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Shifting microbial communities sustain multiyear iron reduction and methanogenesis in ferruginous sediment incubations

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Abstract

Reactive Fe(III) minerals can influence methane (CH₄) emissions by inhibiting microbial methanogenesis or by stimulating anaerobic CH₄ oxidation. The balance between Fe(III) reduction, methanogenesis, and CH₄ oxidation in ferruginous Archean and Paleoproterozoic oceans would have controlled CH₄ fluxes to the atmosphere, thereby regulating the capacity for CH₄ to warm the early Earth under the Faint Young Sun. We studied CH₄ and Fe cycling in anoxic incubations of ferruginous sediment from the ancient ocean analogue Lake Matano, Indonesia, over three successive transfers (500 days in total). Iron reduction, methanogenesis, CH₄ oxidation, and microbial taxonomy were monitored in treatments amended with ferrihydrite or goethite. After three dilutions, Fe(III) reduction persisted only in bottles with ferrihydrite. Enhanced CH₄ production was observed in the presence of goethite, highlighting the potential for reactive Fe(III) oxides to inhibit methanogenesis. Supplementing the media with hydrogen, nickel and selenium did not stimulate methanogenesis. There was limited evidence for Fe(III)-dependent CH₄ oxidation, although some incubations displayed CH₄-stimulated Fe(III) reduction. 16S rRNA profiles continuously changed over the course of enrichment, with ultimate dominance of unclassified members of the order Desulfuromonadales in all treatments. Microbial diversity decreased markedly over the course of incubation, with subtle differences between ferrihydrite and goethite amendments. These results suggest that Fe(III) oxide mineralogy and availability of electron donors could have led to spatial separation of Fe(III)-reducing and methanogenic microbial communities in ferruginous marine sediments, potentially explaining the persistence of CH₄ as a greenhouse gas throughout the first half of Earth history.

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Additional Supporting Information may be found online in the supporting information tab for this article.

1 | INTRODUCTION

Elevated atmospheric methane (CH₄; 100–1,000 ppmv vs. ~2 ppmv in the modern atmosphere) likely played an important role in the first half of Earth history by helping warm Earth's surface temperature enough to sustain liquid water under considerably lower solar radiation (Haqq-Misra, Domagal-Goldman, Kasting, & Kasting, 2008; Kasting, 2005; Pavlov, Kasting, Brown, Rages, & Freedman, 2000; Roberson, Roadt, Halevy, & Kasting, 2011). During this time, the main source of CH₄ was likely hydrogenotrophic methanogenesis (CO₂ + 4 H₂ \rightarrow CH₄ + 2H₂O (Ueno, Yamada, Yoshida, Maruyama, & Isozaki, 2006; Battistuzzi, Feijao, & Hedges, 2004)) from anoxic oceans, which were ferruginous for most of the Archean and Paleoproterozoic eons (Poulton & Canfield, 2011). In these seas, a "ferrous wheel" would have cycled iron from dissolved Fe²⁺ to Fe(III) oxides via microbial photoferrotrophy (and/or abiotic photo-oxidation; Kappler, Pasquero, Konhauser, & Newman, 2005; Crowe, Jones, et al., 2008), and then back to Fe²⁺ via microbial Fe(III) respiration (Craddock & Dauphas, 2011; Johnson, Beard, & Roden, 2008; Konhauser, Newman, & Kappler, 2005; Vargas, Kashefi, Blunt-Harris, & Lovley, 1998).

Ferruginous oceans could have influenced CH₄ cycling by several mechanisms. It is well established that Fe(III)-reducing bacteria have higher affinity for H₂ than hydrogenotrophic methanogens and will therefore outcompete them in the presence of poorly crystalline Fe(III) oxides (e.g., ferrihydrite; Lovley & Phillips, 1987; Lovley & Goodwin, 1988; Zhou, Xu, Yang, & Zhuang, 2014) (note that Fe(III)-reducing bacteria also outcompete acetoclastic methanogens (Lovley & Phillips, 1986), but acetoclastic methanogenesis likely evolved much later in Earth history (Fournier & Gogarten, 2008)). In addition, evidence is accumulating that Fe(III) oxides can mediate or stimulate microbial CH₄ oxidation, either as the direct oxidant (Ettwig et al., 2016; Gal'chenko, 2004), or indirectly by regenerating sulfate by oxidization of reduced sulfur compounds (Sivan, Antler, Turchyn, Marlow, & Orphan, 2014).

The putative microbial metabolism of CH_4 oxidation coupled to Fe(III) reduction is thermodynamically favorable with ferrihydrite

 $(CH_4 + 8Fe(OH)_3 + 15H^+ \rightarrow HCO_3^- + 8Fe^2^+ + 21H_2O; \Delta G_r^0 = -571kJ/mol CH_4)$ and goethite $(CH_4 + 8FeOOH + 15H^+ \rightarrow HCO_3^- + 8Fe^2^+ + 13H_2O; \Delta G_r^0 = -355kJ/mol CH_4)$ as terminal electron acceptors (Caldwell et al., 2008; Zehnder & Brock, 1980). Based on the chemical equations and free energy yields above, we would expect to observe a stoichiometric ratio of 1 CH₄ oxidized per 8 Fe(III) reduced and preferential use of ferrihydrite over goethite as the electron acceptor. Accumulating geochemical evidence for microbial CH₄ oxidation coupled to, or stimulated by, Fe(III) reduction is widespread across modern anoxic ecosystems and anaerobic digester communities (Amos et al., 2012; Beal, House, & Orphan, 2009; Crowe et al., 2011; Egger et al., 2015; Fu et al., 2016; Noroi, Thamdrup, & Schubert, 2013; Riedinger et al., 2014; Rooze, Egger, Tsandev, & Slomp, 2016; Segarra, Comerford, Slaughter, & Joye, 2013; Sivan et al., 2011, 2014; Sturm et al., 2015; Zehnder & Brock, 1980), and a recent study reported simultaneous CH₄ oxidation and ferrihydrite reduction in a 1:8 ratio in an archaea-dominated enrichment culture (Ettwig et al., 2016).

