

1 **Long-term incubations provide insight into the mechanisms of anaerobic**
2 **oxidation of methane in methanogenic lake sediments**

3 Hanni Vigderovich^a, Werner Eckert^b, Michal Elul^a, Maxim Rubin-Blum^c, Marcus Elvert^d, Orit Sivan^a

4 ^aDepartment of Earth and Environmental Science, Ben-Gurion University of the Negev, Beer Sheva, Israel

5 ^bIsrael Oceanographic & Limnological Research, The Yigal Allon Kinneret Limnological Laboratory, Israel

6 ^cIsrael Oceanographic & Limnological Research, Haifa, Israel

7 ^dMARUM - Center for Marine Environmental Sciences and Faculty of Geosciences, University of Bremen,
8 Bremen, Germany

9 *Corresponding author:* Hanni Vigderovich, hannil@post.bgu.ac.il

10 **Abstract**

11 Anaerobic oxidation of methane (AOM) is one of the major processes limiting the release of the
12 greenhouse gas methane from natural environments. In Lake Kinneret (Israel), geochemical profiles
13 and experiments with fresh sediments indicate that iron-coupled AOM (Fe-AOM) sequesters 10-15%
14 of the methane produced in the methanogenic zone (>20 cm sediment depth). The oxidation of methane
15 in this environment was shown to be mediated by a combination of *mcr* gene-bearing archaea and
16 aerobic bacterial methanotrophs. Here, we aimed to investigate the AOM process in terms of various
17 electron acceptors and involved microorganisms during long-term anaerobic sediment slurry
18 incubations (~ 18 months) under controlled conditions. We followed the process with the addition of
19 ¹³C-labeled methane and two stages of incubations: (i) enrichment of the microbial population involved
20 in AOM and (ii) slurry dilution and manipulations, including addition of multiple electron acceptors
21 (metal oxides, nitrate, nitrite and humic substances) and inhibitors for methanogenesis/AOM and sulfate
22 reduction. Carbon isotope measurements in the dissolved inorganic pool in these long-term incubations
23 suggest that considerable AOM consumed 3-8% of the methane produced at a rate of 2.0±0.4 nmol gr⁻¹
24 dry sediment day⁻¹. Carbon isotope measurements in lipids and metagenomic analyses indicate that
25 only anaerobic microbes catalyzed this AOM. Whereas cryptic oxidation of methane by combining
26 archaea and aerobic methanotrophs is feasible in the natural Lake Kinneret sediments, reverse
27 methanogenesis dominates methane turnover in the long-term controlled experiments. Humic
28 substances and iron oxides, but not sulfate, manganese, nitrate, and nitrite, are the likely electron
29 acceptors used during the AOM. Our observations support the contrast between methane oxidation
30 mechanisms in naturally anoxic lake sediments, with potentially co-existing aerobes and anaerobes, and
31 long-term incubations, where anaerobes prevail.

32 Keywords: Anaerobic oxidation of methane (AOM), lake, sediments, dissolved inorganic carbon, stable
33 isotopes, electron acceptor, methanotrophs

34 **1. Introduction**

35 Methane (CH₄) is an effective greenhouse gas (Wuebbles and Hayhoe, 2002) with anthropogenic and
36 natural sources. Natural methane sources contribute about 50% of this gas emission to the atmosphere
37 (Saunio et al., 2020). Aerobic and anaerobic oxidation of methane (AOM) naturally control the release
38 of this gas to the atmosphere (Conrad, 2009; Reeburgh, 2007; Knittel and Boetius, 2009). While sulfate-
39 dependent AOM, catalyzed by ANaerobic MEthanotrophs (ANMEs) 1-3, is widespread mostly in
40 marine sediments (Hoehler et al., 1994; Boetius et al., 2000; Orphan et al., 2001; Treude et al., 2005,
41 2014), in other environments methane oxidation can be coupled to other electron acceptors.

42 AOM coupled to the reduction of iron and manganese oxides has been confirmed in several
43 environments (Beal et al., 2009; Egger et al., 2015; Sivan et al., 2011; Sivan et al., 2014; Segarra et al.,
44 2013; Bar-or et al., 2017; Aromokeye et al., 2020; Su et al., 2020; Mostovaya et al., 2021). Alternative
45 electron acceptors for AOM include other metals, humic substances, nitrate and nitrite. The synthetic
46 analog for humic substances, 9,10-anthraquinone-2,6-disulfonate (AQDS), was shown to serve as a
47 terminal electron acceptor (Scheller et al., 2016; Valenzuela et al., 2017; Bai et al., 2019; Zhang et al.,
48 2019; Fan et al., 2020). Nitrate-dependent AOM has been demonstrated in a consortium of archaea and
49 denitrifying bacteria (Raghoebarsing et al., 2006) and in an enrichment culture of ANME-2d (Haroon
50 et al., 2013; Arshad et al., 2015), whereas nitrite fuels AOM by *Methyloirabilis* (NC-10, Ettwig et al.,
51 2010). ANME-2d and *Methyloirabilis* can also couple AOM to selenite reduction (Luo et al., 2018).
52 The ubiquitous aerobic methanotrophs Methylococcales may oxidize methane and denitrify under
53 hypoxia (Kits et al., 2015), switch to iron reduction (Zheng et al., 2020), or generate oxygen by
54 methanobactins (Dershwitz et al., 2021). The latter study also showed the ability of alphaproteobacterial
55 methanotroph *Methylocystis* sp. strain SB2 to couple methane oxidation and iron reduction.

56 In Lake Kinneret sediments, *in-situ* pore water profiles (Sivan et al., 2011), diagenetic models (Adler
57 et al., 2011) and incubation experiments with fresh sediment slurries (Bar-Or et al., 2017) suggest that
58 iron reduction coupled to AOM (Fe-AOM) removes 10-15% of the produced methane in the deep
59 methanogenic zone (>20 cm below water-sediment interface). Analysis of the microbial community
60 structure revealed that both methanogenic archaea and methanotrophic bacteria are potentially involved
61 in methane oxidation (Bar-Or et al., 2015). Analyses of stable isotopes in fatty acids, the 16S rRNA
62 gene amplicons and metagenomics showed that reverse methanogenesis by archaea and the bacterial
63 type I aerobic methanotrophy by Methylococcales play a role in methane cycling (Bar-Or et al., 2017;
64 Elul et al., 2021). This aerobic methanotrophic activity has been observed in several anoxic
65 hypolimnions and sediments of lakes (Beck et al., 2013; Oswald et al., 2016; Martinez-Cruz et al., 2017;
66 Cabrol et al., 2020), and might be fueled by the presence of oxygen at microlevel up to several meters
67 below the oxycline. However, whether these methanotrophs continue to oxidize methane under strictly
68 anoxic conditions and which electron acceptors are available is still unknown.

69 Here, we used long-term anaerobic incubations to assess the dynamics of methane-oxidizing microbes
70 under anoxic conditions and to quantify various electron acceptors' availability for AOM. For this
71 purpose, we diluted fresh methanogenic sediments from Lake Kinneret with original porewater from
72 the same depth and amended the sediment with ^{13}C -labeled methane, following its oxidation to
73 dissolved inorganic carbon (DIC). Our experiment design consisted of two stages, the first stage
74 included the enrichment of the microbial population involved in AOM, and the second stage involved
75 an additional slurry dilution and several manipulations with multiple electron acceptors and inhibitors.
76 The potential electron acceptors were iron and manganese oxides, nitrate, nitrite and humic substances.
77 We inhibited the *mcr* gene with 2-bromoethanesulfonate (BES), methanogens with acetylene and
78 sulfate reduction and sulfur disproportionation with Na-Molybdate (Nollet et al., 1997; Orembland &
79 Capone, 1988; Lovley & Klug, 1983). We measured methane oxidation rates (by the ^{13}C -DIC
80 enrichment), the electron acceptor characteristics (by their addition or inhibition) and the evaluated
81 changes in microbial diversity over various incubation periods (based on metagenomics and lipid
82 biomarkers). The results from the long-term anaerobic incubations were compared to those of batch and
83 semi-bioreactor experiments that were set up with fresh sediments to follow the changes in methane
84 oxidation mechanisms.

85 **2. Methods**

86 **2.1 Study site**

87 Lake Kinneret (Sea of Galilee) is a warm monomictic freshwater lake located in the North of Israel.
88 The lake is 21 km long and 13 km wide. Its maximum depth is ~42 m at the lake center (station A,
89 Figure S1) and the average depth is 24 m. The lake is thermally stratified from March to December,
90 with the hypolimnion turning anoxic from April. Surface water temperatures range from 15 to 30 °C,
91 and the bottom water temperatures remain between 14-17 °C all year long. The lake sediments are
92 composed mostly of carbonates (40-50%) and clays (20%; Hadas and Pinkas, 1995; Eckert, 2000). The
93 total iron content in the top 40 cm of the sediments is ~3 wt % (Serruya, 1971; Eckert, 2000; Bar-Or et
94 al., 2017). The sediment at the deep methanogenic zone used in this study (~20 cm sediment depth from
95 the water-sediment interface at the lake's center) contains 50% carbonates, 30% clay and 7% iron (Table
96 S1). The porewater's dissolved organic carbon (DOC) concentration increases with depth, ranging from
97 ~6 mg C L⁻¹ at the sediment-water interface to 17 mg C L⁻¹ at 25 cm depth (Adler et al., 2011). Dissolved
98 methane concentrations in the porewater increase sharply from the top sediments to more than 2 mM at
99 15 cm depth and then decrease to 0.5 mM (Adler et al., 2011; Sivan et al., 2011; Bar-Or et al., 2015).

100 **2.2 Experimental set-up**

101 This study compares three incubation strategies (A, B and C) of Lake Kinneret methanogenic sediments
102 amended with original porewater from the same depth, ^{13}C -labeled methane, different potential electron

103 acceptors for AOM (nitrite, nitrate, metal oxides and humic substances) and inhibitors for sulfur cycling
104 and methanogens' activity (Fig. 1):

105 A) Long-term two-stage slurry incubations with a first stage of 1:1 sediment to porewater ratio for three
106 months with high methane content to enrich the microorganisms involved in the AOM. After three
107 months, the slurry was diluted to a 1:3 ratio, then different reactants were added to the incubations,
108 which were monitored for up to 18 months.

109 B) Semi-continuous bioreactor experiments with freshly collected sediments and porewater at a 1:4
110 ratio (respectively), where porewater was exchanged regularly.

111 C) Batch incubation experiments with fresh sampled sediments and porewater at a 1:5 ratio,
112 respectively, and several manipulations (this experimental set-up was described in our previous studies
113 (Bar-Or et al., 2017; Elul et al., 2021)).

114 Here below we describe the experiments. Detailed protocols are found in the supplementary
115 information.

116 2.2.1 Experiment set-up A: Long-term two-stage incubations

117 The sediments for the slurries were collected during several sampling campaigns between 2017 and
118 2019 from the central lake (Station A, Fig. S1) using a gravity corer with 50 cm Perspex cores.
119 Sediments from the methanogenic zone (> 20 cm depth) were diluted with porewater from the
120 methanogenic zone of parallel cores sampled on the same day. The porewater was extracted by
121 centrifugation at 9300 g for 15 minutes, filtered by 0.22 μ M filters into 250 ml glass bottles, sealed with
122 a rubber stopper, and flushed for 30 minutes with N₂.

123 In the first stage, the sediment was diluted with the extracted porewater to create a 1:1 ratio slurry in
124 250 ml glass bottles with a headspace of 70-90 ml under continuous N₂ flushing (Fig. 1). The slurries
125 were flushed with N₂ (99.999%, MAXIMA, Israel) for 30 minutes. Methane was injected using a gas-
126 tight syringe for a final content of 20% in the headspace, where 10% of the injected methane was ¹³C-
127 labeled methane (99%, Sigma-Aldrich). When significant AOM was observed by the increase of $\delta^{13}\text{C}_{\text{DIC}}$
128 after three months (Fig. S2), the incubations were either transferred to the second stage experiments or
129 continued to run with porewater exchange and $\delta^{13}\text{C}_{\text{DIC}}$ values monitored every three months.

130 This study presents ten sets of two-stage incubation experiments with different treatments (electron
131 acceptors/shuttling/inhibitors). They were all set up similarly (protocols in the supplementary
132 information): subsamples (~18 g each) of the pre-incubation slurry were transferred with a syringe
133 under continuous flushing of N₂ gas into 60 ml glass bottles and diluted with fresh anoxic porewater
134 from the methanogenic zone (as described above) to achieve a 1:3 sediment to porewater ratio (Fig. 1),
135 leaving 24 ml of headspace in each experiment bottle. The bottles were crimp-sealed, flushed with N₂

136 gas for 5 minutes, shaken vigorously and flushed again (3 times). ¹³C-labeled methane was added to all
137 the bottles as described in Table 1. The "killed" control bottles in each experiment were autoclaved
138 twice, cooled, and only then were amended with the appropriate treatments and ¹³C-labeled methane.

