1 Complete perchlorate reduction using methane as the sole

2 electron donor and carbon source

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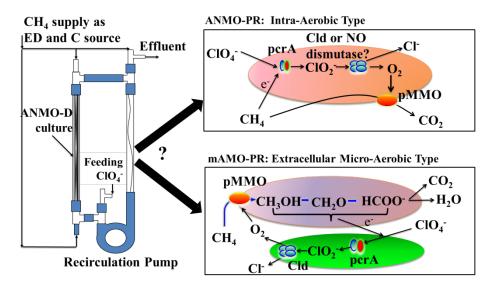
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14 Abstract

15	Using a CH ₄ -based membrane biofilm reactor (MBfR), we studied perchlorate (ClO ₄ ⁻)
16	reduction by a biofilm performing anaerobic methane oxidation coupled to
17	denitrification (ANMO-D). We focused on the effects of nitrate (NO_3^-) and nitrite
18	(NO_2^-) surface loadings on ClO_4^- reduction and on the biofilm community's
19	mechanism for ClO_4^- reduction. The ANMO-D biofilm reduced up to 5 mg/L of
20	ClO_4^- to a non-detectable level using CH_4 as the only electron donor and carbon
21	source when CH_4 delivery was not limiting; NO_3^- was completely reduced as well
22	when its surface loading was ≤ 0.32 g N/m ² -d. When CH ₄ delivery was limiting,
23	NO_3^- inhibited ClO_4^- reduction by competing for the scarce electron donor. NO_2^-
24	inhibited ClO ₄ ⁻ reduction when its surface loading was ≥ 0.10 g N/m ² -d, probably due
25	to cellular toxicity. Although Archaea were present through all stages, Bacteria
26	dominated the ClO ₄ -reducing ANMO-D biofilm, and gene copies of the particulate
27	methane mono-oxygenase ($pMMO$) correlated to the increase of respiratory gene
28	copies. These pieces of evidence support that ClO_4^- reduction by the MBfR biofilm
29	involved chlorite (ClO_2^{-}) dismutation to generate the O_2 needed as a co-substrate for
30	the mono-oxygenation of CH ₄ .
31	Key Words: methane, perchlorate, oxidation, reduction, membrane-biofilm reactor





ANMO-D: Anaerobic methane oxidation coupled to nitrate reduction ANMO-PR: Anaerobic methane oxidation coupled to perchlorate reduction mAMO-PR: micro-Aerobic methane oxidation coupled to perchlorate reduction

35

36 Introduction

Perchlorate (ClO₄) is a strong oxidizing agent that has been widely used in rocket fuel, 37 munitions, and explosives (EPA, 2005).¹ It causes serious health problems by 38 interfering with the production of thyroid hormones needed for growth and 39 development (Coates & Achenbach, 2004).² The typical perchlorate concentration 40 in groundwater is lower than 100 µg/L, but in some cases it can reach concentrations 41 of 20 mg/L or more.^{3,4} Although a nationwide maximum contaminant level (MCL) 42 for ClO_4^- has not yet established by the US EPA, some states have established cleanup 43 levels ranging from 2 to $18 \,\mu \text{g/L}$ for ClO₄⁻ in drinking water (Gu & Coates, 2006).⁵ 44 Nitrate (NO_3^{-}) is an oxyanion commonly co-occurring with ClO_4^{-} in groundwater, for 45 example, at military bases that house rockets (USEPA, 2001).⁶ Because NO_3^- causes 46 methemoglobinemia in infants, the MCL for NO_3^- in drinking water is regulated at 10 47 mg N/L (USEPA, 2009).⁷ NO₃⁻ inhibits ClO_4^- reduction due to competition for the 48 common electron donor when the electron donor is insufficient.^{8,9} 49 Different electron donors have been applied to achieve complete ClO₄⁻ and NO₃⁻ 50 removal by microbiological reduction.¹⁰⁻¹² An interesting example is nitrogen 51 reduction using methane as the sole electron donor and carbon source,^{13, 14} since 52 methane is inexpensive and widely available.¹⁵⁻¹⁷ 53

54	Methane oxidation coupled to denitrification (MO-D) has been extensively studied
55	during the past decade. ^{14, 15, 18-21} Two microbial processes are capable of carrying
56	out MO-D. One is aerobic methane oxidation coupled to denitrification (AMO-D), ^{22,}
57	²³ which is performed by the combined actions of two distinct bacterial groups:
58	methane oxidizers (methanotrophs) and denitrifiers. ^{22, 24, 25} The second is anaerobic

	21.26	
59	methane oxidation coupled to denitrification (ANMO-D). ^{21, 26}	ANMO-D

microorganisms include a bacterial group affiliated with the candidate division NC10
and an archaeal group distantly related to anaerobic methanotrophic archaea.^{19, 27-29}

AMO-D occurs in the presence of O_2 , because methanotrophs require O_2 for the initial mono-oxygenation step. The methanotrophs can release organic intermediates from their catabolism and anabolism, for example, methanol, acetate, and citrate, and the intermediates can be further utilized by denitrifiers as electron donors.^{15, 22, 30} These steps are illustrated schematically in panel A of Figure S1 in Supplemental Information. Although a high concentration of O_2 inhibits denitrification, a certain amount of O_2 is necessary to promote AMO-D.^{30, 31}