Despite the possible importance of coupled Fe(III) and CH₄ cycling in the Archean and Paleoproterozoic eons, long-term studies of Fe(III) reduction under low organic carbon and high CH₄ conditions remain sparse. Lake Matano, Indonesia, is one of the only modern analogues for the ferruginous Archean ocean (Crowe, Jones, et al., 2008). Despite the abundance of Fe(III) oxides that might be expected to suppress methanogenesis, CH₄ accumulates to 1.4 mm in anoxic deep waters (Crowe, Roberts, Weisener, & Fowle, 2007; Crowe, Jones, et al., 2008; Crowe, O'Neill, et al., 2008; Crowe, et al., 2011; Kuntz, Laakso, Schrag, & Crowe, 2015). Methanotrophy is a key carbon fixation process in Lake Matano's oxic-anoxic transition zone, and the dearth of other oxidants (<100 nm nitrate and sulfate) suggests that Fe(III) might be the terminal electron acceptor in methanotrophy (Crowe et al., 2011; Sturm et al., 2015). In this study, we examined the influence of CH₄ and Fe(III) mineral speciation on rates of Fe(III) reduction, methanogenesis, and CH₄ oxidation, and microbial community composition, over three successive dilutions (500 total days of incubation) of anoxic Lake Matano sediments.

2 | MATERIALS AND METHODS

2.1 | Sample collection and storage

A 15-cm sediment core from 200 m water depth in Lake Matano, Sulawesi Island, Indonesia (2°26'S, 121°15'E; in situ sediment temperature ~27°C), was sampled in November 2014 and subsampled at 5-cm increments. Sediments from 0–5 to 5–10 cm depth were fluffy and black, and 10–15 cm was dark gray. Sediments were sealed in gastight bags with no headspace (Hansen, Thamdrup, & Jørgensen, 2000) and stored at 4°C until incubations began in March 2015.

2.2 | Enrichment medium and substrate synthesis

A modified artificial freshwater medium lacking nitrate and sulfate was developed based on the pore water composition of Lake Matano sediments (S.A. Crowe and D.A. Fowle, unpublished work). The medium contained 825 μ M MgCl₂, 550 μ M CaCO₃, 3 mM NaHCO₃, 3.5 μ M K₂HPO₄, 5 μ M Na₂HPO₄, 225 μ M NH₄Cl, 1 μ M CuCl₂, 1.5 μ M Na₂ MoO₄, 2.5 μ M CoCl₂, 23 μ M MnCl₂, 4 μ M ZnCl₂, 9.4 μ M FeCl₃ and mM Na₂ NTA, 0.07 μ M vitamin B₁₂, 0.4 μ M biotin, and 68.5 μ M thiamine. Filter-sterilized vitamin solutions were added after autoclaving. Ferrihydrite (Fe(OH)₃) and goethite (FeOOH) were synthesized as described in Schwertmann & Cornell (1991) and added to enrichments to 10 mM as described below.

2.3 | Inoculation of enrichment and amendments

The sediment was pre-treated for 36 days at 30°C in 100% N_2 headspace to deplete endogenous organic carbon, electron donors, and reactive electron acceptors. After pre-treatment, sediment from the 0- to 5-cm-depth layer was inoculated in a ratio of sediment to medium of 1:5 (v/v) in an anoxic chamber (97% N_2 and 3% H_2 ; Coy Laboratory Products, Grass Lake, MI, USA). Sediment slurry (35 ml) was aliquoted into 70-ml sterile serum bottles, stoppered with sterile butyl stoppers (Geo-Microbial Technologies, Ochelata, OK, USA; pre-boiled in 0.1 N NaOH) and crimped with aluminum seals. Ferric iron was added either as ferrihydrite or goethite to 10 mm. Bottles were purged with 99.99% N_2 for 1 hr, and CH_4 amendments were injected with 10 ml 99.99% CH_4 and 5 ml 99% $^{13}CH_4$ (Cambridge

Isotope Laboratories, Tewksbury, MA, USA). Controls were autoclaved at 121° C for 1 hr on day 0 and again on day 6 of the 1° enrichment. All treatments were duplicated, and bottles were incubated in the dark at 30° C with shaking at 200 rpm.

After 50 days, the volume of all cultures was reduced to 5 ml, and 30 ml of fresh media was added to each bottle, constituting a sixfold dilution. These 2° enrichments were amended with approximately 10 mM of either ferrihydrite or goethite. All bottles were purged with 99.99% N_2 for 1 hr, and all bottles except N_2 controls were injected with 8 ml 99.99% CH_4 and 2 ml 99% $^{13}CH_4$. Controls were autoclaved again at 121°C for 1 hr. DL-Methionine (10 μ M) was added as a sulfur source. After 303 days, cultures were diluted 10-fold with fresh media into new serum bottles (3° enrichment) with the same substrate and headspace composition as the 2° enrichment. A schematic of the incubation and dilutions is shown at the top of Figures 1–3.

After an additional 220 days, goethite-amended N_2 cultures were diluted 25-fold with fresh anoxic media into new serum bottles. Cultures received either 10 mm goethite or no Fe(III). A subset of cultures received 5 ml of 99.99% H_2 (20% headspace) while all others had 100% N_2 headspace. Controls were autoclaved at 121°C. After 48 days, an anoxic solution of nickel (Ni) and selenium (Se) was added to all bottles, yielding final concentrations of 1 μ m Ni and 1 μ m Se.

