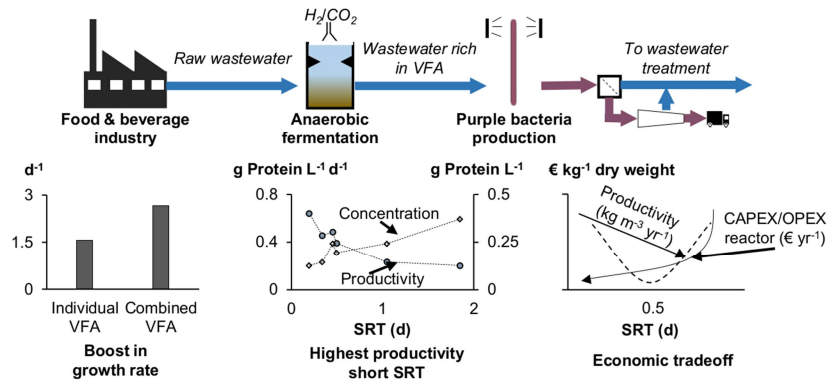


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1 **Volatile fatty acids impacting phototrophic growth kinetics of purple bacteria: paving**  
2 **the way for protein production on fermented wastewater**

3  
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14  
15 **Declaration of interest**

16 None.

17

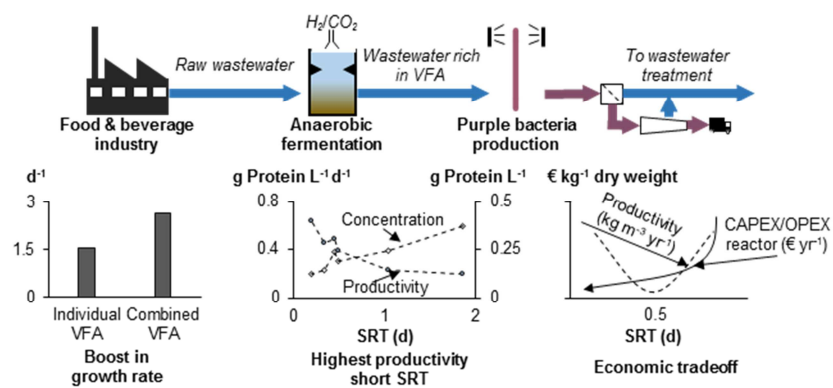
18

19 **Abstract**

20 Nutrient losses in our food chain severely surpass our planetary boundaries. Resource  
21 recovery can contribute to mitigation, for instance through converting wastewater resources to  
22 microbial protein for animal feed. Wastewater typically holds a complex mixture of organics,  
23 posing a challenge to selectively produce heterotrophic biomass. Ensuring the product's  
24 quality could be achieved by anaerobic generation of volatile fatty acids (VFAs) followed by  
25 photoheterotrophic production of purple non-sulfur bacteria (PNSB) with infrared light. This  
26 study aimed to determine the most suitable PNSB culture for VFA conversion and map the  
27 effect of acetate, propionate, butyrate and a VFA mixture on growth and biomass yield. Six  
28 cultures were screened in batch: (i) *Rhodopseudomonas palustris*, (ii) *Rhodobacter*  
29 *sphaeroides*, (iii) *Rhodospirillum rubrum*, (iv) a 3-species synthetic community (i+ii+iii), (v)  
30 a community enriched on VFA holding *Rb. capsulatus*, and (vi) *Rb. capsulatus* (isolate 'v').  
31 The VFA mixture elevated growth rates with a factor 1.3-2.5 compared to individual VFA.  
32 *Rb. capsulatus* showed the highest growth rates: 1.8-2.2 d<sup>-1</sup> (enriched) and 2.3-3.8 d<sup>-1</sup>  
33 (isolated). In a photobioreactor (PBR) inoculated with the *Rb. capsulatus* enrichment,  
34 decreasing sludge retention time (SRT) yielded lower biomass concentrations, yet increased  
35 productivities, reaching 1.7 g dry weight (DW) L<sup>-1</sup> d<sup>-1</sup>, the highest phototrophic rate reported  
36 thus far, and a growth rate of up to 5 d<sup>-1</sup>. PNSB represented 26-57% of the community and the  
37 diversity index was low (3-7), with a dominance of *Rhodopseudomonas* at long SRT and  
38 *Rhodobacter* at short SRT. The biomass yield for all cultures, in batch and reactor cultivation,  
39 approached 1 g COD<sub>Biomass</sub> g<sup>-1</sup> COD<sub>Removed</sub>. An economic estimation for a two-stage approach  
40 on brewery wastewater (load 2427 kg COD d<sup>-1</sup>) showed that 0.5 d SRT allowed for the lowest  
41 production cost (€10 kg<sup>-1</sup> DW; equal shares for capex and opex). The findings strengthen the  
42 potential for a novel two-stage approach for resource recovery from industrial wastewater,  
43 enabling high-rate PNSB production.

44 **Keywords:** short-chain fatty acids, single-cell protein, carboxylate platform, phototrophic  
 45 purple bacteria, carbon capture

46 **Graphical abstract**



## 49 **1 Introduction**

50 The conventional fertilizer–feed–food–fork chain is a cascade of inefficiencies (Steffen et al.,  
51 2015). For every 100 units nitrogen or phosphorus fertilizer applied to the land, only 14 units  
52 are consumed by citizens (Coppens et al., 2016). Upgrading resources from wastewater to  
53 microbial biomass could improve the efficiency of the food chain, with this protein-rich  
54 product used as ingredient in animal feed preparations. In this domain, affordability implies  
55 the need for non-axenic cultivation of microbial communities.

56 Wastewater contains organics, and heterotrophic microorganisms are therefore required to  
57 upgrade these streams to protein. However, wastewater typically holds a variable mixture of  
58 organic molecules of diverse complexity such as polysaccharides, proteins, fatty-acids, *N*-  
59 acetylamino sugars, DNA and polyphenolic structures (Dignac et al., 2000), posing a  
60 challenge to selectively and reproducibly produce heterotrophic biomass. Moreover, from a  
61 market perspective, resource recovery will be most attractive if microbial protein can be  
62 produced controllable with a constant quality (Verstraete et al., 2016). Therefore, a two-stage  
63 approach was conceptually proposed, anaerobically converting organics to volatile fatty acids  
64 (VFAs) followed by microbial protein production (Alloul et al., 2018).

65 Anaerobic fermentation is defined as the conversion of more complex organics to a mixture of  
66 VFA, rich in acetate, propionate and butyrate (Agler et al., 2011, Cagnetta et al., 2016). This  
67 conversion entails hydrolysis of particulate matter, followed by the production of acids  
68 (acidogenesis and acetogenesis) and CO<sub>2</sub> and H<sub>2</sub>. Methanogenesis, i.e. CH<sub>4</sub> formation, is  
69 avoided (Agler et al., 2011). This microbiological process allows to ‘normalize’ the organic  
70 variability over time (Ahn et al., 2001, Banerjee et al., 1999) and the produced VFA with the  
71 corresponding low pH might act as a means to reduce contamination of the influent. The  
72 toxicity of VFA for microorganisms is well described in literature (Sun and O’Riordan 2013)

73 and anaerobic digestion leads to a die-off of pathogens (Wagner et al., 2008). Similar results  
74 might therefore be expected for anaerobic fermentation.

75 Two metabolic types of heterotrophic bacteria are of particular interest for protein production,  
76 namely aerobic heterotrophic bacteria and photoheterotrophic bacteria (Verstraete et al., 2016).  
77 Aerobic heterotrophs have been explored for microbial protein production from the early  
78 1950s (Vriens et al., 1989) and are pioneering in both research and pilot implementation (Lee  
79 et al., 2015, Matassa et al., 2016a). They are characterized by high growth rates ( $2-6\text{ d}^{-1}$ ) and  
80 yields of  $0.44-0.55\text{ g C}_{\text{Biomass}}\text{ g}^{-1}\text{ C}_{\text{Removed}}$  (Tchobanoglous et al., 2003). Photoheterotrophically  
81 grown purple non-sulfur bacteria (PNSB) are also promising for microbial protein. They  
82 perform anoxygenic photosynthesis and use organics as electron donor and carbon source.  
83 Unlike algae, they are not able to use water as electron donor, but require molecules with  
84 lower redox potential than water such as VFAs (Blankenship et al., 1995). They grow at rates  
85 between  $1.51-1.69\text{ d}^{-1}$  for mixed cultures (Hülßen et al., 2014, Kaewsuk et al., 2010) and  
86 between  $0.96-7.10\text{ d}^{-1}$  for pure cultures (Noparatnaraporn et al., 1987, Ponsano et al., 2008).  
87 In addition, substrate-to-biomass organic C conversion yield is around one ( $0.8-1.2\text{ g C}_{\text{biomass}}$   
88  $\text{g}^{-1}\text{ C}_{\text{fed}}$  (Nakajima et al., 1997, van Niel 1944). Compared to aerobic heterotrophs, PNSB  
89 offer an additional steering tool to manage the microbial selectivity due to their unique ability  
90 to grow under anaerobic conditions on infrared (IR) light. In the case of sewage treatment for  
91 instance, with IR irradiation, a PNSB abundance of up to 90% was achieved (Hülßen et al.,  
92 2016a, Hülßen et al., 2016b).

