
168 observed voltage dropped close to the initial (**Fig. 2**), a new feeding cycle
169 could be commenced [34].

170

171 **2.4. Analysis and calculations**

172

173 Cell voltage (U) was measured and monitored through a 100 Ω external
174 resistor by a data acquisition system (National Instruments, USA) in Labview
175 environment. According to Ohm's law, current (I) (and electric power, P) were
176 computed. Cell polarization measurements were carried out by varying the
177 resistors in the external circuit of MFCs between 3.3 kΩ – 10 Ω. From the
178 linear region of voltage vs. polarization current density ($i_{max,P}$) plots, the overall
179 internal cell resistance (R_i) – as the slope of the fitted trendline – could be
180 derived.

181 The energy yield was calculated according to Eq. 1:

182

$$183 \quad Y_S = \frac{E}{m_{\Delta COD} A} \quad (1)$$

184

185 where A is the apparent anode surface area (m^2), E is the cumulated energy
186 (kJ) derived from the integration of $P - t$ curves, $m_{\Delta COD}$ is the quantity of COD
187 removed (gram) during a given cycle. The COD content of particular samples
188 was analyzed in accordance with our previous paper [39] by relying on the
189 standard methods of APHA.

190 The rates of (i) Energy production and (ii) COD removal were computed
191 according to Eqs. 2 and 3, respectively, considering the operation time of
192 given batch cycles (τ):

193

$$194 \quad v_E = \frac{E}{A \tau} \quad (2)$$

195

$$196 \quad v_S = \frac{m_{\Delta COD}}{V \tau} \quad (3)$$

197

198 The effect of substrate concentration on current generation – and thus,
199 intracellular losses – was evaluated by adopting the principles of Monod model
200 [40]. In this model the relation of the two variables (substrate concentration
201 and current density) can be described by Eq. 4.

202

$$203 \quad i = i_{max} \frac{[S]}{K_{S,app} + [S]} \quad (4)$$

204

205 where i denotes the current density (relative to the apparent anode surface
206 area), $K_{S,app}$ is the apparent half-saturation substrate concentration (half-
207 saturation constant) and $[S]$ is the substrate (LPW) concentration. To estimate
208 the kinetic parameters (i_{max} and $K_{S,app}$) the linearized (double-reciprocal) form
209 of Eq. 4 was applied, as represented in Eq. 5.

210

211
$$\frac{1}{i} = \frac{K_{S,app}}{i_{max}[S]} + \frac{1}{i_{max}} \quad (5)$$

212

213 In the model (Eqs. 4 and 5), [S] is given in the unit of e⁻ eq L⁻¹, considering 8 g
214 COD as equivalent of 1 mol e⁻ [40].

215 The determination of mean values and standard deviations/errors for
216 parameters such as i_{max} , $P_{d,max}$, Y_S , v_S , v_E , etc. appearing throughout this work
217 (i.e. **Table 1**) was carried out as detailed in the Supplementary file (**Fig. 1S**).

218

219 **3. Results and Discussion**

220

221 **3.1. Evaluation of bioaugmentation efficiency in MFC – peak current and** 222 **power densities, energy yield**

223

224 In the first part of operation – considered as the acclimation period – 5
225 mM acetate was added in the MFCs as adapting substrate in consecutive
226 cycles until comparable current density profiles in particular reactors were
227 reached over time (after three weeks) [41]. Afterwards, feeding of stabilized
228 MFCs was commenced with LPW and the measurements were dedicated to
229 examine the impact of bioaugmentation. The MFCs were operated with
230 different amounts of LPW in the range of 0.5 – 4 mL (equivalent to 0.88 – 7.04
231 g_{COD} L⁻¹). The most important parameters of each system tested are
232 summarized in **Table 1** (average output values and standard deviations for the
233 individual feeding processes) and the current density profiles can be seen in
234 **Fig. 2**.

235 In terms of highest attainable current and power densities (noticed at the
236 highest LPW supplementation, 7.04 g_{COD} L⁻¹), the MFCs could be ranked in the
237 following order: *Propionibacterium*-MFC (76.2 – 110.3 mA m⁻² / 3.7 – 7.8 mW
238 m⁻², respectively), *Cupriavidus*-MFC (70.6 – 100.1 mA m⁻² / 3.2 – 6.4 mW m⁻²),
239 Control-MFC (66.2 – 102 mA m⁻² / 2.8 – 6.7 mW m⁻²) and *Lactococcus*-MFC
240 (57.6 – 95.1 mA m⁻² / 2.1 – 5.8 mW m⁻²). Interestingly, in the light of already
241 published literature relevant to the latest strain, *L. lactis*, though Freguia et al.

242 [33] achieved proper operation of MFCs using its monoculture to generate
243 current from glucose, no electrogenic activity in MFCs was found by
244 Rosenbaum et al. [42] with *L. lactis* alone on the same substrate. Interestingly,
245 however, co-cultures of *Shewanella oneidensis* and *L. lactis* were able to
246 produce current (64 – 215 mA m⁻²) from this substance [42]. Hence, it can be
247 implied that the behaviour of *L. lactis* is dependent on factors such as the
248 composition of underlying community structure (i.e. the number and features of
249 other bacteria to live and cooperate with), which is likely true for *C. basilensis*
250 and *P. freudenreichii* as well. To assess such aspects (i.e. how the
251 microbiological background of the sludge inoculum influences the integration
252 of particular cultures into the community) the population dynamics should be
253 tracked via molecular biological tools, which should be the subject of a follow-
254 up study.

255 Y_s , as expressed in Eq. 1, is an appropriate response variable to make
256 comparison between the systems from the point of view of cumulative energy
257 recovery efficiency. According to the results, an LPW (substrate)
258 concentration-dependent variation of Y_s was found in all cases, where at low
259 COD loadings (0.88 and 1.76 g_{COD} L⁻¹) only the *Cupriavidus*-MFC could
260 surpass the Control-MFC. In case of *Propioni*-, *Cupriavidus*- and Control-
261 MFCs, nearly equal energy yields (1.59, 1.62 and 1.69 kJ g_{ΔCOD}⁻¹ m⁻²,
262 respectively) could be observed at middle COD loading of 3.52 g_{COD} L⁻¹.
263 Nevertheless, by further increasing the COD loading to the highest value of
264 7.04 g_{COD} L⁻¹, energy yields were significantly improved in all bioaugmented
265 MFCs in comparison with the Control-MFC. As a matter of fact, increment of
266 Y_s relative to the control reactor was 91 %, 47 % and 21 % for *Propioni*-,
267 *Lactococcus*- and *Cupriavidus*-MFCs, respectively (**Table 1**).