2.4 | HCI-extractable Fe²⁺ and Fe³⁺ and soluble Fe²⁺

Samples were taken from each bottle in the anoxic chamber using a 21-gauge needle (BD PrecisionGlideTM) and plastic syringe. Plasticware was stored in the anoxic chamber for at least 24 hr to minimize O_2 sample contamination. For HCl-extractable Fe^{2+} analyses, $100 \, \mu l$ of sediment slurry was extracted with $400 \, \mu l$ 0.6 N HCl in the dark for 1 hr, followed by centrifugation at $10 \, 000 \, g$ for 1 min, injection of $10 \, \mu l$ of supernatant into 990 μl of 10 mM FerroZine reagent in 46 mM HEPES (pH 8.3), and measurement of absorbance at 562 nm (Stookey, 1970). For HCl-extractable Fe^{3+} , $100 \, \mu l$ of sediment slurry was incubated overnight in 0.5 N HCl and 0.25 M NH₂OH-HCl in the dark, followed by centrifugation and measurement as above, and subtraction of HCl-extractable Fe^{2+} as in Kostka and Luther (1994).

2.5 | Methane oxidation

Samples were collected for $\delta^{13}\text{C-DIC}$ analysis by 0.2-µm membrane filtration of medium into crimp top autosampler vials (Thermo Scientific National Target LoVial) and analysis as described in Brandes (2009). Rates of $^{13}\text{CH}_4$ oxidation to $^{13}\text{C-DIC}$ were calculated over the linear period of $\delta^{13}\text{C-DIC}$ increase based on the method in Scheller, Yi, Chadwick, McGlynn, and Orphan (2016). First, the $\delta^{13}\text{C-DIC}$ values were converted into fractional abundances ($^{13}\text{F}=(^{13}\text{C}/^{12}\text{C}+^{13}\text{C})$), and then DIC production from CH₄ oxidation was calculated using the following formula:

$$\Delta[\mathrm{DIC}] = \left(\left([\mathrm{DIC}]_n (^{13} \mathrm{F}_n) \right) - \left([\mathrm{DIC}]_0 (^{13} \mathrm{F}_0) \right) / ^{13} \mathrm{F}_{\mathrm{CH4}} \right),$$

where $[DIC]_n$ and $^{13}F_n$ are equal to the total DIC concentration (mm) and fractional abundance of ^{13}C in the DIC at time n, respectively. $[DIC]_0$ and $^{13}F_0$ are the total DIC concentration (mm) and fractional abundance of DIC at time 0, respectively, and $^{13}F_{CH4}$ is the fractional abundance of ^{13}C in the CH_4 .

2.6 | Headspace methane

Headspace (50 µl) was sampled using a gastight syringe and injected into a gas chromatograph (SRI Instruments 8610C, Torrance, CA, USA) with a HayeSep N column and flame ionization detector to measure headspace CH₄ concentrations. A CH₄ standard (1,000 PPM; Airgas, Radnor Township, PA, USA) was used for calibration.

2.7 | Inductively coupled plasma mass spectrometry

Total dissolved Ni and Se concentrations were measured using inductively coupled plasma mass spectrometry (ICP-MS). To determine the amounts of Ni and Se supplied by the media and Fe(III) oxides, aliquots of media were dispensed in serum bottles, purged with 99.99% N_2 , and amended with 10 mm goethite or ferrihydrite in the same manner as enrichments. Stoppers were penetrated multiple times with 21-gauge stainless steel needles (BD PrecisionGlideTM) to mimic the effect of sampling on enrichment cultures. All samples for ICP-MS were filtered through 0.2- μ m-pore polypropylene syringe filters and diluted in 2% trace metal grade HNO₃ (Fisher Scientific, Inc., Pittsburgh, PA, USA) containing scandium and yttrium as internal standards to account for instrument drift. Calibration standards were prepared from certified stock solutions of Ni (CertiPREP) and Se (BDH), and a blank and calibration standard were measured periodically as quality controls. The measurement detection limits, calculated as three times the standard deviation of the blank (n = 8), were 7 and 128 nm for Ni and Se, respectively.

2.8 | 16S rRNA gene amplicon sequencing

Samples (2 ml) of sediment used for inoculating incubations (hereafter, "sediment inoculum") were taken in February 2015 (prior to pre-treatment) and after incubation for 15 days (1° enrichment), 72 days (2° enrichment) and 469 days (3° enrichment). Nucleic acid was extracted and purified using a MO BIO PowerSoil Isolation Kit following the manufacturer's protocol and MO BIO UltraClean® 15 Purification Kit (MO BIO Laboratories, Carlsbad, CA, USA). 16S rRNA gene amplicons were synthesized from extracted DNA with V4 region-specific barcoded primers F515 and R806 (Caporaso et al., 2011) appended with Illumina-specific adapters according to Kozich, Westcott, Baxter, Highlander, and Schloss (2013) using a Bio-Rad C1000 Touch Thermocycler and QIAGEN Taq PCR Master Mix. Thermal cycling conditions were as follows: initial denaturing at 94°C (5 min), 35 cycles of denaturing at 94°C (40 s), primer annealing at 55°C (40 s), and primer extension at 68°C (30 s). Amplicons were checked for correct size (~400 bp) on a 1% agarose gel and purified using Diffinity RapidTips. Amplicon concentrations were determined on a Qubit[™] (ThermoFisher, Waltham, MA, USA) fluorometer. Amplicons were pooled at equimolar concentrations (4 nmol) and sequenced on an Illumina MiSeq running MISEQ CONTROL software v.2.4.0.4 (Illumina, San Diego, CA, USA) using a 500-cycle MiSeq reagent kit v2 with a 5% PhiX genomic library control, as described by Kozich et al. (2013).

Sequences were deposited as NCBI accession numbers SAMN04532568–04532641 and SAMN05915184–05915222.