93 To date, there are a number of research articles focusing on one-stage, direct, PNSB  
94 production on different types of industrial wastewater and domestic wastewater with pure  
95 cultures and open mixed cultures (Hülßen et al., 2016b, Kornochalart et al., 2014, Liu et al.,  
96 2016, Loo et al., 2013, Lu et al., 2013, Ponsano et al., 2011, Wu et al., 2015). For a two-stage  
97 approach, with anaerobic fermentation preceding the photoheterotrophic reactor, only two

98 approaches are available, starting from solid streams, implementing the fermentation stage as  
99 a liquefaction stage. One study has investigated fecal matter and plant material (Clauwaert et  
100 al., 2017) and another settled/diluted kitchen waste (Wang et al., 2016). On industrial  
101 wastewater from food/beverage, with the organics mainly in solution, such two-stage  
102 approach has not yet been explored. Previous studies that have investigated PNSB in a VFA-  
103 based medium, have studied the VFA biomass yield relationship for *Rhodobacter sphaeroides*  
104 (Nakajima et al., 1997, van Niel 1944), the effect of a VFA mixture on polyhydroxyalkanoate  
105 accumulation with an enrichment community holding *Rhodopseudomonas* sp. (Fradinho et al.,  
106 2014) and the molecular mechanism of VFA assimilation for *Rhodospirillum rubrum* (De  
107 Meur et al., 2018). A broader screening of PNSB for VFA growth kinetics and microbial  
108 selectivity imposed by VFA during non-axenic cultivation has according to the authors'  
109 knowledge not been investigated. Therefore, the first goal of this research was culture  
110 selection by determining the fastest growing PNSB on VFA along with their protein content  
111 and biomass yield. Six different cultures were investigated of which three pure cultures, a 3-  
112 species synthetic community to study potential synergetic effects, a community enriched on a  
113 VFA mixture to obtain a PNSB community specialized in VFA conversion and the isolated  
114 species to verify the findings of the latter. Secondly, this research wanted to unravel whether  
115 individual VFA or a VFA mixture had an impact on growth kinetics since anaerobic  
116 fermented wastewater typically consists of a VFA mixture. Afterwards, the enrichment  
117 community, which had obtained a very good performance during the batch test, was used as  
118 inoculum in a photobioreactor (PBR). Here, it was the objective to investigate the effect of  
119 sludge retention time (SRT) on microbial community dynamics and protein productivity. The  
120 findings of the PBR were finally used as input for an economic based optimization of the  
121 SRT. Production cost for anaerobic fermentation of brewery wastewater, PNSB cultivation,  
122 harvesting and drying were estimated and operational strategies were evaluated.



## 123 **2 Materials and methods**

### 124 **2.1 PSNB strains and communities**

125 To screen for the best PNSB culture for growth on VFA, six cultures were pre-selected: (i)  
126 *Rhodopseudomonas palustris* LMG 18881, (ii) *Rhodobacter sphaeroides* LMG 2827 and (iii)  
127 *Rhodospirillum rubrum* S 1H along with (iv) a 3-species synthetic community (i+ii+iii), (v)  
128 an enrichment community and (vi) *Rb. capsulatus* isolated from the latter through dilution to  
129 extinction. The pure cultures were axenically pre-cultivated in a climate chamber (Snijders  
130 Scientific) with a pre-autoclaved VFA-based medium adapted from Imhoff (2006). The  
131 details are presented in Supplementary Information S1.

132 The enrichment community was obtained by incubating a 1/1/1 volatile suspended solids ratio  
133 (0.1 g VSS L<sup>-1</sup> each) of activated sludge from a sewage treatment plant, activated sludge from  
134 a dairy wastewater treatment plant and sediment from a local pond with a VFA mixture and  
135 IR light to acquire a PNSB culture specialized in VFA conversion. After twenty-four hours of  
136 incubation carotenoid (320 and 380 nm) and bacteriochlorophyll peaks (590, 800 and 880 nm)  
137 appeared in the spectrum. The centrifuged biomass was then characterized by high-throughput  
138 16S rRNA sequencing (Illumina MiSeq; V4 region). The details are presented in  
139 Supplementary Information S2.

### 140 **2.2 Batch experiments**

141 The growth of the six abovementioned cultures (four strains and two communities) was  
142 examined in a 500 mL Erlenmeyer with a working volume of 400 mL through batch  
143 incubation. The experiments for the pure cultures and the 3-species synthetic community were  
144 performed in the climate chamber (conditions *ibid.* Supplementary Information S1). The  
145 experiment with the enrichment community was executed on a bench with IR transmission  
146 filters (conditions *ibid.* Supplementary Information S2). Three individual VFA were tested in  
147 triplicate (equal amount of carbon) along with a VFA mixture for every species and

148 community: (i) 0.73 g acetic acid L<sup>-1</sup>, (ii) 0.54 g propionic acid L<sup>-1</sup>, (iii) 0.60 g butyric acid  
149 L<sup>-1</sup> and (iv) combined VFAs (1/1/1 ratio on carbon mass basis: 0.24 g acetic acid L<sup>-1</sup>, 0.20 g  
150 propionic acid L<sup>-1</sup> and 0.18 g butyric acid L<sup>-1</sup>). The pH was adjusted to 7.0 prior to the  
151 experiment by the addition of 12 M NaOH. The flasks were then inoculated at an optical  
152 density of 0.1 (absorbance at 660 nm). The growth was monitored by measuring the  
153 absorbance at 660 nm. Samples were taken at the start (15 ml) and the end of the test (100  
154 mL).

### 155 **2.3 Photobioreactor experiment**

156 A non-axenic semi-continuous PBR was operated to study the effect of SRT on microbial  
157 community dynamics and protein productivity. The PBR was a vertical tubular vessel with a  
158 working volume of 2.75 L and an external diameter of 12 cm. It was operated for 51 days at a  
159 temperature of 30 ± 2 °C and illuminated with two halogen lamps at a light intensity of 30 W  
160 m<sup>-2</sup> covered with IR transmission filters. Stirring was done with a magnetic stirrer at 720 rpm  
161 (Fisher Scientific, USA). The reactor was inoculated with biomass from the enrichment  
162 community. The headspace of the reactor was flushed with nitrogen gas and the gas outlet  
163 was connected to a nitrogen gas expansion balloon in order to cope with underpressure and  
164 overpressure during withdrawal and fill. Influent and effluent tubing was covered with  
165 aluminum foil to minimize phototrophic growth.

166 A growth experiment of forty-eight hours with a VFA mixture (1/1/1 ratio on carbon mass  
167 basis: 0.73 g acetic acid L<sup>-1</sup>, 0.60 g propionic acid L<sup>-1</sup> and 0.54 g butyric acid L<sup>-1</sup>) was  
168 performed prior to reactor operation to determine the maximal possible biomass concentration  
169 before light limitations occurred (Supplementary Information S3). The maximal biomass  
170 concentration was converted to a VFA concentration at which growth would not be limited by  
171 carbon (i.e. 1 g chemical oxygen demand; COD L<sup>-1</sup> by assuming a yield of 1 g COD<sub>Biomass</sub> g<sup>-1</sup>  
172 COD<sub>VFA</sub>). The COD of the VFA mixture described in subsection 2.2 was adapted to 1 g COD

173 L<sup>-1</sup> and used as influent for the reactor. The pH of the PBR was not controlled, yet the influent  
174 pH was lowered with 12 M HCl in order to have a final pH of 7.0 in the effluent (pH rises due  
175 to VFA consumption).

176 The reactor was operated at six different SRT by altering the volumetric flow rate (L d<sup>-1</sup>). An  
177 overview of reactor conditions during operation is presented in Supplementary Information  
178 S4. The volume exchange ratio was initially set at 13% and later on altered to 25%. One up to  
179 four (four at short SRT) samples were taken daily to monitor the optical density (660nm), pH,  
180 temperature and electrical conductivity. Steady-state conditions were defined as a maximal  
181 change of 10% in optical density (proxy for biomass concentration), pH, temperature and  
182 electrical conductivity for three successive SRT. Samples for further analyses were taken  
183 three sequential SRT after steady-state conditions were reached.

#### 184 **2.4 Analytical procedures**

185 The COD was measured using photometric test kits (Merck, Germany). The biomass yield  
186 was determined by measuring removed COD (i.e. soluble COD<sub>start</sub> minus soluble COD<sub>end</sub>) and  
187 biomass COD (i.e. total COD minus soluble COD). Protein was determined according to an  
188 adapted Lowry procedure (Markwell et al., 1978). VFA were analyzed by High Performance  
189 Liquid Chromatography (Agilent technologies 1200) coupled with a diode array detector (210  
190 nm), a Bio-Rad Aminex<sup>®</sup> column (300 mm 7,8 mm) with Bio-Rad Micro-Guard Cation H  
191 Refill Cartridges and a column temperature of 40 °C. The samples were injected into 50 mM  
192 H<sub>2</sub>SO<sub>4</sub> (Honeywell Fluka<sup>TM</sup>) in deionised water (Arium<sup>®</sup> 611) with an injection volume of 20  
193 µL and a sample flow rate of 0.6 mL min<sup>-1</sup>. Total suspended solids (TSS) and VSS were  
194 determined according to standard methods (Greenberg et al., 1992). Handheld meters were  
195 used to measure pH (Hanna Instruments, USA) and electrical conductivity (Hanna  
196 Instruments, USA).

