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

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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 aceticlastic 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 aceticlastic 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.

1 Introduction

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

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)

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

4 propionate
$$\rightarrow$$
 2 butyrate + 2 acetate (2)

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

46 2 butyrate + 4
$$H_2O \rightarrow$$
 4 acetate + 4 H_2 (3)

The *Smithella* pathway in total:

48 4 propionate + 4
$$H_2O \rightarrow 6$$
 acetate + 4 H_2 (4)

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

4 propionate
$$+ 2 \text{ H}_2\text{O} \rightarrow 7 \text{ CH}_4 + 5 \text{ CO}_2$$
, or (5)

56 4 propionate + 7 sulfate + 11 H⁺
$$\rightarrow$$
 7 HS⁻ + 12 CO₂ + 12 H₂O (6)

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

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

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

Knowledge of carbon isotope fractionation is important for the assessment of the pathways involved in anaerobic degradation of organic matter (Conrad, 2005; Elsner et al., 2005). The δ^{13} C values of organic carbon, acetate and propionate in various soils and sediments were found to be similar (Conrad et al., 2014). The similarity indicates that the enrichment factors (ϵ) of the processes involved in both production and consumption of propionate are probably small. The direct determination of ϵ values in microbial cultures of one propionate-producing and one propionate-consuming bacterium also showed low values (Botsch and Conrad, 2011). However, direct determination of ϵ values in environmental samples is missing. Therefore, we decided to measure isotope fractionation in methanogenic and sulfidogenic paddy soil amended with propionate along with the recording of the production of acetate, CH₄ and CO₂. We also used the treatment with methyl fluoride (CH₃F) to inhibit the consumption of acetate by methanogenic archaea (Janssen and Frenzel, 1997). Recently, we determined the microbial communities in methanogenic and sulfidogenic rice field soils, which were used for assessment of ¹³C isotope fractionation during acetate consumption (Conrad et al., 2021). Here we present analogous data from the same soil suspensions prepared for the propionate degradation experiments.

2 Materials and Methods

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). The main 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%.

The experimental setup was exactly the same as during a previous study on acetate consumption (Conrad et 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 4 weeks. In a second incubation, paddy soil was mixed with autoclaved anoxic H_2O (prepared under N_2) at a ratio of 1:1, was amended with 0.07 g CaSO₄.2 H_2O , and then incubated under N_2 at 25°C for 4 weeks. These two preincubated soil slurries were sampled and stored at -20°C for later molecular analysis (see data in Conrad et al. (2021)). The preincubated soil slurries were also used (in 3 replicates) for the following incubation experiments. Two different sets of incubations were prepared. In the first set (resulting in methanogenic conditions), 5 mL soil slurry preincubated without sulfate was incubated at 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 bottles were the amended with (i) 5 mL H_2O ; (ii) 5 mL H_2O + 4.5 mL H_2O ; (iii) 5 mL 50 mM sodium propionate; (iv) 5 mL 50 mM sodium acetate + 4.5 mL H_2O ; In the second set (resulting in sulfidogenic conditions), 5 mL soil slurry preincubated with sulfate was incubated at 25°C with 40 mL 20 mM potassium phosphate buffer (pH 7.0) in a 150-mL bottle under an atmosphere of H_2O . The amendments were the same as above, but with the addition of 200 H_2O 1 and H_2O 2 suspension corresponding to a concentration of 2.5 M (giving a final concentration of 10 mM sulfate).

