

1 **Evidence for methane production by the marine algae *Emiliana huxleyi***

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## 20 INTRODUCTION

21 Methane (CH<sub>4</sub>), the second important anthropogenic greenhouse gas after CO<sub>2</sub>, is the most  
22 abundant reduced organic compound in the atmosphere and plays a central role in  
23 atmospheric chemistry (IPCC, 2013; Kirschke et al., 2013; Lelieveld et al., 1998). The mixing  
24 ratio of CH<sub>4</sub> in the atmosphere has been increasing from pre-industrial values of around 715  
25 ppbv (parts per billion by volume) to about 1800 ppbv in 2010 (Kirschke et al., 2013). In total,  
26 annual CH<sub>4</sub> emissions from natural and anthropogenic sources amount to 500-600 Tg (10<sup>12</sup> g)  
27 yr<sup>-1</sup>. They derive from various terrestrial and aquatic sources and are balanced primarily by  
28 photochemical oxidation in the troposphere (≈ 80 %), diffusion into the stratosphere and  
29 microbial CH<sub>4</sub> oxidation in soils.

30 Natural sources of atmospheric CH<sub>4</sub> in the biosphere have until recently been attributed to  
31 originate solely from strictly anaerobic microbial processes in wetland soils and rice paddies,  
32 the intestines of termites and ruminants, human and agricultural waste, and from biomass  
33 burning, fossil fuel mining and geological sources including mud volcanoes, vents and seeps.  
34 However, more recent studies have suggested that terrestrial vegetation, fungi and mammals  
35 may also produce CH<sub>4</sub> without an input from methanogens and under aerobic conditions  
36 (Bruhn et al., 2012; Ghyczy et al., 2008; Keppler et al., 2006; Lenhart et al., 2012; Wang et al.,  
37 2013b; Liu et al., 2015). A fraction of these vegetation-derived emissions might be released  
38 directly by *in-situ* formation in plants (Bruhn et al., 2012; Keppler et al., 2009; Wang et al.,  
39 2013a), and it is now apparent that several pathways exist by which CH<sub>4</sub> is generated under  
40 aerobic conditions (Bruhn et al., 2014; Messenger et al., 2009; Wang et al., 2013b). Hence, the  
41 biogeochemical CH<sub>4</sub> cycle appears to be even more complex than previously thought.

42 In particular the biogeochemical cycle of CH<sub>4</sub> in the oceans is still far from being understood.  
43 The world's oceans are considered to be a minor source of CH<sub>4</sub> to the atmosphere with  
44 approximately 0.6-1.2 Tg CH<sub>4</sub> yr<sup>-1</sup> (Rhee et al., 2009). Concentrations of CH<sub>4</sub> in near-surface  
45 waters are often 5–75 % supersaturated with respect to the atmosphere implying a net flux  
46 from the ocean to the atmosphere (Conrad, 2009; Reeburgh, 2007; Scranton and Brewer,  
47 1977). Because the surface ocean is also saturated or slightly supersaturated with oxygen,  
48 which does not favor methanogenesis, the observed CH<sub>4</sub> supersaturation has been termed  
49 the oceanic methane paradox (Kiene, 1991). To explain the source of CH<sub>4</sub> in surface waters, it  
50 has been suggested that methanogenesis takes place in anoxic microenvironments of organic  
51 aggregates (Grossart et al., 2011; Karl and Tilbrook, 1994; Bogard et al., 2014), the guts of  
52 zooplankton or fish (de Angelis and Lee, 1994; Oremland, 1979) and inside bacterial cells  
53 (Damm et al., 2015). It has also been shown that opposite to the conventional view, some  
54 methanogens are remarkably tolerant to oxygen (Angel et al., 2011; Jarrell, 1985).

55 A potential substrate for methanogenesis in such anoxic microniches is  
56 dimethylsulphoniopropionate (DMSP) (Damm et al., 2008; Zindler et al., 2013, Damm et al.,  
57 2015), an algal osmolyte that is abundant in marine phytoplankton and serves as a precursor  
58 for dimethylsulphide (DMS) and dimethylsulphoxide (DMSO) (Stefels et al., 2007; Yoch, 2002)  
59 For example Zindler et al. (2013) measured concentrations of DMS, DMSP, DMSO, and CH<sub>4</sub>, as  
60 well as various phytoplankton marker pigments in the surface ocean along a north-south  
61 transit from Japan to Australia. Positive correlations between DMSP (dissolved) and CH<sub>4</sub>, and  
62 DMSO (particulate and total) and CH<sub>4</sub>, were found along the transit. Based on their data they  
63 concluded that DMSP and DMSO and/or their degradation products serve as substrates for  
64 methanogenic archaea in the western Pacific Ocean.

65 Damm et al. 2010 hypothesized that under N-limitation and a concomitant availability of  
66 phosphorus, marine bacteria use DMSP as a carbon source and thereby release CH<sub>4</sub> as a by-  
67 product and its production could yield energy under aerobic conditions. Methanethiol, a  
68 further potential degradation product of DMSP, may act as a direct precursor of methane in  
69 aerobic environments. By reason of thermodynamic calculations the authors considered that  
70 microorganism can yield energy from the pathway of methanethiol formation operating in its  
71 reverse direction, whereby methane is formed.

72 An alternative non-biological CH<sub>4</sub> formation pathway in seawater might occur via a  
73 photochemical pathway due to the formation of methyl radicals, however photochemical  
74 production of CH<sub>4</sub> in oceans is thought to be negligible under oxic conditions (Bange and Uher,  
75 2005).

76 In addition, Karl et al. (2008) suggested that CH<sub>4</sub> is produced aerobically as a by-product of  
77 methylphosphonate (MPn) decomposition when aerobic marine organisms use  
78 methylphosphonic acid as a source of phosphorus when inorganic sources of this element are  
79 limited. Furthermore, a mechanism has been identified that leads to the formation of CH<sub>4</sub>  
80 from MPn via enzyme-catalytic cleavage of the C-P bond (Kamat et al., 2013). The critical  
81 issue with this pathway is that MPn is not a known natural product, nor has it been detected  
82 in natural systems. However, it was recently shown that the marine archaeon *Nitrosopumilus*  
83 *maritimus* encodes a pathway for MPn biosynthesis and that it produces cell-associated MPn  
84 esters (Metcalf et al., 2012). They argued that these cells could provide sufficient amounts of  
85 MPn precursor to account for the observed CH<sub>4</sub> production in the oxic ocean via the C-P lyase  
86 dependent scenario suggested by Karl et al. (2008). However, it was not possible to explain

87 the supersaturation state of CH<sub>4</sub> in oxic surface water by quantification of produced CH<sub>4</sub> from  
88 dissolved MPn under natural conditions (del Valle and Karl, 2014).

89 It remains equivocal if CH<sub>4</sub> formation from MPn (Karl et al. 2008) or metabolism of DMSP by  
90 methanogens in anoxic microenvironments (Damm et al., 2008; Zindler et al., 2013, Damm et  
91 al., 2015) is sufficient to provide a permanent increase in the concentration of CH<sub>4</sub> in  
92 oxygenated surface waters, or if other pathways are also required to fully explain the CH<sub>4</sub>  
93 oversaturation in oxic waters. In this context it is important to mention that almost 40 years  
94 ago researchers (Scranton and Brewer, 1977; Scranton and Farrington, 1977) already  
95 mentioned the possibility of *in-situ* formation of CH<sub>4</sub> by marine algae. These scientists  
96 measured CH<sub>4</sub> saturation states in open ocean surface waters of the west subtropical North-  
97 Atlantic. They observed 48-67 % higher CH<sub>4</sub> concentrations in surface waters than estimated  
98 from atmospheric equilibrium concentration, with a narrow maximum of CH<sub>4</sub> concentration  
99 in the uppermost part of pycnocline. Since the loss of CH<sub>4</sub> from surface to atmosphere was  
100 calculated to be much larger than diffusion from CH<sub>4</sub> maxima of the pycnocline into the mixed  
101 layer, an *in situ* biological CH<sub>4</sub> formation process within the mixed layer was hypothesized  
102 (Scranton and Farrington, 1977; Scranton and Brewer, 1977). However, direct evidence of  
103 algae-derived CH<sub>4</sub> formation from laboratory experiments with (axenic) algae cultures is still  
104 missing, and the accumulation of CH<sub>4</sub> in the upper water layer has not yet been directly related  
105 to production by algae.