2.9 | 16S rRNA gene amplicon sequence analysis

Demultiplexed amplicon read pairs were quality-trimmed with Trim Galore (Babraham Bioinformatics, http://www.bioinformatics.babraham.ac.uk/projects/) using a base Phred33 score threshold of Q25 and a minimum length cutoff of 100 bp. Reads were then analyzed using mothur (Schloss et al., 2009) following its MiSeq standard operating procedure. Highquality paired reads were merged and screened to remove sequences of incorrect length and those with high numbers of ambiguous base calls. Merged reads were dereplicated and aligned to the ARB SILVA database (release 123; available at http://www.mothur.org/wiki/ Silva reference alignment). Sequences with incorrect alignment and those with homopolymers longer than 8 bp were filtered out. Unique sequences and their frequency in each sample were identified, and then a pre-clustering algorithm was used to further denoise sequences within each sample. Sequences were then chimera-checked using UCHIME (Edgar, Haas, Clemente, Quince, & Knight, 2011). Reads were clustered into operational taxonomic units (OTUs) at 97% similarity based on uncorrected pairwise distance matrices. OTUs were classified using SILVA reference taxonomy database (release 123, available at http://www.mothur.org/wiki/Silva_reference_files). Chao 1 (species richness), phylogenetic diversity, and Shannon index (species evenness) estimates were generated using mothur after normalization to 4,000 sequences per sample.

3 | RESULTS

3.1 | Iron reduction

3.1.1 | **1° Enrichment**—Over the first 10 days of incubation, HCl-extractable Fe²⁺ increased from 10 to 25 mm in ferrihydrite treatments (Figure 1a) and from 10 to 20 mm in goethite treatments (1–2 mm/day; Figure 1b). From day 6 to 10, HCl-extractable Fe³⁺ (7 and 12 mm in ferrihydrite and goethite treatments, respectively) was completely consumed in all bottles except autoclaved controls with ferrihydrite (data not shown). Iron reduction rates were identical with and without CH₄ (Figure 1a,b). Initial autoclaving did not suppress Fe(III) reduction. A second round of autoclaving on day 6 slightly suppressed further activity. From day 10 to day 28, HCl-extractable Fe²⁺ fluctuated in ferrihydrite treatments (Figure 1a) and declined slightly in goethite treatments (Figure 1b). Soluble Fe²⁺ was consistently <1% of HCl-extractable Fe²⁺, and sediment-free controls did not reduce Fe(III) (data not shown).

3.1.2 | **2° Enrichment**—After 1:6 dilution and 10 mm ferrihydrite addition on day 50, HCl-extractable Fe²⁺ increased from 3 to 4 mm over two days and then remained constant through the final time point (day 497) in bottles with and without added CH₄ (Figure 1a). After 10 mm goethite addition on day 50, HCl-extractable Fe²⁺ increased from 2 to 3 mm after a two-day lag period. Thereafter, HCl-extractable Fe²⁺ rose to 4 mm by day 497 in goethite treatments with added CH₄; without CH₄, HCl-extractable Fe²⁺ dropped back to 2 mm (Figure 1b). Autoclaved controls had no activity. Black magnetic minerals formed in all

ferrihydrite treatments except autoclaved controls (Figure 1a). No magnetic minerals formed in goethite treatments (Figure 1b).

3.1.3 | **3° Enrichment**—After 1:10 dilution and addition of 10 mm ferrihydrite on day 352, HCl-extractable Fe²⁺ doubled in the first week in N₂ treatments and bottle CH₄-1 (Figure 1a). Bottle CH₄-2 displayed similar activity after a two-week lag period. Over an additional 100 days (day 466), HCl-extractable Fe²⁺ increased to 2 mm. Goethite treatments and autoclaved controls had minimal activity (Figure 1b). As in the 2° enrichment, magnetic minerals formed in the presence of ferrihydrite (Figure 1a), but not goethite (Figure 1b).

3.2 | Trace metal concentrations

Total dissolved Ni averaged 41 ± 20 nm in fresh basal growth media and was neither affected by Fe(III) oxide additions nor by puncturing of stainless steel needles through stoppers into culture liquid (Table S1). Dissolution of ferrihydrite in HNO₃ liberated significant Ni (2.5 μ m; Table S1). Nickel was higher in enrichment cultures than in basal media: 96–286 and 54–134 nm with ferrihydrite and goethite, respectively; Table S2). Selenium was consistently below the detection limit (<128 nm) in growth media and enrichment culture.

3.3 | Methane production

Goethite treatments consistently displayed higher CH_4 production than those with ferrihydrite (Figure 2). In the 1° goethite enrichment, methanogenesis (13–19 μ M CH_4 /day) coincided with the period of Fe(III) reduction and stopped after HCl-extractable Fe³⁺ was completely consumed on day 10. Methanogenesis persisted throughout the 3° goethite enrichment (3 μ M CH_4 /day). Negligible CH_4 was produced in the presence of ferrihydrite, except for the final time point for bottle N_2 -2 in the 3° enrichment.

In an additional 4° enrichment (days 571–663), we tested the effect of H_2 , Ni, and Se amendments, as well as no Fe(III) controls, on CH₄ production in goethite treatments (Figure S1). As in previous enrichments, Bottle 1 consistently produced more CH₄ (17 μ M CH₄/day) than Bottle 2. There were no significant differences in CH₄ production with and without 20% H₂ headspace and 10 mM goethite. Like in previous enrichments with goethite, minimal Fe(III) reduction was observed (data not shown). No CH₄ was produced in any of the treatments between day 619, when 1 μ M Ni and Se were added, and day 663.

3.4 | Methane oxidation

- **3.4.1** | **1° Enrichment**— 13 C incorporation into DIC began on day 6 in both ferrihydrite and goethite treatments and continued for the remainder of the sampling period (Figure 3a,b). Ferrihydrite treatments showed lower 13 C-DIC enrichment but higher total DIC production (totaling to 1– 2 μ M CH $_4$ oxidized/day) than goethite treatments, which had greatest δ^{13} C enrichment but decreasing DIC concentrations, making calculation of CH $_4$ oxidation rates impossible. Autoclaved controls showed neither 13 C incorporation nor DIC production.
- **3.4.2** | **2° Enrichment**—Both ferrihydrite treatments displayed ¹³C enrichment (Figure 3a), but declining DIC, precluding calculation of CH₄ oxidation rates. Initial pH of 8

declined to 7.6, 6.7 and 6 in the autoclaved, N_2 and CH_4 treatments, respectively. DIC in goethite treatments with CH_4 dropped to undetectable values within three weeks, suggesting sampling or analytical error, which precluded accurate isotopic measurement at these time points. These data are thus not considered further. Autoclaved and N_2 controls did not show pH changes.

