

# Variation of carbon isotope fractionation in hydrogenotrophic methanogenic microbial cultures and environmental samples at different energy status

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## Abstract

Methane is a major product of anaerobic degradation of organic matter and an important greenhouse gas. Its stable carbon isotope composition can be used to reveal active methanogenic pathways, if associated isotope fractionation factors are known. To clarify the causes that lead to the wide variation of fractionation factors of methanogenesis from H<sub>2</sub> plus CO<sub>2</sub> ( $\alpha_{\text{CO}_2-\text{CH}_4}$ ), pure cultures and various cocultures were grown under different thermodynamic conditions. In syntrophic and obligate syntrophic cocultures thriving on different carbohydrate substrates, fermentative bacteria were coupled to three different species of hydrogenotrophic methanogens of the families Methanobacteriaceae and Methanomicrobiaceae. We found that C-isotope fractionation was correlated to the Gibbs free energy change ( $\Delta G$ ) of CH<sub>4</sub> formation from H<sub>2</sub> plus CO<sub>2</sub> and that the relation can be described by a semi-Gauss curve. The derived relationship was used to quantify the average  $\Delta G$  that is available to hydrogenotrophic methanogenic archaea in their habitat, thus avoiding the problems encountered with measurement of low H<sub>2</sub> concentrations on a microscale. Boreal peat, rice field soil, and rumen fluid, which represent major sources of atmospheric CH<sub>4</sub>, exhibited increasingly smaller  $\alpha_{\text{CO}_2-\text{CH}_4}$ , indicating that thermodynamic conditions for hydrogenotrophic methanogens became increasingly more favourable. Vice versa, we hypothesize that environments with similar energetic conditions will also exhibit similar isotope fractionation. Our results, thus, provide a mechanistic constraint for modelling the <sup>13</sup>C flux from microbial sources of atmospheric CH<sub>4</sub>.

**Keywords:** coculture, fractionation, Gibbs free energy, hydrogen, methanogenesis, stable carbon isotopes, thermodynamic limit

Received 6 July 2005; and accepted 23 August 2005

## Introduction

Anoxic environments such as natural wetlands and flooded rice fields contribute approximately a third to the CH<sub>4</sub> budget of the atmosphere (Cicerone & Orem-land, 1988). In these systems microbes almost exclusively produce CH<sub>4</sub> from acetate and H<sub>2</sub>/CO<sub>2</sub>. Hydrogenotrophic methanogenesis ( $4\text{H}_2 + \text{CO}_2 \rightarrow \text{CH}_4 + 2\text{H}_2\text{O}$ ;  $\Delta G^\circ = -130.7 \text{ kJ mol}^{-1}$ ) strongly prefers the isotopically lighter carbon, whereas the isotope effect is less expressed in acetoclastic methanogenesis. This

difference in isotope fractionation can be used for modelling of C-flux in methanogenic environments (Conrad, 2005). In addition, measurements of carbon stable isotope ratios of CH<sub>4</sub> have been proven to be useful for constraining individual atmospheric CH<sub>4</sub> sources and sinks, and in interpreting the atmospheric CH<sub>4</sub> budget (Lowe *et al.*, 1994; Gupta *et al.*, 1996; Bräunlich *et al.*, 2001; Fletcher *et al.*, 2004). Although the <sup>13</sup>C/<sup>12</sup>C-isotopic composition of CH<sub>4</sub> produced from CO<sub>2</sub> reduction has a characteristic signature, the magnitude of fractionation is usually considerably different between many natural systems and methanogenic microbial cultures (Fey *et al.*, 2004; Valentine *et al.*, 2004; Conrad, 2005). The wide variation of mag-

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nitude of fractionation (i.e. fractionation factor  $\alpha$ ) is a problem for modelling of C-fluxes (Conrad, 2005). Several authors observed this discrepancy, and assumed it was because of microbial culturing conditions, which often strongly differ from those in the natural environment (Burke, 1993; Sugimoto & Wada, 1993; Whiticar, 1999). However, the factors determining the wide range of  $\alpha$ -values are basically unknown. Recently, Valentine *et al.* (2004) found that the fractionation of C-isotopes was affected by the supply of H<sub>2</sub> to a CO<sub>2</sub>-reducing culture of *Methanothermobacter marburgensis*, and explained the observation by the differential reversibility hypothesis. They proposed that isotope fractionation in multistep enzymatic processes depends on enzymatic reversibility, which in turn depends on the Gibbs free energy of catabolism. So far, this is the only study measuring both H<sub>2</sub> partial pressures ( $p_{\text{H}_2}$ ) and C-isotope fractionation explicitly, and only two data points exist (Valentine *et al.*, 2004). To control  $p_{\text{H}_2}$  and rigorously study its effect on C-isotope fractionation, we grew cocultures of different H<sub>2</sub>-producing fermenting bacteria and H<sub>2</sub>-consuming methanogenic archaea covering a broad range of  $p_{\text{H}_2}$ . This experimental approach was used to test whether a general function of methanogenic carbon isotope fractionation vs.  $p_{\text{H}_2}$  exists. We found that such a relationship indeed existed, such that  $\alpha$  changed with the Gibbs free energy ( $\Delta G$ ) available to hydrogenotrophic methanogens, and that this relationship was generally valid for the different methanogenic microbial species studied.

Furthermore, we tested whether it is possible to predict  $\alpha$  from  $p_{\text{H}_2}$  measured in samples from natural methanogenic environments. Our results indicate that this is not a reliable option, as the measured values of  $p_{\text{H}_2}$  hardly represent the Gibbs free energy ( $\Delta G$ ) available to hydrogenotrophic methanogens *in situ*. Such a determination has been problematic, as methanogens typically live within micro-aggregates involving steep gradients of  $p_{\text{H}_2}$  (Conrad *et al.*, 1985; Krylova & Conrad, 1998; Hoehler *et al.*, 2001), preventing the exact analysis of energetically relevant H<sub>2</sub> concentrations *in situ*. Instead, we propose that measurement of  $\alpha$  gives a more reliable value of the *in situ*  $\Delta G$  than the measurement of  $p_{\text{H}_2}$ , and show that more free energy is available to the hydrogenotrophic methanogens than suggested by measurement of  $p_{\text{H}_2}$ .

## Material and methods

### Archaeal and bacterial strains

*Methanobacterium bryantii* (DSM 863), *Methanobacterium formicicum* (DSM 1535), *Methanospirillum hungatei* (DSM 864), *M. marburgensis* (DSM 2133), *Acetobacterium woodii*

(DSM 1030), and *Clostridium papyrosolvans* (DSM 2782) were obtained from Deutsche Sammlung für Mikroorganismen und Zellkulturen. *Syntrophobacter fumaroxidans* (DSM 10017) was from our own culture collection (Laboratory of Microbiology, Wageningen University, The Netherlands).

