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

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10 Abstract. Propionate is an important intermediate during the breakdown of organic matter in anoxic flooded paddy 11 soils. Since there are only few experiments on carbon isotope fractionation and the magnitude of the isotopic 12 enrichment factors (ɛ) involved, we measured propionate conversion to acetate, CH4 and CO2 in anoxic paddy 13 soils. Propionate consumption was measured using samples of paddy soil from Vercelli (Italy) and the International 14 Rice Research Institute (IRRI, the Philippines) suspended in phosphate buffer (pH 7.0), both in the absence and 15 presence of sulfate (gypsum), and of methyl fluoride (CH<sub>3</sub>F), an inhibitor of aceticlastic methanogenesis. Under 16 methanogenic conditions, propionate was eventually degraded to CH4 with acetate being a transient intermediate. 17 Butyrate was also a minor intermediate. Methane was mainly produced by aceticlastic methanogenesis. Propionate 18 consumption was inhibited by CH<sub>3</sub>F. Whereas butyrate and CH<sub>4</sub> were <sup>13</sup>C-depleted relative to propionate, acetate 19 and CO<sub>2</sub> were <sup>13</sup>C-enriched. The isotopic enrichment factors ( $\varepsilon_{prop}$ ) of propionate consumption, determined by 20 Mariotti plots, were in a range of -8‰ to -3.5‰. Under sulfidogenic conditions, acetate was also transiently 21 accumulated, but CH<sub>4</sub> production was negligible. Application of CH<sub>3</sub>F hardly affected propionate degradation and 22 acetate accumulation. The initially produced CO2 was <sup>13</sup>C-depleted, whereas the acetate was <sup>13</sup>C-enriched. The 23 values of  $\varepsilon_{prop}$  were -3.5%. It is concluded that degradation of organic carbon via propionate to acetate and CO<sub>2</sub> 24 involves only little isotope fractionation. The results further indicate a major contribution of Syntrophobacter-type 25 propionate fermentation under sulfidogenic conditions and Smithella-type propionate fermentation under 26 methanogenic conditions. This interpretation is consistent with data of the microbial community composition 27 published previously for the same soils.

**Field Code Changed** 

# 28 1 Introduction

20	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 milimolar 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:
42	4 propionate + 8 $H_2O \rightarrow$ 4 acetate + 4 $CO_2$ + 12 $H_2$ (1)
43	Propionate degradation by non-randomizing Smithella proceeds by dismutation of propionate:
44	4 propionate $\rightarrow$ 2 butyrate + 2 acetate (2)
45	Butyrate is then syntrophically converted (e.g., by Syntrophomonas (McInerney et al., 1981)):
46	2 butyrate + 4 H <sub>2</sub> O $\rightarrow$ 4 acetate + 4 H <sub>2</sub> (3)
47	The Smithella pathway in total:
48	4 propionate + 4 $H_2O \rightarrow 6$ acetate + 4 $H_2$ (4)
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:
55	4 propionate + 2 H <sub>2</sub> O $\rightarrow$ 7 CH <sub>4</sub> + 5 CO <sub>2</sub> , or (5)
56	4 propionate + 7 sulfate + 11 H <sup>+</sup> $\rightarrow$ 7 HS <sup>-</sup> + 12 CO <sub>2</sub> + 12 H <sub>2</sub> O (6)
57	Note, that the relative production of acetate and H <sub>2</sub> is different for Syntrophobacter and Smithella fermentation,
58	being 1:3 and 3:2, respectively. Therefore, aceticlastic 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 CO<sub>2</sub>, and 66 subsequently oxidize the accumulated acetate to CO<sub>2</sub>. 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  $\delta^{13}$ 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  $(\varepsilon)$  of the processes involved in both production and consumption of 75 propionate are probably small. The direct determination of  $\varepsilon$  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  $\varepsilon$  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<sub>4</sub> and CO<sub>2</sub>. We also used the treatment with methyl fluoride (CH<sub>3</sub>F) 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 13C 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.

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#### 85 2 Materials and Methods