268 Overall, from the process evaluation considering peak current and
269 power densities as well as energy yield, it would appear that the obligate
270 anaerobic *P. freudenreichii* was the most promising among the strains for
271 augmentation under the experimental circumstances provided.

272

273 **3.2. On energy production and COD removal rates**

274

275 Trends of Coulombic efficiency (CE) (derived in accordance with Logan
276 et al. [3] considering the amount of COD removed) as a function of COD
277 loading can be observed in **Fig. 3**, which implies that bioaugmentation had an
278 advantageous effect on CE at every operating point. The difference between
279 CEs was less pronounced at the lowest COD loading where CE obtained to be
280 about 1.3 – 1.8 %. Nevertheless, by increasing the COD loading, the
281 increment in CE values of the augmented MFCs became more and more
282 emphasized compared to the control and by reaching the highest loading (7.04
283 $\text{g}_{\text{COD}} \text{L}^{-1}$), the *Propionibacterium*-, *Cupriavidus*- and *Lactococcus*-MFC
284 exceeded the CE of Control-MFC by 129, 35 and 50 % (with corresponding
285 CEs of 3.88 ± 0.21 %, 2.28 ± 0.1 % and 2.52 ± 0.14 % versus 1.69 ± 0.12 %),
286 respectively. CEs in the same order of magnitude had been obtained in our
287 previous work, demonstrating a sequential anaerobic treatment process
288 (biohydrogen fermentation – biogas generation – microbial fuel cell) for the
289 enhancement of overall energy recovery from LPW as feedstock [37].

290 To assess the MFC efficiency, not only the total achievable energy
291 yields (product) and COD (substrate) removals are to be considered but
292 corresponding rates as well since the process should be accomplished within
293 a reasonable time. Consequently, an evaluation based on reaction rate-like
294 variables defined in Eqs. 2 and 3 is of importance.

295 As it is depicted in **Fig. 4**, similar relationship could be established
296 between energy production rate (v_E) and COD loadings for all MFCs until 3.52
297 $\text{g}_{\text{COD}} \text{L}^{-1}$ concentration. However, at the highest COD dose (7.04 $\text{g}_{\text{COD}} \text{L}^{-1}$), v_E in
298 the bioaugmented cells was declined and hence, a peak v_E value could be
299 noted within the COD range investigated. The phenomena that decreased v_E
300 was observed in case of bioaugmented cells at 7.04 $\text{g}_{\text{COD}} \text{L}^{-1}$ (than at 3.52 g_{COD}
301 L^{-1}) is attributed to the nonlinear increase of operation times for batch cycles. It
302 is also to notice in **Fig. 4** that there was a considerable difference of v_E
303 between the systems at 3.52 $\text{g}_{\text{COD}} \text{L}^{-1}$ concentration, leading to a 47 % faster

304 energy recovery rate by the most efficient *Propionibacterium*-MFC compared
305 to the control (non-bioaugmented) reactor (482 and 327 J m⁻² d⁻¹,
306 respectively). However, at highest COD addition (7.04 g_{COD} L⁻¹), more or less
307 similar v_E was found for all MFCs This suggests the existence of substrate
308 (COD) saturation range where although more organic matter is available, the
309 reaction rate is not further enhanced in a proportional way due to fully
310 exploited capacity of exoelectrogens present in MFCs. A basically similar
311 discussion can be made concerning the data related to COD (substrate)
312 removal rates (v_S), as illustrated in **Fig. 5**. The fact that tendencies in v_E and v_S
313 are analogous can be explained by concurrent product (energy) formation and
314 substrate (COD in LPW feedstock) consumption. In essence, at 3.52 g_{COD} L⁻¹,
315 the maximum v_S was attained with the *Propionibacterium*-MFC, being 31 %
316 higher than for the Control-MFC (5.52 and 4.2 g L⁻¹ d⁻¹, respectively). The v_S
317 values (between 1 – 5.52 g L⁻¹ d⁻¹, depending on the actual COD loading) are
318 comparable to the relevant literature, where for example Raghavulu et al. [22]
319 demonstrated v_S of 0.41 g L⁻¹ d⁻¹ by using *S. haliotis* as augmentation species.
320 In another publication, v_S of 0.59 g L⁻¹ d⁻¹ could be reached with *P. aeruginosa*-
321 augmented MFCs, which was 11 % greater than the non-augmented system
322 demonstrating $v_S = 0.53$ g L⁻¹ d⁻¹ [24]. Moreover, phenol-utilizing (pure culture)
323 *C. basilensis*-MFC could be characterized by roughly one order of magnitude
324 lower COD removal rate ($v_S \approx 0.05$ g L⁻¹ d⁻¹) [32].

325 It is noteworthy that v_E and v_S are representative for a whole batch cycle,
326 during which, however, various stages of both product formation and substrate
327 removal can be distinguished. These, in particular, include consecutive phases
328 of (i) increasing, (ii) maximal (steady-state) and (iii) decreasing current
329 production and simultaneous COD elimination rates. Among them, the main
330 point of interest is the steady-state with maximal (i) current production and (ii)
331 substrate utilization rates, where various intra- and extracellular
332 mechanism/factors play a role [40]. Thus, in the next sections, the MFC data
333 collected under steady-state conditions will be processed. Firstly (Section

334 3.3.), a kinetic approach will be applied to get an insight to intracellular losses
335 related to reaction rate and bioconversion capacity of exoelectrogens [40].
336 Secondly (Section 3.4.), polarization results will be presented to evaluate
337 extracellular losses [40].

338

339 **3.3. Monod model for substrate utilization kinetics – intracellular losses**

