

This item is the archived peer-reviewed author-version of:

Preferential photoassimilation of volatile fatty acids by purple non-sulfur bacteria : experimental kinetics and dynamic modelling

Reference:

Segura Paloma Cabecas, De Meur Quentin, Alloul Abbas, Tanghe Audrey, Onderwater Rob, Vlaeminck Siegfried, Vande Wouwer Alain, Wattiez Ruddy, Dewasme Laurent, Leroy Baptiste.- Preferential photoassimilation of volatile fatty acids by purple non-sulfur bacteria : experimental kinetics and dynamic modelling

Biochemical engineering journal - ISSN 1369-703X - 186(2022), 108547

Full text (Publisher's DOI): <https://doi.org/10.1016/J.BEJ.2022.108547>

To cite this reference: <https://hdl.handle.net/10067/1927410151162165141>

1 **Preferential photoassimilation of volatile fatty acids by purple non-sulfur bacteria:**

2 **Experimental kinetics and dynamic modelling**

3 Paloma Cabecas Segura¹, Quentin De Meur¹, Abbas Alloul², Audrey Tanghe ³, Rob
4 Onderwater³, Siegfried E. Vlaeminck², Alain Vande Wouwer ⁴ , Ruddy Wattiez¹, Laurent
5 Dewasme⁴, Baptiste Leroy¹

6

7 ¹Laboratory of Proteomics and Microbiology, University of Mons, 7000 Mons, Belgium.

8 ²Research Group of Sustainable Energy, Air and Water Technology, Department of Bioscience
9 Engineering, University of Antwerp, Groenenborgerlaan 2020, Antwerpen, Belgium.

10 ³Materia Nova ASBL, Parc Initialis, Avenue Copernic 3, 7000 Mons, Belgium

11 ⁴Systems, Estimation, Control and Optimization Group (SECO), University of Mons, 7000
12 Mons, Belgium

13

14

15

16

17 **Corresponding author:**

18 E-mail : baptiste.leroy@umons.ac.be

19 Phone : +32-65-372271

20 Address: Laboratory of Proteomic and Microbiology, UMONS, Avenue du champs de Mars, 6
21 (Pentagone 3B), 7000, Mons, Belgium)

22 **Highlights :**

- 23 • *Rs. rubrum* sequentially assimilates different volatile fatty acids.
- 24 • The inhibition phenomenon probably occurs to avoid a substrate cycle.
- 25 • Macroscopic dynamic models were developed to describe multiple substrate uptake.
- 26 • Sequential assimilation of VFA is best described using an inhibition constant.

27

28

29

30

31

32

33

34

35

36

37

38

39

40

41

42 **Abstract :**

43 Purple non-sulfur bacteria (PNSB) are known for their metabolic versatility and thrive
44 as anoxygenic photoheterotrophs. In environmental engineering and resource recovery, cells
45 would grow on mixtures of volatile fatty acids (VFA) generated by anaerobic fermentation of
46 waste streams. In this study, we aim to better understand the behavior of *Rhodospirillum*
47 *rubrum*, a model PNSB species, grown using multiple VFA as carbon sources. We highlighted
48 that assimilation of individual VFA follows a sequential pattern. Based on observations in other
49 PNSB, this seems to be specific to isocitrate lyase-lacking organisms. We hypothesized that the
50 inhibition phenomenon could be due to the regulation of the metabolic fluxes in the substrate
51 cycle between acetoacetyl-CoA and crotonyl-CoA. Developed macroscopic dynamic models
52 showed a good predictive capability for substrate competition for every VFA mixture
53 containing acetate, propionate, and/or butyrate. These novel insights provide valuable input for
54 better design and operation of PNSB-based waste treatment solutions.

55

56

57

58

59

60

61

62

63

64 **Keywords :**

65 resource recovery, wastewater treatment, parameter estimation, *Rhodospirillum rubrum*

66

67

68

69

70

71

72

73

74

75

76

77

78

79

80

81

82

83

84 **Nomenclature:**

85 $\mu_{\max_{\text{ace}}}$, $\mu_{\max_{\text{prop}}}$, $\mu_{\max_{\text{val}}}$, $\mu_{\max_{\text{but}}}$, $\mu_{\max_{\text{VFA}}}$, μ_{\max} : maximal uptake rates (h^{-1})

86 K_{ace} , K_{prop} , K_{but} , K_{val} : half-saturation constants (g/g)

87 $K_{i_{\text{ace}}}$, $K_{i_{\text{prop}}}$, $K_{i_{\text{prop+ace}}}$: inhibition constants (g/l)

88 C_{ace} , C_{prop} , C_{but} , C_{val} , C_{hb} , C_{hv} , h_{bc} , h_{vc} : substrate to biomass yield coefficients (g/g)

89 [Ace], [Prop], [But], X : concentrations of acetate, propionate, butyrate and biomass

90

91

92

93

94

95

96

97

98

99

100

101

102

103

104

105 **1. Introduction**

106 Over the past few years, the interest in a more circular economy has been rising, suggesting
107 that waste should be treated as secondary feedstocks, being valorized and upcycled. Following
108 this principle, an important research and development effort was undertaken to develop
109 sustainable environmental biotechnologies to valorize organic in wastewater or solid waste in
110 higher added value products such as biofuel, bioplastics, or microbial proteins [1–3]. Purple
111 non-sulfur bacteria (PNSB) can offer solutions to phototrophically upcycle organic resources
112 from waste streams to several added-value compounds such as fertilizers, for feed,
113 polyhydroxyalkanoates (PHA) for bioplastics, pigments, coenzyme Q10 or H₂ [4–6].
114 Nevertheless, waste streams contain usually a mixture of carbon compounds such as
115 polysaccharides, fatty acids, DNA, or polyphenolic structures which make it challenging to
116 develop a universal, robust and reproducible process [7]. Conversion of such complex organic
117 matrices, for instance through fermentation, into simpler molecules such as volatile fatty acids
118 (VFAs), is a way to increase the process feasibility and transferability. VFAs can then be
119 subsequently easily assimilated by organisms such as PNSB [8]. The most found VFAs in
120 fermented waste streams are acetate, butyrate, isobutyrate, propionate, valerate, and isovalerate.
121 Even though, the VFA speciation depends on many factors such as substrate, temperature, pH,
122 or hydraulic and solid retention times, the most abundant produced VFAs being usually acetate,
123 butyrate, propionate, and valerate [9–14].

124 For the synthesis of PHA in PNSB, a precise understanding of the assimilation processes
125 of VFAs is required as the carbon source used to support the bacterial growth directly impacts
126 the composition of the polymer [15,16]. In this frame, some mathematical models were
127 developed to describe the assimilation in microbial cultures of some volatile fatty acid mixtures
128 [17–21]. Substrate uptake rate varies significantly depending on the composition of the mix
129 and, in addition, the biomass may exhibit some substrate consumption preferences. It was, for

130 instance, observed that *Plasticicumulans acidivorans* preferentially consume butyrate over
131 acetate [19]. Therefore, the development of such a model requires the observation and
132 understanding of the microbial strain metabolic behavior in presence of different VFA mixtures,
133 as well as an insight into the metabolic pathways involved in their assimilation. C₂ compounds
134 such as acetate are usually assimilated through the glyoxylate shunt [22]. However, some PNSB
135 are lacking the isocitrate lyase (ICL⁻ organisms) which is the key enzyme of this pathway. In
136 this case, the net assimilation of C₂ compounds requires the use of alternative anaplerotic
137 pathways. The ethylmalonyl-CoA (EMC) pathway has been proposed as a key alternative
138 pathway to fulfill this role in several ICL⁻ PNSB including *Rs. rubrum* [23–25](Figure S2). In
139 *Rs. rubrum*, the propionyl-CoA resulting from the EMC is first carboxylated in methylmalonyl-
140 CoA before being converted to succinate [23–25]. When propionate is used as a carbon source,
141 its assimilation equivalently occurs via methylmalonyl-CoA [26](Figure S2). The assimilation
142 of butyrate (C₄) under photoheterotrophic conditions, remains poorly described even though
143 this molecule is known to support photoheterotrophic growth in most of the PNSB. A recent
144 study carried out in our lab on *Rs. rubrum* has underlined the use of the EMC pathway for
145 butyrate assimilation but also of the newly proposed methylbutanoyl-CoA
146 (MBC) pathway [27](Figure S2). Most studies performed with PNSB to better characterize the
147 photoheterotrophic assimilation pathways of VFAs were achieved on cell cultures supplied with
148 a unique carbon source. From that perspective, the obtained data can only result in a metabolic
149 adaptation to the defined carbon source which can significantly affect metabolism and genome
150 as we recently demonstrated in the context of the long-term adaptation of *Rs. rubrum* to the
151 assimilation of acetate [28]. However, when referring to an environmental context but also an
152 industrial application where waste streams are used as substrates, the probability that PNSB
153 grows with a unique carbon source is fairly low. Some studies already highlighted a synergic

154 effect on the uptake rate of VFAs or a delay in this uptake when VFAs were simultaneously
155 present in the medium using PNSB [5,6].

156 This study aims to experimentally unravel the simultaneous uptakes pattern of individual
157 VFAs in a mixture using *Rs. rubrum* as a model organism and to propose a mathematical model
158 able to describe such phenomenon. In addition, the PNSB *Rhodobacter capsulatus*, and
159 *Rhodobacter sphaeroides*, often retrieved in open communities for wastewater treatment were
160 also investigated regarding their assimilatory behavior for blends of VFAs.