# 86 2.1 Paddy soils and incubation conditions

The soil samples were from the research stations in Vercelli, Italy and the International Rice research Institute
(IRRI) in the Philippines. Sampling and soil characteristics were described before (Liu et al., 2018b). <u>The main</u>
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
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<sub>2</sub>O at a ratio of 1:1 and incubated under N<sub>2</sub> at 25°C for 93 4 weeks. In a second incubation, paddy soil was mixed with autoclaved anoxic H<sub>2</sub>O (prepared under N<sub>2</sub>) at a ratio 94 of 1:1, was amended with 0.07 g CaSO<sub>4</sub>.2H<sub>2</sub>O, and then incubated under N<sub>2</sub> 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 mlmL 98 soil slurry preincubated without sulfate was incubated at 25°C with 40 mlmL 20 mM potassium phosphate buffer 99 (pH 7.0) in a 150-mlmL bottle under an atmosphere of N2. The bottles were the amended with (i) 5 mlmL H2O; 100 (ii) 5 mlmL H<sub>2</sub>O + 4.5 mlmL CH<sub>3</sub>F; (iii) 5 mlmL 50 mM sodium propionate; (iv) 5 mlmL 50 mM sodium acetate 101 + 4.5 mlmL CH<sub>3</sub>F. In the second set (resulting in sulfidogenic conditions), 5 mlmL soil slurry preincubated with 102 sulfate was incubated at 25°C with 40 mlmL 20 mM potassium phosphate buffer (pH 7.0) in a 150-mlmL bottle 103 under an atmosphere of N<sub>2</sub>. The amendments were the same as above, but with the addition of 200  $\mu$ l of a CaSO<sub>4</sub> 104 suspension corresponding to a concentration of 2.5 M (giving a final concentration of 10 mM sulfate).

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106 *2.2 Chemical and isotopic analyses* 

107	Gas samples for analysis of partial pressures of CH <sub>4</sub> and CO <sub>2</sub> were taken from the headspace of the incubation $\int_{-\infty}^{+}$	1
108	bottles after vigorous manual shaking for about 30 s using a gas-tight pressure-lock syringe, which had been	A.
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 ${}^{13}C/{}^{12}C$ 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 ( $\delta^{13}C$ ) relative to the Vienna Peedee Belemnite standard having a $^{13}C/^{12}C$ ratio ( $R_{standard}$ ) of 0.01118:	il.
117	$\delta^{13}C=10^3~(R_{sample}/R_{standard}-1).~The~precision~of~the~GC-C-IRMS~was~\pm~0.2\%,~that~of~the~HPLC-IRMS~was~\pm~0.2\%,~that~0.2\%,~t$	۱ <u>ا</u>
118	0.3‰.	l
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120	2.3 Calculations	
121	Millimolar concentrations of CH <sub>4</sub> were calculated from the mixing ratios (1 ppmy = $10^{-6}$ bar) measured in the	

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Millimolar concentrations of CH<sub>4</sub> were calculated from the mixing ratios (1 ppmv =  $10^{-6}$  bar) measured in the gas phase of the incubation bottles: 1000 ppmv CH<sub>4</sub> correspond to 0.09 µmol per mlmL of liquid. Note, that this is the total amount of CH<sub>4</sub> in the gas phase relative to the liquid phase.

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

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

126 also expressed as  $\varepsilon \equiv 1000 (1 - \alpha)$  in permil. The carbon isotope enrichment factor  $\varepsilon_{prop}$  associated with propionate

(7)

(8)

(9)

127 consumption was calculated from the temporal change of  $\delta^{13}$ C of propionate as described by Mariotti et al. 128 (Mariotti et al., 1981) from the residual reactant

129  $\delta_{\rm r} = \delta_{\rm ri} + \varepsilon \left[ \ln(1-f) \right]$ 

130

where  $\delta_{ri}$  is the isotopic composition of the reactant (propionate) at the beginning, and  $\delta_r$  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  $(0 \le f_{prop} \le 1)$ . Linear regression of  $\delta^{13}$ C of propionate against  $\ln(1 - f)$  yields

133  $\epsilon_{prop}$  as the slope of best fit lines. The regressions of  $\delta^{13}$ C of propionate were done for data in the range of  $f_{prop} < \delta^{13}$ C of propionate were done for data in the range of  $f_{prop} < \delta^{13}$ C of propionate were done for data in the range of  $f_{prop} < \delta^{13}$ C of propionate were done for data in the range of  $f_{prop} < \delta^{13}$ C of propionate were done for data in the range of  $f_{prop} < \delta^{13}$ C of propionate were done for data in the range of  $f_{prop} < \delta^{13}$ C of propionate were done for data in the range of  $f_{prop} < \delta^{13}$ C of propionate were done for data in the range of  $f_{prop} < \delta^{13}$ C of propionate were done for data in the range of  $f_{prop} < \delta^{13}$ C of propionate were done for data in the range of  $f_{prop} < \delta^{13}$ C of propionate were done for data in the range of  $f_{prop} < \delta^{13}$ C of propionate were done for data in the range of  $f_{prop} < \delta^{13}$ C of propionate were done for data in the range of  $f_{prop} < \delta^{13}$ C of propionate were done for data in the range of  $f_{prop} < \delta^{13}$ C of propionate were done for data in the range of  $f_{prop} < \delta^{13}$ C of propionate were done for data in the range of  $f_{prop} < \delta^{13}$ C of propionate were done for data in the range of  $f_{prop} < \delta^{13}$ C of propionate were done for data in the range of  $f_{prop} < \delta^{13}$ C of propionate were done for data in the range of  $f_{prop} < \delta^{13}$ C of propionate were done for data in the range of  $f_{prop} < \delta^{13}$ C of propionate were done for data in the range of  $f_{prop} < \delta^{13}$ C of propionate were done for data in the range of  $f_{prop} < \delta^{13}$ C of propionate were done for data in the range of  $f_{prop} < \delta^{13}$ C of propionate were done for data in the range of  $f_{prop} < \delta^{13}$ C of propionate were done for data in the range of  $f_{prop} < \delta^{13}$ C of propionate were done for data in the range of  $f_{prop} < \delta^{13}$ C of propionate were done for data in the range of  $f_{prop} < \delta^{13}$ C of propionate were done for data in the range of  $f_{prop} < \delta^{13}$ C of propio