106 The aim of our study was to quantify *in-situ* CH<sub>4</sub> formation from marine algae such as  
107 coccolithophores and to identify precursor compounds of CH<sub>4</sub> via <sup>13</sup>C labelling techniques.  
108 Therefore, we used *Emiliania huxleyi*, a widely distributed, prolific alga. The coccolithophore

109 blooms including *E. huxleyi* are the major regional source of DMS release to the atmosphere  
110 (Holligan et al., 1993). Specific goals in this study were (I) to measure CH<sub>4</sub> production of a  
111 biogeochemically important marine phytoplankton, (II) to screen for methanogenic archaea  
112 or bacteria and (III) to identify methyl sulfides, such as the amino acid methionine, that play a  
113 role in metabolic pathways of algae - as possible precursors for CH<sub>4</sub>.

114

## 115 **MATERIAL & METHODS**

### 116 **Culture media and culture conditions**

117 Monoclonal cultures of *E. huxleyi* [RCC1216; <http://roscoff-culture-collection.org/>] were  
118 grown in full batch mode (Langer et al. 2013) in sterile filtered (0.2 µm) seawater (Helgoland,  
119 North Sea) enriched with phosphate, nitrate, trace metals and vitamins according to F/2  
120 (Guillard and Ryther, 1962). Main cultures were inoculated with 3500 cells ml<sup>-1</sup>, sampled from  
121 a pre-culture grown in dilute batch mode (Langer et al. 2009). Final cell densities of the main  
122 cultures were approximately 1 × 10<sup>6</sup> cells ml<sup>-1</sup>.

123

124 To investigate algae-derived CH<sub>4</sub> formation a closed-chamber system was used. Hence 2l flasks  
125 (Schott, Germany) filled with 1800 ml sterile filtered sea water and with 480 ml headspace  
126 volume were used in our investigations. The flasks were sealed with lids (GL 45, PP, 2 port,  
127 Duran Group) equipped with two three-way-ports (Discofix<sup>®</sup>-3, B-Braun), where one port was  
128 used for water and the other port (fitted with a sterile filter, 0.2 µm; PTFE, Satorius) for gas  
129 sampling. The cells were grown on a day/night cycle of 16/8 h at 20°C and a light intensity of

130  $\approx 450 \mu\text{E}$  over a 10 day period. Initial dissolved inorganic carbon (DIC) of the culture medium  
131 was  $2235 \mu\text{mol l}^{-1}$  (for details on DIC measurements see Langer et al. 2009).

132

133 The different treatments and number of replicates are provided in Table 1. To increase the  
134 detectability of  $\text{CH}_4$ -formation and to exclude a possible contamination with  $\text{CH}_4$  from the  
135 surrounding air,  $^{13}\text{C}$ -labelled bicarbonate ( $\text{NaH}^{13}\text{CO}_3$ , 99 % purity, Sigma-Aldrich, Germany)  
136 was added to the cultures. Bicarbonate (Bic) was used as C-source for biomass production. To  
137 gain a  $^{13}\text{C}$ -enrichment of 1 % of the total inorganic C ( $\text{CO}_2$ ,  $\text{HCO}_3^-$ , and  $\text{CO}_3^{2-}$ ),  $22.35 \mu\text{mol l}^{-1}$   
138  $\text{NaH}^{13}\text{CO}_3$  was added, leading to a theoretical  $\delta^{13}\text{C}$  value of 882 ‰.

139 We used two different control treatments: 1) Algae cultures without  $^{13}\text{C}$ -Bic and 2) sea water  
140 with  $^{13}\text{C}$ -Bic.

141 To test methionine (Met) as a precursor of algae-derived  $\text{CH}_4$ , Met where only the sulfur-  
142 bound methyl-group was  $^{13}\text{C}$ -labelled (R-S- $^{13}\text{CH}_3$ , 99 % enriched,  $1 \mu\text{mol l}^{-1}$ ) was added to the  
143 cultures. Met has previously been identified as a methyl-group donor for  $\text{CH}_4$  biosynthesis in  
144 higher plants and fungi (Lenhart et al. 2012, 2015). Moreover, marine algae use Met to  
145 produce DMSP, DMS and DMSO, substances that can be released into seawater and known to  
146 act as precursors for abiotic  $\text{CH}_4$  production.

147

#### 148 **Sample collection and analysis**

149 Samples were taken daily from day 4 until day 10 (see Table 1). Prior to day 4, algae biomass  
150 was too low to allow measurement of changes in  $\text{CH}_4$  mixing ratio.

151 For GC-FID/ECD and CF-IRMS analysis samples of headspace (30 ml) were taken from each  
152 flask. GC-samples were measured within 24h after sampling while GC-IRMS samples were  
153 stored in 12 ml exetainers until  $^{13}\text{C-CH}_4$  measurements were carried out.

154 After gas sampling, samples of medium (25 ml) from each flask were also taken for cell density  
155 determination. These samples were supplemented with 0.15 ml Lugol solution (Utermöhl,  
156 1958) and stored in 50 ml Falcon tubes at 4°C. In order to maintain atmospheric pressure  
157 within the flask, surrounding air was allowed to enter via an orifice fitted with a sterile filter  
158 to avoid bacterial contamination. Variable amounts of water and headspace volume as well  
159 as inflow of surrounding air were all taken into consideration when  $\text{CH}_4$  production rates were  
160 calculated.

161 Cell density was determined via a Hemocytometer (Thoma-Kammer with 256 fields, 0.0025  
162  $\text{mm}^2 \times 0.1 \text{ mm}$ ; Laboroptik Ltd, UK).

163

#### 164 **Gas chromatography**

165 Gas samples were analysed for  $\text{CH}_4$  mixing ratio within 24 h on a gas chromatograph  
166 (Shimadzu GC-14B, Kyoto, Japan) fitted with a flame ionization detector (FID) operating at 230  
167 °C with  $\text{N}_2$  as carrier gas ( $25 \text{ ml min}^{-1}$ ) (Kammann et al., 2009). The GC column (PorapakQ, Fa.  
168 Millipore, Schwallbach, mesh 80/100) was 3.2 m long and 1/8 inch in diameter. The length of  
169 the precolumn was 0.8 m. The GC gas flow scheme and automated sampling was that of  
170 Mosier and Mack (1980) and Loftfield (1997), and peak area integration was undertaken with  
171 the Software PeakSimple, version 2.66. The standard deviation (s.d.) of the mean of six  
172 atmospheric air standard samples was below 0.2 % for  $\text{CH}_4$ .

173



174 **Continuous flow isotope ratio mass spectrometry (CF-IRMS) for measurement of  $\delta^{13}\text{C}$  values**  
175 **of  $\text{CH}_4$**

176 Headspace gas from exetainers was transferred to an evacuated sample loop (40 mL).  
177 Interfering compounds were separated by GC and  $\text{CH}_4$  trapped on Hayesep D. The sample was  
178 then transferred to the IRMS system (ThermoFinnigan Delta<sup>plus</sup> XL, Thermo Finnigan, Bremen,  
179 Germany) via an open split. The working reference gas was carbon dioxide of high purity  
180 (carbon dioxide 4.5, Messer Griesheim, Frankfurt, Germany) with a known  $\delta^{13}\text{C}$  value of -  
181 23.64 ‰ relative to Vienna Pee Dee Belemnite (V-PDB). All  $\delta^{13}\text{C}$  values of  $\text{CH}_4$  were corrected  
182 using three  $\text{CH}_4$  working standards (isometric instruments, Victoria, Canada) calibrated against  
183 IAEA and NIST reference substances. The calibrated  $\delta^{13}\text{C}$ - $\text{CH}_4$  values of the three working  
184 standards were  $-23.9 \pm 0.2$  ‰,  $-38.3 \pm 0.2$  ‰ and  $-54.5 \pm 0.2$  ‰. Samples were routinely analysed  
185 three times ( $n = 3$ ) and the average standard deviations of the CF-IRMS measurements were  
186 in the range of 0.1 to 0.3 ‰.

187 All  $^{13}\text{C}/^{12}\text{C}$  -isotope ratios are expressed in the conventional  $\delta$  notation in per mil [‰] versus  
188 V-PDB, using the following equation (Eq. 1):

$$189 \quad \delta^{13}\text{C} = \left( \frac{(^{13}\text{C}/^{12}\text{C})_{\text{sample}}}{(^{13}\text{C}/^{12}\text{C})_{\text{standard}}} \right) - 1. \quad (\text{Eq. 1})$$

190 To determine the  $\delta^{13}\text{C}$  signature of the  $\text{CH}_4$  source, the Keeling-plot method was applied  
191 (Keeling, 1958).

192

193 **Microbial investigations**

194 **DNA extraction and real-time PCR**

195 Samples for DNA extraction were taken from the stem culture (RCC 1216) during the  
196 stationary growth phase ( $2 \times 10^6$  cells ml<sup>-1</sup>). After DNA extraction, realtime PCR was used to  
197 detect mcrA-genes, which are solely found in methanogenic archaea. As positive proof,  
198 aliquots of the samples were supplemented with a defined cell density of  
199 *Methanothermobacter marburgensis* (either  $10^4$  or  $10^7$  cells ml<sup>-1</sup>).