2.2 Chemical and isotopic analyses

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

2.3 Calculations

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

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

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

also expressed as $\varepsilon \equiv 1000~(1-\alpha)$ in permil. The carbon isotope enrichment factor ε_{prop} associated with propionate consumption was calculated from the temporal change of $\delta^{13}C$ of propionate as described by Mariotti et al. (Mariotti et al., 1981) from the residual reactant

$$\delta_{\rm r} = \delta_{\rm ri} + \varepsilon \left[\ln(1-f) \right] \tag{8}$$

where δ_{ri} is the isotopic composition of the reactant (propionate) at the beginning, and δ_r is the isotopic composition of the residual propionate, both at the instant when f is determined. f_{prop} is the fractional yield of the products based on the consumption of propionate $(0 < f_{prop} < 1)$. Linear regression of δ^{13} C of propionate against $\ln(1 - f)$ yields ϵ_{prop} as the slope of best fit lines. The regressions of δ^{13} C of propionate were done for data in the range of $f_{prop} < 0.7$. The linear regressions were done individually for each experimental replicate (n = 3) and were only accepted if $r^2 > 0.9$. The ϵ values resulting from the replicate experiments were then averaged (\pm SE).

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

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

with $\delta^{13}C_{CH4} = \delta^{13}C$ of total CH₄ produced, $\delta^{13}C_{CH4\text{-mc}} = \delta^{13}C$ of CH₄ produced from hydrogenotrophic methanogenesis, which is equivalent to the CH₄ produced in the presence of CH₃F, and $\delta^{13}C_{CH4\text{-ma}} = \delta^{13}C$ of CH₄ produced from aceticlastic methanogenesis. The $\delta^{13}C_{CH4\text{-ma}}$ was approximated from the $\delta^{13}C$ of acetate in the presence of CH₃F assuming that the methyl group of acetate was depleted in ¹³C by 8% (Conrad et al., 2014) and that the enrichment factor ($\epsilon_{CH4,ac\text{-methyl}}$) for CH₄ being produced from acetate-methyl was between 0 and -20%.

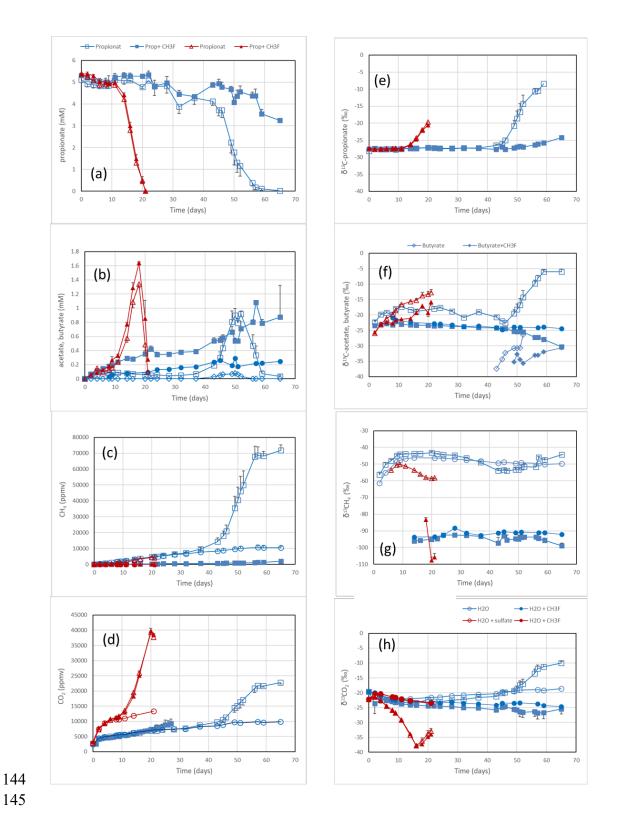


Figure 1: Propionate conversion to acetate, butyrate, CH_4 and CO_2 in suspensions of paddy soil from Vercelli (Italy) after addition of propionate without sulfate (blue squares) or propionate plus sulfate (gypsum) (red triangles) without CH_3F (open symbols) or with CH_3F (closed symbols). Controls 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 CH_4 (1 ppmv = 10^{-6} bar), (d) mixing ratios of CO_2 , (e) $\delta^{13}C$ of propionate, (f) $\delta^{13}C$ of acetate and butyrate, (g) $\delta^{13}C$ of CH_4 , and (h) $\delta^{13}C$ of CO_2 . Means \pm SE.