134 0.7. The linear regressions were done individually for each experimental replicate (n = 3) and were only accepted

135 if  $r^2 > 0.9$ . The  $\epsilon$  values resulting from the replicate experiments were then averaged ( $\pm$  SE).

136The fraction ( $f_{H2}$ ) of CH4 derived from hydrogenotrophic methanogenesis was determined as described before137(Conrad et al., 2010) using

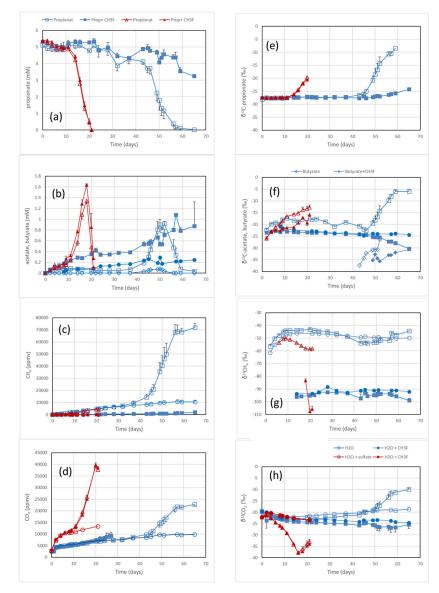
138  $f_{H2} = (\delta^{13}C_{CH4} - \delta^{13}C_{CH4-ma})/(\delta^{13}C_{CH4-mc} - \delta^{13}C_{CH4-ma})$ 

139 with  $\delta^{13}C_{CH4} = \delta^{13}C$  of total CH<sub>4</sub> produced,  $\delta^{13}C_{CH4-mc} = \delta^{13}C$  of CH<sub>4</sub> produced from hydrogenotrophic 140 methanogenesis, which is equivalent to the CH<sub>4</sub> produced in the presence of CH<sub>3</sub>F, and  $\delta^{13}C_{CH4-ma} = \delta^{13}C$  of CH<sub>4</sub>

141 produced from aceticlastic methanogenesis. The  $\delta^{13}C_{CH4-ma}$  was approximated from the  $\delta^{13}C$  of acetate in the

presence of CH<sub>3</sub>F assuming that the methyl group of acetate was depleted in <sup>13</sup>C by 8‰ (Conrad et al., 2014) and

143 that the enrichment factor ( $\varepsilon_{CH4,ac-methyl}$ ) for CH<sub>4</sub> being produced from acetate-methyl was between 0 and -20%.



147Figure 1: Propionate conversion to acetate, butyrate, CH4 and CO2 in suspensions of paddy soil from Vercelli148(Italy) after addition of propionate without sulfate (blue squares) or propionate plus sulfate (gypsum) (red triangles)149without CH3F (open symbols) or with CH3F (closed symbols). Controls with addition of only water (blue or red150circles) are only shown occasionally. The panels show the temporal change of (a) concentrations of propionate,151(b) concentrations of acetate and butyrate (blue diamonds), (c) mixing ratios of CH4 (1 ppmv = 10<sup>-6</sup> bar), (d) mixing152ratios of CO2, (e)  $\delta^{13}$ C of propionate, (f)  $\delta^{13}$ C of acetate and butyrate, (g)  $\delta^{13}$ C of CH4, and (h)  $\delta^{13}$ C of CO2. Means153± SE.