161

162 2. **Materials and methods**

163 2.1. *Rs. rubrum* tests: Strain and culture conditions

164 *Rhodospirillum rubrum* S1H (ATCC15903) is grown in a medium used for
165 photoheterotrophic culture conditions, that is based on the basal salt medium of Segers &
166 Verstraete described by Suhaimi *et al.* [29,30], supplemented with NH_4Cl (35 mM) as the
167 nitrogen source, NaHCO_3 (3 mM or 50mM), and biotin (0.06 mM). Six different mixtures of
168 VFAs were used as carbon sources: [1] acetate (35 mM), butyrate (10 mM) and propionate (5
169 mM) in the 7:4:1.5 carbon ratio, [2] acetate (31 mM) and propionate (20.6 mM), [3] acetate (31
170 mM) and butyrate (15.5 mM), [4] propionate (20.6 mM) and butyrate (15.5 mM), and [5]
171 acetate (12mM) and butyrate (15.5mM), [6] propionate (15mM) and butyrate(15,5mM).
172 Mixtures [2], [3], [4] have a carbon ratio between VFAs of 1:1. Succinate (31 mM) was used
173 as carbon source in control conditions, at the same net carbon concentration (124 mM). The pH
174 was adjusted to 6.9. *Rs. rubrum* was grown under anaerobic phototrophic conditions in 50 ml
175 sealed serum flasks under $50 \mu\text{mole.m}^{-2}.\text{sec}^{-1}$ of light intensity supplied by halogen lamps
176 (Sencys; 10 W; 100 lumens; 2,650 K), at 30°C with a rotary shaking at 200 rpm. Each culture
177 condition was achieved with five biological replicates, except for the cultivation experiments.
178 Nitrogen gas was used to purge oxygen from the upper gas phase and the flasks were

179 hermetically sealed. The cultures were inoculated at a starting OD_{680nm} between 0.4 and 0.5 and
180 the growth was monitored following the turbidity at OD_{680nm}. A correlation curve between OD
181 and biomass dry weight was used to convert OD_{680nm} in biomass content.

182 2.2. *Rs. rubrum* tests: Monitoring of VFAs consumption

183 The monitoring of *Rs. rubrum* VFA consumption was performed on culture supernatants
184 obtained by centrifugation at 16000 g for 10 min at 4°C and stored at -20°C before analysis.
185 Aliquots (100 µl) of culture supernatant were analyzed by HPLC-refractometry (Waters 2695
186 separation Module; Waters 2414 Refractive Index Detector). The separation was done in
187 isocratic mode using a Shodex Sugar SH1011 column (300 mm x 8 mm) with 5 mM H₂SO₄ as
188 the mobile phase. The amounts of acetate, propionate, butyrate, and succinate were determined
189 by integrating their specific peaks and comparing area under the curve with external standards.

190 2.3. *Rb. sphaeroides* and *Rb. capsulatus* tests: Strains, culture conditions, and 191 analytical methods

192 *Rhodobacter sphaeroides* (LMG 2827), *Rhodobacter capsulatus*, and an enriched
193 community which was dominated by 80% by *Rhodobacter capsulatus* were used as bacterial
194 strains to extend the study to other PNSB. Batch experiments involving *Rhodobacter*
195 *sphaeroides* and *Rhodobacter capsulatus* were carried out in 500 mL Erlenmeyer with a
196 working volume of 500 mL in triplicates and closed with a screw cap containing a silicone
197 rubber polytetrafluoroethylene protected septum (Duran, Germany). The headspace was
198 flushed for 1 minute with nitrogen gas. Stirring was performed with a multipoint stirrer at 300
199 rpm (Thermo Scientific, USA). The batch experiment for *Rb. sphaeroides* was performed in a
200 climate chamber (Snijders Scientific) at 28 °C equipped with 36 W fluorescent lamps (Sylvania,
201 Germany) at a light intensity of 30 W m⁻². The batch test for the enriched *Rb. capsulatus* was
202 tested at 29 ± 1 °C, illuminated with 500 W halogen lamps (Philips, Netherlands) at a light
203 intensity of 30 W m⁻² and covered with infrared transmission filters (Bay Plastics, UK).

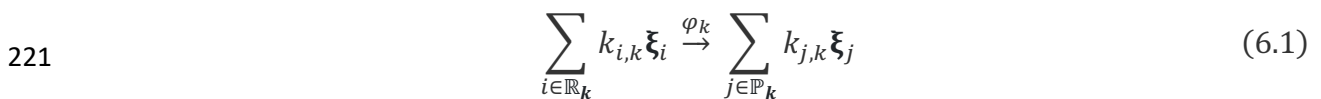
204 Samples of 10 ml were taken for further analyses and to monitor the growth
 205 spectrophotometrically at 660 nm. VFAs were analyzed by High-Performance Liquid
 206 Chromatography (Agilent technologies 1200) coupled with a diode array detector (210 nm), a
 207 Bio-Rad Aminex[®] column (300 mm 7,8mm) with Bio-Rad Micro-Guard Cation H Refill
 208 Cartridges and a column temperature of 40°C. The samples were injected into 50 mM H₂SO₄
 209 (Honeywell Fluka[™]) in deionized water (Arium[®] 611) with an injection volume of 20 μL and
 210 a sample flow rate of 0.6 ml min⁻¹.

211 2.4. Statistical analysis on biological experiments

212 All graphs were plotted using the software GraphPad Prism (Version 6.01). The results
 213 were reported as the means and standard deviations computed by GraphPad Prism 6.01. To
 214 evaluate the significance of changes, t-tests or ANOVA were performed with a threshold set at
 215 0.05. Significance difference was marked using different lower cases. These analyses
 216 were performed using MATLAB software (version R2018a 9.4.0.813654).

217 2.5. Mathematical model identification procedure

218 The proposed modeling approach is based on a priori knowledge of the metabolic
 219 network and data obtained from the in-lab experiments described in section 3.2. The considered
 220 bioprocess can be described by the following macroscopic reaction scheme:



222 with $k \in [1, M]$ where M is the number of reactions, \mathbb{R}_k and \mathbb{P}_k denote respectively the set of
 223 reactants and products in the reaction k . The parameters $k_{i,k}$, and $k_{j,k}$ are pseudo stoichiometric
 224 coefficients while φ_k is the corresponding reaction rate. Applying mass balance to (1), the
 225 following ordinary differential equation (ODE) system is obtained, describing the variations of
 226 each species concentration with time:

227
$$\frac{d\xi(t)}{dt} = K \varphi(\xi, \theta_\varphi, t) \quad (6.2)$$

228 where K is the pseudo-stoichiometric matrix and θ_φ the kinetic parameters. Parameter
 229 estimation is conducted by minimizing a least-squares criterion measuring the distance between
 230 model-simulated data ξ_m and experimental measurements ξ_{exp} as in:

231
$$J(\theta) = (\xi_m(\theta) - \xi_{exp(\theta)})^T Q^{-1} (\xi_m(\theta) - \xi_{exp(\theta)}) \quad (6.3)$$

232 where θ is the parameter vector (gathering stoichiometric and kinetic parameters).

233 The initial state ξ_0 is a vector of length N times n_{exp} , n_{exp} being the number of used
 234 experiments and N the size of vector ξ , that is, the number of studied macroscopic species (in
 235 the current case, biomass and VFAs). $\xi_{0,0}$ is guessed using the experimental measurements at
 236 time $t = 0$. Q is the measurement error covariance matrix designed as a diagonal matrix
 237 (assuming independent distributions of the measurement errors) containing the squares of the
 238 maximum concentration levels of each species, a good practice rule to normalize the distances
 239 calculated in (3) when the error distributions are not assumed to be known a priori. Parameter
 240 identification is performed with the MATLAB optimizer “fmincon” which uses an interior point
 241 method and tolerates upper and lower bounds on the assumed unknown parameters, allowing
 242 to reduce the search space and accelerate the optimization. The latter option can indeed be
 243 achieved since, for most of the kinetic and stoichiometric parameters, the general order of
 244 magnitude can be easily guessed from previous studies, or some trivial data analyzes.

245 The data sets obtained from three biological replicates are used for each substrate
 246 operating condition (VFAs mixture) for parameter estimation and model direct validation.
 247 Supplementary data coming from a separate experiment are used for model cross-validation.
 248 Local parameter identifiability and sensitivity analysis are addressed based on the Fisher
 249 Information Matrix (FIM), which can be computed as follows :

250
$$\text{FIM} = \sum_{tk=1}^{n_{meas}} \left(\xi_{\theta,i}(t_k, \hat{\theta}) \right)^T Q^{-1} \xi_{\theta,i}(t_k, \hat{\theta}) \quad (6.4)$$

251 where t_k is the sampling time and n_{meas} is the number of samples. An estimate of the parameter
 252 estimation error covariance matrix can be obtained from the Cramer-Rao bound as follows:

253
$$\hat{P} > \sigma^2 \text{FIM}^{-1} \quad (6.5)$$

254 with σ^2 being the a posteriori estimate of the measurement error variance calculated from the
 255 residual cost function at the optimum:

256
$$\sigma^2 = \frac{J^*}{n_{exp} * N_{meas} - n_{\theta}} \quad (6.6)$$

257 where N_{meas} is the total number of measurements (n_{meas} times N) and n_{θ} is the number of
 258 estimated parameters. Model fitting is assessed by the root mean square error (RMSE).