### Cultivation

Cocultures of the fermenting organisms *C. papyrosolvans* and *A. woodii* were each grown with methanogenic partner organisms (*M. bryantii* or *M. hungatei*) in glass bottles (500 mL; Müller Krempel, Bülach, Switzerland) with 250 mL as culture volume on phosphate-buffered mineral medium with defined bicarbonate concentrations (0.16–2.8 mM) under N<sub>2</sub>. The composition was (in g L<sup>-1</sup> unless otherwise indicated as follows): KH<sub>2</sub>PO<sub>4</sub>, 1.9; Na<sub>2</sub>HPO<sub>4</sub> · 2H<sub>2</sub>O, 6.4; NH<sub>4</sub>Cl, 0.3; MgCl<sub>2</sub> · 6H<sub>2</sub>O, 0.1; NaCl, 0.3; KCl, 0.15; CaCl<sub>2</sub> · 2H<sub>2</sub>O, 0.055; Na<sub>2</sub>S · 9H<sub>2</sub>O, 0.24; trace element solution, 2 mL (Chin *et al.*, 1998); alkaline trace element solution 1 mL (Stams *et al.*, 1993); vitamin solution, 1 mL (Wolin *et al.*, 1963); resazurine at 0.5% (wt/vol), 1 mL; pH adjusted to 7.2. Concentrations of glucose, cellobiose, and cellulose (expressed as anhydroglucose;  $M_w = 162 \text{ g mol}^{-1}$ ) were 4.44, 2.34, and 4.94 mM, respectively. Cocultures of *S. fumaroxidans* and *M. hungatei* or *M. formicicum* were grown in glass bottles (1000 mL; Müller Krempel) under N<sub>2</sub> on the medium described by Stams *et al.* (1993) containing 38 mM bicarbonate and 30 mM propionate. *M. marburgensis* was grown in a flow-through batch reactor on mineral medium with H<sub>2</sub>/CO<sub>2</sub>/N<sub>2</sub> gas mixtures of 80/20/0% and 5/20/75%, respectively (Schönheit *et al.*, 1980). Rapid gas flow guaranteed that  $\leq 3\%$  of the CO<sub>2</sub> supplied was converted to CH<sub>4</sub>. This assured that inflow and outflow of CO<sub>2</sub> had nearly the same isotope composition and proper calculation of  $\alpha_{\text{CO}_2\text{-CH}_4}$  was possible. Cultures were grown at 30 °C, except for those with *S. fumaroxidans* and *M. marburgensis* grown at 37 °C and 65 °C, respectively.

### Environmental samples

Rice field soil samples were collected from rice fields of the Italian Rice Research Institute in Vercelli, Italy. Soil slurries, which represent the water-saturated conditions after flooding in rice fields, were prepared at a weight ratio of 1 : 1 with the addition of 4 mg rice straw g<sup>-1</sup> soil and anoxically incubated at 30 °C. The peat was sampled at the Lakkasuo mire complex in central Finland (61°48'N, 24°19'E, ca. 150 m altitude), from a mesotrophic fen (MES), an oligotrophic fen (OLI), and an ombrotrophic bog (OMB) and anoxically incubated at 10 °C (Galand *et al.*, 2005). Methanogenesis from

acetate was inhibited with CH<sub>3</sub>F (Janssen *et al.*, 1997). The concentration of CH<sub>3</sub>F was optimized as described (Conrad & Klose, 1999), applying 1% in rice field soil and 2% in peat soil. Selective inhibition by CH<sub>3</sub>F was earlier confirmed by stoichiometrical correlation of accumulated acetate with the deficit in CH<sub>4</sub> production (Frenzel & Bosse, 1996; Conrad & Klose, 1999). At higher CH<sub>3</sub>F concentration hydrogenotrophic methanogenesis is partially inhibited (Conrad & Klose, 1999), but does not change its carbon isotope fractionation (Conrad, 2005). Rumen fluid was sampled at the slaughterhouse of Marburg and immediately afterwards incubated under N<sub>2</sub> at 30 °C or H<sub>2</sub>/CO<sub>2</sub> (4:1) at 25 °C.

### Analytical methods

CH<sub>4</sub> and CO<sub>2</sub> were analysed by GC-FID (Shimadzu, Kyoto, Japan). CO<sub>2</sub> was detected upon conversion to CH<sub>4</sub> with a methanizer (Ni-catalyst at 350 °C, Chrompack, Middelburg, Netherlands). H<sub>2</sub> was analysed by GC-TCD (Shimadzu) and a HgO-to-Hg conversion detector (RGD2; Trace Analytical, Menlo Park, CA, USA) (Seitz *et al.*, 1990). Stable isotope analysis of <sup>13</sup>C/<sup>12</sup>C in gas samples was performed using a gas chromatograph combustion isotope ratio mass spectrometer (GC-C-IRMS) system (Thermoquest, Bremen, Germany). For principle operation see Brand (1996). Briefly, after conversion of CH<sub>4</sub> to CO<sub>2</sub> in the Finnigan Standard GC Combustion Interface III, isotope ratios were detected with the IRMS (Finnigan MAT model delta plus, Thermoquest, Bremen, Germany) (Fey *et al.*, 2004). Reference gas was CO<sub>2</sub> (99.998% purity; Messer-Griessheim, Düsseldorf, Germany), calibrated with the working standard methylstearate (Merck, Darmstadt, Germany). The latter was intercalibrated at the Max-Planck-Institut für Biogeochemie, Jena, Germany (courtesy of Dr W. A. Brand) against NBS22 and USGS 24 and reported in the delta notation vs. V-PDB:

$$\delta^{13}\text{C} = 10^3(R_{\text{sa}}/R_{\text{st}} - 1)$$

with  $R$

$$= {}^{13}\text{C}/{}^{12}\text{C} \text{ of sample (sa) and standard (st), respectively.}$$

The precision of repeated analysis was  $\pm 0.2\text{‰}$  when 1.3 nmol CH<sub>4</sub> was injected.

### Calculations

Fractionation factors for methanogenesis from H<sub>2</sub>/CO<sub>2</sub> are defined according to Hayes (1993):

$$\alpha_{\text{CO}_2\text{-CH}_4} = (\delta^{13}\text{C}_{\text{CO}_2} + 1000)/(\delta^{13}\text{C}_{\text{CH}_4} + 1000). \quad (1)$$

The isotopic signature for a newly formed CH<sub>4</sub> ( $\delta_n$ ) was calculated from the isotopic signatures at two time

points  $t=1$  ( $\delta_1$ ) and  $t=2$  ( $\delta_2$ ) by the following mass balance equation:

$$\delta_2 = f_n \delta_n + (1 - f_n) \delta_1 \quad (2)$$

with  $f_n$  being the fraction of the newly formed CH<sub>4</sub> relative to the total at  $t=2$ .

Gibbs free energies for methanogenesis in the methanogenic system were calculated from the standard Gibbs free energies of formation (Thauer *et al.*, 1977) and the Nernst equation:

$$\Delta G = \Delta G^\circ + RT \ln \frac{p_{\text{CH}_4}}{p_{\text{H}_2}^4 p_{\text{CO}_2}}. \quad (3)$$

$\Delta G^\circ$  was corrected for a temperature of 25 °C using the Van't Hoff equation.  $\alpha$  was corrected following the algorithm given by Whiticar *et al.* (1986), which is based on carbon isotope exchange equilibria (Richet *et al.*, 1977):

