

Fractionation of stable carbon isotopes during microbial propionate consumption in anoxic rice paddy soils

Ralf Conrad¹, Peter Claus¹

¹Max Planck Institute for Terrestrial Microbiology, Karl-von-Frisch-Str. 10, 35043 Marburg, Germany

Correspondence to: Ralf Conrad (Conrad@mpi-marburg.mpg.de)

Abstract. Propionate is an important intermediate during the breakdown of organic matter in anoxic flooded paddy soils. Since there are only few experiments on carbon isotope fractionation and the magnitude of the isotopic enrichment factors (ϵ) involved, we measured propionate conversion to acetate, CH₄ and CO₂ in anoxic paddy soils. Propionate consumption was measured using samples of paddy soil from Vercelli (Italy) and the International Rice Research Institute (IRRI, the Philippines) suspended in phosphate buffer (pH 7.0), both in the absence and presence of sulfate (gypsum), and of methyl fluoride (CH₃F), an inhibitor of acetoclastic methanogenesis. Under methanogenic conditions, propionate was eventually degraded to CH₄ with acetate being a transient intermediate. Butyrate was also a minor intermediate. Methane was mainly produced by acetoclastic methanogenesis. Propionate consumption was inhibited by CH₃F. Whereas butyrate and CH₄ were ¹³C-depleted relative to propionate, acetate and CO₂ were ¹³C-enriched. The isotopic enrichment factors (ϵ_{prop}) of propionate consumption, determined by Mariotti plots, were in a range of -8‰ to -3.5‰. Under sulfidogenic conditions, acetate was also transiently accumulated, but CH₄ production was negligible. Application of CH₃F hardly affected propionate degradation and acetate accumulation. The initially produced CO₂ was ¹³C-depleted, whereas the acetate was ¹³C-enriched. The values of ϵ_{prop} were -3.5‰. It is concluded that degradation of organic carbon via propionate to acetate and CO₂ involves only little isotope fractionation. The results further indicate a major contribution of *Syntrophobacter*-type propionate fermentation under sulfidogenic conditions and *Smithella*-type propionate fermentation under methanogenic conditions. This interpretation is consistent with data of the microbial community composition published previously for the same soils.

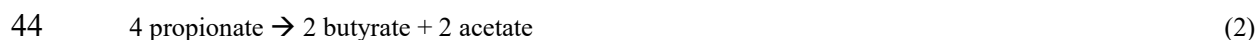
28 1 Introduction

29 Propionate is a common intermediate of organic matter degradation in anoxic paddy soils. In the absence of
30 sulfate reduction or methanogenesis propionate may accumulate to millimolar concentrations (Conrad et al., 2014;
31 Glissmann and Conrad, 2000; Nozoe, 1997). Under methanogenic conditions propionate is degraded by
32 fermentation. Several different biochemical pathways are conceivable for propionate fermentation (Textor et al.,
33 1997). The major fermentation pathways are those by *Syntrophobacter* (Boone and Bryant, 1980) and *Smithella*
34 (Liu et al., 1999) both members of Deltaproteobacteria. *Syntrophobacter* operates the methylmalonyl-CoA
35 pathway, which results in randomization of the carbon positions of propionate (Houwen et al., 1991). This pathway
36 can also be found in *Desulfotomaculum* sp. and *Pelotomaculum* sp. (Chen et al., 2005; DeBok et al., 2005; Imachi
37 et al., 2002; Plugge et al., 2002), and apparently exists in many anoxic environments (Imachi et al., 2006; Krylova
38 et al., 1997; Schink, 1985). *Smithella*, on the other hand, operates a dismutation pathway, which does not result in
39 randomization (DeBok et al., 2001). This pathway has also been found in many anoxic environments (Gan et al.,
40 2012; Lueders et al., 2004; Xia et al., 2019).

41 Propionate degradation by randomizing *Syntrophobacter* proceeds via succinate in the following way:



43 Propionate degradation by non-randomizing *Smithella* proceeds by dismutation of propionate:



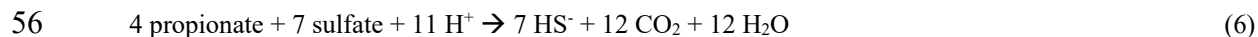
45 Butyrate is then syntrophically converted (e.g., by *Syntrophomonas* (McInerney et al., 1981)):



47 The *Smithella* pathway in total:



49 Propionate fermentation is thermodynamically endergonic under standard conditions and therefore, requires
50 syntrophic microbial partners that further convert the fermentation products. Under methanogenic conditions, the
51 syntrophic partners are methanogenic archaea, which consume the products acetate and H₂. Under sulfidogenic
52 conditions sulfate-reducing bacteria replace the methanogens. Propionate can also be directly oxidized to CO₂ by
53 propionate-degrading sulfate reducers. The overall reaction stoichiometry is the same for *Syntrophobacter* and
54 *Smithella*:



57 Note, that the relative production of acetate and H₂ is different for *Syntrophobacter* and *Smithella* fermentation,
58 being 1:3 and 3:2, respectively. Therefore, acetate-producing methanogenesis contributes relatively more than
59 hydrogenotrophic methanogenesis, when propionate is fermented by *Smithella* rather than *Syntrophobacter*. Under
60 methanogenic conditions, propionate degradation in anoxic paddy soils operates close to the thermodynamic limits
61 (Krylova and Conrad, 1998; Yao and Conrad, 2001). These restrictions are more severe for *Syntrophobacter* than
62 for *Smithella* (Dolfing, 2013).

63 Using paddy soil from Italy and the Philippines Liu and coworkers (Liu et al., 2018a; Liu and Conrad, 2017)
64 have recently shown that propionate consumption under sulfidogenic conditions is mainly achieved by
65 *Syntrophobacter* species or other Syntrophobacteraceae, which first oxidize propionate to acetate and CO₂, and
66 subsequently oxidize the accumulated acetate to CO₂. They also showed that *Smithella* was probably involved in

67 methanogenic propionate degradation. The involvement of *Smithella* has also been shown for other paddy soils
68 and sediments (Gan et al., 2012; Lueders et al., 2004; Xia et al., 2019). Since we used in the present study the same
69 soils as Liu and coworkers (Liu et al., 2018a; Liu and Conrad, 2017), we assumed that propionate degradation was
70 achieved by the same microorganisms.

71 Knowledge of carbon isotope fractionation is important for the assessment of the pathways involved in
72 anaerobic degradation of organic matter (Conrad, 2005; Elsner et al., 2005). The $\delta^{13}\text{C}$ values of organic carbon,
73 acetate and propionate in various soils and sediments were found to be similar (Conrad et al., 2014). The similarity
74 indicates that the enrichment factors (ϵ) of the processes involved in both production and consumption of
75 propionate are probably small. The direct determination of ϵ values in microbial cultures of one propionate-
76 producing and one propionate-consuming bacterium also showed low values (Botsch and Conrad, 2011). However,
77 direct determination of ϵ values in environmental samples is missing. Therefore, we decided to measure isotope
78 fractionation in methanogenic and sulfidogenic paddy soil amended with propionate along with the recording of
79 the production of acetate, CH_4 and CO_2 . We also used the treatment with methyl fluoride (CH_3F) to inhibit the
80 consumption of acetate by methanogenic archaea (Janssen and Frenzel, 1997). Recently, we determined the
81 microbial communities in methanogenic and sulfidogenic rice field soils, which were used for assessment of ^{13}C
82 isotope fractionation during acetate consumption (Conrad et al., 2021). Here we present analogous data from the
83 same soil suspensions prepared for the propionate degradation experiments.

84

85 **2 Materials and Methods**

86 *2.1 Paddy soils and incubation conditions*

87 The soil samples were from the research stations in Vercelli, Italy and the International Rice research Institute
88 (IRRI) in the Philippines. Sampling and soil characteristics were described before (Liu et al., 2018b). The main
89 soil characteristics will be given. The Italian soil is a sandy loam with a pH of 5.75, total C of 1.1% and total N
90 of 0.08%. The Philippine soil is a silt loam with a pH of 6.3, total C of 1.9% and total N of 0.2%.