259 **3. Results and discussion**

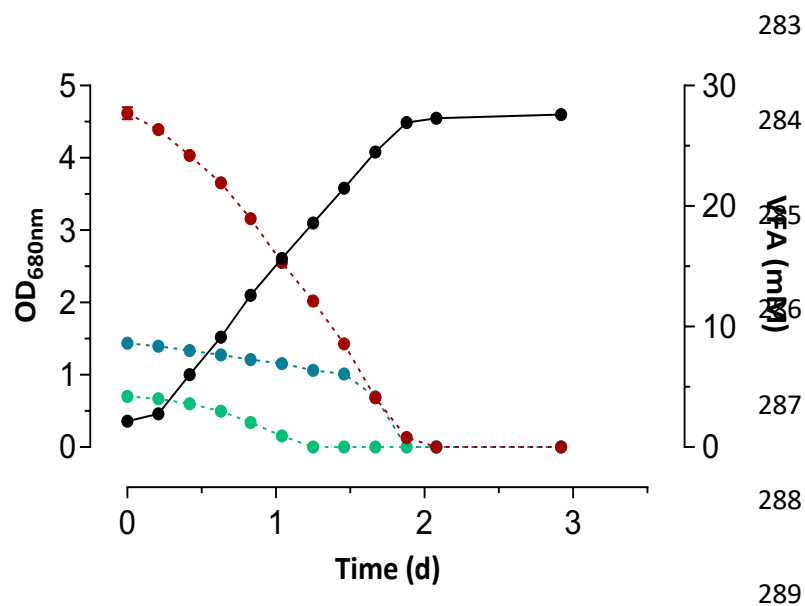
260 In this first experimental study, the composition of the mixture of VFAs and the
 261 proportion of each component was based on the composition of the effluents of a thermophilic
 262 anaerobic fermentation pilot plan operated in the context of the micro-ecological life support
 263 system alternative MELiSSA program from ESA [31]. *Rs. rubrum* was first grown in a culture
 264 medium supplemented with a mixture of VFAs containing acetate, butyrate, and propionate in
 265 a 7:4:1.5 carbon ratio and 3 mM of bicarbonate. The growth rate in the control (succinate) and
 266 the VFAs mixture conditions did not show any significant difference but for the same nominal
 267 124 mM of carbon present in the medium, the biomass level reached at the end of the culture
 268 was higher with the mixture of VFAs than in the control condition (Figure 1).

269

270

271 3.1. Delayed butyrate uptake in presence of acetate and propionate

272 Unexpectedly, while the use of acetate as a single carbon source under
273 photoheterotrophic conditions with low bicarbonate levels is characterized by a long lag phase
274 (100h)[28](Figure S1), we observed that both the growth and the acetate assimilation started
275 without delay with a VFA mixture largely dominated by acetate. The monitoring of the VFA
276 concentrations in the culture medium during the growth showed that acetate assimilation rate
277 was high from the beginning of the experiment. The assimilation rate of propionate was low
278 during the first days and then quickly increased. For butyrate, on the other hand, the assimilation
279 rate was low until day 1.5 and increased only after the complete assimilation of propionate and
280 a significant decrease of acetate concentration (Figure 1). The VFA consumptions followed a
281 sequential pattern of assimilation, butyrate being significantly consumed essentially at the end
282 of the experiment.



290 **Figure 1:** Monitoring of the growth(●) and VFA consumptions in cultures of *Rs. rubrum*
291 cultivated with a mix of acetate(●), propionate(●), and butyrate(●) as carbon sources (7:4:1.5
292 carbon ration).

293

294 3.2. Butyrate assimilation is delayed by the presence of acetate and/or propionate

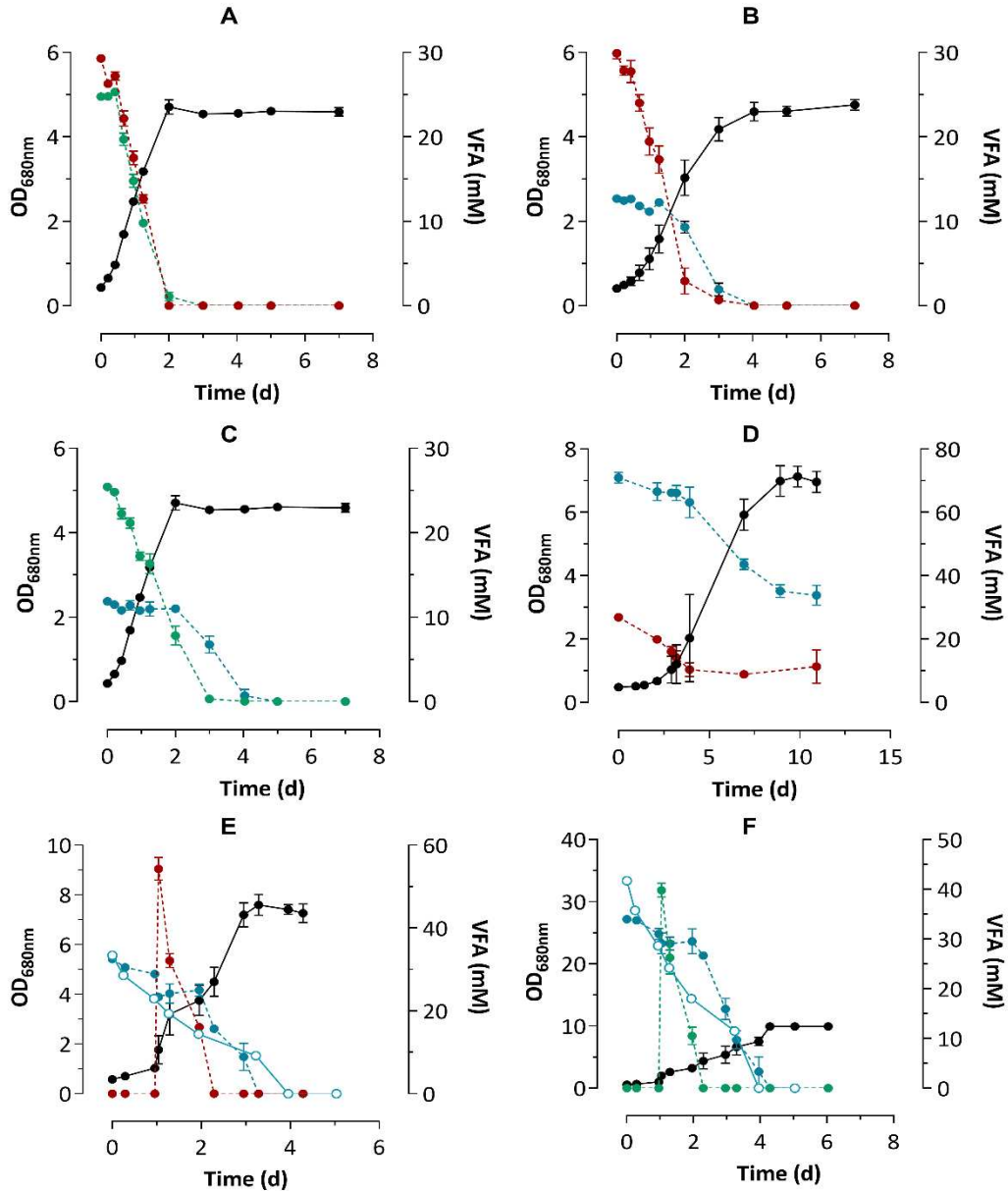
295 To define if acetate or propionate inhibits assimilation of butyrate, *Rs. rubrum* was
296 grown in a culture medium supplemented with 3 different binary mixtures each containing only
297 two of the three VFAs in the same equivalent net carbon concentrations (for a total of 124 mM
298 net of carbon) and 3mM of bicarbonate (Figure 2). Culture of *Rs. rubrum* using acetate,
299 propionate, butyrate as sole carbon source were used as controlled (Figure S1). To confirm the
300 absence of deleterious interaction observed here before, the assimilation profile of a mixture of
301 acetate and propionate was first explored. Both VFAs were simultaneously consumed during
302 the growth phase and propionate could be completely assimilated with the 3mM of bicarbonate
303 supplemented to the medium (Figure 2A). Considering the assimilation pathways of propionate
304 [26] and acetate [24] in *Rs. rubrum*, this observation appeared to be quite logical since the
305 propionyl-CoA is a metabolic intermediate of the ethylmalonyl-CoA pathway used to
306 photoassimilate acetate.

307 Regarding the assimilation of a mixture of acetate and butyrate, no lag-phase and no
308 dependency on bicarbonate supplementation were observed in the growth (Figure 2B). Similar
309 observations could be made on the culture growing with propionate and butyrate as carbon
310 sources (Figure 2. C). In both cases, butyrate consumption was delayed until the concentration
311 of the other VFA has dropped down. To confirm this observation, the assimilation rates of each
312 VFA at different steps of the growth period were calculated (Figure 3). It can be observed that
313 when VFAs are alone or in a mixture in which no sequential assimilation is observed, as, in the
314 case of a mixture of acetate and propionate, no change in the assimilation rate is observed along
315 the biomass growth period. In contrast, a significant difference in the VFA uptake is observed
316 in a mix exhibiting their sequential assimilation (Figure 3). In the case of the mixture of acetate
317 and butyrate, it can be noticed that while at the beginning of the culture acetate has a
318 significantly higher uptake rate than butyrate, during the second part of the cultivation the

319 opposite phenomenon is observed (Figure 3). Similar observations can be made for the mixture
320 of propionate and butyrate or the mixture of acetate, propionate, and butyrate. This tends to
321 confirm the sequential assimilation of VFAs when butyrate is assimilated in presence of acetate
322 or propionate. A fifth mixture was used containing acetate (25 mM) and excess butyrate (70
323 mM). In the medium containing this mixture, the NH_4Cl concentration was adjusted to 85 mM
324 according to the net carbon concentration (330 mM) to respect the C/N ratio.