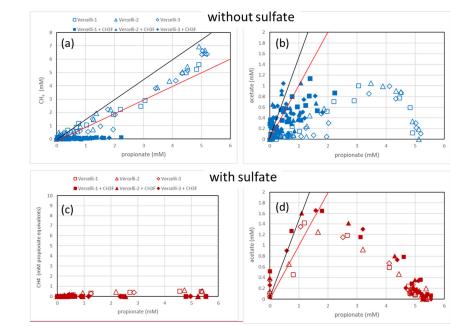
# 155 3 Results

156 3.1 Conversion of propionate under methanogenic and sulfidogenic conditions

157 Incubation of buffered suspensions of rice field soil from Vercelli (Fig. 1) and the IRRI (Fig. S1) resulted in 158 similar patterns of propionate degradation to acetate, CH4 and CO2. Under methanogenic conditions in the absence 159 of sulfate, propionate degradation started after a lag phase of about 20 d (Fig. 1a) resulting in the production of 160 acetate (Fig. 1b), CH4 (Fig. 1c) and CO2 (Fig. 1d). The formation of acetate, CH4 and CO2 in the absence of 161 propionate was only very small. The accumulation of acetate was only transient, except when aceticlastic 162 methanogenesis was inhibited by CH<sub>3</sub>F (Fig. 1b). Similar observations were made in IRRI soil (Fig. S1a-d). The 163 production of CH4 was roughly equimolar to the consumption of propionate, but was nearly zero when aceticlastic 164 methanogenesis was inhibited by CH3F (Fig. 2a). Under these conditions, acetate accumulated to nearly equimolar 165 amounts with the consumed propionate (Fig. 2b), but in IRRI soil acetate accumulation was less than equimolar 166 (Fig. S2b). Butyrate was also a transient intermediate of propionate degradation and was produced and consumed 167 simultaneously with acetate (Fig. 1b, S1b). However, the accumulated concentrations were small (<0.1 mM).

168In the presence of sulfate, propionate degradation started after a lag phase of only about 10 days (Fig. 1a)169resulting in the accumulation of acetate (Fig. 1b) and the production of  $CO_2$  (Fig. 1d), but  $CH_4$  production was170close to zero (Fig. 1c). Similar results were obtained with IRRI soil (Fig. S1a-d). The accumulated acetate was171equimolar (slightly less than equimolar in the IRRI soil (Fig. S2d)) to the consumption of propionate (Fig. 2d), but172 $CH_4$  was not accumulated (Fig. 2c). Addition of  $CH_3F$  had no effect. Butyrate was not detected. The accumulated173acetate was subsequently degraded resulting in further production of  $CO_2$  (Fig. 1b,d).





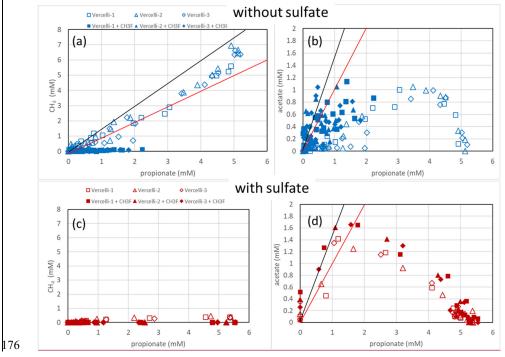


Figure 2: Balance of (a, c) produced CH<sub>4</sub> and (b, d) produced acetate against the consumed propionate under (a, b) methanogenic and (c, d) sulfidogenic conditions in paddy soil from Vercelli (Italy). The open and closed symbols denote conditions in the absence and the presence of CH<sub>3</sub>F, respectively. The black and red lines in panel (a) indicate aceticlastic methanogenesis after generation of acetate by either *Smithella* (equ.4) and or *Syntrophobacter* (equ.1), respectively. The black and red lines in panel (b and d) indicate transient acetate production by *Smithella* and *Syntrophobacter*, respectively. The different symbols indicate three different replicates.

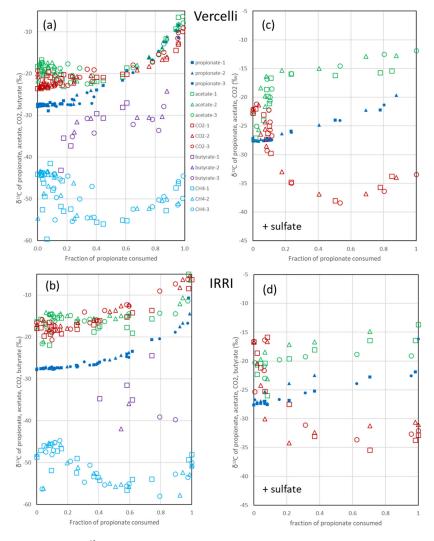
# 185 3.2 Isotope fractionation during propionate degradation

186 After onset of propionate degradation, the 813C of propionate (Fig. 1e) and acetate (Fig. 1f) increased indicating 187 that the light isotope was preferentially consumed. The  $\delta^{13}$ C values of CO<sub>2</sub> also increased (Fig. 1h). The same was 188 the case for butyrate (Fig. 1f). Similar results were obtained with IRRI soil (Fig. S1e-h). When aceticlastic 189 methanogenesis was inhibited by CH<sub>3</sub>F, the  $\delta^{13}$ C values of these compounds increased only slightly or decreased 190 (Fig. 1e,f,h). However, the  $\delta^{13}$ C of CH<sub>4</sub> was much more negative (30-50‰) in the presence than in the absence of 191 CH<sub>3</sub>F (Fig. 1g). The  $\delta^{13}$ C values of CH<sub>4</sub> in unamended soil (H<sub>2</sub>O control) were similar to those in propionate 192 amended soil (Fig. 1g). To visualize the change of the metabolic <sup>13</sup>C content of the metabolic products relative to 193 the substrates, the  $\delta^{13}$ C values were plotted against the increasing fractions ( $f_{prop}$ ) of propionate consumed both in 194 soil from Vercelli (Fig.3a) and the IRRI (Fig.3b). The patterns of  $\delta^{13}$ C values against the  $f_{prop}$  indicated kinetic 195 isotope fractionation. Note that the  $\delta^{13}$ C values of acetate and CO<sub>2</sub> were higher than those of propionate, whereas 196the values of butyrate and  $CH_4$  were lower (Fig.3a,b). The  $\delta^{13}C$  of  $CH_4$  decreased until about 40% of the propionate197had been consumed, and then increased again to its initial (low) values (-50% to -45%) (Fig.3a,b).