91 The experimental setup was exactly the same as during a previous study on acetate consumption (Conrad et
92 al., 2021). Paddy soil was mixed with autoclaved anoxic H_2O at a ratio of 1:1 and incubated under N_2 at 25°C for
93 4 weeks. In a second incubation, paddy soil was mixed with autoclaved anoxic H_2O (prepared under N_2) at a ratio
94 of 1:1, was amended with 0.07 g $\text{CaSO}_4 \cdot 2\text{H}_2\text{O}$, and then incubated under N_2 at 25°C for 4 weeks. These two
95 preincubated soil slurries were sampled and stored at -20°C for later molecular analysis (see data in Conrad et al.
96 (2021)). The preincubated soil slurries were also used (in 3 replicates) for the following incubation experiments.
97 Two different sets of incubations were prepared. In the first set (resulting in methanogenic conditions), 5 mL soil
98 slurry preincubated without sulfate was incubated at 25°C with 40 mL 20 mM potassium phosphate buffer (pH
99 7.0) in a 150-mL bottle under an atmosphere of N_2 . The bottles were amended with (i) 5 mL H_2O ; (ii) 5 mL
100 H_2O + 4.5 mL CH_3F ; (iii) 5 mL 50 mM sodium propionate; (iv) 5 mL 50 mM sodium acetate + 4.5 mL CH_3F . In
101 the second set (resulting in sulfidogenic conditions), 5 mL soil slurry preincubated with sulfate was incubated at
102 25°C with 40 mL 20 mM potassium phosphate buffer (pH 7.0) in a 150-mL bottle under an atmosphere of N_2 . The
103 amendments were the same as above, but with the addition of 200 μl of a CaSO_4 suspension corresponding to a
104 concentration of 2.5 M (giving a final concentration of 10 mM sulfate).

105

106 2.2 Chemical and isotopic analyses

107 Gas samples for analysis of partial pressures of CH₄ and CO₂ were taken from the headspace of the incubation
108 bottles after vigorous manual shaking for about 30 s using a gas-tight pressure-lock syringe, which had been
109 flushed with N₂ before each sampling. Soil slurries were sampled, centrifuged and filtered through a 0.2 μm
110 cellulose membrane filter and stored frozen at -20°C for later fatty acid analysis. Chemical and isotopic analyses
111 were performed as described in detail previously (Goevert and Conrad, 2009). Methane was analyzed by gas
112 chromatography (GC) with flame ionization detector. Carbon dioxide was analyzed after conversion to CH₄ with
113 a Ni catalyst. Stable isotope analyses of ¹³C/¹²C in gas samples were performed using GC-combustion isotope ratio
114 mass spectrometry (GC-C-IRMS). Propionate, butyrate and acetate were measured using high-performance liquid
115 chromatography (HPLC) linked via a Finnigan LC IsoLink to an IRMS. The isotopic values are reported in the
116 delta notation (δ¹³C) relative to the Vienna Pee Dee Belemnite standard having a ¹³C/¹²C ratio (R_{standard}) of 0.01118:
117 δ¹³C = 10³ (R_{sample}/R_{standard} - 1). The precision of the GC-C-IRMS was ± 0.2‰, that of the HPLC-IRMS was ±
118 0.3‰.

119

120 2.3 Calculations

121 Millimolar concentrations of CH₄ were calculated from the mixing ratios (1 ppmv = 10⁻⁶ bar) measured in the
122 gas phase of the incubation bottles: 1000 ppmv CH₄ correspond to 0.09 μmol per mL of liquid. Note, that this is
123 the total amount of CH₄ in the gas phase relative to the liquid phase.

124 Fractionation factors for reaction A → B are defined after Hayes (Hayes, 1993) as:

$$125 \alpha_{A/B} = (\delta_A + 1000) / (\delta_B + 1000) \quad (7)$$

126 also expressed as ε ≡ 1000 (1 - α) in permil. The carbon isotope enrichment factor ε_{prop} associated with propionate
127 consumption was calculated from the temporal change of δ¹³C of propionate as described by Mariotti et al.
128 (Mariotti et al., 1981) from the residual reactant

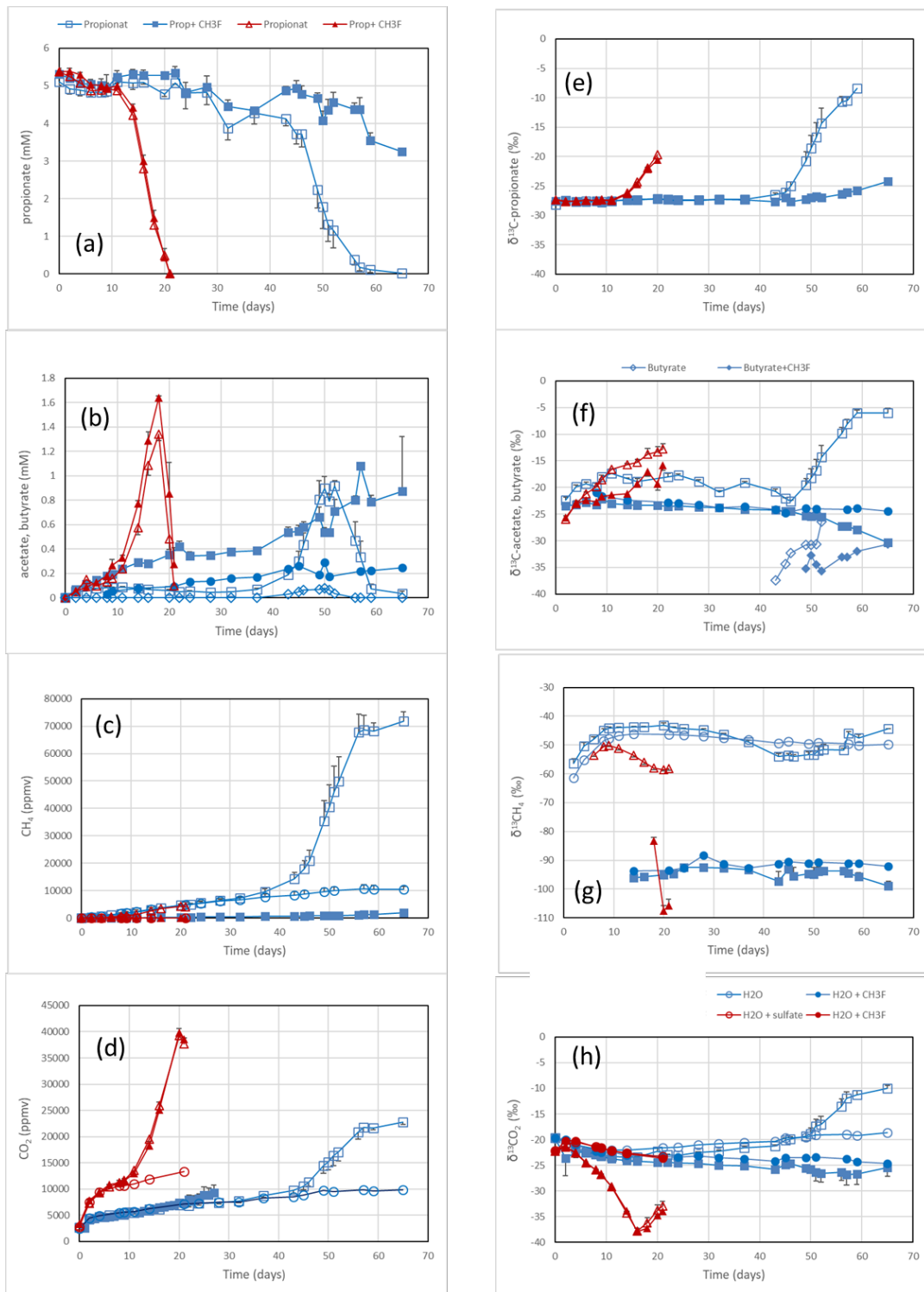
$$129 \delta_r = \delta_{ri} + \varepsilon [\ln(1-f)] \quad (8)$$

130 where δ_{ri} is the isotopic composition of the reactant (propionate) at the beginning, and δ_r is the isotopic composition
131 of the residual propionate, both at the instant when *f* is determined. *f*_{prop} is the fractional yield of the products based
132 on the consumption of propionate (0 < *f*_{prop} < 1). Linear regression of δ¹³C of propionate against ln(1 - *f*) yields
133 ε_{prop} as the slope of best fit lines. The regressions of δ¹³C of propionate were done for data in the range of *f*_{prop} <
134 0.7. The linear regressions were done individually for each experimental replicate (n = 3) and were only accepted
135 if r² > 0.9. The ε values resulting from the replicate experiments were then averaged (± SE).

