



Dominance of Diffusive Methane Emissions From Lowland Headwater Streams Promotes Oxidation and Isotopic Enrichment

Andrew L. Robison^{1,2*}, Wilfred M. Wollheim¹, Clarice R. Perryman³, Annie R. Cotter¹, Jessica E. Mackay¹, Ruth K. Varner³, Paige Clarizia³ and Jessica G. Ernakovich¹

¹Department of Natural Resources and the Environment, University of New Hampshire, Durham, NH, United States, ²Stream Biofilm and Ecosystem Research Laboratory, École Polytechnique Fédérale de Lausanne, Lausanne, Switzerland, ³Department of Earth Sciences, University of New Hampshire, Durham, NH, United States

OPEN ACCESS

Edited by:

Daniel F. McGinnis,
Université de Genève, Switzerland

Reviewed by:

Mark Alexander Lever,
ETH Zürich, Switzerland
Changhui Wang,
Shanxi Agricultural University, China

*Correspondence:

Andrew L. Robison
andrew.robison@epfl.ch

Specialty section:

This article was submitted to
Biogeochemical Dynamics,
a section of the journal
Frontiers in Environmental Science

Received: 08 October 2021

Accepted: 30 December 2021

Published: 31 January 2022

Citation:

Robison AL, Wollheim WM,
Perryman CR, Cotter AR, Mackay JE,
Varner RK, Clarizia P and
Ernakovich JG (2022) Dominance of
Diffusive Methane Emissions From
Lowland Headwater Streams
Promotes Oxidation and
Isotopic Enrichment.
Front. Environ. Sci. 9:791305.
doi: 10.3389/fenvs.2021.791305

Inland waters are the largest natural source of methane (CH₄) to the atmosphere, yet the contribution from small streams to this flux is not clearly defined. To fully understand CH₄ emissions from streams and rivers, we must consider the relative importance of CH₄ emission pathways, the prominence of microbially-mediated production and oxidation of CH₄, and the isotopic signature of emitted CH₄. Here, we construct a complete CH₄ emission budgets for four lowland headwater streams by quantifying diffusive CH₄ emissions and comparing them to previously published rates of ebullitive emissions. We also examine the isotopic composition of CH₄ along with the sediment microbial community to investigate production and oxidation across the streams. We find that all four streams are supersaturated with respect to CH₄ with diffusive emissions accounting for approximately 78–100% of total CH₄ emissions. Isotopic and microbial data suggest CH₄ oxidation is prevalent across the streams, depleting approximately half of the dissolved CH₄ pool before emission. We propose a conceptual model of CH₄ production, oxidation, and emission from small streams, where the dominance of diffusive emissions is greater compared to other aquatic ecosystems, and the impact of CH₄ oxidation is observable in the emitted isotopic values. As a result, we suggest the CH₄ emitted from small streams is isotopically heavy compared to lentic ecosystems. Our results further demonstrate streams are important components of the global CH₄ cycle yet may be characterized by a unique pattern of cycling and emission that differentiate them from other aquatic ecosystems.

Keywords: methane, streams, isotopes, methanogens, methane oxidation, emissions

INTRODUCTION

Since the start of the industrial revolution, atmospheric concentrations of methane (CH₄) have increased nearly threefold (Saunio et al., 2020). This atmospheric enrichment of CH₄ is caused predominantly by anthropogenic activities; however, aquatic ecosystems comprise roughly half of all global CH₄ emissions (Rosentreter et al., 2021). Most research into CH₄ emissions and cycling in freshwater systems has occurred in wetlands, lakes, and impoundments, but recent studies have started to highlight the role of rivers and streams in the global CH₄ budget (Stanley et al., 2016; Zhang

et al., 2020). Rosentreter et al. (2021) estimate that streams and rivers emit approximately $30.5 \text{ Tg CH}_4 \text{ yr}^{-1}$ globally, which represents about 20% of the total annual emissions from lakes ($151 \text{ Tg CH}_4 \text{ yr}^{-1}$) or wetlands ($149 \text{ Tg CH}_4 \text{ yr}^{-1}$; Saunois et al., 2020). Methane cycling in small headwater streams, which comprise a majority of river network length (Bishop et al., 2008), represents a gap in our understanding of aquatic CH_4 dynamics. Headwater streams are known to be disproportionately important in the emissions of carbon dioxide within river networks (Hotchkiss et al., 2015; Marx et al., 2017), yet their relative importance regarding CH_4 remains undefined. Additionally, most investigations of CH_4 focus on quantifying CH_4 concentration or flux rather than how CH_4 cycles through the ecosystem (Stanley et al., 2016). As a result, more comprehensive measurements of CH_4 emissions and an improved understanding of CH_4 cycling are needed to accurately include streams in regional and global CH_4 budgets.