## 197 **2.5 Molecular microbial and phylogenetic analyses**

198 Molecular microbial analysis was used to characterize the most abundant PNSB in the  
199 enrichment, characterize the isolated species of the latter and follow-up richness, Shannon  
200 index, diversity and PNSB abundance during reactor operation. Phylogenetic relationships  
201 between the 16S rRNA gene sequence of the isolated species was done in order to classify the  
202 species.

203 A sample for microbial analysis was taken after four days of incubation. The most dominant  
204 species of the enrichment community was isolated using the dilution to extinction method.  
205 Biomass samples of the PBR were collected at three successive SRT when steady-state was  
206 reached. DNA was extracted using a PowerFecal® DNA isolation kit (QIAGEN, Germany)  
207 according to the manufacturer's instruction. The microbial community was analyzed using  
208 high-throughput 16S amplicon sequencing and a dual index strategy as described by Kozich et  
209 al., (2013). A detailed description can be found in Supplementary Information S5.

210 For the isolate, PCR products were sequenced using Sanger sequencing with both 8F and  
211 1525R primers (Genetic Service Facility, Antwerp). Editing, complete overlapping and  
212 consensus construction was performed with the Geneious software. The consensus sequences  
213 were annotated with sequences in the EZBioCloud database (Chunlab Inc., Korea).

214 MEGA7 software was used to conduct phylogenetic analysis (Kumar et al., 2016). The  
215 phylogenetic tree was built using the Maximum Likelihood method based on the General  
216 Time Reversible model (Nei and Kumar 2000). The percentage of trees in which the  
217 associated taxa clustered together is shown next to the branches. Initial trees for the heuristic  
218 search were obtained automatically by applying the Maximum Parsimony method. In total,  
219 1000 bootstrap replications were performed to test for branch robustness.

## 220 **2.6 Statistical analyses**

221 Multiple comparisons were performed using the parametric analysis of variance test and post-  
222 hoc pairwise comparisons using the Tukey's range test. Homogeneity of variances was  
223 verified by executing a Levene's test and normality of data residuals using the Shapiro–Wilk  
224 normality test. The non-parametric Kruskal-Wallis rank sum test and post-hoc pairwise  
225 comparisons using the Mann-Whitney U test (*p-values* were adjusted using the Benjamini-  
226 Hochberg correction) were performed when normality was rejected. A significance level of  $p$   
227  $< 0.05$  was chosen. All analyses were performed in R using RStudio (RStudio®, USA) for  
228 Windows (R Core Team 2017).

## 229 **2.7 Economic estimation**

230 An economic evaluation was performed to validate the optimal SRT for PNSB production,  
231 according to a four-step methodology which is thoroughly described in Supplementary  
232 Information S6. The following parameters were taken into account such as construction,  
233 piping, PBR, circulation pump, ultrafiltration unit, centrifuge, spray dryer etc. An overview of  
234 all cost parameters is presented in Supplementary Information Table S1. After PNSB  
235 production, wastewater still need to be treated to reach effluent discharge limits. The  
236 wastewater treatment cost that are prevented due to PNSB production (remove part of COD)  
237 are taken into account in the model, yet valorization of biogas is not included. Process steps  
238 that are animal specific such as nucleic acid removal, endotoxin removal and pasteurization  
239 were not taken into account in the estimation. In addition, our process is an open systems and  
240 thus potentially prone to pathogens. Therefore, we opt for a multiple barrier principle as  
241 discussed by Alloul et al., (2018) which entails the following steps: i) inactivation of  
242 pathogens during anaerobic fermentation, ii) solid/liquid separation after anaerobic  
243 fermentation iii) selective culture conditions (PNSB on infrared light), iv) drying and  
244 pasteurization of biomass.

245 The primary goal of this model was to determine the best SRT to operate the PBR. It was not  
246 intended to determine an accurate production cost of PNSB for brewery wastewater on full-  
247 scale which than can be used for other research. This cost estimation ought to be seen as a  
248 decision making tool for R&D and the corresponding production scheme can be used as  
249 example for further research.

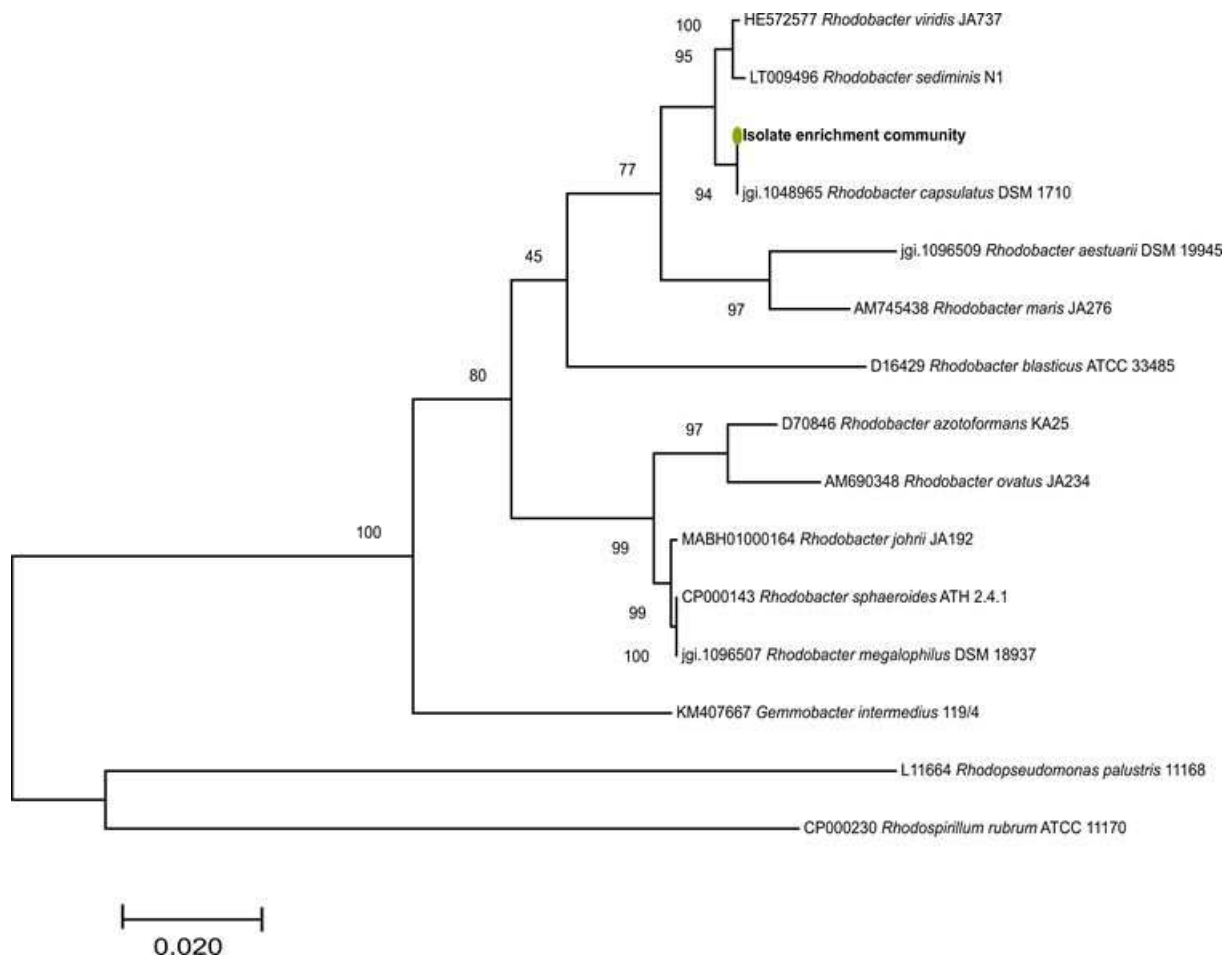
### 250 **3 Results and discussion**

#### 251 **3.1 Enrichment community**

252 The results of the microbial community analysis of the PNSB enrichment can be found in  
253 Supplementary Information S5 and show an almost complete dominance of PNSB (85%  
254 relative abundance). The community was dominated by an amplicon sequence variant with a  
255 relative abundance of 80% which was classified as *Rhodobacter/Gemmobacter* (EZBioCloud  
256 database). Two other PNSB genera were also present although in low abundance (3%  
257 *Rhodopseudomonas*, 2% *Pararhodospirillum*). Of note, the short V4 region of the 16S rRNA  
258 gene probably does not allow to differentiate between some *Rhodobacter/Gemmobacter*  
259 species. To further characterize the most abundant PNSB and determine whether the amplicon  
260 sequence variant corresponds to *Rhodobacter* or *Gemmobacter*, a dilution to extinction  
261 method was applied to obtain an isolate which was subjected to Sanger sequencing of the full  
262 16S rRNA gene (~1500 bp). Phylogenetic analysis of this sequence together with other  
263 closely related species showed that this isolate could be classified as a *Rhodobacter*  
264 *capsulatus* strain (Figure 1).