198 Under sulfidogenic conditions, only very little CH<sub>4</sub> was produced. Similarly as under methanogenic conditions,

199 the  $\delta^{13}$ C of propionate (Fig. 1e) and of acetate (Fig. 1f) increased after onset of propionate degradation indicating 200 that the light isotope was preferentially consumed. However, the  $\delta^{13}$ C values of CO<sub>2</sub> decreased during the first 10-201 15 days when acetate was accumulated (Fig. 1h, S1h). Inhibition of aceticlastic methanogenesis by CH<sub>3</sub>F had no 202 effect on the  $\delta^{13}$ C of propionate and CO<sub>2</sub>, but the values of acetate increased less than in the absence of CH<sub>3</sub>F (Fig. 203 1f). Also,  $\delta^{13}C$  of CH<sub>4</sub> was lower in the presence than in the absence of CH<sub>3</sub>F (Fig. 1g), but the amounts of CH<sub>4</sub> 204 produced were only very small (Fig. 1c). The values of  $\delta^{13}$ C of propionate and acetate increased with increasing 205  $f_{prop}$  (Fig. 3c,d). The  $\delta^{13}$ C of acetate was generally by about 5-10% higher than the  $\delta^{13}$ C of propionate but also 206 increased with  $f_{prop}$  indicating kinetic isotope fractionation. However, the  $\delta^{13}$ C of CO<sub>2</sub> did not increase, but instead 207 decreased after onset of propionate degradation reaching about -35‰ when 50% of the propionate had been 208 consumed and acetate accumulation had reached a maximum (Fig. 3c,d). Thereafter,  $\delta^{13}C$  of CO<sub>2</sub> increased or 209 became constant.

210 Mariotti plots of the  ${}^{13}C$  of propionate as function of  $f_{prop}$  could be created for methanogenic and sulfidogenic 211 incubation conditions, the latter both in the absence and the presence of CH<sub>3</sub>F (Fig. 4). The lines were straight even 212 when more than 70% of the propionate was consumed. Nevertheless, enrichment factors ( $\epsilon$ ) were determined only 213 for  $f_{prop} < 0.7$  and for regressions giving  $r^2 > 0.9$ . The  $\varepsilon_{prop}$  values were determined for each individual incubation 214 and then averaged over the replicates (n = 2-3). The results for Vercelli and IRRI soils are summarized in Fig. 5. 215 The average \$\varepsilon\_{prop}\$ values under methanogenic conditions were about -8‰ for Vercelli and about -3.5‰ for IRRI 216 soil. The average  $\epsilon_{prop}$  values under sulfidogenic conditions were around -3.5‰ in both soils and irrespectively 217 whether CH<sub>3</sub>F was present or not.



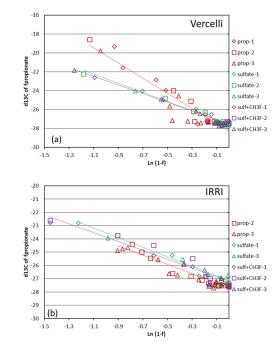
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**Figure 3:** Change of  $\delta^{13}$ C of propionate, acetate, butyrate, CO<sub>2</sub> and CH<sub>4</sub> relative to the fraction of propionate consumed ( $f_{prop}$ ) under (a, b) methanogenic and (c, d) sulfidogenic conditions in paddy soil from (a, c) Vercelli (Italy) and (b, d) the IRRI (the Philippines). The different symbols indicate three different replicates.

223 *3.3 Hydrogenotrophic methanogenesis* 

The difference in the  $\delta^{13}$ C of CH<sub>4</sub> in the presence and the absence of CH<sub>3</sub>F was used together with the  $\delta^{13}$ C of acetate to roughly estimate the percentage of CH<sub>4</sub> derived from H<sub>2</sub>/CO<sub>2</sub> versus acetate (Fig. S3). The percentage fractions of hydrogenotrophic methanogenesis (*f*<sub>*H*2</sub>) in Vercelli soil reached a maximum after 40-50 d when acetate concentrations also reached a maximum (Fig. S3a) and then decreased strongly. The same was the case in IRRI soil after around 35 d (Fig. S3b). When assuming a reasonable isotopic enrichment factor of  $\varepsilon_{CH4,ac-methyl} = -15\%$ ,