340

341 The current generation and its kinetics are determined by two main
342 factors at intracellular level (where the electrons are conveyed from the
343 electron donor molecule to the outer membrane proteins or secreted shuttle
344 molecules) [40]. Firstly, substrate degradation takes place and reduced
345 intracellular charge carrier molecules (NADH) are formed [43]. Afterwards,
346 processes with the involvement of electron transport chain govern the
347 electrons to the starting point of extracellular electron transfer. The former step
348 can be described by the Monod model (Eq. 4), which correlates the current
349 density with substrate concentration [43]. Therefore, plotting maximal (steady-
350 state) current densities vs. substrate concentration allows studying related
351 (intracellular) energy losses. It is to note that experimental results obtained at
352 $0.44 \text{ g}_{\text{COD}} \text{ L}^{-1}$ were added to make the analysis via the Monod model more
353 reliable. The double-reciprocal interpretation (Eq. 5) of Monod model is
354 depicted in **Fig. 6**. Based on the slope of trendlines fitted for the bioaugmented
355 and non-bioaugmented (control) MFCs, kinetic parameters (i_{max} and $K_{\text{S,app}}$)
356 could be delivered. As it can be drawn from **Table 2**, comparable i_{max} values
357 were found for all systems ($109.9\text{-}120.5 \text{ mA m}^{-2}$). This, together with **Fig. 7**
358 confirms the implications made in Section 3.3. regarding the existence of a
359 substrate saturation range where the highest COD loading ($7.04 \text{ g}_{\text{COD}} \text{ L}^{-1}$,
360 which is $882 \text{ e}^{-} \text{ meq L}^{-1}$ according to Eq. 5) belongs to. As for $K_{\text{S,app}}$ listed in
361 **Table 2**, the MFC augmented with *P. freudenreichii* demonstrated the lowest
362 value with $67.7 \text{ e}^{-} \text{ meq L}^{-1}$, followed by *C. basilensis*-MFC ($73.5 \text{ e}^{-} \text{ meq L}^{-1}$),
363 Control-MFC ($91 \text{ e}^{-} \text{ meq L}^{-1}$) and *Lactococcus*-MFC ($99.4 \text{ e}^{-} \text{ meq L}^{-1}$). In
364 essence, obtaining a lower $K_{\text{S,app}}$ is advantageous from a reaction rate point of
365 view. Thus, the energy production rate (v_{E}) achieved in *Propionibacterium*-
366 MFC (compared to the other reactors) can be likely associated with the low
367 $K_{\text{S,app}}$ value, helping to maintain relatively higher electricity generation even at
368 lower substrate (COD) concentrations in accordance with the Monod model
369 (zero-order kinetics). Overall, bioaugmentation with the aid of selected pure

370 bacterial cultures such as *P. freudenreichii* and *C. basilensis* could effectively
371 decrease the limiting substrate concentration in MFCs. Once high (close to
372 maximal) v_E is kept at lower $[S]$, the intracellular losses ascribed to the only
373 partly exploited capacity of active exoelectrogens (causing limitation of
374 reaction rate) in MFC can be reduced [40].

375 In fact, $K_{S,app}$ and i_{max} values obtained in this work are somewhat lower
376 compared to other MFC research studies using components such as acetate,
377 ethanol or propionate, probably due to the complex structure of LPW
378 feedstock. For instance, $K_{S,app} = 19 \text{ e}^- \text{ meq L}^{-1}$ ($i_{max} = 2200 \text{ mA m}^{-2}$) was
379 observed in case of acetate-utilizing MFC [45]. In another work, $K_{S,app} = 0.18 -$
380 $58 \text{ e}^- \text{ meq L}^{-1}$ was documented for ethanol substrate [46]. In microbial
381 electrolysis cell (MEC) mode, Torres et al. [47] reported half-saturation
382 constants of 22, 5.3 and $3.8 \text{ e}^- \text{ meq L}^{-1}$ for acetate, ethanol and propionate,
383 respectively, while maximal current densities varied between approximately
384 $1.8 - 9 \text{ A m}^{-2}$.

385

386 **3.4. Cell polarization characteristics – extracellular losses**

387

388 Basically, polarization techniques can be applied to describe the system
389 from extracellular processes (and related potential or energy losses) [3].
390 These, on the anode side, cover (i) the transfer of electrons to the conductive
391 biofilm matrix and/or soluble shuttle molecules in the bulk phase and (ii) the
392 charge transport (conductive or diffusive) to the anode surface, where the
393 electrode reaction takes place. By varying the external resistance in the MFC
394 electrical circuit and measuring the cell voltage subsequently, polarization and
395 power density curves (U vs. i and P_d vs. i , respectively) can be registered (**Fig.**
396 **8**). Based on these data, the actual internal resistance (R_i) of an MFC is
397 estimated [3]. Considering the polarization chart (taken in steady-state at 7.04
398 gCOD L^{-1} LPW concentration), (i) activation polarization, (ii) ohmic and (iii)
399 concentration polarization regions could be identified in each MFC. The open
400 circuit voltages (OCV) were comparable, spanning $425 - 442 \text{ mV}$. As for R_i in

401 the bioaugmented MFCs, values belonging to *Lactococcus*-,
402 *Propionibacterium*- and *Cupriavidus*-MFC were noted such as 347 Ω , 341 Ω
403 and 348 Ω (with $R^2 > 0.98$). In the Control-MFC, the corresponding value was
404 higher (383 Ω).

405 The comparable voltages occurring at low current densities (**Fig. 8**) can
406 be explained by the restricted passage of electrons through the circuit, caused
407 by high external resistor [44]. Therefore, from these similar values, a
408 resistance value can be assumed above which the global reaction rate in MFC
409 (ending with proton reduction at the cathode by electrons captured and
410 delivered from the anode) will be independent of the microbial reduction rate of
411 charge-carrying redox components. By lowering the external resistance,
412 continuous voltage drop and simultaneously increasing current density can be
413 observed, where more oxidized-form electron carriers are present and
414 implicitly, the marked role of electro-active microbial metabolism becomes
415 apparent. Moreover, i in various MFCs can be properly distinguished at lower
416 (external) resistances, as indicated by **Fig. 8**. In general, the bioaugmented
417 MFCs generated higher maximal polarization current density ($i_{max,P}$) than the
418 Control-MFC did. Expressed in numbers, $i_{max,P}$ of 110, 116 and 127 mA m^{-2}
419 could be reached in *Lactococcus*-, *Cupriavidus*- and *Propionibacterium*-MFCs,
420 respectively, where the latest case demonstrates 21 % increment relative to
421 the non-augmented system (105 mA m^{-2}).

422 The significantly positive effect of bioaugmentation on MFC performance
423 could be recognized on grounds of maximal power densities ($P_{d,max}$, **Fig. 8**) to
424 be ordered as follows: 6.6 mW m^{-2} (Control-MFC), 7.9 mW m^{-2} (*Cupriavidus*-
425 MFC), 8.2 mW m^{-2} (*Lactococcus*-MFC), 9.8 mW m^{-2} (*Propionibacterium*-MFC).
426 Thus, in this aspect too, the enrichment of microbial consortia by *P.*
427 *freudenreichii* was the most advantageous strategy to improve
428 bioelectrochemical system efficiency. The findings presented are in agreement
429 with the literature, where Raghavulu et al. [22] attained OCV of 378 mV using
430 *S. haliotis* for bioaugmentation with R_i , $i_{max,P}$ and $P_{d,max}$ of 300 Ω , 320 mA m^{-2}
431 and 29.6 mW m^{-2} , respectively. In addition, bioaugmentation of MFCs with *P.*

432 *aeruginosa* resulted in relatively high OCV (418 mV) and maximal power
433 density (69.9 mW m⁻²) with polarization current density of ~ 450 mA m⁻² [24].
434 The results of Reiche et al. [28] for *P. freudenreichii*-driven MFC are also
435 comparable to ours with *Propionibacterium*-MFC, realizing OCV of 485 mV
436 and $P_{d,max}$ of 14.9 mW m⁻² [28]. In MFCs operated with monoculture of *C.*
437 *basilensis* as exoelectrogenic biocatalyst, Friman et al. [32] could observe
438 OCV of about 250 mV and $P_{d,max}$ of 10 mW m⁻², which coincide well with our
439 values in *Cupriavidus*-MFC (OCV = 425 mV and $P_{d,max} = 7.9$ mW m⁻²).