3 Results

3.1 Conversion of propionate under methanogenic and sulfidogenic conditions

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

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

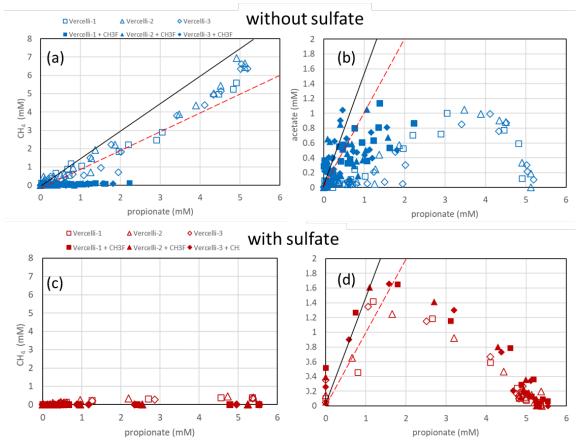


Figure 2: Balance of (a, c) produced CH₄ and (b, d) produced acetate against the consumed propionate under (a, b) methanogenic and (c, d) sulfidogenic conditions in paddy soil from Vercelli (Italy). The open and closed symbols denote conditions in the absence and the presence of CH₃F, respectively. The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

3.2 Isotope fractionation during propionate degradation

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

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

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

Figure 3: Change of δ^{13} C of propionate, acetate, butyrate, CO₂ and CH₄ 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.

3.3 Hydrogenotrophic methanogenesis

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The difference in the δ^{13} C of CH₄ in the presence and the absence of CH₃F was used together with the δ^{13} C of acetate to roughly estimate the percentage of CH₄ derived from H₂/CO₂ versus acetate (Fig. S3). The percentage fractions of hydrogenotrophic methanogenesis (f_{H2}) 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 $\varepsilon_{\text{CH4,ac-methyl}} = -15\%$, which is in-between the $\varepsilon_{\text{CH4,ac-methyl}}$ of aceticlastic *Methanosaeta* (Penning et al., 2006; Valentine et al., 2004) and *Methanosarcina* species (Gelwicks et al., 1994; Goevert and Conrad, 2009), the average f_{H2} values were 0% for Vercelli soil and 20% for IRRI soil (Fig. S3c).

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-1.5

(b)

-1.3

-1.1

-0.9

-0.7

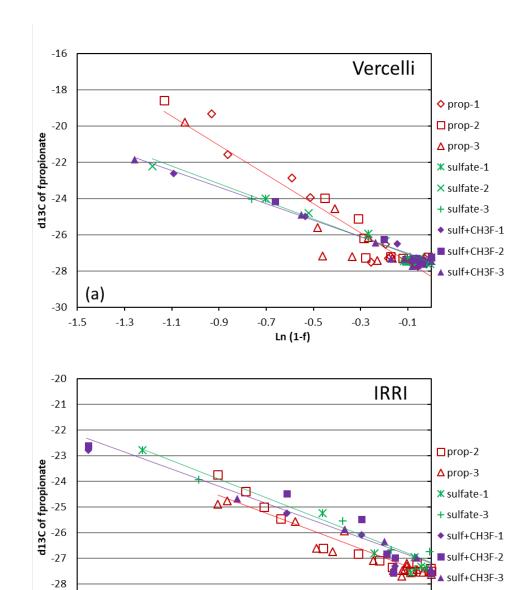


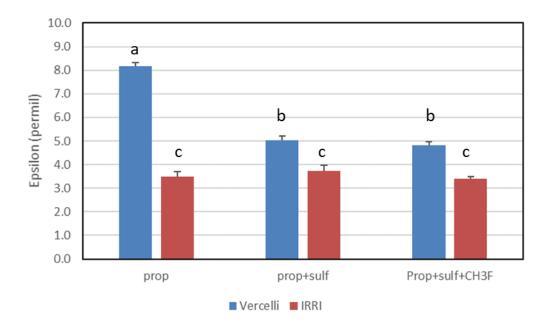
Figure 4: Mariotti plots of propionate consumption under methanogenic and sulfidogenic (\pm CH₃F) conditions in paddy soil from (a) Vercelli and (b) the IRRI. The different symbols indicate three different replicates; the lines give the results of linear regression averaged over the replicates.