265 Due to dominance of *Rb. capsulatus* in the enrichment, we expected that growth performance  
266 of this strain would be above par. Growth rates for *Rb. capsulatus* between 4.9-5.4 d<sup>-1</sup> are  
267 described in literature (Willison 1988), which are lower than values found for *Rb. sphaeroides*  
268 (7.4 d<sup>-1</sup>) (Sangkharak and Prasertsan 2007). However, both results are for pure cultures

269 derived from mutant-strains. It is therefore difficult to draw conclusive statement about the  
 270 growth performance of *Rb. capsulatus* on VFA. Therefore, in section 3.2 we will compare  
 271 growth rates for several culture on individual and combined VFAs.



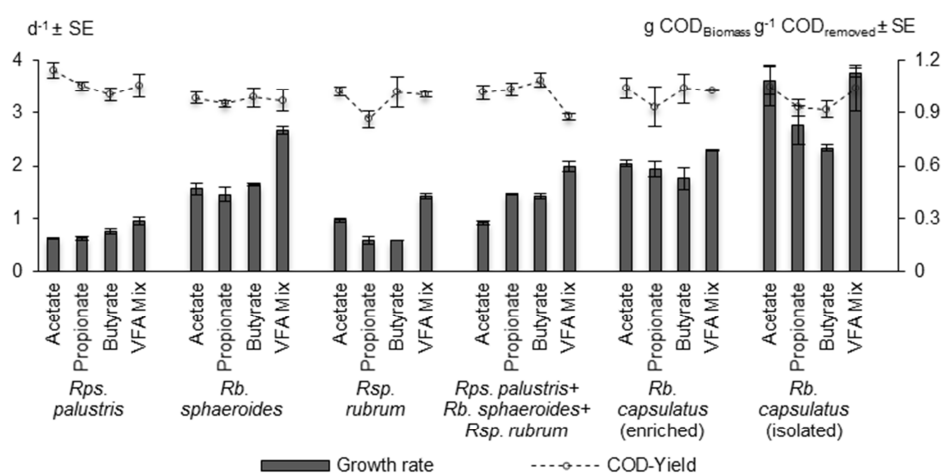
272  
 273 **Figure 1** Phylogenetic relationship between the 16S rRNA gene sequence of an isolate of the  
 274 enrichment, the three tested species, other species with 100, 99 and 98% similarity blast, and  
 275 other *Rhodobacter capsulatus* strains (EZBioCloud database)

### 276 3.2 PNSB growth kinetics for individual and combined VFA

277 Batch experiments were performed to explore the effect of individual and combined VFAs on  
 278 the biomass yield and growth rate, and screen within the six cultures for the most productive,  
 279 i.e. fastest growing microorganism on VFA.

280 Figure 2 presents the biomass yield which was overall between 0.9-1.1 g COD<sub>Biomass</sub> g<sup>-1</sup>  
 281 COD<sub>Removed</sub>. There was no significant difference ( $p > 0.05$ ) between individual VFA or

282 individual and combined VFAs or between different species and communities. PNSB have a  
 283 biomass yield equal to  $1 \text{ g COD}_{\text{biomass}} \text{ g}^{-1} \text{ COD}_{\text{removed}}$  and even higher due to photo-  
 284 assimilation of a highly reduced electron donor (i.e. propionate and butyrate) along with  $\text{CO}_2$   
 285 that serves as electron sink to compensate for excess reducing power (Blankenship et al.,  
 286 1995). Nakajima et al., (1997) and van Niel (1944) have also observed a biomass yield of  
 287 respectively 0.8 and between 0.8-0.9  $\text{g COD}_{\text{Biomass}} \text{ g}^{-1} \text{ COD}_{\text{Removed}}$ , yet these researchers only  
 288 studied *Rb. sphaeroides*. The near-perfect substrate-to-biomass conversion is a critical  
 289 advantage of PNSB compared to aerobic heterotrophs as roughly a double production  
 290 potential is possible with a given amount of COD in the wastewater (yield  $0.57 \text{ g COD}_{\text{biomass}}$   
 291  $\text{g}^{-1} \text{ COD}_{\text{removed}}$ ; Tchobanoglous et al., 2003).



292  
 293 **Figure 2** Overview of growth rate (left y-axis) and yield (right y-axis) for six cultures for  
 294 individual and combined volatile fatty acids (VFAs). Error bars show standard error; COD:  
 295 chemical oxygen demand.

296 For the growth rate, a change in individual VFA had no effect, as observed for all species and  
 297 communities (Figure 2). The most remarkable observation during the batch test is the boost in  
 298 growth rate when PNSB are fed with combined VFAs. This enhanced growth rate effect was  
 299 noticeable for *Rps. palustris*, *Rb. sphaeroides*, *Rsp. rubrum* and the 3-species synthetic



300 community. The boost in growth rate was between 1.3-2.5 times higher compared too  
301 individual VFA, most pronounced for *Rsp. rubrum*. The absolute value for the enriched *Rb.*  
302 *capsulatus* and the isolated *Rb. capsulatus* was higher for combined VFAs, yet there was no  
303 significant difference between acetate and the VFA mixture ( $p > 0.05$ ). The study of  
304 Nakajima et al., (1997) did not observe a growth rate enhancement effect for individual VFA  
305 (acetate, propionate and butyrate) compared to three different combined VFA ratios (1/1/1;  
306 2/1/1 and 12/2/1 g carbon for respectively acetate, propionate and butyrate). This is probably  
307 due to the lack of sampling points during the exponential phase resulting in flattening down of  
308 the exponential correlation. Moreover, only *Rb. sphaeroides* was tested without any  
309 replicates, which makes it difficult to draw general conclusions. More recently, the study of  
310 Fradinho et al., (2014) has observed higher growth rates (consumption rate multiplied by  
311 biomass yield) for combined VFAs ( $1.53 \text{ d}^{-1}$ ) compared to acetate ( $1.22 \text{ d}^{-1}$ ) in line with our  
312 results. They studied the effect of a VFA mixture (4/1/1 g carbon) on polyhydroxyalkanoate  
313 accumulation with an enrichment community holding *Rhodopseudomonas* sp. The effect of  
314 combined VFAs is probably due to the presence of acetate as co-substrate. Without acetate the  
315 consumption of propionate and butyrate is limited by an internal acetyl-CoA deficit (Fradinho  
316 et al., 2014). Overall, our study is the first broader screening of PNSB for growth on VFA.  
317 The results highlight that a VFA mixture, and therefore likely fermented wastewater, is a  
318 more suitable substrate to produce PNSB at high rate compared to individual VFA. However,  
319 the VFA ratio can change for a fermentate depending on the operational conditions (Lee et al.,  
320 2014). Future research should therefore further unravel the effects of different VFA carbon  
321 ratios.

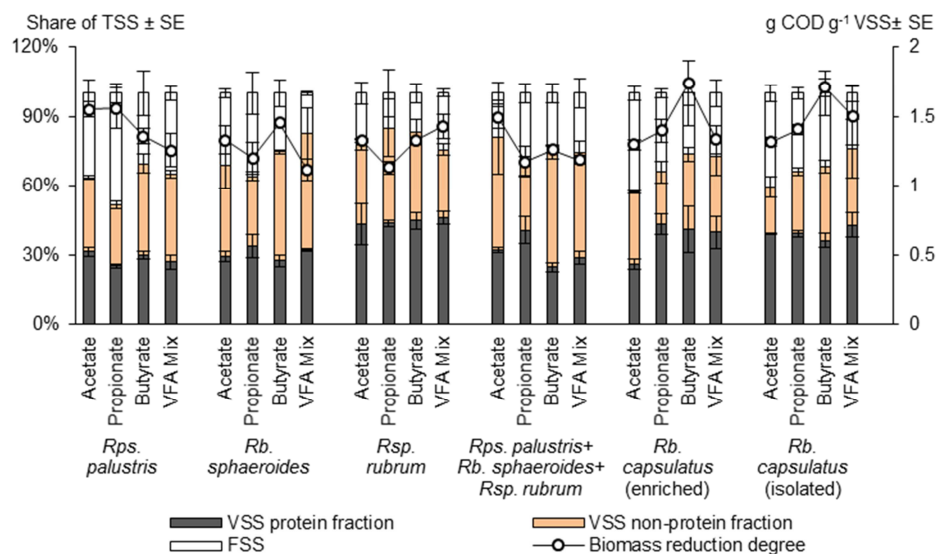
322 The isolated *Rb. capsulatus* had the highest growth rates for all individual and combined  
323 VFAs. Therefore, this culture might have the best potential to grow on fermentate. Due to the  
324 context of the application, which is resource recovery on wastewater, we choose to proceed

325 with a non-axenic reactor and non-axenic culture, which is the enriched *Rb. capsulatus*.  
326 Growth rates of the isolated *Rb. capsulatus* were 1.3-1.8 times higher compared to the  
327 enriched *Rb. capsulatus*. A similar observation can be made for the 3-species synthetic  
328 community which was the mixture of *Rps. palustris*, *Rb. sphaeroides*, *Rsp. rubrum*. Growth  
329 rates of *Rb. sphaeroides* (best individual performance of three species) for acetate and the  
330 combined VFAs were respectively 1.4 and 1.7 times higher compared to the 3-species  
331 synthetic community. This compares to the results of the *Rb. capsulatus* isolate which also  
332 had higher growth rates compared to the enrichment community (mixed culture). PNSB  
333 species share the same resources, therefore a competitive interaction in communities could be  
334 present which might result in detrimental effects for all species (Little et al., 2008).