440 In this work, the selected electro-active bacteria were known as
441 producers of electron shuttle molecules (Section 1.) and therefore, a process
442 via such soluble compounds can be supposed. This argument seems to be
443 supported by the current density values documented in this investigation ($i_{max,P}$
444 in the order of 10² mA m⁻²), implying the more likely occurrence of mediated
445 (diffusion controlled) electron transport rather than a direct contact mechanism
446 [40].

447

448 **4. Conclusions**

449

450 In this study, bioaugmentation process and its effect on microbial fuel
451 cell performance were investigated by several electro-active bacterial cultures.
452 Considering the electric outputs (i.e. current and power density) and energy
453 yield, the bioaugmented MFCs were more efficient at higher COD loadings
454 than the control. The analysis of energy production and COD removal rates
455 revealed an optimum COD loading. Besides, substrate saturation and the
456 existence of zero-order kinetics region at the highest substrate concentration
457 were confirmed by applying Monod model. $K_{S,app}$ values could be significantly
458 decreased in case of *Propionibacterium*- and *Cupriavidus*-MFC compared to
459 the control. Polarization measurements indicated the positive impact of
460 bioaugmentation on extracellular losses and enhanced electron shuttle
461 mechanism could be presumed. In conclusion, microbial augmentation can be
462 considered as a promising strategy to improve microbial fuel cells. After

463 examination of systems behavior from various points of views,
464 *Propionibacterium freudenreichii* was found as the most advantageous strain
465 among those tested for bioaugmentation in the experiments.

466

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468

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479

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647

648 **Table 1 – Stationary electric outputs and energy yield at different COD**
 649 **loadings for bioaugmented and control MFCs.**

COD loading (g _{COD} L ⁻¹)		<i>Propionibacterium</i> - MFC	<i>Cupriavidus</i> - MFC	<i>Lactococcus</i> - MFC	Control- MFC
0.88		76.2 ± 1.98	70.6 ± 1.23	57.6 ± 3.61	66.2 ± 3.27
1.76	i_{max}	87.7 ± 2.46	81.3 ± 1.76	76.5 ± 1.37	80.2 ± 2.52
3.52	(mA m ⁻²)	109.7 ± 0.86	95.8 ± 1.42	91.4 ± 1.98	92.3 ± 1.55
7.04		110.3 ± 0.76	100.1 ± 1.94	95.1 ± 1.55	102 ± 4.61
0.88		3.7 ± 0.18	3.2 ± 0.11	2.1 ± 0.24	2.8 ± 0.25
1.76	$P_{d,max}$	4.9 ± 0.26	4.2 ± 0.19	3.8 ± 0.13	4.1 ± 0.25
3.52	(mW m ⁻²)	7.7 ± 0.12	5.9 ± 0.17	5.4 ± 0.22	5.5 ± 0.17
7.04		7.8 ± 0.13	6.4 ± 0.24	5.8 ± 0.18	6.7 ± 0.57
0.88		1.33 ± 0.05	1.54 ± 0.06	0.86 ± 0.04	1.45 ± 0.11
1.76	Y_s	1.19 ± 0.05	1.63 ± 0.09	1.15 ± 0.06	1.53 ± 0.17
3.52	(kJ g _{ΔCOD} ⁻¹ m ⁻²)	1.59 ± 0.09	1.62 ± 0.09	1.43 ± 0.09	1.69 ± 0.09
7.04		3.62 ± 0.20	2.29 ± 0.09	2.77 ± 0.16	1.89 ± 0.13

650

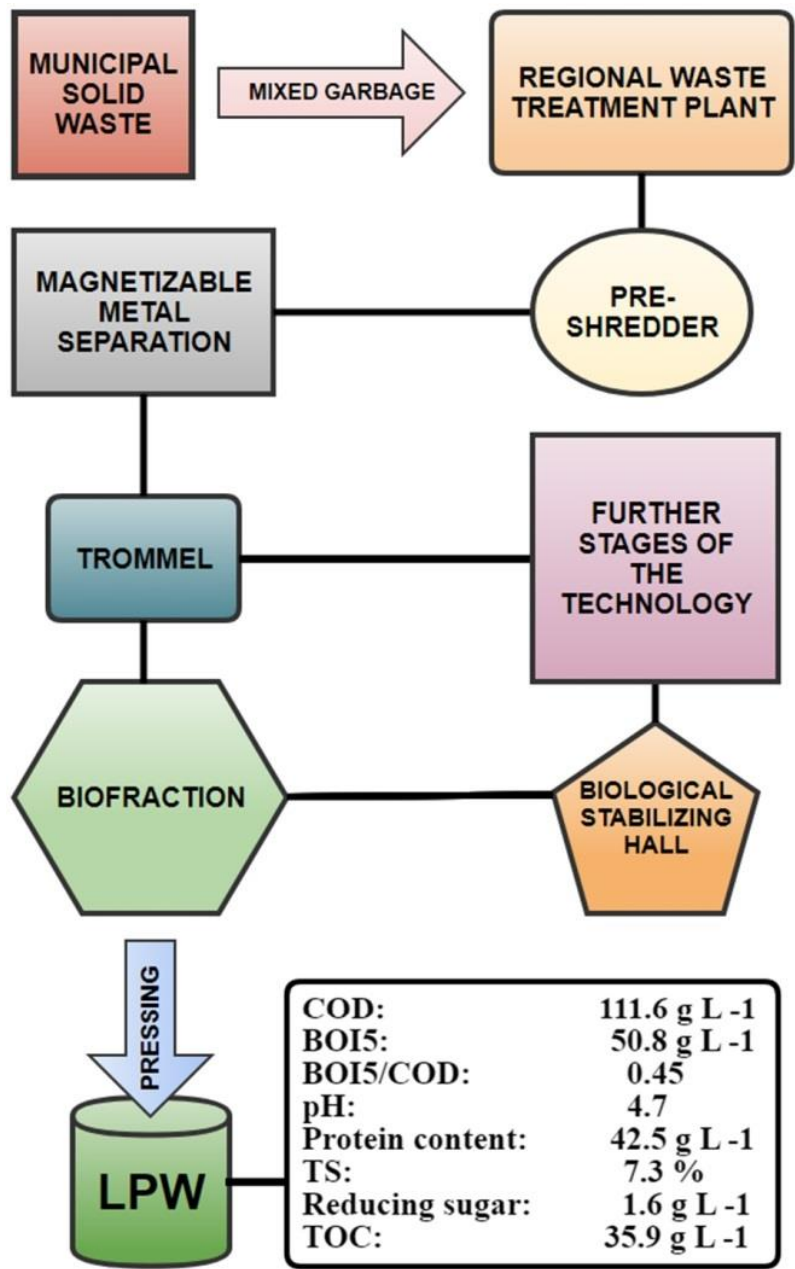
651 **Table 2 – Kinetic parameters and R-squared value of the fitted Monod**
652 **model.**