325

326



327

328 **Figure 2:** Monitoring of the growth and VFA consumptions in cultures of *Rs. rubrum* S1H
 329 cultivated in binary mixtures of VFAs: (A) acetate and propionate (n=3), (B) acetate and
 330 butyrate (n=3), (C) propionate and butyrate (n=3) (D) acetate with an excess of butyrate (n=3),
 331 (E) butyrate and spike of acetate (n=3), (F) butyrate and spike of propionate (n=3). The
 332 production of biomass (●) and the concentration of acetate (●), butyrate (●), and propionate (●)
 333 are presented. Control with consumption of butyrate by *Rs. rubrum* in presence of 50 mM of
 334 bicarbonate (○).

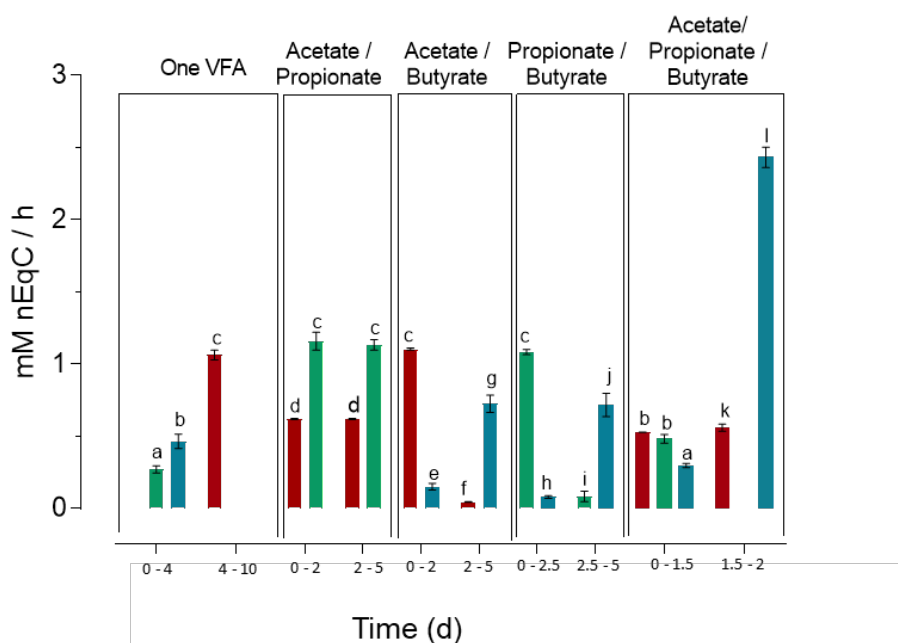
335 To determine if the observed phenomenon is ruled by the concentration levels of acetate
336 and butyrate, *Rs. rubrum* was cultivated with a medium supplemented with acetate and butyrate
337 in a carbon ratio of 1:5, respectively. Despite the excess of butyrate in the culture medium, the
338 preferential assimilation of acetate over butyrate is still appearing (Figure 2D), suggesting that
339 this mechanism is not related to the higher concentration of acetate. It is worth noting that
340 acetate assimilation rate was lower in this condition which is unexplained today and should
341 require further investigation. As expected from the higher net carbon concentration, the culture
342 reached very high biomass levels. Nevertheless, the growth stopped before the complete
343 assimilations of butyrate and acetate. Considering that the carbon/nitrogen balance has been
344 conserved in medium preparation, the depletion of other compounds such as phosphorus may
345 explain this phenomenon. To test whether or not the latter could be linked to a substrate
346 preference, acetate and propionate were supplemented in the medium during the exponential
347 phase of culture initially growing on butyrate as the main carbon source (Figure 2E and 2F).
348 Culture growing with butyrate as the sole carbon source required excess of bicarbonate to grow
349 therefore those experiments were carried with 50mM of bicarbonate in the culture medium. As
350 shown in Figures 2.E and 2.F, butyrate assimilation stopped when acetate or propionate spikes
351 occurred, indicating a possible substrate preference or inhibition at an enzymatic level. This
352 inhibition seemed, however, to be limited to high acetate concentrations since butyrate
353 assimilation started before the complete consumption of acetate. The inhibitory effect of acetate
354 on butyrate assimilation could, in some way, be expected. For their assimilation, acetate and
355 butyrate share some metabolic intermediates between acetyl-CoA and crotonyl-CoA as well as
356 the ethylmalonyl-CoA pathway.

357 When acetate is the carbon source, acetyl-CoA is converted into crotonyl-CoA, while the
358 latter serves as a substrate to produce acetyl-CoA when *Rs. rubrum* grows on butyrate(Figure
359 S2). The resulting metabolic mismatch is mainly related to the differential expression of two

360 enzymes. The first one, Rru_A3079, specific to the photoassimilation of butyrate, catalyzes the
361 conversion of the (S)-3-hydroxybutyryl-CoA into acetoacetyl-CoA. This enzyme has been
362 observed to be up-regulated in butyrate condition (fold change: 1.9, P -value: $5e-8$) [27], and
363 downregulated in acetate condition (fold change: 0.3, P -value: $4e-4$)[24]. The second enzyme,
364 Rru_A0273, is involved in the acetate assimilation and is responsible for the conversion of
365 acetoacetyl-CoA into (R)-3-hydroxybutyryl-[28]. It is strongly downregulated in butyrate
366 conditions (fold change: 0.3, P -value: $4e-7$)[27]. Moreover, the data obtained from a mutant
367 fitness assay performed on a transposon mutant library of *Rs. rubrum* S1H showed that the gene
368 Rru_A2964, coding for the enzyme catalyzing the interconversion of the crotonyl-CoA and the
369 (R)-3-hydroxybutyryl-CoA, is essential for the strain survival under acetate condition[28],
370 whereas the mutations knocking this gene out appeared to be beneficial in butyrate condition.
371 Depending on the available carbon source, the bacterium seems therefore to differentially
372 regulate the expression of two routes, avoiding a cycle between acetoacetyl-CoA and crotonyl-
373 CoA. Interestingly, Rru_A1835, the enzyme involved in the conversion of the butyryl-CoA into
374 crotonyl-CoA (and therefore directly responsible for the entry of the butyrate in the central
375 metabolism), is strongly downregulated by acetate (fold change: 0.2, P -value: $2e-4$)[24],
376 whatever the initial concentration of butyrate (Figure S2).

377 The inhibitory effect of acetate is therefore maybe not only the result of a metabolic
378 mismatch but could also result from direct regulation of the expression of enzymes specifically
379 related to the butyrate photoassimilation. The inhibitory effect of propionate on the assimilation
380 of butyrate is more difficult to interpret. As propionyl-CoA is an intermediate of both the EMC
381 and MBC pathways, being probably both used for the butyrate assimilation [27], it could be
382 hypothesized that its presence in high quantity could induce negative feedback on those
383 upstream metabolic pathways, preventing the butyrate assimilation (Figure S2). Alternatively,
384 one could also hypothesize that the downregulation of the expression of genes required for

385 butyrate assimilation is due to metabolites that are both produced during acetate and propionate
 386 assimilation.

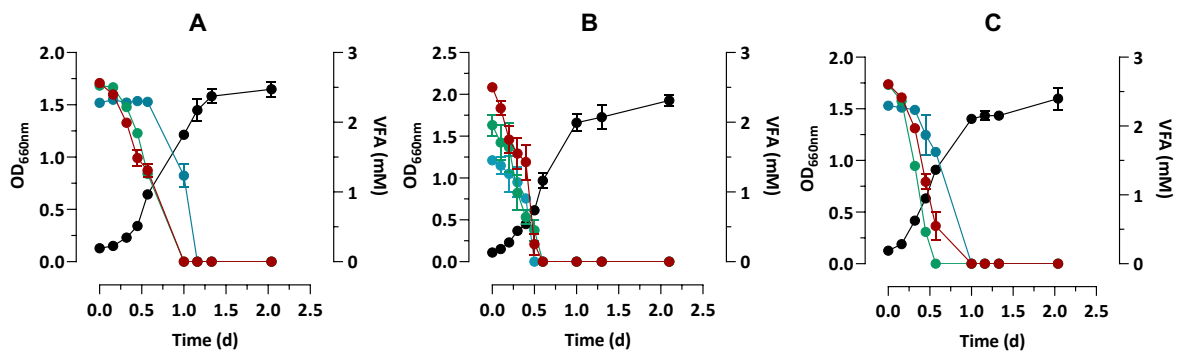


387
 388 **Figure 3:** Substrate consumption rate during the different phases of culture of *Rs. rubrum* using
 389 acetate (■), propionate (■), butyrate (■), or a mixture of those VFAs as a carbon source. Lower
 390 case marked significant difference (p-value<0.05; n=3)