136 The fraction (*f*_{H2}) of CH₄ derived from hydrogenotrophic methanogenesis was determined as described before
137 (Conrad et al., 2010) using

$$138 f_{H2} = (\delta^{13}C_{CH4} - \delta^{13}C_{CH4-ma}) / (\delta^{13}C_{CH4-mc} - \delta^{13}C_{CH4-ma}) \quad (9)$$

139 with δ¹³C_{CH4} = δ¹³C of total CH₄ produced, δ¹³C_{CH4-mc} = δ¹³C of CH₄ produced from hydrogenotrophic
140 methanogenesis, which is equivalent to the CH₄ produced in the presence of CH₃F, and δ¹³C_{CH4-ma} = δ¹³C of CH₄
141 produced from acetoclastic methanogenesis. The δ¹³C_{CH4-ma} was approximated from the δ¹³C of acetate in the
142 presence of CH₃F assuming that the methyl group of acetate was depleted in ¹³C by 8‰ (Conrad et al., 2014) and
143 that the enrichment factor (ε_{CH4,ac-methyl}) for CH₄ being produced from acetate-methyl was between 0 and -20‰.



144
 145
 146
 147
 148
 149
 150
 151
 152

Figure 1: Propionate conversion to acetate, butyrate, CH₄ and CO₂ in suspensions of paddy soil from Vercelli (Italy) after addition of propionate without sulfate (blue squares) or propionate plus sulfate (gypsum) (red triangles) without CH₃F (open symbols) or with CH₃F (closed symbols). Controls with addition of only water (blue or red circles) are only shown occasionally. The panels show the temporal change of (a) concentrations of propionate, (b) concentrations of acetate and butyrate (blue diamonds), (c) mixing ratios of CH₄ (1 ppmv = 10⁻⁶ bar), (d) mixing ratios of CO₂, (e) δ¹³C of propionate, (f) δ¹³C of acetate and butyrate, (g) δ¹³C of CH₄, and (h) δ¹³C of CO₂. Means ± SE.

153

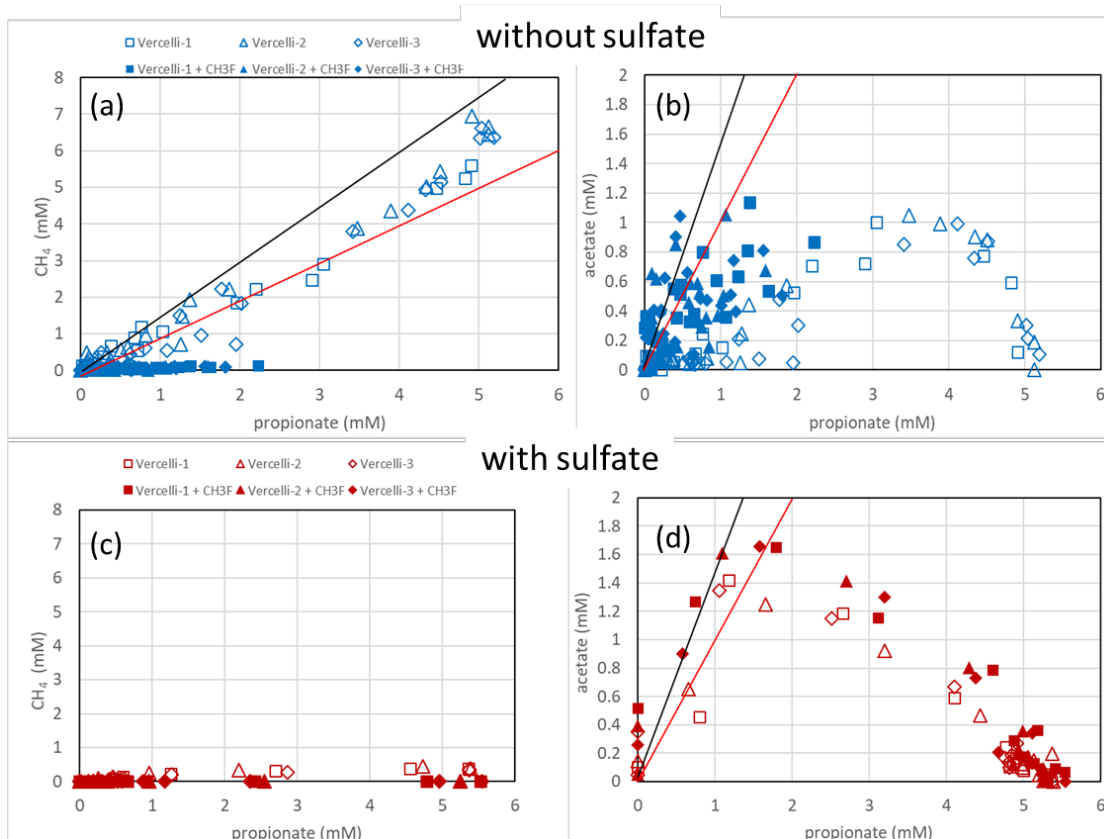
154 3 Results

155 3.1 Conversion of propionate under methanogenic and sulfidogenic conditions

156 Incubation of buffered suspensions of rice field soil from Vercelli (Fig. 1) and the IRRI (Fig. S1) resulted in
157 similar patterns of propionate degradation to acetate, CH₄ and CO₂. Under methanogenic conditions in the absence
158 of sulfate, propionate degradation started after a lag phase of about 20 d (Fig. 1a) resulting in the production of
159 acetate (Fig. 1b), CH₄ (Fig. 1c) and CO₂ (Fig. 1d). The formation of acetate, CH₄ and CO₂ in the absence of
160 propionate was only very small. The accumulation of acetate was only transient, except when acetate
161 methanogenesis was inhibited by CH₃F (Fig. 1b). Similar observations were made in IRRI soil (Fig. S1a-d). The
162 production of CH₄ was roughly equimolar to the consumption of propionate, but was nearly zero when acetate
163 methanogenesis was inhibited by CH₃F (Fig. 2a). Under these conditions, acetate accumulated to nearly equimolar
164 amounts with the consumed propionate (Fig. 2b), but in IRRI soil acetate accumulation was less than equimolar
165 (Fig. S2b). Butyrate was also a transient intermediate of propionate degradation and was produced and consumed
166 simultaneously with acetate (Fig. 1b, S1b). However, the accumulated concentrations were small (<0.1 mM).

167 In the presence of sulfate, propionate degradation started after a lag phase of only about 10 days (Fig. 1a)
168 resulting in the accumulation of acetate (Fig. 1b) and the production of CO₂ (Fig. 1d), but CH₄ production was
169 close to zero (Fig. 1c). Similar results were obtained with IRRI soil (Fig. S1a-d). The accumulated acetate was
170 equimolar (slightly less than equimolar in the IRRI soil (Fig. S2d)) to the consumption of propionate (Fig. 2d), but
171 CH₄ was not accumulated (Fig. 2c). Addition of CH₃F had no effect. Butyrate was not detected. The accumulated
172 acetate was subsequently degraded resulting in further production of CO₂ (Fig. 1b,d).

173



174

175 **Figure 2:** Balance of (a, c) produced CH₄ and (b, d) produced acetate against the consumed propionate under (a,
176 b) methanogenic and (c, d) sulfidogenic conditions in paddy soil from Vercelli (Italy). The open and closed
177 symbols denote conditions in the absence and the presence of CH₃F, respectively. The black and red lines in panel
178 (a) indicate aceticlastic methanogenesis after generation of acetate by either *Smithella* (equ.4) or *Syntrophobacter*
179 (equ.1). The black and red lines in panel b and d indicate transient acetate production by *Smithella* and
180 *Syntrophobacter*, respectively. The different symbols indicate three different replicates.