139 Electron acceptors were added either as a powder (hematite, magnetite, clay, MnO₂, humic substances)
140 or in dissolved form in double-distilled water (DDW) (KNO₃ and NaNO₂). The involvement of sulfur
141 cycling was tested by inhibition with Na-molybdate (Lovley and Klug, 1983), while the other electron
142 acceptors were tested for their potential participation by their addition to the slurries. AQDS was added
143 as an analog for humic substances, which was previously shown to serve as a terminal electron acceptor
144 for AOM and electron shuttling for iron reduction (e.g., Scheller et al., 2016; Sivan et al., 2016).
145 Amorphous iron (Fe(OH)₃) was prepared in the lab by dissolving FeCl₃ in DDW, then titrated with
146 NaOH 1.5 N up to pH 7 and was added to the bottles by injection. The final concentration of each
147 addition is detailed in Table 1. The ¹³C-labeled methane was injected to all experiment bottles at the
148 beginning of each experiment (unless mentioned otherwise) using a gas-tight syringe from a stock bottle
149 filled with ¹³C-labeled methane gas (which was replaced with saturated NaCl solution). Three different
150 inhibitors were added to three different experiments: molybdate, BES and acetylene. Molybdate was
151 added to experiment No. 1 to detect the feasibility of an active sulfur cycle. BES was added to
152 experiment No. 8 at the start of the experiment. Acetylene was added to experiment No. 9. It was
153 injected during the experiment to two bottles at different timepoints after ¹³C enrichment in the DIC
154 was observed (Table 1).

155 All live treatments were set up in duplicates or triplicates and we present the average with an error bar.
156 In two experiments, only one "killed" control bottle was set up. The slurry was prioritized for other
157 treatments since the killed controls showed repetitive no activity for numerous previous experiments.
158 The humic substrate experiment used natural (humic) substance that were extracted from a different
159 lake. One experiment was set up without any additional electron acceptor to assess the rate of
160 methanogenesis in the two-stage slurries. Porewater was sampled anaerobically for δ¹³C_{DIC} and
161 dissolved Fe(II) measurements in duplicates (2 ml), and methane was measured from the headspace.
162 Variations in the δ¹³C_{DIC} values between the experiments resulted from different amounts of ¹³C-labeled
163 methane injected at the start of each experiment.

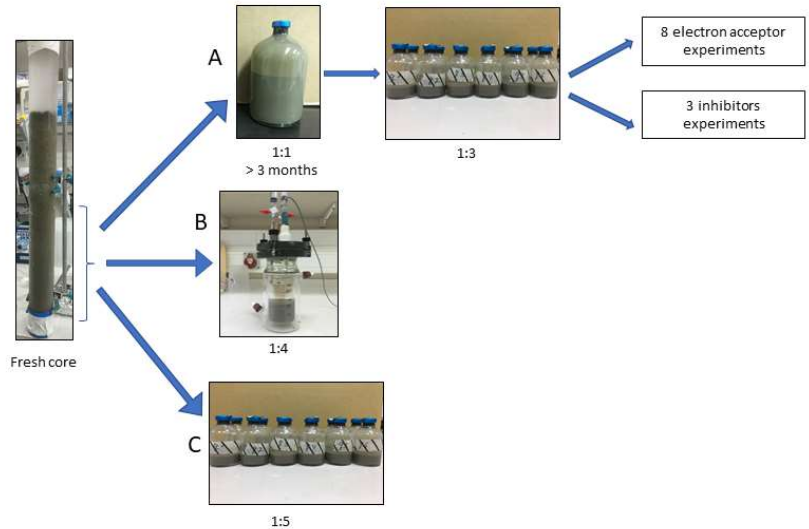
164 2.2.2 Experiment set-up B: Semi-bioreactor

165 Semi-bioreactors regularly monitored the redox state at close-to-natural *in-situ* conditions for 15 months
166 in freshly collected sediments. Two 0.5 L semi-bioreactors (Fig. 1) (LENZ, Weinheim, Germany) were
167 set up with fresh sediments from the methanogenic zone (25 - 40 cm) of Lake Kinneret Station A
168 immediately after their collection. Both reactors were filled headspace-free with a slurry at a 1:4
169 sediment - pore water ratio. One of the bioreactors was amended with 10 mM hematite and the second
170 without it, serving as a control. To dissolve ¹³C-labeled methane in the porewater, 15 ml of porewater

171 were replaced with 15 ml of methane gas (a mixture of $^{12}\text{CH}_4$ and $^{13}\text{CH}_4$) to produce methane-only
 172 headspace for 24 hours. The reactors were shaken repeatedly during those hours. After 24 hours, the
 173 gas was replaced with anoxic porewater, so that there was no headspace at all. This resulted in lower
 174 methane concentrations than the batch experiments (0.2 mM vs. ~2 mM, respectively). Redox potential
 175 was monitored continuously by electrode (Metrohm, Herisau, Switzerland) to verify anoxic conditions
 176 and determine the redox state throughout the incubation period. The bioreactors were subsampled
 177 weekly to bi-weekly, and the sample volume (5-10 ml) was replaced immediately by preconditioned
 178 anoxic (flushed with N_2 gas for 15 minutes) porewater from the methanogenic zone. As outlined below,
 179 samples were analyzed for dissolved Fe(II), CH_4 , and $\delta^{13}\text{C}_{\text{DIC}}$. Additional subsamples for metagenome
 180 and lipid analyses were taken at the beginning of the experiment and on days 151 and 382, respectively.

181 2.2.3 Experiment set-up C: Fresh batch experiment

182 Sediments for this experiment were collected in August 2013 at Station A, similar to the sediments for
 183 the pre-incubations. The sediments below 26 cm depth were diluted under anaerobic conditions with
 184 porewater from the same depth to reach a 1:5 sediment to porewater ratio. The slurry was divided into
 185 60 ml experiment bottles with 40 ml slurry in each bottle. The sampling and experimental set-up details
 186 are described in Bar-Or et al., 2017. Here we present the results of $\delta^{13}\text{C}_{\text{DIC}}$, metagenome and lipid
 187 analyses of two treatments: natural (with only ^{13}C -labeled methane) and hematite. The experiment ran
 188 for 15 months.



189

190 Figure 1: Flow diagram of the experimental design. Three types of experiments were set up from sediments of
 191 the methanogenic zone (below 20 cm): A. Two-stage slurry experiments with diluted pre-incubated slurries and
 192 porewater (1:3 sediment to porewater ratio). Ten experiments were set up this way, 8 of them with different
 193 electron acceptors for 6-18 months, and three different inhibitors for 12-18 months (to one experiment, both
 194 electron acceptors and an inhibitor were added). B. Semi-bioreactor experiment with freshly collected
 195 sediments. C. Fresh batch experiment -slurry experiment with freshly collected sediments (Bar-Or et al., 2017).

Table 1: Specific details of the three types of experiments: two-stage, semi-bioreactor and fresh batch experiments.

Experiment serial number (SN)	Experiment	Treatment	# of bottles	CH ₄ [ml]	¹³ CH ₄ [ml]	Fe ₂ O ₃ [mM]	Fe ₃ O ₄ [mM]	Fe(OH) ₃ [mM]	MnO ₂ [mM]	NO ₂ [mM]	NO ₃ [mM]	AQDS [mM]	Humic substances [mM]	PCA [mM]	Fe-bearing nontronite (day) [gr]	N ₂ -molybdate [mM]	BES [mM]	Acetylene [μL]	Temp (°C)	Duration [day]	Comments
1	Hematite	¹³ CH ₄ , ¹³ C ₁₃ H ₄ +hematite	2	1	1	10													20	201	The methane that was added at the beginning of the experiment was not labelled, so ¹³ C-labeled methane was added after 105 days. N ₂ -molybdate was added to one of the bottles on day 365
2	Magnetite	¹³ CH ₄ , ¹³ CH ₄ +magnetite, ¹³ CH ₄ +Fe(OH) ₃ , Killed+ ¹³ CH ₄ +magnetite	2	1	1	10	10	10									1		16	447	N ₂ -molybdate was added to one of the bottles on day 365
3	MnO ₂	¹³ CH ₄ , ¹³ CH ₄ +MnO ₂	2	1.2	1.2				10										20	201	200 μL ¹³ CH ₄ was added on day 1, then another 1 ml. was added on day 24.
4	Nitrate	¹³ CH ₄ +NO ₃ (high conc.), ¹³ CH ₄ +hematite, ¹³ CH ₄ +NO ₃ (high conc.)+hematite, ¹³ CH ₄ +NO ₃ (low conc.)+hematite, Killed+ ¹³ CH ₄ +NO ₃ (high conc.)+hematite	2	1	0.5	12					1								20	306	
5	Nitrite	¹³ CH ₄ +NO ₂ (high conc.)+hematite, ¹³ CH ₄ +NO ₂ (low conc.)+hematite, Killed+ ¹³ CH ₄ +NO ₂ (high conc.)+hematite	2	1	0.5	10				0.5									20	493	
6	AQDS	¹³ CH ₄ +AQDS, ¹³ CH ₄ +AQDS+hematite, Killed+ ¹³ CH ₄ +AQDS	2	1	1	10						5							20	264	
7	Natural humic acids and clay	¹³ CH ₄ , ¹³ CH ₄ +hematite, ¹³ CH ₄ +humic acid, ¹³ CH ₄ +clay, Killed+ ¹³ CH ₄ +hematite	2	1	1	10							0.5						20	169	The head space of the experiment bottles was flushed with N ₂ on day 51 and ¹³ CH ₄ was added. This was done in order to match the the clay bottles.
8	Bromoethanesulfonate (BES)	¹³ CH ₄ +hematite, ¹³ CH ₄ +hematite+BES, ¹³ CH ₄ +hematite	2	9	1	10											20		20	493	
9	Acetylene	¹³ CH ₄ +hematite+acetylene, Killed+ ¹³ CH ₄ +hematite	2	1	0.5	10												120	20	321	Acetylene was injected to each bottle at different time point during the experiment.
10	No electron acceptor	No additions, ¹³ CH ₄	3	1	1														20	147	
	Semi-bioreactor	¹³ CH ₄ , ¹³ CH ₄ +hematite		15		10													16	345	
	Freshly collected sediment exp.	¹³ CH ₄ , ¹³ CH ₄ +hematite		0.05															16	677	
				0.05		20													20	467	

198 **2.3 Analytical methods**

199 Measurements of $\delta^{13}\text{C}_{\text{DIC}}$ were performed on a DeltaV Advantage Thermo Scientific isotope-ratio mass-
200 spectrometer (IRMS). Results are reported referent to the Vienna Pee Dee Belemnite (VPDB) standard.
201 For these measurements about 0.3 ml of filtered (0.22 μm) porewater was injected into a 12 ml glass
202 vial with He atmosphere and 10 μl of H_3PO_4 85% to acidify all the DIC species to CO_2 (g). The
203 headspace autosampler (CTC Analytics. Type PC PAL) took a gas sample from the vials and measured
204 the $\delta^{13}\text{C}_{\text{DIC}}$ of the sample on the GasBench interface with a precision of ± 0.1 ‰. DIC was measured on
205 the IRMS using the peak height and a precision of 0.05 mM. Dissolved Fe(II) concentrations were
206 measured using the ferrozine method (Stookey, 1970) by a spectrophotometer at 562 nm wavelength
207 with a detection limit of 1 $\mu\text{mol L}^{-1}$. Methane concentrations were measured from the headspace. A 100
208 μL headspace sample was taken with a gas-tight syringe and was analyzed by a focus gas
209 chromatograph (GC) equipped with a flame ionization detector (FID) with a detection limit of 0.005
210 μmol . Bottles to which acetylene was added were measured similarly for ethylene to determine the
211 acetylene turnover with the N cycle.

212 A sub-set of samples (Table 3) was investigated for the assimilation of ^{13}C -labeled methane into polar
213 lipid-derived fatty acids (PLFAs) and intact ether lipid-derived hydrocarbons. A total lipid extract
214 (TLE) was obtained from 0.4 to 1.6 g of the freeze-dried sediment or incubated sediment slurry using a
215 modified Bligh and Dyer protocol (Sturt et al., 2004). Before extraction, 1 μg of 1,2-diheneicosanoyl-
216 *sn*-glycero-3-phosphocholine and 2-methyloctadecanoic acid were added as internal standards. PLFAs
217 in the TLE were converted to fatty acid methyl esters (FAMES) using saponification with KOH/MeOH
218 and derivatization with BF_3/MeOH (Elvert et al., 2003). Intact archaeal ether lipids in the TLE were
219 separated from the apolar archaeal lipid compounds using preparative liquid chromatography (Meador
220 et al., 2014) followed by ether cleavage with BBR_3 in dichloromethane forming hydrocarbons (Lin et
221 al., 2010). Both FAMES and ether-cleaved hydrocarbons were analyzed by GC-mass spectrometry (GC-
222 MS; Thermo Finnigan Trace GC coupled to a Trace MS) for identification and GC-IRMS (Thermo
223 Scientific Trace GC coupled via a GC Isolink interface to a Delta V Plus) for determination of $\delta^{13}\text{C}$
224 values using the column and temperature program settings described by Aepfler et al. (2019). The $\delta^{13}\text{C}$
225 values are reported with an analytical precision better than 1‰ as determined by long-term
226 measurements of an *n*-alkane standard with known isotopic composition of each compound. Reported
227 fatty acid isotope data are corrected for the introduction of the methyl group during derivatization by
228 mass balance calculation similar to equation 1 using the measured $\delta^{13}\text{C}$ value of each FAME and the
229 known isotopic composition of methanol as input parameters.