200 The DNA extraction was carried out according to (Bürgmann et al., 2001). 1 ml of the algae  
201 culture was transferred into a 2 ml vial containing 200 µl of Zirconia-silica beads (Roth) and  
202 centrifuged for 20 minutes ( $1.3 \times 10^4$  U min<sup>-1</sup>; 20°C). Afterwards, 850 µl of the supernatant was  
203 replaced with extraction buffer (Bürgmann et al., 2001) and beaten for 50 s (Retsch, type  
204 MM2). After centrifugation the supernatant was transferred to another vial (2 ml, Eppendorf,  
205 Germany), mixed with 850 µl phenol/chloroform/iso-amyl-alcohol-solution (Roth) and again  
206 centrifuged for 5 minutes ( $1.3 \times 10^4$  U min<sup>-1</sup>; 20°C). The water phase was supplemented with  
207 800 µl phenol, mixed and centrifuged again. Afterwards, the water phase was transferred in a  
208 new vial, mixed with 800 µl precipitating buffer (PEG) and centrifuged for 60 min ( $1.3 \times 10^4$  U  
209 min<sup>-1</sup>; 20°C). The pellet was washed with 800 µl ethanol (75%; -20°C, centrifuged for 10 min  
210 at  $1.3 \times 10^4$  U min<sup>-1</sup>; 20°C) and air-dried in the laboratory. For elution and storage of the pellet  
211 we used 20 µl nuclease-free water.

212 Real-time PCR was carried out according to Kampmann et al. (2012) with a Rotor-Gene 3000  
213 (Fa. Corbett Research, Australia) by using Absolute™ QPCR SYBR® Green Mix (ABgene). For the  
214 detection of mcrA-Genes we used the primer  
215 (ML forward: 5'GGTGGTGTMGATTACACARTAYGCWACAGC-3'; ML reverse:

216 5' AACTAYCCWAACTAYGCAATGAA-3'), which encodes the  $\alpha$ -subunit of the methyl-CoM-  
217 reductase, that solely occurs in methanogenic archaea (Luton et al., 2002).

218 The real-time PCR reference standards were produced according to Kampmann et al. (2012).  
219 By using the standard solution ( $5.5 \times 10^7$  DNA copies  $\mu\text{l}^{-1}$ ) dilution with nuclease-free water  
220 was accomplished down to  $5.5 \times 10^1$  copies per  $\mu\text{l}^{-1}$ . All standards and regular samples taken  
221 from the flasks were analyzed with four repetitions.

222 Quality assurance of the real-time PCR-product was achieved by melt curve analysis and  
223 gelelectrophoresis using the fluorescent stain GelRedTM (Biotium).

224

#### 225 **Cultivation approach**

226 In addition to real-time PCR, a cultivation/enrichment procedure (Kampmann et al., 2012) was  
227 conducted to screen for methanogenic archaea in algae cultures. The enrichment medium  
228 (Widdel and Bak, 1992) was modified for marine conditions by adding  $320 \text{ mmol l}^{-1}$  NaCl;  $16$   
229  $\text{mmol l}^{-1}$   $\text{MgCl}_2$  and  $1 \text{ mmol l}^{-1}$   $\text{NaHCO}_3$ . At day 10 an aliquot (5 ml) of each cultivation flask was  
230 transferred into injection flasks (Ochs, Bovenden-Lenglern, Germany) with the enrichment-  
231 medium (50 ml) and acetate (10 mM), methanol (5 mM) was added and in the gas phase  $\text{H}_2$   
232 and  $\text{CO}_2$  (90:10) was provided as substrates. Incubation was carried out over a period of 6  
233 weeks at  $20^\circ\text{C}$  in the dark.

234

#### 235 **CH<sub>4</sub> mass**

236 The mass of CH<sub>4</sub> ( $m_{CH_4}$ ) per flask was calculated via the ideal gas law from the corrected CH<sub>4</sub>  
237 mixing ratio (ppmv), where the changing volume of water and headspace and the inflow of  
238 surrounding air were all considered, according to Eqn. 3:

239

$$240 \quad m_{CH_4} = \frac{p}{R \times T} \times c_{CH_4} \times V \times M_{CH_4} \quad (3)$$

241

242 Where p = pressure, T = temperature, R = ideal gas constant, V = volume, and  $M_{CH_4}$  =  
243 molweight CH<sub>4</sub>. The solubility of CH<sub>4</sub> in the water phase was calculated according to  
244 Wiesenburg and Guinasso (Wiesenburg and Guinasso Jr, 1979) based on the headspace-CH<sub>4</sub>  
245 mixing ratio, temperature and salinity of the water phase.

246

#### 247 **Calculation of CH<sub>4</sub> production**

248 The low CH<sub>4</sub> mixing ratios produced by *E. huxleyi* during the exponential growth phase  
249 precluded the determination of CH<sub>4</sub> production during this period. Therefore we calculated  
250 production from day 7 to day 10, a period representing the transition from exponential to  
251 stationary phase. This growth phase features changing growth rates and cellular CH<sub>4</sub> quotas,  
252 rendering the dilute batch method of calculating production inapplicable (Langer et al. 2013).  
253 We followed the recommendation of Langer et al. (2013) and calculated incremental (daily)  
254 CH<sub>4</sub> production:

$$255 \quad P_{inc} = q_{inc} \times \mu_{inc} \quad (4)$$

256 with  $P_{inc}$  = incremental  $CH_4$  production [ $ng\ CH_4\ cell^{-1}\ day^{-1}$ ],  $q_{inc}$  = incremental cellular  $CH_4$   
257 quota [ $ng\ CH_4\ cell^{-1}$ ],  $\mu_{inc}$  = incremental growth rate [ $day^{-1}$ ]

258 Incremental growth rate was calculated according to:

$$259 \quad \mu_{inc} = LN(t_1) - LN(t_0) \quad (5)$$

260 with  $t_1$  = cell density on the day  $q_{inc}$  was determined,  $t_0$  = cell density on the previous day. We  
261 present average  $P_{inc}$  (STDEV).

262 In order to compare  $CH_4$  production to literature data it was necessary to normalize to cellular  
263 particulate organic carbon (POC) quota, as opposed to cell. The POC normalized  $CH_4$   
264 production is termed “methane emission rate” in the following. Since it was not possible to  
265 measure cellular POC quota on a daily basis, we used a literature value determined for the  
266 same strain under similar culture conditions, i.e.  $10.67\ pg\ POC\ cell^{-1}$  (Langer et al. 2009). We  
267 are aware of the fact that cellular POC quota is likely to change alongside other element  
268 quotas when approaching stationary phase, but this change is well below an order of  
269 magnitude (Langer et al. 2013). For our purpose this method is therefore sufficiently accurate  
270 to determine POC normalized  $CH_4$  production.

271

## 272 **Statistics**

273 To test for significant differences in cell density,  $CH_4$  mixing ratio and  $CH_4$  content between  
274 the treatments, two-way ANOVA (considering repeated measurements) and a Post-Hoc-Test  
275 (Fisher LSD-Test; alpha 5 %) was used.

276

## 277 **RESULTS**

### 278 **Algae growth**

279 Cell density and growth of the cultures are presented in Figure 2a, b over the whole incubation  
280 period for all treatments. The initial cell density at time 0 ( $t_0$ ) was  $3.5 \times 10^3$  cells  $\text{ml}^{-1}$  in all  
281 flasks. At day 10 cell density reached its maximum value with  $1.37 \times 10^6$  cells  $\text{ml}^{-1}$  (algae),  $0.82$   
282  $\times 10^6$  cells  $\text{ml}^{-1}$  (algae +  $^{13}\text{C}$ -Bic) and  $1.24 \times 10^6$  cells  $\text{ml}^{-1}$  (algae +  $^{13}\text{C}$ -Met). The exponential  
283 growth rates ( $\mu$ ) were  $0.85 \pm 0.2 \text{ d}^{-1}$  for "algae +  $^{13}\text{C}$ -Met",  $0.98 \pm 0.1 \text{ d}^{-1}$  for "algae +  $^{13}\text{C}$ -Bic",  
284 and  $1.06 \pm \text{d}^{-1}$  for the control "algae" (n.s.,  $p = 0.286$ ). Significant differences in cell density  
285 between the treatments only occurred at days 9 and 10, where the cell density of the control  
286 "algae" was higher than in the treatments where  $^{13}\text{C}$ -Bic or  $^{13}\text{C}$ -Met was added.

### 287 **Methane mixing ratio**

288 Initial headspace- $\text{CH}_4$  mixing ratios measured at day 4 were in the range of 1899 to 1913 ppbv  
289 for all treatments including the controls without algae. From day 4 to day 7 headspace- $\text{CH}_4$   
290 mixing ratios slightly increased in all flasks. Therefore, no significant differences in the  $\text{CH}_4$ -  
291 mixing ratios occurred between the treatments. After day 8  $\text{CH}_4$  mixing ratios in the flasks  
292 containing algae were significantly higher compared to the controls without algae (Fig. 2c, d).  
293 The highest  $\text{CH}_4$  mixing ratios at day 10 corresponded to  $2102 \pm 62$  ppbv (algae +  $^{13}\text{C}$ -Met),  
294  $2138 \pm 42$  ppbv (algae +  $^{13}\text{C}$ -Bic) and  $2119 \pm 25$  ppbv (algae).

295 Hence, from day 4 to day 10 the  $\text{CH}_4$  mixing ratios increased by about 192 ppbv (algae +  $^{13}\text{C}$ -  
296 Met), 49 ppbv (sea water +  $^{13}\text{C}$ -Met), 235 ppbv (algae +  $^{13}\text{C}$ -Bic) and 67 ppbv (sea water +  $^{13}\text{C}$ -  
297 Bic), respectively.

298 **Stable carbon isotope values of methane**

299 The  $\delta^{13}\text{C}$  signature of headspace- $\text{CH}_4$  ( $\delta^{13}\text{CH}_4$  value) is presented in Figure 2e, f. Addition of  
300  $^{13}\text{C}$ -Bic did not affect  $\text{CH}_4$  production of algae, but the  $\delta^{13}\text{CH}_4$  value was clearly different from  
301 that of the control “algae”. The initial value of  $-47.9 \pm 0.2$  ‰ increased to  $44 \pm 13$  ‰ whereas  
302 in the controls “seawater +  $^{13}\text{C}$ -Bic” and “algae” no change in the  $\delta^{13}\text{CH}_4$  value was observed.

303 Addition of  $^{13}\text{C}$ -Met did not affect algal  $\text{CH}_4$  formation, but it increased the  $\delta^{13}\text{CH}_4$  signature  
304 from  $-46.35 \pm 0.84$  ‰ to  $59.1 \pm 25.3$  ‰ (day 8). In the treatment “ $^{13}\text{C}$ -Met”, where only  
305 isotopically labelled Met was added to sterile filtered sea water, a small increase from  $-48.0 \pm$   
306  $0.3$  to  $-38.1 \pm 2.3$  ‰ (at day 10) was observed.

307 Based on the initial amount of  $^{13}\text{C}$ -Bic and the total amount of  $^{13}\text{CH}_4$  at the end of the  
308 incubation period,  $88.3 \pm 17.2$  pmol of  $22.4$   $\mu\text{mol}$   $^{13}\text{C}$ -Bic were converted to  $^{13}\text{CH}_4$ . For Met,  
309 this was  $78.5 \pm 18.6$  pmol of the initial  $1.8$   $\mu\text{mol}$   $^{13}\text{C}$ -Met.

310 The Keeling-plots to determine the  $^{13}\text{C}$  values of the  $\text{CH}_4$  source are presented in (Fig. 3). For  
311 the bicarbonate treatment (“Algae +  $^{13}\text{C}$ -Bic”), the mean  $\delta^{13}\text{CH}_4$  value of the  $\text{CH}_4$  source was  
312  $811.9 \pm 89.9$  ‰, which is close to the calculated  $\delta^{13}\text{C}$  value of  $881.5$  ‰ after the addition of  
313  $\text{NaH}^{13}\text{CO}_3$ .

314 For the treatment “Algae +  $^{13}\text{C}$ -Met” we applied the Keeling-plot method only for the period  
315 from day 5 to day 7, as the increase in the  $\delta^{13}\text{C}$  values were not linear after day 7. For this  
316 treatment, the  $\delta^{13}\text{C}$  values of the  $\text{CH}_4$  source range between  $967$  and  $2979$  ‰.

317

318 The correlation between the growth of the algae cultures and the total amount of CH<sub>4</sub> in the  
319 flasks (headspace + water phase) is presented in Figure 4. For the treatment “algae + <sup>13</sup>C-Bic”  
320 (Fig. 4a) there is an exponential correlation between cell density and CH<sub>4</sub>-content ( $r^2 = 0.994$ ).  
321 Whereas for the treatment “algae + <sup>13</sup>C-Met” (Fig. 4b) a linear correlation was observed ( $r^2 =$   
322  $0.995$ ).

323

324 The daily CH<sub>4</sub> content in the flasks for days 8, 9 and 10 is shown in Figure 5. For all flasks the  
325 CH<sub>4</sub> content exceeded the CH<sub>4</sub> content of the respective control, with a continuous increase  
326 of the CH<sub>4</sub> content in the flasks containing algae. At day 10, the difference between “algae +  
327 <sup>13</sup>C-Bic” and “sea water + <sup>13</sup>C-Bic” and between “algae + <sup>13</sup>C-Met” and “sea water + <sup>13</sup>C-Met”  
328 was  $65 \pm 16$  and  $54 \pm 22$  ng, respectively.

329

330 The CH<sub>4</sub> production of algae presented in Table 2 shows no major differences between the  
331 treatments. Furthermore for all treatments, the daily CH<sub>4</sub> production rates did not change over  
332 time (Fig. 6).

333

### 334 **Microbial investigations**

335 Via real-time PCR no mcrA-genes could be detected in the flasks containing the CH<sub>4</sub>-producing  
336 algae cultures. Whereas the positive control in which the algae culture was supplemented  
337 with  $10^4$  and  $10^7$  cells ml<sup>-1</sup> of the methanogenic archaea *Methanothermobacter*  
338 *marburgensis*,  $9.4 \cdot 10^4$  and  $4.6 \cdot 10^6$  mcrA-gene copies ml<sup>-1</sup> have been detected, respectively.



339 With the cultivation approach, where an aliquot of each flask was taken at day 10 and  
340 transferred in the media for enrichment of methanogenic archaea, no CH<sub>4</sub> production was  
341 observed after the 6 week incubation period. In case of a successful enrichment of  
342 methanogenic archaea, the CH<sub>4</sub>-mixing ratio in the headspace would increase over time.

343

## 344 **DISCUSSION**

345 Our results of the CH<sub>4</sub> mixing ratio and stable isotope measurements provide unambiguous  
346 evidence that *E. huxleyi* produces CH<sub>4</sub>. In the following we will discuss the relationship  
347 between CH<sub>4</sub> production and growth of the algae, stable isotope measurements, potential  
348 precursor compounds, and the exclusion of methanogenic archaea. Finally, we will discuss the  
349 implications of our results for the methane paradox in oxic waters.

### 350 **Growth and CH<sub>4</sub> production**

351 Over the course of the exponential growth phase headspace CH<sub>4</sub> mixing ratios in treatments  
352 containing *E. huxleyi* were not measurably different from the control treatments. Therefore it  
353 was not possible to determine CH<sub>4</sub> production in the exponential growth phase. However, we  
354 conclude that *E. huxleyi* produces CH<sub>4</sub> throughout all growth phases as will be detailed in the  
355 following. In the transitional growth phase leading up to stationary phase we calculated  
356 incremental CH<sub>4</sub> production (daily). The transitional phase features declining growth rate and  
357 often increasing cellular carbon quotas (Langer et al. 2013). Also cellular CH<sub>4</sub> quotas did  
358 increase (data not shown). On the other hand, CH<sub>4</sub> production remained constant within the  
359 measurements of error, displaying a slight downward trend when approaching stationary

360 phase (Fig. 6). Therefore we conclude that CH<sub>4</sub> production is not a feature of senescent cells  
361 only, but probably is operational in all growth phases. This is interesting in the context of the  
362 ecology and biogeochemistry of *E. huxleyi*. Contrary to the traditional assumption that *E.*  
363 *huxleyi* production in the field is dominated by late summer bloom events, it was recently  
364 shown that non-bloom production in spring contributes significantly to yearly average  
365 production and therefore bloom events are not exceptionally important in biogeochemical  
366 terms (Schiebel et al. 2011). Since senescent cells in field samples are mainly a feature of late  
367 bloom stages, the exclusive production of CH<sub>4</sub> by such cells would confine any contribution of  
368 *E. huxleyi* to the oceanic CH<sub>4</sub> budget to a relatively short, and biogeochemically less important,  
369 period. However from results found in this study we would propose that *E. huxleyi* produces  
370 CH<sub>4</sub> during all growth phases as part of its normal metabolism. If our findings are confirmed  
371 and supported by other research groups this has considerable implications as it would render  
372 this species a prolific aerobic producer of CH<sub>4</sub> on a par with, for example, terrestrial plants  
373 (Bruhn et al., 2012).

#### 374 **Methane emission rates**

375 To calculate CH<sub>4</sub> emission rates of *E. huxleyi*, we normalized CH<sub>4</sub> production to cellular  
376 particulate organic carbon (POC) content (see Material and Methods). The CH<sub>4</sub> emissions were  
377 0.7 µg POC g<sup>-1</sup> d<sup>-1</sup>, or 30 ng g<sup>-1</sup> POC h<sup>-1</sup> (mean for all treatments, n = 8).

378 In this study the main aim was (as a proof of principle) to unambiguously provide evidence  
379 that *E. huxleyi* are able to produce methane under aerobic conditions and without the help of  
380 microorganisms.

381 However, we suggest that CH<sub>4</sub> emission rates of *E. huxleyi* algae are different under changing  
382 environmental conditions, e.g. temperature, light intensity or nutrient supply. The effect of  
383 changing environmental parameters should be the focus of future investigations.

384 For comparison CH<sub>4</sub> emission rates presented so far for terrestrial plants range from 0.3 to  
385 370 ng g<sup>-1</sup> DW (dry weight) h<sup>-1</sup> (Keppler et al., 2006; Wishkerman et al., 2011; Lenhart et al.,  
386 2015; Brüggemann et al., 2009).

387

#### 388 **Inorganic and organic precursors of CH<sub>4</sub>**

389 Based on the addition of bicarbonate (<sup>13</sup>C-Bic, 1 % enrichment), which is the principal carbon  
390 source for growth of algae, and the measurements of δ<sup>13</sup>CH<sub>4</sub> values it was possible to clearly  
391 identify bicarbonate as the principal carbon precursor of CH<sub>4</sub> in *E. huxleyi*.

392 In the flasks where algae were supplemented with <sup>13</sup>C-Bic, a significant increase in δ<sup>13</sup>CH<sub>4</sub>  
393 values occurred over the incubation period, which shows that algae use bicarbonate as  
394 precursor carbon (C) for CH<sub>4</sub> production. As expected, in the controls flasks “algae” where no  
395 <sup>13</sup>C-Bic was added and the control “sea water + <sup>13</sup>C-Bic” without algae, no change in δ<sup>13</sup>CH<sub>4</sub>  
396 values was observed. The initial δ<sup>13</sup>C value of the bicarbonate in the treatment “algae + <sup>13</sup>C-  
397 bic” (+882 ‰) is within the range of the source δ<sup>13</sup>CH<sub>4</sub> values obtained via the Keeling-plot  
398 method (+812 ±90 ‰). Even though there might be kinetic isotope fractionations involved in  
399 each of the several steps during organic matter formation these data clearly indicate that  
400 bicarbonate is the principle inorganic carbon precursor of CH<sub>4</sub> produced in algae.

401 Bicarbonate is taken up by the algae via autotrophic C fixation (Burns and Beardall, 1987) and  
402 might therefore - during several steps of metabolism i.e. formation of organic compounds -  
403 lead to the formation of CH<sub>4</sub>. Probably, it will be used as an unspecific C source in many  
404 different metabolic pathways, e.g. the synthesis of lignin, pectin, and cellulose (Kanehisa et  
405 al., 2014) – components already known as CH<sub>4</sub> precursors from terrestrial plants, where via  
406 methyl group cleavage CH<sub>4</sub> can be produced (Keppler et al., 2008; Bruhn et al., 2009; Vigano  
407 et al., 2009). However, lignin and pectin are not commonly found in marine algae such as *E.*  
408 *huxleyi*. For these organisms sulphur bonded methyl groups such as thioethers, sulfoxides and  
409 sulfonium salts (methionine, S-adenosylmethionine SAM, dimethylsulfoniopropionate DMSP,  
410 dimethyl sulfoxide DMSO, dimethyl sulfide DMS) are of much more interest. For our  
411 experiments, we used <sup>13</sup>C positionally labelled Met where only the sulfur-bond methyl group  
412 (–S-CH<sub>3</sub>) was 99 % enriched in <sup>13</sup>C. Our choice of this compound was partly due to its  
413 commercial availability but more importantly because it is known to be involved in a number  
414 of metabolic pathways and transmethylation reactions (Stefels, 2000, Bruhn et al. 2012).

415 In contrast to the ubiquitous C-source bicarbonate –which can also be used to build Met in  
416 algae (Stefels, 2000) – Met is incorporated in specific metabolic pathways. Algae use part of  
417 the Met for protein synthesis, in *E. huxleyi* it is also involved in the synthesis of DMSP, a main  
418 precursor of DMS and DMSO.

419 The clear increase in δ<sup>13</sup>CH<sub>4</sub> values of headspace-CH<sub>4</sub> in the treatment “algae +<sup>13</sup>C-Met” (Fig.  
420 2e, f) shows that the methyl thiol group of Met is a direct CH<sub>4</sub> precursor. The Keeling-plot  
421 results (Fig. 3) show higher variability for Met than for Bic. However, Met is almost certainly  
422 not the only precursor of CH<sub>4</sub>, as the headspace-CH<sub>4</sub> mixing ratios increased (Fig. 2d), while

423 the  $^{13}\text{C}$  values of headspace- $\text{CH}_4$  showed a saturation curve (Fig. 2f). This indicates either a  
424 shift from Met to other  $\text{CH}_4$  precursors, or to the use of newly synthesized, non-labelled Met.  
425 Based on the initial amount and the total amount of  $^{13}\text{CH}_4$  formed at the end of the incubation,  
426 only a small fraction (79 pmol, i.e. 4.0 ‰) of the initial added  $^{13}\text{C}$ -Met (1.8  $\mu\text{mol}$ ) was converted  
427 to  $^{13}\text{CH}_4$ . The formation of  $\text{CH}_4$  from  $^{13}\text{C}$ -Met explains roughly about 3 % of the total amount  
428 of  $\text{CH}_4$  formed throughout the incubation period. Possibly, the formation of potential  
429 precursors of  $\text{CH}_4$  may change under various climatic conditions, leading to varying  $\text{CH}_4$   
430 production rates in different pathways.

431 This observation is in line with the findings of Lenhart and colleagues who demonstrated the  
432 sulphur-bound methyl group of Met as a precursor for  $\text{CH}_4$  in plants (Lenhart et al., 2015) and  
433 fungi (Lenhart et al., 2012). The linear increase in headspace- $\text{CH}_4$  mixing ratio (Fig. 2d) together  
434 with the non-linear increase in  $\delta^{13}\text{CH}_4$  signature (Fig. 1f) indicates that the pool of  $^{13}\text{C}$ -Met was  
435 either exhausted or was diluted by newly synthesized, non  $^{13}\text{C}$  enriched Met.

436 In addition, we also found an indication for a chemical  $\text{CH}_4$  formation pathway in the sea water  
437 with Met as methyl-group donor as a small increase in  $^{13}\text{CH}_4$  values in the control treatment  
438 "sea water +  $^{13}\text{C}$ -Met" was observed (Fig. 2f). This  $\text{CH}_4$  formation pathway is approximately  
439 10-fold lower when compared to the treatment "algae +  $^{13}\text{C}$ -Met" and is only observed in the  
440 isotopic experiment, but not when only  $\text{CH}_4$  mixing ratio is considered (Fig. 2d). However, this  
441 observation is in line with some previous findings (Althoff et al., 2010; Althoff et al., 2014),  
442 who showed that abiotic formation of  $\text{CH}_4$  due to the degradation of methionine or ascorbic  
443 acid by light or oxidants such as iron minerals is possible. In the case of methionine it was

444 shown that the sulphur-bound methyl group of Met was the carbon precursor for CH<sub>4</sub> (Althoff  
445 et al. 2014).

#### 446 **Potential implications for the occurrence of CH<sub>4</sub> in oxic marine waters**

447 Several hypotheses with regard to the occurrence of the seasonal and spatial CH<sub>4</sub>  
448 oversaturation in oxic surface waters (Bange et al., 1994; Forster et al., 2009; Owens et al.,  
449 1991) have been postulated. They include CH<sub>4</sub> formation from methanogenic archaea in  
450 anoxic microsites (Karl and Tilbrook, 1994), or CH<sub>4</sub> formation via the C-P-lyase pathway from  
451 methylphosphonate (Karl et al., 2008).

452 In the ocean, both CH<sub>4</sub> production by methanogens and consumption via methanotrophic  
453 bacteria occur simultaneously. Therefore, CH<sub>4</sub> production can exceed estimated CH<sub>4</sub>  
454 production rates when based solely on CH<sub>4</sub> mixing ratio measurements (Reeburgh, 2007). To  
455 provide a noteworthy contribution to oceanic CH<sub>4</sub> production, precursors must either be  
456 available in high abundance or be continually synthesized. Algae-derived methylated sulphur  
457 compounds such as Met, DMSP, DMS, and DMSO are ubiquitous in the ocean but show a high  
458 spatial and temporal variability with high mixing ratios in algal blooms. Therefore, they are  
459 potential compounds that might be involved in CH<sub>4</sub> formation in the oceans (Keppler et al.,  
460 2009; Althoff et al., 2014). The involvement of methyl moieties from methylated sulfur  
461 compounds in CH<sub>4</sub> biosynthesis might therefore play an important role in pelagic CH<sub>4</sub>  
462 production. Mixing ratios of DMS and DMSP in sea water during algal blooms were reported  
463 in the range of 0.82 to 8.3 nmol l<sup>-1</sup> and 1.25 to 368 nmol<sup>-1</sup>, respectively (Matrai and Keller,  
464 1993).

465 The CH<sub>4</sub> emission rates of *E. huxleyi* may also occur by a second formation pathway, where  
466 DMSP is first converted to DMS and subsequently oxidized to DMSO (Bentley and Chasteen,  
467 2004).

468 However, several studies have afforded evidence for a CH<sub>4</sub> formation pathway via methyl  
469 radicals (Althoff et al., 2014; Eberhardt and Colina, 1988; Herscu-Kluska et al., 2008), leading  
470 to the hypothesis that algae-derived DMSO can also act as a precursor of CH<sub>4</sub> in oxic seawater  
471 (Althoff et al., 2014). A correlation between Met and DMSP synthesis was provided by Gröne  
472 and Kirst (1992) who showed that supplementation of *Tetraselmis subcordiformis* with 100 µg  
473 l<sup>-1</sup> Met yielded a 2.6-fold increase in DMSP. For *E. huxleyi*, DMSO mixing ratios in the stationary  
474 growth phase can reach 0.1 pg per cell (Simo et al., 1998). Assuming that a similar DMSO  
475 mixing ratio were to be found in our study, this would mean that in every 4 × 10<sup>3</sup> DMSO  
476 molecules per day must be transferred to CH<sub>4</sub> to explain the observed increase in CH<sub>4</sub>.  
477 Moreover, a positive correlation was observed between Chlorophyll a and CH<sub>4</sub>, as well as  
478 between DMSP or DMSO and CH<sub>4</sub> (Zindler et al., 2013).

479

## 480 **Conclusions and Outlook**

481 Our study provides the first isotope evidence that marine algae such as *E. huxleyi* produce CH<sub>4</sub>  
482 with bicarbonate and the sulfur-bound methyl group of Met as C precursors. Our results based  
483 on real-time PCR and enrichment of methanogenic Archaea make it highly unlikely that there  
484 is a contribution of Archaea to the observed CH<sub>4</sub> production. It is of interest to note that it is  
485 almost 40 years since algae were suggested as a possible direct source of CH<sub>4</sub> in the ocean  
486 (Scranton and Brewer, 1977; Scranton and Farrington, 1977). Thus despite the scientific

487 endeavors of numerous research groups over a considerable period of time the explanation  
488 for the frequently monitored CH<sub>4</sub> oversaturation of oxic surface waters in oceans and fresh  
489 water lakes is still a topic of debate (Zindler et al., 2013; Tang et al., 2014; Damm et al., 2008).  
490 Since our results unambiguously show that the common coccolithophore *E. huxleyi* is able to  
491 produce CH<sub>4</sub> *per se* under oxic conditions we thus suggest that algae living in marine  
492 environments might contribute to the regional and temporal oversaturation of surface waters.  
493 However, our results of the laboratory experiments should be confirmed by field  
494 measurements in the ocean.

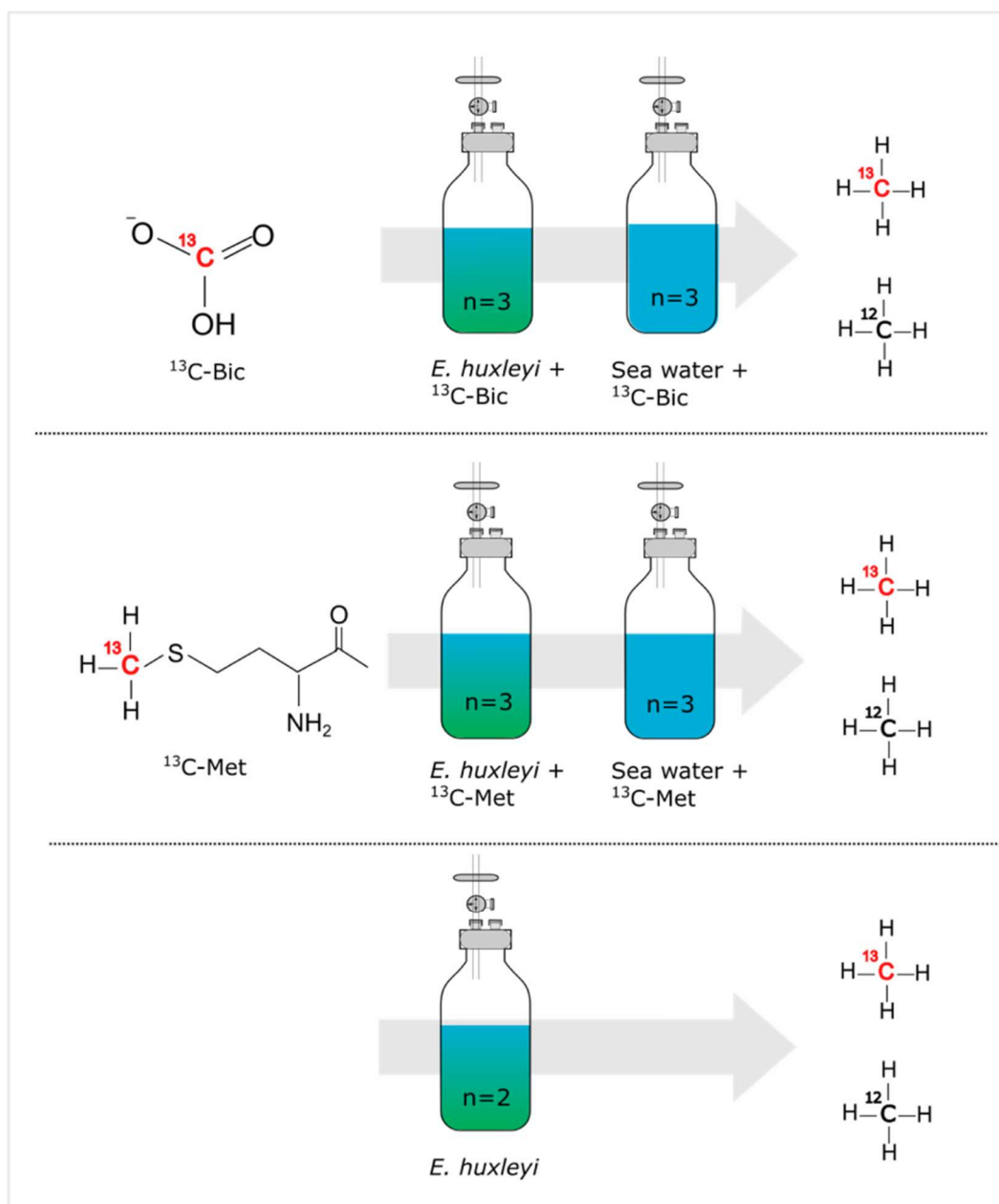
495 We would encourage further studies in this research area to make use of stable isotope  
496 techniques together with field measurements as we consider such an approach well suited for  
497 the elucidation of the pathways involved in CH<sub>4</sub> formation in oceanic waters.

498

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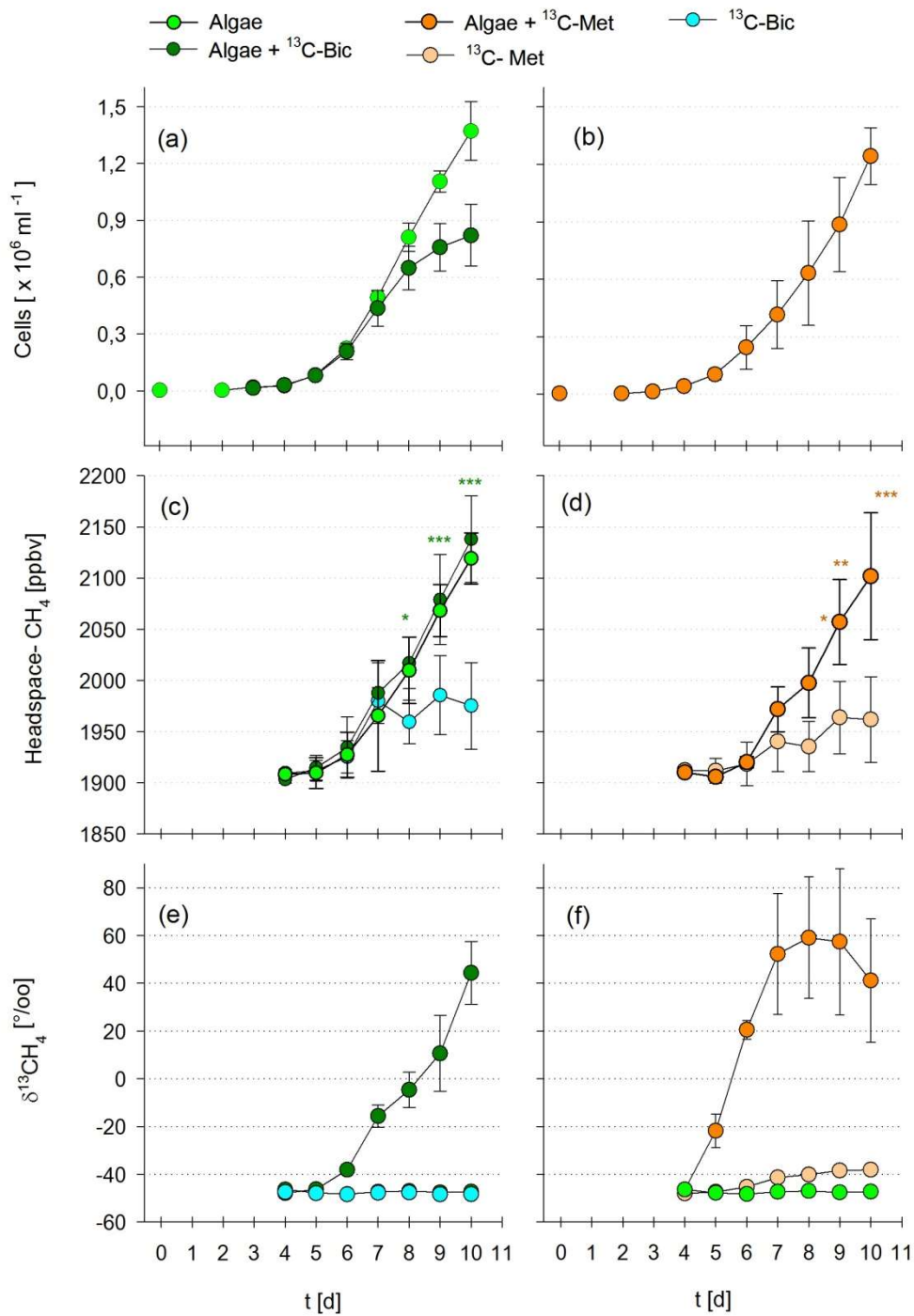




508

509 **Fig. 1:** Experimental. The potential precursors of  $\text{CH}_4$ ,  $^{13}\text{C}$ -labelled bicarbonate ( $^{13}\text{C-Bic}$ ) or a  
 510 position-specific  $^{13}\text{C}$ -labelled methionine ( $^{13}\text{C-Met}$ ) were added to the flasks containing either  
 511 a culture of *E. huxleyi* or sea water only.

512



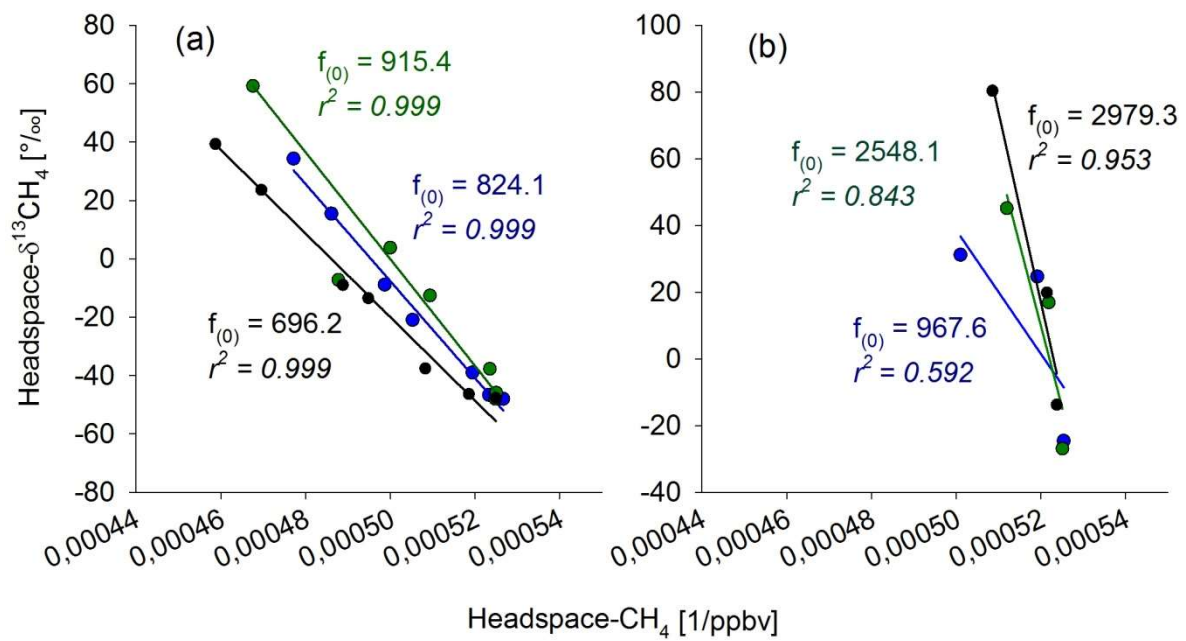
513

514 **Fig. 2:** Culture cell density when algae grown in seawater (n = 2) supplemented with (a) Bic or  
 515 (b) Met (n = 3) and headspace CH<sub>4</sub> mixing ratio for cultures supplemented with (c) Bic or (d)  
 516 Met.  $\delta^{13}\text{C-CH}_4$  values after addition of (e) <sup>13</sup>C-Bic and (f) <sup>13</sup>C-Met (n = 3; error bars mark the

517 standard deviation). Stars mark the significance between “algae +  $^{13}\text{C}$ -Bic” and “sea water +  $^{13}\text{C}$ -Bic” or between “algae +  $^{13}\text{C}$ -Met” and “sea water +  $^{13}\text{C}$ -Met”, respectively, with \* $p \leq 0.05$ ;  
 518  $^{13}\text{C}$ -Bic” or between “algae +  $^{13}\text{C}$ -Met” and “sea water +  $^{13}\text{C}$ -Met”, respectively, with \* $p \leq 0.05$ ;  
 519 \*\* $p \leq 0.01$ ; \*\*\* $p \leq 0.001$ .

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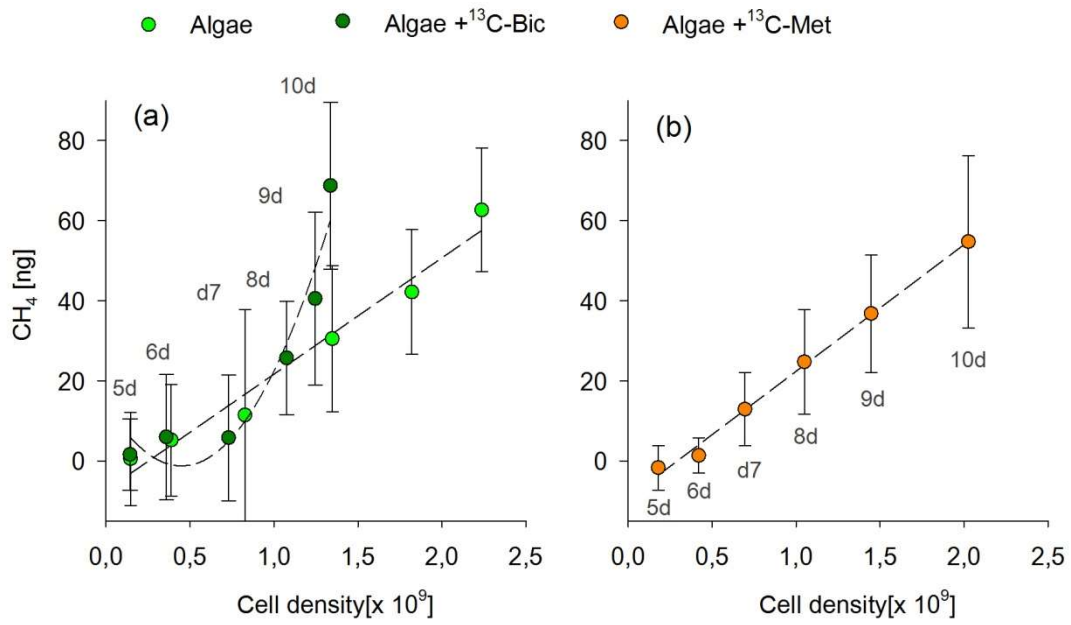
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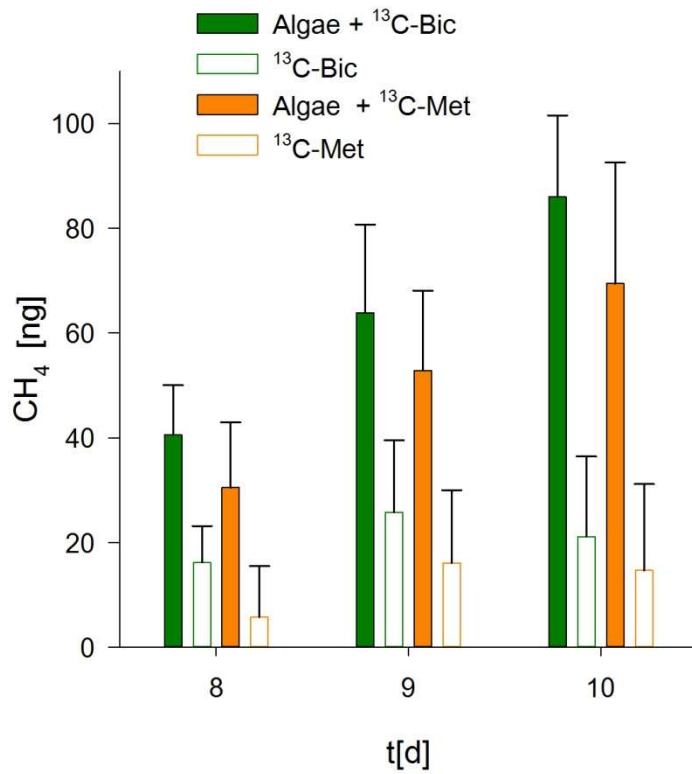
523 **Fig. 3:** Keeling-plots for the treatment (a) “algae +  $^{13}\text{C}$ -Bic” and (b) “algae +  $^{13}\text{C}$ -Met”, where  
 524  $f_{(0)}$  refers to the  $^{13}\text{C}$  value of the  $\text{CH}_4$ -source.

525



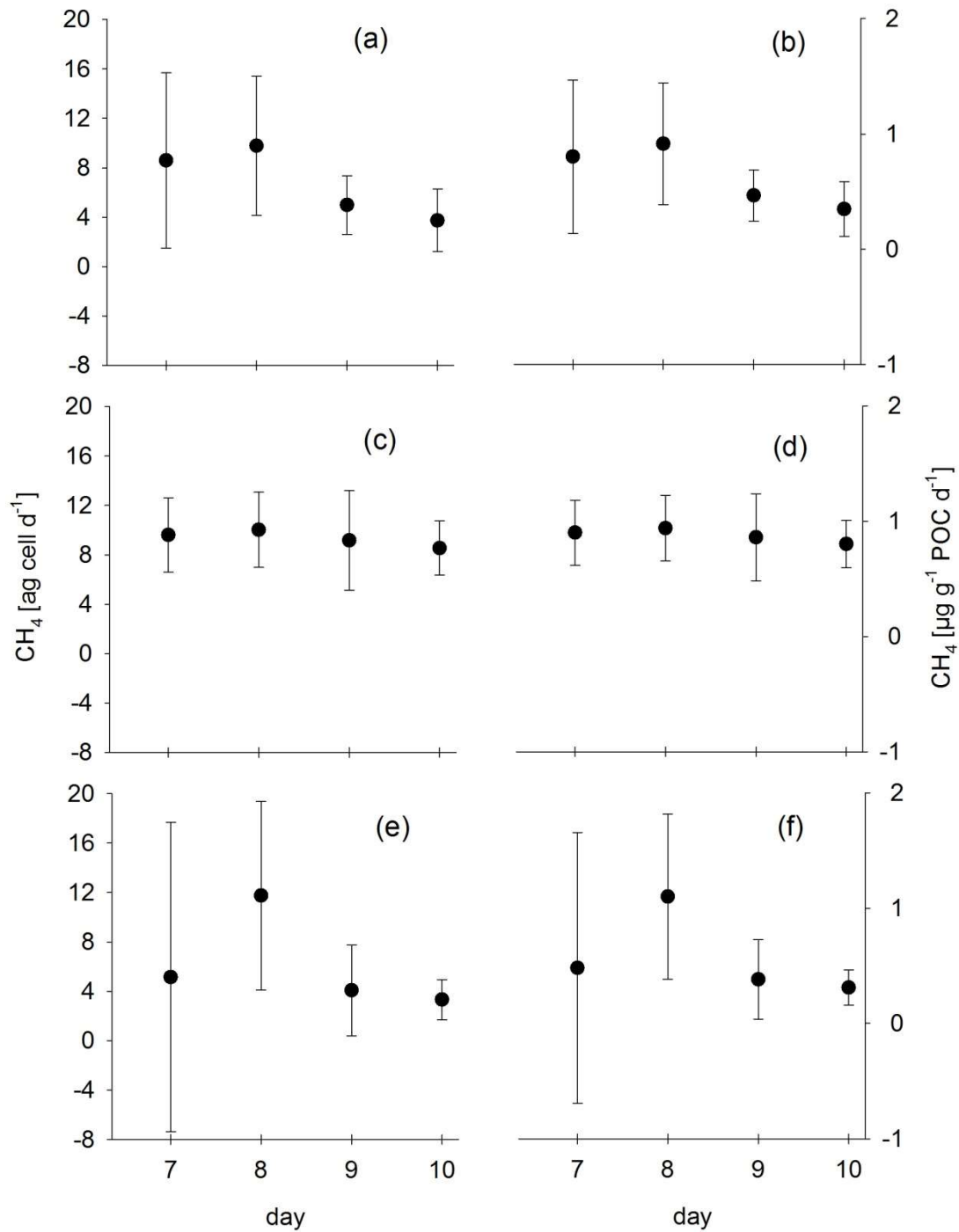
526

527 **Fig. 4:** Correlation between cell density per flask and CH<sub>4</sub> content (sum of headspace and  
 528 water phase) for the coccolithophore *E. huxleyi* (a) in seawater only (n = 2), supplemented  
 529 with (a) <sup>13</sup>C-labelled bicarbonate (Bic) or (b) methionine (Met) (n = 3); error bars mark the  
 530 standard deviation; d = day of incubation.



531

532 **Fig. 5:** Mean CH<sub>4</sub> content (sum of headspace and water phase) in the flasks of *E. huxleyi*  
 533 supplemented with either bicarbonate or methionine (n = 3) and the respective control  
 534 without algae (n = 2) measured at days 8, 9 and 10; error bars show the standard deviation.



535

536 **Fig. 6:** Daily CH<sub>4</sub> production of *E. huxleyi* for days 7 to 10 (a, c, e) on a per cell basis and (b, d,  
 537 f) relative to particulate organic carbon (POC) separately for the treatments (a, b) *E. huxleyi* +  
 538 <sup>13</sup>C-Bic (n = 3), *E. huxleyi* + <sup>13</sup>C-Met (n = 3), and *E. huxleyi* (n = 2). Values are presented as means  
 539 with the standard deviation.

540

541 **Tables:**542 **Tab. 1:** Overview of sample collection during the incubation of *E. huxleyi*.

Day		0	1	2	3	4	5	6	7	8	9	10
Headspace	CH <sub>4</sub>					x	x	x	x	x	x	x
	δ <sup>13</sup> CH <sub>4</sub>					x	x	x	x	x	x	x
Water	cell density	x			x	x	x	x	x	x	x	x

543

544 **Tab. 2:** Mean daily CH<sub>4</sub> production rates of *E. huxleyi* (\*n = 2; \*\*n = 3) determined between545 days 7 and 10, ag = attogramm = 10<sup>-18</sup>.

Treatment	CH <sub>4</sub> [ag cell <sup>-1</sup> d <sup>-1</sup> ]	CH <sub>4</sub> [μg g <sup>-1</sup> POC d <sup>-1</sup> ]
<i>E. huxleyi</i> + <sup>13</sup> C-Bic**	6.8 ± 4.1	0.63 ± 0.39
<i>E. huxleyi</i> + <sup>13</sup> C-Met**	9.3 ± 2.6	0.88 ± 0.24
<i>E. huxleyi</i> *	6.1 ± 3.7	0.57 ± 0.35

546

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