MFC type	i_{max} (mA m ⁻²)	$K_{S,app}$ (e ⁻ meq L ⁻¹)	R ² (-)
<i>Propionibacterium</i> -MFC	120.5	67.7	0.988
<i>Cupriavidus</i> -MFC	111.1	73.5	0.999
<i>Lactococcus</i> -MFC	109.9	99.4	0.990
Control-MFC	112.4	91.0	0.988

653

654 Fig. 1

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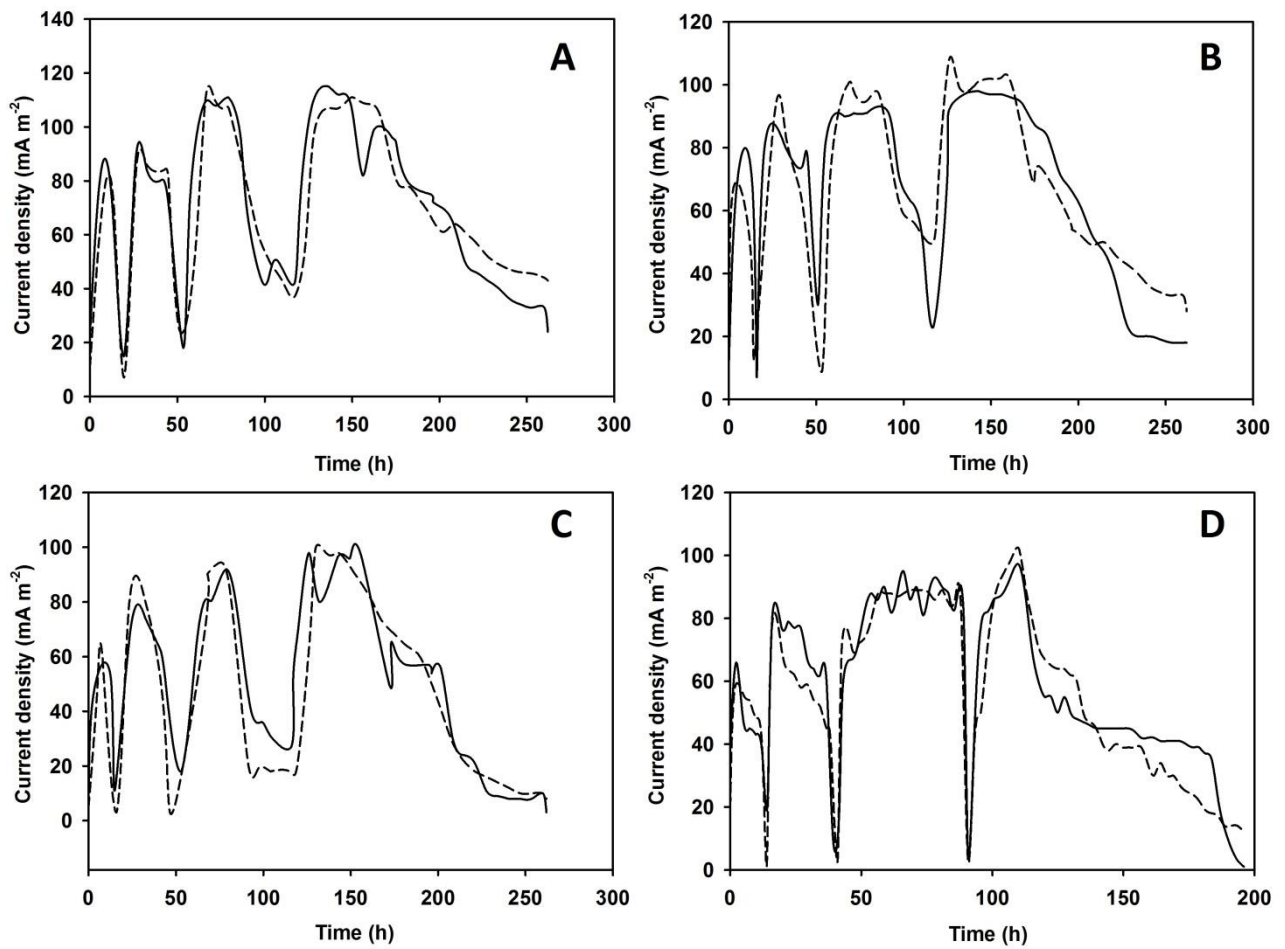
658

Process flow diagram of LPW preparation.

659

660

661 Fig. 2



662

663

664 **Current density profiles of the different MFCs (progress curves of**

665 **replicates are shown). A: *Propionibacterium*-MFC; B: *Cupriavidus*-MFC; C:**

666 ***Lactococcus*-MFC; D: Control-MFC. In all MFC, the order of substrate (LPW)**