- 229 which is in-between the  $\varepsilon_{CH4,ac-methyl}$  of aceticlastic *Methanosaeta* (Penning et al., 2006; Valentine et al., 2004) and
- 230 Methanosarcina species (Gelwicks et al., 1994; Goevert and Conrad, 2009), the average f<sub>H2</sub> values were 0% for

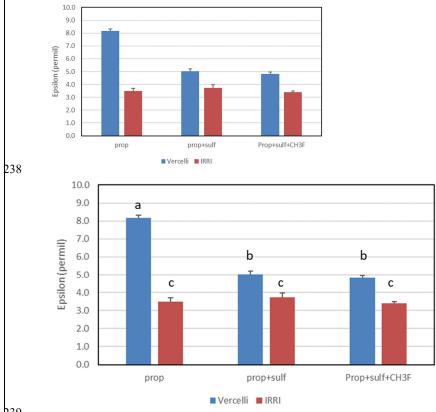


231 Vercelli soil and 20% for IRRI soil (Fig. S3c).

232 233

234 Figure 4: Mariotti plots of propionate consumption under methanogenic and sulfidogenic (± CH<sub>3</sub>F) conditions in

- 235 paddy soil from (a) Vercelli and (b) the IRRI. The different symbols indicate three different replicates; the lines
- 236 give the results of linear regression averaged over the replicates.



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240 241 242 243 Figure 5: Isotopic enrichment factors (Eprop, given as negative values) in paddy soils without and with addition of sulfate (gypsum) and  $CH_3F$ . Means  $\pm$  SE. The differences between the incubations were examined using Hukey's post hoc test of a one-way analysis of variance (ANOVA). Different letter son top of bars indicate significant difference (P < 0.05) between the data.

#### 245 4 Discussion

246 Pathway of propionate degradation

247 Our results showed that propionate was degraded via acetate as main transient intermediate finally resulting in 248 the production of CH<sub>4</sub> and CO<sub>2</sub> under methanogenic and CO<sub>2</sub> under sulfidogenic conditions. These results are 249 consistent with previous observations by Liu and Conrad (Liu and Conrad, 2017) using the same paddy soils. 250 Stable isotope probing and correlation network analysis of the microbial communities have shown that propionate 251 degradation is accomplished by both Syntrophopbacter and Smithella species (Gan et al., 2012; Liu and Conrad, 252 2017; Lueders et al., 2004). The present study showed that propionate degradation under methanogenic conditions 253 was consistent with the major operation of the Smithella pathway. The main argument for this conclusion is the 254 observation that butyrate was a transient intermediate of propionate degradation, albeit at low concentrations (Fig. 255 1, S1). In the Smithella pathway butyrate is further fermented to acetate and H<sub>2</sub>. However, production of H<sub>2</sub> is 256 smaller in the Smithella than in the Syntrophobacter pathway, while production of acetate is larger. Indeed,

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257 aceticlastic methanogenesis explained all the propionate-driven methanogenesis in the paddy soils (Fig. 2a, S2a). 258 The average hydrogenotrophic methanogenesis by contrast contributed almost zero in Vercelli soil and only about 259 20% in IRRI soil (Fig. S3c). The relatively larger contribution of aceticlastic than hydrogenotrophic 260 methanogenesis to methanogenic propionate degradation supports the conclusion that the Smithella pathway was 261 dominating over the Syntrophobacter pathway. Arguments against the Smithella pathway are that the accumulated 262 CH<sub>4</sub> amounted to less than the expected 1.75 mole per mole propionate consumed in Vercelli soil (Fig. 2a) and 263 even less in IRRI soil (Fig. S2a). With inhibition of aceticlastic methanogenesis, acetate accumulation in Vercelli 264 soil accounted for about 1 mole acetate per mole propionate, being in a range that is compatible with propionate 265 fermentation by either Smithella or Syntrophobacter (Fig. 2b). In IRRI soil however, acetate accumulation 266 accounted for less than 1 mole acetate per mole propionate (Fig. S2b). Note, however, that the accumulation of 267 acetate reflects only that part of propionate fermentation, which was not inhibited by CH<sub>3</sub>F. Our conclusion that 268 propionate was degraded mainly by Smithella under methanogenic conditions is consistent with the microbial 269 community structure in the paddy soils from Vercelli and IRRI, which contains not only Syntrophobacter species 270 but also Smithella together with Syntrophomonas, which is able to ferment butyrate (Liu and Conrad, 2017).