391 3.3. Sequential VFA uptake is restricted to ICL- organisms

392
 393 To compare VFA assimilation patterns in *Rs. rubrum* with other strains of PNSB,
 394 *Rhodobacter sphaeroides*, *Rhodobacter capsulatus*, and an enriched non-axenic culture holding
 395 *Rhodobacter capsulatus* as the majoritarian strain was grown on a mixture of volatile fatty acids
 396 containing acetate, propionate, and butyrate. Monitoring of VFA consumptions by *Rhodobacter*
 397 *sphaeroides* highlighted an assimilation pattern similar to what was observed in *Rs. rubrum*
 398 (Figure 4A) with butyrate being assimilated at high rate only after concentration of acetate and
 399 propionate decreased. In *Rh. sphaeroides*, there was a clear delay in butyrate assimilation which
 400 was not consumed before day 1. Conversely, VFAs were almost simultaneously assimilated
 401 (Figure 4B) by *Rb. capsulatus* and no phase of “low-rate” assimilation of butyrate could be

402 observed. For the enriched non-axenic culture holding *Rhodobacter capsulatus* as majoritarian
 403 strain (Figure 4C), a very short phase of low-rate assimilation of butyrate was observed, but the
 404 assimilation rate of butyrate quickly increased not waiting for full consumption of acetate and
 405 butyrate as observed for *Rh. sphaeroides*. The sequential assimilation of VFAs, with high rate
 406 assimilation of butyrate being observed only when acetate/propionate concentration dropped,
 407 could therefore be a characteristic of ICL⁻ organisms such as *Rs. rubrum* and *Rb. sphaeroides*
 408 that are known to depend on the EMC pathway for the assimilation of acetate. The possible
 409 metabolic mismatch described before could also explain why only ICL⁻ organisms which rely
 410 on the EMC pathway for the assimilation of acetate are unable to concomitantly assimilate
 411 acetate and butyrate. On the other hand, microorganisms using the glyoxylate shunt for acetate
 412 assimilation can simultaneously assimilate butyrate, as observed for *Rh. capsulatus* in the
 413 present study.



414
 415
 416 **Figure 4:** Monitoring of the growth (●) and VFA consumption in batch cultures of multiple
 417 purple bacteria cultivated in an Erlenmeyer with acetate(●), butyrate (●), and propionate(●) as
 418 carbon source. Cultures of (A) *Rb. sphaeroides*, (B) *Rb. capsulatus* and (C) an enriched culture
 419 of *Rb. Capsulatus*.

420

421

422 3.4. Development of a dynamic model describing *Rs. rubrum* substrate uptake
 423 preferences from a mixture of VFAs

424 The aim is to develop the simplest mathematical mode that described a pattern of
 425 assimilation of VFA by *Rs. rubrum* based on the experimental data presented and proteomic
 426 analysis carried in mixture production and presented in Cabecas-Segura *et al.* 2021[32]. To do
 427 so, the first step is to formulate a hypothesis on the kinetic structure. In a second step, it must
 428 be ensured that all the phenomena assumed to be present in the data sets are significantly
 429 identifiable in practice, proceeding to a practical parametric identifiability analysis. If the latter
 430 does not provide satisfactory results, the factors containing the non-identifiable parameters
 431 should be removed, or new data better highlighting the corresponding phenomenon should be
 432 considered. In this case, if the confidence interval of the parameter is to wide, the model will
 433 be reduced.

434 **Table 1:** Selected kinetic structure to describe the assimilation of multiple VFAs by *Rs. rubrum*

Case	Carbon source 1	Carbon source 2
Mix acetate propionate Case A.1	$\varphi_1 = X \cdot \mu_{\max_{Ace}} \cdot \frac{[Ace]}{K_{Ace} + [Ace]}$	$\varphi_2 = X \cdot \mu_{\max_{Prop}} \cdot \frac{[Prop]}{K_{Prop} + [Prop]}$
Mix acetate butyrate Case B.1	$\varphi_1 = X \cdot \mu_{\max_{Ace}} \cdot \frac{[Ace]}{K_{Ace} + [Ace]}$	$\varphi_2 = X \cdot \mu_{\max_{But}} \cdot \frac{[But]}{K_{But} + [But]} \cdot \frac{K_{i_{Ace}}}{K_{i_{Ace}} + [Ace]}$
Mix propionate butyrate Case C.1	$\varphi_2 = X \cdot \mu_{\max_{Prop}} \cdot \frac{[Prop]}{K_{Prop} + [Prop]}$	$\varphi_2 = X \cdot \mu_{\max_{But}} \cdot \frac{[But]}{K_{But} + [But]} \cdot \frac{K_{i_{Prop}}}{K_{i_{Prop}} + [Prop]}$
Mix acetate propionate butyrate Case D.1	$\varphi_1 = X \cdot \mu_{\max_{Ace}} \cdot \frac{[Ace]}{K_{Ace} + [Ace]}$	$\varphi_2 = X \cdot \mu_{\max_{Prop}} \cdot \frac{[Prop]}{K_{Prop} + [Prop]}$
	Carbon source 3	
	$\varphi_3 = X \cdot \mu_{\max_{But}} \cdot \frac{[But]}{K_{But} + [But]} \cdot \frac{K_{i_{Ace}}}{K_{i_{Ace}} + [Ace]} \cdot \frac{K_{i_{Prop}}}{K_{i_{Prop}} + [Prop]}$	

435

436 In the case of the assimilation of multiple VFAs by *Rs. rubrum*, two different scenarios
437 that are the concomitant and the sequential assimilation of VFAs must be considered. As such
438 different kinetic structures will be tested to describe those phenomena the corresponding mass
439 balance are presented in supplementary data in table S1. The precision of the several proposed
440 models will be assessed based on the residual of the cost function (J) as defined in (3). To
441 describe the assimilation of VFAs that do not exhibit any preferential uptake effect, such as
442 acetate and propionate, a first model is proposed, considering their assimilations via distinct
443 metabolic pathways (Table 1). As reported in table 2, the resulting fitting is satisfactory when
444 acetate and propionate are used as substrates (J=0.25). However, it is not the case when acetate
445 and butyrate are used as carbon sources (J=6.49). Various kinetic models were thus tested to
446 describe the sequential assimilation of VFAs, namely a discontinuous switch from one VFA
447 consumption to the other (Table S.2 β) or the introduction of an inhibition factor (Table S.2 γ).
448 The best-predicting model considers the inhibition of butyrate assimilation by acetate as
449 indicated by the optimization residual reported in Table S.3. This kinetic structure could be
450 extended to the scenario using propionate and butyrate as carbon sources (J=0.23) and to three
451 VFA mixtures with a double inhibition phenomenon (J=0.42). This tends to corroborate the
452 hypothesis formulated earlier stating that the sequential assimilation is due to an inhibition
453 phenomenon at the enzymatic level rather than a switch from a carbon source to another. The
454 addition of an inhibition factor to the kinetic structure of VFA uptake has notably already been
455 proposed to explain comparable phenomena occurring at the enzymatic level [21].

456

457

458

459

460

461 **Table 2:** Parameter estimations and their confidence intervals for several models describing the
 462 assimilations of VFA mixtures.

Case	Mix acetate propionate (A.1)		Mix acetate butyrate (B.1)		Mix propionate butyrate (C.1)		Mix acetate propionate butyrate (D.1)	
	Value	CI (%)	Value	CI (%)	Value	CI (%)	Value	CI (%)
$\mu_{\max_{\text{Ace}}}$	0.08	108	0.13	114	-	-	2.11	6
$\mu_{\max_{\text{Prop}}}$	0.04	238	-	-	0.02	730	-	-
$\mu_{\max_{\text{But}}}$	-	-	0.03	488	0,02	884	0.04	371
K_{Ace}	0.99	9	1.24	11	-	-	0.13	98
K_{Prop}	0.32	26	-	-	0,1	167	17.96	10
K_{But}	-	-	0.04	339	0,2	80	0.71	18
$K_{i_{\text{Ace}}}$	-	-	0.73	20	-	-	2.38	5
$K_{i_{\text{Prop}}}$	-	-	-	-	0.22	74	0.02	653
C_{Ace}	0.85	10	0.82	18	-	-	0.51	25
C_{Prop}	2.02	4	-	-	1.22	13	0.45	28
C_{But}	-	-	0.37	40	3.44	5	1.10	11
J	0.25		0.42		0.23		0.42	

463

464 The second step in the development of the mathematical model is to ensure the
 465 parametric identifiability which will be assessed via the calculation of the 95% confidence
 466 intervals inferred from the Fisher Information Matrix (FIM) in (5). The confidence intervals
 467 related to the first proposed models are not acceptable regarding the VFA maximum uptake
 468 rates (μ_{\max}) (Table 2). Different model reductions not presented here were achieved. However,
 469 only the application of a sole and identical maximum uptake rate for all carbon sources of the
 470 culture medium (Table 3) led to the obtention of parameter estimations with acceptable
 471 confidence intervals (Table 4). Network interconnections between the assimilation pathways of
 472 each substrate could explain this phenomenon observed at the macroscopic level.

473

474

475 **Table 3:** Kinetic structures to describe the assimilations of multiple VFAs by *Rs. rubrum* after
 476 model reduction.