230 For the metagenomic analyses, total genomic DNA was extracted from the semi-bioreactor experiment
231 (duplicates a and b), pre-incubation slurries ($^{13}\text{CH}_4$ -only control, $^{13}\text{CH}_4$ + hematite) and their respective
232 initial slurries (t0), using the DNeasy PowerLyzer PowerSoil Kit (QIAGEN). Genomic DNA was eluted

233 using 50 μ l of elution buffer and stored at -20°C . Metagenomics libraries were prepared at the
 234 sequencing core facility at the University of Illinois at Chicago using Nextera XT DNA library
 235 preparation kit (Illumina, USA). 19-40 million 2×150 bp paired-end reads per library were sequenced
 236 using Illumina NextSeq500. Metagenomes were co-assembled from concatenated reads of all
 237 metagenomic libraries with Spades V3.12 (Bankevich et al., 2012; Nurk et al., 2013), following
 238 decontamination, quality filtering (QV= 10) and adapter-trimming with the BBDuk tool from the
 239 BBDuk suite (Bushnell B, <http://sourceforge.net/projects/bbmap/>). Downstream analyses, including
 240 reading coverage estimates, automatic binning with maxbin (Wu et al., 2014) and metabat2 (Kang et
 241 al., 2019) bin refining with DAS tool (Sieber et al., 2018), were performed within the SqueezeMeta
 242 framework (Tamames and Puente-Sánchez, 2019). GTDB-Tk was used to classify the metagenome-
 243 assembled genomes (MAGs) based on Genome Taxonomy Database release 95 (Parks et al., 2021).
 244 The principal component analysis biplot was constructed with Past V4.03 (Hammer et al., 2001).

245 Methanogenesis rate was calculated from temporal changes in methane concentration in a representative
 246 pre-incubated slurry experiment (Fig. S3). The amount of methane oxidized was calculated by a simple
 247 mass balance calculation according to equations 1 and 2:

$$248 \quad x \times F^{13}\text{CH}_4 + (1 - x) \times \text{FDI}^{13}\text{C}_i = \text{FDI}^{13}\text{C}_f \quad (1)$$

$$249 \quad [\text{CH}_4]_{\text{ox}} = x \times [\text{DIC}]_f \quad (2)$$

250 The final DIC pool comprises two end members; the initial DIC pool and the oxidized ^{13}C - CH_4 . The
 251 term x denotes the fraction of oxidized ^{13}C - CH_4 , while $1-x$ denotes the fraction of the initial DIC pool
 252 out of the final DIC pool. $F^{13}\text{CH}_4$ is the fraction of ^{13}C out of the total CH_4 at t_0 , $\text{FDI}^{13}\text{C}_i$ is the fraction
 253 of ^{13}C out of the total DIC at t_0 , and $\text{FDI}^{13}\text{C}_f$ is the fraction of ^{13}C out of the total DIC at t -final. $[\text{CH}_4]_{\text{ox}}$
 254 is the amount (concentration in pore water) of the methane oxidized throughout the full incubation
 255 period, and $[\text{DIC}]_f$ is the DIC concentration at t -final. It was assumed that the isotopic composition of
 256 the labeled CH_4 did not change significantly throughout the incubation period.

257 **3. Results**

258 In ten sets of slurry incubation experiments, we followed the progress of the methane oxidation process
 259 in (type A) long-term two-stage incubations from Lake Kinneret methanogenic sediments (Figs. 2 and
 260 3) by monitoring the changes $\delta^{13}\text{C}_{\text{DIC}}$ values, metagenomic and specific isotope lipid analyses. We also
 261 followed methane oxidation in a semi-bioreactor system (type B) with freshly collected sediments with
 262 or without the addition of hematite (Fig. 2). The results were compared to fresh batch slurry incubations
 263 (type C) from the same methanogenic zone, presented by Bar-Or et al. (2017) and Elul et al. (2021).

264 **3.1 Geochemical trends in the two-stage experiments**

265 In the two-stage experiments (type A), there was a conversion of ^{13}C -methane to ^{13}C -DIC in all the
266 natural non-killed slurries, indicating significant AOM (Figs. 2 and 3). The $\delta^{13}\text{C}_{\text{DIC}}$ values of the natural
267 sediment amended only with ^{13}C -methane treatments (the "methane-only" control) reached up to 743‰,
268 even with the low abundance of microbial populations in these sediments. Average AOM rate in the
269 methane-only controls was $2.0 \pm 0.4 \text{ nmol gr}^{-1} \text{ dry sediment day}^{-1}$ (Table 2). At the same time,
270 methanogenesis occurred with a net methanogenesis rate of $\sim 25 \text{ nmol gr dry sediment}^{-1} \text{ day}^{-1}$ (Fig. S3,
271 Table S2, Table 2.). The two-stage geochemical experiments tested first the potential of several electron
272 acceptors to perform and stimulate the AOM process, as detailed below.

273 3.1.1 Metals as electron acceptors

274 Iron and manganese oxides were added as potential electron acceptors. The addition of hematite to three
275 different treatments increased the $\delta^{13}\text{C}_{\text{DIC}}$ values with time and reached up to 694‰ (Fig. 2), similarly
276 to the natural (methane-only) controls. The average AOM rate in those treatments was $1.0 \pm 0.3 \text{ nmol gr}$
277 $\text{dry sediment}^{-1} \text{ day}^{-1}$ (Table 2). Magnetite amendments resulted in a minor increase of $\delta^{13}\text{C}_{\text{DIC}}$ values
278 compared to the methane-only controls (200‰ and 265‰, respectively, Fig. 3A) with an AOM rate of
279 $1.8 \text{ nmol gr dry sediment}^{-1} \text{ day}^{-1}$. Amorphous iron amendments resulted in only a 22‰ increase in
280 $\delta^{13}\text{C}_{\text{DIC}}$ and a lower AOM rate ($0.1 \text{ nmol gr dry sediment}^{-1} \text{ day}^{-1}$, Fig. 3A and Table 2). The addition of
281 iron-bearing clay nontronite did not cause any increase in the $\delta^{13}\text{C}_{\text{DIC}}$ values but dissolved Fe(II)
282 concentrations increased compared to the natural methane-only control (Fig. 3B, Fig. 4). No AOM was
283 detected 200 days following the addition of MnO_2 based on $\delta^{13}\text{C}_{\text{DIC}}$ estimates, whereas the $\delta^{13}\text{C}_{\text{DIC}}$
284 values of the methane-only controls reached over 500‰ (Fig. 3F).

285 3.1.2 Sulfate as an electron acceptor

286 The involvement of sulfate in the AOM of two-stage incubations was tested to detect the feasibility of
287 an active cryptic sulfur cycle, even with the absence of detectable sulfate in the methanogenic sediments
288 (Holmkvist et al., 2011). It was quantified directly by adding Na-molybdate, an inhibitor of sulfate
289 reducers and sulfur disproportionators, to the methane-only controls and slurries amended with
290 magnetite (Fig. 3A). This addition did not change the increase of $\delta^{13}\text{C}_{\text{DIC}}$ with time, and thus the AOM
291 rates, similar to the observation in the fresh batch incubations (Bar-O et al., 2017).

292 3.1.3 Nitrate and nitrite as electron acceptors

293 Nitrate and nitrite involvement in the AOM was tested to detect the feasibility of an active cryptic
294 nitrogen cycle, even with the absence of detectable nitrate and nitrite in the sediments. Nitrate was
295 added at two different concentrations (0.2 and 1 mM, Fig. 3C) to the two-stage slurries amended with
296 hematite, as these concentrations were shown previously to promote AOM in other settings (Ettwig et
297 al., 2010). Hematite addition alone increased the $\delta^{13}\text{C}_{\text{DIC}}$ values by $\sim 200\%$ during the 306 days of the

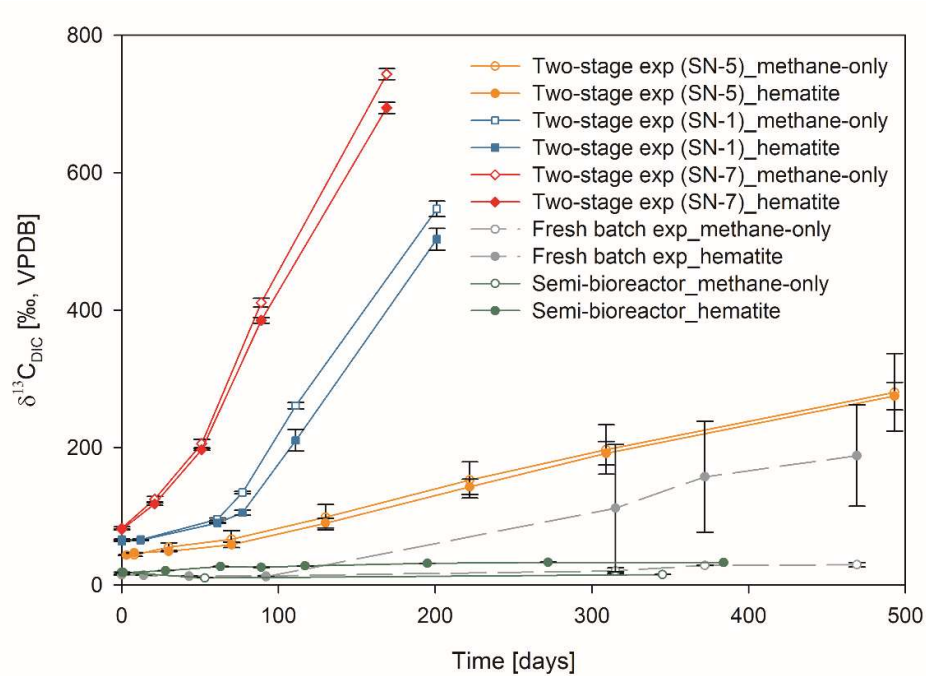
298 experiment. The $\delta^{13}\text{C}_{\text{DIC}}$ in the bottles with the addition of 1 mM nitrate, with and without hematite
299 (Fig. 3C; the data points of the two treatments are on top of each other) decreased from 43‰ at the
300 beginning of the experiment to 35‰ after 306 days. The $\delta^{13}\text{C}_{\text{DIC}}$ in the bottles with the addition of 0.2
301 mM nitrate and hematite increased by 27‰ at the end of the experiment. We also observed no increase
302 in $\delta^{13}\text{C}_{\text{DIC}}$ during the first 222 days following the addition of 0.5 mM of nitrite (Fig. 3D), then $\delta^{13}\text{C}_{\text{DIC}}$
303 increased by 19‰ until the incubation was terminated. The respective AOM rate was 0.2 nmol gr dry
304 sediment⁻¹ day⁻¹ (Table 2). Following the addition of 0.1 mM nitrite, $\delta^{13}\text{C}_{\text{DIC}}$ increased only after 130
305 days and reached 158‰ at day 493. The respective AOM rate was 0.5 nmol gr dry sediment⁻¹ day⁻¹. In
306 the methane-only controls, $\delta^{13}\text{C}_{\text{DIC}}$ value reached a maximum of 330‰.

307 3.1.4 Organic compounds as electron acceptors

308 Two of the two-stage incubation experiments were amended with synthetic and natural organic electron
309 acceptors to test the potential of organic electron acceptors. The addition of AQDS to slurries with and
310 without hematite decreased the $\delta^{13}\text{C}_{\text{DIC}}$ values during the entire experiment duration (Fig. 3E). The
311 dissolved Fe(II) showed an increase of 50 μM in these treatments, whereas without AQDS there was
312 an increase of 20 μM (Fig. S4). We further tested the effect of naturally occurring humic substances
313 using those isolated from a different natural lake. The results show that the $\delta^{13}\text{C}_{\text{DIC}}$ values did not change
314 at the beginning of the experiments (Fig. 3B), while a steep increase of ~ 90 μM in their Fe(II)
315 concentrations were observed (Fig. 4). After 20 days, the $\delta^{13}\text{C}_{\text{DIC}}$ values of these slurries started to
316 increase dramatically from 84‰ to 150‰ with an AOM rate of 1.2 nmol gr dry sediment⁻¹ day⁻¹ (Fig.
317 3B, Table 2). We observed a mirrored trend of the dissolved Fe(II) concentrations to that of $\delta^{13}\text{C}_{\text{DIC}}$
318 with a steep increase during the first 20 days followed by a decrease of 37 μM (Fig. 4).