69 ANMO-D can follow two pathways, illustrated schematically in panels B and C of Figure S1. Raghoebarsing et al. hypothesized the "Reverse Methanogesis" pathway, 70 which involves the combined action of Archaea and denitrifying bacteria.¹⁹ The 71 72 Archaea carry out reverse methanogenesis to generate H_2 that is shuttled to the denitrifying bacteria that respire NO_2^- to N_2 . Haroon et al. reported that Archaea 73 74 population ANME-2d (Methanoperedens nitroreducens) catalyzed CH₄ oxidation by methylcoenzyme M reductase (mcrABCDG) through a reverse-methanogenesis 75 pathway using NO_3^- as their terminal electron acceptor and generating NO_2^- ; the 76 NC10 bacteria then reduced NO₂⁻ to N_2 .¹⁴ 77

The second ANMO-D pathway is called the "Intra-Aerobic" pathway,^{21, 32} and only
one denitrifying microorganism (*Candidatus Methylomirabilis oxyfera*) was involved.

Denitrification was carried out by stepwise reduction of NO₂⁻ to NO using nitrate reductase (*narGHJI*) and nitrite reductase (*nirSJFD/GH/L*); then, NO was disproportionated to produce O₂ intracellularly and N₂ using an unknown dismutase enzyme. The O₂ was then used by the same microorganism as a co-substrate for methane mono-oxygenation by a membrane-bound particulate methane mono-oxygenase (*pMMO*).

While the true mechanism of ANMO-D is unresolved, it is thermodynamically feasible, as shown by redox equations 1 and 2 for NO_3^- or NO_2^- as the terminal electron acceptor:¹⁹

89
$$5CH_4 + 8NO_3^- + 8H^+ = 5CO_2 + 4N_2 + 14H_2O$$
 $\triangle G^{0'} = -765 \text{ KJmol}^{-1}CH_4$ (1)

90
$$3CH_4 + 8NO_2^- + 8H_7 = 3CO_2 + 4N_2 + 10H_2O$$
 $\triangle G^{0'} = -928 \text{ KJmol}^{-1}CH_4$ (2)

91 When perchlorate (ClO_4) is the electron acceptor, a similar reaction between ClO_4

92 and
$$CH_4$$
 also is thermodynamically feasible: ³³

93
$$CH_4 + ClO_4^- = HCO_3^- + Cl^- + H_2O$$
 $\triangle G^{0'} = -792 \text{ KJmol}^{-1}CH_4$ (3)

The pathway for dissimilatory perchlorate reduction begins with reduction of ClO₄⁻ to 94 chlorite (ClO₂, catalyzed by perchlorate reductase, *pcrA*) and ends with dismutation 95 of ClO_2^- to yield chloride (Cl⁻, catalyzed by chlorite dismutase, *cld*) and molecular 96 oxygen (O_2) , which is essential for the methane oxidation.² Miller et al. confirmed 97 complete CH₄ removal coupled with ClO₂⁻ dismutation by a mixture of methanotrophs 98 and the perchlorate-reducing bacterium *Dechloromonas agitate* CKB.³⁴ The 99 methanotrophs used extracellular O_2 derived from disproportion of ClO_2^- by D. 100 agitate CKB to oxidize CH₄ aerobically; thus, it was an extracellular-aerobic pathway 101

102	that required that the substrate for dismutation, ClO_2^- , be supplied. The mixed
103	culture did not oxidize CH_4 when supplied with ClO_4^- or ClO_3^- ; thus, they concluded
104	that O_2 produced via ClO_4^- reduction was unavailable for the aerobic methanotrophs.
105	So far, no study has successfully reduced ClO_4^- using CH_4 as the sole electron donor
106	and carbon source. However, thermodynamics and the reality that most denitrifiers
107	are able to reduce ClO_4^- (using either nitrate or perchlorate reductase) means that a
108	ANMO-D or AMO-D culture has the possibility to reduce ClO_4^- using CH_4 as electron
109	donor and carbon source.
110	The H ₂ -based membrane biofilm reactor (MBfR) has been applied successfully for
111	microbial removal of oxidized contaminants, including NO_3^- and $ClO_4^{11, 35-37}$ The
112	non-porous walls of hollow-fiber membranes transfer H_2 directly to a biofilm of
113	H_2 -oxidizing bacteria that reduce one or more electron acceptors. ³⁵ The use of
114	"bubbleless" membranes and the rapid oxidation of H_2 in the biofilm allow nearly 100%
115	utilization of H_2 , preventing H_2 losses to the atmosphere or effluent liquid. ^{36, 38}
116	The MBfR also could provide a means for the safe and efficient supply of CH ₄ to
117	drive ANMO-D, AMO-D, and ClO_4^- reduction. Supporting the concept, Sun et al
118	reported that an aerobic methane-based MBfR removed up to 97% of NO ₃ ⁻ applied at
119	a concentration of 30 mg N/L, 39 and Shi et al achieved 86 mg N/m ² -d NO ₃ ⁻ removal in
120	an anaerobic MBfR provided with CH_4 as carbon source and electron donor. ⁴⁰
10.	
121	The objective of this study was to evaluate ClO_4^- reduction in a CH ₄ -based MBfR.

122	Specifically, we studied the reduction patterns of NO_2^- , NO_3^- , and ClO_4^- when we
123	exposed the biofilm to different relative loadings. We quantified the
124	CH ₄ -permeation coefficient through the membrane wall to determine if the delivery
125	rate of CH_4 was limiting, and we used quantitative real-time PCR (qPCR) to monitor
126	how the abundances of functional genes key to respiration reactions were affected by
127	the acceptor loadings. Based on several types of evidence, we are able to provide
128	mechanistic interpretation about what controls ClO_4^- reduction by the biofilm and the
129	likely pathways by which NO_3^- and ClO_4^- reductions occur when CH_4 is the sole
130	electron donor.

131 Materials and Methods

132 *CH*₄ permeation

133	Steady-state CH ₄ -permeation experiments were carried out in the same system (shown
134	schematically in Supporting Information (SI) Figure S2) Tang et al. used for
135	quantifying H_2 permeation. ⁴¹ Deionized water was pumped through the serum bottle
136	(total volume of 1.6×10^{-4} m ³ , liquid volume of 0.6×10^{-4} m ³) at a flow rate of $7.2 \times$
137	10^{-4} m ³ /d. The hollow fibers in the serum bottle were pressurized with CH ₄ (99.99%)
138	purity, Shanghai Gas Company, China) at a pressure of 1.0 bar (14.5 psi). CH_4
139	diffused through the hollow-fiber wall, dissolved in the water, and partitioned into the
140	headspace. A magnetic stirring bar ensured complete mixing of the liquid and that
141	CH_4 was rapidly partitioned to the gas phase. We took the headspace gas samples to
142	measure its CH_4 partial pressure. Steady state was achieved when the CH_4 partial
143	pressure was stable for at least 40 hydraulic retention times (HRTs). ⁴¹ We then
144	calculated the CH ₄ permeability of the membrane fiber based on the method in Tang
145	et al. ⁴¹ The equations and experimental parameters for the CH ₄ -permeation test are
146	summarized in the Supplementary Information (Table S1 & S2).

147 MBfR Setup

We used a two-column MBfR system similar to Zhao et al.⁸ The MBfR had
composite hollow fibers (hydrophobic microporous polyethylene fiber, 280-µm o.d.,
and a 180-µm i.d., pore size 0.1-0.15 µm) manufactured by Mitsubishi Rayon (Model

151	MHF-200TL, Mitsubishi, Ltd., Japan). The fibers were glued into a CH ₄ -supply
152	manifold at the bottom of the MBfR column, and the top of each fiber was sealed.
153	The total volume of the MBfR was 65 mL, and the total membrane surface area was
154	7.0 cm^2 . The MBfR was completely mixed by recirculation with a peristaltic pump
155	(Longer Pump, model 1515X, Longer Precision Pump Co, Ltd, China) at 100
156	mL/min.

157 Start up and continuous operation of the MBfR

158 We inoculated the MBfR with 10 mL of ANMO-D culture (original maintained

anaerobic) donated by Dr. Wei Xiang Wu at Zhejiang University (China) and enriched

160 the community by recirculating a mineral salt medium (described below) containing 2

161 mg N/L NO₂⁻ for 2 days. To accumulate enough biomass, we fed the MBfR with 2

 $162 \text{ mg N/L of NO}_2^-$ continuously for 40 days, when complete NO $_2^-$ reduction was

163 achieved. To investigate ClO_4^- reduction in the presence of NO_2^- , we fed the MBfR

164 with ClO_4^- and NO_2^- at influent concentrations of 1 mg/L and 2 mg N/L, respectively,

165 in Stage 1. Since NO₃⁻ may inhibit ClO_4^- reduction at high surface loadings (> 0.6 g

166 N/m²-d) or improve ClO₄⁻ removal at medium (0.1-0.6 g N/m²-d) or small (<0.1 g

167 N/m^2 -d) loadings in a H₂-based MBfR,⁹ we then systematically changed the influent

168 concentrations of NO_3^- and ClO_4^- in Stages 2 through 6. We allowed each stage to

reach steady state, which was defined as effluent concentrations stable (<10%

170 variation) for a minimum of three HRTs. The influent concentrations were: Stage

171 2: 1 mg/L ClO₄⁻; Stage 3: 1 mg/L ClO₄⁻ and 1.1 mg N/L of NO₃⁻; Stage 4: 1

 NO₃⁻; and Stage 6: 5 mg/L ClO₄⁻. To investigate ClO₄⁻ reduction in the presence of NO₂⁻ after high ClO₄⁻-removal efficiency had been achieved, we fed the MBfR 5 mg/L of ClO₄⁻ and 5 mg N/L of NO₂⁻ in Stage 7. Actual influent concentrations varied slightly from the targets and are presented in Table S3 and Figure 1. The influent feeding rate was 0.5 mL/min (HRT of 130 min), the CH₄ pressure was 10 psi (0.69 bar) for Stage 1-3 and 15 psi (1.03 bar) for the latter stages, and the temperature was 29±1°C for all experiments. The medium pH was adjusted to 7.0±0.2 with hydrochloric acid and contained the following mineral salts (analytical grade or purer) per L of demineralized water: CaCl₂ 1 mg, NaHCO₃ 0.3 g, MgSO₄•7H₂O 5 mg, KaH₂PO₄ 0.2 g, Na₂HPO₄•12H₂O 0.4 g, 1 mL acid trace element solution (HCl 100 mM, 2.085 g of FeSO₄•7H₂O, 68 mg of ZnSO₄•7H₂O, 14 mg of H₃BO₃, 120 mg of CoCl₂•6H₂O, 500 mg of MnCl₂•4H₂O, 320 mg of CuSO₄, 95 mg of NiCl₂•6H₂O per liter), and 1 mL alkaline trace element solution (NaOH 10 mM, 67 mg of SeO₂, 50 mg of Na₂WO₄•2H₂O, 242 mg of Na₂MOO₄•2H₂O per liter). The medium was de-gassed with N₂ to maintain an anaerobic condition. 	172	mg/L ClO ₄ ⁻ and 11.3 mg N/L of NO ₃ ⁻ ; Stage 5: 1 mg/L ClO_4^- and 4.5 mg N/L of
 mg/L of ClO₄ and 5 mg N/L of NO₂ in Stage 7. Actual influent concentrations varied slightly from the targets and are presented in Table S3 and Figure 1. The influent feeding rate was 0.5 mL/min (HRT of 130 min), the CH₄ pressure was 10 psi (0.69 bar) for Stage 1-3 and 15 psi (1.03 bar) for the latter stages, and the temperature was 29±1°C for all experiments. The medium pH was adjusted to 7.0±0.2 with hydrochloric acid and contained the following mineral salts (analytical grade or purer) per L of demineralized water: CaCl₂ 1 mg, NaHCO₃ 0.3 g, MgSO₄•7H₂O 5 mg, KaH₂PO₄ 0.2 g, Na₂HPO₄•12H₂O 0.4 g, 1 mL acid trace element solution (HCl 100 mM, 2.085 g of FeSO₄•7H₂O, 68 mg of ZnSO₄•7H₂O, 14 mg of H₃BO₃, 120 mg of CoCl₂•6H₂O, 500 mg of MnCl₂•4H₂O, 320 mg of CuSO₄, 95 mg of NiCl₂•6H₂O per liter), and 1 mL alkaline trace element solution (NaOH 10 mM, 67 mg of SeO₂, 50 mg of Na₂WO₄•2H₂O, 242 mg of Na₂MOO₄•2H₂O per liter). The medium was de-gassed with N₂ to maintain an anaerobic condition. 	173	NO ₃ ; and Stage 6: 5 mg/L ClO_4 . To investigate ClO ₄ ⁻ reduction in the presence of
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 solution (HCl 100 mM, 2.085 g of FeSO₄•7H₂O, 68 mg of ZnSO₄•7H₂O, 14 mg of H₃BO₃, 120 mg of CoCl₂•6H₂O, 500 mg of MnCl₂•4H₂O, 320 mg of CuSO₄, 95 mg of NiCl₂•6H₂O per liter), and 1 mL alkaline trace element solution (NaOH 10 mM, 67 mg of SeO₂, 50 mg of Na₂WO₄•2H₂O, 242 mg of Na₂MoO₄•2H₂O per liter). The medium was de-gassed with N₂ to maintain an anaerobic condition. During Stage 4, the CH₄ supply was accidently lost for 48 hours, and we immediately 	181	grade or purer) per L of demineralized water: CaCl ₂ 1 mg, NaHCO ₃ 0.3 g,
 H₃BO₃, 120 mg of CoCl₂•6H₂O, 500 mg of MnCl₂•4H₂O, 320 mg of CuSO₄, 95 mg of NiCl₂•6H₂O per liter), and 1 mL alkaline trace element solution (NaOH 10 mM, 67 mg of SeO₂, 50 mg of Na₂WO₄•2H₂O, 242 mg of Na₂MoO₄•2H₂O per liter). The medium was de-gassed with N₂ to maintain an anaerobic condition. During Stage 4, the CH₄ supply was accidently lost for 48 hours, and we immediately 	182	MgSO ₄ •7H ₂ O 5 mg, KaH ₂ PO ₄ 0.2 g, Na ₂ HPO ₄ •12H ₂ O 0.4 g, 1 mL acid trace element
 NiCl₂•6H₂O per liter), and 1 mL alkaline trace element solution (NaOH 10 mM, 67 mg of SeO₂, 50 mg of Na₂WO₄•2H₂O, 242 mg of Na₂MoO₄•2H₂O per liter). The medium was de-gassed with N₂ to maintain an anaerobic condition. During Stage 4, the CH₄ supply was accidently lost for 48 hours, and we immediately 	183	solution (HCl 100 mM, 2.085 g of FeSO ₄ •7H ₂ O, 68 mg of ZnSO ₄ •7H ₂ O, 14 mg of
 mg of SeO₂, 50 mg of Na₂WO₄•2H₂O, 242 mg of Na₂MoO₄•2H₂O per liter). The medium was de-gassed with N₂ to maintain an anaerobic condition. During Stage 4, the CH₄ supply was accidently lost for 48 hours, and we immediately 	184	H ₃ BO ₃ , 120 mg of CoCl ₂ •6H ₂ O, 500 mg of MnCl ₂ •4H ₂ O, 320 mg of CuSO ₄ , 95 mg of
 medium was de-gassed with N₂ to maintain an anaerobic condition. During Stage 4, the CH₄ supply was accidently lost for 48 hours, and we immediately 	185	NiCl ₂ •6H ₂ O per liter), and 1 mL alkaline trace element solution (NaOH 10 mM, 67
188 During Stage 4, the CH ₄ supply was accidently lost for 48 hours, and we immediately	186	mg of SeO ₂ , 50 mg of Na ₂ WO ₄ •2H ₂ O, 242 mg of Na ₂ MoO ₄ •2H ₂ O per liter). The
	187	medium was de-gassed with N_2 to maintain an anaerobic condition.
substituted N_2 gas to keep the fibers pressurized. To re-evaluate the response found	188	During Stage 4, the CH_4 supply was accidently lost for 48 hours, and we immediately
	189	substituted N_2 gas to keep the fibers pressurized. To re-evaluate the response found

- in Stage 4, we intentionally stopped the CH_4 supply for 30 hours in Stage 5. The
- CH_4 supply was reinstated when the removal percentages were zero for NO₃⁻ and
- ClO_4^- .

194	We measured the CH_4 partial pressure (P_{CH4}) of gas samples using a gas
195	chromatograph (Agilent Technologies GC system, model 7890A, Agilent
196	Technologies Inc., U.S.A) equipped with a flame ionization detector and a packed
197	column (30 m long, 0.32 mm i.d., 0.5 μ m thickness, cross-linked polydimethysiloxane
198	film, J&W scientific, U.S.A.). N_2 was the carrier gas fed at a constant pressure of
199	0.96 bar and a constant flow rate of 0.065 m^3/d , and the temperature conditions for
200	injection and detector were 200 and 260°C, respectively. Analytical grade CH_4 was
201	used for standard calibration curves and for the experiments.
202	We took liquid samples from the MBfR with 5-mL syringes and filtered them
203	immediately through a 0.2- μ m membrane filter (LC+PVDF membrane, Shanghai
204	Xinya, China). We assayed for NO_3^- and NO_2^- using ion chromatography (Metrohm
205	833 Basic IC plus, Switzerland) with an A-Supp-5 column, an eluent containing 3.2
206	mM NaHCO ₃ , 1.0 mM Na ₂ CO ₃ , and 5% Acetone in a flow rate of 1 mL/min. ClO ₄ ⁻
207	was measured using ion chromatography (Metrohm 833 Basic IC plus, Switzerland)
208	with an AS 16 column and AG 16 pre-column, eluent concentration of 35 mM KOH,
209	and a 1.5 mL/min flow rate. Dissolved O_2 was measured with a dissolved oxygen
210	probe (Starter, model 300D, Ohaus Instruments Company, Germany), and the
211	concentrations for ~0.2 mg/L for the influent and ≤ 0.1 mg/L for the effluent. The pH
212	values of the influent and effluent were measured by a pH meter (Seven Easy, Mettler
213	Toledo, Switzerland) and were between 7.4 and 7.7 for all stages.

215 We calculated the NO₃⁻ and ClO₄⁻ removal fluxes (g/m^2 -d) according to:

216
$$J = (S^{o}-S)Q/A$$
 (4)

- 217 in which S° and S are the influent and effluent NO_3^- or ClO_4^- concentration (g/L), Q is
- the influent flow rate to the MBfR system (L/d), and A is the membrane surface area
- 219 (m^2) . The CH₄ flux was calculated from the removal fluxes and reaction
- stoichiometries shown in equations 5 through 7.42

$$NO_{2}^{-} + 0.828CH_{4} + H^{+} = 0.04CO_{2} + 0.42N_{2} + 0.158C_{5}H_{7}O_{2}N + 1.6H_{2}O$$
(5)

$$NO_{3}^{-} + 1.2CH_{4} + H^{+} = 0.2CO_{2} + 0.4N_{2} + 0.2C_{5}H_{7}O_{2}N + 2.2H_{2}O$$
(6)

$$ClO_{4}^{-} + 1.613CH_{4} + 0.175NO_{3}^{-} + 0.175H^{+} = Cl^{-} + 0.737CO_{2} + 2.7H_{2}O + 0.175C_{5}H_{7}O_{2}N$$
(7)

- 221 We compared the actual CH₄ flux to the maximum CH₄ flux that can be delivered
- through the composite hollow fiber at the applied CH₄ pressure to indicate if CH₄
- delivery was limiting.⁴¹

224 Biofilm Sampling and DNA extraction

225 We collected biofilm samples when the reactor reached a steady state for all stages

- except Stage 2. Sparging with N_2 gas at the sampling point to preclude O_2 exposure,
- we cut off one ~10-cm-long section from the coupon fiber and then sealed the
- remaining by tying the end into a knot. We then extracted DNA using the DNeasy
- Blood & Tissue Kit (Qiagen, USA) as previously described by Zhao et al (2011).³⁸

We used plasmids containing target fragments as positive controls and to produce 232 calibration curves.⁸ The primers and qPCR conditions were the same as previously 233 described for pcrA – reductase for ClO_4^{-43} narG – reductase for NO_3^{-44} nirS – 234 reductase for NO_2^{-45} , mcrA – formation of methane from most of methanogens, ⁴⁶ 235 $pMMO - CH_4$ mono-oxygenase,⁴⁷ the *16S rRNA* gene for bacteria,⁴⁸ and the *16S rRNA* 236 for archaea.⁴⁹ We used the SYBR Premix Ex Taq Kit (Takara Bio Inc, Japan) and 237 performed qPCR as previously described by Zhao et al. (2011).⁸ The copy numbers 238 of each gene were calculated by comparison to standard curves. Negative controls 239 included water instead of template DNA in the PCR reaction mix. We performed 240 triplicate PCR reactions for all samples and negative controls. The slopes of the 241 plasmid standard curves and efficiency values for quantification by qPCR are shown 242 in Table S4. 243

Results and Discussion

Methane permeability

247	Figure 1 shows the headspace pressures during the CH ₄ -permeation experiment.
248	Steady state was achieved at ~15 hours for the composite fiber. The CH_4
249	permeability for the composite fiber was 1.03×10^{-7} m ³ CH ₄ at standard temperature
250	and pressure - m membrane thickness/ m^2 hollow fiber surface area - d - bar. This
251	permeability is about 10-fold smaller than the H_2 permeability for the same composite
252	fiber and temperature. ⁴¹ Although the Henry's law constant of CH_4 is only slightly
253	smaller than for H_2 , its mass-to-mole ratio (16 g/mol) is about 8 times greater than for
254	H_2 (2 g/mol), making the CH_4 molecule bulkier and more slowly diffusing through the
255	membrane wall.

256 Perchlorate reduction in the presence of nitrate and nitrite

257	Figure 2-A shows the influent and effluent concentrations of NO_2^- , NO_3^- , and ClO_4^-
258	for the entire set of experiments, and Figure 2-B shows the corresponding removal
259	percentages. ClO_4^- reduction could be achieved when CH_4 was the sole electron
260	donor and carbon source. 100% reductions occurred in Stages 2, 3, 5, and 6, and
261	major partial reduction was achieved in Stages 1 and 7. Though Miller et al
262	established a link between ClO_2^- and CH_4 consumption in soils and mixed cultures by
263	D. agitate CKB and methanotrophs (Methylococcus capsulatus Bath or
264	Methylomicrobium album BG8) using acetate as the electron donor and carbon source,

they did not find any upstream connection between ClO_4^- or ClO_3^- reduction and 265 methane oxidation.³⁴ They concluded that oxygen generation during perchlorate 266 267 reduction was negligible or unavailable for aerobic methanotrophs. In contrast, our results clearly show that the MBfR biofilm was able to reduce ClO₄⁻ using CH₄ as the 268 sole electron donor. This success of coupling perchlorate reduction with anaerobic 269 methane oxidation suggests that reduction of other anions might also be coupled to 270 anaerobic methane oxidation. Hence, it would be interesting to explore whether 271 bromate, selenate, chromate and other anion contaminants could be reduced in similar 272 273 CH₄-based systems.

274 Comparison among Stages 1, 2, 6, and 7 shows that NO_2^- inhibited CIO_4^- reduction:

275 ClO₄⁻ reduction was complete when NO₂⁻ was absent in the influent in Stages 2 and 6,

but it decreased to < 50% when NO₂⁻ was present in the influent at a surface loadings

277 of 0.1-0.4 g N/m²-d (1.69 \pm 0.006 mg N/L for Stages 1, 5.22 \pm 0.13 mg N/L for Stage 7

in the influent, respectively).

279 Comparison among Stages 2, 3, 4, 5, and 6 shows that NO_3^- also inhibited CIO_4^-

reduction, but only at high NO_3^- surface loadings. When the NO_3^- surface loadings

were $<0.32\pm0.003$ g N/m²-d (Stages 2, 3, 5, and 6), ClO₄⁻ and NO₃⁻ reductions were

282 complete. However, when the NO_3^- surface loading increased to 0.78 ± 0.09 g N/m²-d

- in Stage 4, ClO₄⁻ reduction dropped to \leq 5%, with NO₃⁻ reduction declining to \leq 85%.
- 284 This trend is consistent with Tang et al,⁹ who used biofilm modeling to quantify the

impact of NO_3^- loading on perchlorate reduction when H_2 was the electron donor.

High NO_3^- loading slowed ClO_4^- reduction by competing for the common donor (H₂ for Tang et al.⁹ and CH₄ here).

288	The MBfR accidently lost its CH ₄ supply for 48 hours (days 75-77), and we provided
289	N_2 gas to keep the fibers pressurized (Figure S3-A). NO_3^- removal dropped sharply
290	to 2% before the CH_4 supply was recovered, but it returned to 70% within 12 hours.
291	However, ClO_4^- removal remained low (2%) after recovery of the CH_4 supply,
292	although it recovered to 100% in Stage 5, when the nitrate loading was smaller. To
293	reinforce that methane was the electron donor responsible for perchlorate and nitrate
294	reduction, we repeated the CH_4 -loss experience during Stage 5 by intentionally
295	replacing the CH_4 supply with N_2 gas for 30 hours beginning on day 94 (Figure S3-B).
296	NO_3^- and ClO_4^- removals dropped to 0 within 12 hours for ClO_4^- and 24 hours for
297	NO_3 , but both returned to 100% after the CH_4 supply was recovered.
298	We calculated the consumption fluxes of NO_2^- , NO_3^- , and ClO_4^- , along with the
298 299	We calculated the consumption fluxes of NO ₂ ⁻ , NO ₃ ⁻ , and ClO ₄ ⁻ , along with the stoichiometric fluxes of CH ₄ (from equations 5 – 7). The fluxes are summarized in
299	stoichiometric fluxes of CH_4 (from equations 5 – 7). The fluxes are summarized in
299 300	stoichiometric fluxes of CH_4 (from equations 5 – 7). The fluxes are summarized in Table 1 for each steady state. One important comparison is between the actual CH_4
299 300 301	stoichiometric fluxes of CH_4 (from equations 5 – 7). The fluxes are summarized in Table 1 for each steady state. One important comparison is between the actual CH_4 consumption and the maximum possible CH_4 flux. The maximum fluxes were 57.9
299 300 301 302	stoichiometric fluxes of CH_4 (from equations 5 – 7). The fluxes are summarized in Table 1 for each steady state. One important comparison is between the actual CH_4 consumption and the maximum possible CH_4 flux. The maximum fluxes were 57.9 mmol CH_4/m^2 -d for Stages 1 – 4 and 86.8 mmol CH_4/m^2 -d for Stages 5-7, both
 299 300 301 302 303 	stoichiometric fluxes of CH ₄ (from equations 5 – 7). The fluxes are summarized in Table 1 for each steady state. One important comparison is between the actual CH ₄ consumption and the maximum possible CH ₄ flux. The maximum fluxes were 57.9 mmol CH ₄ /m ² -d for Stages 1 – 4 and 86.8 mmol CH ₄ /m ² -d for Stages 5-7, both calculated from the K _m of CH ₄ of the composite fiber for the experimental conditions.
 299 300 301 302 303 304 	stoichiometric fluxes of CH ₄ (from equations 5 – 7). The fluxes are summarized in Table 1 for each steady state. One important comparison is between the actual CH ₄ consumption and the maximum possible CH ₄ flux. The maximum fluxes were 57.9 mmol CH ₄ /m ² -d for Stages 1 – 4 and 86.8 mmol CH ₄ /m ² -d for Stages 5-7, both calculated from the K _m of CH ₄ of the composite fiber for the experimental conditions. The maximum CH ₄ delivery flux for Stages 1, 2, 3, 5, 6, and 7 was substantially

307	$(47.5\pm7.20 \text{ mmol/m}^2\text{-d})$ was close to the maximum CH ₄ flux (57.9 mmol CH ₄ /m ² -d).
308	In Stage 4, the effluent concentration of nitrate was stable at 1.39 ± 0.21 mg N/L for 2
309	days before the methane supply was lost (Figure S3). Based on stoichiometry, the
310	maximum methane-delivery rate could remove NO_3^- at a maximum flux of 0.67 g
311	N/m^2 -d, which corresponds to 100% removal of an influent concentration at 10.2 mg
312	N/L. The actual influent concentration was 11.3 ± 0.40 mg N/L in Stage 4, which
313	explains the partial NO_3^- removal and that competition of CH_4 is why ClO_4^- reduction
314	remained very low throughout Stage 4.

Because donor limitation was not an issue for Stages 1 and 7, the negative impact of NO₂⁻ on ClO₄⁻ reduction probably was due to toxicity of NO₂⁻, not to competition for CH₄. Kluber & Conrad reported that methanogenesis activity could be significantly inhibited by adding NO₂^{-.50} King & Schnell reported that NO₂⁻ could inhibit the methane oxidation by methanotrophs, and the inhibition was inversely proportional to headspace methane concentrations.⁵¹

321 Functional Community Structure through Functional Gene Analysis

Figure 3 shows the 16S rRNA gene copies for Bacteria and Archaea, functional-gene copy numbers, and fluxes of the tested electron acceptors through all stages. The copy number of the *pcrA* gene gradually increased from Stage 1 to Stage 5, and this was parallel to overall increasing flux of $ClO_4^- + NO_3^-$ and accumulation of more bacteria, illustrated by the increasing gene copies for the 16S rRNA gene. The copy number of the *pcrA* gene decreased when the flux of all electron acceptors significantly decreased in Stage 6 (due to the absence of NO₃⁻ and NO₂⁻), but increased again when NO₂⁻ was re-introduced at a flux of 0.39 ± 0.01 g N/m²-d in Stage 7. Since most denitrifying bacteria (DB) are able to reduce ClO₄⁻ and may harbor the *pcrA* gene,^{52, 53} it is not surprising that the abundance of the *pcrA* gene was significantly related with the NO₃⁻ flux in our study (Table S5), as well as in previous MBfR studies with H₂.^{8, 54, 55}

334 Similar to the *pcrA* gene, *nirS* and *narG* genes gradually increased from Stage 1 to Stage 5, though the fluxes of $NO_3^++NO_2^-$ decreased from Stage 4 to 5; again, the 335 increases likely were due to accumulating bacteria overall. When NO_3^- and NO_2^- 336 337 were absent in Stage 6, the *nirS* and *narG* abundances dropped by 0.5 to 1 order of magnitude. Also similar with the *pcrA* trend, the *nirS* and *narG* genes increased in 338 Stage 7 when NO_2^{-1} was re-introduced into the MBfR system. However, the *narG* 339 340 abundance was similar to *nirS* in Stage 1, when NO₂⁻ was fed at a low loading, and became much lower than nirS in Stage 7 when NO_2^- was fed at a higher loading. 341 Because the *NarG* gene is not selective for all DB,⁵⁶ *nirS* is mostly used to quantify 342 the DB.⁵⁷ 343

Overall, Bacteria (16S rRNA gene) were ~2 orders of magnitudes higher than Archaea
(Archaeal 16S rRNA gene) through all stages, suggesting that Bacteria dominated
Archaea. While the abundances of *mcrA* and *pMMO* genes were about the same in
Stages 1 and 3, the *pMMO* gene increased much more by Stage 5 and in parallel to the
large increase in the flux of CH₄. The *mcrA* and *pMMO* genes decreased in Stage 6,

349	when NO_3^- and NO_2^- were absent in the system, resulting in a much lower CH_4 flux.
350	The <i>pMMO</i> gene abundance returned to its Stage 5 level with the increase of CH_4 flux
351	Stage 7, but the <i>mcrA</i> gene remained low in Stage 7.
352	Archaea are necessary for the "Reverse Methanogesis" ANMO-D pathway, as they
353	produce electrons for denitrification. The low abundance of Archaea (Fig. 3)
354	supports the "Reverse Methanogenesis" was not important in the ClO ₄ -reducing,
355	CH ₄ -oxidizing biofilm. Further support is given in Figure 4, which shows that the
356	gene copies of mcrA and respiration genes had no correlation.
357	Intracellularly generated O ₂ is essential for the "Intra-Aerobic Type" ANMO-D
358	pathway, in which Candidatus M. oxyfera (or a similar methanotroph) oxidizes CH ₄
359	via an initial mono-oxygenation reaction. Figure 4 shows that the gene copies of
360	<i>pMMO</i> correlated to the gene copies for $narG+nirS+pcrA$, which supports an essential
361	role of O_2 generation associated with ClO_4^- reduction. This association is logical if
362	the key bacteria reducing ClO_4^- used a chlorite dismutase in a manner similar to NO
363	disproportionation in denitrification. ²¹ If O ₂ were produced and consumed
364	intracellularly, ClO ₄ ⁻ reduction occurred via an "Intra-Aerobic Type" ANMO-PR
365	pathway, which is illustrated in Figure 5-A.
366	Rikken et al found that O_2 was released extracellularly during complete ClO_4^-
367	reduction. ⁵⁸ Thus, another possibility is that CH_4 oxidation was coupled to ClO_4^-
368	reduction by a mixture of methanotrophs and perchlorate-reducing bacteria using
369	<i>pMMO</i> and <i>pcrA</i> separately. Miller et al. reported that a variety of

370	methane-oxidizing bacteria, e.g., M. capsulatus Bath, M. album BG8, and M.
371	<i>trichlsporium</i> OB3b, were able to utilize O_2 released from the disproportion of ClO_2^-
372	by dissmilatory perchlorate-reducing bacteria. ³⁴ Sun et al. reported that, ³⁹ in an
373	AMO-D process, co-existing methanotrophs consumed O ₂ preferentially, creating a
374	micro-aerobic environment conducive for denitrification. In addition, the
375	methanotrophs released organic intermediates that served as electron donors for
376	denitrification. ^{15, 22, 30} We name this potential mechanism "micro-Aerobic Methane
377	Oxidation coupled to Perchlorate Reduction," or "mAMO-PR". It is illustrated in
378	Figure 5-B.
379	
517	In summary, we found that the biofilm in an MBfR was able to reduce up to 5 mg/L of
380	In summary, we found that the biofilm in an MBfR was able to reduce up to 5 mg/L of ClO_4^- to non-detectable levels using CH_4 as the only electron donor and carbon source
380	ClO_4^- to non-detectable levels using CH_4 as the only electron donor and carbon source
380 381	ClO_4^- to non-detectable levels using CH_4 as the only electron donor and carbon source in the presence of NO_3^- at a surface loading of ≤ 0.32 g N/m ² -d. While NO_3^- at high
380 381 382	ClO_4^- to non-detectable levels using CH_4 as the only electron donor and carbon source in the presence of NO_3^- at a surface loading of ≤ 0.32 g N/m ² -d. While NO_3^- at high surface loadings (e.g., 0.78 g N/m ² -d) inhibited ClO_4^- reduction due to electron-donor

to the increase of respiratory gene copies, while *mcrA* did not; thus, the CH₄-oxidizing

387 biofilm likely respired ClO_4^- by a pathway that involved generating O_2 using ClO_2^-

dismutation, with the O₂ utilized as a co-substrate for the mono-oxygenation of CH₄.

389 Two options are possible: (1) ANMO-PR via a single strain producing and utilizing

390 intracellular O₂, and (2) mAMO-PR, in which ClO₄⁻-reducing bacteria produce

391 extracellular O_2 by ClO_2^- dismutation, while methanotrophs uses O_2 as a co-substrate

392	to initiate oxidation of CH_4 . This study shows that it is feasible to use methane as an
393	electron donor to biologically remove perchlorate, which is a new option for
394	perchlorate reduction and a new application for the MBfR. Should further study
395	demonstrate that methane is a versatile electron donor, like hydrogen, for reducing
396	oxidized contaminants in water and wastewater treatment, then methane could be used
397	as an inexpensive electron donor.

398 Acknowledgments

- 399 Authors greatly thank "The National Key Technology R&D Program
- 400 (2014ZX07101-012)", "National Natural Science Foundation of China (Grant No.
- 401 21107091, Grant No. 21377109)", and "National High Technology Research and
- 402 Development Program of China (2013AA06A205)" for their financial support.

403 Supporting Information Available

- 404 Table S1-5 and Figure S1-3. This material is available free of charge via the Internet
- 405 at http://pubs.asc.org.

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