335 The results of this study are also relevant for regenerative life support systems such as the  
336 MELiSSA loop, currently using the PNSB *Rsp. rubrum* for VFA conversion (Clauwaert et al.,  
337 2017). Our results point towards *Rb. capsulatus* as a more suitable culture for intensified  
338 protein production. Indeed, *Rb. capsulatus* is 2.6 times faster compared to *Rsp. rubrum*, and  
339 will allow for a more compact and lightweight bioreactor, a key advantage for space missions.  
340 However, also other factors ought to be taken into account for culture selection such as the  
341 broader metabolic capabilities, production of metabolites and the ability to maintain genetic  
342 stability under more severe radiation (Clauwaert et al., 2017).

343 Figure 3 presents an overview of the biomass composition (protein, non-protein VSS and  
344 fixed suspended solids; FSS). No general trend could be observed in protein content between  
345 individual or individual and combined VFAs. The overall protein content of *Rsp. rubrum* was  
346 significant higher than *Rps. palustris*, *Rb. sphaeroides* and equal to the enriched *Rb.*  
347 *capsulatus* and isolated *Rb. capsulatus* and the 3-species synthetic community. Vрати (1984)  
348 observed protein-levels that were between 1 and 2 times higher than our results. Growth in a  
349 medium with carbohydrates and a higher nitrogen concentration might potentially affect the

350 protein content. Vradi (1984) used clarified cow dung slurry as medium which contained 4 g  
 351 L<sup>-1</sup> carbohydrates and 10 times more nitrogen than our medium. On the contrary, a VFA-  
 352 based medium was used in our tests. The degree of biomass reduction or biomass COD:VSS  
 353 ratio was on average 1.35 g COD g<sup>-1</sup> VSS (Figure 3), which is comparable with values of  
 354 activated sludge (1.20-1.49 g COD g<sup>-1</sup> VSS; Contreras et al., 2002).



355 **Figure 3** Biomass composition of all batch experiments (left y-axis), as breakdown of total  
 356 suspended solids (TSS) into volatile suspended solids (VSS) protein fraction, VSS non-  
 357 protein fraction, fixed suspended solids (FSS) and as biomass reduction degree (right y-axis).  
 358 Error bars show standard errors.

### 360 3.3 Maximizing the protein productivity in a PBR on a VFA mixture

361 A non-axenic semi-continuous PBR was operated as chemostat, and the effect of SRT on  
 362 microbial community dynamics and protein productivity (g protein L<sup>-1</sup> d<sup>-1</sup>) was studied. In the  
 363 batch experiments, the *Rb. capsulatus* cultures had growth rates of 3.8 d<sup>-1</sup> on combined VFAs.  
 364 Therefore, it was expected that PNSB would be washed out at the corresponding dilution rate,  
 365 representing a SRT below 0.3 d. However, a shorter SRT of even 0.19 d (4.6 h) was achieved  
 366 and *Rhodobacter* was still abundantly present (**Figure 4**).

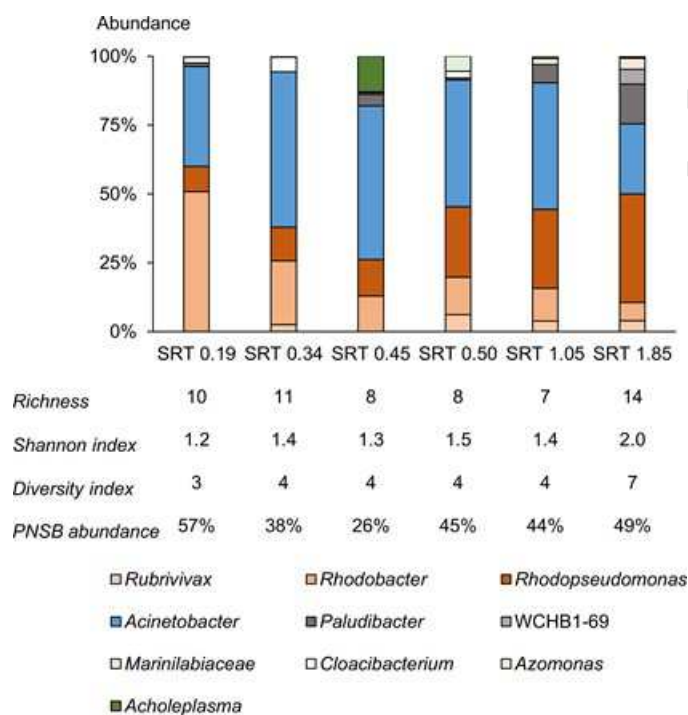
367 It was anticipated that implementation of high-rate conditions would impose an additional  
368 selective pressure on the microbial community favoring PNSB (cf. fast growth on VFA and  
369 IR light), leading to washout of slow growing non-PNSB microorganisms. The minimal  
370 attainable SRT of a species (i.e.  $1/\mu_{\max}$ ) is during washout higher than the actual SRT imposed  
371 on the reactor. Therefore, production of new cells will be lower than the amount that is  
372 washed out (Saikaly and Oerther 2004). This hypothesis was in line with the findings of  
373 Matassa et al., (2016b). They have operated a continuous bioreactor using hydrogen-oxidizing  
374 bacteria for protein production at a SRT of 10 h. The microbial community was almost  
375 completely dominated by hydrogen-oxidizing bacteria. However, washout effects might be  
376 more complex for mixed cultures than for pure cultures. For example, the SRT at which  
377 washout occurs can differ because the available COD for each species will be lowered by the  
378 presence of other species (Saikaly and Oerther 2004).

379 The results of 16S rRNA gene sequencing showed relatively high PNSB dominance during  
380 operation (relative abundance around 27-60%; Figure 4), albeit lower than the PNSB  
381 abundance in the inoculum enrichment community (85%, Supplementary Information Figure  
382 S3). Species richness was highest at a SRT of 2 d, decreased to 8 at a SRT of 0.5 d and finally  
383 increased again to 10-11 at SRT values between 0.2-0.3 d. No extra selective pressure  
384 favoring PNSB growth could be observed at shorter SRT based on the results of the diversity  
385 index. However, a selective microbial community (low diversity index) with relatively high  
386 PNSB dominance could be maintained over several SRT in the PBR.

387 Three PNSB genera could be detected namely *Rhodobacter*, *Rhodopseudomonas* and  
388 *Rubrivivax* of which the former two were most abundant. The SRT had an effect on relative  
389 abundance between different PNSB genera. The genus *Rhodopseudomonas* was most  
390 dominant at long SRT, while *Rhodobacter* at short SRT. A clear distinction between the  
391 enriched *Rb. capsulatus* community and the microbial community during reactor operation is

392 the strong presence of *Acinetobacter* (relative abundance around 25-56%). Members of this  
 393 genus were also found in other studies investigating open mixed PNSB cultures with relative  
 394 abundances of 17.3% (Wang et al., 2016), 0.05-0.6% (Hülßen et al., 2016a) and qualitative  
 395 detected through DGGE followed by sequencing (Izu et al., 2001). These gram-negative  
 396 bacteria are known for their ability to accumulate phosphate, as employed in the enhanced  
 397 biological phosphorus removal process (Kim and Pagilla 2000). As phosphate accumulating  
 398 organisms are typically stimulated with alteration in anaerobic and aerobic conditions. It is  
 399 possible that oxygen entered the PBR during withdrawal of the effluent (higher headspace to  
 400 volume ratio).

401

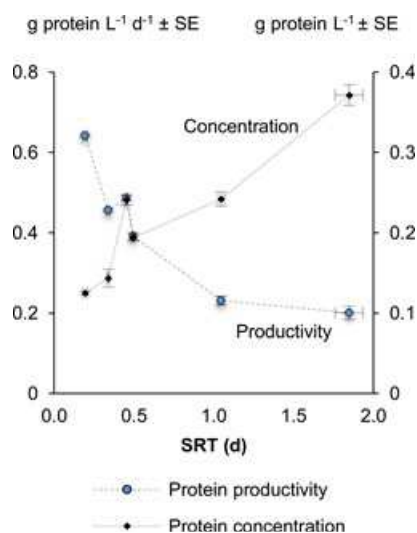


402

403 **Figure 4** Effect of sludge retention time (SRT) on microbial community composition,  
 404 richness, Shannon index, diversity index which is the exponential of the Shannon index and  
 405 purple non-sulfur bacteria (PNSB) abundance during reactor operation. PNSB genera  
 406 *Rhodobacter*, *Rhodopseudomonas* and *Rubrivivax* are all marked in orange colors.