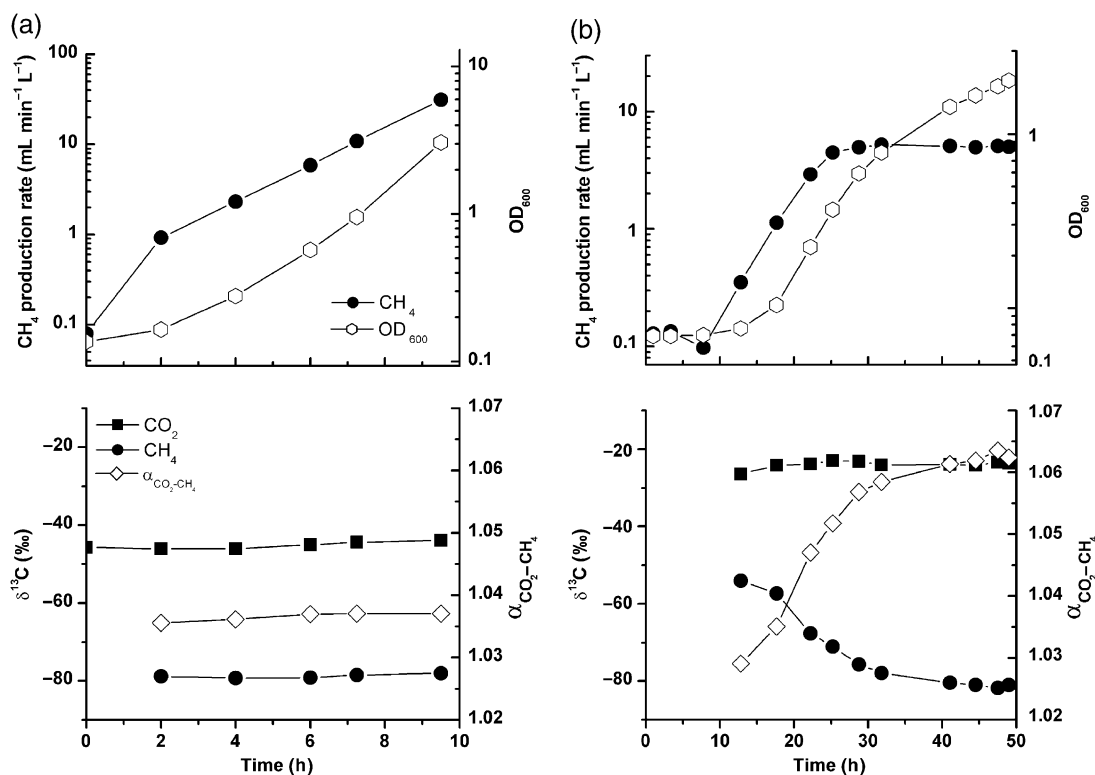
$$10^2 \ln \alpha_T = 2.92(10^3/T) - 2.96, \quad (4)$$

where  $T$  is the formation temperature in K. Correction was then made by the difference between the  $\alpha_T$  calculated for 25 °C and the growth temperatures of the cultures (adjustment of  $\alpha_{\text{CO}_2\text{-CH}_4}$  was 0.0017 for 30 °C, 0.0041 for 37 °C, and 0.0123 for 65 °C).

## Results

### Pure culture experiments

Pure cultures of *M. marburgensis* were grown in a flow-through batch bioreactor on different H<sub>2</sub>/CO<sub>2</sub>/N<sub>2</sub> gas mixtures (80/20/0% and 5/20/75%; Fig. 1). Almost parallel increase of optical density and CH<sub>4</sub> concentration indicated nonlimited growth conditions in the high-H<sub>2</sub> reactor (Fig. 1a), whereas in the low-H<sub>2</sub> reactor (Fig. 1b) two phases of exponential growth were displayed. In the first 30 h the CH<sub>4</sub> production rate and OD<sub>600</sub> increased exponentially (nonlimited growth). After that, however, the growth rate decreased, and the increase in CH<sub>4</sub> production rate declined and stabilized at 5 mL min<sup>-1</sup> (litre of H<sub>2</sub>/CO<sub>2</sub>/N<sub>2</sub>)<sup>-1</sup>. Such behavior is typical for hydrogenotrophic growth of methanogens, which uncouple CH<sub>4</sub> production from biomass production when the supply of H<sub>2</sub> is not limiting (Schönheit *et al.*, 1980). During transition of the CH<sub>4</sub> production rate to a stable value the H<sub>2</sub> partial pressures ( $p_{\text{H}_2}$ ) measured in the gas exhaust decreased and stabilized after 30 h (data not shown), demonstrating that H<sub>2</sub> limited growth conditions established. Values of  $\alpha_{\text{CO}_2\text{-CH}_4}$  ranged between 1.034 and 1.039 under nonlimited conditions (Fig. 1a, b), and increased in the reactor with low H<sub>2</sub> concentration (Fig. 1b) to constant values between 1.062 and 1.064, when H<sub>2</sub> became limiting.



**Fig. 1** Growth curve and  $\text{CH}_4$  production of *Methanothermobacter marburgensis* in flow-through bioreactor experiments with  $\text{H}_2/\text{CO}_2/\text{N}_2$  gas mixtures of (a) 80/20/0% and (b) 5/20/75% (upper diagrams). Time course of  $\delta^{13}\text{C}_{\text{CH}_4}$ ,  $\delta^{13}\text{C}_{\text{CO}_2}$ , and  $\alpha$  is shown in the lower diagrams of (a) and (b). Rapid gas flow guaranteed that  $\leq 3\%$  of the  $\text{CO}_2$  supplied was converted to  $\text{CH}_4$ . Unlimited availability of  $\text{H}_2$  (a) resulted in low and stable  $\alpha$ -values, whereas limitation of  $\text{H}_2$  (b), as indicated by slow growth and stagnant  $\text{CH}_4$  production rate during the late growth phase, resulted in high  $\alpha$ -values.

### Coculture experiments

Influence of  $p_{\text{H}_2}$  on carbon isotope fractionation was then tested in a closed system, where equilibrium of gases between headspace and liquid phase establishes. To create different  $p_{\text{H}_2}$  growth conditions for hydrogenotrophic methanogens, syntrophic (wide range of  $p_{\text{H}_2}$ ), and obligate syntrophic cocultures (narrow  $p_{\text{H}_2}$  range) were chosen for our experiments. In syntrophic cocultures of *C. papyrosolvans* and *M. bryantii* glucose and cellobiose were completely fermented by *C. papyrosolvans* within 200 h to acetate, lactate, ethanol, formate,  $\text{H}_2$ , and  $\text{CO}_2$ . Only  $\text{H}_2$  and  $\text{CO}_2$  were further converted to  $\text{CH}_4$  by the methanogenic archaeon *M. bryantii* and accumulated exponentially in the system (Fig. 2). Partial pressures of the intermediate  $\text{H}_2$  increased transiently to more than 2.5 kPa (Fig. 2). Cellulose, by contrast, allowed only a relatively low fermentation rate limited by the hydrolysis of cellulose. In these cultures, hydrogenotrophic  $\text{CH}_4$  production kept the  $p_{\text{H}_2}$  low ( $p_{\text{H}_2} < 0.1$  kPa) and  $\text{CH}_4$  was linearly produced. Carbon isotope composition of  $\text{CO}_2$  and newly formed  $\text{CH}_4$ , (Eqn (2)), from which  $\alpha_{\text{CO}_2-\text{CH}_4}$  was calculated, (Eqn

(1)), is shown in Fig 2. Calculation of  $\delta^{13}\text{C}$  of  $\text{CH}_4$  was made evaluating each data point separately, as the determination is very sensitive, if changes of the measured  $\delta^{13}\text{C}$  of  $\text{CH}_4$  (accumulated) and of the  $\text{CH}_4$  partial pressure are small. Changes of isotope ratio and concentration that were close to the analytical error of GC-IRMS or GC-FID measurement were therefore not included in the data evaluation. In the cocultures growing on glucose or cellobiose,  $\text{CH}_4$  formed by *M. bryantii* at high  $p_{\text{H}_2}$  exhibited low carbon isotope fractionation ( $\alpha_{\text{CO}_2-\text{CH}_4} < 1.04$ ), whereas at low  $p_{\text{H}_2}$   $\alpha$ -values increased. In contrast, on cellulose,  $\alpha$ -values were relatively high from the beginning on ( $\alpha_{\text{CO}_2-\text{CH}_4} = 1.044$ ) and stabilized after 500 h at  $\alpha_{\text{CO}_2-\text{CH}_4} = 1.076$  with  $p_{\text{H}_2} = 10$  Pa (Fig. 2).