	Carbon source 1	Carbon source 2
Mix acetate propionate Case A.2	$\varphi_1 = X \cdot \mu_{\max_{Ace+Prop}} \cdot \frac{[Ace]}{K_{Ace} + [Ace]}$	$\varphi_2 = X \cdot \mu_{\max_{Ace+Prop}} \cdot \frac{[Prop]}{K_{Prop} + [Prop]}$
Mix acetate butyrate Case B.2	$\varphi_1 = X \cdot \mu_{\max_{Ace+but}} \cdot \frac{[Ace]}{K_{Ace} + [Ace]}$	$\varphi_2 = X \cdot \mu_{\max_{Ace+But}} \cdot \frac{[But]}{K_{But} + [But]} \cdot \frac{[Ace]}{K_{iAce} + [Ace]}$
Mix propionate butyrate Case C.2	$\varphi_2 = X \cdot \mu_{\max_{Prop+But}} \cdot \frac{[Prop]}{K_{Prop} + [Prop]}$	$\varphi_2 = X \cdot \mu_{\max_{Prop+But}} \cdot \frac{[But]}{K_{But} + [But]} \cdot \frac{[Prop]}{K_{iProp} + [Prop]}$
Mix acetate propionate butyrate Case D.2	$\varphi_1 = X \cdot \mu_{\max_{3VFA}} \cdot \frac{[Ace]}{K_{Ace} + [Ace]}$	$\varphi_2 = X \cdot \mu_{\max_{3VFA}} \cdot \frac{[Prop]}{K_{Prop} + [Prop]}$
	Carbon source 3	
	$\varphi_3 = X \cdot \mu_{\max_{3VFA}} \cdot \frac{[But]}{K_{But} + [But]} \cdot \frac{[Ace]}{K_{iAce} + [Ace]} \cdot \frac{[Prop]}{K_{iProp} + [Prop]}$	

477

478 Indeed, propionyl-CoA and acetyl-CoA share a common metabolic assimilation
 479 pathway as propionyl-CoA is the final product of the EMC, i.e. the anaplerotic pathway
 480 involved in acetyl-CoA assimilation [24,33]. Therefore acetyl-CoA and propionyl-CoA
 481 assimilations occur via the same metabolic road, that converts propionyl-CoA in succinyl-CoA
 482 via a methylmalonyl intermediate before it enters into the TCA cycle [24,33]. In addition, the
 483 assimilation of butyrate occurs via the EMC and the MBC pathways which exhibit propionyl-
 484 CoA and acetyl-CoA as intermediates. Therefore, it can be assumed that the maximum rates of
 485 acetate, propionate and butyrate assimilations could be the same.

486

487

488

489 **Table 4:** Parameter estimations and their confidence intervals for several models describing the
 490 assimilations of VFA mixtures after model reduction

Case	Mix acetate propionate (A.1)		Mix acetate butyrate (B.2)		Mix propionate butyrate (C.2)		Mix acetate propionate butyrate (D.2)	
	Value	CI (%)	Value	CI (%)	Value	CI (%)	Value	CI (%)
$\mu_{\max_{\text{VFA}}}$	0.14	46	0.10	151	0.05	157	0.003	9234
K_{ace}	1.56	4	1.03	15	0.82	19	27.3	1
K_{prop}	60.85	1	0.56	28	0.22	33	0.000	37231 91
K_{but}	-	-	-	-	-	-	3552	0,01
Ki_{ace}	-	-	0.24	65	-	-	0.004	7455
Ki_{prop}	-	-	-	-	0.10	74	0.007	4448
C_{ace}	0.66	10	0.88	18	-	-	56.76	1
C_{prop}	2.00	1	-	-	0.90	8	61.15	1
C_{but}	-	-	0.35	45	18.86	10	40709	0.001
J	0.24		0.40		0.23		17.5	

491

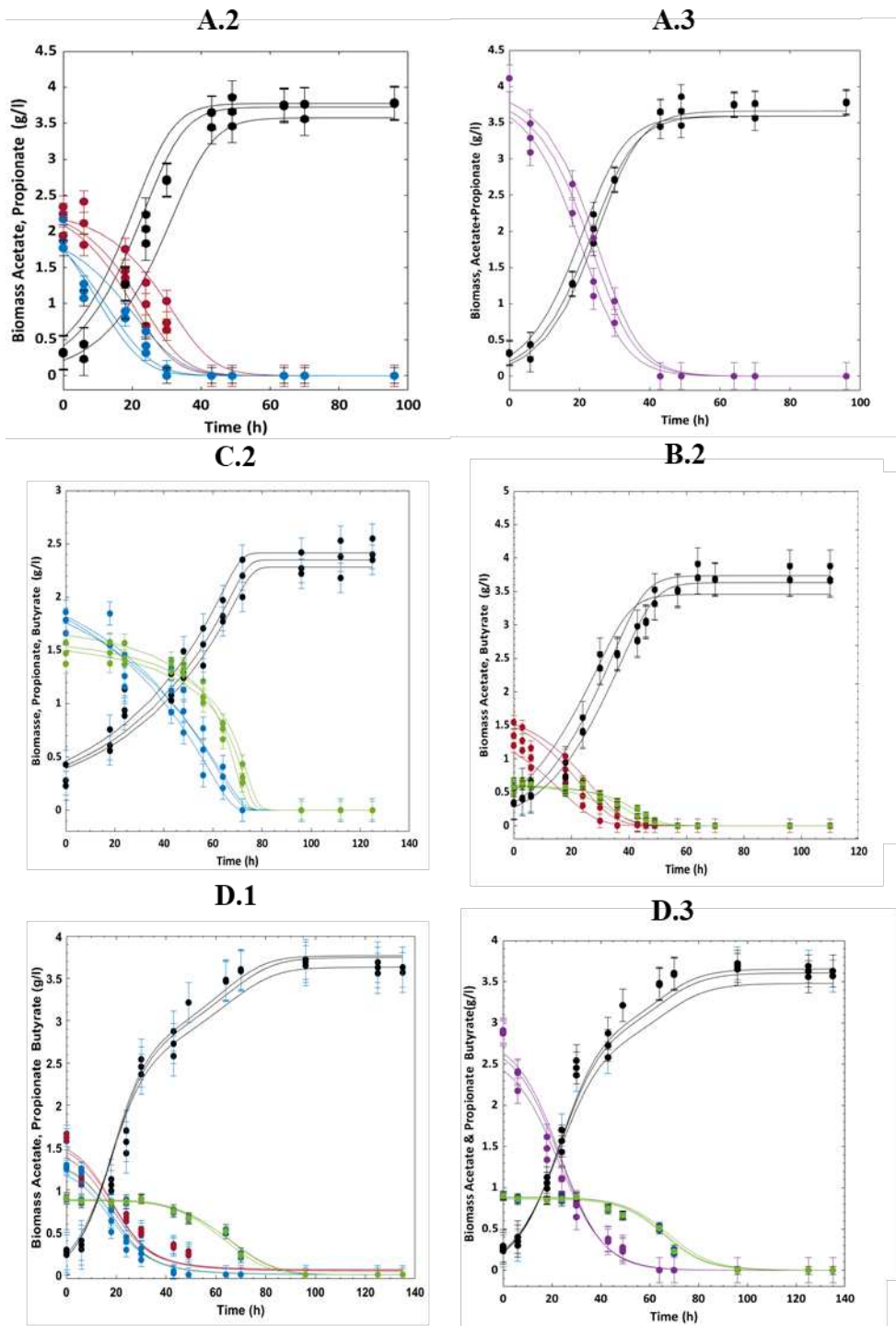
492 Regarding the mixture of three VFAs, the inhibition parameters present large confidence
 493 intervals (Table 4). This could be explained by the fact that propionate and acetate are
 494 simultaneously assimilated and as such, it is not possible to differentiate the specific inhibition
 495 induced by each substrate separately. The same observation can be made regarding the
 496 saturation constants. A model reduction is therefore proposed, removing one saturation and one
 497 inhibition constant. All the resulting possible combinations were tested, and the results of the
 498 parameter estimations are presented in supplementary data in Table S.4. This model reduction
 499 improves the fitting but provides a poor sensitivity of the inhibition and saturation constants,
 500 which induces a probable over-parametrization (Table S.4). Unfortunately, no contributive
 501 information from the metabolic pathways allows discriminating among the several possibilities
 502 to suppress inhibition or saturation constants.

503 **Table 5:** Kinetic structures to describe the assimilations of multiple VFAs by *Rs. rubrum* when
 504 acetate and propionate are considered as one sole substrate.

	Carbon source 1	Carbon source 2
Mix acetate and propionate Case A.3	$\varphi_1 = X \cdot \mu_{\max_{Ace+Prop}} \cdot \frac{[Ace] + [Prop]}{K_{Ace+Prop} + [Ace] + [Prop]}$	
Mix acetate propionate and butyrate Case D.3	$\varphi_1 = X \cdot \mu_{\max_{3\ VFA}} \cdot \frac{[Ace] + [Prop]}{K_{Ace+Prop} + [Ace] + [Prop]}$	
	Carbon source 3	
	$\varphi_2 = X \cdot \mu_{\max_{3\ VFA}} \cdot \frac{[But]}{K_{But} + [But]} \cdot \frac{[Prop] + [Ace]}{K_{i_{Prop+Ace}} + [Ace] + [Prop]}$	

505

506 This is comparable to the phenomenon observed during model development for the
 507 description of the uptakes of acetate and propionate. A model describing the uptakes of acetate,
 508 butyrate, and propionate considering butyrate on one hand, and a combination of acetate and
 509 propionate, on the other hand, was thus challenged (Table 5). Interestingly, the resulting fitting
 510 is the best and the parameter estimation is also the most accurate even if the inhibition constant
 511 still presents a relatively poor sensitivity which can be explained by its relatively low value,
 512 therefore difficult to identify with the available data (Table 6).