181

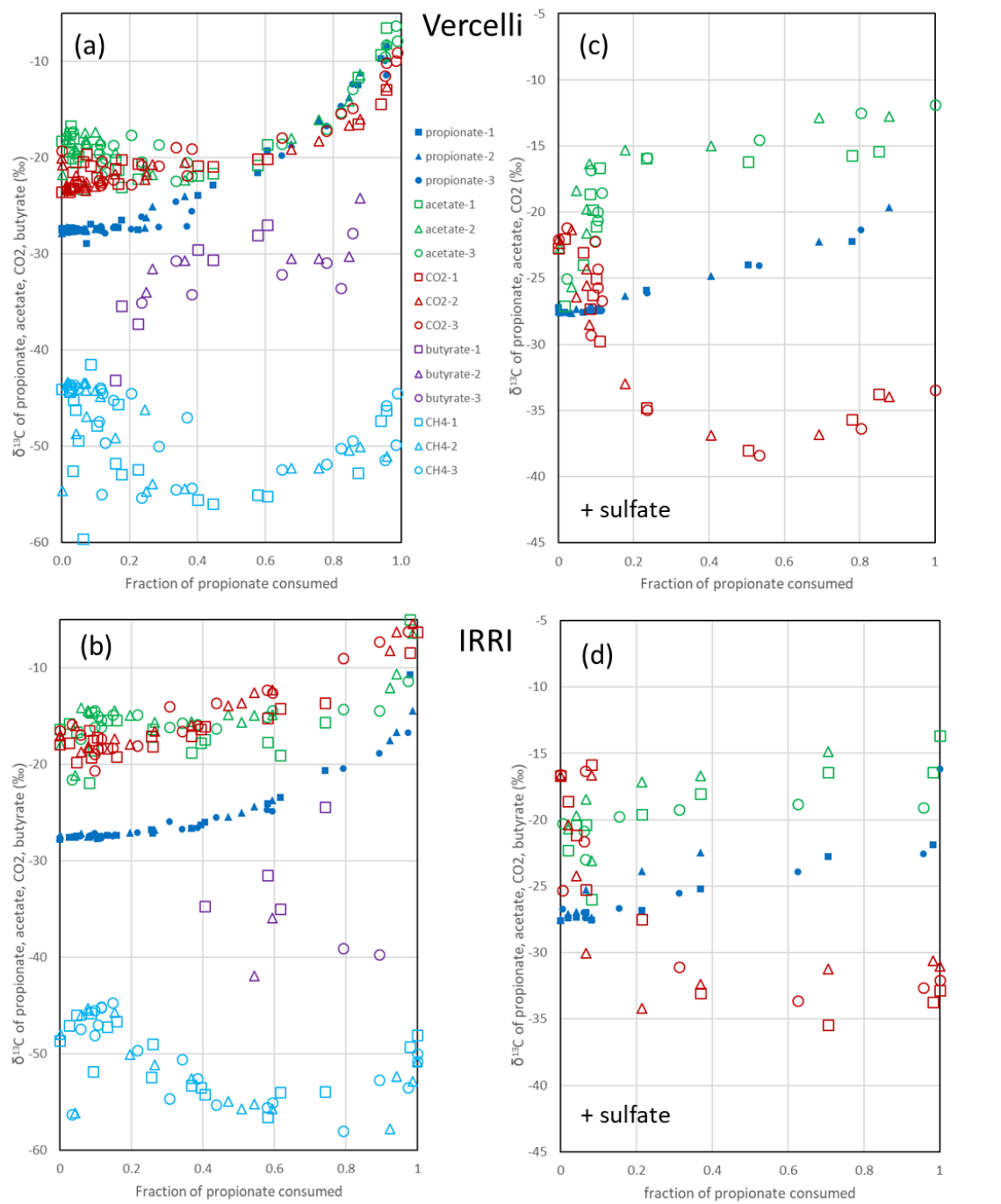
182 3.2 Isotope fractionation during propionate degradation

183 After onset of propionate degradation, the $\delta^{13}\text{C}$ of propionate (Fig. 1e) and acetate (Fig. 1f) increased indicating
184 that the light isotope was preferentially consumed. The $\delta^{13}\text{C}$ values of CO₂ also increased (Fig. 1h). The same was
185 the case for butyrate (Fig. 1f). Similar results were obtained with IRRI soil (Fig. S1e-h). When aceticlastic
186 methanogenesis was inhibited by CH₃F, the $\delta^{13}\text{C}$ values of these compounds increased only slightly or decreased
187 (Fig. 1e,f,h). However, the $\delta^{13}\text{C}$ of CH₄ was much more negative (30-50‰) in the presence than in the absence of
188 CH₃F (Fig. 1g). The $\delta^{13}\text{C}$ values of CH₄ in unamended soil (H₂O control) were similar to those in propionate
189 amended soil (Fig. 1g). To visualize the change of the metabolic ^{13}C content of the metabolic products relative to
190 the substrates, the $\delta^{13}\text{C}$ values were plotted against the increasing fractions (f_{prop}) of propionate consumed both in
191 soil from Vercelli (Fig.3a) and the IRRI (Fig.3b). The patterns of $\delta^{13}\text{C}$ values against the f_{prop} indicated kinetic
192 isotope fractionation. Note that the $\delta^{13}\text{C}$ values of acetate and CO₂ were higher than those of propionate, whereas
193 the values of butyrate and CH₄ were lower (Fig.3a,b). The $\delta^{13}\text{C}$ of CH₄ decreased until about 40% of the propionate
194 had been consumed, and then increased again to its initial (low) values (-50‰ to -45‰) (Fig.3a,b).

195 Under sulfidogenic conditions, only very little CH₄ was produced. Similarly as under methanogenic conditions,
196 the $\delta^{13}\text{C}$ of propionate (Fig. 1e) and of acetate (Fig. 1f) increased after onset of propionate degradation indicating
197 that the light isotope was preferentially consumed. However, the $\delta^{13}\text{C}$ values of CO₂ decreased during the first 10-
198 15 days when acetate was accumulated (Fig. 1h, S1h). Inhibition of aceticlastic methanogenesis by CH₃F had no
199 effect on the $\delta^{13}\text{C}$ of propionate and CO₂, but the values of acetate increased less than in the absence of CH₃F (Fig.
200 1f). Also, $\delta^{13}\text{C}$ of CH₄ was lower in the presence than in the absence of CH₃F (Fig. 1g), but the amounts of CH₄
201 produced were only very small (Fig. 1c). The values of $\delta^{13}\text{C}$ of propionate and acetate increased with increasing
202 f_{prop} (Fig. 3c,d). The $\delta^{13}\text{C}$ of acetate was generally by about 5-10‰ higher than the $\delta^{13}\text{C}$ of propionate but also
203 increased with f_{prop} indicating kinetic isotope fractionation. However, the $\delta^{13}\text{C}$ of CO₂ did not increase, but instead
204 decreased after onset of propionate degradation reaching about -35‰ when 50% of the propionate had been
205 consumed and acetate accumulation had reached a maximum (Fig. 3c,d). Thereafter, $\delta^{13}\text{C}$ of CO₂ increased or
206 became constant.

207 Mariotti plots of the ^{13}C of propionate as function of f_{prop} could be created for methanogenic and sulfidogenic
208 incubation conditions, the latter both in the absence and the presence of CH₃F (Fig. 4). The lines were straight even
209 when more than 70% of the propionate was consumed. Nevertheless, enrichment factors (ϵ) were determined only
210 for $f_{prop} < 0.7$ and for regressions giving $r^2 > 0.9$. The ϵ_{prop} values were determined for each individual incubation
211 and then averaged over the replicates (n = 2-3). The results for Vercelli and IRRI soils are summarized in Fig. 5.
212 The average ϵ_{prop} values under methanogenic conditions were about -8‰ for Vercelli and about -3.5‰ for IRRI

213 soil. The average ϵ_{prop} values under sulfidogenic conditions were around -3.5‰ in both soils and irrespectively
 214 whether CH_3F was present or not.