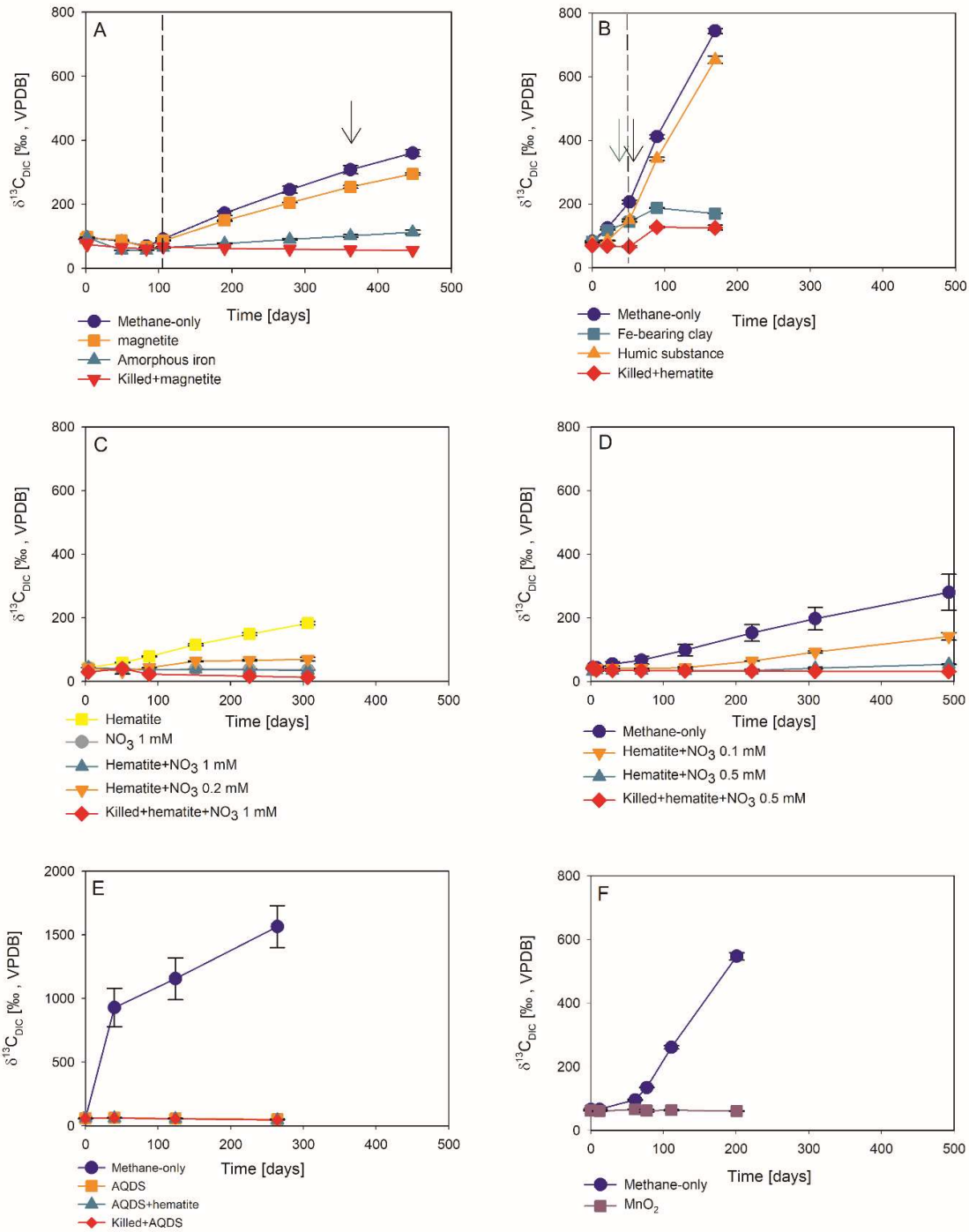
319 To evaluate which metabolic processes drive AOM, we analyzed $\delta^{13}\text{C}_{\text{DIC}}$ following the addition of
320 inhibitors: i) BES, a specific inhibitor for methanogens and ANME's *mcrA* genes and ii) acetylene, a
321 non-specific inhibitor for methanogens. Both cases showed complete inhibition of labeled ¹³C-DIC
322 production following the addition, similarly to the killed control (Fig. 5). Acetylene can also inhibit
323 nitrogen cycling in some cases; however, this has been shown to result in the production of ethylene
324 (Oremland and Capone, 1988). In our case, no ethylene was detected, supporting the inhibition only of
325 methanogens' activity.

326



327

328 Figure 2: Comparison of $\delta^{13}\text{C}_{\text{DIC}}$ values among the three types of experiments: A) three two-stage slurry
 329 experiments; B) the semi-bioreactor experiment; and C) slurry batch experiment with freshly collected sediments
 330 (Bar-Or et al., 2017). In each experiment, two treatments are shown, with hematite (filled symbol) and without
 331 (empty symbols) hematite addition. The error bars represent the average deviation of the mean of
 332 duplicate/triplicate bottles.



333

334 Figure 3: The potential of different electron acceptors for AOM in Lake Kinneret in the pre-incubated long-term
 335 slurry experiments with the following treatments: (A) with and without the addition of magnetite and amorphous
 336 iron ($\text{Fe}(\text{OH})_3$). The dashed line represents the addition of ^{13}C -labeled CH_4 . The black arrow represents the
 337 addition of Na-molybdate as an inhibitor for sulfate reduction. (B) with clay and natural humic substance. The
 338 green arrow represents the time clay was added to the relevant bottles, the dashed line represents the time the
 339 headspace of the bottles was flushed again with N_2 , and the black arrow represents the second injection of 1 mL
 340 of ^{13}C -labeled methane. (C) with the addition of hematite and two different concentrations of nitrate. (D) with the

341 addition of hematite and two different concentrations of nitrite. (E) with the addition of AQDS. (F) with and
 342 without the addition of ¹³C -labeled methane was added to all the bottles (specific details on each experiment can
 343 be found in Table 1). Error bars represent the average of the absolute deviations of data points from their mean.

344

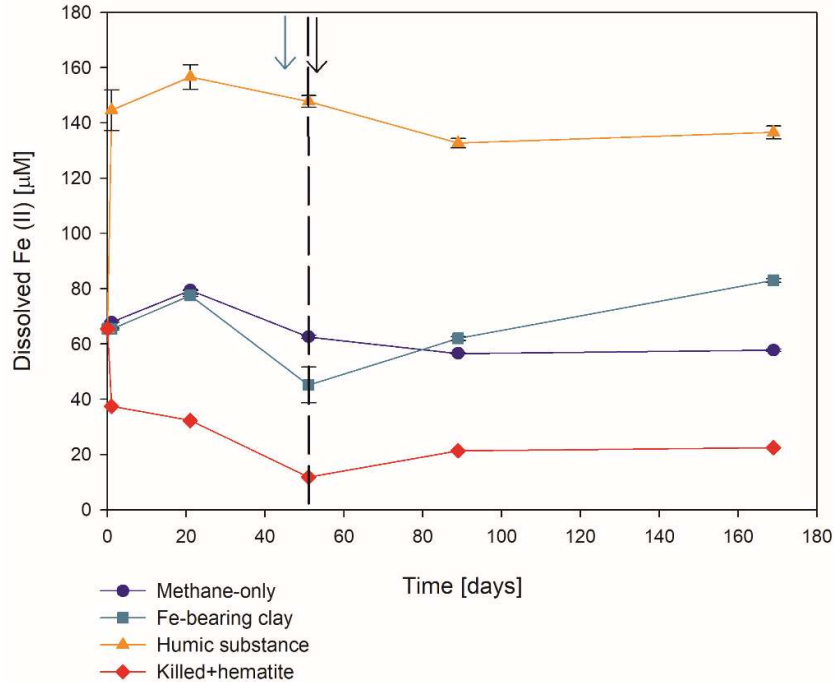
345 Table 2: Methanogenesis and AOM rates in experiment A (two-stage slurries) amended with ¹³C-labeled methane
 346 and different electron acceptors (methanogenesis rate was calculated in one of the experiments and was assumed
 347 to be similar in all of them).

Experiment serial number (SN)	Treatment	Methanogenesis rate [nmol/gr dry sediment X day]	AOM rate [nmol/gr dry sediment X day]	AOM/methanogenesis [%]
10	methane only	24.8	1.1	4.4
1	methane only	24.8	1.6	6.4
	methane+hematite	24.8	0.5	2.1
2	methane only	24.8	2.4	8.2
	methane+magnetite	24.8	1.8	6.3
	methane+amorphous iron	24.8	0.1	0.5
7	methane only	24.8	1.4	6.4
	methane+hematite	24.8	1.3	6.0
	methane+humics	24.8	1.2	5.4
5	methane only	24.8	1.0	4.6
	methane+hematite	24.8	1.0	4.6
	methane+hematite+nitrite 0.5 mM	24.8	0.2	0.8
	methane+hematite+nitrite 0.1 mM	24.8	0.5	2.1

348

349

350



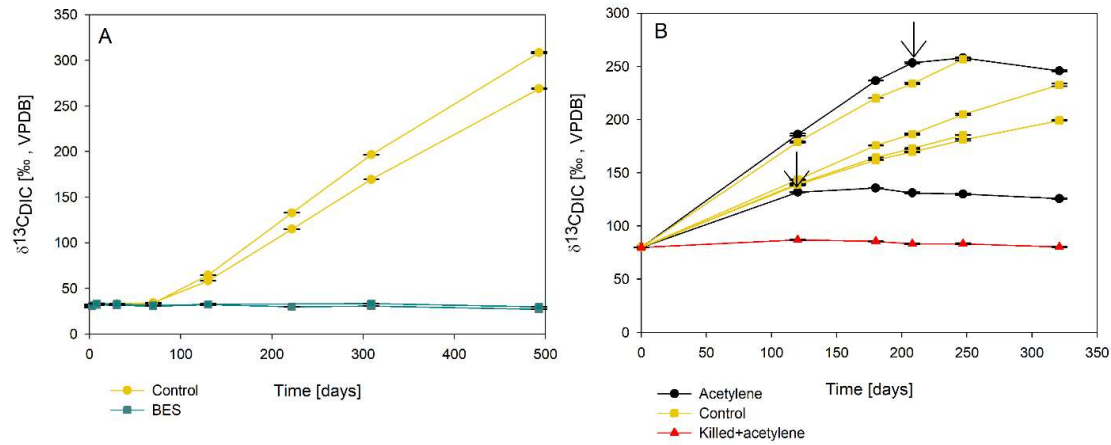
351

352 Figure 4: The dissolved Fe(II) change in the two-stage experiment No. 7 containing clay, natural humic acid, and
 353 PCA. The green arrow represents the time clay was added to the specific bottles and those bottles flushed with
 354 N₂, the dashed line represents the time the rest of the bottles were flushed, and the black arrow represents the time
 355 ¹³CH₄ was added again. Error bars represent the average of the absolute deviations of data points from their mean.

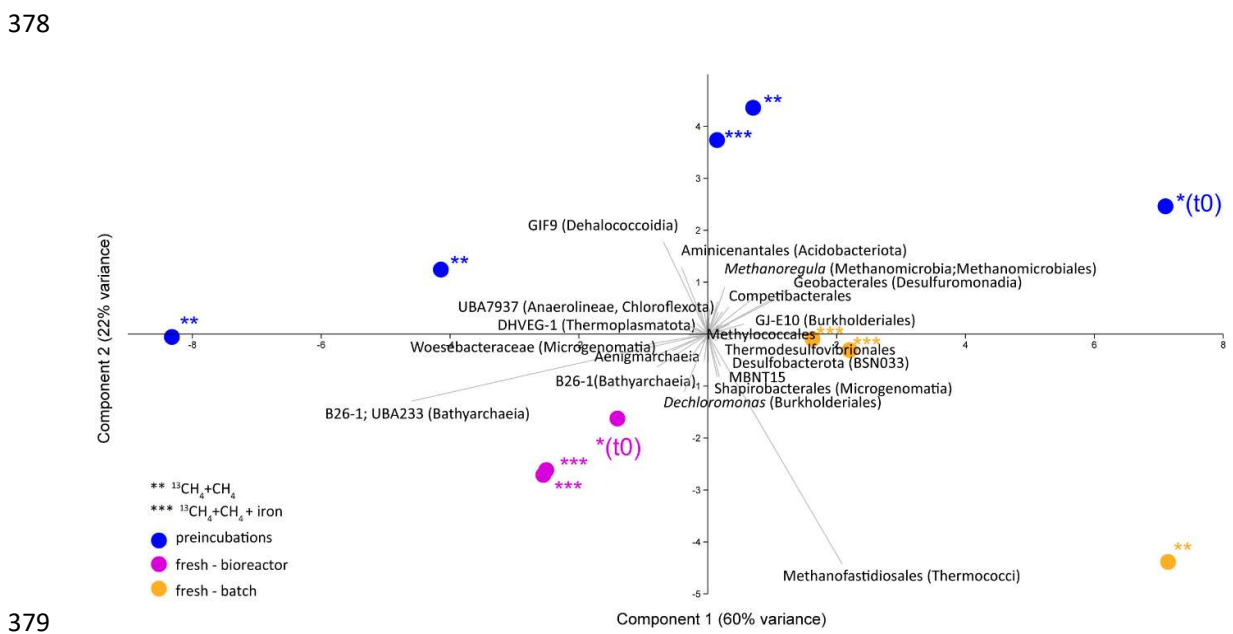
356 3.2 Microbial dynamics

357 Analyses of taxonomy and coverage of metagenome-assembled genomes suggest that in the pre-
 358 incubated slurries, Bathyarchaea are the dominant archaea, together with putative methanogens such
 359 as Methanofastidiales (Thermococci), Methanoregulaceae (Methanomicrobia) and Methanotrichales
 360 (Methanosarcinia) (Supplementary coverage table). Bonafide ANME (ANME-1) were detected at
 361 substantial coverage of approximately 1 (the 27th most abundant out of 195 MAGs) in all the treatments.
 362 Among bacteria, sulfate reducers Desulfobacterota and Thermodesulfobivibrionales (Nitrospirota) were
 363 prominent together with the GIF9 Dehalococcoida lineage, which is known to metabolize chlorinated
 364 compounds in lake sediments (Biderre-Petit et al., 2016). Some Methylospirales (NC10) were found
 365 (average coverage of 0.32±0.06), and no Methanoperedens were detected. Methylococcales
 366 methanotrophs were found in the natural sediments and fresh batch and bioreactor incubations (average
 367 of 0.34±0.02), as opposed to the average coverage of 0.09±0.04 in long-term incubations.
 368 Methylococcales comprised *Methylothermophilum*, *Methylothermobacter* and *Methylobacter* genera
 369 (Supplementary coverage table). The methylophilic partners of aerobic methanotrophs,
 370 *Methylothermophilum*, were found in fresh batch and bioreactor incubations, where *Methylothermobacter* was found,
 371 in line with previous studies showing their association (Beck et al., 2013). Principal component analysis

372 shows the grouping of long-term pre-incubated slurries, semi-bioreactor incubations, and fresh batch
 373 experiments (Fig. 6), emphasizing microbial dynamics over time.