513

514 **Figure 5:** Experimental monitoring (dot) and simulated value (line), of biomass growth(●),
 515 acetate (●), propionate(●), and butyrate (●) consumptions in cultures of *Rs. rubrum* S1H. The
 516 substrate shown in (●) is composed of acetate and propionate. The bars represent the
 517 confidence intervals at 95% calculated from the a posteriori measurement error variance from
 518 (6).

519 In figure 5 are presented in the measured data and the respective predictions of the
 520 calibrated models describing scenarios A.2, A.3, B.2, C.2, D.2, and D.3 that exhibit both good
 521 fitting and satisfactory parameter estimation.

522

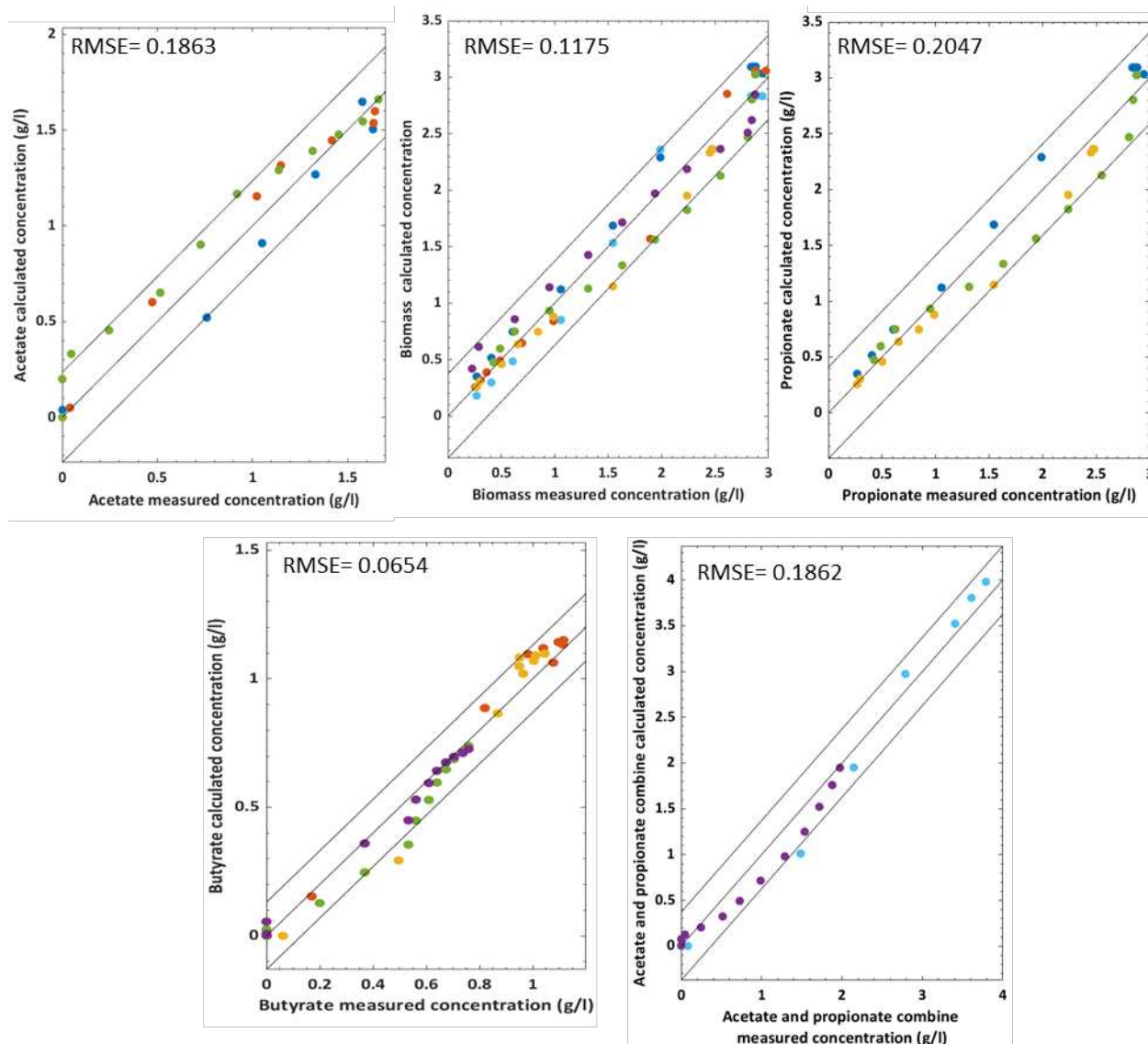
523 **Table 6:** Parameter estimations and their confidence intervals for several models describing the
 524 assimilations of VFA mixtures when acetate and propionate are considered as one substrate

	Mix acetate and propionate (A.3)		Mix acetate propionate and butyrate (D.3)	
	Value	CI (%)	Value	CI %
μ_{maxVF}	0.47	29	2.61	4
K_{VFA}	10.98	1	67.14	0.2
K_{but}	-	-	94.87	0.1
$K_{\text{ace+Pro}}$	-	-	0.025	491
C_{VFA}	1.08	13	0.95	10
C_{but}	-	-	1.36	7
J	0.086		0.33	

525

526 The best models were cross-validated using a data set obtained from an independent
 527 experiment. In the following, each state prediction is quantitatively assessed by its RMSE
 528 (Figure 6). The resulting small values confirm the good predictive capability of the model
 529 whatever the considered substrate combination. It can also be noticed that all predicted state
 530 trajectories are included in the 95% confidence intervals. The mixture of 3 VFAs as carbon
 531 source however shows the lowest predictive capability and parameter sensitivity. The
 532 development of those models brought a precise representation of multiple VFA uptakes by *Rs.*
 533 *rubrum* and in particular the different inhibition phenomena via a kinetic structure based on
 534 inhibition factors. In addition, the best representation is obtained when the assimilation occurs
 535 via the same pathway. Those macroscopic models present an added explicative value for the
 536 process in addition to a predictive capacity and, by extension, could serve as a digital twin basis

537 for process control and optimization [34,35]. Those models could also be useful in the
 538 development of software sensors for online VFA monitoring based on biomass measurement to
 539 ensure maximal biomass growth rate or prevent accumulation at inhibitory levels of implied
 540 metabolites.



541

542 **Figure 6:** Correlation between model predictions and measured data for each case scenario
 543 describing the assimilation of mixtures of VFAs by *Rs. rubrum*. Each color represents a specific
 544 case as follows: Case A.2(●), Case A.3(●), Case B.2(●), Case C.2 (●), Case D.1(●), Case
 545 D.3(●)

546 **4. Conclusion**

547 To better understand and control processes involving several VFAs, macroscopic
548 mechanistic models were developed in *Rs. rubrum*, to provide an accurate prediction of
549 substrate competition occurring when a mixture of acetate, propionate, and butyrate is used in
550 cultures of *Rs. rubrum*. The good predictive capability of the proposed models and the related
551 parameter accuracy, independently of the VFA mixture content, are quite encouraging and these
552 satisfactory results, therefore, open the door to a better design and operation of PNSB-based
553 industrial biotechnology for resource recovery, enabling high overall VFA assimilation
554 efficiencies without jeopardizing the overall process rates.

555

556 **Declaration of interest :** None

557 **Funding sources :**

558 This work was supported by the Belgian Fund for Scientific Research (Grand Equipment-F.R.S-
559 FNRS); the Concerted Research Action ARC project [P. Cabecas, PHASYN, 2017]; the CDR
560 -FNRS [B. Leroy, Redox homeostasis in purple bacteria]; the Research Foundation Flanders
561 [A.Alloul, 12W0522N]; the funded by the European Union's Horizon 2020 Research and
562 Innovation program on project 'Saraswati 2.0' [821427, 2020], the IOF via project PurpleRace
563 [A. Alloul, 40207].

564

565

566

567

568

569 **5. References**

- 570 [1] G. Moretto, F. Valentino, P. Pavan, M. Majone, D. Bolzonella, Optimization of urban
571 waste fermentation for volatile fatty acids production, *Waste Manag.* 92 (2019) 21–29.
- 572 [2] F. Valentino, G. Moretto, L. Lorini, D. Bolzonella, P. Pavan, M. Majone, Pilot-Scale
573 Polyhydroxyalkanoate Production from Combined Treatment of Organic Fraction of
574 Municipal Solid Waste and Sewage Sludge, *Ind. Eng. Chem. Res.* 58 (2019) 12149–
575 12158. <https://doi.org/10.1021/acs.iecr.9b01831>.
- 576 [3] W. Verstraete, P. Clauwaert, S.E. Vlaeminck, Used water and nutrients: recovery
577 perspectives in a ‘panta rhei’ context, *Bioresour. Technol.* 215 (2016) 199–208.
- 578 [4] D.-H.H. Kim, J.-H.H. Lee, Y. Hwang, S. Kang, M.-S.S. Kim, Continuous cultivation of
579 photosynthetic bacteria for fatty acids production, *Bioresour. Technol.* 148 (2013) 277–
580 282. <https://doi.org/10.1016/j.biortech.2013.08.078>.
- 581 [5] J.C. Fradinho, A. Oehmen, M.A.M.M. Reis, Photosynthetic mixed culture
582 polyhydroxyalkanoate (PHA) production from individual and mixed volatile fatty acids
583 (VFAs): Substrate preferences and co-substrate uptake, *J. Biotechnol.* 185 (2014) 19–
584 27. <https://doi.org/10.1016/j.jbiotec.2014.05.035>.
- 585 [6] A. Alloul, S. Wuyts, S. Lebeer, S.E. Vlaeminck, Volatile fatty acids impacting
586 phototrophic growth kinetics of purple bacteria: Paving the way for protein production
587 on fermented wastewater, *Water Res.* 152 (2019) 138–147.
588 <https://doi.org/10.1016/j.watres.2018.12.025>.
- 589 [7] M.-F. Dignac, P. Ginestet, D. Rybacki, A. Bruchet, V. Urbain, P. Scribe, Fate of
590 wastewater organic pollution during activated sludge treatment: nature of residual
591 organic matter, *Water Res.* 34 (2000) 4185–4194.
592 [https://doi.org/https://doi.org/10.1016/S0043-1354\(00\)00195-0](https://doi.org/https://doi.org/10.1016/S0043-1354(00)00195-0).
- 593 [8] A. Alloul, R. Ganigué, M. Spiller, F. Meerburg, C. Cagnetta, K. Rabaey, S.E.