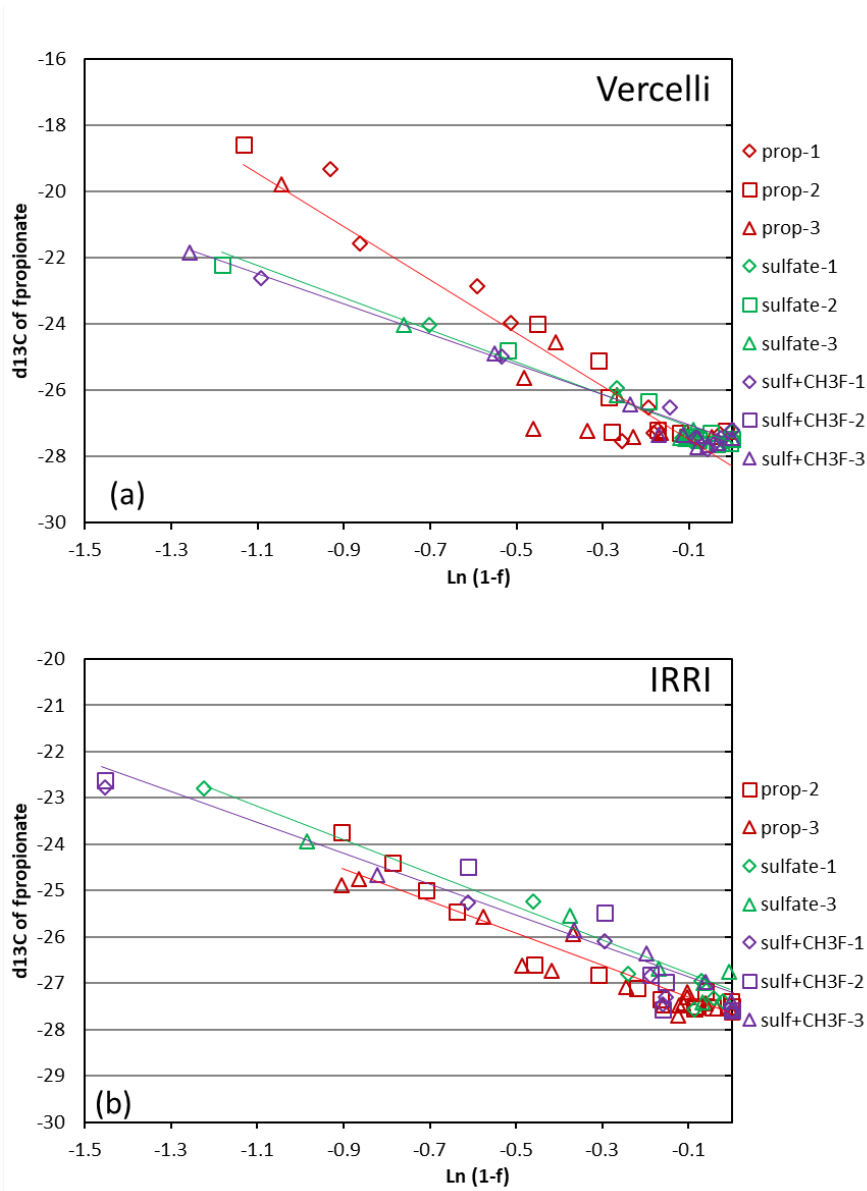


215
 216 **Figure 3:** Change of $\delta^{13}C$ of propionate, acetate, butyrate, CO_2 and CH_4 relative to the fraction of propionate
 217 consumed (f_{prop}) under (a, b) methanogenic and (c, d) sulfidogenic conditions in paddy soil from (a, c) Vercelli
 218 (Italy) and (b, d) the IIRRI (the Philippines). The different symbols indicate three different replicates.

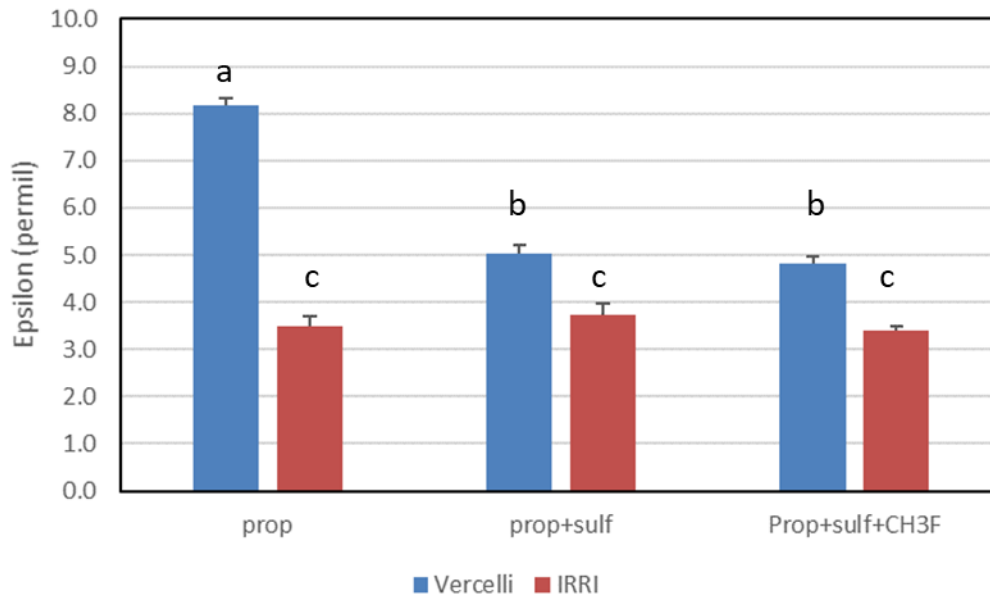
219
 220 **3.3 Hydrogenotrophic methanogenesis**

221 The difference in the $\delta^{13}C$ of CH_4 in the presence and the absence of CH_3F was used together with the $\delta^{13}C$ of
 222 acetate to roughly estimate the percentage of CH_4 derived from H_2/CO_2 versus acetate (Fig. S3). The percentage
 223 fractions of hydrogenotrophic methanogenesis (f_{H_2}) in Vercelli soil reached a maximum after 40-50 d when acetate

224 concentrations also reached a maximum (Fig. S3a) and then decreased strongly. The same was the case in IRRI
 225 soil after around 35 d (Fig. S3b). When assuming a reasonable isotopic enrichment factor of $\epsilon_{\text{CH}_4, \text{ac-methyl}} = -15\%$,
 226 which is in-between the $\epsilon_{\text{CH}_4, \text{ac-methyl}}$ of acetivlastic *Methanosaeta* (Penning et al., 2006; Valentine et al., 2004) and
 227 *Methanosarcina* species (Gelwicks et al., 1994; Govert and Conrad, 2009), the average f_{H_2} values were 0% for
 228 Vercelli soil and 20% for IRRI soil (Fig. S3c).
 229



230
 231
 232 **Figure 4:** Mariotti plots of propionate consumption under methanogenic and sulfidogenic ($\pm \text{CH}_3\text{F}$) conditions in
 233 paddy soil from (a) Vercelli and (b) the IRRI. The different symbols indicate three different replicates; the lines
 234 give the results of linear regression averaged over the replicates.
 235



236
 237 **Figure 5:** Isotopic enrichment factors (ϵ_{prop} , given as negative values) in paddy soils without and with addition of
 238 sulfate (gypsum) and CH_3F . Means \pm SE. The differences between the incubations were examined using Hukey's
 239 post hoc test of a one-way analysis of variance (ANOVA). Different letter son top of bars indicate significant
 240 difference ($P < 0.05$) between the data.