407 Figure 5 presents an overview of the effect of SRT on protein concentration (g protein L<sup>-1</sup>)  
408 and protein productivity (g protein L<sup>-1</sup> d<sup>-1</sup>, protein concentration multiplied by dilution rate).  
409 A general trend that can be observed is the decrease in steady-state protein concentration due  
410 to higher biomass washout imposed by shorter SRT, yet the protein productivity increased.  
411 The highest protein productivity was 0.64 g protein L<sup>-1</sup> d<sup>-1</sup> or expressed in dry weight 1.7 g  
412 TSS L<sup>-1</sup> d<sup>-1</sup> achieved at a SRT of 4.6 h. Carlozzi and Sacchi (2001) have also obtained high  
413 photoheterotrophic productivities (1.1-1.4 g TSS L<sup>-1</sup> d<sup>-1</sup>) in an outdoor PBR with *Rps.*  
414 *palustris* as inoculum on a VFA-based medium, yet results are still lower than our findings  
415 probably due to natural light-dark cycles. To conclude, the TSS productivity of 1.7 g TSS L<sup>-1</sup>  
416 d<sup>-1</sup> is according to the authors' knowledge the highest reported in a PBR for a PNSB  
417 enrichment community and the first research to realize high-rate PNSB production (short  
418 SRT).

419 An important parameter in designing a PBR on fermented wastewater is the biomass specific  
420 removal rate, which was  $4.5 \pm 1.4$  g COD<sub>Removed</sub> g<sup>-1</sup> COD<sub>Biomass</sub> d<sup>-1</sup> (SRT 4.6 h). In literature  
421 values for acetate, propionate and butyrate of respectively 1.5-2.4, 2.0-1.3, 0.1-1.7 g  
422 COD<sub>Removed</sub> g<sup>-1</sup> COD<sub>Biomass</sub> d<sup>-1</sup> are reported (Fradinho et al., 2014, Puyol et al., 2017). These  
423 high specific removal rates make PNSB competitive to high-rate aerobic heterotrophic  
424 bacteria (1-6 g COD<sub>Removed</sub> g<sup>-1</sup> COD<sub>Biomass</sub> d<sup>-1</sup>) (Meerburg et al., 2015). However, the attainable  
425 biomass concentrations are higher for aerobic heterotrophs (not limited by light), and  
426 therefore the volumetric COD removal rates will also be higher, compared to phototrophically  
427 grown PNSB.



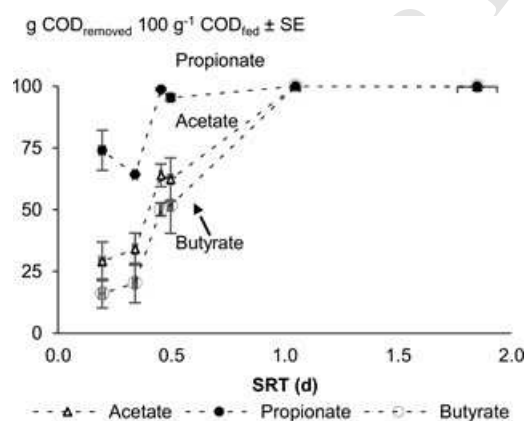
428  
429

430 **Figure 5** Effect of sludge retention time (SRT) on protein productivity (left y-axis) and  
431 protein concentration (right y-axis) for photoheterotrophic biomass production on VFA  
432 mixture (1/1/1 ratio on carbon mass basis: 0.73 g acetic acid L<sup>-1</sup>, 0.60 g propionic acid L<sup>-1</sup> and  
433 0.54 g butyric acid L<sup>-1</sup>) a semi-continuous photobioreactor (PBR). Error bars show standard  
434 error.

435 Figure 6 presents an overview of the VFA removal efficiency. All VFA are consumed at a  
436 SRT between 1-2 d, yet acetate and butyrate are washed out at a SRT of 0.5 d and lower. On  
437 the contrary, the removal efficiency for propionate remained almost unchanged at a SRT of  
438 around 0.5 d and was still 74 g COD<sub>Removed</sub> 100 g<sup>-1</sup> COD<sub>Fed</sub> at a SRT of 4.6 h. Hence, these  
439 results show that the microbial community had a preference for propionate over butyrate and  
440 acetate. De Meur et al., (2018) has studied binary VFA mixtures with *Rsp. rubrum* and  
441 observed a lag for butyrate uptake for the mixtures acetate/butyrate and propionate/butyrate,  
442 yet acetate and propionate were simultaneous assimilated. The lag in uptake is explained by  
443 De Meur et al., (2018) as a metabolic mismatch due to the fact that acetate and butyrate share  
444 acetyl-CoA and crotonyl-CoA as metabolic intermediates. It probably exists for  
445 microorganisms that are isocitrate lyase negative (ICL<sup>-</sup>) and use other pathways for VFA  
446 assimilation (ethylmalonyl-CoA pathway) for example *Rsp. rubrum* and *Rb. sphaeroides*

447 (Alber et al., 2006, Kornberg and Lascelles 1960). It is difficult to extrapolate the findings of  
 448 De Meur et al., (2018) to our results because it is not possible to verify which PNSB are  
 449  $ICL^+/ICL^-$  on a genus level and it is not known how non-PNSB species contribute to the  
 450 specific VFA removal. More research is still required to unravel this at metabolic level for  
 451 different PNSB. Recently Puyol et al., (2017) have published a phototrophic model for  
 452 domestic wastewater treatment in anaerobic conditions. The effect of acetate/propionate on  
 453 butyrate uptake is not included, yet our findings show that it can be crucial. New models  
 454 should take these effects into account. These findings are also key when operating a PBR on  
 455 fermented or on synthetic medium as acetate and butyrate washout can occur at short  
 456 hydraulic retention time (HRT) or high loading rates.

457 Full VFA removal ( $100 \text{ g COD}_{\text{Removed}} 100 \text{ g}^{-1} \text{ COD}_{\text{Fed}}$ ) at 1 and 2 d SRT showed that PNSB  
 458 were actually underfed and thus probably higher volumetric productivities are achievable at  
 459 these SRT. This was initially not observed because the soluble COD at a SRT of 1 and 2 d  
 460 was around  $230 \text{ mg COD L}^{-1}$ , yet VFA analysis showed that the VFA concentration was zero.  
 461 The excess in soluble COD could possibly be due to the production of extracellular enzymes  
 462 (e.g. vitamin lipase, protease, amylase etc. ; Chumpol et al., 2017).



463  
 464 **Figure 6** Volatile fatty acid (VFA) removal efficiency for acetate, propionate and butyrate as  
 465 function of the sludge retention time (SRT). VFA mixture 1/1/1 ratio on carbon mass basis:



466 0.73 g acetic acid L<sup>-1</sup>, 0.60 g propionic acid L<sup>-1</sup> and 0.54 g butyric acid L<sup>-1</sup>. Error bars show  
467 standard errors (SE).

### 468 **3.4 Economic tradeoff to determine optimal SRT**

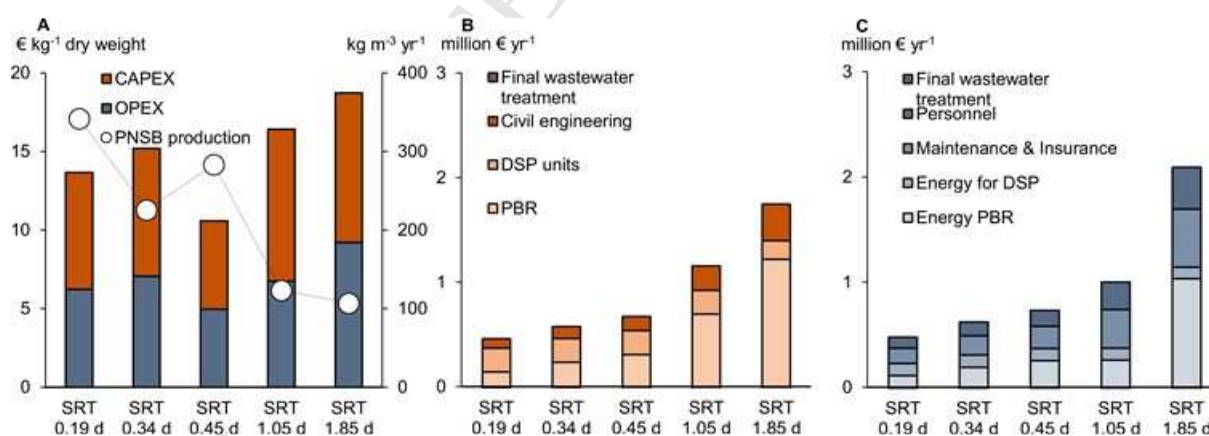
469 The reactor cultivation showed an opposite influence of SRT on protein productivity and  
470 harvestability (biomass concentration; Figure 5). A cost estimation on brewery wastewater  
471 corrected for normal costs of wastewater treatment (flow 1150 m<sup>3</sup> d<sup>-1</sup> and COD-load 2427 kg  
472 COD d<sup>-1</sup>) was therefore performed to determine the optimum SRT for reactor operation,  
473 including anaerobic fermentation, the PBR, downstream processing (harvesting and drying)  
474 and final wastewater treatment after PNSB production to reach discharge limits. A process  
475 scheme is presented in Supplementary Information S6 which depicts the production of PNSB  
476 and the brewery wastewater treatment system after PNSB production.