374
 375 Figure 5: The change of $\delta^{13}\text{C}_{\text{DIC}}$ values with time in two long-term sediment slurry incubations amended with
 376 hematite and ^{13}C -labeled methane. (A) with/out BES and (B) with/out acetylene. Black arrows represent the time
 377 at which acetylene was injected to the experiment bottle. The error bars are smaller than the symbols.



379
 380 Figure 6: Principal component analysis comparing three types of samples: long-term pre-incubated slurries (blue
 381 – experiment A), semi-bioreactor (pink – experiment B) and fresh batch experiments (orange – experiment C).
 382 One asterisk represents t0, two asterisks denote methane-only treatments, three asterisks represent hematite
 383 treatment.

384
 385 **3.3 Lipid analysis**

386 The $\delta^{13}\text{C}$ values of the archaeol-derived isoprenoid phytane in the long-term pre-incubated samples
 387 were between -5 and -17‰ and thus showed a ^{13}C -enrichment between 15-27‰ relative to the original
 388 sediment, indicative of methane-derived carbon assimilation by archaea (Table 3). This was less
 389 pronounced for acyclic biphytane, dominantly derived from caldarchaeol, which showed a ^{13}C -
 390 enrichment of 5-10‰. For bacterial-derived fatty acids, the shift in $\delta^{13}\text{C}$ -values of up to 10‰ relative
 391 to the original sediment was in a similar range but would have been expected to be much higher if
 392 aerobic methanotrophs were active as was previously indicated by the extreme ^{13}C -enrichment of up to
 393 1,650‰ observed in freshly incubated batch samples (Bar-Or et al., 2017).

394 Table 3: The $\delta^{13}\text{C}$ values (in ‰) of fatty acids and isoprenoid hydrocarbons from different experiments compared
 395 to values obtained from the original sediment in the methanogenic zone.

Description	Temperature (°C)	Sampling (days)	Fatty acids		Hydrocarbons	
			C _{16:1ω9/8/7}	C _{16:1ω5}	Phytane	Biphytane
Pre-incubated slurry + $^{13}\text{CH}_4$ +hematite	20	411	-40	-43	-17	-23
Pre-incubated slurry + $^{13}\text{CH}_4$ (bottle A)	20	411	-40	-43	-13	-24
Pre-incubated slurry + $^{13}\text{CH}_4$ (bottle B)	20	1227	-36	-41	-5	-38
^a Fresh batch experiment+ $^{13}\text{CH}_4$ +hematite	20	470	610	1600	-14	-28
Semi-bioreactor+ $^{13}\text{CH}_4$ +hematite	16	382	n.d.	n.d.	n.d.	n.d.
Original sediment (28-30 cm)	14		-44	-51	-32	-33

^a Bar-Or et al., 2017

n.d. – Not detected

396

397 **4. Discussion**

398 **4.1 AOM is maintained in long-term two-stage incubation experiments**

399 Our previous porewater profiles of Lake Kinneret indicate that microbial sulfate reduction dominates
 400 the anoxic hypolimnion and the surface sediments, while methanogenesis is confined to the sediments
 401 below the sulfate boundary (Adler et al., 2011; Sivan et al., 2011; Bar-Or et al., 2015). The *in-situ*
 402 geochemical and microbial diversity profiles, as well as geochemical and metagenomic analyses of
 403 batch incubations with fresh sediments, provided evidence for Fe-AOM in the deep methanogenic zone
 404 below 20 cm depth (Adler et al., 2011; Sivan et al., 2011; Bar-Or et al., 2015; Bar-Or et al., 2017; Elul
 405 et al., 2021; Fig. 2). The profiles and the incubations showed an unexpected presence of aerobic
 406 bacterial methanotrophs together with anaerobic microorganisms, such as methanogens and iron
 407 reducers, in the anoxic sediments. They suggested that both *mcr* gene-bearing archaea and aerobic
 408 bacterial methanotrophs mediate methane oxidation. In this study analyses of ^{13}C -DIC derived from
 409 ^{13}C -labeled methane suggest that considerable AOM takes place also in the long-term incubations, even
 410 after the two stages and the low abundance of the microbial populations. Below, we characterize this
 411 AOM process in these incubation experiments.

412 **4.1 Potential electron acceptors for AOM in the long-term two-stage incubation experiments**

413 The pre-incubated long-term incubations data show a sharp increase in the $\delta^{13}\text{C}_{\text{DIC}}$ values of both natural
414 (methane-only) and hematite amendments in the two-stage incubations (Fig. 2). However, no difference
415 in $\delta^{13}\text{C}_{\text{DIC}}$ between the two treatments was observed following the addition of hematite as the electron
416 acceptor. This differs from the experiments B and C observations with fresh sediment, where the
417 addition of hematite showed higher values than the methane-only treatment (Fig. 2; Bar-Or et al., 2017).
418 This was particularly dramatic in the batch slurries (experiment C), but it was also significant in the
419 semi-bioreactor (experiment B). We believe that the difference in the bioreactors would have been more
420 pronounced if methane concentrations were higher, but it is still significant. The results suggest that
421 either hematite lacks the potential to stimulate the AOM activity during long-term experiments or the
422 presence of enough natural Fe(III) in the sediments to sustain the maximum potential of Fe-AOM.

423 Measurements of $\delta^{13}\text{C}_{\text{DIC}}$ show that additions of magnetite, amorphous iron, ferric iron from clays and
424 manganese oxide in the two-stage incubations result in a less pronounced increase in the $\delta^{13}\text{C}_{\text{DIC}}$ values
425 compared to the methane only controls (Figs. 2 and 3), reducing the AOM signal. One possible
426 explanation is that these metal oxides may inhibit AOM, either directly or by a preference for
427 organoclastic iron reduction over Fe-AOM, which adds isotopically light carbon from the organics
428 rather than heavy carbon from the ^{13}C -labeled methane. Using mass-balance estimations in the methane-
429 only treatments and the amorphous iron ones and considering the DIC concentrations and $\delta^{13}\text{C}_{\text{DIC}}$ values
430 of the methane-only treatments at the beginning of the experiment (6 mM and 60‰, respectively) and
431 the values at the end (6.5 mM and 360‰, respectively), about 0.5 mM of the DIC was added by AOM
432 of methane with $\delta^{13}\text{C}$ of ~4000‰. The DIC and $\delta^{13}\text{C}_{\text{DIC}}$ values of the amorphous iron treatment at the
433 beginning of the experiment were 5.4 mM and 60‰, respectively, and the values at the end were 6.1
434 mM and 120‰, respectively. Assuming the same $\delta^{13}\text{C}$ of the added methane of 4000‰ and $\delta^{13}\text{C}_{\text{TOC}}$ of
435 -30‰ (Sivan et al., 2011), 0.1 mM of the DIC should derive from AOM and 0.6 mM from organoclastic
436 metabolism. This means that adding amorphous iron to the system decreased the AOM activity and
437 encouraged the oxidation of other organic compounds rather than methane. Intrinsic microbes,
438 particularly the commonly detected ex-deltaproteobacterial lineages such as Geobacterales, may
439 catalyze Fe(III) metal reduction, regardless of AOM. Manganese oxides are found in very low
440 abundance in Lake Kinneret sediments (0.1 %, Table S1). Thus, their role in metal-AOM is likely
441 minimal.

442 Sulfate concentrations in the methanogenic Lake Kinneret sediments are low (< 5 μM , Bar-Or et al.,
443 2015; Elul et al., 2021). Sulfide concentrations have also been reported to be minor (<0.3 μM , Sivan et
444 al., 2011). However, since pyrite and FeS precipitate in the top sediments, cryptic cycling via pyrite or
445 FeS may replenish sulfate available for AOM (Bottrell et al., 2000). Na-molybdate addition to the two-
446 stage slurries, including those amended with and without magnetite, did not change the $\delta^{13}\text{C}_{\text{DIC}}$
447 dynamics, which remained similar to those from before the inhibitor's addition (Fig. 3A). This is in line

448 with the fresh batch sediment slurries (Bar-Or et al., 2017) and hints that sulfate is not a potent electron
449 acceptor for AOM in this environment. Furthermore, although sulfate-reducing bacteria were abundant,
450 none of these reducers belonged to the known clades of ANME partners (Supplementary coverage
451 table).

452 The nitrate and nitrite concentrations are undetectable in the porewater of Lake Kinneret sediments
453 (Nüsslein et al., 2001; Sivan et al., 2011), but they may occur as an intermediate product of ammonium
454 oxidation coupled to iron reduction. We thus assessed the role of nitrate and nitrite as electron acceptors
455 in the two-stage slurries. The results indicate that the addition of nitrate delayed AOM and likely
456 promoted denitrification. This is consistent with the fact that ANME-2d was not found. In the case of
457 nitrite, even low concentrations appeared to delay the increase in $\delta^{13}\text{C}_{\text{DIC}}$ values, suggesting that
458 organoclastic denitrification outcompetes AOM, and nitrite-AOM is not prominent in the two-stage
459 incubations, despite the occurrence of *Methylomirabilia* (Figs. 3C, D).

460 Humic substances may promote AOM by continuously shuttling electrons to metal oxides (Valenzuela
461 et al., 2019). Humic substances were not measured directly in Lake Kinneret sediments, but the DOC
462 concentrations in porewater at the methanogenic depth were high (~1.5 mM, Adler et al., 2011),
463 suggesting that they play a role in AOM. The addition of the synthetic humic analogs AQDS did not
464 cause any enrichment in ^{13}C -DIC, but an increase of the dissolved Fe(II) concentrations compared to
465 the methane-only treatments. This may be explained by AQDS acting as an electron shuttle in
466 organoclastic iron reduction, producing isotopically light carbon that masks the AOM signal (Fig. 3E,
467 Fig. S4). Yet, the natural humic substances may support AOM, as was suggested by Valenzuela et al.
468 (2017). In our incubations, the natural humic substances promoted oxidation of organic matter and iron
469 reduction at first, probably by shuttling electrons from organic compounds other than methane to natural
470 iron oxides in the sediments (Figs. 3B and 4). Then, when the availability of the iron oxides or the
471 organic matter decreased, humic substances likely facilitated AOM (Fig. 3B).

472 Overall, our long-term batch experiments, which included different electron acceptors, indicate that
473 sulfate, nitrate, nitrite and Mn-oxides do not support AOM in Lake Kinneret methanogenic sediments.
474 The potential electron acceptors are natural humic substrates with or without iron minerals that are
475 abundant in the sediment and preferably react with methane rather than with other organics. The
476 involvement of iron oxides in the AOM will be further explored after removing natural iron oxides from
477 the sediments to simulate iron limitation.

478 **4.2 Main microbial players in the long-term pre-incubated slurries**

479 Methane oxidation in the pre-incubated Lake Kinneret sediments is likely mediated by either ANMEs
480 or methanogens, as the addition of BES, a specific inhibitor for methanogens and ANME's *mcrA* genes,
481 and acetylene immediately stopped the AOM, similarly to the killed bottles, and the BES addition to
482 fresh sediment experiments (Bar-Or et al., 2017) (Fig. 5). Apart from methane-metabolizing organisms,

483 acetylene can inhibit nitrogen cycling, resulting in ethylene production (Oremland and Capone, 1988).
484 This is not the case in our incubations, as no ethylene was produced. The increase in $\delta^{13}\text{C}$ values in
485 phytane and biphytane (Table 3) also indicates active archaeal methanogens or ANMEs (Wegener et
486 al., 2008; Kellermann et al., 2012; Kurth et al., 2019).