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

277 278 4 propionate + 3 sulfate + 3  $\text{H}^+ \rightarrow$  3  $\text{HS}^-$  + 4 acetate + 4  $\text{CO}_2$  + 4  $\text{H}_2\text{O}$ (10)279 It was interesting, that CH<sub>3</sub>F was not only a strong inhibitor of aceticlastic methanogenesis (which was 280 expected), but also a relatively strong inhibitor of propionate fermentation, but only under methanogenic but not 281 under sulfidogenic conditions. Inhibition of propionate fermentation under methanogenic conditions has been 282 observed before in three different paddy soils and has been interpreted as being due to the adverse thermodynamic 283 conditions when acetate accumulates (Conrad et al., 2014). However, this interpretation cannot be true, since 284 accumulation of acetate also occurred under sulfidogenic conditions, where CH<sub>3</sub>F did not inhibit propionate 285 degradation. In fact it is mainly the accumulation of H<sub>2</sub> rather than acetate, to which propionate degradation is 286 thermodynamically sensitive. This is the reason why the Smithella pathway is less sensitive to thermodynamic 287 inhibition than the Syntrophobacter pathway (Dolfing, 2013). However, CH3F did not inhibit H2 consumption by 288 methanogens, as seen by the low  $\delta^{13}$ C of CH<sub>4</sub> in the presence of CH<sub>3</sub>F. Furthermore, the first step of the Smithella-289 type propionate fermentation does not produce any H<sub>2</sub> and therefore, propionate should-in the presence of CH<sub>3</sub>F 290 should at least be fermented to butyrate and acetate, which however, was not the case. Hence, the reason why CH<sub>3</sub>F 291 inhibited propionate fermentation under methanogenic but not under sulfidogenic conditions remains unknown. 292 Perhaps it is Smithella being more sensitive to CH<sub>3</sub>F than Syntrophobacter. 293

293

294 Fractionation during propionate degradation

295 The isotopic fractionation of propionate apparently followed Raleigh distillation that is characteristic for kinetic 296 isotope fractionation in a closed system. The isotopic enrichment factor, which was determined from Mariotti plots, 297 was in the range of  $\varepsilon_{prop}$  = -8‰ to -3.5‰, which is less than the enrichment factor for methanogenic acetate 298 consumption, which has been found to be  $\varepsilon_{ac} = -21\%$  to -17% (Conrad et al., 2021). The  $\varepsilon_{prop}$  values are on the 299 same order as those predicted from  $\delta^{13}$ C values of propionate, acetate and organic carbon measured in various 300 methanogenic soils and sediments (Conrad et al., 2014). Propionate degradation resulted in the formation of <sup>13</sup>C-301 enriched acetate and CO2 and <sup>13</sup>C-depleted butyrate and CH4. The formation of <sup>13</sup>C-depleted butyrate can be 302 explained by kinetic isotope effect with the preferential utilization of <sup>13</sup>C-depleted propionate in the initial 303 dismutation reaction by Smithella. However, the production of 13C-enriched acetate cannot be explained by a linear 304 kinetic isotope effect. We assume that the dismutation of propionate is a branch point (Fry, 2003; Hayes, 2001), at 305 which the carbon flow is split into the production of <sup>13</sup>C-enriched acetate and <sup>13</sup>C-depleted butyrate. At the branch 306 point the carbon isotope flow shows a preferential flow of <sup>12</sup>C into the product generated by the reaction with the 307 larger fractionation factor, which would be butyrate. The further conversion of butyrate should produce acetate 308 that is depleted in <sup>13</sup>C. This acetate together with the acetate produced from propionate dismutation should result 309 in the  $\delta^{13}$ C-acetate that is observed. The total acetate pool initially had a  $\delta^{13}$ C that was up to 10% heavier than the 310  $\delta^{13}$ C of propionate. In the end, the  $\delta^{13}$ C values were about equal. The observation that acetate was  $^{13}$ C-enriched 311 relative to propionate is consistent with  $\delta^{13}C$  data in various soils and sediments (Conrad et al., 2014) reporting 312 that acetate is on the average enriched by 6% relative to propionate. Acetate was further converted to CH<sub>4</sub> and to 313  $CO_2$ . In Vercelli soil, the  $\delta^{13}C$  of  $CH_4$  was about 25-35% lighter than the  $\delta^{13}C$  of acetate. In IRRI soil,  $^{13}C$  depletion 314 was even larger (30-40%). In both soils, the isotopic enrichment factors for acetate consumption were in a range 315 of -12‰ to -17‰ and for CH4 production from acetate in a range of -37‰ to -27‰ (Conrad et al., 2021). 316 Considering that a certain percentage (albeit small) of CH4 was formed from CO2 reduction by hydrogenotrophic 317 methanogenesis, which displays relatively negative enrichment factors (see the  $\delta^{13}C$  of CH<sub>4</sub> in the presence of 318 CH<sub>3</sub>F, Fig. 1g), the observed difference in  $\delta^{13}$ C of CH<sub>4</sub> versus acetate is reasonable. In *Smithella* fermentation, the 319 only CO2 production occurs during the fermentation of butyrate and the aceticlastic conversion of acetate. In both 320 cases CO<sub>2</sub> should be <sup>13</sup>C-depleted relative to the substrates. Note, that this was not the case. Unfortunately, the <sup>13</sup>C 321 contents of the individual C atoms of propionate, butyrate and acetate are not known. The <sup>13</sup>C content in the 322 different C positions might also affect the  $\delta^{13}$ C of CH<sub>4</sub> and CO<sub>2</sub>, which are formed. It is also possible that besides 323 Smithella fermentation, the Syntrophobacter fermentation contributed to propionate degradation. In summary, the 324 detailed process of isotope fractionation during the pathway of propionate degradation is unclear. However, the 325 magnitude of the enrichment factors involved was relatively small, being on the order of <10%.