477 The share of capital- and operational expenditure (CAPEX and OPEX) relative to total cost  
478 was respectively between 51-59% and 41-49%. In general, both CAPEX and OPEX were  
479 dominated by the cultivation stage, i.e. the PBR (Figure 7). With decreasing SRT, biomass  
480 production increased from 107 to 342 tonnes m<sup>-3</sup> yr<sup>-1</sup>. Three production zones can be  
481 recognized: (i) high productivity yet high expenses (SRT < 0.5 d), (ii) most economical  
482 production (SRT of 0.5 d and cost of € 10 kg<sup>-1</sup> dry weight; DW) and (iii) low productivity and  
483 high expenses (SRT > 0.5 d).

484 The production cost of PNSB is comparable to microalgae (€ 5.96-12.6 kg<sup>-1</sup> DW) (Acien et  
485 al., 2012, Norsker et al., 2011), yet could be reduced if sunlight would be used. The CAPEX  
486 for the artificial light system is not costly (0.5-1.5%), yet the energy for light is between 13-  
487 28% of the OPEX. Sunlight PNSB production could theoretically lower the production cost to  
488 € 9 kg<sup>-1</sup> DW. PNSB production on sunlight is possible at high productivities as proven by  
489 (Carlozzi et al., 2006). However for temperate climates it should be demonstrated that these  
490 high productivities are still reached during colder and darker months.

491 PNSB production will reduce the COD concentration and therefore a portion of the  
 492 wastewater treatment cost will be exempted. Overall, savings due to avoided treatment cost  
 493 were between € 29,000-68,000 yr<sup>-1</sup>, which accounts to 13-31% of the wastewater treatment  
 494 costs without PNSB production. More savings are possible if productivities further increased  
 495 because final wastewater treatment can then be simplified to aerobic treatment and no  
 496 additional digester is needed. Decoupling SRT from HRT by a membrane bioreactor for  
 497 example (Hülßen et al., 2016b) could increase the productivity due to higher volumetric  
 498 removal rates imposed by a higher biomass concentration. For now, wastewater treatment  
 499 after PNSB production with merely aerobic treatment was not possible because the effluent  
 500 COD concentration remained too high (1.54 g COD L<sup>-1</sup>) and nutrients too low (14 mg N L<sup>-1</sup>  
 501 and 0.3 mg P L<sup>-1</sup>). If PNSB production was optimized, only aerobic treatment would be  
 502 required. This would result in € 130,000 yr<sup>-1</sup> extra savings for the treatment system (no  
 503 nutrient addition taken in to account).

504 Overall, this cost assessment shows that a SRT of 0.5 d is most optimal for PNSB production,  
 505 and production costs could potentially be optimized by sunlight.



506  
 507 **Figure 7** Economic tradeoff based on the production cost of photoheterotrophic biomass (left  
 508 Y axis, A) as function of decrease in productivity (right Y axis, A) and increase in  
 509 harvestability for longer sludge retention time (SRT). Distribution of capital expenditure  
 510

511 (CAPEX, B) and operational expenditure (OPEX, C). DSP: downstream processing, PBR:  
512 photobioreactor

#### 513 **4 Conclusions**

514 (i) Combining the VFAs acetate, propionate and butyrate resulted in a boost in growth  
515 rate compared to the individual VFA, for all tested PNSB cultures. Anaerobic fermentate of  
516 wastewater consists of a VFA mixture, and might result in the same effect.

517 (ii) PBR experiments showed that propionate was the preferential VFA source when  
518 dosing combined VFAs. There is hence no use in overdosing acetate or butyrate in a synthetic  
519 medium or to steer fermentation to butyrate production. Acetate and butyrate washout can be  
520 problematic for a PBR at short HRT or at high loading-rates.

521 (iii) PNSB have a near-perfect substrate-to-biomass COD conversion when grown on  
522 VFA. This roughly doubles the production potential compared to aerobic heterotrophs.

523 (iv) This research is first to screen a broader range of PNSB for growth on VFA. From six  
524 PNSB cultures, the isolated and enriched *Rhodobacter capsulatus* obtained the highest growth  
525 rates.

526 (v) A biomass productivity of  $1.7 \text{ g DW L}^{-1} \text{ d}^{-1}$  was achieved in the PBR, the highest rate  
527 reported for a phototrophic PNSB enrichment community with a focus on production.

528 (vi) Biomass-specific removal rates of  $4.5 \pm 1.4 \text{ g COD}_{\text{Removed}} \text{ g}^{-1} \text{ COD}_{\text{Biomass}} \text{ d}^{-1}$  were  
529 observed in the PBR. PNSB are therefore competitive to aerobic heterotrophic bacteria  
530 (alternative source of microbial protein) on a biomass-specific basis, yet cannot compete on a  
531 volumetric basis.

532 (vii) PNSB are able to cope with very short SRT. Their relative abundance was between  
533 26-60% and the community diversity index was low (3-7). VFA and IR therefore seem  
534 suitable tools to enhance selectivity in a non-axenic PBR.

535 (viii) *Rb. capsulatus* might be a more suitable culture than *Rsp. rubrum* for regenerative life  
536 support systems for space with a focus on compact reactors.

537 (ix) The reactor cultivation showed an opposite influence of SRT (0.19-1.85 d) on protein  
538 productivity and harvestability. An economic estimation for PNSB production enabled to the  
539 derivation of 0.5 days SRT as the most cost-efficient for biomass production, at a cost of € 10  
540 kg<sup>-1</sup> DW.