487 Using the isotopic composition of specific lipids and metagenomics, we identified a considerable
488 abundance of aerobic methanotrophs and methylotrophs in the fresh sediments, but not in the pre-
489 incubation slurries (Table 3, Fig. 6), suggesting a minor role of these lineages in the latter. The
490 metagenomic data (Fig. 6, Supplementary coverage table) also indicate that Bathyarchaeia, which might
491 be involved in methane metabolism (Evens et al., 2015), were enriched in the bioreactor incubations,
492 yet their role in Lake Kinneret AOM remains to be evaluated. ANME-1 are likely mediators of AOM in
493 these sediments, although methane oxidation via the reverse methanogenesis is feasible for some
494 methanogens in Lake Kinneret sediments (Elul et al., 2021). We also observed changes in abundance
495 of bacterial degraders of organic matter and necromass: for example, GIF9 Dehalococcoidia, which can
496 metabolize complex organics under methanogenic conditions (Cheng et al., 2019; Hug et al., 2013),
497 were most abundant in long-term incubations (Fig. 6, Supplementary coverage table).

498 **4.3 Mechanism of methane oxidation in the long-term incubations – AOM versus back flux**

499 Our results indicate net methanogenesis in the two-stage incubation experiments with an average rate
500 of $25 \text{ nmol gr}^{-1} \text{ dry sediment day}^{-1}$ (Table 2, Fig. S3 and Table S2), similar to fresh incubation
501 experiments (Bar-Or et al., 2017), despite the overall increasing trend of $\delta^{13}\text{C}_{\text{DIC}}$ values resulting from
502 potential methane turnover (Figs. 2 and 3). A likely explanation for both signals is an interplay between
503 methane production and oxidation, with the latter triggered by reverse methanogenesis in bona fide
504 ANMEs or some methanogens (Hallam et al., 2004; Timmers et al., 2017). Due to the overall production
505 of methane and the lack of intensive stimulation of AOM by any electron acceptor, the significant
506 increase in $\delta^{13}\text{C}_{\text{DIC}}$ values could theoretically result from carbon back flux during methanogenesis,
507 which is feasible in environments close to thermodynamic equilibrium (Gropp et al., 2021). We used
508 DIC mass balance calculations to determine whether back flux can be accounted for in the incubations.
509 Based on equations 1 and 2, 3-8% of the ^{13}C -methane should be converted into DIC to reach the
510 observed ^{13}C -enrichment. These estimates are orders of magnitude higher than the previously reported
511 0.001-0.3% values for methanogenesis back flux in cultures (Zehnder and Brock, 1979; Moran et al.,
512 2005), and in the same range of 3.2 to 5.5% of back flux observed in ANME-enrichment cultures (Holler
513 et al., 2011). In contrast, modeling approaches from AOM-dominated marine sediment samples and
514 associated ANME enrichment cultures indicated the absence of net methanogenesis (Yoshinaga et al.,
515 2014; Chuang et al., 2019; Meister et al., 2019; Wegener et al., 2021). Thus, it is unlikely that back flux
516 alone can account for the methane-DIC conversion in Lake Kinneret sediments. Moreover, just back
517 flux in marine methanogenic sediment with similar net methanogenesis rates and abundant methane-

518 metabolizing archaea did not yield any significant ¹³C-enrichment in the DIC pool following sediment
519 incubations (Sela-Adler et al., 2015; Amiel, 2018; Vigderovich et al., 2019; Yorshensky, 2019) (Table
520 S3). Therefore, methanogenesis back flux alone seems less likely to sustain the observed DIC values
521 than active AOM.

522 **Conclusions**

523 The geochemical and microbial profiles together with fresh sediment incubations showed evidence for
524 Fe-AOM in the methanogenic zone of Lake Kinneret, which removes about 10-15% of the produced
525 methane (Adler et al., 2011; Sivan et al., 2011). Anaerobic archaea appear to carry out methane turnover
526 in these reduced sediments by reverse methanogenesis, but aerobic Methylococcales may be involved
527 in methane oxidation, which is in line with other evidence of aerobic bacterial activity in the deep anoxic
528 hypolimnion of lakes and their shallow sediments (Beck et al., 2013; Oswald et al., 2016; Martinez-
529 Cruz et al., 2017; Cabrol et al., 2020). The simultaneous presence of aerobes and anaerobes in nature,
530 even 20 meters below the thermocline and oxycline, may result from trace amounts of oxygen trapped
531 in nano-niches or even in mineral layers (Wang et al., 2018), even if sensitive sensors do not detect
532 them. This oxygen portion may not be removed by purging at the beginning of our experiments but is
533 rather slowly used by the methanotrophs for their survival. However, after several incubation stages
534 and intensive purging for a prolonged time, only archaea remained active and were involved in methane
535 turnover, which was most likely coupled to the reduction of electron acceptors such as humic substances
536 and iron.

537 **Competing interests.** The authors declare that they have no conflict of interest.

538 **Acknowledgements**

539 We would like to thank B. Sulimani and O. Tzabari from the Yigal Allon Kinneret Limnological
540 Laboratory for their onboard technical assistance. We thank all of O. Sivan's lab members for their help
541 during sampling, and especially to N. Lotem for the help with the mass balance calculations and
542 discussions and to E. Eliani-Russak for her technical assistance. Many thanks to K. Hachmann from M.
543 Elvert's lab for his help during lipid analysis and to J. Gropp for insightful discussions on the back flux.
544 This work was supported by the ERC consolidator grant (818450) and the Israel Science Foundation
545 (857-2016) of O. Sivan. Funding for M. Elvert was provided by the Deutsche Forschungsgemeinschaft
546 (DFG) (49926684) and EXC 2077 (390741601). Funding for M. Rubin-Blum was provided by the Israel
547 Science Foundation (913/19), the U.S.-Israel Binational Science Foundation (2019055) and Ministry of
548 Science and Technology (1126), and H. Vigderovich was supported by the student fellowship of the
549 Israeli water authority.

550

551 **References**

- 552 Adler, Michal, Eckert, W., & Sivan, O. (2011). Quantifying rates of methanogenesis and methanotrophy in Lake
553 Kinneret sediments (Israel) using porewater profiles. *Limnology and Oceanography*, *56*(4), 1525–1535.
554 <https://doi.org/10.4319/lo.2011.56.4.1525>
- 555 Aepfler, R. F., Bühring, S. I., & Elvert, M. (2019). Substrate characteristic bacterial fatty acid production based
556 on amino acid assimilation and transformation in marine sediments. *FEMS Microbiology Ecology*, *95*(10),
557 1–15. <https://doi.org/10.1093/femsec/fiz131>
- 558 Amiel, N. (2018). *Authigenic magnetite in deep sediments*. MsC thesis, Ben Gurion University of the Negev.
- 559 Aromokeye, D. A., Kulkarni, A. C., Elvert, M., Wegener, G., Henkel, S., Coffinet, S., Eickhorst, T., Oni, O. E.,
560 Richter-Heitmann, T., Schnakenberg, A., Taubner, H., Wunder, L., Yin, X., Zhu, Q., Hinrichs, K. U.,
561 Kasten, S., & Friedrich, M. W. (2020). Rates and Microbial Players of Iron-Driven Anaerobic Oxidation
562 of Methane in Methanic Marine Sediments. *Frontiers in Microbiology*, *10*(January), 1–19.
563 <https://doi.org/10.3389/fmicb.2019.03041>
- 564 Arshad, A., Speth, D. R., De Graaf, R. M., Op den Camp, H. J. M., Jetten, M. S. M., & Welte, C. U. (2015). A
565 metagenomics-based metabolic model of nitrate-dependent anaerobic oxidation of methane by
566 *Methanoperedens*-like archaea. *Frontiers in Microbiology*, *6*(DEC), 1–14.
567 <https://doi.org/10.3389/fmicb.2015.01423>
- 568 Bai, Y. N., Wang, X. N., Wu, J., Lu, Y. Z., Fu, L., Zhang, F., Lau, T. C., & Zeng, R. J. (2019). Humic substances
569 as electron acceptors for anaerobic oxidation of methane driven by ANME-2d. *Water Research*, *164*,
570 114935. <https://doi.org/10.1016/j.watres.2019.114935>
- 571 Bankevich, A., Nurk, S., Antipov, D., Gurevich, A. a., Dvorkin, M., Kulikov, A. S., Lesin, V. M., Nicolenko, S.
572 I., Pham, S., Pribelski, A. D., Sirotkin, A. V., Vyahhi, N., Tesler, G., Aleksyev, A. M., & Pevzner, P. a.
573 (2012). SPAdes: A New Genome Assembly Algorithm and Its Applications to Single-Cell Sequencing.
574 *Journal of Computational Biology*, *19*(5), 455–477. <https://doi.org/10.1089/cmb.2012.0021>
- 575 Bar-Or, I., Ben-Dov, E., Kushmaro, A., Eckert, W., & Sivan, O. (2015). *Methane-related changes in*
576 *prokaryotes along geochemical profiles in sediments of Lake Kinneret (Israel) Methane-related changes*
577 *in prokaryotes along geochemical profiles in sediments of Lake Kinneret (Israel)*. (August).
578 <https://doi.org/10.5194/bg-12-2847-2015>
- 579 Bar-Or, I., Elvert, M., Eckert, W., Kushmaro, A., Vigderovich, H., Zhu, Q., Ben-Dov, E., & Sivan, O. (2017).
580 Iron-Coupled Anaerobic Oxidation of Methane Performed by a Mixed Bacterial-Archaeal Community
581 Based on Poorly Reactive Minerals. *Environmental Science & Technology*, *51*, 12293–12301.
582 <https://doi.org/10.1021/acs.est.7b03126>
- 583 Beal, E. J., House, C. H., & Orphan, V. J. (2009). Manganese-and Iron-Dependent Marine Methane Oxidation.
584 *Science (New York, N.Y.)*, *325*(5937), 184–187. <https://doi.org/10.1126/science.1169984>
- 585 Beck, D. A. C., Kalyuzhnaya, M. G., Malfatti, S., Tringe, S. G., del Rio, T. G., Ivanova, N., Lidstrom, M. E., &

586 Chistoserdova, L. (2013). A metagenomic insight into freshwater methane-utilizing communities and
587 evidence for cooperation between the Methylococcaceae and the Methylophilaceae. *PeerJ*, 2013(1), 1–23.
588 <https://doi.org/10.7717/peerj.23>

589 Biderre-Petit, C., Dugat-Bony, E., Mege, M., Parisot, N., Adrian, L., Moné, A., Denonfoux, J., Peyretailade, E.,
590 Debroas, D., Boucher, D., Peyret, P. (2016). Distribution of Dehalococcoidia in the anaerobic deep water
591 of a remote meromictic crater lake and detection of Dehalococcoidia-derived reductive dehalogenase
592 homologous genes. *PLoS ONE*, 11(1), 1–19. <https://doi.org/10.1371/journal.pone.0145558>

593 Boetius, A., Ravensschlag, K., Schubert, C. J., Rickert, D., Widdel, F., Gieseke, A., Amann, R., Jørgensen, B.B.,
594 Witte, U., & Pfannkuche, O. (2000). A marine microbial consortium apparently mediating AOM. *Nature*,
595 407(October), 623–626.

596 Bottrell, S. H., Parkes, R. J., Cragg, B. A., & Raiswell, R. (2000): Isotopic evidence for anoxic pyrite oxidation
597 and stimulation of bacterial sulphate reduction in marine sediments, *J. Geol. Soc. London*, 157, 711–714.
598 <https://doi.org/10.1144/jgs.157.4.711>.

599 Cabrol, L., Thalasso, F., Gandois, L., Sepulveda-Jauregui, A., Martinez-Cruz, K., Teisserenc, R., Tananaev, N.,
600 Tveit, A., Svenning, M. M., & Barret, M. (2020). Anaerobic oxidation of methane and associated
601 microbiome in anoxic water of Northwestern Siberian lakes. *Science of the Total Environment*, 736,
602 139588. <https://doi.org/10.1016/j.scitotenv.2020.139588>

603 Cheng, L., Shi, S. bao, Yang, L., Zhang, Y., Dolfing, J., Sun, Y. ge, Liu, L., Li, Q., Tu, B., Dai, L., Shi, Q., &
604 Zhang, H. (2019). Preferential degradation of long-chain alkyl substituted hydrocarbons in heavy oil under
605 methanogenic conditions. *Organic Geochemistry*, 138. <https://doi.org/10.1016/j.orggeochem.2019.103927>

606 Chuang, P. C., Yang, T. F., Wallmann, K., Matsumoto, R., Hu, C. Y., Chen, H. W., Lin, S., Sun, CH., Li, HC.,
607 Wang, Y., & Dale, A. W. (2019). Carbon isotope exchange during anaerobic oxidation of methane (AOM)
608 in sediments of the northeastern South China Sea. *Geochimica et Cosmochimica Acta*, 246, 138–155.
609 <https://doi.org/10.1016/j.gca.2018.11.003>

610 Conrad, R. (2009). The global methane cycle: Recent advances in understanding the microbial processes
611 involved. *Environmental Microbiology Reports*, 1(5), 285–292. <https://doi.org/10.1111/j.1758-2229.2009.00038.x>

613 Dershwitz, P., Bandow, N. L., Yang, J., Semrau, J. D., McEllistrem, M. T., Heinze, R. A., Fonseca, M.,
614 Ledesma, J. C., Jennett, J. R., DiSpirito, A. M., Athwal, N. S., Hargrove, M. S., Bobik, T. A., Zischka, H.,
615 & DiSpirito, A. A. (2021). Oxygen Generation via Water Splitting by a Novel Biogenic Metal Ion-
616 Binding Compound. *Applied and Environmental Microbiology*, 87(14), 1–14.
617 <https://doi.org/10.1128/aem.00286-21>

618 Eckert, T. (2000). The Influence of Chemical Stratification in the Water Column on Sulfur and Iron Dynamics
619 in Pore Waters and Sediments of Lake Kinneret, Israel. *M.Sc. Thesis*, University of Bayreuth, Germany.