326 Under sulfidogenic conditions, propionate was most probably degraded by Syntrophobacter spp., first to 327 acetate, then finally to CO2 (Liu et al., 2018a; Liu and Conrad, 2017). The carbon isotope fractionation of 328 propionate consumption was with an enrichment factor of  $\varepsilon_{prop} = -3.5\%$  comparatively small. Propionate was 329 eventually converted to two carbon products of which one was depleted (the CO2) and the other was enriched (the 330 acetate) in <sup>13</sup>C. In case of Syntrophobacter-type degradation, acetate and CO<sub>2</sub> are produced from the conversion of 331 pyruvate, which is generated in the methylmalonyl-CoA pathway. In this pathway, CO2 is first consumed by the 332 conversion of propionyl-CoA to methylmalonyl-CoA and then produced by the conversion of oxaloacetate to 333 pyruvate. Pyruvate is finally converted to acetate and CO2, which should both be 13C-depleted with respect to 334 pyruvate (DeNiro and Epstein, 1977). However, both acetate and CO2 were initially <sup>13</sup>C-enriched relative to 335 propionate (about 2-5%), and then changed in opposite directions with acetate becoming increasingly <sup>13</sup>C-enriched 336 and CO2 becoming increasingly 13C-depleted until the time, when acetate accumulation had reached a maximum 337 (Fig. 5). Then,  $\delta^{13}$ C of both acetate and CO<sub>2</sub> increased together with the increase of  $^{13}$ C of propionate (Fig. 5). 338 Increase of  $\delta^{13}C$  of acetate is often explained by consumption, especially through aceticlastic methanogenesis 339 (Heuer et al., 2010; Heuer et al., 2009). However, hardly any CH4 was produced under sulfidogenic conditions and 340 the 13C enrichment occurred during the phase of acetate accumulation. Therefore, the enrichment likely happened 341 during acetate production from propionate degradation. The increasing <sup>13</sup>C-depletion of CO<sub>2</sub> can also not be 342 explained by consumption but only by the production from propionate. Hence, isotope fractionation during the 343 conversion of propionate, in particular during the conversion of propionate to pyruvate is unclear. We assume 344 complications during the carboxylation and decarboxylation reactions. Unfortunately, we hardly found any 345 literature data on the isotope fractionation of propionate fermentation. A coculture of Syntrophobacter 346 fumaroxidans with Methanobacterium formicicum exhibited marginal propionate fractionation with  $\varepsilon_{\text{prop}} = 0.9\%$ 347 and the formation of acetate, that was slightly <sup>13</sup>C-enriched (about 5‰) (Botsch and Conrad, 2011), similarly as 348 observed here. In summary, the mechanism of isotope fractionation during the conversion of propionate is not 349 completely clear, but the magnitude of isotope fractionation is quite low.

### 351 5 Conclusions

350

352 Propionate degradation under sulfidogenic conditions was explained by the metabolism of Syntrophobacteraceae, 353 which in a first step converted propionate to <sup>13</sup>C-enriched acetate and <sup>13</sup>C-depleted CO<sub>2</sub> By contrast, propionate 354 degradation under methanogenic conditions was at least partially due to metabolism by Smithella, which in a first 355 step converted propionate to <sup>13</sup>C-enriched acetate and <sup>13</sup>C-depleted butyrate. However, the isotopic enrichment 356 factors ( $\varepsilon_{prop}$ ) of propionate consumption in two paddy soils were generally very low (-8% to -3.5%) both under 357 methanogenic and sulfidogenic conditions. This low range is consistent with literature values of  $\delta^{13}$ C, collected 358 for propionate, acetate and organic carbon in various soils and sediments (Conrad et al., 2014). Fractionation of 359 propionate carbon actually seems to be smaller than fractionation of acetate, which is at least two times larger 360 (Conrad et al., 2021). Hence, degradation of organic carbon via propionate to acetate and CO2 apparently involves 361 only little isotope fractionation being on the order of <10%. By contrast, further degradation of acetate and CO<sub>2</sub> 362 (+H2) to CH4 involves substantial isotope fractionation. This is also the case for chemolithotrophic acetate 363 production (Conrad et al., 2014). 364

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# 365 Supplement link

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367 Author contribution: RC designed the experiments, evaluated the data and wrote the manuscript, PC conducted368 the experiments.

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

- 371
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