3.4.3 | **3° Enrichment**—Bottle CH_4 -2 with ferrihydrite was the only treatment with significant ^{13}C incorporation into DIC over the first 15 days (Figure 3a). Over the same interval, DIC increased in both ferrihydrite-amended bottles (yielding CH_4 oxidation rates of 32 and 7 μ M/day in bottles 1 and 2, respectively) and pH dropped from 8.2 to 7.1 and 7.9 in bottles 1 and 2, respectively. By day 470, ^{13}C enrichment and DIC concentrations in both ferrihydrite-amended bottles had returned to a level similar to that at the start of the 3° enrichment. Autoclaved controls did not exhibit any change in DIC and pH. Goethite treatments had initial DIC concentrations (3–5 mM) higher than those in previous enrichments. In the goethite-amended autoclaved controls and bottle CH_4 -1, DIC concentrations dropped over the 3° enrichment. Only goethite-amended bottle CH_4 -2 increased in DIC, without concurrent ^{13}C enrichment. Large DIC variability implies that reported rates may be underestimates if declining pH led to outgassing of ^{13}C -DIC into the headspace CO_2 pool.

3.5 | Microbial taxonomy

- **3.5.1** | **Inoculum**—16S rRNA gene amplicons from the sediment inoculum were dominated by Bathyarchaeota (25%), formerly Miscellaneous Crenarchaeotal Group, and unclassified Archaea (11%; Figure 4).
- **3.5.2** | **1° Enrichment**—Species richness, evenness, and phylogenetic diversity decreased relative to the inoculum in all treatments (Figure 4). Geobacteraceae (Deltaproteobacteria) became dominant (22–36%) in all ferrihydrite treatments (Figure 4a); the dominant OTU had 97% similarity to *Geothermobacter* sp. Ferrihydrite-amended bottle CH₄-1 was enriched in Betaproteobacteria, specifically Comamonadaceae (17%) and Rhodocyclaceae (9%; Figure 4a). Bathyarchaeota persisted in goethite treatments (11–25%; Figure 4b).
- **3.5.3** | **2° Enrichment**—All treatments declined further in species richness, evenness, and phylogenetic diversity. Unclassified Desulfuromondales dominated both ferrihydrite and goethite enrichments (34–68%). The dominant OTU had 98% similarity to *Geobacter hephaestius/Geobacter lovleyi*. Geobacteraceae declined in ferrihydrite enrichments (2%–18%; Figure 4a). Campylobacteraceae (Epsilonproteobacteria), a trace constituent of the inoculum and 1° enrichment, were enriched in goethite treatments with CH₄ (23%–40%); the dominant OTU had 98% similarity to *Sulfurospirillum barnesii* (Figure 4b). The most abundant methanogenic Euryarchaeota family, Methanobacteriaceae, comprised 1%–2% and 6%–7% of sequences in ferrihydrite and goethite treatments, respectively; the dominant OTU had 100% similarity to *Methanobacterium flexile*. Bathyarchaeota were depleted compared to the 1° enrichment (Figure 4).

3.5.4 | **3° Enrichment**—Species richness, evenness, and phylogenetic diversity continued to decline (Figure 4a,b). Unclassified Desulfuromondales dominated goethite treatments (32%–76%; Figure 4b) and were less abundant in ferrihydrite treatments (18%–38%); as in the 2° enrichment, the dominant OTU had 98% identity to *G. hephaestius/G. lovleyi*. Rhodocyclaceae were more abundant in ferrihydrite treatments with CH₄ (14%–15%) than with N₂ (2%–4%; Figure 4a); the dominant OTU had 100% similarity to *Azospira oryzae/Dechlorosoma suillum*. Peptococcaceae (Firmicutes) were most abundant in bottle CH₄-2 with ferrihydrite (30%); the dominant OTU had 96% similarity to uncultured members of the genus *Thermincola*. Syntrophaceae (Deltaproteobacteria) were enriched in all ferrihydrite treatments (11%–16%) and goethite treatments with N₂ (8%–15%); the dominant OTU had 97% similarity to *Smithella propionica*. Methanobacteriaceae comprised 1%–4% of sequences in goethite treatments and were absent from ferrihydrite treatments; as in the 2° enrichment, the dominant OTU had 100% identity to *M. flexile*.

4 | DISCUSSION

4.1 | Fe(III) reduction rates in long-term ferruginous sediment incubations

Initial rates of HCl-extractable Fe^{2+} production (1–2 mm/day) in the 1° enrichment were similar to those from freshwater wetlands with organic carbon as the electron donor (Jensen, Thamdrup, Rysgaard, Holmer, & Fossing, 2003; Kostka, Roychoudhury, & Van Cappellen, 2002; Roden & Wetzel, 2002). Despite replenishment of Fe(III) substrates, activity declined with each successive transfer, likely reflecting organic carbon limitation. The next most thermodynamically favorable electron donor, H_2 , could have been supplied by fermenters (such as Syntrophaceae in the 3° enrichment), but would ultimately still require a source of organic carbon. Some of our incubations display evidence for CH_4 , the next most thermodynamically favorable electron donor, as a source of electrons for Fe(III) reduction (e.g., higher Fe^{2+} yields with CH_4 addition with ferrihydrite in 1° and 3° enrichments, and with goethite in the 2° enrichment; see further discussion below).

Higher Fe(III) reduction rates were maintained on ferrihydrite than goethite, consistent with its higher energetic yield and (typically) greater surface area. Magnetic mineral formation was likely due to adsorption of Fe^{2+} onto ferrihydrite followed by solid-state conversion of ferrihydrite to magnetite (Hansel et al., 2003). As the HCl-extraction method does not dissolve magnetite and magnetite-adsorbed Fe^{2+} (Poulton & Canfield, 2005), it is possible that our Fe(III) reduction rates based on HCl-extractable Fe(II) production were underestimates of the total Fe(III) reduction.