- 594 Vlaeminck, Capture–Ferment–Upgrade: A Three-Step Approach for the Valorization
595 of Sewage Organics as Commodities, *Environ. Sci. Technol.* 52 (2018) 6729–6742.
596 <https://doi.org/10.1021/acs.est.7b05712>.
- 597 [9] S.-J. Lim, B.J. Kim, C.-M. Jeong, Y.H. Ahn, H.N. Chang, Anaerobic organic acid
598 production of food waste in once-a-day feeding and drawing-off bioreactor, *Bioresour.*
599 *Technol.* 99 (2008) 7866–7874.
- 600 [10] K. Komemoto, Y.G. Lim, N. Nagao, Y. Onoue, C. Niwa, T. Toda, Effect of
601 temperature on VFA's and biogas production in anaerobic solubilization of food waste,
602 *Waste Manag.* 29 (2009) 2950–2955.
- 603 [11] P. Zhang, Y. Chen, Q. Zhou, Waste activated sludge hydrolysis and short-chain fatty
604 acids accumulation under mesophilic and thermophilic conditions: effect of pH, *Water*
605 *Res.* 43 (2009) 3735–3742.
- 606 [12] J. Jiang, Y. Zhang, K. Li, Q. Wang, C. Gong, M. Li, Volatile fatty acids production
607 from food waste: Effects of pH, temperature, and organic loading rate, *Bioresour.*
608 *Technol.* 143 (2013) 525–530.
609 <https://doi.org/https://doi.org/10.1016/j.biortech.2013.06.025>.
- 610 [13] W.S. Lee, A.S.M. Chua, H.K. Yeoh, G.C. Ngoh, A review of the production and
611 applications of waste-derived volatile fatty acids, *Chem. Eng. J.* 235 (2014) 83–99.
- 612 [14] S. Dahiya, O. Sarkar, Y. V Swamy, S.V. Mohan, Acidogenic fermentation of food
613 waste for volatile fatty acid production with co-generation of biohydrogen, *Bioresour.*
614 *Technol.* 182 (2015) 103–113.
- 615 [15] H. Brandl, R.A. Gross, R.W. Lenz, R. Lloyd, R.C. Fuller, The accumulation of poly(3-
616 hydroxyalkanoates) in *Rhodobacter sphaeroides*, *Arch. Microbiol.* 155 (1991) 337–
617 340. <https://doi.org/10.1007/bf00243452>.
- 618 [16] H. Brandl, E.J. Knee, R.C. Fuller, R.A. Gross, R.W. Lenz, Ability of the phototrophic

- 619 bacterium *Rhodospirillum rubrum* to produce various poly (β -hydroxyalkanoates):
620 Potential sources for biodegradable polyesters, *Int. J. Biol. Macromol.* 11 (1989) 49–
621 55. [https://doi.org/10.1016/0141-8130\(89\)90040-8](https://doi.org/10.1016/0141-8130(89)90040-8).
- 622 [17] J.M.L. Dias, A. Oehmen, L.S. Serafim, P.C. Lemos, M.A.M. Reis, R. Oliveira,
623 Metabolic modelling of polyhydroxyalkanoate copolymers production by mixed
624 microbial cultures, *BMC Syst. Biol.* 2 (2008) 59.
- 625 [18] Y. Jiang, M. Heibly, R. Kleerebezem, G. Muyzer, M.C.M. van Loosdrecht, Metabolic
626 modeling of mixed substrate uptake for polyhydroxyalkanoate (PHA) production,
627 *Water Res.* 45 (2011) 1309–1321. <https://doi.org/10.1016/j.watres.2010.10.009>.
- 628 [19] L. Marang, Y. Jiang, M.C.M. van Loosdrecht, R. Kleerebezem, Butyrate as preferred
629 substrate for polyhydroxybutyrate production, *Bioresour. Technol.* 142 (2013) 232–
630 239.
- 631 [20] J. Tamis, L. Marang, Y. Jiang, M.C.M. van Loosdrecht, R. Kleerebezem, Modeling
632 PHA-producing microbial enrichment cultures—towards a generalized model with
633 predictive power, *N. Biotechnol.* 31 (2014) 324–334.
- 634 [21] X. Wang, G. Carvalho, M.A.M. Reis, A. Oehmen, Metabolic modeling of the substrate
635 competition among multiple VFAs for PHA production by mixed microbial cultures, *J.*
636 *Biotechnol.* 280 (2018) 62–69.
- 637 [22] H.L. Korneberg, H.A. Krebs, Synthesis of Cell Constituents from C2-Units by a
638 Modified Tricarboxylic Acid Cycle, *Nature.* 179 (1957) 988–991.
639 <https://doi.org/10.1038/179988a0>.
- 640 [23] T.J. Erb, I.A. Berg, V. Brecht, M. Müller, G. Fuchs, B.E. Alber, Synthesis of C5-
641 dicarboxylic acids from C2-units involving crotonyl-CoA carboxylase/reductase: the
642 ethylmalonyl-CoA pathway, *Proc. Natl. Acad. Sci.* 104 (2007) 10631–10636.
- 643 [24] B. Leroy, Q. De Meur, C. Moulin, G. Wegria, R. Wattiez, New insight into the

644 photoheterotrophic growth of the isocitrate lyase-lacking purple bacterium
645 *Rhodospirillum rubrum* on acetate, *Microbiology*. 161 (2015) 1061–1072.
646 <https://doi.org/10.1099/mic.0.000067>.

647 [25] K. Schneider, R. Peyraud, P. Kiefer, P. Christen, N. Delmotte, S. Massou, J.-C. Portais,
648 J.A. Vorholt, The ethylmalonyl-CoA pathway is used in place of the glyoxylate cycle
649 by *Methylobacterium extorquens* AM1 during growth on acetate, *J. Biol. Chem.* 287
650 (2012) 757–766.

651 [26] M. Knight, The photometabolism of propionate by *Rhodospirillum rubrum*, *Biochem.*
652 *J.* 84 (1962) 170.

653 [27] Q. De Meur, A. Deutschbauer, M. Koch, G. Bayon-Vicente, P. Cabecas Segura, R.
654 Wattiez, B. Leroy, New perspectives on butyrate assimilation in *Rhodospirillum*
655 *rubrum* S1H under photoheterotrophic conditions, *BMC Microbiol.* (2020) 1–20.

656 [28] Q. De Meur, A. Deutschbauer, M. Koch, R. Wattiez, B. Leroy, Genetic plasticity and
657 ethylmalonyl coenzyme A pathway during acetate assimilation in *Rhodospirillum*
658 *rubrum* S1H under photoheterotrophic conditions, *Appl. Environ. Microbiol.* 84 (2018)
659 e02038-17.

660 [29] L. Segers, W. Verstraete, Conversion of organic acids to H₂ by *Rhodospirillaceae*
661 grown with glutamate or dinitrogen as nitrogen source, *Biotechnol. Bioeng.* 25 (1983)
662 2843–2853.

663 [30] M. Suhaimi, J. Liessens, W. Verstraete, NH₄⁺/4-N assimilation by *Rhodobacter*
664 *capsulatus* ATCC 23782 grown axenically and non-axenically in N and C rich media, *J.*
665 *Appl. Bacteriol.* 62 (1987) 53–64.

666 [31] ESA, Engineering of the waste compartment - ESA contract 15689/01/NL/ND
667 TECHNICAL NOTE 71.9.4 – Confidential report, 2005.

668 [32] P. Cabecas Segura, Q. De Meur, A. Tanghe, R. Onderwater, L. Dewasme, R. Wattiez,

669 B. Leroy, Effects of Mixing Volatile Fatty Acids as Carbon Sources on *Rhodospirillum*
670 *rubrum* Carbon Metabolism and Redox Balance Mechanisms, *Microorg.* . 9 (2021).
671 <https://doi.org/10.3390/microorganisms9091996>.

672 [33] R.K. Clayton, Photosynthetic metabolism of propionate in *Rhodospirillum rubrum*,
673 *Arch. Mikrobiol.* 26 (1957) 29–31.

674 [34] J.H. Lee, Model predictive control: Review of the three decades of development, *Int. J.*
675 *Control. Autom. Syst.* 9 (2011) 415–424.

676 [35] S.J. Qin, T.A. Badgwell, A survey of industrial model predictive control technology,
677 *Control Eng. Pract.* 11 (2003) 733–764.

678

679

680

681

682

683

684

685

686

687

688

689

690