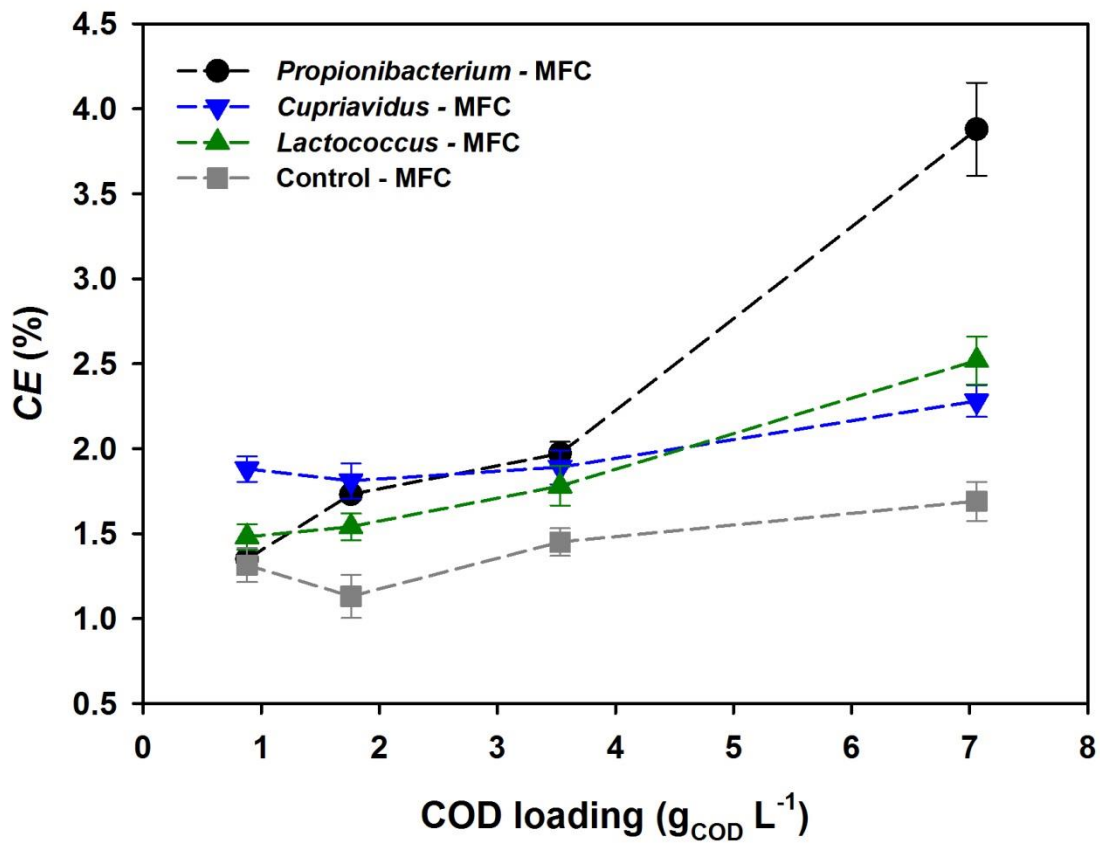
667 **loadings was the following: 0.5; 1; 2 and 4 mL.**

668

669

670 Fig. 3

671



672

673

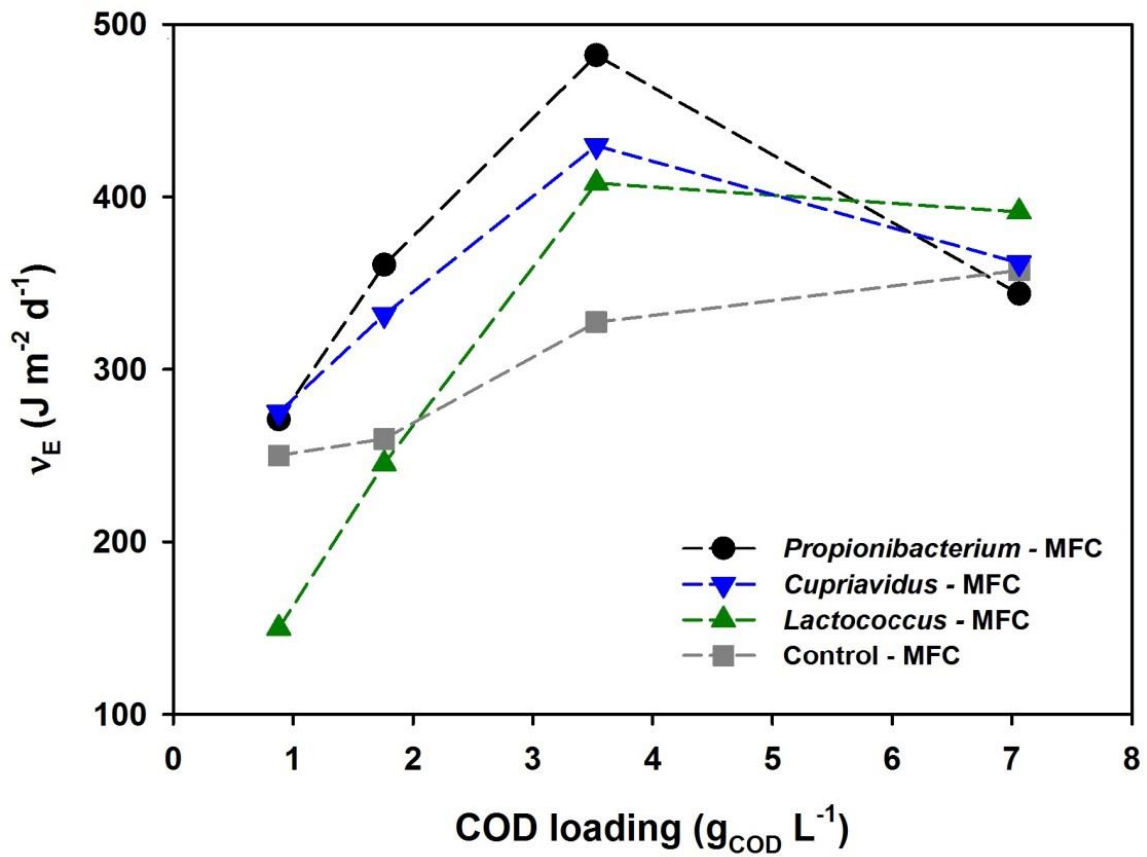
674 **Coulombic efficiency as a function of COD loading. -●-: *Propionibacterium*-**