Additional syntrophic cocultures (*C. papyrosolvans* with *M. hungatei*, *A. woodii* with *M. bryantii*, and *A. woodii* with *M. hungatei*) were analysed the same way. Trends in these cocultures were the same as shown in Fig. 2 for *C. papyrosolvans* and *M. bryantii* and are, therefore, not shown individually, but are later included in the data analysis. Isotope fractionation at very low  $p_{\text{H}_2}$  was investigated with the obligate syntrophic



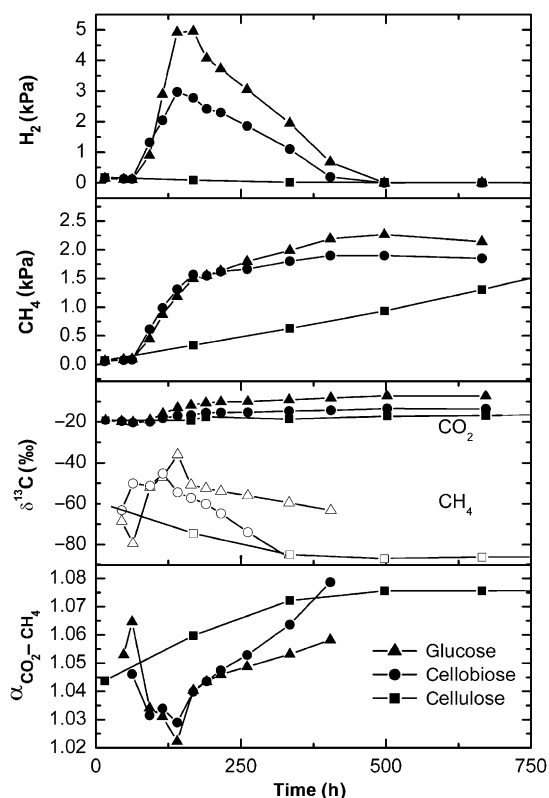


Fig. 2 Change of fractionation factor  $\alpha$ ,  $\delta^{13}\text{C}_{\text{CH}_4}$ ,  $\delta^{13}\text{C}_{\text{CO}_2}$ , and partial pressures of  $\text{H}_2$  and  $\text{CH}_4$  in a coculture of *Clostridium papyrosolvens* and *Methanobacterium bryantii*. High  $p_{\text{H}_2}$  correspond to low  $\alpha$  and vice versa. With cellulose, *M. bryantii* operated close to its threshold for  $\text{H}_2$  exhibiting very high isotope fractionation.

cocultures of *S. fumaroxidans* coupled to either *M. hungatei* or *M. formicicum*. Propionate oxidation by *S. fumaroxidans* was only possible at low  $p_{\text{H}_2}$  equivalent to a  $\Delta G$  of about  $-15 \text{ kJ mol}^{-1}$ , similarly as observed before (Scholten & Conrad, 2000). This low  $p_{\text{H}_2}$  was maintained by the methanogen, which, in turn, was also only provided with little free energy ( $\Delta G \approx -40$  to  $-20 \text{ kJ mol}^{-1}$ ). In contrast, the cocultures, which were not obligate syntrophic, operated at a higher  $p_{\text{H}_2}$ , and thus allowed generally more free energy ( $\Delta G \approx -100$  to  $-50 \text{ kJ mol}^{-1}$ ) to the methanogenic partner.

#### Data analysis of cocultures

Although each individual coculture showed a good correlation between the values of  $p_{\text{H}_2}$  and  $\alpha_{\text{CO}_2-\text{CH}_4}$  measured during the course of growth ( $r^2 > 0.82$ ), there was no general relationship between  $p_{\text{H}_2}$  and  $\alpha_{\text{CO}_2-\text{CH}_4}$  across all the different coculture experiments. Instead, a good correlation was found between the catabolic  $\Delta G$  of the methanogenic reaction and the  $\alpha_{\text{CO}_2-\text{CH}_4}$  of the reaction determined in the same incubation (Figs 3

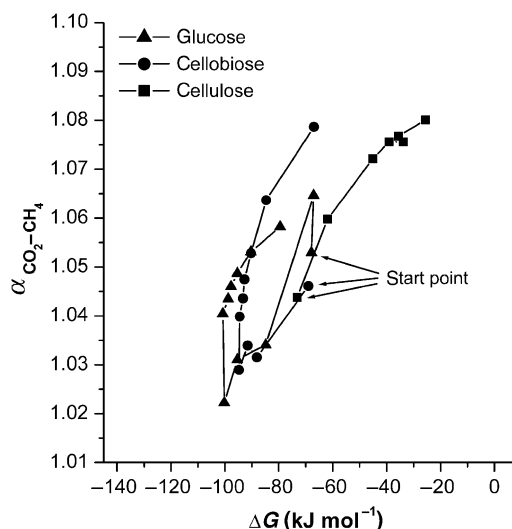
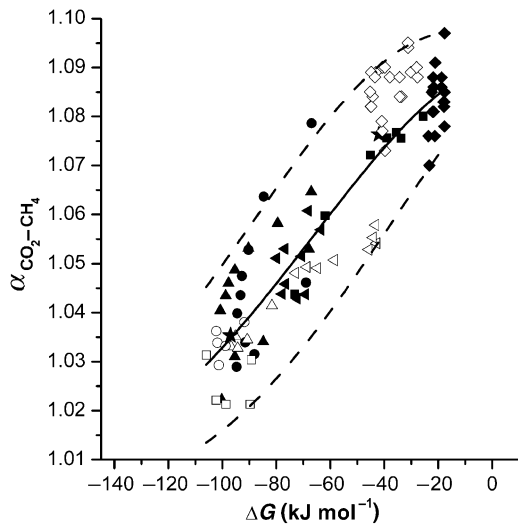


Fig. 3  $\alpha$ - $\Delta G$  relation in coculture of *Clostridium papyrosolvens* and *Methanobacterium bryantii* growing on different substrates, process and isotope data of which are shown in Fig. 2.  $\Delta G$  and  $\alpha_{\text{CO}_2-\text{CH}_4}$  well correlate and show a decrease and later increase for glucose and cellobiose, whereas for cellulose there is a steady increase.

and 4). Values of  $\Delta G$  were calculated of the actual conditions using measured  $p_{\text{H}_2}$ ,  $p_{\text{CH}_4}$ , and  $p_{\text{CO}_2}$ , and the Nernst (Eqn (3)). Figure 3 shows the relation for the coculture of *C. papyrosolvens* and *M. bryantii* (Fig. 2). After inoculation  $\Delta G$  for the coculture was  $-70 \text{ kJ mol}^{-1}$ . During the course of the experiment  $\Delta G$  decreased in cocultures with glucose and cellobiose to  $-100 \text{ kJ mol}^{-1}$ , concomitant with a decrease in  $\alpha_{\text{CO}_2-\text{CH}_4}$  (1.02–1.03). With decrease in  $p_{\text{H}_2}$  and therefore increase in  $\Delta G$ ,  $\alpha_{\text{CO}_2-\text{CH}_4}$  also increased again. In the coculture with cellulose, where  $\Delta G$  steadily increased with time,  $\alpha_{\text{CO}_2-\text{CH}_4}$  increased simultaneously to values up to 1.08.