-0.5

Ln (1-f)

-0.3

-0.1



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Figure 5: Isotopic enrichment factors (ε_{prop} , given as negative values) in paddy soils without and with addition of sulfate (gypsum) and CH₃F. 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.

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4 Discussion

Pathway of propionate degradation

Our results showed that propionate was degraded via acetate as main transient intermediate finally resulting in the production of CH₄ and CO₂ under methanogenic and CO₂ under sulfidogenic conditions. These results are consistent with previous observations by Liu and Conrad (Liu and Conrad, 2017) using the same paddy soils. Stable isotope probing and correlation network analysis of the microbial communities have shown that propionate degradation is accomplished by both Syntrophopbacter and Smithella species (Gan et al., 2012; Liu and Conrad, 2017; Lueders et al., 2004). The present study showed that propionate degradation under methanogenic conditions was consistent with the major operation of the Smithella pathway. The main argument for this conclusion is the observation that butyrate was a transient intermediate of propionate degradation, albeit at low concentrations (Fig. 1, S1). In the Smithella pathway butyrate is further fermented to acetate and H2. However, production of H2 is smaller in the Smithella than in the Syntrophobacter pathway, while production of acetate is larger. Indeed, aceticlastic methanogenesis explained all the propionate-driven methanogenesis in the paddy soils (Fig. 2a, S2a). The average hydrogenotrophic methanogenesis by contrast contributed almost zero in Vercelli soil and only about 20% in IRRI soil (Fig. S3c). The relatively larger contribution of aceticlastic than hydrogenotrophic methanogenesis to methanogenic propionate degradation supports the conclusion that the Smithella pathway was dominating over the Syntrophobacter pathway. Arguments against the Smithella pathway are that the accumulated CH₄ amounted to less than the expected 1.75 mole per mole propionate consumed in Vercelli soil (Fig. 2a) and even less in IRRI soil (Fig. S2a). With inhibition of aceticlastic methanogenesis, acetate accumulation in Vercelli soil accounted for about 1 mole acetate per mole propionate, being in a range that is compatible with propionate fermentation by either Smithella or Syntrophobacter (Fig. 2b). In IRRI soil however, acetate accumulation accounted for less than 1 mole acetate per mole propionate (Fig. S2b). Note, however, that the accumulation of acetate reflects only that part of propionate fermentation, which was not inhibited by CH₃F. Our conclusion that propionate was degraded mainly by *Smithella* under methanogenic conditions is consistent with the microbial community structure in the paddy soils from Vercelli and IRRI, which contains not only *Syntrophobacter* species 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 CO₂, or syntrophically as under methanogenic conditions, but with subsequent oxidation of H₂ 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 CO₂, and subsequently oxidized the accumulated acetate to CO₂. 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

4 propionate + 3 sulfate + 3 H⁺
$$\rightarrow$$
 3 HS⁻ + 4 acetate + 4 CO₂ + 4 H₂O (10)

It was interesting, that CH₃F was not only a strong inhibitor of aceticlastic methanogenesis (which was expected), but also a relatively strong inhibitor of propionate fermentation, but only under methanogenic but not under sulfidogenic conditions. Inhibition of propionate fermentation under methanogenic conditions has been observed before in three different paddy soils and has been interpreted as being due to the adverse thermodynamic conditions when acetate accumulates (Conrad et al., 2014). However, this interpretation cannot be true, since accumulation of acetate also occurred under sulfidogenic conditions, where CH₃F did not inhibit propionate degradation. In fact it is mainly the accumulation of H₂ rather than acetate, to which propionate degradation is thermodynamically sensitive. This is the reason why the *Smithella* pathway is less sensitive to thermodynamic inhibition than the *Syntrophobacter* pathway (Dolfing, 2013). However, CH₃F did not inhibit H₂ consumption by methanogens, as seen by the low δ^{13} C of CH₄ in the presence of CH₃F. Furthermore, the first step of the *Smithella*-type propionate fermentation does not produce any H₂ and therefore, propionate in the presence of CH₃F should at least be fermented to butyrate and acetate, which however, was not the case. Hence, the reason why CH₃F inhibited propionate fermentation under methanogenic but not under sulfidogenic conditions remains unknown. Perhaps it is *Smithella* being more sensitive to CH₃F than *Syntrophobacter*.