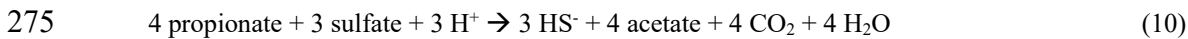
241
 242 **4 Discussion**

243 *Pathway of propionate degradation*

244 Our results showed that propionate was degraded via acetate as main transient intermediate finally resulting in
 245 the production of CH_4 and CO_2 under methanogenic and CO_2 under sulfidogenic conditions. These results are
 246 consistent with previous observations by Liu and Conrad (Liu and Conrad, 2017) using the same paddy soils.
 247 Stable isotope probing and correlation network analysis of the microbial communities have shown that propionate
 248 degradation is accomplished by both *Syntrophobacter* and *Smithella* species (Gan et al., 2012; Liu and Conrad,
 249 2017; Lueders et al., 2004). The present study showed that propionate degradation under methanogenic conditions
 250 was consistent with the major operation of the *Smithella* pathway. The main argument for this conclusion is the
 251 observation that butyrate was a transient intermediate of propionate degradation, albeit at low concentrations (Fig.
 252 1, S1). In the *Smithella* pathway butyrate is further fermented to acetate and H_2 . However, production of H_2 is
 253 smaller in the *Smithella* than in the *Syntrophobacter* pathway, while production of acetate is larger. Indeed,
 254 aceticlastic methanogenesis explained all the propionate-driven methanogenesis in the paddy soils (Fig. 2a, S2a).
 255 The average hydrogenotrophic methanogenesis by contrast contributed almost zero in Vercelli soil and only about
 256 20% in IRRI soil (Fig. S3c). The relatively larger contribution of aceticlastic than hydrogenotrophic
 257 methanogenesis to methanogenic propionate degradation supports the conclusion that the *Smithella* pathway was
 258 dominating over the *Syntrophobacter* pathway. Arguments against the *Smithella* pathway are that the accumulated
 259 CH_4 amounted to less than the expected 1.75 mole per mole propionate consumed in Vercelli soil (Fig. 2a) and
 260 even less in IRRI soil (Fig. S2a). With inhibition of aceticlastic methanogenesis, acetate accumulation in Vercelli
 261 soil accounted for about 1 mole acetate per mole propionate, being in a range that is compatible with propionate
 262 fermentation by either *Smithella* or *Syntrophobacter* (Fig. 2b). In IRRI soil however, acetate accumulation

263 accounted for less than 1 mole acetate per mole propionate (Fig. S2b). Note, however, that the accumulation of
264 acetate reflects only that part of propionate fermentation, which was not inhibited by CH₃F. Our conclusion that
265 propionate was degraded mainly by *Smithella* under methanogenic conditions is consistent with the microbial
266 community structure in the paddy soils from Vercelli and IRRI, which contains not only *Syntrophobacter* species
267 but also *Smithella* together with *Syntrophomonas*, which is able to ferment butyrate (Liu and Conrad, 2017).

268 Under sulfidogenic conditions, propionate can be oxidized in different ways, either directly by sulfate reducers
269 forming acetate and CO₂, or syntrophically as under methanogenic conditions, but with subsequent oxidation of
270 H₂ and acetate by sulfate reducers. Using the same paddy soils, Liu and coworkers (Liu et al., 2018a; Liu and
271 Conrad, 2017) recently showed that under sulfidogenic conditions propionate consumption was mainly achieved
272 by *Syntrophobacter* spp., which first oxidized propionate to acetate and CO₂, and subsequently oxidized the
273 accumulated acetate to CO₂. These were exactly the processes observed in the present study, where propionate
274 degradation initially resulted in almost equimolar accumulation of acetate (Fig. 2d) according to



276 It was interesting, that CH₃F was not only a strong inhibitor of aceticlastic methanogenesis (which was
277 expected), but also a relatively strong inhibitor of propionate fermentation, but only under methanogenic but not
278 under sulfidogenic conditions. Inhibition of propionate fermentation under methanogenic conditions has been
279 observed before in three different paddy soils and has been interpreted as being due to the adverse thermodynamic
280 conditions when acetate accumulates (Conrad et al., 2014). However, this interpretation cannot be true, since
281 accumulation of acetate also occurred under sulfidogenic conditions, where CH₃F did not inhibit propionate
282 degradation. In fact it is mainly the accumulation of H₂ rather than acetate, to which propionate degradation is
283 thermodynamically sensitive. This is the reason why the *Smithella* pathway is less sensitive to thermodynamic
284 inhibition than the *Syntrophobacter* pathway (Dolfing, 2013). However, CH₃F did not inhibit H₂ consumption by
285 methanogens, as seen by the low δ¹³C of CH₄ in the presence of CH₃F. Furthermore, the first step of the *Smithella*-
286 type propionate fermentation does not produce any H₂ and therefore, propionate in the presence of CH₃F should at
287 least be fermented to butyrate and acetate, which however, was not the case. Hence, the reason why CH₃F inhibited
288 propionate fermentation under methanogenic but not under sulfidogenic conditions remains unknown. Perhaps it
289 is *Smithella* being more sensitive to CH₃F than *Syntrophobacter*.

290

291 *Fractionation during propionate degradation*

292 The isotopic fractionation of propionate apparently followed Raleigh distillation that is characteristic for kinetic
293 isotope fractionation in a closed system. The isotopic enrichment factor, which was determined from Mariotti plots,
294 was in the range of ε_{prop} = -8‰ to -3.5‰, which is less than the enrichment factor for methanogenic acetate
295 consumption, which has been found to be ε_{ac} = -21‰ to -17‰ (Conrad et al., 2021). The ε_{prop} values are on the
296 same order as those predicted from δ¹³C values of propionate, acetate and organic carbon measured in various
297 methanogenic soils and sediments (Conrad et al., 2014). Propionate degradation resulted in the formation of ¹³C-
298 enriched acetate and CO₂ and ¹³C-depleted butyrate and CH₄. The formation of ¹³C-depleted butyrate can be
299 explained by kinetic isotope effect with the preferential utilization of ¹³C-depleted propionate in the initial
300 dismutation reaction by *Smithella*. However, the production of ¹³C-enriched acetate cannot be explained by a linear
301 kinetic isotope effect. We assume that the dismutation of propionate is a branch point (Fry, 2003; Hayes, 2001), at

302 which the carbon flow is split into the production of ^{13}C -enriched acetate and ^{13}C -depleted butyrate. At the branch
303 point the carbon isotope flow shows a preferential flow of ^{12}C into the product generated by the reaction with the
304 larger fractionation factor, which would be butyrate. The further conversion of butyrate should produce acetate
305 that is depleted in ^{13}C . This acetate together with the acetate produced from propionate dismutation should result
306 in the $\delta^{13}\text{C}$ -acetate that is observed. The total acetate pool initially had a $\delta^{13}\text{C}$ that was up to 10‰ heavier than the
307 $\delta^{13}\text{C}$ of propionate. In the end, the $\delta^{13}\text{C}$ values were about equal. The observation that acetate was ^{13}C -enriched
308 relative to propionate is consistent with $\delta^{13}\text{C}$ data in various soils and sediments (Conrad et al., 2014) reporting
309 that acetate is on the average enriched by 6‰ relative to propionate. Acetate was further converted to CH_4 and to
310 CO_2 . In Vercelli soil, the $\delta^{13}\text{C}$ of CH_4 was about 25-35‰ lighter than the $\delta^{13}\text{C}$ of acetate. In IRRI soil, ^{13}C depletion
311 was even larger (30-40‰). In both soils, the isotopic enrichment factors for acetate consumption were in a range
312 of -12‰ to -17‰ and for CH_4 production from acetate in a range of -37‰ to -27‰ (Conrad et al., 2021).
313 Considering that a certain percentage (albeit small) of CH_4 was formed from CO_2 reduction by hydrogenotrophic
314 methanogenesis, which displays relatively negative enrichment factors (see the $\delta^{13}\text{C}$ of CH_4 in the presence of
315 CH_3F , Fig. 1g), the observed difference in $\delta^{13}\text{C}$ of CH_4 versus acetate is reasonable. In *Smithella* fermentation, the
316 only CO_2 production occurs during the fermentation of butyrate and the aceticlastic conversion of acetate. In both
317 cases CO_2 should be ^{13}C -depleted relative to the substrates. Note, that this was not the case. Unfortunately, the ^{13}C
318 contents of the individual C atoms of propionate, butyrate and acetate are not known. The ^{13}C content in the
319 different C positions might also affect the $\delta^{13}\text{C}$ of CH_4 and CO_2 , which are formed. It is also possible that besides
320 *Smithella* fermentation, the *Syntrophobacter* fermentation contributed to propionate degradation. In summary, the
321 detailed process of isotope fractionation during the pathway of propionate degradation is unclear. However, the
322 magnitude of the enrichment factors involved was relatively small, being on the order of <10‰.