Figure 4 summarizes the results of all the different syntrophic and obligate syntrophic cocultures tested. We calculated the values of  $\alpha_{\text{CO}_2-\text{CH}_4}$  and  $\Delta G$  separately for each replicate of the cocultures to increase precision. Syntrophic cocultures (with *C. papyrosolvens* or *A. woodii* as fermenting partner) were active within a wide range of  $\Delta G$ , as well as  $\alpha_{\text{CO}_2-\text{CH}_4}$ , whereas obligate syntrophic cocultures (*S. fumaroxidans* as fermenting partner) could only operate within a narrow range of  $\Delta G$  ( $-45$  to  $-17 \text{ kJ mol}^{-1}$ ). Both, *M. bryantii* and *M. hungatei*, displayed the relation over a wide range of  $\Delta G$  and  $\alpha_{\text{CO}_2-\text{CH}_4}$ . *M. formicicum* was only used in the obligate syntrophic cocultures and was, therefore, restricted to the high  $\Delta G$  values, characteristic for these cultures. Pure culture data of *M. marburgensis* (Fig. 1) could not be included in the figure because of lack of the *in situ*  $p_{\text{H}_2}$  in the flow-through system necessary for the calcu-



**Fig. 4**  $\alpha$ - $\Delta G$  relation in cocultures of  $H_2$ -producing fermenting and  $H_2$ -consuming methanogenic microorganisms; the line is a fit (using Origin Microcal™) to the Gauss function ( $y = y_0 + A \exp[-(x - x_c)^2 / (2w^2)] = 1 + 0.0919 \exp[-(x - 11.8376)^2 / 12170]$ ).  $\blacktriangle$ , *Clostridium papyrosolvens*/*Methanobacterium bryantii* (glucose);  $\bullet$ , *C. papyrosolvens*/*M. bryantii* (cellobiose);  $\blacksquare$ , *C. papyrosolvens*/*M. bryantii* (cellulose);  $\blacktriangleleft$ , *Acetobacterium woodii*/*M. bryantii* (glucose);  $\triangle$ , *C. papyrosolvens*/*Methanospirillum hungatei* (glucose);  $\circ$ , *C. papyrosolvens*/*M. hungatei* (cellobiose);  $\square$ , *C. papyrosolvens*/*M. hungatei* (cellulose);  $\triangleleft$ , *A. woodii*/*M. hungatei* (glucose);  $\blacklozenge$ , *Syntrophobacter fumaroxidans*/*Methanobacterium formicicum* (propionate);  $\diamond$ , *S. fumaroxidans*/*M. hungatei* (propionate);  $\star$ , *M. marburgensis* ( $H_2/CO_2$ ; data from Valentine *et al.* (2004))

lation of  $\Delta G$ . Yet, two data points from Valentine *et al.* (2004) are included and they fit the observed  $\alpha$ - $\Delta G$  relation well. A Gauss function, Eqn (5), was used to fit the data (justification see Discussion) and resulted in a goodness of fit of  $r^2 = 0.77$ :

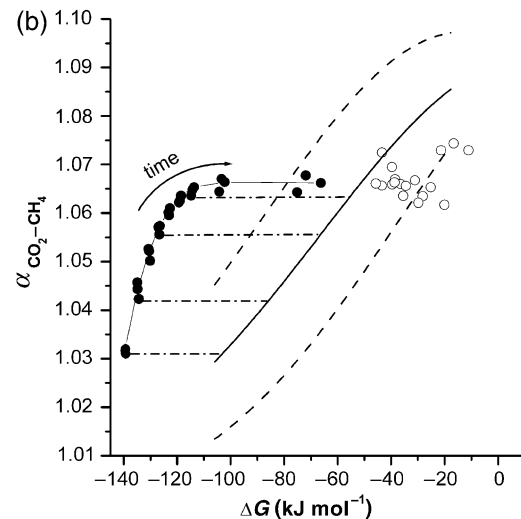
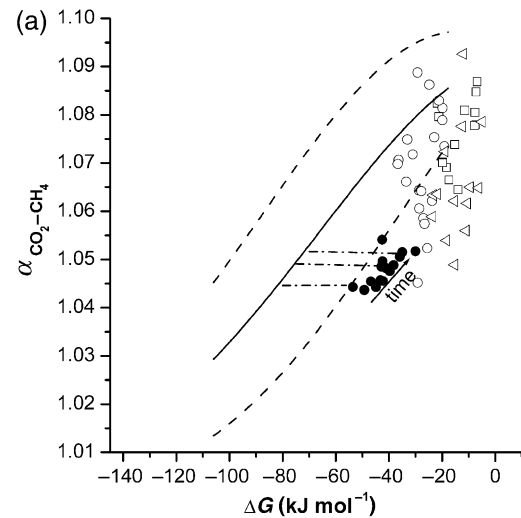
$$y = y_0 + A \exp[-(x - x_c)^2 / (2w^2)] \quad (5)$$

with  $y_0 = 1$ ;  $A = 0.0919 \pm 0.0104$ ;  $x_c = 11.8376 \pm 23.5222$ ;  $w = 78.0067 \pm 14.1127$ .

Because of the individual treatment of replicates no errors bars can be shown for the data points, but we calculated the 95% confidence range for the  $\alpha$ - $\Delta G$  relation (included in Fig. 4).

#### Incubations of environmental samples

To test the implication of the observed  $\alpha$ - $\Delta G$  relation for natural methanogenic systems, samples from different environments were incubated under anoxic conditions. As in the cocultures, only time points with active methanogenesis were used in the data analysis, which was performed as described for the cocultures. In the  $\alpha$ - $\Delta G$  relation diagrams of the environmental incubations



**Fig. 5**  $\alpha$ - $\Delta G$  relation in different environmental samples including regression curve and confidence lines from Fig. 4; arrows indicate the time course of incubation, and dash-dotted lines show the difference between the apparent  $\Delta G$  (symbols) and the actual  $\Delta G$  (regression curve) of hydrogenotrophic methanogenesis. The actual  $\Delta G$  available to methanogens in their environmental habitat was up to  $60 \text{ kJ mol}^{-1}$  more negative than that suggested from the apparent  $\Delta G$  determined from bulk measurement of  $p_{H_2}$  (difference shown by horizontal dash-dotted lines). However, in rumen fluid incubated under  $H_2/CO_2$  it was more positive. (a) Rice field soil and peat ( $\bullet$ , rice field soil;  $\square$ , peat OLI;  $\circ$ , peat MES;  $\triangleleft$ , peat OMB). (b) Rumen fluid ( $\circ$ , rumen ( $N_2$ );  $\bullet$ , rumen ( $H_2/CO_2$ )).