4.2 | Fe(III) oxide mineralogy controls methane production and methanogen taxonomy

Our observation of higher rates of methanogenesis in goethite vs. ferrihydrite amendments is consistent with prior results showing that bacteria that reduce ferrihydrite better outcompete methanogenic archaea for H₂ and acetate than those that reduce more crystalline Fe(III) oxides, including goethite (Hori, Müller, Igarashi, Conrad, & Friedrich, 2010; Lovley & Goodwin, 1988; Lovley & Phillips, 1987; Roden & Wetzel, 1996; Zhou et al., 2014). This outcompetition is also broadly supported by taxonomic shifts in our enrichment cultures. In particular, anaerobic heterotrophs such as *Geothermobacter* sp. (Kashefi, Holmes, Baross, &

Lovley, 2003) were enriched in ferrihydrite treatments by day 15 and may have outcompeted other microbes for organic carbon sources.

Higher abundances of Methanobacteriaceae (0.1%-1% and 1%-4% on days 15 and 469, respectively) in goethite than ferrihydrite treatments (0.1%) suggest that CH₄ in goethite treatments came from the substrates used by Methanobacteriaceae (H_2/CO_2 , formate, or CO). Addition of H₂ did not stimulate additional methanogenesis in the 4° amendment, implying another limiting substrate or growth condition. The ferrihydrite treatment (bottle 2) that produced CH₄ by day 469 contained 3% Methanosaetaceae; the most dominant OTU had 98% similarity to *Methanosaeta concilii*, in agreement with observations from the Lake Matano water column (Crowe et al., 2011). *Methanosaeta* spp. produce CH₄ from acetate, or from H_2/CO_2 via direct interspecies electron transfer with *Geobacter* (Rotaru et al., 2014).

4.3 | Fe(III)-dependent CH₄ oxidation

Enrichments were established under conditions thought to be favorable for Fe(III)-dependent CH_4 oxidation, with Fe(III) oxides and CH_4 as the most abundant electron acceptors and donors, respectively. In the 1° enrichment, incorporation of $^{13}CH_4$ into DIC overlapped with the second phase of Fe(III) reduction (days 6–10), but calculating the stoichiometry of CH_4 oxidized to Fe(III) reduced posed a challenge due to similar rates of Fe(III) reduction with and without added CH_4 and in autoclaved controls. ^{13}C -DIC enrichment from the back reaction of hydrogenotrophic methanogenesis (Zehnder & Brock, 1979) was ruled out because CH_4 oxidation continued after Fe(III) reduction and methanogenesis stopped at day 10. Therefore, CH_4 was likely oxidized by an electron acceptor other than Fe(III) (e.g., O_2 , Mn(IV), NO_x^- , SO_4^{2-}) in the 1° enrichment, likely supplied by residual sediment or inadvertent introduction of air.

During the first 15 days of the 3° enrichment, rates of CH₄ oxidation and HCl-extractable Fe²⁺ production were similar (~10–20 μ M/day) and roughly consistent with the low rates presented in Ettwig et al. (2016) that yielded a 1:8 ratio. However, the lack of multiple time points for the interval of simultaneous Fe(III) reduction and CH₄ oxidation, as well as similar initial rates of Fe(III) reduction with and without CH₄ throughout this interval, prevents us from attributing this activity to Fe(III)-dependent CH₄ oxidation with high confidence.

It is notable that the two incubations with the highest rates of CH_4 oxidation (ferrihydrite bottle 1 in 1° and 3° enrichments) were also the only treatments with very different microbial community compositions relative to other bottles in the same enrichment. In the 1° enrichment, ferrihydrite bottle 1 was enriched in Betaproteobacteria (Comamonadaceae and Rhodocyclaceae). In the 3° enrichment, the CH_4 -1 sample had less Peptococcaceae than other ferrihydrite incubations. By day 469, the betaproteobacterium *A. oryzae/D. suillum*, a member of the Rhodocyclaceae family, was more abundant in both of the CH_4 vs. N_2 treatments. The potential role of this microbe in CH_4 cycling remains unclear, as laboratory cultures of this species are not known to oxidize CH_4 . Notably, related members of the Betaproteobacteria, including the genera *Azospira* and *Comamonas* found here, are typically facultative anaerobes that can use alternative electron acceptors like NO_3^- , NO_2^- , or

perchlorate (Reinhold-Hurek & Hurek, 2015; Willems, 2014). As such, these Betaproteobacteria are poised to respond to enhanced electron acceptor supply that accompanies pulse of O₂. Anecdotally, Betaproteobacteria are frequently associated with environments that are characterized by fluctuating redox conditions and periodic exposure to O₂ (Converse, McKinley, Resch, & Roden, 2015). Thus, their growth in our incubations may be a response to trace O₂ introduction. It is also possible that the growth of novel organisms capable of high rates of Fe(III)-dependent CH₄ oxidation was inhibited by other unidentified factors, potentially related to the batch-style incubations, the use of butyl rubber stoppers (Niemann et al., 2015), or the lack of a critical substrate in the enrichment medium.

4.4 | Effect of Fe(III) oxide and carbon substrates on microbial community diversity

The microbial community underwent multiple shifts over the 500-day incubation, with an overall decrease in species members, evenness, and phylogenetic diversity, likely in response to declining organic carbon. By the 3° enrichment, species evenness was consistently lower in each goethite-amended treatment than in the respective ferrihydrite-amended treatment. This could mean that the greater energetic yield of ferrihydrite reduction fosters higher diversity, or that the higher reactivity of ferrihydrite allowed it to be utilized by more organisms than goethite.