620 Egger, M., Rasigraf, O., Sapart, C. J., Jilbert, T., Jetten, M. S. M., Röckmann, T., van der Veen, C., Bändä, N.,
621 Kartal, B., Ettwig, K. F., & Slomp, C. P. (2015). Iron-mediated anaerobic oxidation of methane in

- 622 brackish coastal sediments. *Environmental Science and Technology*, 49(1), 277–283.
 623 <https://doi.org/10.1021/es503663z>
- 624 Elul, M., Rubin-Blum, M., Ronen, Z., Bar-Or, I., Eckert, W., & Sivan, O. (2021). Metagenomic insights into the
 625 metabolism of microbial communities that mediate iron and methane cycling in Lake Kinneret sediments.
 626 *Biogeosciences Discussions*, 1–24. <https://doi.org/10.5194/bg-2020-329>
- 627 Elvert, M., Boetius, A., Knittel, K., & Jørgensen, B. B. (2003). Characterization of specific membrane fatty
 628 acids as chemotaxonomic markers for sulfate-reducing bacteria involved in anaerobic oxidation of
 629 methane. *Geomicrobiology Journal*, 20(4), 403–419. <https://doi.org/10.1080/01490450303894>
- 630 Ettwig, Katharina F, Butler, M. K., Le Paslier, D., Pelletier, E., Mangenot, S., Kuypers, M. M. M., Schreiber, F.,
 631 Dutilh, B. E., Zedelius, J., de Beer, D. Gloerich, J., Wessels, H. J. C. T., van Alen, T., Luesken, F., Wu,
 632 M. L., van de Pas-Schoonen K. T., Op den Camp, H. J. M., Jansen-Megens, E. M., Francoijs, KJ.,
 633 Stunnenberg, H., Weissenbach, J., Jetten, M. S. M., & Strous, M. (2010). Nitrite-driven anaerobic
 634 methane oxidation by oxygenic bacteria. *Nature*, 464(7288), 543–548.
 635 <https://doi.org/10.1038/nature08883>
- 636 Evans, P. N., Parks, D. H., Chadwick, G. L., Robbins, S. J., Orphan V. J., Golding, S. D., & Tyson, G. W.
 637 (2015). *Science*. 350(6259), 434–438. <http://doi.org/10.1126/science.aac7745>.
- 638
 639 Fan, L., Dippold, M. A., Ge, T., Wu, J., Thiel, V., Kuzyakov, Y., & Dorodnikov, M. (2020). Anaerobic
 640 oxidation of methane in paddy soil: Role of electron acceptors and fertilization in mitigating CH₄ fluxes.
 641 *Soil Biology and Biochemistry*, 141, 107685. <https://doi.org/10.1016/j.soilbio.2019.107685>
- 642 Gropp, J., Iron, M. A., & Halevy, I. (2021). Theoretical estimates of equilibrium carbon and hydrogen isotope
 643 effects in microbial methane production and anaerobic oxidation of methane. *Geochimica et*
 644 *Cosmochimica Acta*, 295, 237–264. <https://doi.org/10.1016/j.gca.2020.10.018>
- 645 Hadas, O., & Pinkas, R. (1995). Sulphate reduction in the hypolimnion and sediments of Lake Kinneret , Israel.
 646 *Freshwater Biology*, (33), 63–72.
- 647 Hallam, S. J., Putnam, N., Preston, C. M., Detter, J. C., Rokhsar, D., Richardson, P. H., & DeLong, E. F. (2004).
 648 Reverse methanogenesis: Testing the hypothesis with environmental genomics. *Science*, 305(5689),
 649 1457–1462. <https://doi.org/10.1126/science.1100025>
- 650 Hammer, Ø., Harper, D. A. T., & Ryan, P. D. (2001) Past: paleontological statistics software package for
 651 education and data analysis. *Paleontologia-Electronica*. 4 (1), 9.
 652
- 653 Haroon, M. F., Hu, S., Shi, Y., Imelfort, M., Keller, J., Hugenholtz, P., Yuan, Z., & Tyson, G. W. (2013).
 654 Anaerobic oxidation of methane coupled to nitrate reduction in a novel archaeal lineage. *Nature*,
 655 500(7464), 567–570. <https://doi.org/10.1038/nature12375>
- 656 Hoehler, T. M., Alperin, M. J., Albert, D. B., & Martens, C. S. (1994). Field and laboratory, evidence for a
 657 methane-sulfate reducer consortium.pdf. *Global Biogeochemical Cycles*, 8(4), 451–463.
- 658 Holler, T., Wegener, G., Niemann, H., Deusner, C., Ferdelman, T. G., Boetius, A., Brunner, B., & Widdel, F.
 659 (2011). Carbon and sulfur back flux during anaerobic microbial oxidation of methane and coupled sulfate

- 660 reduction. *Proceedings of the National Academy of Sciences of the United States of America*, 108(52).
661 <https://doi.org/10.1073/pnas.1106032108>
- 662 Holmkvist, L., Ferdelman, T. G., & Jørgensen, B. B. (2011). A cryptic sulfur cycle driven by iron in the
663 methane zone of marine sediment (Aarhus Bay, Denmark). *Geochimica et Cosmochimica Acta*, 75(12),
664 3581–3599. <https://doi.org/10.1016/j.gca.2011.03.033>
- 665 Hug, L. A., Castelle, C. J., Wrighton, K. C., Thomas, B. C., Sharon, I., Frischkorn, K. R., Williams, K. H.,
666 Tringe, S. G., & Banfield, J. F. (2013). Community genomic analyses constrain the distribution of
667 metabolic traits across the Chloroflexi phylum and indicate roles in sediment carbon cycling. *Microbiome*,
668 1(1), 1–17. <https://doi.org/10.1186/2049-2618-1-22>
- 669 Kang, D. D., Li, F., Kirton, E., Thomas, A., Egan, R., An, H., & Wang, Z. (2019). MetaBAT 2: An adaptive
670 binning algorithm for robust and efficient genome reconstruction from metagenome assemblies. *PeerJ*,
671 2019(7), 1–13. <https://doi.org/10.7717/peerj.7359>
- 672 Kellermann, M. Y., Wegener, G., Elvert, M., Yoshinaga, M. Y., Lin, Y. S., Holler, T., Mollar, P. X., Knittel K.,
673 & Hinrichs, K. U. (2012). Autotrophy as a predominant mode of carbon fixation in anaerobic methane-
674 oxidizing microbial communities. *Proceedings of the National Academy of Sciences of the USA* 109(47),
675 19321-19326. doi:10.1073/pnas.1208795109.
- 676 Kits, K. D., Klotz, M. G., & Stein, L. Y. (2015). Methane oxidation coupled to nitrate reduction under hypoxia
677 by the Gammaproteobacterium *Methylomonas denitrificans*, sp. nov. type strain FJG1. *Environmental*
678 *Microbiology*, 17(9), 3219–3232. <https://doi.org/10.1111/1462-2920.12772>
- 679 Knittel, K., & Boetius, A. (2009). Anaerobic oxidation of methane: Progress with an unknown process. *Annual*
680 *Review of Microbiology*, 63, 311–334. <https://doi.org/10.1146/annurev.micro.61.080706.093130>
- 681 Kurth, J.M., Nadine T Smit, Stefanie Berger, Stefan Schouten, Mike S M Jetten, Cornelia U Welte, Anaerobic
682 methanotrophic archaea of the ANME-2d clade feature lipid composition that differs from other ANME
683 archaea, *FEMS Microbiology Ecology*, Volume 95, Issue 7, July 2019, fiz082.
- 684 Li, X., Hou, L., Liu, M., Zheng, Y., Yin, G., Lin, X., Cheng, L., Li, Y., & Hu, X. (2015). Evidence of Nitrogen
685 Loss from Anaerobic Ammonium Oxidation Coupled with Ferric Iron Reduction in an Intertidal Wetland.
686 *Environmental Science and Technology*, 49(19), 11560–11568. <https://doi.org/10.1021/acs.est.5b03419>
- 687 Lin, Y. S., Lipp, J. S., Yoshinaga, M. Y., Lin, S. H., Elvert, M., & Hinrichs, K. U. (2010). Intramolecular stable
688 carbon isotopic analysis of archaeal glycosyl tetraether lipids. *Rapid Communications in Mass*
689 *Spectrometry*, 24(19), 2817–2826. <https://doi.org/10.1002/rcm.4707>
- 690 Lovley, D. R., & Klug, M. J. (1983). Sulfate reducers can outcompete methanogens at freshwater sulfate
691 concentrations. *Applied and Environmental Microbiology*, 45(1), 187–192.
692 <https://doi.org/10.1128/aem.45.1.187-192.1983>
- 693 Luo, J. H., Chen, H., Hu, S., Cai, C., Yuan, Z., & Guo, J. (2018). Microbial Selenate Reduction Driven by a
694 Denitrifying Anaerobic Methane Oxidation Biofilm. *Environmental Science and Technology*, 52(7),

695 4006–4012. <https://doi.org/10.1021/acs.est.7b05046>

696 Martinez-cruz, K., Leewis, M., Charold, I., Sepulveda-jauregui, A., Walter, K., Thalasso, F., & Beth, M. (2017).
697 Science of the Total Environment Anaerobic oxidation of methane by aerobic methanotrophs in sub-
698 Arctic lake sediments. *Science of the Total Environment*, 607–608, 23–31.
699 <https://doi.org/10.1016/j.scitotenv.2017.06.187>

700 Meador, T. B., Gagen, E. J., Loscar, M. E., Goldhammer, T., Yoshinaga, M. Y., Wendt, J., Thomm, M., &
701 Hinrichs, K. U. (2014). Thermococcus kodakarensis modulates its polar membrane lipids and elemental
702 composition according to growth stage and phosphate availability. *Frontiers in Microbiology*, 5(JAN), 1–
703 13. <https://doi.org/10.3389/fmicb.2014.00010>

704 Meister, P., Liu, B., Khalili, A., Böttcher, M. E., & Jørgensen, B. B. (2019). Factors controlling the carbon
705 isotope composition of dissolved inorganic carbon and methane in marine porewater: An evaluation by
706 reaction-transport modelling. *Journal of Marine Systems*, 200(August), 103227.
707 <https://doi.org/10.1016/j.jmarsys.2019.103227>

708 Moran, J. J., House, C. H., Freeman, K. H., & Ferry, J. G. (2005). Trace methane oxidation studied in several
709 Euryarchaeota under diverse conditions. *Archaea*, 1(5), 303–309. <https://doi.org/10.1155/2005/650670>

710 Mosrovaya, A., Wind-Hansen, M., Rousteau, P., Bristow, L. A., & Thamdrup, B. (2021) Sulfate- and iron-
711 dependent anaerobic methane oxidation occurring side-by-side in freshwater lake sediments. *Limnology
712 and Oceanography*. <https://doi.org/10.1002/lno.11988>

713 Nollet, L., Demeyer, D., & Verstraete, W. (1997). Effect of 2-bromoethanesulfonic acid and Peptostreptococcus
714 productus ATCC 35244 addition on stimulation of reductive acetogenesis in the ruminal ecosystem by
715 selective inhibition of methanogenesis. *Applied and Environmental Microbiology*, 63(1), 194–200.
716 <https://doi.org/10.1128/aem.63.1.194-200.1997>

717 Nurk, S., Bankevich, A., & Antipov, D. (2013). Assembling genomes and mini-metagenomes from highly
718 chimeric reads. *Research in Computational Molecular Biology*, 158–170. [https://doi.org/10.1007/978-3-
719 642-37195-0](https://doi.org/10.1007/978-3-642-37195-0)

720 Nüsslein, B., Chin, K. J., Eckert, W., & Conrad, R. (2001). Evidence for anaerobic syntrophic acetate oxidation
721 during methane production in the profundal sediment of subtropical Lake Kinneret (Israel). *Environmental
722 Microbiology*, 3(7), 460–470. <https://doi.org/10.1046/j.1462-2920.2001.00215.x>

723 Orembland, R. S., & Capone, D. G. (1988). *Use of "Specific" Inhibitors in Biogeochemistry and Microbial
724 Ecology* (Vol. 10). <https://doi.org/10.2307/4514>

725 Orphan, V. J., House, C. H., & Hinrichs, K. (2001). Methane-Consuming Archaea Revealed by Directly
726 Coupled Isotopic and Phylogenetic Analysis. *Science*, 293(July), 484–488.
727 <https://doi.org/10.1126/science.1061338>

728 Oswald, K., Milucka, J., Brand, A., Hach, P., Littmann, S., Wehrli, B., Albersten, M., Daims, H., Wagner, M.,
729 Kuypers, M. M. M., Schubert, C. J., & Milucka, J. (2016). Aerobic gammaproteobacterial methanotrophs

- 730 mitigate methane emissions from oxic and anoxic lake waters. *Limnology and Oceanography*, 61, S101–
731 S118. <https://doi.org/10.1002/lno.10312>
- 732 Parks, D. H., Chuvochina, M., Rinke, C., Mussig, A. J., Chaumeil, P.-A., & Hugenholtz, P. (2021) GTDB: an
733 ongoing census of bacterial and archaeal diversity through a phylogenetically consistent, rank
734 normalized and complete genome-based taxonomy. *Nucleic Acids Research*, 202, 1-10.
735 <http://doi.org/10.1093/nar/gkab776>
- 736 Raghoebarsing, A. A., Pol, A., Van De Pas-Schoonen, K. T., Smolders, A. J. P., Ettwig, K. F., Rijpstra, W. I. C.,
737 Schouten, S., Sinninghe Damsté, J. S., Op den Camp, H. J. M., Jetten, M. S. M., & Strous, M. (2006). A
738 microbial consortium couples anaerobic methane oxidation to denitrification. *Nature*, 440(7086), 918–
739 921. <https://doi.org/10.1038/nature04617>
- 740 Reeburgh, W. S. (2007). Oceanic Methane Biogeochemistry. *ChemInform*, 38(20), 486–513.
741 <https://doi.org/10.1002/chin.200720267>
- 742 Saunio, M., Stavert, A. R., Poulter, B., Bousquet, P., Canadell, J. G., Jackson, R. B., Raymond, P. A.,
743 Dlugokencky, E. J., Houweling, S., Patra, P. K., Ciais, P., Arora, V. K., Bastviken, D., Bergamaschi, P.,
744 Blake, D. R., Brailsford, G., Bruhwiler, L., Carlson, K. M., Carrol, M., Castaldi, S., Chandra, N.,
745 Crevoisier, C., Crill, P. M., Covey, K., Curry, C. L., Etiope, G., Frankenberg, C., Gedney, N., Hegglin, M.
746 I., Höglund-Isaksson, L., Hugelius, G., Ishizawa, M., Ito, A., Janssens-Maenhout, G., Jensen, K. M., Joos,
747 F., Kleinen, T., Krummel, P. B., Langenfelds, R. L., Laruelle, G. G., Liu, L., Machida, T., Maksyutov, S.,
748 McDonald, K. C., McNorton, J., Miller, P. A., Melton, J. R., Morino, I., Müller, J., Murguía-Flores, F.,
749 Naik, V., Niwa, Y., Noce, S., O'Doherty, S., Parker, R. J., Peng, C., Peng, S., Peters, G. P., Prigent, C.,
750 Prinn, R., Ramonet, M., Regnier, P., Riley, W. J., Rosentreter, J. A., Segers, A., Simpson, I. J., Shi, H.,
751 Smith, S. J., Steele, L. P., Thornton, B. F., Tian, H., Tohjima, Y., Tubiello, F. N., Tsuruta, A., Viovy, N.,
752 Voulgarakis, A., Weber, T. S., van Weele, M., van der Werf, G. R., Weiss, R. F., Worthy, D., Wunch, D.,
753 Yin, Y., Yoshida, Y., Zhang, W., Zhang, Z., Zhao, Y., Zheng, B., Zhu, Q., Zhu, Q., and Zhuang, Q.: The
754 Global Methane Budget 2000–2017, *Earth Syst. Sci. Data*, 12, 1561–1623, [https://doi.org/10.5194/essd-](https://doi.org/10.5194/essd-12-1561-2020)
755 [12-1561-2020](https://doi.org/10.5194/essd-12-1561-2020), 2020.
- 756 Scheller, S., Yu, H., Chadwick, G. L., McGlynn, S. E., & Orphan, V. J. (2016). Artificial electron acceptors
757 decouple archaeal methane oxidation from sulfate reduction. *Science*, 351(6274), 1754–1756.
758 <https://doi.org/10.1126/science.aad7154>
- 759 Segarra, K. E. a, Comerford, C., Slaughter, J., & Joye, S. B. (2013). Impact of electron acceptor availability on
760 the anaerobic oxidation of methane in coastal freshwater and brackish wetland sediments. *Geochimica et*
761 *Cosmochimica Acta*, 115, 15–30. <https://doi.org/10.1016/j.gca.2013.03.029>
- 762 Sela-Adler, M., Herut, B., Bar-Or, I., Antler, G., Eliani-Russak, E., Levy, E., Makovsky, Y., & Sivan, O.
763 (2015). Geochemical evidence for biogenic methane production and consumption in the shallow
764 sediments of the SE Mediterranean shelf (Israel). *Continental Shelf Research*, 101, 117–124.
765 <https://doi.org/10.1016/j.csr.2015.04.001>
- 766 Serruya, C. (1971). Lake Kinneret: the nutrient chemistry of the Sediments. *Limnology and Oceanography*,

767 16(May), 510–521.

768 Shuai, W., & Jaffé, P. R. (2019). Anaerobic ammonium oxidation coupled to iron reduction in constructed
769 wetland mesocosms. *Science of the Total Environment*, 648, 984–992.
770 <https://doi.org/10.1016/j.scitotenv.2018.08.189>

771 Sieber, C. M. K., Probst, A. J., Sharrar, A., Thomas, B. C., Hess, M., Tringe, S. G., & Banfield, J. F. (2018).
772 Recovery of genomes from metagenomes via a dereplication, aggregation and scoring strategy. *Nature*
773 *Microbiology*, 3(7), 836–843. <https://doi.org/10.1038/s41564-018-0171-1>

774 Sivan, O., Adler, M., Pearson, A., Gelman, F., Bar-Or, I., John, S. G., & Eckert, W. (2011). Geochemical
775 evidence for iron-mediated anaerobic oxidation of methane. *Limnology and Oceanography*, 56(4), 1536–
776 1544.

777 Sivan, O., Antler, G., Turchyn, A. V., Marlow, J. J., & Orphan, V. J., (2014). Iron oxides stimulate sulfate-
778 driven anaerobic methane oxidation in seeps. PNAS. 111, E4139-E4147.
779 <http://doi.org/10.1073/pnas.1412269111>

780 Sivan, O., Shusta, S., & Valentine, D. L. (2016). Methanogens rapidly transition from methane production to
781 iron reduction. *Geobiology*, 190–203. <https://doi.org/10.1111/gbi.12172>

782 Stookey, L. L. (1970). Ferrozine-a new spectrophotometric reagent for iron. *Analytical Chemistry*, 42(7), 779–
783 781. <https://doi.org/10.1021/ac60289a016>

784 Sturt, H. F., Summons, R. E., Smith, K., Elvert, M., & Hinrichs, K. U. (2004). Intact polar membrane lipids in
785 prokaryotes and sediments deciphered by high-performance liquid chromatography/electrospray
786 ionization multistage mass spectrometry - New biomarkers for biogeochemistry and microbial ecology.
787 *Rapid Communications in Mass Spectrometry*, 18(6), 617–628. <https://doi.org/10.1002/rem.1378>

788 Su, G., Zopfi, J., Yao, H., Steinle, L., Niemann, H., & Lehmann, M. F. (2020). Manganese/iron-supported
789 sulfate-dependent anaerobic oxidation of methane by archaea in lake sediments. *Limnology and*
790 *Oceanography*, 65(4), 863–875. <https://doi.org/10.1002/lno.11354>

791 Tamames, J., & Puente-Sánchez, F. (2019). SqueezeMeta, A Highly Portable, Fully Automatic Metagenomic
792 Analysis Pipeline. *Frontiers in Microbiology*, 9. <https://doi.org/10.3389/fmicb.2018.03349>

793 Timmers, P. H. A., Welte, C. U., Koehorst, J. J., Plugge, C. M., Jetten, M. S. M., & Stams, A. J. M. (2017).
794 Reverse Methanogenesis and Respiration in Methanotrophic Archaea. *Archaea*, 2017(Figure 1).
795 <https://doi.org/10.1155/2017/1654237>

796 Treude, T., Krause, S., Maltby, J., Dale, A. W., Coffin, R., & Hamdan, L. J. (2014). Sulfate reduction and
797 methane oxidation activity below the sulfate-methane transition zone in Alaskan Beaufort Sea continental
798 margin sediments: Implications for deep sulfur cycling. *Geochimica et Cosmochimica Acta*, 144, 217–
799 237. <https://doi.org/10.1016/j.gca.2014.08.018>

800 Treude, T., Niggemann, J., Kallmeyer, J., Wintersteller, P., Schubert, C. J., Boetius, A., & Jørgensen, B. B.
801 (2005). Anaerobic oxidation of methane and sulfate reduction along the Chilean continental margin.

802 *Geochimica et Cosmochimica Acta*, 69(11), 2767–2779. <https://doi.org/10.1016/j.gca.2005.01.002>

803 Valenzuela, E. I., Avendaño, K. A., Balagurusamy, N., Arriaga, S., Nieto-Delgado, C., Thalasso, F., &
804 Cervantes, F. J. (2019). Electron shuttling mediated by humic substances fuels anaerobic methane
805 oxidation and carbon burial in wetland sediments. *Science of the Total Environment*, 650, 2674–2684.
806 <https://doi.org/10.1016/j.scitotenv.2018.09.388>

807 Valenzuela, E. I., Prieto-Davó, A., López-Lozano, N. E., Hernández-Eligio, A., Vega-Alvarado, L., Juárez, K.,
808 García-González, A. S., López, M. G., & Cervantes, F. J. (2017). Anaerobic methane oxidation driven by
809 microbial reduction of natural organic matter in a tropical wetland. *Applied and Environmental*
810 *Microbiology*, 83(11), 1–15. <https://doi.org/10.1128/AEM.00645-17>

811 Vigderovich, H., Liang, L., Herut, B., Wang, F., Wurgaft, E., Rubin-Blum, M., & Sivan, O. (2019). Evidence
812 for microbial iron reduction in the methanogenic sediments of the oligotrophic SE Mediterranean
813 continental shelf. *Biogeosciences Discussions*, 1–25. <https://doi.org/10.5194/bg-2019-21>

814 Wang, L., Miao, X., Ali, J., Lyu, T., & Pan, G. (2018). Quantification of Oxygen Nanobubbles in Particulate
815 Matters and Potential Applications in Remediation of Anaerobic Environment. *ACS Omega*, 3(9), 10624–
816 10630. <https://doi.org/10.1021/acsomega.8b00784>

817 Wegener G, Niemann H, Elvert M, Hinrichs K-U, Boetius A (2008). Assimilation of methane and inorganic
818 carbon by microbial communities mediating the anaerobic oxidation of methane. *Environmental*
819 *Microbiology* 10(9), 2287-2298. doi: 10.1111/j.1462-2920.2008.01653.x.

820 Wegener, G., Gropp, J., Taubner, H., Halevy, I., & Elvert, M. (2021). Sulfate-dependent reversibility of
821 intracellular reactions explains the opposing isotope effects in the anaerobic oxidation of methane. *Science*
822 *Advances*, 7(19), 1–14. <https://doi.org/10.1126/sciadv.abe4939>

823 Wu, Y.W., Tang, Y.-H., Tringe, S. G., Simmons, B. A., & Singer, S. W. (2014). MaxBin: an automated binning
824 method to recover individual genomes from metagenomes using. *Microbiome*, 2(26), 4904–4909.
825 Retrieved from <https://microbiomejournal.biomedcentral.com/articles/10.1186/2049-2618-2-26>

826 Wuebbles, D. J., & Hayhoe, K. (2002). Atmospheric methane and global change. *Earth-Science Reviews*, 57(3–
827 4), 177–210. [https://doi.org/10.1016/S0012-8252\(01\)00062-9](https://doi.org/10.1016/S0012-8252(01)00062-9)

828 Yorshansky, O. (2019). *Iron Reduction in Deep Marine Sediments of the Eastern Mediterranean Continental*
829 *Shelf and the Yaqon Estuary*. MsC thesis, Ben Gurion University of the Negev.

830 Yoshinaga, M. Y., Holler, T., Goldhammer, T., Wegener, G., Pohlman, J. W., Brunner, B., Kuypers, M. M. M.,
831 Hinrichs, K. U., & Elvert, M. (2014). Carbon isotope equilibration during sulphate-limited anaerobic
832 oxidation of methane. *Nature Geoscience*, 7(3), 190–194. <https://doi.org/10.1038/ngeo2069>

833 Zehnder, a J., & Brock, T. D. (1979). Methane formation and methane oxidation by methanogenic bacteria.
834 *Journal of Bacteriology*, 137(1), 420–432.

835 Zhang, X., Xia, J., Pu, J., Cai, C., Tyson, G. W., Yuan, Z., & Hu, S. (2019). Biochar-Mediated Anaerobic
836 Oxidation of Methane. *Environmental Science and Technology*, 53(12), 6660–6668.

837 <https://doi.org/10.1021/acs.est.9b01345>

838 Zheng, Y., Wang, H., Liu, Y., Zhu, B., Li, J., Yang, Y., Qin, W., Chen, L., Wu, X., Chistoserdova, L., & Zhao,
839 F. (2020). Methane-Dependent Mineral Reduction by Aerobic Methanotrophs under Hypoxia.

840 *Environmental Science and Technology Letters*, 7(8), 606–612. <https://doi.org/10.1021/acs.estlett.0c00436>

841