675 **MFC; -▼-: *Cupriavidus*-MFC; -▲-: *Lactococcus*-MFC; -■-: Control-MFC.**

676

677

678 Fig. 4



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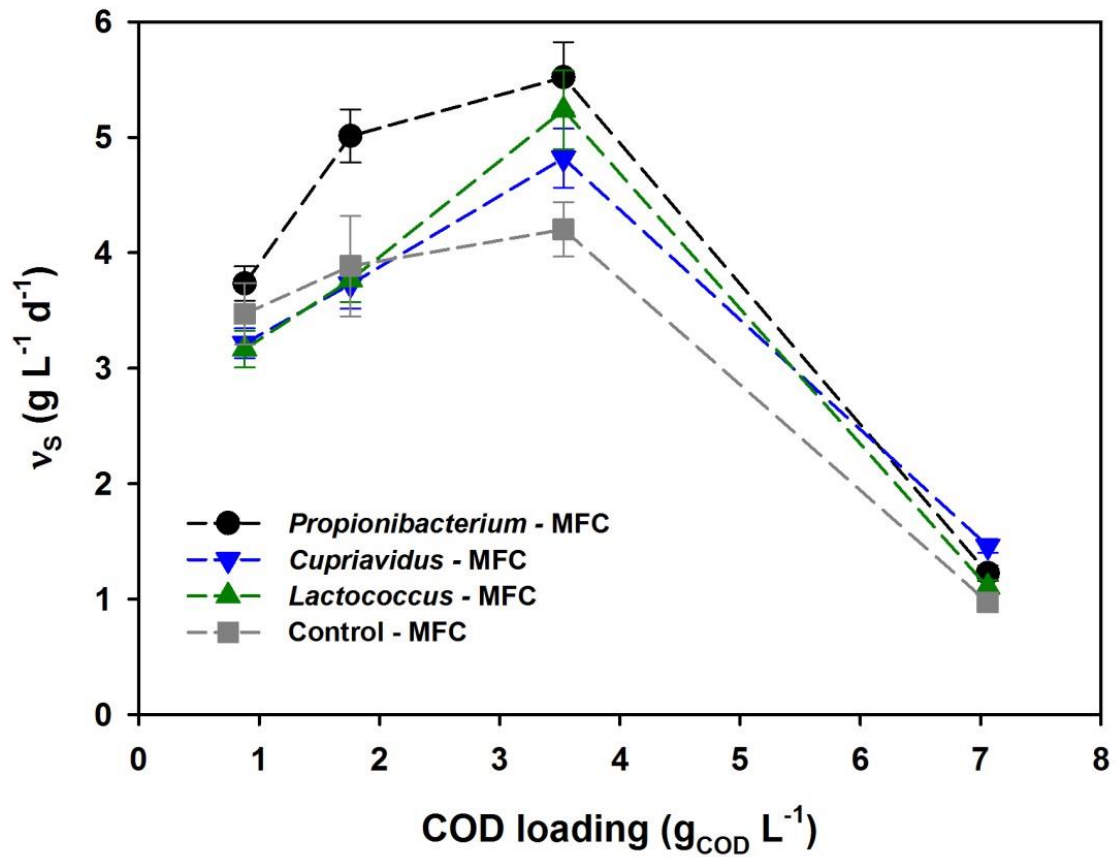
680

681 **Energy production rate of bioaugmented and control MFCs as a function**
682 **of COD loading. -●-: *Propionibacterium*-MFC; -▼-: *Cupriavidus*-MFC; -▲-:**
683 ***Lactococcus*-MFC; -■-: Control-MFC.**

684

685 Fig. 5

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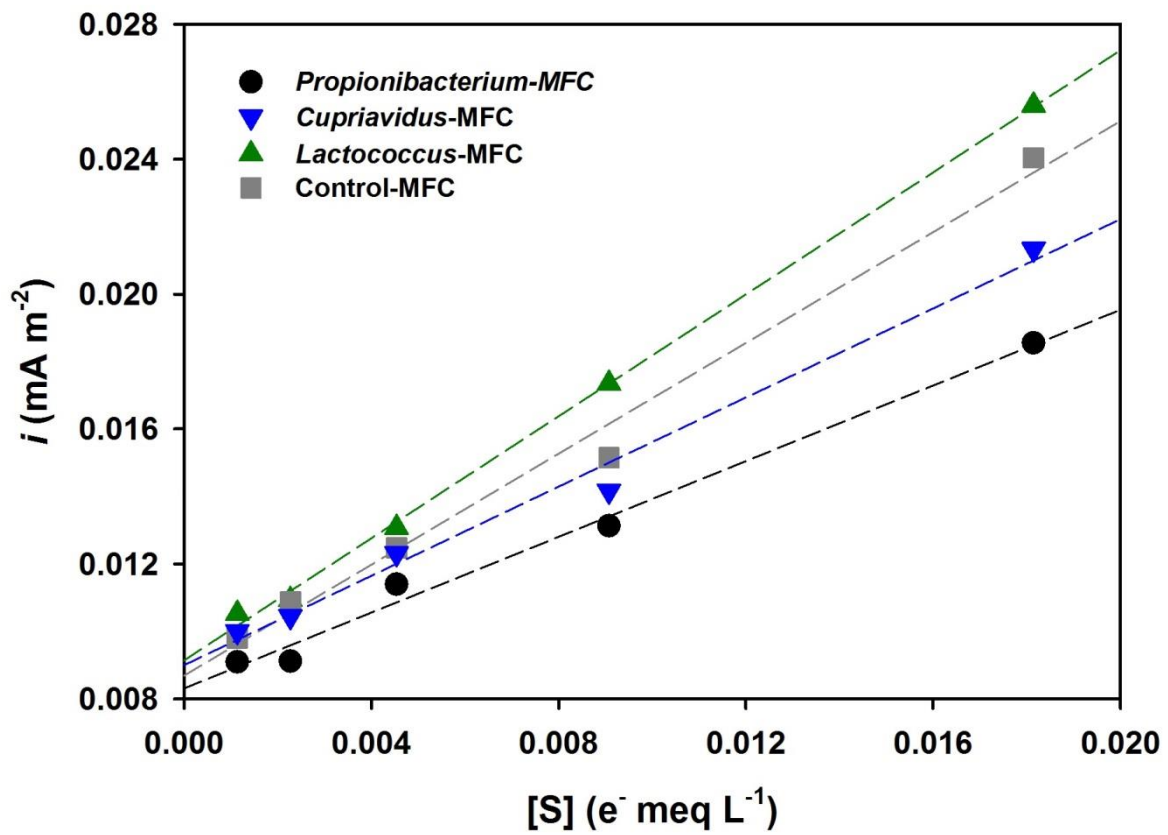
689 **COD removal rate of bioaugmented and control MFCs as a function of**