Fractionation during propionate degradation

The isotopic fractionation of propionate apparently followed Raleigh distillation that is characteristic for kinetic isotope fractionation in a closed system. The isotopic enrichment factor, which was determined from Mariotti plots, was in the range of $\varepsilon_{prop} = -8\%$ to -3.5%, which is less than the enrichment factor for methanogenic acetate consumption, which has been found to be $\varepsilon_{ac} = -21\%$ to -17% (Conrad et al., 2021). The ε_{prop} values are on the same order as those predicted from δ^{13} C values of propionate, acetate and organic carbon measured in various methanogenic soils and sediments (Conrad et al., 2014). Propionate degradation resulted in the formation of 13 C-enriched acetate and CO_2 and 13 C-depleted butyrate and CH_4 . The formation of 13 C-depleted butyrate can be explained by kinetic isotope effect with the preferential utilization of 13 C-depleted propionate in the initial dismutation reaction by *Smithella*. However, the production of 13 C-enriched acetate cannot be explained by a linear kinetic isotope effect. We assume that the dismutation of propionate is a branch point (Fry, 2003; Hayes, 2001), at

which the carbon flow is split into the production of ¹³C-enriched acetate and ¹³C-depleted butyrate. At the branch point the carbon isotope flow shows a preferential flow of ¹²C into the product generated by the reaction with the larger fractionation factor, which would be butyrate. The further conversion of butyrate should produce acetate that is depleted in ¹³C. This acetate together with the acetate produced from propionate dismutation should result in the δ^{13} C-acetate that is observed. The total acetate pool initially had a δ^{13} C that was up to 10% heavier than the δ^{13} C of propionate. In the end, the δ^{13} C values were about equal. The observation that acetate was 13 C-enriched relative to propionate is consistent with δ¹³C data in various soils and sediments (Conrad et al., 2014) reporting that acetate is on the average enriched by 6% relative to propionate. Acetate was further converted to CH₄ and to CO_2 . In Vercelli soil, the $\delta^{13}C$ of CH₄ was about 25-35% lighter than the $\delta^{13}C$ of acetate. In IRRI soil, ^{13}C depletion was even larger (30-40%). In both soils, the isotopic enrichment factors for acetate consumption were in a range of -12% to -17% and for CH₄ production from acetate in a range of -37% to -27% (Conrad et al., 2021). Considering that a certain percentage (albeit small) of CH₄ was formed from CO₂ reduction by hydrogenotrophic methanogenesis, which displays relatively negative enrichment factors (see the δ^{13} C of CH₄ in the presence of CH₃F, Fig. 1g), the observed difference in δ^{13} C of CH₄ versus acetate is reasonable. In *Smithella* fermentation, the only CO₂ production occurs during the fermentation of butyrate and the aceticlastic conversion of acetate. In both cases CO₂ should be ¹³C-depleted relative to the substrates. Note, that this was not the case. Unfortunately, the ¹³C contents of the individual C atoms of propionate, butyrate and acetate are not known. The ¹³C content in the different C positions might also affect the δ^{13} C of CH₄ and CO₂, which are formed. It is also possible that besides Smithella fermentation, the Syntrophobacter fermentation contributed to propionate degradation. In summary, the detailed process of isotope fractionation during the pathway of propionate degradation is unclear. However, the magnitude of the enrichment factors involved was relatively small, being on the order of <10%.