323 Under sulfidogenic conditions, propionate was most probably degraded by *Syntrophobacter* spp., first to
324 acetate, then finally to CO_2 (Liu et al., 2018a; Liu and Conrad, 2017). The carbon isotope fractionation of
325 propionate consumption was with an enrichment factor of $\epsilon_{\text{prop}} = -3.5\text{‰}$ comparatively small. Propionate was
326 eventually converted to two carbon products of which one was depleted (the CO_2) and the other was enriched (the
327 acetate) in ^{13}C . In case of *Syntrophobacter*-type degradation, acetate and CO_2 are produced from the conversion of
328 pyruvate, which is generated in the methylmalonyl-CoA pathway. In this pathway, CO_2 is first consumed by the
329 conversion of propionyl-CoA to methylmalonyl-CoA and then produced by the conversion of oxaloacetate to
330 pyruvate. Pyruvate is finally converted to acetate and CO_2 , which should both be ^{13}C -depleted with respect to
331 pyruvate (DeNiro and Epstein, 1977). However, both acetate and CO_2 were initially ^{13}C -enriched relative to
332 propionate (about 2-5‰), and then changed in opposite directions with acetate becoming increasingly ^{13}C -enriched
333 and CO_2 becoming increasingly ^{13}C -depleted until the time, when acetate accumulation had reached a maximum
334 (Fig. 5). Then, $\delta^{13}\text{C}$ of both acetate and CO_2 increased together with the increase of ^{13}C of propionate (Fig. 5).
335 Increase of $\delta^{13}\text{C}$ of acetate is often explained by consumption, especially through aceticlastic methanogenesis
336 (Heuer et al., 2010; Heuer et al., 2009). However, hardly any CH_4 was produced under sulfidogenic conditions and
337 the ^{13}C enrichment occurred during the phase of acetate accumulation. Therefore, the enrichment likely happened
338 during acetate production from propionate degradation. The increasing ^{13}C -depletion of CO_2 can also not be
339 explained by consumption but only by the production from propionate. Hence, isotope fractionation during the
340 conversion of propionate, in particular during the conversion of propionate to pyruvate is unclear. We assume

341 complications during the carboxylation and decarboxylation reactions. Unfortunately, we hardly found any
342 literature data on the isotope fractionation of propionate fermentation. A coculture of *Syntrophobacter*
343 *fumaroxidans* with *Methanobacterium formicicum* exhibited marginal propionate fractionation with $\epsilon_{\text{prop}} = 0.9\text{‰}$
344 and the formation of acetate, that was slightly ^{13}C -enriched (about 5‰) (Botsch and Conrad, 2011), similarly as
345 observed here. In summary, the mechanism of isotope fractionation during the conversion of propionate is not
346 completely clear, but the magnitude of isotope fractionation is quite low.

347

348 **5 Conclusions**

349 Propionate degradation under sulfidogenic conditions was explained by the metabolism of *Syntrophobacteraceae*,
350 which in a first step converted propionate to ^{13}C -enriched acetate and ^{13}C -depleted CO_2 . By contrast, propionate
351 degradation under methanogenic conditions was at least partially due to metabolism by *Smithella*, which in a first
352 step converted propionate to ^{13}C -enriched acetate and ^{13}C -depleted butyrate. However, the isotopic enrichment
353 factors (ϵ_{prop}) of propionate consumption in two paddy soils were generally very low (-8‰ to -3.5‰) both under
354 methanogenic and sulfidogenic conditions. This low range is consistent with literature values of $\delta^{13}\text{C}$, collected
355 for propionate, acetate and organic carbon in various soils and sediments (Conrad et al., 2014). Fractionation of
356 propionate carbon actually seems to be smaller than fractionation of acetate, which is at least two times larger
357 (Conrad et al., 2021). Hence, degradation of organic carbon via propionate to acetate and CO_2 apparently involves
358 only little isotope fractionation being on the order of <10‰. By contrast, further degradation of acetate and CO_2
359 (+ H_2) to CH_4 involves substantial isotope fractionation. This is also the case for chemolithotrophic acetate
360 production (Conrad et al., 2014).

361

362 **Supplement link**

363

364 **Author contribution:** RC designed the experiments, evaluated the data and wrote the manuscript, PC conducted
365 the experiments.

366

367 **Competing interests:** The authors declare that they have no conflict of interests.

368

369 **Acknowledgements**

370 We thank the Fonds der Chemischen Industrie for financial support.

371

372 **References**

373

374 Boone, D. R. and Bryant, M. P.: Propionate-degrading bacterium, *Syntrophobacter wolinii* sp. nov.
375 gen. nov., from methanogenic ecosystems, Appl. Environ. Microbiol., 40, 626-632, 1980.

376 Botsch, K. C. and Conrad, R.: Fractionation of stable carbon isotopes during anaerobic production
377 and degradation of propionate in defined microbial cultures, Org. Geochem., 42, 289-295, 2011.

- 378 Chen, S. Y., Liu, X. L., and Dong, X. Z.: *Syntrophobacter sulfatireducens* sp. nov., a novel
379 syntrophic, propionate-oxidizing bacterium isolated from UASB reactors, *Int. J. Syst. Evol.*
380 *Microbiol.*, 55, 1319-1324, 2005.
- 381 Conrad, R.: Quantification of methanogenic pathways using stable carbon isotopic signatures: a
382 review and a proposal, *Org. Geochem.*, 36, 739-752, 2005.
- 383 Conrad, R., Claus, P., and Casper, P.: Stable isotope fractionation during the methanogenic
384 degradation of organic matter in the sediment of an acidic bog lake, Lake Grosse Fuchskuhle,
385 *Limnol. Oceanogr.*, 55, 1932-1942, 2010.
- 386 Conrad, R., Claus, P., Chidthaisong, A., Lu, Y., Scavino, A., Liu, Y., Angel, R., Galand, P., Casper,
387 P., Guerin, F., and Enrich-Prast, A.: Stable carbon isotope biogeochemistry of propionate and
388 acetate in methanogenic soils and lake sediments, *Org. Geochem.*, 73, 1-7, 2014.
- 389 Conrad, R., Liu, P., and Claus, P.: Fractionation of stable carbon isotopes during acetate
390 consumption by methanogenic and sulfidogenic microbial communities in rice paddy soils and
391 lake sediments, *Biogeosciences*, 18, 6533-6546, 2021.
- 392 DeBok, F. A. M., Harmsen, H. J. M., Plugge, C. M., DeVries, M. C., Akkermans, A. D. L., DeVos,
393 W. M., and Stams, A. J. M.: The first true obligately syntrophic propionate-oxidizing bacterium,
394 *Pelotomaculum schinkii* sp. nov., co-cultured with *Methanospirillum hungatei*, and emended
395 description of the genus *Pelotomaculum*, *Int. J. Syst. Evol. Microbiol.*, 55, 1697-1703, 2005.
- 396 DeBok, F. A. M., Stams, A. J. M., Dijkema, C., and Boone, D. R.: Pathway of propionate oxidation
397 by a syntrophic culture of *Smithella propionica* and *Methanospirillum hungatei*, *Appl. Environ.*
398 *Microbiol.*, 67, 1800-1804, 2001.
- 399 DeNiro, M. J. and Epstein, S.: Mechanism of carbon isotope fractionation associated with lipid
400 synthesis, *Science*, 197, 261-263, 1977.
- 401 Dolfing, J.: Syntrophic propionate oxidation via butyrate: a novel window of opportunity under
402 methanogenic conditions, *Appl. Environ. Microbiol.*, 79, 4515-4516, 2013.
- 403 Elsner, M., Zwank, L., Hunkeler, D., and Schwarzenbach, R. P.: A new concept linking observable
404 stable isotope fractionation to transformation pathways of organic pollutants [review], *Environ.*
405 *Sci. Technol.*, 39, 6896-6916, 2005.
- 406 Fry, B.: Steady state models of stable isotopic distributions, *Isotopes Environ. Health Studies*, 39,
407 219-232, 2003.
- 408 Gan, Y., Qiu, Q., Liu, P., Rui, J., and Lu, Y.: Syntrophic oxidation of propionate in rice field soil at
409 15 and 30°C under methanogenic conditions, *Appl. Environ. Microbiol.*, 78, 4923-4932, 2012.