(Fig. 5) the regression curve of the culture experiments and its 95% confidence lines are also included. In incubations of rice field soil and peat, acetoclastic methanogenesis was inhibited with methyl fluoride (Janssen & Frenzel, 1997) to avoid changes of the  $\delta^{13}C$  of  $CH_4$  by  $CH_4$  produced from acetate (Fig. 5a). This approach has recently been proposed for the determi-

nation of  $\alpha_{\text{CO}_2\text{-CH}_4}$ , (Conrad, 2005) and successfully tested in eutrophic lake sediment (Chan *et al.*, 2005) and rice field soil (unpublished data). In rice field soil,  $\Delta G$  ( $-54$  to  $-30$   $\text{kJ mol}^{-1}$ ) and  $\alpha_{\text{CO}_2\text{-CH}_4}$  (1.043–1.053) increased simultaneously with time. The peat incubations plotted over a wide range of  $\alpha_{\text{CO}_2\text{-CH}_4}$  (1.045–1.087) and at high  $\Delta G$  ( $-37$  to  $-7$   $\text{kJ mol}^{-1}$ ). In the rice field soil and peat system, data followed the  $\alpha$ - $\Delta G$  relation only for  $\alpha_{\text{CO}_2\text{-CH}_4} > 1.065$ , whereas for  $\alpha_{\text{CO}_2\text{-CH}_4} < 1.065$ , data were on the right side of the  $\alpha$ - $\Delta G$  relation. Using the  $\alpha$ - $\Delta G$  relation and the  $p_{\text{H}_2}$  measured in the environmental samples would thus predict (via  $\Delta G$  that decreases with  $p_{\text{H}_2}$ ) much higher  $\alpha$  than actually measured. On the other hand, the  $\alpha$ - $\Delta G$  relation and the measured  $\alpha$  would predict much lower (more negative) values of  $\Delta G$  than calculated from measured  $p_{\text{H}_2}$ .

We also used rumen fluid, as in this environment  $\text{CH}_4$  is almost exclusively produced from  $\text{H}_2/\text{CO}_2$  (Sharp *et al.*, 1998) and, thus, inhibition of acetoclastic methanogenesis with methyl fluoride is not necessary. Incubation of rumen fluid was conducted under  $\text{N}_2$  and  $\text{H}_2/\text{CO}_2$  headspace (Fig. 5b). For incubations under  $\text{N}_2$  headspace data clustered in a relatively confined area ( $\Delta G = -11$  to  $-46$   $\text{kJ mol}^{-1}$ ;  $\alpha_{\text{CO}_2\text{-CH}_4} = 1.061$ – $1.074$ ) within the 95% confidence limit of the  $\alpha$ - $\Delta G$  relation. In contrast, the  $\text{H}_2/\text{CO}_2$  headspace created very low  $\Delta G$  values at the beginning ( $-140$   $\text{kJ mol}^{-1}$ ), which continuously decreased, when  $\text{CH}_4$  was produced by hydrogenotrophic archaea. Values of  $\alpha_{\text{CO}_2\text{-CH}_4}$  strongly increased initially (1.032–1.067) and stabilized at a similar  $\alpha_{\text{CO}_2\text{-CH}_4}$  as observed in the incubation under  $\text{N}_2$ . Simultaneously, bulk  $p_{\text{H}_2}$  decreased so that the measured  $\Delta G$  gradually changed from  $-145$  to  $-75$   $\text{kJ mol}^{-1}$ , similarly as observed in rumen fluid incubated under a  $\text{N}_2$  atmosphere. Hence, for rumen incubated under  $\text{H}_2/\text{CO}_2$ , data first plotted on the left side of the  $\alpha$ - $\Delta G$  relation curve, but later approached and finally crossed the curve reaching the cluster of the rumen incubation under  $\text{N}_2$ . Again, the  $\alpha$ - $\Delta G$  relation together with the measured  $p_{\text{H}_2}$  did not allow prediction of  $\alpha$ . However, using the  $\alpha$ - $\Delta G$  relation together with the  $\alpha$  measured after stabilization would predict a reasonable value of  $\Delta G$  available to the hydrogenotrophic methanogens.

## Discussion

### Pure and coculture experiments

Different hydrogenotrophic methanogenic archaea were grown under varying growth conditions to elucidate the factors causing the fractionation of carbon isotopes during  $\text{CH}_4$  production from  $\text{H}_2/\text{CO}_2$ . In pure

cultures of *M. marburgensis* (Fig. 1) we could show, that  $\alpha_{\text{CO}_2\text{-CH}_4}$  depended on the  $p_{\text{H}_2}$ . Values of  $\alpha_{\text{CO}_2\text{-CH}_4}$  increased with decreasing  $p_{\text{H}_2}$ , confirming the results by Valentine *et al.* (2004). To quantify the observed correlation we used cocultures in closed systems, where partial pressures of the gases involved in the methanogenic reaction could be quantified with a high precision.  $\alpha_{\text{CO}_2\text{-CH}_4}$  produced from *M. bryantii* in coculture with *C. papyrosolvens* at high  $p_{\text{H}_2}$  was similar as reported for pure cultures of *Methanobacterium* spp. (Belyaev *et al.*, 1983) when grown on  $\text{H}_2/\text{CO}_2$  at a ratio of 4:1 as required for stoichiometric conversion to  $\text{CH}_4$ . However, with decreasing  $p_{\text{H}_2}$ , as for *M. marburgensis* in the flow-through reactor,  $\alpha_{\text{CO}_2\text{-CH}_4}$  strongly increased. The cocultures demonstrated, that the relation of  $\alpha$  to  $p_{\text{H}_2}$  was not specific to pure culture data, and that the coculture experiments were not biased by the different mode of  $\text{H}_2/\text{CO}_2$  gas transfer (i.e. between different microbes within the culture medium or between the gas phase and the culture medium). In addition, we found that the different isotopic composition of glucose and cellobiose,  $\delta^{13}\text{C} = -10.85 \pm 0.05\text{‰}$  and  $-25.23 \pm 0.47\text{‰}$ , respectively, resulted in a different isotopic composition of the pool of  $\text{CO}_2$ , but did not influence the value of  $\alpha_{\text{CO}_2\text{-CH}_4}$  (Fig. 2). This observation shows that isotopic equilibrium was reached between the  $\text{CO}_2$  produced by fermentation and the  $\text{CO}_2$  in the medium.

### $\alpha$ - $\Delta G$ relation

Although we found a good correlation of  $\alpha_{\text{CO}_2\text{-CH}_4}$  with  $p_{\text{H}_2}$  for the single cocultures, the lack of a general dependency across the different coculture experiments indicated that  $p_{\text{H}_2}$  was not the only factor determining  $\alpha_{\text{CO}_2\text{-CH}_4}$ . The catabolic  $\Delta G$ , however, did yield for all data a good correlation ( $r^2 = 0.77$ ). Valentine *et al.* (2004) recently observed that a culture of *M. marburgensis* exhibited a larger fractionation ( $\alpha_{\text{CO}_2\text{-CH}_4} = 1.064$ ) at low (310 Pa) and a lower fractionation ( $\alpha_{\text{CO}_2\text{-CH}_4} = 1.031$ ) at high (80 000 Pa)  $p_{\text{H}_2}$  (data points included in Fig. 4), and proposed that this difference is because of the extent of enzymatic reversibility that is controlled by the  $\Delta G$  of the methanogenic reaction. They further suggested that the differential reversibility would act in concert with the intracellular availability of  $\text{CO}_2$ . High metabolic rates would decrease the intracellular concentration of  $\text{CO}_2$ , effectively creating an irreversible flux of  $\text{CO}_2$  to  $\text{CH}_4$ . If then  $\text{CO}_2$  was quantitatively consumed, consequently, no fractionation would occur by methanogenesis ( $\alpha_{\text{CO}_2\text{-CH}_4} = 1.0$ ) and only a slight fractionation by transport into the cell would be observed.