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Under sulfidogenic conditions, propionate was most probably degraded by Syntrophobacter spp., first to acetate, then finally to CO2 (Liu et al., 2018a; Liu and Conrad, 2017). The carbon isotope fractionation of propionate consumption was with an enrichment factor of $\varepsilon_{prop} = -3.5\%$ comparatively small. Propionate was eventually converted to two carbon products of which one was depleted (the CO₂) and the other was enriched (the acetate) in ¹³C. In case of Syntrophobacter-type degradation, acetate and CO₂ are produced from the conversion of pyruvate, which is generated in the methylmalonyl-CoA pathway. In this pathway, CO2 is first consumed by the conversion of propionyl-CoA to methylmalonyl-CoA and then produced by the conversion of oxaloacetate to pyruvate. Pyruvate is finally converted to acetate and CO₂, which should both be ¹³C-depleted with respect to pyruvate (DeNiro and Epstein, 1977). However, both acetate and CO₂ were initially ¹³C-enriched relative to propionate (about 2-5%), and then changed in opposite directions with acetate becoming increasingly ¹³C-enriched and CO₂ becoming increasingly ¹³C-depleted until the time, when acetate accumulation had reached a maximum (Fig. 5). Then, δ^{13} C of both acetate and CO₂ increased together with the increase of 13 C of propionate (Fig. 5). Increase of δ^{13} C of acetate is often explained by consumption, especially through aceticlastic methanogenesis (Heuer et al., 2010; Heuer et al., 2009). However, hardly any CH₄ was produced under sulfidogenic conditions and the ¹³C enrichment occurred during the phase of acetate accumulation. Therefore, the enrichment likely happened during acetate production from propionate degradation. The increasing ¹³C-depletion of CO₂ can also not be explained by consumption but only by the production from propionate. Hence, isotope fractionation during the conversion of propionate, in particular during the conversion of propionate to pyruvate is unclear. We assume

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

Conclusions

Propionate degradation under sulfidogenic conditions was explained by the metabolism of *Syntrophobacteraceae*, which in a first step converted propionate to 13 C-enriched acetate and 13 C-depleted CO₂. By contrast, propionate degradation under methanogenic conditions was at least partially due to metabolism by *Smithella*, which in a first step converted propionate to 13 C-enriched acetate and 13 C-depleted butyrate. However, the isotopic enrichment factors (ϵ_{prop}) of propionate consumption in two paddy soils were generally very low (-8‰ to -3.5‰) both under methanogenic and sulfidogenic conditions. This low range is consistent with literature values of δ^{13} C, collected for propionate, acetate and organic carbon in various soils and sediments (Conrad et al., 2014). Fractionation of propionate carbon actually seems to be smaller than fractionation of acetate, which is at least two times larger (Conrad et al., 2021). Hence, degradation of organic carbon via propionate to acetate and CO₂ apparently involves only little isotope fractionation being on the order of <10‰. By contrast, further degradation of acetate and CO₂ (+H₂) to CH₄ involves substantial isotope fractionation. This is also the case for chemolithotrophic acetate production (Conrad et al., 2014).