All of the most enriched taxa in our enrichments comprised 0.1% of the inoculum community and have members that reduce Fe(III) in laboratory cultures. Within those taxa, the most abundant OTUs were closely related to organisms capable of Fe(III) reduction (Geothermobacter sp., G. hephaestius/G. lovleyi, Thermincola sp., and S. barnesii) (Kashefi et al., 2003; Stolz et al., 1999; Zavarzina et al., 2007). Desulfuromonadales was the only metal-reducing taxon that was continuously present in all enrichments (3%-11%, 34%-53%, and 18%–76% in the, 1°, 2°, and 3° enrichment, respectively). Other taxa differed significantly in their abundance over the course of incubation. Geobacteraceae was enriched at day 15 with ferrihydrite (22%–36%) but had declined in abundance by day 72 (8%–18%). Still other taxa, including Rhodocyclaceae and Peptococcaceae, were enriched in the presence of ferrihydrite at day 469. In goethite treatments, Campylobacteraceae (23%–40%) were enriched at day 72, but were minimal at day 469. The succession of different metalreducing taxa may be due to the changing availability of electron donors (e.g., H₂ and organic C). Enrichment of Syntrophaceae, known for their syntrophic fermentative interactions, suggests the establishment of syntrophy in the 3° enrichment in response to depletion of electron donors.

4.5 | Nickel sources

Enrichment cultures contained ~2–10× more total dissolved Ni than the basal growth medium. The inoculum (~60 nm in Lake Matano deep water; Crowe, O'Neill, et al., 2008) would not have significantly contributed to the Ni pool past the 1° enrichment, and repeated needle exposure had no effect on Ni concentrations. The Ni source to enrichment cultures was likely partial ferrihydrite dissolution, as ferrihydrite readily scavenges Ni from solution (Zegeye et al., 2012), while its dissolution liberates Ni (Table S1; Crowe, O'Neill, et al., 2007). Slow Ni leaching from silicate glass during extended contact between microbes and

the serum bottles could have contributed another source of Ni in microbial enrichments vs. abiotic controls (Hausrath, Liermann, House, Ferry, & Brantley, 2007).

4.6 | Geobiological implications

Our results point to a mineralogical control on Fe(III) reduction, methanogenesis, and microbial community composition and diversity, under conditions of severe organic carbon limitation. These conditions likely existed in Archean and Paleoproterozoic oceans with relatively low amounts of primary production (Farquhar, Zerkle, & Bekker, 2011; Knoll, Bergmann, & Strauss, 2016). We posit that the relative abundance and distribution of Fe(III) phases in marine sediments would have impacted methanogenesis rates in the Archean and Paleoproterozoic. Sediments below shallow water columns were likely fed by abundant amorphous Fe(III) from photoferrotrophic activity, resulting in rapid sedimentation of amorphous Fe(III) phases (e.g., ferrihydrite). These Fe(III) oxides could have supported diverse Fe(III)-reducing communities that outcompeted other taxa such as methanogens for limited carbon and nutrients.

Conversely, slow deposition and aging of ferrihydrite to goethite could have limited both the abundance and diversity of Fe(III)-reducing microbes in sediments, allowing for more organic carbon remineralization via methanogenesis than Fe(III) reduction, as recently calculated for Lake Matano (Crowe et al., 2011; Kuntz et al., 2015). In the open ocean, organic carbon and Fe(III) would likely have been consumed before reaching sediments, leaving behind more crystalline Fe(III) phases. Importantly, the role of Fe(III)-driven CH_4 oxidation appears limited given our experimental results, although we cannot rule out this pathway given that some of our data suggest it may operate at low rates.

Availability of trace metal nutrients is another important consideration in potential controls on ancient CH₄ and Fe cycling. Measurements of Ni/Fe ratios in ancient marine sediments indicate that total dissolved Ni decreased from ~400 nm before 2.7 Ga to 200 nm between 2.7 and 2.5 Ga, to modern levels of 2–11 nm at ~0.5 Ga, assuming that the Fe(III) minerals in Archean sediments were of biological origin (Eickhoff et al., 2014; Konhauser et al., 2009, 2015). It is likely that abundant Ni would have been bound and sequestered in Fe(III) oxide-rich sediments. Rapid and widespread Archean redox cycling of Fe(III) could have served as constant source of Ni for methanogenic communities. The influence of changing availability of Se (Stüeken, Buick, & Anbar, 2015) and other trace nutrients on methanogenesis rates through time remains open for further exploration.

Overall, our results support a model for a sustained CH_4 greenhouse in the Archean and Paleoproterozoic due to emissions from ferruginous oceans with spatially segregated habitats of bacterial reduction of reactive Fe(III) oxides and methanogenesis in the presence of less reactive Fe(III) phases. Rates of Fe(III) deposition, aging, and recrystallization may thus have played an important role in regulating the preservation of sedimentary Fe(III), the production of CH_4 , and the ecology and diversity of the biosphere during the first half of Earth history. By the mid-Proterozoic, rising seawater sulfate likely stimulated anaerobic CH_4 oxidation, thereby minimizing marine CH_4 emissions and the CH_4 greenhouse (Olson, Reinhard, & Lyons, 2016).