Our results are consistent with this hypothesis demonstrating that  $\alpha$  is indeed a function of the  $\Delta G$ . With respect to energetics of methanogenesis, it is important

to note that methanogens activate  $\text{CO}_2$  by forming formyl methanofuran. This reaction is endergonic under standard conditions and becomes exergonic only when coupled to the late reaction steps that conserve energy (e.g. methyl transferase and heterodisulfide reductase; Thauer, 1998). A low value of the  $\Delta G$  (large negative values) of methanogenesis from  $\text{H}_2 + \text{CO}_2$  thus facilitates activation of  $\text{CO}_2$  by also lowering the  $\Delta G$  of this single reaction, decreases reversibility, and results in low  $\alpha_{\text{CO}_2\text{-CH}_4}$ . A high value of the  $\Delta G$  (low negative values), on the other hand, impedes activation of  $\text{CO}_2$ , increases  $\Delta G$  of this reaction, and results in high  $\alpha_{\text{CO}_2\text{-CH}_4}$ . For activation of  $\text{CO}_2$  a certain amount of free energy is always required, since methanogenesis can only proceed if this step is made exergonic. With increase of  $\Delta G$ , less free energy will be available to drive each step of the multireaction process, further increasing the reversibility and, therefore, the fractionation of the reactions. Finally a maximum of  $\alpha$  is reached at the thermodynamic limit of the microorganisms (Fig. 4), which in theory is equivalent to  $\frac{1}{4} - \frac{1}{3}$  ATP or about  $-15$  to  $-25 \text{ kJ mol}^{-1}$  for mesophilic methanogens (Yao & Conrad, 1999; Hoehler *et al.*, 2001; Schink & Stams, 2003; Müller, 2004). As methanogenesis cannot proceed above  $\Delta G = 0$ , a semi-Gauss curve was used for correlation. The maximum fractionation is solely determined by  $\Delta G$ , as isotopic equilibrium of the  $\text{CO}_2$  pools outside and inside the cell can be assumed at the low metabolic rates associated with high  $\Delta G$ . However, irreversible  $\text{CO}_2$  flow becomes more important with decreasing  $\Delta G$  and the concomitant increase of the metabolic rate, and finally governs the  $\alpha$ - $\Delta G$  relation. At very low  $\Delta G$  the hypothetical limit of zero fractionation ( $\alpha_{\text{CO}_2\text{-CH}_4}$ ) will be reached, if all  $\text{CO}_2$  entering the cell is quantitatively metabolized. The semi-Gauss curve accounts for this boundary condition predicted by irreversible  $\text{CO}_2$  flow.

As explained above, the observed relation (Fig. 4) is partly explained by the reversibility hypothesis depending on the free energy of the reaction (Valentine *et al.*, 2004). For H-isotopes a similar correlation (i.e. a free-energy dependence of the kinetic isotope effect) was found, showing a bell-shaped curve with its maximum at  $\Delta G = 0$  (Sühnel, 1990; Cook, 1991). There, the experimentally observed correlations in proton-transfer reactions could be theoretically explained both by the Marcus and quantum-statistical mechanical model. To our knowledge similar investigations on carbon isotopes have not been performed yet, so this interesting finding for H-isotopes should be evaluated for the element carbon. In addition, methanogenesis is a multireaction process. This further complicates the calculation of carbon isotope fractionation, as concentrations of reactants and products of each single reaction of methanogenesis have to be known to determine  $\Delta G$ . There-

fore, we have to point out that the similarity between the theoretical free energy dependence of the kinetic isotope effect observed for H-isotopes and the  $\alpha$ - $\Delta G$  relation observed for hydrogenotrophic methanogenesis in our study is probably only circumstantial. However, the observed relationship between  $\alpha_{\text{CO}_2\text{-CH}_4}$  and  $\Delta G$  presented here appeared to be valid for all the four different methanogenic microbial species (from three genera and two families) tested (Fig. 4), emphasizing that it is a general feature of C-isotope fractionation in hydrogenotrophic methanogenesis. Although two of the tested methanogens (*M. bryantii* and *M. hungatei*) follow the  $\alpha$ - $\Delta G$  relation over the whole range of  $\alpha_{\text{CO}_2\text{-CH}_4}$  and  $\Delta G$  observed, it cannot be dismissed that specifics of enzymes – like individual binding sites and activation energies – are different in other hydrogenotrophic methanogens and might change the  $\alpha$ - $\Delta G$  relation to a certain extent. This could be the case, if for example *K*- and *r*-strategists had developed differently fractionating sets of enzymes depending on  $p_{\text{H}_2}$ . However, regarding carbon isotope fractionation, none of this has been reported yet, and the archaea tested in this study are also typical representatives of hydrogenotrophic methanogens in natural environments (Grosskopf *et al.*, 1998; Koizumi *et al.*, 2004; Chan *et al.*, 2005).

#### *Application of the $\alpha$ - $\Delta G$ relation to environmental systems*

Studies determining  $\Delta G$  in natural systems from measurements of  $p_{\text{H}_2}$ ,  $p_{\text{CH}_4}$  and  $p_{\text{CO}_2}$  have indicated that hydrogenotrophic methanogens operate at the thermodynamic limit ( $\Delta G = -15$  and  $-25 \text{ kJ mol}^{-1}$ ). Although  $\text{CH}_4$  formation may be active at  $\Delta G \approx 0$  (Jackson & McInerney, 2002), such activity does not support growth, which in chemostat culture required  $\Delta G \approx -36 \text{ kJ mol}^{-1}$  (Seitz *et al.*, 1990). The severe energetic limitation in natural systems is in contrast to observations that the composition of methanogenic communities in the environment can change with time implying growth (Peters & Conrad, 1996; Lueders & Friedrich, 2002; Krüger *et al.*, 2005). Determination of *in situ*  $\Delta G$  is dependent on the measurement of realistic  $p_{\text{H}_2}$ . The actual *in situ*  $p_{\text{H}_2}$  is at the microsite where the methanogens live, which, however, may not be represented by the  $p_{\text{H}_2}$  that is typically measured in the bulk fluid or in the equilibrated gas phase of the whole environmental sample. In natural environments, methanogens are closely associated with  $\text{H}_2$ -forming bacteria so that interspecies- $\text{H}_2$ -transfer can occur between juxtaposed partners (Conrad *et al.*, 1985; Krylova & Conrad, 1998; Hoehler *et al.*, 2001). Hence, methanogens may experience higher  $p_{\text{H}_2}$  (i.e. more negative  $\Delta G$ ) than



deduced from bulk measurements, which may overestimate the effective  $\Delta G$  (i.e. less negative).