690 **COD loading. -●-: *Propionibacterium*-MFC; -▼-: *Cupriavidus*-MFC; -▲-:**

691 ***Lactococcus*-MFC; -■-: Control-MFC.**

692

693 Fig. 6



694

695

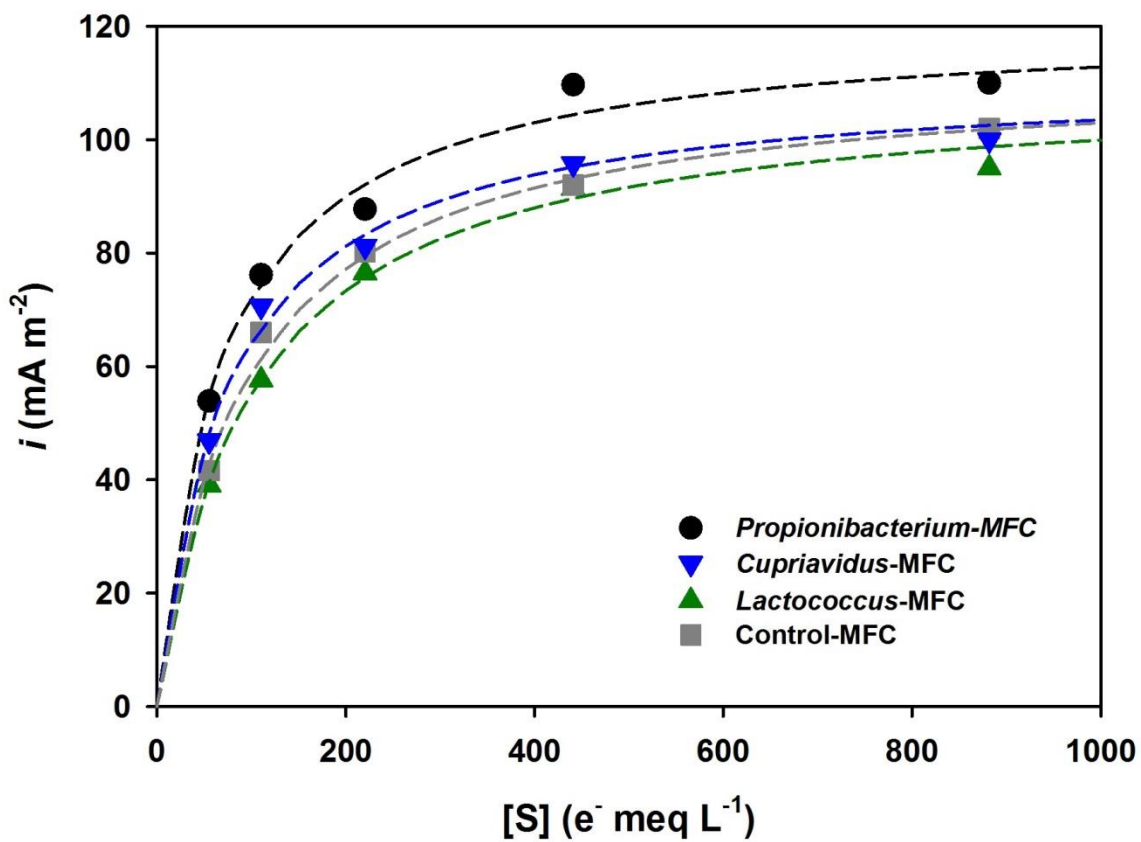
696 **Double-reciprocal plot of the Monod model for estimating the kinetic**

697 **parameters. -●-: *Propionibacterium*-MFC; -▼-: *Cupriavidus*-MFC; -▲-:**

698 ***Lactococcus*-MFC; -■-: Control-MFC.**

699

700 Fig. 7



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702

703 **Monod kinetics of the bioaugmented and control MFCs. -●-:**

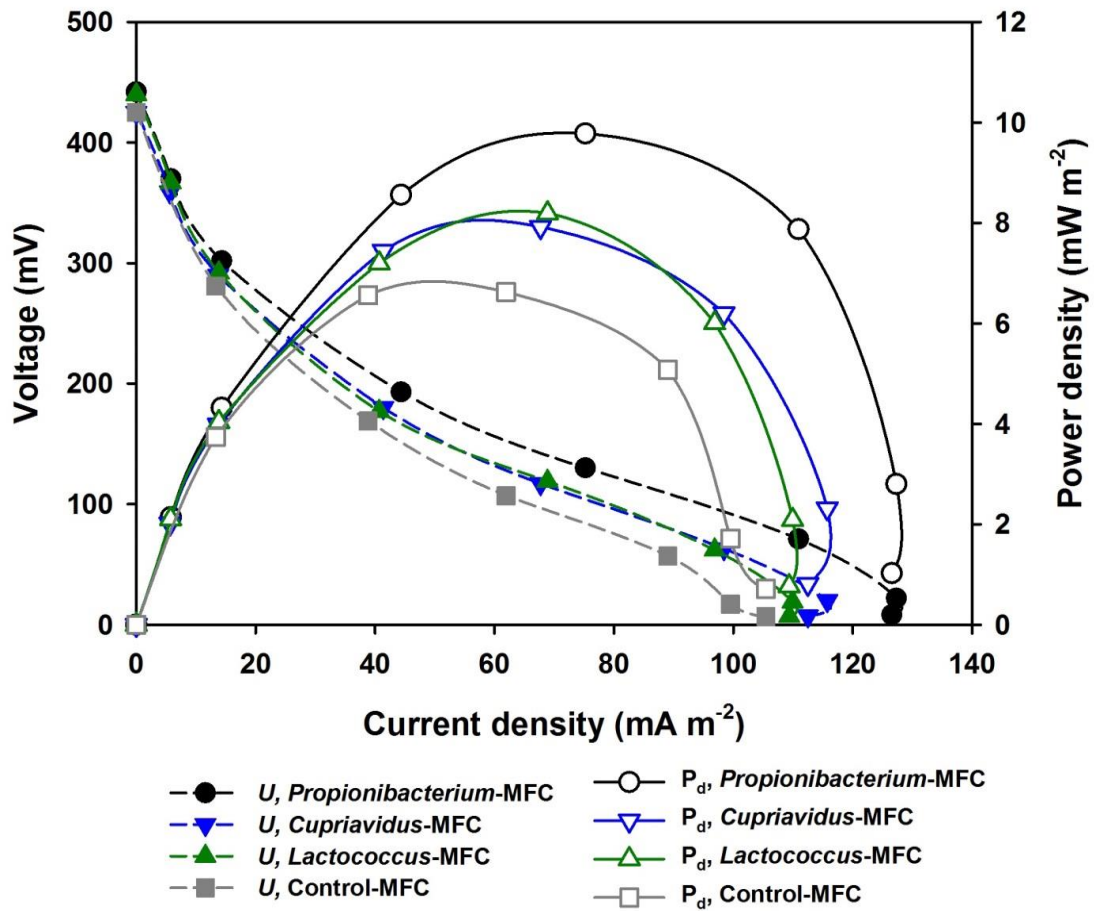
704 *Propionibacterium*-MFC; -▼-: *Cupriavidus*-MFC; -▲-: *Lactococcus*-MFC; -■-:

705

Control-MFC.

706

707 Fig. 8



708

709

710 **Polarization curves and power density plots for different MFCs.** -●-: *U*,
 711 *Propionibacterium*-MFC; -▼-: *U*, *Cupriavidus*-MFC; -▲-: *U*, *Lactococcus*-MFC;
 712 -■-: *U*, Control-MFC; -○-: P_d , *Propionibacterium*-MFC; -▽-: P_d , *Cupriavidus*-
 713 MFC; -△-: P_d , *Lactococcus*-MFC; -□-: P_d , Control-MFC.

714