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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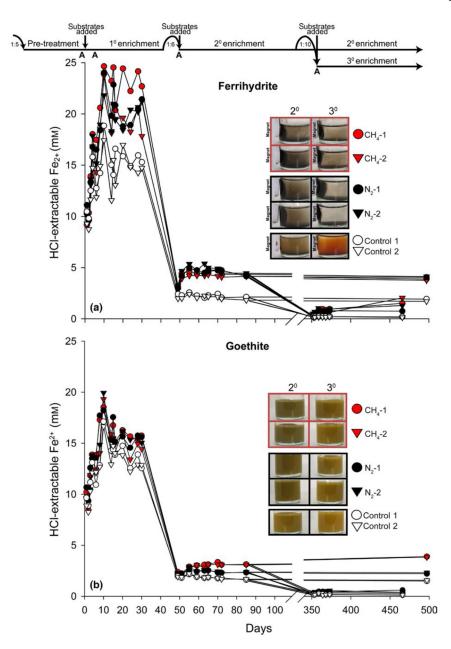


FIGURE 1. HCl-extractable Fe^{2+} for sediment enrichments with (a) ferrihydrite and (b) goethite over 497 days. Timeline at top shows transfer dates and dilution ratios. "A" represents days that controls were autoclaved. Red and black symbols represent treatments with and without CH_4 , respectively. White symbols represent autoclaved controls. All treatments were run in duplicate (circle and triangle symbols). Photographs depict 2° and 3° enrichment bottles on day 497 with evidence for magnetic mineral formation in live treatments amended with ferrihydrite

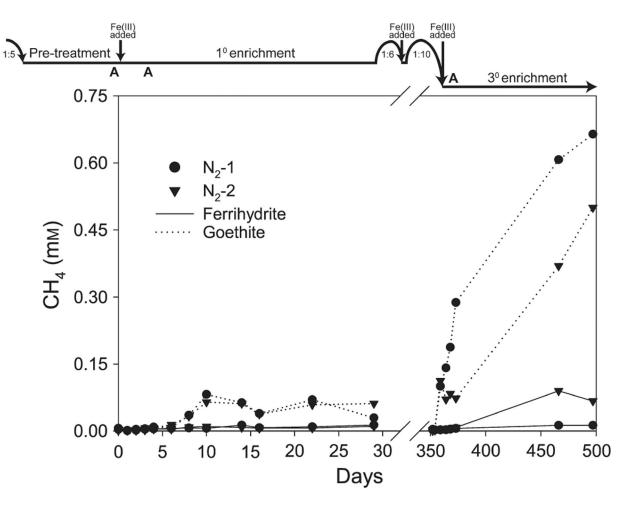


FIGURE 2. Accumulation of CH_4 in the headspace of sediment enrichments. Timeline at top shows transfer dates and dilution ratios. Solid and dotted lines represent ferrihydrite and goethite treatments, respectively. All treatments were run in duplicate (circle and triangle symbols). Original headspace was 100% N_2

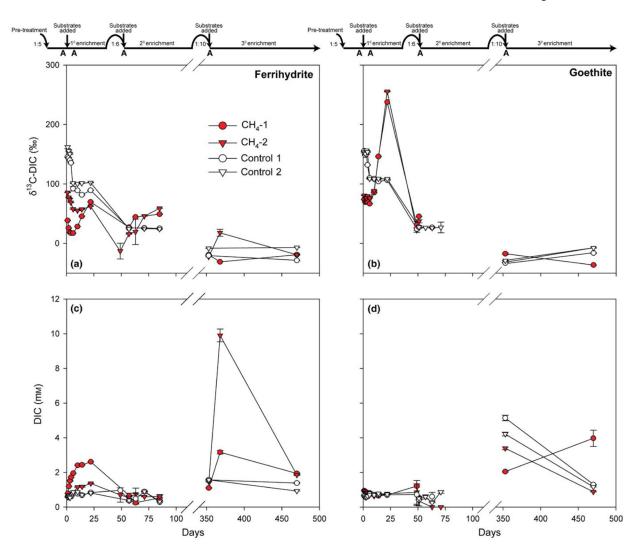


FIGURE 3.

Dissolved inorganic carbon (DIC) isotopic composition and concentration for sediment enrichments amended with $^{13}\text{CH}_4$ and either (a,c) ferrihydrite or (b,d) goethite. Timeline at top shows transfer dates and dilution ratios. "A" represents days that controls were autoclaved. Red and white symbols represent live treatments and autoclaved controls, respectively. Errors bars represent standard deviation of triplicate measurements. Calculated methane oxidation rates for the 1° enrichment were 1.7 and 1.1 μm CH₄/day ferrihydrite bottles 1 and 2, respectively, and 0.2 and 0.8 μm CH₄/day for goethite bottles 1 and 2, respectively. Isotopic data are not plotted for DIC concentrations 0.5 mm. Rate calculations were not possible for the 2° enrichment due to low/variable DIC

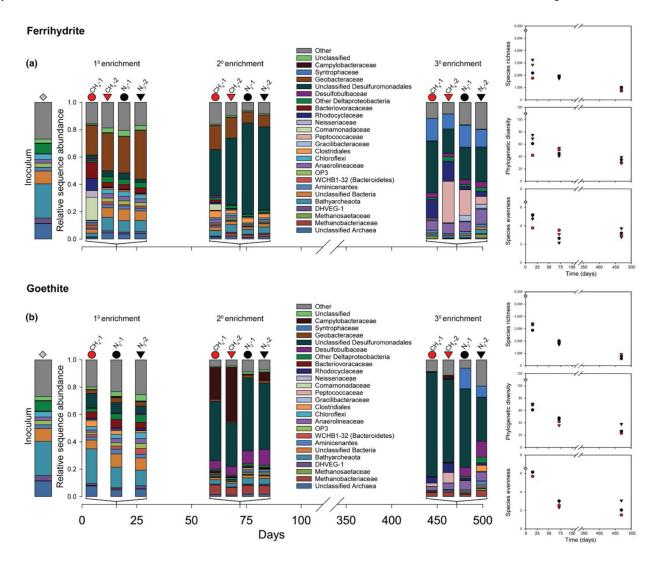


FIGURE 4. 16S rRNA gene diversity and phylogenetic diversity for inoculum and sediment enrichments amended with (a) ferrihydrite and (b) goethite. Samples were taken on day 15 (1 $^{\circ}$ enrichment), 72 (2 $^{\circ}$ enrichment), and 469 (3 $^{\circ}$ enrichment). Red and black symbols represent treatments with and without CH₄, respectively. Gray diamonds represent inoculum samples. All treatments were run in duplicate (circle and triangle symbols). Species richness, phylogenetic diversity, and species evenness for the sediment inoculum and enrichments normalized to 4,000 sequences per sample are shown to the right of bar charts