Having an alternative way to determine  $\Delta G$  (i.e. the  $\alpha$ - $\Delta G$  relation), we tested the hypothesis that the  $\Delta G$  effective for the methanogens in natural systems may be lower than expected from  $p_{\text{H}_2}$  measurements. During incubation of rumen fluid the  $p_{\text{H}_2}$  experienced by the methanogens is probably different than that measured in the headspace, as microbial aggregates are formed (see above) and rumen methanogens to a large part intimately associate with protozoa (Sharp *et al.*, 1998) creating  $p_{\text{H}_2}$  gradients.  $p_{\text{H}_2}$  gradients lead to a measured  $p_{\text{H}_2}$  that does not agree with the  $p_{\text{H}_2}$  experienced by the methanogenic organism. In the case of  $\text{H}_2/\text{CO}_2$  in the headspace, a lower  $p_{\text{H}_2}$  was present in close proximity to the methanogens, which consumed  $\text{H}_2$ . In the case of a  $\text{N}_2$  headspace, it was the opposite, since the products of fermentation, which are the substrates of methanogenesis, were produced by fermenting organisms next to the methanogenic archaea. The methanogens, therefore, had already depleted the  $p_{\text{H}_2}$  before it could be quantified in the bulk fluid or the headspace in equilibrium with the bulk liquid. An exception to this rule was some of the peat samples (Fig. 5a;  $\alpha_{\text{CO}_2\text{-CH}_4} > 1.065$ ), in which degradable organic substrates and thus  $\text{H}_2$  production rates were presumably low. In these incubations metabolic activity of methanogens kept the  $p_{\text{H}_2}$  close to the threshold of hydrogenotrophic methanogenesis, that the  $\Delta G$  determined from bulk  $p_{\text{H}_2}$  measurements approached the  $\alpha$ - $\Delta G$  relation close to the maximum  $\alpha$  (i.e. a gradient between the  $p_{\text{H}_2}$  experienced by the archaea and the  $p_{\text{H}_2}$  measured in the headspace was reduced to a minimum; Fig. 5a). Collectively, these results are consistent with our hypothesis that  $\alpha_{\text{CO}_2\text{-CH}_4}$  is a good predictor of the  $\Delta G$  available to the methanogens *in situ*. The values of  $\Delta G$  determined the conventional way, (Eqn (3)), and using the  $\alpha$ - $\Delta G$  relation are exemplarily shown by horizontal dash-dotted lines for the incubation of rumen under  $\text{H}_2/\text{CO}_2$  headspace and rice field soil and show a difference of up to  $60 \text{ kJ mol}^{-1}$  (Fig. 5). We propose that the more reliable  $\Delta G$  values were those extrapolated from the  $\alpha$ - $\Delta G$  relation, and that a large fraction of the methanogens actually experienced a more negative  $\Delta G$  than indicated by the measurement of bulk  $p_{\text{H}_2}$ .

The  $\alpha$ - $\Delta G$  relation found in our coculture experiments may overcome the problem of biased  $p_{\text{H}_2}$  measurements in natural methanogenic environments, as the average  $\Delta G$  available to the hydrogenotrophic methanogenic community can now be constrained by the C-isotope fractionation. The resulting *in situ*  $\Delta G$  values show that hydrogenotrophic methanogens often thrive at  $\Delta G$  of  $-40$  to  $-100 \text{ kJ mol}^{-1}$ . These  $\Delta G$  values are lower than those derived from measurements of  $p_{\text{H}_2}$  in the bulk of

environmental samples (i.e. about  $-20 \text{ kJ mol}^{-1}$  in anoxic rice field soil; Yao & Conrad, 1999), and are sufficient for the synthesis of 1/2 to 1 ATP per reaction. Thus, there is a chance to solve the dilemma of earlier observations where relatively high  $\Delta G$  values, hardly sufficient for active growth of methanogenic populations, were associated with changes in community composition or even net growth of methanogenic populations (Peters & Conrad, 1996; Lueders & Friedrich, 2002; Krüger *et al.*, 2005).

## Conclusions

We have shown that the fractionation of stable C-isotopes during hydrogenotrophic methanogenesis is related to the Gibbs free energy of catabolism, thus confirming the differential reversibility hypothesis of Valentine *et al.* (2004). The existence of a  $\alpha$ - $\Delta G$  relation that is common to all species of  $\text{H}_2/\text{CO}_2$ -consuming methanogens implies that the mechanistic basis for stable C-isotope fractionation is the same. This implication is quite likely, since the enzymology of hydrogenotrophic methanogenesis is rather uniform (Thauer, 1998), but should be confirmed by more detailed research.

We have also shown that the  $\alpha$ - $\Delta G$  relation can be used to determine the free energy available for hydrogenotrophic methanogenesis *in situ* without the need to measure the notoriously low  $\text{H}_2$  partial pressures, which may be biased. Comparison of  $\Delta G$  calculated from the  $\alpha$ - $\Delta G$  relation and by the traditional way may be used to interpret the microenvironment of methanogenic microbial consortia with respect to the  $\text{H}_2$  gradient experienced by the microbiota. Such a knowledge, even when approximate, helps judge the *in situ* living conditions of the methanogenic microbiota, a metabolic microbial group with high impact on global atmospheric chemistry and physics in the past (Kasting, 1993), the present (Lelieveld & Crutzen, 1992), and the future (Wang *et al.*, 2004).

Accepting a common  $\alpha$ - $\Delta G$  relation for hydrogenotrophic methanogenesis, this would have implications for the interpretation of stable isotope signatures measured in natural environments. Values of  $\delta^{13}\text{C}$  in environmental  $\text{CH}_4$  and  $\text{CO}_2$  are frequently interpreted with respect to the predominant path of  $\text{CH}_4$  formation. Calculating an apparent  $\alpha$  ( $\alpha_c = (\delta^{13}\text{CO}_2 + 10^3)/(\delta^{13}\text{CH}_4 + 10^3)$ ) it is generally assumed that low  $\alpha_c < 1.055$  and high  $\alpha_c > 1.065$  are indicative of the predominant operation of acetotrophic and hydrogenotrophic methanogenesis, respectively (Whiticar *et al.*, 1986; Whiticar, 1999). However, hydrogenotrophic methanogenesis may also cause low  $\alpha_c$ , if the methanogens operate in a microenvironment with plenty of free energy

available (e.g. eutrophic anoxic systems), in which H<sub>2</sub> production by fermenting bacteria is high. Hence, one should be more cautious when interpreting  $\delta^{13}\text{C}$  values without determining the fractionation factor of hydrogenotrophic methanogenesis separately.

In addition, the  $\alpha$ - $\Delta G$  relation explains the wide variation of  $\alpha_{\text{CO}_2\text{-CH}_4}$  in the environment. Although we cannot use the relation to precisely quantify C-isotope fractionation, we can better predict  $\alpha_{\text{CO}_2\text{-CH}_4}$  in different types of environments. We tested three environments (peat, rice field soil, and rumen), which are important for global CH<sub>4</sub>, showing that the relatively narrow range of energy conditions in these environments determines the observed range of  $\alpha$ -values. These findings agree with environmental studies, where low values of  $\alpha_{\text{CO}_2\text{-CH}_4}$  were found in the high- $p_{\text{H}_2}$  environment rumen ( $\alpha_{\text{CO}_2\text{-CH}_4} = 1.031$ ; Levin *et al.*, 1993), whereas under energy-limiting conditions as in bogs values of  $\alpha_{\text{CO}_2\text{-CH}_4}$  were close to 1.07 (Lansdown *et al.*, 1992; Avery *et al.*, 1999). Because of the relatively favorable energy conditions for hydrogenotrophic methanogens in rice field soils, values of  $\alpha$  there usually show intermediate isotope fractionation ( $\alpha_{\text{CO}_2\text{-CH}_4} = 1.045\text{--}1.060$ ; Sugimoto & Wada, 1993; Tyler *et al.*, 1997; Bilek *et al.*, 1999). With the  $\alpha$ - $\Delta G$  relation as theoretical background in combination with experimental data from methanogenic environments, we may better predict values of  $\alpha_{\text{CO}_2\text{-CH}_4}$  for CH<sub>4</sub>-producing environments for which no detailed isotope studies exist. Our results thus provide a mechanistic constraint for modelling the  $^{13}\text{C}$  flux from microbial sources of atmospheric CH<sub>4</sub>.

### Acknowledgements

We thank P. Claus for excellent technical assistance during analysis of  $^{13}\text{C}$  data and W. A. Brand for isotopic analysis of the carbohydrate substrates. This work was supported by the Research Council for Earth and Life Sciences (ALW) with financial aid from the Netherlands Organization for Scientific Research (NWO).

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