Supplement link

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

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372 References

- Boone, D. R. and Bryant, M. P.: Propionate-degrading bacterium, Syntrophobacter wolinii sp. nov.
- 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
- 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
- degradation of organic matter in the sediment of an acidic bog lake, Lake Grosse Fuchskuhle,
- 385 Limnol. Oceanogr., 55, 1932-1942, 2010.
- 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
- 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.
- DeBok, F. A. M., Harmsen, H. J. M., Plugge, C. M., DeVries, M. C., Akkermans, A. D. L., DeVos,
- 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
- description of the genus *Pelotomaculum*, Int. J. Syst. Evol. Microbiol., 55, 1697-1703, 2005.
- DeBok, F. A. M., Stams, A. J. M., Dijkema, C., and Boone, D. R.: Pathway of propionate oxidation
- 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
- methanogenic conditions, Appl. Environ. Microbiol., 79, 4515-4516, 2013.
- Elsner, M., Zwank, L., Hunkeler, D., and Schwarzenbach, R. P.: A new concept linking observable
- 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 aceticlastic
- 411 methanogenesis, Appl. Environ. Microbiol., 60, 467-472, 1994.
- 412 Glissmann, K. and Conrad, R.: Fermentation pattern of methanogenic degradation of rice straw in
- anoxic paddy soil, FEMS Microbiol. Ecol., 31, 117-126, 2000.
- 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 ¹³C 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.
- Heuer, V. B., Krüger, M., Elvert, M., and Hinrichs, K. U.: Experimental studies on the stable carbon
- isotope biogeochemistry of acetate in lake sediments, Org. Geochem., 41, 22-30, 2010.
- Heuer, V. B., Pohlman, J. W., Torres, M. E., Elvert, M., and Hinrichs, K. U.: The stable carbon
- isotope biogeochemistry of acetate and other dissolved carbon species in deep subseafloor
- sediments at the northern Cascadia Margin, Geochim. Cosmochim. Acta, 73, 3323-3336, 2009.
- Houwen, F. P., Dijkema, C., Stams, A. J. M., and Zehnder, A. J. B.: Propionate metabolism in
- anaerobic bacteria determination of carboxylation reactions with 13C-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
- thermopropionicum gen. nov., sp. nov., an anaerobic, thermophilic, syntrophic propionate-
- oxidizing bacterium, Int. J. Syst. Evol. Microbiol., 52, 1729-1735, 2002.
- Imachi, H., Sekiguchi, Y., Kamagata, Y., Loy, A., Qiu, Y. L., Hugenholtz, P., Kimura, N., Wagner,
- M., Ohashi, A., and Harada, H.: Non-sulfate-reducing, syntrophic bacteria affiliated with
- Desulfotomaculum cluster I are widely distributed in methanogenic environments, Appl.
- 435 Environ. Microbiol., 72, 2080-2091, 2006.
- Janssen, P. H. and Frenzel, P.: Inhibition of methanogenesis by methyl fluoride studies of pure and
- 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
- soil, FEMS Microbiol. Ecol., 26, 281-288, 1998.

- Krylova, N. I., Janssen, P. H., and Conrad, R.: Turnover of propionate in methanogenic paddy soil,
- 442 FEMS Microbiol. Ecol., 23, 107-117, 1997.
- Liu, P., Pommerenke, B., and Conrad, R.: Identification of Syntrophobacteraceae as major acetate-
- 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
- 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
- methanogenic microbial communities in two paddy soils and one desert soil, Soil Biol. Biochem.,
- 450 124, 236-244, 2018b.
- Liu, Y. T., Balkwill, D. L., Aldrich, H. C., Drake, G. R., and Boone, D. R.: Characterization of the
- anaerobic propionate-degrading syntrophs Smithella propionica gen. nov., sp. nov. and
- 453 Syntrophobacter wolinii, Int. J. Syst. Bacteriol., 49, 545-556, 1999.
- 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.:
- Experimental determination of nitrogen kinetic isotope fractionation: some principles;
- 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.
- nov. sp. nov., an anaerobic, syntrophic, fatty acid-oxidizing bacterium, Appl. Environ.
- 462 Microbiol., 41, 1029-1039, 1981.
- Nozoe, T.: Effects of methanogenesis and sulfate-reduction on acetogenetic oxidation of propionate
- and further decomposition of acetate in paddy soil, Soil Sci. Plant Nutr., 43, 1-10, 1997.
- 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.
- Plugge, C. M., Balk, M., and Stams, A. J. M.: Desulfotomaculum thermobenzoicum subsp.
- 469 thermosyntrophicum subsp. nov., a thermophilic, syntrophic, propionate-oxidizing, spore-
- 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
- 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