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551

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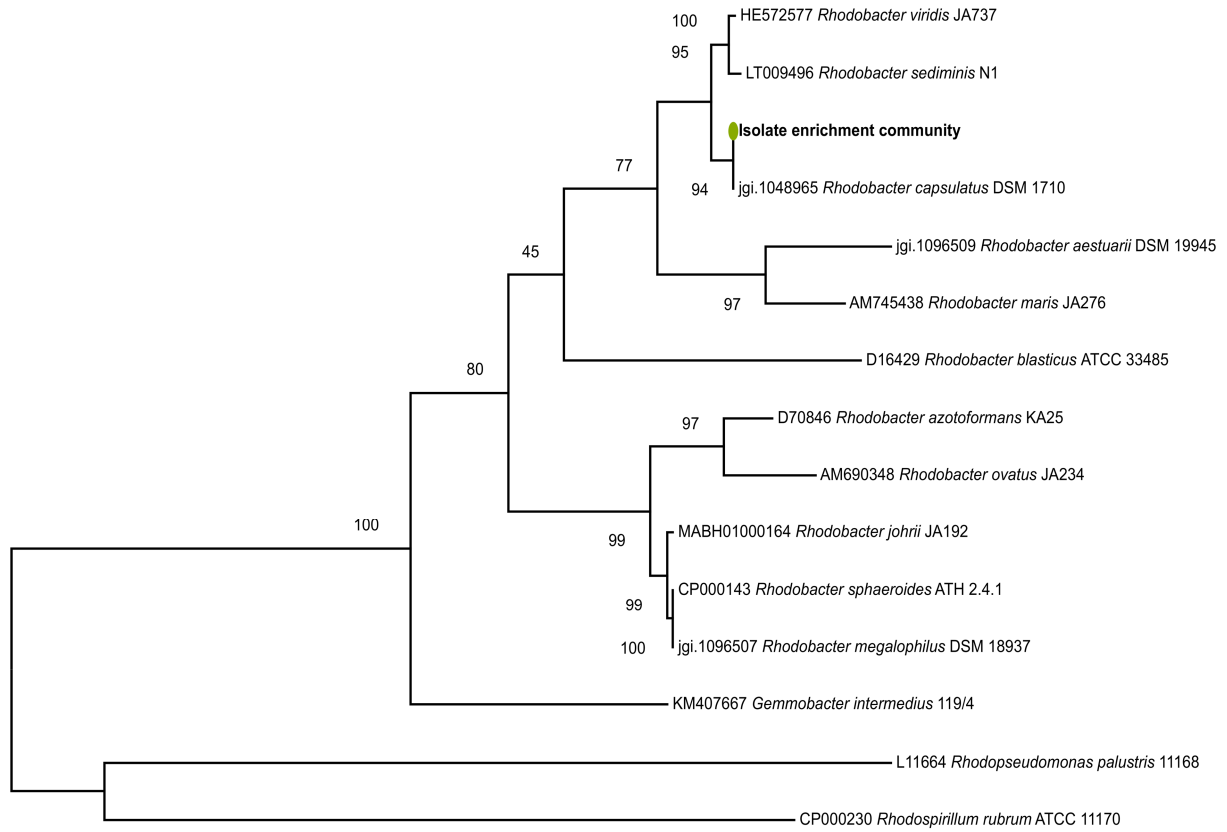
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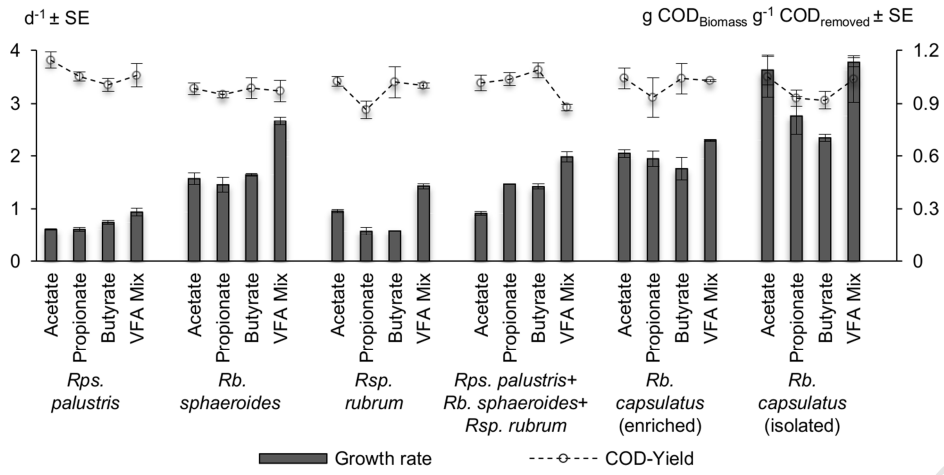
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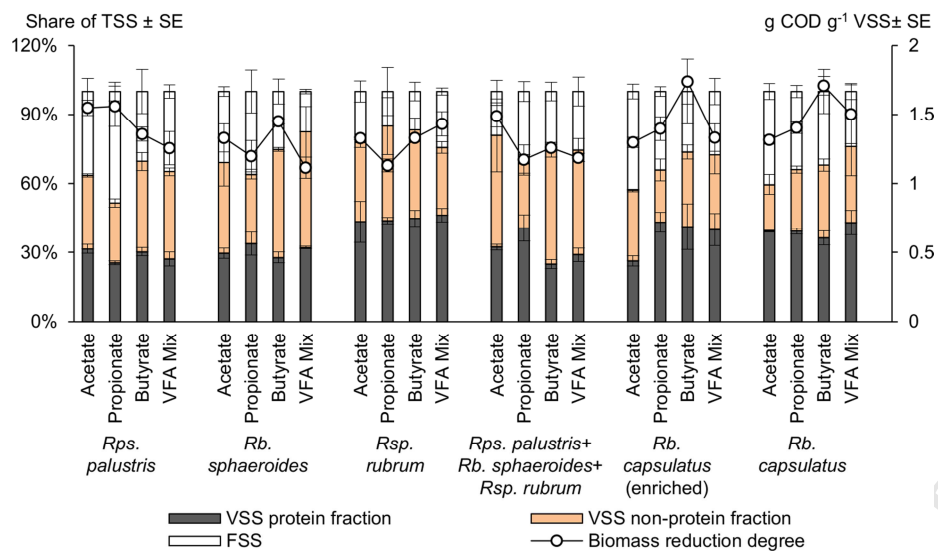


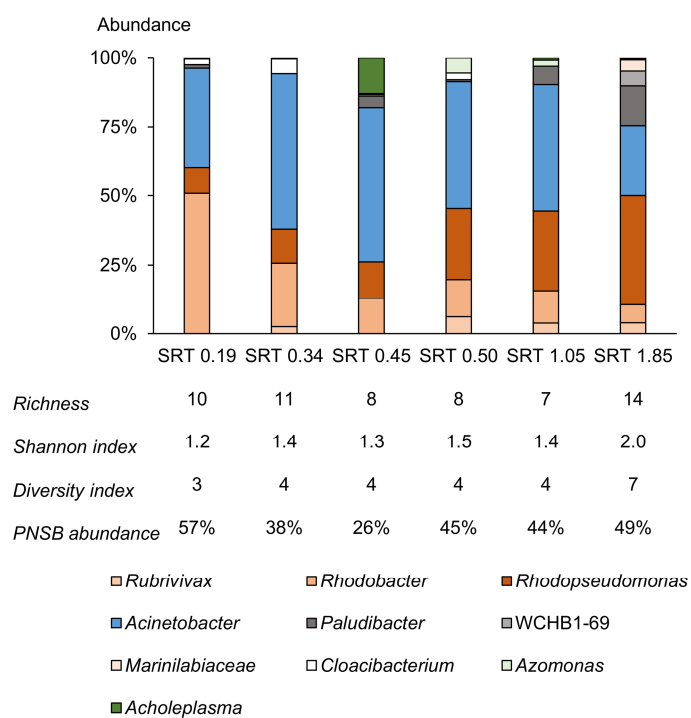


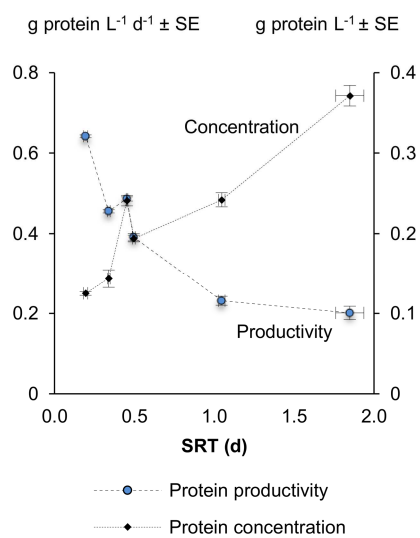
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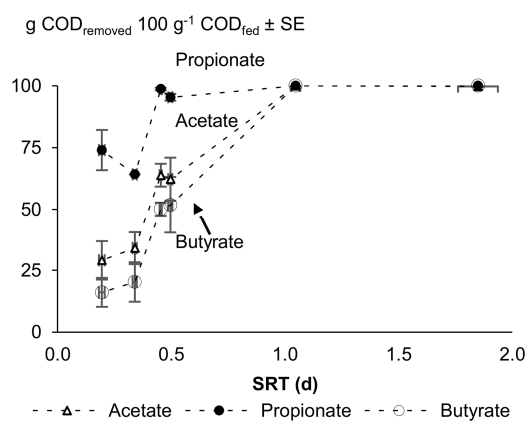
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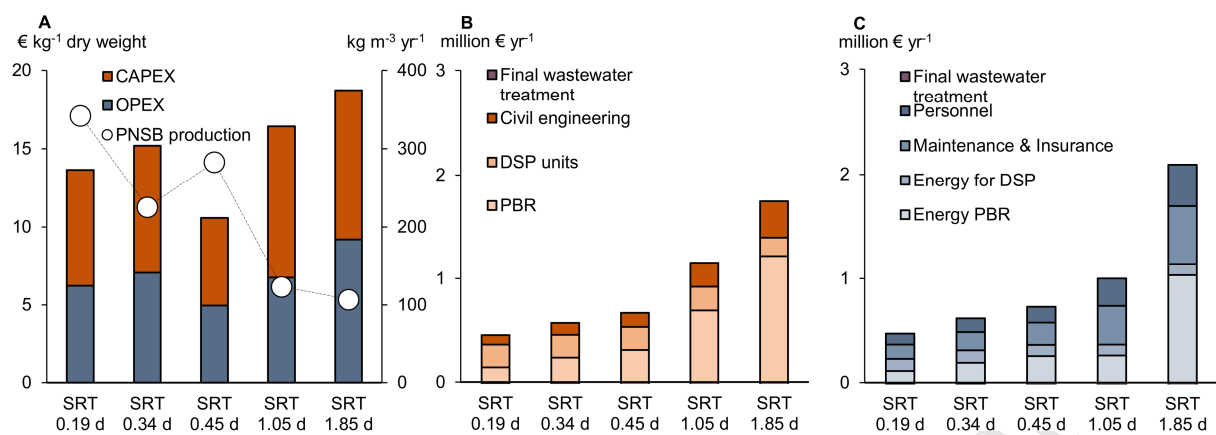












## Highlights

- Mixed rather than individual VFA boost growth of purple non-sulfur bacteria (PNSB)
- From six PNSB cultures, *Rhodobacter capsulatus* obtained the highest growth rates
- Highest volumetric productivity (1.7 g dry weight L<sup>-1</sup> d<sup>-1</sup>) reported for PNSB
- PNSB have a 1-to-1 or higher substrate-to-biomass conversion when grown on VFA
- PNSB most cost effective at sludge age of 0.5d based on model of brewery wastewater



**Declaration of interests**

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

The authors declare the following financial interests/personal relationships which may be considered as potential competing interests: