

1 **We would like to thank all reviewers for their critical comments, which we think helped to improve**
2 **the quality and clarity of this manuscript. We hope our responses and adaptations are adequate to**
3 **accept this manuscript for publication in Biogeosciences. Please find our detailed responses below.**

4
5 **The uploaded revised manuscript includes the reviewer changes as well as revised English spelling**
6 **and grammar.**

7 **Ronald Oremland, Referee #1**

8 Received and published: 15 September 2015

9 The manuscript by Maltby et al. examines the processes of methanogenesis and
10 sulfate-reduction along a transect of seafloor transiting a near-shore depositional-rich
11 region to an offshore, deeper sediment locale. The work was done on the Peruvian
12 shelf, a region of high productivity and oxygen-minimum zone/anoxic bottom waters.
13 The work also acquired a number of relevant parameters along with the bioassays.
14 The main finding was that the shallow sediments nearest shore had high rates of
15 methanogenesis at the sediment surface, implying a contribution of non-competitive
16 substrate precursors (e.g., methylated amines and methylated sulfides) as precursors of methane.
17 Methanol was added as a proxy non-competitive substrate while molybdate was employed to detect
18 use of competitive substrates to underscore this point. Although considerable work has been done
19 along these lines in salt-marsh sediments and hypersaline systems, very little has been done in open
20 marine systems like the one described here.

21 I do not have substantive technical criticisms, but offer the following points to strengthen and clarify
22 the manuscript:

23 page 14871, lines 26-27: the logic here is not obvious that H₂/acetate increase with
24 depth as organic matter becomes more recalcitrant.

25 **Authors Reply:** We agree with the reviewer and decided to delete this sentence to avoid confusion.

26 page 14872, lines 27 - 28: some statement should be made about probable sources of non-
27 competitive substrate precursors, such as degradation of organic osmolytes (e.g., DMSP, betaine).

28
29 **Authors Reply:** We thank the reviewer for this comment and added this information in the text.

30
31 page 14878, lines 2-3: since the sediments were mixed with bottom water, which contained abundant
32 sulfate, these rates may underestimate the potential of deeper sediment regions where sulfate is
33 low.

34
35 **Authors Reply:** The focus of the paper was to determine methanogenesis activity within the sulfate
36 reduction zone, i.e., in the presence of sulfate. Within the investigated sediment (0-30 cmbsf) sulfate
37 was always above 9 mM in situ. We are therefore confident that the detected methanogenesis activity
38 reflects its potential under the given environmental conditions.

39
40 page 14879, line 1 (and elsewhere, page 14885 bottom): what percentage of the added 10 mM
41 methanol went to CH₄ (plus CO₂) in the incubations?

42
43 **Authors Reply:** We expect that close to 100 % of the methanol was converted to CH₄ (plus CO₂). We
44 did similar experiments in another study where we followed the conversion of methanol by ¹³C-

45 Labeling. However, as we did not conduct these experiments in the current study, we rather do not
46 want to make any assumptions.

47
48 Minor corrections:
49 page 14870, line 1: co-occurrence (concurrence implies an agreement)

50
51 **Authors Reply: Done**

52
53 page 14870, line 10: multiple cores (not multicorer cores)

54
55 **Authors Reply: Done**

56
57 page 14870, line 23: decrease (not decline)

58
59 **Authors Reply: Done**

60
61 page 14873, line 15: ...an environment where both...(no comma)

62
63 **Authors Reply: Done**

64
65 page 14875, line 11: a 5 m steel barrel (not "steal" unless the authors actually pilfered the corer from
66 another lab)

67
68 **Authors Reply: Done. We would never steal from our colleagues ;-)**

69
70 page 14877, line 24 - 25: sliced into 5 cm

71
72 **Authors Reply: Done**

73
74 page 14878, line 28: molybdate

75
76 **Authors Reply: Done**

77
78 page 14882, line 2: a grey color

79
80 **Authors Reply: Done**

81
82 page 14887, line 19: co-occurred

83
84 **Authors Reply: Done**

85
86 page 14889, line 23: were (not "where")

87
88 **Authors Reply: Done**

89
90 page 14895, line 10: were (not "was")

91
92 **Authors Reply: Done**

93
94
95
96
97

98

99 **Anonymous Referee #2**

100 Received and published: 19 October 2015

101 The manuscript of Maltby et al. describes rates of sulfate reduction and methanogenesis
102 were measured in various radiotracer incubations. The study highlights the role
103 of methanogenesis in near-surface sediments (here termed shallow methanogenesis)
104 in overall carbon mineralization. Methodologically the study is extremely well designed
105 and the experimental setup is flawless.

106
107 The only flaw that I see in this paper is in the treatment of the bag incubations in relation to the
108 whole-core incubations. While whole core incubations are next best thing to in-situ experiments with
109 benthic landers (which come with their own set of problems and limitations), bag experiments for
110 rate measurements will definitely give results that are different to measurements on intact
111 sediment cores. Numerous studies have reported the effects of structural disturbance
112 on turnover rates. Although the bag experiments were only performed in order to study
113 the effect of various substrate additions, especially non-competitive substrates, the measured
114 rates are presented in a way that the reader might get the impression that these
115 rates are actually comparable to the whole core incubation data. I would therefore suggest
116 to stress the differences between the whole core and bag incubations and discuss
117 the limitations of the different techniques.

118
119 **Authors Reply:** We thank the reviewer for this helpful comment. There is indeed a difference
120 between the sulfate reduction rates (whole core method) and the net methanogenesis rates (slurry
121 incubations with anoxic deep water). The additional experiments with addition of substrate (slurry
122 incubations with artificial seawater) are marked by a different title: “potential methanogenesis”,
123 which stresses the difference compared to net methanogenesis and sulfate reduction. However, we
124 agree that we have to point out the differences in net methanogenesis rates and sulfate reduction
125 rates during our comparison. Therefore, we added this information to the discussion.

126
127 Minor comments:

128
129 p14872, line 26: Why do these conditions favour methanogenesis, anoxia and fresh organic matter
130 are also perfect conditions for sulfate reduction

131
132 **Authors Reply:** Methanogens have a high sensitivity to oxygen (sulfate reducers tolerate oxygen
133 much better). We argue that the depletion of oxygen in the bottom water (and with that also
134 absence of bioirrigation) allow methanogens to colonize and thrive close to (or at the) sediment-
135 water interface. We added a few words to clarify this point.

136
137 p14873, line 2: As far as I know Limfjorden sediment is permanently anoxic, at least below the upper
138 few mm, only the oxygen concentration in the water column changes over the year. I think
139 this sentence should be rephrased to avoid confusion.

140
141 **Authors Reply:** We agree with the reviewer and changed the sentence.

142
143
144 p14875 line 8 and 15: Why did you process the samples in two different cold rooms with different
145 temperatures?

146

147 **Authors Reply:** This was a matter of space. The cores from the multicores were all processed in a 9°C
148 cold laboratory container, which was used by different scientific parties on board. When we
149 processed the gravity core, space was limited so we moved to the 4 °C cold room (a storage room),
150 which was not acclimated to 9°C.

151
152 p14875 line 11: I still think that you paid for the barrel on your corer and did not steal
153 it...

154
155 **Authors Reply:** Done.

156
157 p14878, line 21: What do you mean by "transferred completely"? Did you do a
158 quantitative transfer or did you fill the bottle without headspace?

159
160 **Authors Reply:** We filled the bottles without headspace. To avoid confusion, we changed the
161 sentence.

162
163 p14879, line 27f: Section 2.3 describes porewater sampling, not rate measurements. What do you
164 mean by "according to the above scheme"? Did you use a slurry? How did you get the sediment
165 into the glass syringes? Or do you mean the old Jørgensen glass barrels (Glass tube
166 with syringe plunger)?

167
168 **Authors Reply:** "According to the above scheme" refers to the sampled sediment depths, not the
169 type of measurements. We changed the wording to make it clearer.
170 By "glass syringes" we indeed meant the Jørgensen glass barrels. We changed the sentence
171 accordingly.

172
173 p14880, line 9f: Why did you do change your technique? I always thought that the old one was just
174 fine?

175
176 **Authors Reply:** Absolutely. It is basically the same method just that for the methanogenesis rate
177 calculation you need the total DIC concentration and not the total methane concentration (which
178 you need for AOM calculation). We therefore did DIC analyses instead of gas chromatography.

179
180 p14887, line 12: Why didn't you use for example the SO₄ or DIC PW profile to align the cores?
181 Comparison between the topmost Gravity Core sample and the MUC cores should give you a
182 reasonable estimate how much sediment was blown off by the Gravity Core

183
184 **Authors Reply:** We did look at the SO₄ profiles to check if the statement of ~ 20 cm fits in our case.
185 However, as the sampling intervals in the gravity core were rather large, we did not feel comfortable
186 to align the cores.

187
188 p14889, line 13-15: Please show the data, this could be important.

189
190 **Authors Reply:** As the deep sediment layers are not the focus of the presented study, we do not
191 think adding the iron data will change the view on our findings and rather distract from our story. We
192 therefore refrain from showing this data.

193
194 p14890, line 21: To me the term "transport velocity" implies an active movement, which would only
195 be important in zones with active fluid flow. Here we are talking about purely diffusive systems and I
196 would recommend sticking to those to avoid confusion.

197

198 **Authors Reply:** In this context (introducing the SMTZ) we actually meant both diffusive and advective
199 transport. We changed the wording to "flux" as a more neutral term, which considers both diffusive
200 and advective transport.
201

202

203 **Microbial methanogenesis in the sulfate-reducing zone of surface sediments traversing the**
204 **Peruvian margin**

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211

212 **Abstract**

213 We studied the concurrence of methanogenesis and sulfate reduction in surface sediments (0-25 cm
214 below seafloor) at six stations (70, 145, 253, 407, 990 and 1024 m) along the Peruvian margin (12°S).

215 This oceanographic region is characterized by high carbon export to the seafloor creating an extensive
216 oxygen minimum zone (OMZ) on the shelf, both factors that could favor surface methanogenesis.

217 Sediments sampled along the depth transect traversed areas of anoxic and oxic conditions in the
218 bottom-near water. Net methane production (batch incubations) and sulfate reduction (³⁵S-sulfate

219 radiotracer incubation) were determined in the upper 0-25 cmbsf of ~~multiple~~ cores from all
220 stations, while deep hydrogenotrophic methanogenesis (> 30 cmbsf, ¹⁴C-bicarbonate radiotracer

221 incubation) was determined in two gravity cores at selected sites (78 and 407 m). Furthermore,
222 stimulation (methanol addition) and inhibition (molybdate addition) experiments were carried out to
223 investigate the relationship between sulfate reduction and methanogenesis.

224 Highest rates of methanogenesis and sulfate reduction in the surface sediments, integrated over 0-25
225 cmbsf, were observed on the shelf (70-253 m, 0.06-0.1 mmol m⁻² d⁻¹ and 0.5-4.7 mmol m⁻² d⁻¹,

226 respectively), while lowest rates were discovered at the deepest site (1024 m, 0.03 and 0.2 mmol m⁻² d⁻¹,
227 respectively). The addition of methanol resulted in significantly higher surface methanogenesis

228 activity, suggesting that the process was mostly based on non-competitive substrates, i.e., substrates
229 not used by sulfate reducers. In the deeper sediment horizons, where competition was probably

230 relieved due to the decrease~~line~~ of sulfate, the usage of competitive substrates was confirmed by the
231 detection of hydrogenotrophic activity in the sulfate-depleted zone at the shallow shelf station (70 m).
232 Surface methanogenesis appeared to be correlated to the availability of labile organic matter (C/N
233 ratio) and organic carbon degradation (DIC production), both of which support the supply of
234 methanogenic substrates. A negative correlation ~~between~~ methanogenesis rates ~~and~~ dissolved
235 oxygen in the bottom-near water was not obvious, however, anoxic conditions within the OMZ might
236 be advantageous for methanogenic organisms at the sediment-water interface.

237 Our results revealed a high relevance of surface methanogenesis on the shelf, where the ratio between
238 surface to deep (below sulfate penetration) methanogenic activity ranged between 0.13 and 10⁵. In
239 addition, methane concentration profiles indicated a partial release of surface methane into the water
240 column as well as a partial consumption of methane by anaerobic methane oxidation (AOM) in the
241 surface sediment. The present study suggests that surface methanogenesis might play a greater role in
242 benthic methane budgeting than previously thought, especially for fueling AOM above the sulfate-
243 methane transition zone.

244

245

246 *Keywords: Oxygen minimum zone, organic matter, competition, anaerobic oxidation of methane,*
247 *emission*

248

249

250 **1. Introduction**

251 Microbial methanogenesis represents the terminal step of organic matter degradation in marine
252 sediments (Jørgensen, 2006). The process is entirely restricted to a small group of prokaryotes within
253 the domain of the Archaea (Thauer, 1998). Methanogens produce methane from a narrow spectrum of
254 substrates, primarily carbon dioxide (CO₂) and hydrogen (H₂) (hydrogenotrophic pathway), as well as
255 acetate (acetoclastic pathway) (Zinder, 1993). In addition, methanol or methylated compounds such as
256 methylamine ~~can be~~ utilized (methylotrophic pathway) (Oremland & Polcin, 1982; Buckley et al.,
257 2008; Zinder, 1993; King et al., 1983). Substrates for methanogenesis are produced during
258 depolymerization and fermentation of organic macromolecules (e.g., sugars, vitamins, amino acids) to
259 smaller monomeric products (Jørgensen, 2006; Schink & Zeikus, 1982; Neill et al., 1978; Donnelly &
260 Dagley, 1980).

261 Acetoclastic and hydrogenotrophic methanogenesis are predominantly found in deeper sediment zones
262 below sulfate penetration, owing to the ~~more effective utilization of~~ ~~competition with sulfate reducers~~
263 ~~that outcompete methanogens for~~ H₂ and acetate ~~by sulfate reducers~~ due to their higher substrate
264 affinity (Oremland & Polcin 1982; Jørgensen 2006). ~~Furthermore, CO₂/H₂ and acetate are the more~~
265 ~~abundant substrates in deeper sediments as degradability of organic matter, and with it the substrate~~
266 ~~variety and availability, decreases with increasing sediment depth (Jørgensen, 2006).~~

267 Methanogens ~~can~~ avoid competition with sulfate reducers by the utilization of non-competitive
268 substrates, such as methanol or methylamines (Oremland & Polcin, 1982; King et al., 1983).
269 Facilitated by the usage of such non-competitive substrates, sulfate reduction and methanogenesis
270 were found to co-occur in sulfate-containing salt marsh sediments (Oremland et al., 1982; Buckley et
271 al., 2008; Senior et al., 1982). Concurrent activity of sulfate reduction and methanogenesis in the
272 marine environment has mostly been postulated for organic-rich sediments (Mitterer, 2010; Jørgensen
273 & Parkes, 2010; Treude et al., 2009, 2005a; Hines & Buck, 1982; Crill & Martens, 1986); however,
274 ~~details research on the~~ magnitude and environmental controls of surface methanogenesis ~~are is~~ still
275 ~~poorly understood sparse~~ (Holmer & Kristensen, 1994; Ferdelman et al., 1997).

276 In a study from Eckernförde Bay, southwestern Baltic Sea, considerable in vitro methanogenic activity
277 was observed in samples taken from 5 to 40 cm sediment depth (Treude et al. 2005). Although in vitro

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278 activity was measured in sulfate-free setups, methanogenic activity coincided with zones of in-situ
279 sulfate reduction. The authors concluded a coexistence of the two types of organisms, which could be
280 enabled through either the usage of non-competitive substrates, dormancy of methanogens until phases
281 of sulfate depletion, and/or temporal or spatial heterogeneity in the sediments. Eckernförde Bay
282 sediments feature a high input of organic matter due to a shallow water depth (~30 m) and pronounced
283 phytoplankton blooms in spring, summer, and fall (Smetacek, 1985). Furthermore, seasonal hypoxia
284 ($O_2 < 90 \mu M$) or even anoxia ($O_2 = 0 \mu M$) occur in the deep layers of the water column caused by
285 stratification and degradation of organic matter (Bange et al. 2011). Oxygen-depleted conditions in
286 the bottom water together with frequent input of fresh organic matter possibly favors methanogenesis
287 in surface sediment by offering reduced conditions and non-competitive substrates. As non-
288 competitive substrates can be derived from organic osmolytes such as betaine or
289 dimethylsulfoniopropionate (DMSP), a high load of organic matter (e.g. through sedimentation of
290 phytoplankton blooms) can increase the availability of non-competitive substrates (Zinder, 1993; **Van**
291 **Der Maarel & Hansen, 1997**). In accordance, methanogenesis activity was observed within the sulfate-
292 reducing zone of organic-rich sediments and seasonally hypoxic sediments from the seasonally
293 hypoxic Limfjorden sound, Northern Denmark (Jørgensen & Parkes, 2010; Jørgensen, 1977).

294
295 The environmental relevance of surface methanogenesis is hitherto unknown. Its closeness to the
296 sediment-water interface makes it a potential source for methane emissions into the water column,
297 unless the methane is microbially consumed before escaping the sediment (Knittel & Boetius, 2009).
298 Methane escapes the sediment either by diffusion or, when methane saturation is exceeded, in the form
299 of gas bubbles (Whiticar, 1978; Wever & Fiedler, 1995; Judd et al., 1997; Dimitrov, 2002). The
300 fraction How much of the released methane released to the water column that reaches the atmosphere
301 mainly depends on water depth, as methane is also consumed within the water column through aerobic
302 microbial oxidation (Reeburgh, 2007; Valentine et al., 2001). Thus, shallow coastal areas have higher
303 methane emission potentials than the open ocean (Bange et al., 1994) and a greater potential to
304 contribute to methane-dependent. Once in the atmospheric warming, methane acts as a very potent
305 greenhouse gas (IPCC, 2014).

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306

307 In the present study we focused on the upwelling region off the Peruvian coast, which is another
308 ~~excellent~~ example of an environment, where both factors that potentially favor surface methanogenesis
309 convene, i.e., a high export of organic carbon and low dissolved oxygen concentrations in the bottom
310 water. This upwelling region represents one of the most productive systems in the world oceans,
311 creating one of the most intense oxygen minimum zones (OMZ, Kamykowski & Zentara 1990;
312 Pennington et al. 2006). Oxygen concentrations in waters impinging on the seafloor are below 20 μM
313 or even reach anoxia. Research on surface sediment methanogenesis in upwelling regions is ~~rare~~
314 scarce and its potential role in the carbon cycling of the Peruvian OMZ is completely unknown. In a
315 study from the central Chilean upwelling area (87 m water depth, 0.5-6 cm sediment depth), ~~low~~
316 methane production ~~rates of methane was were~~ detected despite high sulfate reduction activity, when
317 ~~offering~~ the non-competitive substrate trimethylamine was offered (Ferdelman et al., 1997). The
318 authors concluded that the prevailing methanogens were competing with sulfate reducers for H_2 and
319 with acetogens for methylamines, explaining the overall low methanogenesis activity observed
320 (Ferdelman et al., 1997).

321 Even though the Chilean and Peruvian OMZs are connected, commonly known as OMZ in the
322 Eastern South Pacific Ocean (ESP) (Fuenzalida et al., 2009), the core of the ESP-OMZ is centered
323 off Peru with an upper boundary at < 100 m and a vertical distribution to > 600 m versus a thinner
324 OMZ band off Chile constrained between 100-400 m water depth (Fuenzalida et al., 2009). ~~The~~
325 anoxic conditions in the water column of the OMZ core (and therewith a lack of bioirrigating
326 macrofauna introducing oxygen into the sediments (Kristensen, 2000)) together with the high export
327 rates of labile organic carbon to the seafloor (Reimers & Suess, 1983; Dale et al., 2015) provide
328 favorable conditions for methanogenesis activity in surface sediments, thus increasing the potential for
329 benthic methane emissions.

330 Here, we provide first insights into surface methanogenesis in sediment cores (< 30 cmbsf =
331 centimeters ~~below~~ seafloor) taken along the Peruvian shelf and margin. We hypothesize that
332 methanogenesis coexists with sulfate reduction through the utilization of non-competitive substrates.
333 In addition, we postulate that surface methanogenesis depends on the quantity and quality (=

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334 freshness) of organic carbon, and the concentrations of dissolved oxygen in the bottom water. We
335 therefore expect spatial variability of surface methanogenesis along the continental shelf and margin.
336 The observed methanogenic activity will be compared to methane concentrations in the bottom-near
337 water to discuss the potential relevance of surface methanogenesis for methane emissions into the
338 pelagic zone.

339

340 **2. Material and Methods**

341 **2.1 Study site and sediment sampling**

342 Samples were taken during the R.V. Meteor cruise M92 between 5. Jan and 3. Feb 2013 along a depth
343 transect off the Peruvian coast from the shelf (~70 m) to the continental [margin slope](#) (~1000 m). The
344 transect was located in the central part of the ESP-OMZ (Fuenzalida et al., 2009) at 12°S. Further
345 hydrographic details on the study area can be found elsewhere (Dale et al., 2015).

346 Sediment cores for the determination of near-surface methanogenesis were collected at six stations
347 along the depth transect at 70, 145, 253, 407, 776 and 1024 m water depth (Fig.1), using a multiple
348 corer with a mounted camera (TV-MUC). The MUC held seven cores (length: 60 cm, inner diameter:
349 10 cm) and covered an area of ~1 m². If necessary, a second MUC was deployed at the same station,
350 thus sediment cores could originate from different MUC casts. Station numbers were assigned in
351 accordance with Dale et al., (2015). After retrieval, sediment cores were transferred to a ~9°C cold
352 room and processed at the same day.

353 In addition to the MUC, a gravity corer was deployed at two stations (78 and 407 m) for determining
354 deep methanogenesis. The total core length was 400 cm and 206 cm, respectively. The gravity corer
355 was equipped with a 260 kg weight and a 5 m [steel](#) barrel (diameter: 14 cm). The replaceable core
356 liner (PVC, diameter: 12.5 cm) was housed within the barrel and fixed with a core catcher. After
357 retrieval, sediment cores from the gravity corer were sliced into 1-m sections, capped on both sides,
358 and brought to the cold room (4°C) for further processing. Relevant station details for MUC and
359 gravity cores are summarized in Table 1.

360

361 **2.2 Water column sampling**

362 CTD/Rosette water column casts were conducted at the same station as sediment coring (for details
363 see Table 1). Temperature and oxygen data ~~were~~ taken from Dale et al. 2015.

364 For the analysis of methane concentrations in the bottom-near water, water was sampled ca. 1.5 m
365 above the seafloor from 10 L Niskin bottles mounted on ~~the a-CTD/Rosette water sampler~~. The
366 collected water was filled bubble-free into 60 ml vials (triplicates), each vial containing 3 pellets of
367 sodium hydroxide (NaOH, ~0.3 M per vial) to stop microbial activity and force dissolved gas into the
368 headspace. After closing the vials with a butyl rubber stopper and a crimp seal, 10 ml of water was
369 removed with a N₂-flushed 10 ml syringe and replaced with N₂ gas from a second syringe to create a
370 headspace in the sampling vials. Samples were stored and transported at room temperature until
371 further processing.

372 In the home laboratory, 100 µl of the headspace volume was injected into a Shimadzu GC-2014 gas
373 chromatograph equipped with a flame ionization detector and a HaySep-T 100/120 column (Length 3
374 m, diameter: 2 mm). Gases were separated isothermally at 75°C with helium carrier gas. Methane
375 concentrations were calibrated against methane standards (Scotty gases). The detection limit was 0.1
376 ppm with a precision of 2 %.

377

378 **2.3 Porewater geochemistry**

379 Porewater sampling for MUC cores has been previously described by Dale et al., (2015). In short, one
380 MUC core per station was subsampled in an argon-filled glove bag, to preserve redox sensitive
381 constituents.

382 The gravity cores at St. 1 (78 m) and St. 8 (407 m) were subsampled at 10-12 different sediment
383 depths (depending on core length) resulting in depth intervals of 20-33 cm. Before sampling, the
384 plastic core liner was cut open with an electric saw at the specific depths. Porewater was extracted by
385 using anoxic (flushed with argon), wetted rhizons (Rhizosphere Research Products, Seeberg-Elverfeldt
386 et al., 2005).

387 Sulfate concentrations were determined by ion chromatography (Methrom 761) as described
388 previously by Dale et al., (2015).

389 For DIC analysis, 1.8 ml of porewater was transferred into a 2 ml glass vial, fixed with 10 µl saturated
390 mercury chloride solution and crimp sealed. Samples were stored at 4°C until further processing in the
391 home laboratory. DIC concentration was determined as CO₂ with a multi N/C 2100 analyzer (Analytik
392 Jena) following the manufacturer's instructions. Therefore the sample was acidified with phosphoric
393 acid and the outgassing CO₂ was measured. The detection limit was 20 µM with a precision of 2-3%.

394

395 **2.4 Sediment porosity and particulate organic carbon/nitrogen**

396 Methodology and data for porosity, particulate organic carbon (POC) and particulate organic nitrogen
397 (PON) have been previously described by Dale et al., (2015).

398 In short, wet sediment samples were taken from the porewater MUC core and the gravity cores for
399 determination of porosity from the weight difference of wet and freeze-dried sediment. POC and PON
400 were analyzed with a Carlo-Erba element analyzer (NA 1500). Ratios of POC:PON were calculated by
401 division.

402

403 **2.5 Sediment methane**

404 For sediment methane concentration, one MUC core per station was sliced in 2 cm intervals until 20
405 cm depth, followed by 5 cm intervals until the end of the core (maximum depth = 48 cm). Gravity
406 cores were subsampled according to the above scheme (see 2.3). From each sampled sediment layer, 2
407 cm³ sediment were transferred into a 15 ml serum glass vial containing 5 ml of NaOH (2.5% w/w).
408 The vial was closed with a butyl stopper, crimp sealed and shaken thoroughly to stop microbial
409 activity and to force all methane into the headspace. Vials were stored upside down at room
410 temperature until measurement in the home laboratory.

411 Sediment methane concentration was determined by injecting 0.1 ml of headspace volume into a
412 Shimadzu GC-2014 gas chromatograph as described under section 2.2.

413

414 **2.6 Net methanogenesis activity in MUC cores**

415 Sediment from MUC cores was used to determine net methanogenesis, which is defined as the sum of
416 total methane production and consumption, including all available methanogenic substrates in the
417 sediment. Net methanogenesis was determined by measuring the linear increase of methane
418 concentration in the headspace of closed incubation vials over time. Therefore, one MUC core per
419 station was sliced into 5 cm intervals, transferring 10 cm³ of sediment in triplicates into a N₂-flushed
420 60 ml serum glass vials. The sediment core lengths ranged from between 25 up to 48 cm, resulting in
421 maximum 10 depth intervals. Ten ml of anoxic deep water overlying each MUC core was added to the
422 vial and the slurry was mixed under a constant N₂ stream (Hungate, 1950) before being sealed with a
423 butyl rubber stopper and crimped. The sediment slurry was repeatedly flushed with N₂ through the
424 stopper to ensure guarantee fully anoxic conditions. The vials were incubated in the dark and at 9°C,
425 which reflected the average in situ temperature along the depth transect (see Table 1). The first gas
426 chromatographic measurement was done directly after preparation of the vials, by injecting 100 µl of
427 headspace sample into the gas chromatograph. The on-board Hewlett Packard-5890 gas
428 chromatograph was equipped with a flame ionization detector and a HaySep-T 100/120 column
429 (Length 3m, diameter: 2mm). Gases were separated isothermally at 75°C with helium carrier gas.
430 Methane concentrations were calibrated against methane standards. The detection limit was 1 ppm
431 with a precision of < 5%. Measurements were done in 2-4 day-intervals over a total incubation time of
432 ~2 weeks.

433

434 **2.7 Potential non-competitive and competitive methanogenesis in sediment slurries from MUC** 435 **cores**

436 Sediment slurry experiments were conducted with sediment from St. 1 (70 m) to examine the
437 interaction between sulfate reduction and methanogenesis, as this station revealed highest microbial
438 activity of sulfate reduction and methanogenesis. On board, the sediment core was sliced in 5 cm
439 intervals. Sediment from the 0-5 cm interval and the 20-25 cm interval was transferred completely into
440 250 ml glass bottles, which were then closed without headspace (filled to top) with a butyl rubber
441 stopper and screw cap. Until further treatment, sediment was stored at 4°C on board and later in a 1°C
442 cold room on shore.

443 Approximately 6 months after the cruise, sediment slurries from both depth intervals were prepared by
444 mixing 5 ml sediment in a 1:1 ratio with artificial, fully marine seawater (Widdel & Bak, 1992) before
445 further manipulations.

446 In total, three different treatments, each in triplicates, were prepared per depth: 1) sulfate-rich (28
447 mM), serving as a control 2) sulfate-rich plus molybdate (22 mM) from now on referred to as
448 molybdate-treatment, and 3) sulfate-rich plus methanol (10 mM) from now on referred to as methanol-
449 treatment.

450 Molybdate was used as an enzymatic inhibitor for sulfate reduction (Oremland & Capone, 1988).

451 Methanol is a known non-competitive substrate used by methanogens, but not by sulfate reducers
452 (Oremland & Polcin, 1982), which makes it suitable to examine non-competitive methanogenesis.

453 The sediment slurries were incubated at 9°C in the dark for 23 days and headspace concentration of
454 methane was measured repeatedly over time on a gas chromatograph. Therefore, 100 µl of headspace
455 was removed from the gas vials and injected into a Shimadzu gas chromatograph (GC-2014) equipped
456 with a methanizer (inactive), a packed Haysep-D column and a flame ionization detector. The column
457 temperature was 80°C and the helium flow was set to 12 ml min⁻¹. Methane concentrations were
458 measured against methane standards. The detection limit was 0.1 ppm with a precision of <5%. Rates
459 were determined from the linear increase of methane concentration over time. Due to differences in
460 the linear increase between the three treatments, rates were determined at two different time points: the
461 first period of incubation includes the starting point (day 0) until day 5, the second period includes day
462 8 to day 23 (Supplement information, Fig. S1).

463 Student's t-test (independent, two-tailed, $\alpha = 0.05$) was applied to detect significant differences
464 between the three different treatments.

465

466 **2.8 Gross hydrogenotrophic methanogenesis activity in gravity cores**

467 For the determination of surface to deep methanogenesis activity in gravity cores the radiotracer
468 technique using ¹⁴C-bicarbonate was applied (Jørgensen, 1978). With this method only
469 hydrogenotrophic methanogenesis from CO₂/H₂ can be determined, which is the expected main
470 pathway in deeper sediment layers.

471 Sampled sediment depths were according to the [sampling above](#) scheme [described under section 2.3.](#)
472 [\(see 2.3\)](#). Circa 5 cm³ of sediment was sampled in triplicates into glass [tubes equipped with syringe](#)
473 [plungers syringes](#) and then sealed headspace-free with butyl rubber stoppers. Then, ¹⁴C-bicarbonate-
474 tracer (dissolved in water, pH = 8-9, injection volume 6 µl, activity 222 kBq, specific activity 1.85-
475 2.22 GBq/mmol) was injected through the stopper. The vials were incubated for 48 hours at 9°C
476 before the reaction was stopped by transferring the sediment into 50 ml glass vials filled with 20 ml
477 NaOH (2.5%), closed with butyl rubber stoppers and shaken thoroughly. Five controls were produced
478 from various sediment depths by injecting the radiotracer directly into the NaOH with sediment.
479 In the home laboratory, ¹⁴C- methane production was determined with the slightly modified method by
480 Treude et al., (2005a) used for the determination of anaerobic oxidation of methane. The method was
481 identical, except [that](#) no unlabeled methane was determined by gas chromatography. Instead, DIC
482 values were used to calculate hydrogenotrophic methane production (= CO₂ reduction):

$$MG\ rate = \frac{{}^{14}\text{CH}_4 * [\text{DIC}]}{({}^{14}\text{CH}_4 + {}^{14}\text{C-DIC}) * t}$$

486 The methanogenesis rate (*MG rate*) is expressed in nmol CH₄ cm⁻³ sediment d⁻¹, ¹⁴CH₄ is the activity
487 of produced ¹⁴CH₄, ¹⁴C-DIC is the activity of residual radioactive dissolved organic carbon (DIC= CO₂
488 + HCO₃⁻ + CO₃²⁻), [DIC] is the concentration of dissolved organic carbon in nmol cm⁻³ sediment, and t
489 is the incubation time in days.

490

491 **2.9 Sulfate reduction in MUC cores**

492 One MUC core per station was used for the determination of sulfate reduction. First, two replicate
493 push cores (length 30 cm, inner diameter 2.6 cm) were subsampled from one MUC core. The actual
494 core length varied from 23-25 cm total length. Then, 6 µl (~150 kBq) of carrier-free ³⁵SO₄²⁻
495 radiotracer (dissolved in water, specific activity 37 TBq mmol⁻¹) was injected into the replicate
496 pushcores in 1 cm intervals according to the whole-core injection method [of Jørgensen, \(1978\)](#). Push

497 cores were incubated for ca. 12 h at 9°C. After incubation, bacterial activity was stopped by slicing the
498 push core into 1-cm intervals and transferring each sediment layer into 50 ml plastic centrifuge tubes
499 filled with 20 ml zinc acetate (20% w/w). Controls were done in triplicates from different depths.
500 Here, the sediment was first fixed with zinc acetate before adding the tracer. Rates for sulfate
501 reduction were determined using the cold chromium distillation procedure according to Kallmeyer et
502 al., (2004).
503 The yielded sulfate reduction rates have to be treated with caution, due to long (up to 3 half-life times
504 of ^{35}S) and unfrozen storage. Storage of sulfate reduction samples without freezing has recently been
505 shown to result in the re-oxidation of ^{35}S -sulfides, which results in an underestimation of sulfate
506 reduction rates (Røy et al., 2014). During this reaction, zinc sulfide (Zn^{35}S) and iron sulfide (Fe^{35}S)
507 are re-oxidized to sulfate by reactive Fe(III), which originates from the reaction of Fe^{2+} with oxygen.
508 Fe^{2+} is released during the gradual conversion of FeS to ZnS, which has the lower solubility product.
509 Still, we do trust the relative distribution of activity along depth profiles and consider a potential
510 underestimation of absolute rates.

511

512 **3. Results**

513 **3.1 Water column oxygen and methane concentration**

514 Dissolved oxygen in the bottom water was below detection limit from St.1 (70 m) to St. 8 (407 m),
515 subsequently increasing with water depth to 53 μM at the deepest site (see Table 1 and Dale et al.,
516 2015). At the shallowest St. 1 (70 m) the water was turbid and smelled of sulfide.

517 Dissolved methane concentrations in the bottom water were high on the shelf (St.1-6, 70-253 m) and
518 10 fold lower at the deeper sites (St. 8-10, 407-1024 m; Table 1). The highest measured methane
519 concentration was detected at St. 6 (253 m, ~80 nM) and lowest concentrations were detected at St. 10
520 (1024 m, ~4 nM).

521 **3.2 Sediment core description**

522 A detailed sediment description for the porewater geochemistry cores has been already published in
523 detail by Dale et al., (2015). In short, sediments revealed a grey color with a black surface layer at

524 St. 1 (70 m), a dark olive green color at St. 4-8 (145-407 m), and a green-brown color at St. 9 and 10
525 (770 -1024 m). Sediment texture was soft and fluffy at St. 1-6 (70-253 m), and was less soft at the
526 deeper sites. St. 8 (407 m) revealed a fluffy surface layer followed by a dense clay layer > 2 cmbsf
527 sediment depth. In addition, phosphorite nodules were found at the sediment surface (0-2 cmbsf) of St.
528 8 (407 m).

529 Mats of the sulfur oxidizing bacteria *Thioploca spp.* (Gallardo, 1977) were visible at the sediment
530 surface at St.1-6 (70-253 m), with the densest mat at St. 1 (70 m) continuously decreasing with
531 increasing water depth. Sheaths of *Thioploca* were visible until 20-30 cmbsf at St. 1, 4 and 6 (70-253
532 m).

533 Foraminifera could be observed at the sediment surface of St. 8 (407 m), St. 9 (770 m) and St. 10
534 (1024 m). St. 8 (407 m) showed a thick layer of foraminifera ooze on the sediment surface (0-3 cmbsf)
535 while St. 9 (770 m) and St. 10 (1024 m) showed only scattered foraminifera at the sediment surface (0-
536 5 cmbsf).

537 Macrofauna (large polychaetes, oligochaetes, ophiuroids) were restricted to the sites below the OMZ
538 at St. 9 (770 m) and St. 10 (1024 m), where deep waters were oxygenated. However, small snails (~ 1
539 cm) were observed at St. 8 (407 m).

540

541 **3.3 Geochemical parameters in MUC cores**

542 Porewater and solid phase geochemistry of sediments retrieved by the MUC cores are shown in Fig. 2.
543 Surface sediment (0-0.5 cmbsf) POC content increased along the continental shelf from 1.6 wt % at
544 the shallow St. 1 (70 m) to a maximum of 15 wt % at St. 8 (253 m). Surface POC content decreased
545 again with increasing water depth showing the lowest POC content at St. 10 (1024 m, 2 wt %). While
546 POC content showed more or less stable profiles throughout the sediment core at St. 1 (70 m, around 3
547 wt %), St. 9 (770 m, around 4 wt %), and St. 10 (1024 m, around 3 wt %), POC content was stable
548 only in the upper ~ 10 cmbsf at St. 4 (150 m, around 10 wt %) and St. 6 (253 m, around 15 wt %),
549 followed by a decrease until the deepest sampled depth (2 wt % and 9 wt %, respectively). At St. 8
550 (407 m), POC content increased with sediment depth below 3 cmbsf (from 4 wt % to 9 wt %), which

551 consisted of dense clay (see above). In the upper 3 cmbsf, POC decreased from ~ 7 wt % to ~ 4 wt %,
552 which was the sediment layer with a more fluffy appearance.

553 The sediment surface C/N ratio was lowest at St. 1 (70 m, 6.2) and increased along the continental
554 shelf showing the highest surface C/N ratio at St. 10 (1024 m, 11). St. 8 (407 m) was exceptional, as it
555 showed slightly lower surface C/N ratio (8) as at St. 6 (253 m, 9). St. 8 (407 m) was also the only site
556 showing an increase of 4 units in the upper 0-5 cmbsf, followed by stable ratios around 12 throughout
557 the rest of the core. St. 1 and 4 (70 and 145 m) showed shallower increases in C/N ratio in the upper ~
558 2 cmbsf and upper 1 cmbsf, respectively, followed by stable ratios around 10 until the bottom of the
559 core. At St. 9 and 10 (770 and 1024 m), C/N ratios ranged around 11 and 12, respectively.

560 The highest increase in methane concentration was observed at St. 1 (70 m). Here, methane increased
561 linearly from the surface (1 μM) to the bottom of the core (100 μM). All other stations showed either
562 no clear trend (St. 4= 145 m) or only slight methane increases with depth. At St. 9 (770 m), even a
563 decrease in methane concentration was observed from the surface to the bottom of core.

564 Besides St. 1 (70 m), which showed a strong decrease in sulfate (SO_4^{2-}) concentration with depth from
565 about 28 mM at the top to about 9 mM at the bottom of the core (43 cmbsf), all other stations showed
566 SO_4^{2-} concentrations > 25 mM throughout the cores. At St. 4, 6 and 9 (145, 253, 770 m), SO_4^{2-} showed
567 very slight decrease with depth from about 28 mM at the top to about 25 mM at the bottom of the core.
568 Porewater SO_4^{2-} concentrations were stable around 28 mM throughout the core at St. 8 and 10 (407
569 and 1024 m).

570 Dissolved inorganic carbon (DIC) concentration increased with depth at St. 1- 6 (70 -253 m). St. 1 (70
571 m) showed the steepest increase with depth, showing the lowest DIC concentration at the top (2.3
572 mM) and the highest at the deepest sampled depth (21.6 mM). At St. 4 (153 m), maximum
573 concentration was reached at ~ 23 cmbsf with 4 mM. St. 6 (253 m) showed maximum concentration at
574 the deepest sampled depth with 9 mM. St. 8 and 9 (407 and 770 m) showed stable DIC concentrations
575 around 2.3 mM throughout the core. No DIC data was available for St. 10 (1024 m).

576

577 **3.4 Net methanogenesis and gross sulfate reduction in MUC cores**

578 Maximum net methanogenesis rates (Fig. 2) were detected at St. 1 (70 m, $1.1 \pm 0.5 \text{ nmol cm}^{-3} \text{ d}^{-1}$, 20-
579 25 cmbsf) and St. 6 (253 m, $1.3 \pm 0.65 \text{ nmol cm}^{-3} \text{ d}^{-1}$, 25-30 cmbsf). At all other stations,
580 methanogenesis was mostly below $0.5 \text{ nmol cm}^{-3} \text{ d}^{-1}$ throughout the cores. St. 8 (407 m) showed
581 methanogenesis activity only in the top 10 cmbsf with the maximum at 5-10 cmbsf ($0.2 \pm 0.5 \text{ nmol cm}^{-3} \text{ d}^{-1}$).
582 At St. 9 and 10 (770 and 1024 m), maximum methanogenesis activity was found in the surface
583 layer (0-5 cmbsf) with $0.3 \pm 0.4 \text{ nmol cm}^{-3} \text{ d}^{-1}$ and $0.4 \pm 0.6 \text{ nmol cm}^{-3} \text{ d}^{-1}$, respectively. St. 10 (1024 m)
584 also showed high average methanogenesis at 10-15 cmbsf ($1.5 \pm 2.5 \text{ nmol cm}^{-3} \text{ d}^{-1}$), which was caused
585 by a single high replicate ($4.3 \text{ nmol cm}^{-3} \text{ d}^{-1}$). In the following, e.g., integration of rates, we will
586 exclude this single high replicate, which will be further elaborated in the discussion.

587 At all stations beside St. 9 (770 m), sulfate reduction activity was highest in the 0-1 cmbsf horizon,
588 followed by a sharp decrease in activity of 20-90% in the subsequent 1-2 cmbsf horizon. Highest
589 measured rates at 0-1 cmbsf were observed at St. 4 (145 m, $290 \text{ nmol cm}^{-3} \text{ d}^{-1}$), followed by St. 1 (70
590 m, $270 \text{ nmol cm}^{-3} \text{ d}^{-1}$). Surface (0-1 cmbsf) sulfate reduction activity decreased from St. 4 (145 m) to
591 St. 8 (407 m) with concomitant increase in water depth. St. 9 (770 m) was the only site without a
592 surface sulfate reduction maximum. Here, highest rates were found at 7 cmbsf ($11.2 \text{ nmol cm}^{-3} \text{ d}^{-1}$).
593 St. 6, 8 and 9 (253, 407, and 770 m) showed a second but smaller maximum of sulfate reduction
594 activity. At St. 6 (253 m), this second maximum was situated at 20.5 cmbsf ($6.2 \text{ nmol cm}^{-3} \text{ d}^{-1}$). St. 8
595 and 9 (407 and 770 m) showed additional maxima at 4.5 cmbsf ($3.1 \text{ nmol cm}^{-3} \text{ d}^{-1}$) and 2.5 cmbsf (1.5
596 $\text{nmol cm}^{-3} \text{ d}^{-1}$), respectively. At St. 9 (770 m), sulfate reduction activity was not detectable at most
597 depth > 10 cmbsf. At St. 10 (1024 m), no sulfate reduction activity was detectable throughout the entire
598 core. At St. 9 and 10 (770 and 1024 m) we cannot exclude that sulfate reduction was present but
599 undetectable due to long, unfrozen storage of the samples (see 2.7).

600 Figure- 32 shows an overview of integrated methanogenesis and sulfate reduction rates (over the upper
601 0-25 cm) along the depth transect on the Peruvian margin. Highest integrated surface methanogenesis
602 activity was detected on the shelf (70, 145 and 253 m) with $0.1 \pm 0.03 \text{ mmol m}^{-2} \text{ d}^{-1}$, $0.06 \pm 0.02 \text{ mmol}$
603 $\text{m}^{-2} \text{ d}^{-1}$, and $0.07 \pm 0.01 \text{ mmol m}^{-2} \text{ d}^{-1}$, respectively. St. 8 (407 m) revealed the lowest integrated
604 methanogenesis rate of all sites ($0.02 \pm 0.00 \text{ mmol m}^{-2} \text{ d}^{-1}$). St. 9 (770 m) and St. 10 (1024 m) showed
605 integrated methanogenesis activity around $0.03 \pm 0.02 \text{ mmol m}^{-2} \text{ d}^{-1}$, respectively.

606 Integrated sulfate reduction activity decreased along the continental margin with increasing water
607 depth, revealing the highest activity at the St. 1 (70 m, 4.7 mmol m⁻² d⁻¹) and the lowest activity at St.
608 9 (770 m, 0.2 mmol m⁻² d⁻¹). Please note again, that integrated sulfate reduction rates are probably
609 underestimated due to long, unfrozen storage of the samples (see 2.7).

610

611 **3.5 Potential competitive and non-competitive methanogenesis in sediment slurries from MUC** 612 **cores**

613 Results from the sediment slurry experiments, in which we added either the sulfate reduction inhibitor
614 molybdate, the non-competitive substrate methanol, or no additives (control), are shown in Fig. 43.

615 During the first phase of incubation, all three treatments showed rates within the same order of
616 magnitude. Nevertheless, potential methanogenesis rates were significantly higher ($p < 0.05$) in all
617 treatments in the shallow sediment horizon (0-5 cmbsf) compared to the deep horizon (20-25 cmbsf).
618 In addition, potential methanogenesis was always significantly higher in the molybdate and methanol
619 treatment compared to the control.

620 During the second phase of the incubation (day 8-23), potential methanogenesis showed a different
621 pattern. Rates in the methanol treatment were 350 and 4 times higher compared to the control and
622 molybdate treatment in the 0-5 cm horizon and the 20-25 cm horizon, respectively ($p < 0.05$). Control
623 and molybdate treatments showed no significant difference ($p > 0.05$) in the shallow and deep horizon.

624

625 **3.6 Geochemical parameters and gross hydrogenotrophic methanogenesis activity in gravity** 626 **cores**

627 At the shallow St. 1 (78 m), POC concentration slightly decreased with depth, from ~4 wt % at the
628 surface to about 2-3 wt % at the bottom of the core (385 cmbsf, Fig. 5). At St. 8 (407 m), POC
629 concentrations were slightly higher with values ranging around 8-9 wt % in the upper 120 cmbsf, and
630 then decreasing with depth. The C/N ratio at St. 1 (78 m) remained around 10 throughout the core,
631 while it showed slightly higher values around 12 throughout the core at St. 8 (407 m).

632 At St. 1 (78 m), the methane concentration increased with depth from 0.1 mM at the surface to the
633 highest measured concentration at 165 cmbsf (~5 mM), followed by a decrease to ~ 2 mM at 198
634 cmbsf. Methane concentration stayed around 2 mM until the deepest measured depth (385 cmbsf).
635 Methane concentrations at St. 8 (407 m) ranged from 14 to 17 μM in the upper 120 cmbsf, then
636 increased to a maximum of 36 μM at 180 cmbsf, followed by a decrease to 28 μM at the deepest
637 sampled depth (195 cmbsf).
638 SO_4^{2-} concentration at St. 1 (78 m) decreased with depth with the highest concentration (10 mM) at the
639 shallowest measured sediment depth (33 cmbsf) and the lowest concentration at 350 cmbsf (0.16 mM).
640 At St. 8 (407 m), SO_4^{2-} concentration decreased slightly from ~28 mM at the shallowest measured
641 sediment depth (20 cmbsf) to ~24 mM at 145 cmbsf, followed by stable concentrations around 25 mM
642 until the bottom of the core.
643 DIC concentrations were 5-8 times higher at St. 1 (78 m) compared to St. 8 (407 m) and increased
644 with sediment depth from ~21 mM at 33 cmbsf to ~39 mM at 385 cmbsf. DIC concentrations at St. 8
645 (407 m) could only be measured at distinct sediment depths due to limited amounts of porewater but
646 still revealed a slight increase with sediment depth (from ~3 mM to ~5 mM).
647 Hydrogenotrophic methanogenesis at St. 1 (78 m) was present but low below 66 cmbsf until it reached
648 a peak between 300 and 400 cmbsf ($0.7 \text{ nmol cm}^{-3} \text{ d}^{-1}$). In contrast, no hydrogenotrophic
649 methanogenesis activity was detected at St. 8 (407 m).

650

651 **4. Discussion**

652 **4.1 Concurrent activity of methanogenesis and sulfate reduction in surface sediments**

653 Before we discuss the distribution of methanogenesis in the collected sediment cores, it has to be
654 pointed out that the top soft sediment layer (ca. 0-20 cm) of gravity cores is often disturbed or even
655 lost during the coring procedure. Hence, surface parameters in the gravity cores should not be directly
656 compared to the respective depth layers in MUC cores. According to this likely offset, we will use the
657 term "surface methanogenesis/sediments" when referring to MUC cores and "deep
658 methanogenesis/sediments" when referring to gravity cores.

659 We would further like the reader to keep in mind that we will compare two different types of rate
660 determinations: radiotracer incubations of undisturbed sediments (deep hydrogenotrophic
661 methanogenesis, surface sulfate reduction) and sediment slurry incubations (surface total
662 methanogenesis). While the first method preserves the natural heterogeneity of the sediment, the latter
663 homogenizes and dilutes sediment ingredients and organisms, which could have both negative and
664 positive effects on the natural activity. As we are mainly interested in the vertical distribution of these
665 processes within the sediment, these comparisons are justifiable.

666 In the present study, methanogenesis and sulfate reduction ~~co-~~occurred in surface sediments along
667 the entire depth transect (70-1024 m) on the Peruvian margin (12°S). Methanogenesis activity was
668 detected in sediment layers that revealed high porewater sulfate concentrations and sulfate reduction
669 activity (besides St. 10, where sulfate reduction was undetectable). Even though absolute sulfate
670 reduction rates were most likely underestimated, we trust relative distribution pattern in the sediment
671 and along the continental margin.

672 As the competition between methanogens and sulfate reducers for H₂ and acetate was probably never
673 relieved, the detected surface methanogenesis was most likely based on non-competitive substrates
674 such as methanol or methylated compounds including methylated amines or methylated sulfides
675 (Oremland & Polcin, 1982; Oremland & Taylor, 1978; Kiene et al., 1986). Likewise, in a study off
676 Chile (0-6 cm sediment depth, 87 m water depth), surface methanogenesis was found to be coupled to
677 the non-competitive substrate trimethylamine, and not to CO₂/H₂ or acetate, in sediments where sulfate
678 and sulfate reduction was abundant (Ferdelman et al., 1997).

679 Non-competitive substrate utilization by methanogens in the present study was further confirmed by a
680 significant increase of potential methanogenesis after the addition of methanol to sediment slurries
681 from St. 1 (70 m) (Fig. 4 B). The delayed response of methanogenesis after methanol addition
682 (Supplement, Fig. S1), however, suggests that the present microbial methanogenic community was not
683 primarily feeding on methanol. Potentially, other non-competitive substrates like dimethyl sulfides
684 were utilized predominantly. While most methylotrophic methanogens are able to use both methanol
685 and methylated amines, growth on dimethyl sulfide appears to be restricted to only a few
686 methylotrophic species (Oremland et al., 1989). Dimethyl sulfides could have ~~accumulated~~~~build-up~~

687 during the long storage time (~ 6 months) before experimentation. Even though methylated sulfur
688 compounds (e.g., dimethyl sulfide or methanliol) are mainly produced by organisms in the marine
689 photic zone (e.g., Andreae & Raemdonck 1983), it was recently postulated that these compounds may
690 also be generated through nucleophilic attack by sulfide on methyl groups in the sedimentary organic
691 matter (Mitterer, 2010). As sulfate reduction was a predominant process in the sediment, it could have
692 delivered sufficient sulfide to produce methylated sulfur compounds. Consequently, results from the
693 sediment slurry experiments might not reflect the activity of the in situ methanogenic community as
694 we cannot exclude community shifts as a response to the availability of alternative substrates that were
695 produced during the long storage.

696 The utilization of the competitive substrates H₂ and acetate by the methanogens ~~occurs~~ probably only
697 occurs when sulfate reducers are inhibited. Accordingly, potential methanogenesis rates in the
698 molybdate treatment of the sediment slurry experiment were significantly higher in the two studied
699 horizons (0-5 and 20-25 cmbsf) compared to the controls during the first phase of the incubation (day
700 0-5), indicating the usage of competitive substrate facilitated by the inhibition of sulfate reduction.
701 However, in the second phase (day 8-23) of the incubation, rates were much lower in both the control
702 and molybdate treatment and did not show significant differences in both horizons (p>0.05). In this
703 second phase, methane production might have slowed down due to depletion of electron donors.
704 Hydrogenotrophic methanogenesis in the gravity core from St. 1 (78 m) showed no activity at depths
705 where porewater sulfate concentrations were >0.7 mM. Instead activity peaked where porewater
706 sulfate was lowest (0.16 mM at 350 cmbsf), supporting the above conclusions regarding competition
707 within the sulfate zone. The observation that sulfate was never completely depleted in the porewater
708 until the bottom of the gravity core, in combination with an increase of iron (II) in the porewater at
709 depths > 200 cmbsf (data not shown), hint to the presence of a cryptic sulfur cycle that is responsible
710 for deep formation of sulfate (Holmkvist et al., 2011; Treude et al., 2014) .

711 In comparison, surface net methanogenesis activity along the Peruvian margin was similar to
712 activities found off Chile at 87 m water depth (0-0.6 nmol cm⁻³ d⁻¹) (Ferdelman et al., 1997). The
713 slightly higher rates determined in our study (St.1= 70 m; 0.4-1.7 nmol cm⁻³ d⁻¹) could be related to
714 different approaches, as our rates represent the sum of net methanogenesis from all available

715 substrates in the sediment, while rates off Chile were based only on CO₂, acetate, and trimethylamine
716 utilization. Hence, total methanogenesis could have been easily underestimated, if methanogenesis
717 was supplied by other substrates, which is not unlikely, as methylotrophic methanogens, which are
718 able to use methanol or methylated amines, were the dominant type of methanogens in these sediments
719 (Ferdelman et al., 1997). Interestingly, the authors detected a high number of acetogens,
720 ~~implying~~ ~~ieating~~ that acetogenesis competed for methylamines or other methylated compounds
721 (Ferdelman et al., 1997). A competition with acetogens for methylated substrates is conceivable for
722 our study, but would require further corroboration studies.
723

724 4.2 Surface vs deep methanogenesis

725 Maximum single net surface methanogenesis activities detected in our study (0.3-4.3 nmol cm⁻³ d⁻¹)
726 were found to be at the very low end ~~of~~ or even one order of magnitude lower than organic-rich,
727 sulfate-depleted sediments (9.8-37 nmol cm⁻³ d⁻¹, 0-40 cmbsf, Treude et al., 2005a, 10-17 nmol cm⁻³
728 d⁻¹, 0-30 cmbsf, Schmaljohann 1996, 100-300 nmol cm⁻³ d⁻¹, 0-30 cmbsf, Crill & Martens, 1983, 1986,
729 100-400 nmol cm⁻³ d⁻¹, 0-3 cmbsf, Alperin et al. 1992). To estimate the overall relevance of surface
730 methanogenesis within the sulfate zone compared to deep methane production, we estimated the deep
731 methane production in our study and compiled an overview of published deep methane production
732 data from the sulfate-free zone of organic-rich sediments (Table 2). For this comparison, the deep
733 methane production was assumed to equal the flux of methane into the sulfate-methane-transition zone
734 (SMTZ), where it is consumed by anaerobic oxidation of methane (AOM). Within the SMTZ, both
735 sulfate and methane are depleted steeply as a result of AOM, thus dividing the sulfate-reducing zone
736 above from the methanogenic zone below. The SMTZ is the main niche for AOM in marine
737 sediments, acting as an important filter for upwards migrating methane (Knittel & Boetius, 2009). The
738 SMTZ can be found at decimeters to tens of meters below the seafloor, depending on the burial rate of
739 reactive organic matter, the depth of the methane production zone, and the transport velocity flux of
740 methane and sulfate as well as their consumption rates (Knittel & Boetius, 2009).
741 In the present study, a SMTZ was only detected in the gravity core taken at St. 1 (78 m; Fig. 5), where
742 it was located between 66 and 99 cmbsf, i.e., below the penetration depth of the MUC cores. We

743 estimated a methane flux (= deep methane production) into the SMTZ (from 99 to 66 cmbsf)
744 according to Iversen & Jørgensen, (1993) using a seawater methane-diffusion coefficient from Schulz,
745 (2006) which was corrected for porosity resulting in a sediment-diffusion coefficient for methane of
746 $D_s = 1.325 \times 10^{-5} \text{ cm}^2 \text{ s}^{-1}$ at 15 °C. The resulting deep methane production ($0.8 \text{ mmol m}^{-2} \text{ d}^{-1}$) was
747 slightly higher (ratio of 0.13, surface vs. deep) but still in the same magnitude as the integrated surface
748 methanogenesis at St. 1 (70 m; $0.1 \text{ mmol m}^{-2} \text{ d}^{-1}$). Compared to a different study from the Peruvian
749 OMZ, the ratio between shallow (0.07 to $0.1 \text{ mmol m}^{-2} \text{ d}^{-1}$, this study) vs. deep (8.9×10^{-8} to 2.2×10^{-7}
750 $\text{mmol m}^{-2} \text{ d}^{-1}$; Arning et al., 2012) methanogenesis on the shelf (150-250 m) was 3.2×10^5 to
751 1.1×10^6 . Both examples highlight the significance of surface methanogenesis, especially on the
752 Peruvian shelf. On the lower Peruvian slope (~3800 m water depth), deep methanogenesis increased
753 (up to $0.017 \text{ mmol m}^{-2} \text{ d}^{-1}$; Arning et al., 2012). In contrast, surface methanogenesis at the deeper St.
754 10 (1024 m) was lower ($0.02 \text{ mmol m}^{-2} \text{ d}^{-1}$) compared to the shelf indicating a decreasing relevance of
755 surface methanogenesis along the margin with increasing relevance of deep methanogenesis. The
756 decrease of surface methanogenesis with increasing water depth might be correlated to the decreasing
757 organic carbon content and freshness in the sediment (Fig. 6), ~~as which will be~~ further discussed in
758 section 4.4.

759 In comparison with other organic-rich sediments (Table 2), surface methanogenesis off Peru was in
760 the same order of magnitude as most reported deep methanogenesis (e.g., off Namibia, off Chile,
761 Limfjorden). The only exception was Eckernförde Bay (Baltic Sea), where surface methanogenesis
762 off Peru was less than 15% of deep methanogenesis. Eckernförde Bay has a water depth of only ~30 m
763 with high carbon export, featuring extremely high methanogenesis activity below the SMTZ, causing
764 supersaturation and methane gas ebullition (Whiticar, 2002; Treude et al., 2005a).

765 **4.3 Potential consumption and emission of surface methane**

766 Due to its closeness to the sediment-water interface, surface methanogenesis along the Peruvian
767 margin could lead to methane emissions from the sediment into the water column. A short diffusion
768 distance, especially in the top most sediment layers, might facilitate a partial escape of methane from
769 consumption by microbes. As surface methanogenesis decreased with water depth (Fig. 3), the

770 methane emission potential appears to be highest on the shelf. Sediment methane concentrations in the
771 0-2 sediment horizon of all sites along the margin were always higher than bottom-near water methane
772 concentrations (~1.5 m above seafloor; Table 1, Fig. 2), hinting towards an efflux of methane from the
773 sediment. However, more precise profiling of methane at the sediment-water interface would be
774 necessary to confirm this hypothesis. Still, most of the sediment methane profiles suggest methane
775 consumption close to the seafloor to some extent, which would reduce the amount of emitted methane
776 (Fig. 2). AOM might act as an important methane filter at the sediment surface of the shelf stations,
777 where anoxic conditions dominated, while aerobic oxidation might prevail at the deeper stations below
778 the OMZ (St. 9 and 10). The presence of methane oxidation above the SMTZ of organic-rich
779 sediments has been reported earlier (Treude et al., 2005a, 2005b), and could indeed be fueled by
780 surface methanogenesis. An immediate oxidation of the produced methane would explain why
781 sediment methane profiles did not necessarily correlate with peaks in surface methanogenesis (see,
782 e.g., St 6, 253 m). The importance of AOM for the reduction of methane emissions from surface
783 methanogenesis remains speculative, as explicit data is missing. On the basis of our findings, however,
784 we suggest to consider surface methanogenesis as a possible driver for AOM above the SMTZ in
785 earlier and future studies.

786

787 **4.4 Factors controlling methanogenesis along the Peruvian margin**

788 For this discussion, we excluded the high integrated methane production observed in one of the
789 replicates at station 10 (1024 m); as we do not think that the detected activity ($0.23 \text{ mmol m}^{-2} \text{ d}^{-1}$) is
790 representative for this deep site, especially as sediment POC content was lowest at station 10
791 compared to the other stations (<4%, Fig. 2). The outlier might have been caused by additional carbon
792 sources in the sediment, e.g., from fecal pellets or organic carbon released from dead infauna, thus
793 stimulating below-surface microbial activities during our incubations (Ziervogel et al., 2014; Bertics et
794 al., 2013).

795

796 **4.4.1 Oxygen**

797 Oxygen is an important controlling factor, as methanogenesis is an oxygen- and redox-sensitive
798 process (Oremland, 1988). Some methanogens can tolerate oxygen exposure for several hours before
799 they die, ~~although however,~~ no methane ~~is will be~~ produced in the presence of oxygen (Zinder, 1993).
800 Comparing integrated surface methanogenesis (over 0-25 cmbsf) from the shallowest to the deepest
801 station (Fig. 3), highest rates ($> 0.05 \text{ mmol m}^{-2} \text{ d}^{-1}$) were detected on the shelf (St. 1, 4 and 6=70, 145,
802 253 m), where oxygen concentrations were below detection (Fig.6), providing advantageous
803 conditions for methanogenesis, particularly at the very sediment surface, where normally aerobic
804 respiration dominates (Jørgensen, 2006). Below the OMZ, integrated methanogenesis decreased.
805 Bioturbating macrofauna and megafauna (e.g., mussels, polychaetes, oligochaetes) were observed at
806 these sites (St. 9 and 10, 770 and 1024 m) (Mosch et al. 2012), which could have transported oxygen
807 into deeper sediment layer (Orsi et al., 1996), thus leading to less reduced conditions ($> -200 \text{ mV}$)
808 unsuitable for methanogens (Oremland, 1988). However, integrated methanogenesis was lowest at St.
809 8 (407 m), which still revealed anoxic bottom water. Thus, oxygen might just be advantageous but not
810 the driving factor for surface methanogenesis.

811

812 4.4.2 Organic matter

813 The ~~probably~~ most important factor controlling benthic methanogenesis activity is ~~probably~~ the POC
814 content of the sediment, as it determines the substrate availability and variety, and can thus relieve the
815 competitive situation between methanogens and sulfate reducers (Holmer & Kristensen, 1994; Treude
816 et al., 2009). ~~Consequently, Hence, we would expect~~ high methanogenesis ~~rates may be expected~~
817 ~~along the Peruvian margin~~ at sites with high organic carbon load, ~~along the Peruvian margin. However~~
818 ~~Conversely,~~ integrated methanogenesis rates ~~didare~~ not correlat~~ing~~ with sediment POC content (Fig.
819 6). While POC content was increasing from St. 1 (70 m) to St. 6 (253 m), followed by a decrease until
820 St. 10 (1024 m), integrated methanogenesis showed rather a decreasing trend with increasing water
821 depth. This deviation ~~from expectations~~ might be caused by another factor, ~~such as not only the~~
822 ~~quantity of~~ organic matter ~~is important for microbial degradation but also its~~ quality, i.e., freshness.
823 Numerous studies have shown that the quality of the organic matter is important for the rate and

824 magnitude of microbial organic matter degradation (Westrich & Berner, 1984; Canfield, 1994; Amon
825 et al., 2001; Middelburg, 1989).
826 Integrated methanogenesis and C/N ratios (indicating the freshness of organic matter) were negatively
827 correlated along the Peruvian margin (Fig. 6), suggesting that fresh, labile organic matter is
828 advantageous for surface methanogenesis. As methanogens consume mostly short, monomeric
829 substrates, they depend on other microbial groups to break down large organic macromolecules
830 (Zinder, 1993). Hence, labile organic matter offers an important supply of methanogenic substrates.
831 In agreement with this hypothesis, highest integrated methanogenesis rates were observed at St. 1 (70
832 m), which revealed the freshest organic matter (lowest C/N, Fig. 6) and the highest POC
833 remineralization rates along the Peruvian margin (Dale et al., 2015). The degradation of organic matter
834 within the water column was probably limited at St. 1 (70 m) due to anoxic conditions and high
835 sedimentation rates (Dale et al., 2015); hence, labile organic matter accumulated at the seafloor,
836 thereby increasing the benthic POC degradation and resulting in high substrate availability and variety
837 for the methanogenic community.

838 Nevertheless, lowest methanogenesis ~~rates~~ ~~were~~ measured at St. 8 (407 m), which was neither the
839 site of the highest C/N ratio, lowest POC content (Fig. 6), or the lowest POC mineralization (Dale et
840 al., 2015). In this particular case, methanogenesis was most likely controlled by the sediment
841 properties. Methanogenesis activity was undetectable below 10 cmbsf, which coincided with a very
842 dense and sticky clay layer. The POC profile at St. 8 (407 m) revealed lower concentrations in the
843 upper 5 cmbsf, followed by an increase with depth, suggesting that either the organic matter at this
844 station was resistant to microbial attack (indicated by the increase in C/N) or that microbes were not as
845 frequent/active in the dense clay layer as at the surface. Similarly, sulfate reduction and microbial
846 nitrogen fixation (Gier et al., 2015) (~~Gier et al., submitted~~) showed very low activity at this site (Fig.
847 2).

848

849 **5. Conclusion**

850 The present study demonstrated that methanogenesis coincides with sulfate reduction in surface
851 sediments (< 30 cmbsf) along the Peruvian margin. The competition with sulfate reducers was
852 partially relieved due to the high load of organic carbon allowing both groups to show concurrent
853 activity through the utilization of non-competitive substrates by the methanogens.

854 The significance of surface methanogenesis was high on the shelf, where ratios between surface and
855 deep methanogenesis ~~were~~ around 0.13 (this study) or even as high as $\sim 10^5$ (compared to Arning et
856 al. 2012), and decreased with increasing water depth. Accordingly, we assume that potential methane
857 emissions into the water column, indicated by a higher methane concentration at the sediment surface
858 compared to the bottom water, should be highest on the shelf, where surface methane production rates
859 were highest. Our results further hint towards a partial consumption of methane before reaching the
860 sediment-water interface, probably by anaerobic oxidation of methane (AOM). In this case, surface
861 methanogenesis might act as important supplier of methane for AOM above the SMTZ, which has
862 been largely ~~overlooked previously seen before~~.

863 We postulate that the dominant factor controlling surface methanogenesis is the availability of
864 (primarily labile) organic matter. The high load of organic carbon and resulting high organic carbon
865 mineralization rates secure the supply ~~offer~~ of methanogenic substrates, especially on the shelf, which
866 mitigates the competition between sulfate reducers and methanogens. Anoxic conditions in the
867 overlying water might be advantageous for the oxygen-sensitive process of methanogenesis, but does
868 not appear to primarily control benthic rates, as they change within the anoxic zones.

869 Interestingly, organic matter made available by bioturbating infauna (e.g., fecal pellets or dead
870 organisms) could be an important additional factor facilitating methanogenesis in surface sediments.

871 As shown in this study, methanogenesis rates vary strongly in bioturbated sediments below the OMZ,
872 sometimes exceeding all other observed methanogenic rates.

873 Future studies should seek to (1) identify methanogens and their metabolic capabilities in surface
874 sediments, (2) determine the direct interaction between surface methanogenesis and AOM, ~~and~~ (3)
875 evaluate the effect of organic matter hot spots on total benthic surface methanogenesis in organic-rich
876 sediments.

877

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888

889 **Author contribution**

890 J.M. and T.T. designed the experiments. J.M. carried out all methanogenesis experiments, T.T.
891 conducted sulfate reduction measurements. Porewater measurements of MUC cores were coordinated
892 by A.D. and S.S. J.M. prepared the manuscript with contributions of all co-authors.

893

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1080 **Figure Captions**

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1082 **Figure 1:** Location of sampling sites off Peru along the depth transect at 12° S. Source: Schlitzer, R.,
1083 Ocean Data View, <http://odv.awi.de>, 2014

1084 **Figure 2:** Profiles of particulate organic carbon (POC), C/N ratio, methane (CH₄), sulfate (SO₄²⁻), DIC
1085 (dissolved inorganic carbon), net methanogenesis (MG) rates and sulfate reduction (SR) rates in the
1086 MUC cores along the depth transect. For MG, triplicates (symbols) and mean (solid line) are shown.
1087 For SR, duplicates are shown. Data points from the overlaying water in the MUC core (OLW) are set
1088 to 0 cm. Note deviant scale dimension for MG at St. 6 and for SR at St. 1 and 2.

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1090 **Figure 3:** Integrated methanogenesis and sulfate reduction rates (0-25 cm) along the depth transect.
1091 For methanogenesis rates (black bars), average values are shown with standard deviation. Note for St.
1092 10 a mean from two replicates is shown without standard deviation (pattern-filled bar) and the outlier
1093 is shown separately (cross). For sulfate reduction rates (blue bars), means from two replicates are
1094 shown without standard deviation.

1095 **Figure 4:** Potential methanogenesis rates in sediment slurry experiments from the two sediment
1096 intervals (0-5 cm and 20-25 cm) at St. 1 (70 m). The first phase of the incubation shows rates
1097 calculated from day 0 to 5 (A), while the second phase of the incubation summarizes the rates from
1098 day 8-23 (B). "Control" is the treatment with sulfate-rich (28 mM) artificial seawater medium, "plus
1099 Mb" is the treatment with sulfate-rich artificial seawater medium plus methylcobalamin (Mb, 22mM), and
1100 "plus Meth" is defined as the treatment with sulfate-rich artificial seawater medium plus methanol
1101 (Meth, 10 mM). Per treatment, average values are shown with standard deviation. Please note the
1102 split-up in the diagram in part B and the different x-axis for methanogenesis

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1104 **Figure 5:** Profiles of particulate organic carbon (POC), C/N ratio, methane (CH₄), sulfate (SO₄²⁻),
1105 dissolved inorganic carbon (DIC), and hydrogenotrophic methanogenesis (MG) rates in the gravity

1106 cores at two stations within the depth transect. For MG, triplicates (symbols) and mean (solid line) are
1107 shown.

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1109 **Figure 6:** Bottom-near water methane (CH_4) and oxygen (O_2) concentrations along the depth transect
1110 (above). Surface sediment particulate organic carbon (POC) content and C/N ratio together with
1111 integrated methanogenesis (MG) rates (0-25 cmbsf) along the depth transect (below). For MG rates,
1112 averages are shown with standard deviation beside St. 10, where a mean from two replicates is shown
1113 (see text). Please note the secondary y-axis.

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1117 **Tables**

1118 Table 1: Stations, instruments, chemical/physical parameters in the bottom-near water, and analyses
 1119 applied to samples along the depth transect on the Peruvian margin (12°S). For abbreviations see
 1120 footnote.

Station No	Instrumen t	Latitude (S)	Longitude (W)	Water depth (m)	O ₂ (μM)	Temp. (°C)	CH ₄ (nM)	Type of analysis
1	MUC 13	12°13.492	77°10.511	70				All
	MUC 38	12°13.517	77°10.084	70				SE
	GC 8	12°14.500	77°9.611	78				GC-All
	CTD 9	12°13.535	77°10.522	73	bdl	14	38.6	WC
4	MUC 10	12°18.704	77°17.790	145				All
	CTD 14	12°18.697	77°18.004	145	bdl	13.4	24.4	WC
6	MUC 5	12°23.321	77°24.176	253				Gas+PW
	MUC 6	12°23.322	77°24.181	253				nMG
	CTD 6	12°24.904	77°26.314	305	bdl	12	79.6	WC
8	MUC 23	12°27.198	77°29.497	407				Gas+ PW
	MUC 24	12°27.197	77°29.497	407				nMG
	GC 3	12°27.192	77°29.491	407				GC-All
	CTD 37	12°29.502	77°29.502	407	bdl	10.6	7.3	WC
9	MUC 17	12°31.374	77°35.183	770				Gas+ PW
	MUC 18	12°31.373	77°35.184	770				nMG
	CTD 27	12°31.327	77°35.265	770	19	5.5	8.4	WC
10	MUC 28	12°35.377	77°40.975	1024				Gas+ PW
	MUC 29	12°35.377	77°40.976	1024				nMG
	CTD 11	12°34.863	77°38.954	1010	53	4.4	3.9	WC

1121 MUC = multicorer, GC= gravity corer, CTD = CTD/Rosette, bdl= below detection limit (5μM), All = methane
 1122 gas analysis, porewater analysis, net methanogenesis analysis, SE = slurry experiment, GC-All= analysis for
 1123 gravity cores including methane gas analysis, porewater analysis, hydrogenotrophic methanogenesis analysis,
 1124 WC= Water column analyses, Gas = methane gas analysis, PW= porewater analysis, nMG= net methanogenesis
 1125 analysis,
 1126

Table 2: Comparison of deep methanogenesis in organic-rich sediments from different regions with surface methanogenesis ($0.02\text{-}0.1 \text{ mmol m}^{-2} \text{ d}^{-1}$) determined in the present study. The ratio range was achieved by dividing the lowest surface by the highest deep and the highest surface by the lowest deep methanogenic activity, respectively.

	Water Depth (m)	Depth of SMTZ (mbsf)	Methane flux into the SMTZ = integrated deep methanogenesis ($\text{mmol m}^{-2} \text{ d}^{-1}$)	Ratio between surface methanogenesis (present study) and deep methanogenesis	Reference
Namibia (SE Atlantic)	1312-2060	3-10	0.07-0.15	0.13-1.43	Niewöhner et al., (1998)
Eckernförde Bay (SW Baltic Sea)	25-28	0.5-1.5	0.66-1.88	0.01-0.15	Treude et al., (2005a)
Chile (SE Pacific)	797-2746	3-4	0.068-0.13	0.15-1.47	Treude et al., (2005b)
Limfjorden (North Sea)	7-10	1-1.5	0.076	0.03-1.32	Jørgensen & Parkes, (2010)
Peru (SE Pacific)	150-3819	2-50	2.2×10^{-7} -0.017	$1.18\text{-}4.55 \times 10^5$	Arning et al., (2012)
Peru (SE Pacific)	70-1024	0.7-1	0.8	0.03-0.13	present study

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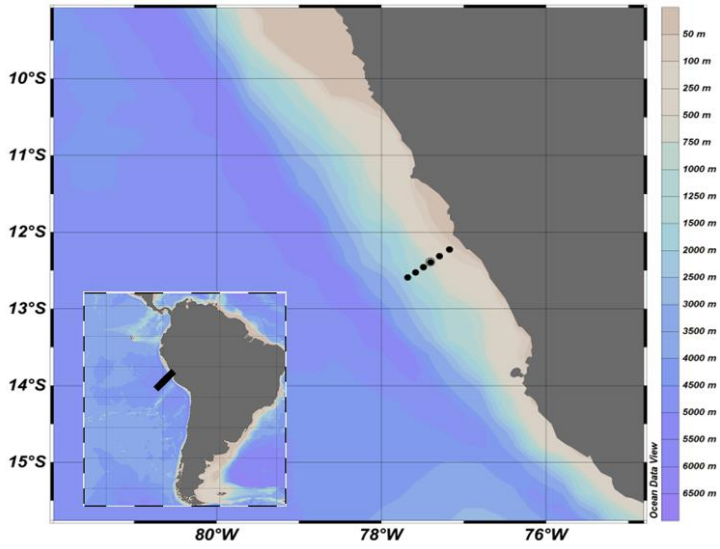
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1132 **Figures**

1133 **Figure 1**



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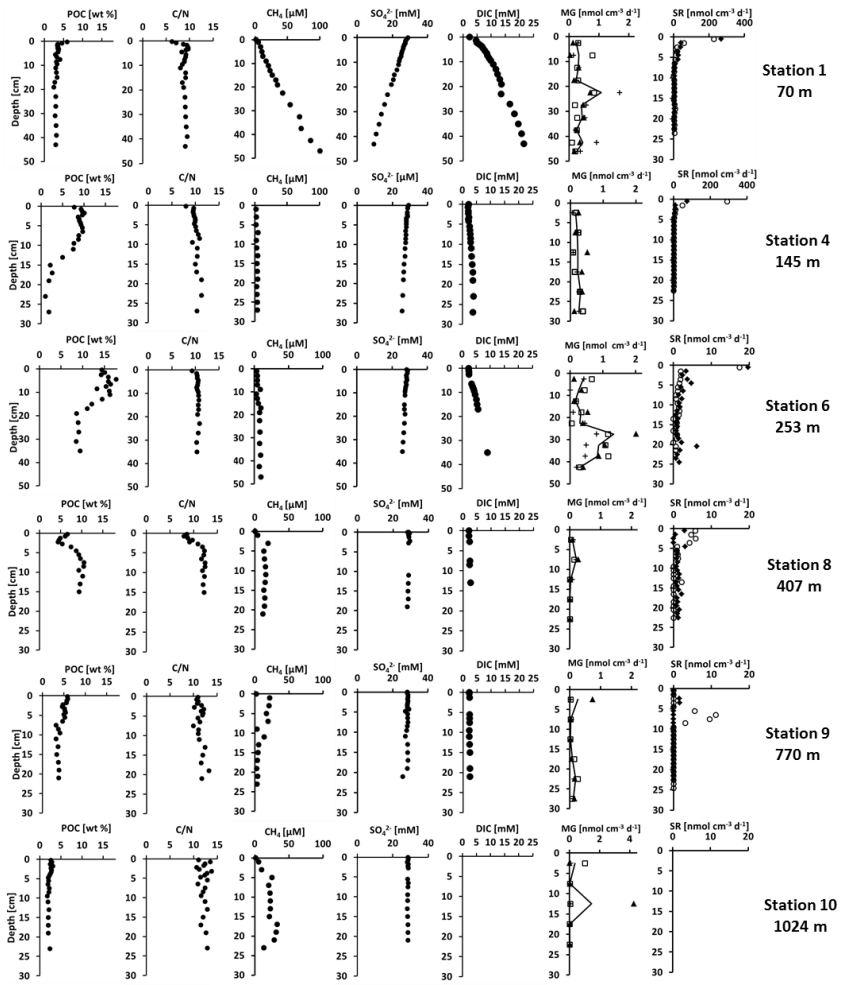
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1148 **Figure 2**



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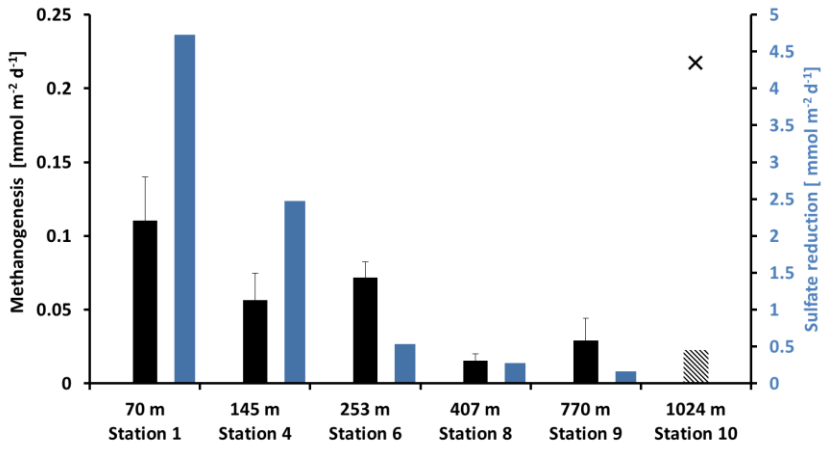
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1156 **Figure 3**



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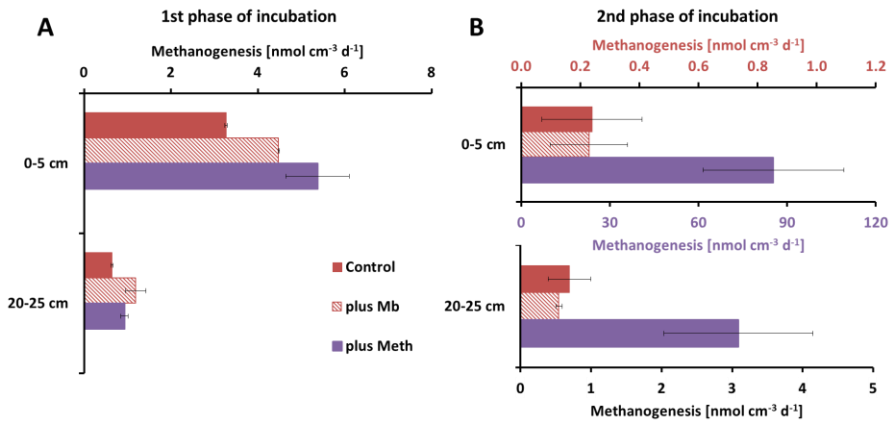
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1173 **Figure 4**



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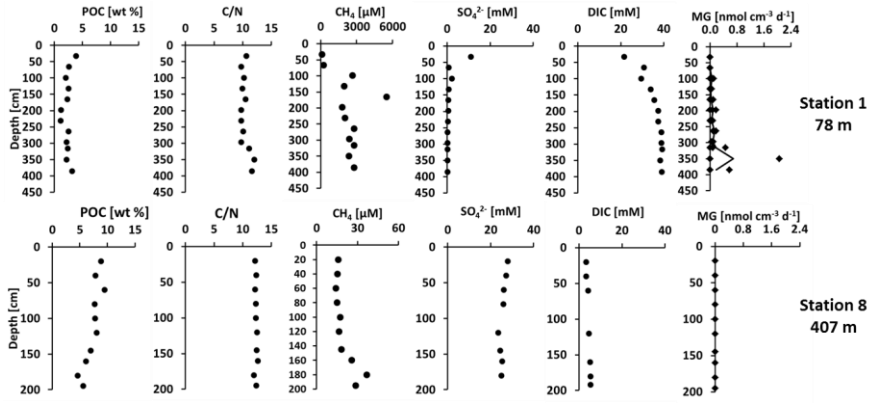
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1191 **Figure 5**



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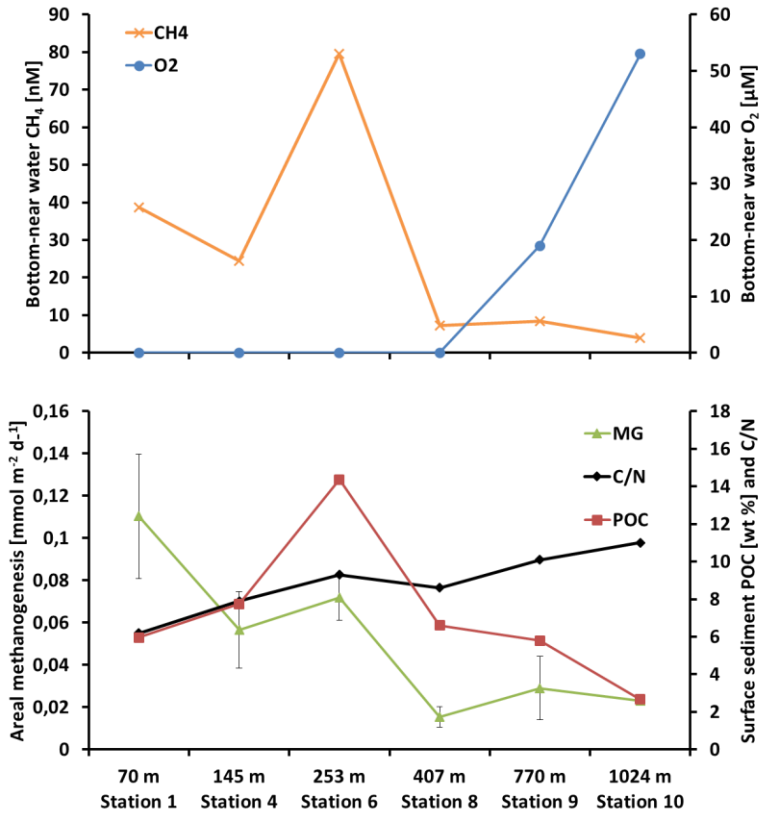
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1207 Figure 6



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