- 410 Gelwicks, J. T., Risatti, J. B., and Hayes, J. M.: Carbon isotope effects associated with acetoclastic
411 methanogenesis, *Appl. Environ. Microbiol.*, 60, 467-472, 1994.
- 412 Glissmann, K. and Conrad, R.: Fermentation pattern of methanogenic degradation of rice straw in
413 anoxic paddy soil, *FEMS Microbiol. Ecol.*, 31, 117-126, 2000.
- 414 Goevert, D. and Conrad, R.: Effect of substrate concentration on carbon isotope fractionation during
415 acetoclastic methanogenesis by *Methanosarcina barkeri* and *M. acetivorans* and in rice field soil,
416 *Appl. Environ. Microbiol.*, 75, 2605-2612, 2009.
- 417 Hayes, J. M.: Factors controlling ^{13}C contents of sedimentary organic compounds: principles and
418 evidence, *Mar. Geol.*, 113, 111-125, 1993.
- 419 Hayes, J. M.: Fractionation of carbon and hydrogen isotopes in biosynthetic processes, *Stable*
420 *Isotope Geochemistry*, 43, 225-277, 2001.
- 421 Heuer, V. B., Krüger, M., Elvert, M., and Hinrichs, K. U.: Experimental studies on the stable carbon
422 isotope biogeochemistry of acetate in lake sediments, *Org. Geochem.*, 41, 22-30, 2010.
- 423 Heuer, V. B., Pohlman, J. W., Torres, M. E., Elvert, M., and Hinrichs, K. U.: The stable carbon
424 isotope biogeochemistry of acetate and other dissolved carbon species in deep seafloor
425 sediments at the northern Cascadia Margin, *Geochim. Cosmochim. Acta*, 73, 3323-3336, 2009.
- 426 Houwen, F. P., Dijkema, C., Stams, A. J. M., and Zehnder, A. J. B.: Propionate metabolism in
427 anaerobic bacteria - determination of carboxylation reactions with ^{13}C -NMR spectroscopy,
428 *Biochim. Biophys. Acta*, 1056, 126-132, 1991.
- 429 Imachi, H., Sekiguchi, Y., Kamagata, Y., Hanada, S., Ohashi, A., and Harada, H.: *Pelotomaculum*
430 *thermopropionicum* gen. nov., sp. nov., an anaerobic, thermophilic, syntrophic propionate-
431 oxidizing bacterium, *Int. J. Syst. Evol. Microbiol.*, 52, 1729-1735, 2002.
- 432 Imachi, H., Sekiguchi, Y., Kamagata, Y., Loy, A., Qiu, Y. L., Hugenholtz, P., Kimura, N., Wagner,
433 M., Ohashi, A., and Harada, H.: Non-sulfate-reducing, syntrophic bacteria affiliated with
434 *Desulfotomaculum* cluster I are widely distributed in methanogenic environments, *Appl.*
435 *Environ. Microbiol.*, 72, 2080-2091, 2006.
- 436 Janssen, P. H. and Frenzel, P.: Inhibition of methanogenesis by methyl fluoride - studies of pure and
437 defined mixed cultures of anaerobic bacteria and archaea, *Appl. Environ. Microbiol.*, 63, 4552-
438 4557, 1997.
- 439 Krylova, N. I. and Conrad, R.: Thermodynamics of propionate degradation in methanogenic paddy
440 soil, *FEMS Microbiol. Ecol.*, 26, 281-288, 1998.

- 441 Krylova, N. I., Janssen, P. H., and Conrad, R.: Turnover of propionate in methanogenic paddy soil,
442 FEMS Microbiol. Ecol., 23, 107-117, 1997.
- 443 Liu, P., Pommerenke, B., and Conrad, R.: Identification of *Syntrophobacteraceae* as major acetate-
444 degrading sulfate reducing bacteria in Italian paddy soil, Environ. Microbiol., 20, 337-354,
445 2018a.
- 446 Liu, P. F. and Conrad, R.: *Syntrophobacteraceae*-affiliated species are major propionate-degrading
447 sulfate reducers in paddy soil, Environ. Microbiol., 19, 1669-1686, 2017.
- 448 Liu, P. F., Klose, M., and Conrad, R.: Temperature effects on structure and function of the
449 methanogenic microbial communities in two paddy soils and one desert soil, Soil Biol. Biochem.,
450 124, 236-244, 2018b.
- 451 Liu, Y. T., Balkwill, D. L., Aldrich, H. C., Drake, G. R., and Boone, D. R.: Characterization of the
452 anaerobic propionate-degrading syntrophs *Smithella propionica* gen. nov., sp. nov. and
453 *Syntrophobacter wolinii*, Int. J. Syst. Bacteriol., 49, 545-556, 1999.
- 454 Lueders, T., Pommerenke, B., and Friedrich, M. W.: Stable-isotope probing of microorganisms
455 thriving at thermodynamic limits: Syntrophic propionate oxidation in flooded soil, Appl.
456 Environ. Microbiol., 70, 5778-5786, 2004.
- 457 Mariotti, A., Germon, J. C., Hubert, P., Kaiser, P., Letolle, R., Tardieux, A., and Tardieux, P.:
458 Experimental determination of nitrogen kinetic isotope fractionation: some principles;
459 illustration for the denitrification and nitrification processes, Plant and Soil, 62, 413-430, 1981.
- 460 McInerney, M. J., Bryant, M. P., Hespell, R. B., and Costerton, J. W.: *Syntrophomonas wolfei* gen.
461 nov. sp. nov., an anaerobic, syntrophic, fatty acid-oxidizing bacterium, Appl. Environ.
462 Microbiol., 41, 1029-1039, 1981.
- 463 Nozoe, T.: Effects of methanogenesis and sulfate-reduction on acetogenic oxidation of propionate
464 and further decomposition of acetate in paddy soil, Soil Sci. Plant Nutr., 43, 1-10, 1997.
- 465 Penning, H., Claus, P., Casper, P., and Conrad, R.: Carbon isotope fractionation during acetoclastic
466 methanogenesis by *Methanosaeta concilii* in culture and a lake sediment, Appl. Environ.
467 Microbiol., 72, 5648-5652, 2006.
- 468 Plugge, C. M., Balk, M., and Stams, A. J. M.: *Desulfotomaculum thermobenzoicum* subsp.
469 *thermosyntrophicum* subsp. nov., a thermophilic, syntrophic, propionate-oxidizing, spore-
470 forming bacterium, Int. J. Syst. Evol. Microbiol., 52, 391-399, 2002.
- 471 Schink, B.: Mechanisms and kinetics of succinate and propionate degradation in anoxic freshwater
472 sediments and sewage sludge, J. Gen. Microbiol., 131, 643-650, 1985.

- 473 Textor, S., Wendisch, V. F., DeGraaf, A., Mueller, U., Linder, M. I., Linder, D., and Buckel, W.:
474 Propionate oxidation in *Escherichia coli* - evidence for operation of a methylcitrate cycle in
475 bacteria, Arch. Microbiol., 168, 428-436, 1997.
- 476 Valentine, D. L., Chidthaisong, A., Rice, A., Reeburgh, W. S., and Tyler, S. C.: Carbon and hydrogen
477 isotope fractionation by moderately thermophilic methanogens, Geochim. Cosmochim. Acta, 68,
478 1571-1590, 2004.
- 479 Xia, X. X., Zhang, J. C., Song, T. Z., and Lu, Y. H.: Stimulation of *Smithella*-dominated propionate
480 oxidation in a sediment enrichment by magnetite and carbon nanotubes, Environ. Microbiol.
481 Reports, 11, 236-248, 2019.
- 482 Yao, H. and Conrad, R.: Thermodynamics of propionate degradation in anoxic paddy soil from
483 different rice-growing regions, Soil Biol. Biochem., 33, 359-364, 2001.
484
485