The fundamental controls of CH_4 cycling are consistent across aquatic ecosystems, thus we can apply much of what is known about lentic systems to study stream ecosystems. In general, CH_4 is produced via methanogenesis by archaea under anoxic conditions (Chowdhury and Dick, 2013), and the inundated sediments of aquatic environments are generally well-suited for this because of their low redox conditions (e.g., Wik et al., 2018). A key process that consumes CH_4 in surface waters is biological oxidation by bacterial and archaeal methanotrophs, which can function under oxic or anoxic conditions (Conrad, 2009). Oxidation of CH_4 is prevalent in aquatic ecosystems because methanotrophs are generally more productive where there are strong opposing gradients of CH_4 and oxygen, common to aquatic ecosystems (Chowdhury and Dick, 2013). The oxidation of CH_4 can be an important process mitigating CH_4 emissions from aquatic systems (Bastviken et al., 2008; Sawakuchi et al., 2016).

Streams are largely differentiated from lentic waterbodies by the presence of flow, which may affect CH_4 cycling via turbulent mixing and oxygenation of the water column and benthic environment (Trimmer et al., 2010). Previously, the mixing of oxic water into stream sediments was thought to limit methanogenesis, but stream sediments have been shown to support methanogenesis (Sanders et al., 2007; Bodmer et al., 2020). High rates of CH_4 oxidation have also been observed in streams (Shelley et al., 2017), attributed to the strong redox gradient at the sediment-water interface and the introduction of oxygenated waters into the sediments via hyporheic flow (Hampton et al., 2020). However, the relatively rapid mixing of streams and rivers due to flow compared to lentic systems may reduce the proportion of CH_4 oxidized before emissions (Sawakuchi et al., 2016). That is, higher turbulence and gas exchange in streams and rivers may result in less depletion of the CH_4 pool *via* oxidation because of reduced transport time. This may explain why complete oxidation of the dissolved CH_4 pool has been observed in lakes (Bastviken et al., 2008), but not in streams and rivers (Sawakuchi et al., 2016).

The relative importance of CH_4 production and oxidation also affects the isotopic composition of CH_4 (Chanton, 2005; Chanton et al., 2006). The $\delta^{13}\text{C}\text{-CH}_4$ and $\delta\text{D}\text{-CH}_4$ signature of emitted

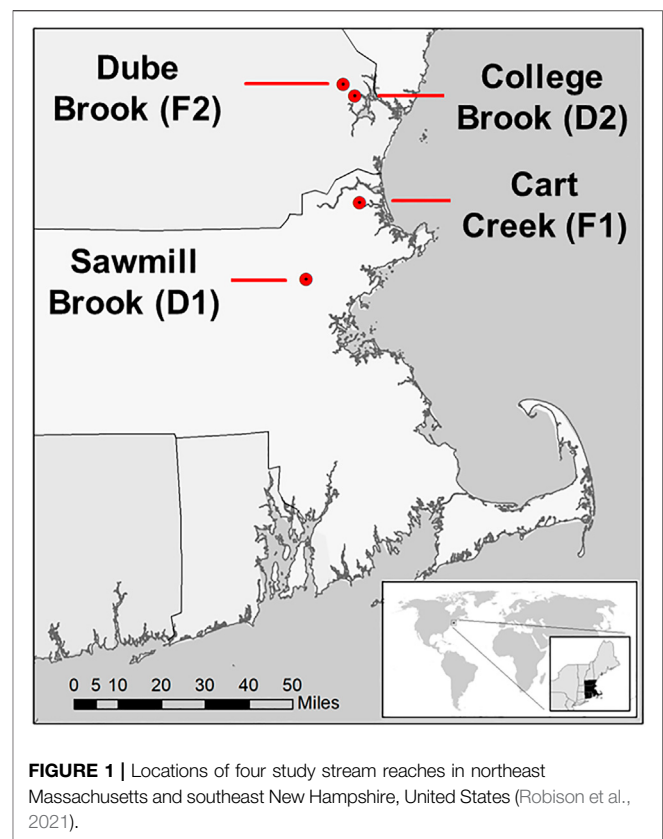


FIGURE 1 | Locations of four study stream reaches in northeast Massachusetts and southeast New Hampshire, United States (Robison et al., 2021).

CH_4 from a particular source (e.g., wetlands, landfills) can be used in atmospheric mixing models to constrain source contributions to bulk atmospheric CH_4 (Schwietzke et al., 2016). For example, CH_4 oxidation exerts a fractionation pattern of $\delta^{13}\text{C}\text{-CH}_4$ and $\delta\text{D}\text{-CH}_4$ enrichment compared to the source signature of the CH_4 (Chanton et al., 2005). While the exact fractionation can vary significantly (Conrad, 2005), the change in $\delta^{13}\text{C}\text{-CH}_4$ and $\delta\text{D}\text{-CH}_4$ values typically range from approximately 5–20‰ and 50–250‰, respectively (Whiticar, 1999; Wang et al., 2016). The emission pathway by which gaseous CH_4 leaves aquatic environment also has implications for the isotopic values of the emitted gas (Chanton, 2005). For example, Chanton et al. (1989) established that the $\delta^{13}\text{C}\text{-CH}_4$ values found within released bubbles were not significantly different from those of the reservoir of CH_4 held within the sediment. Meanwhile, CH_4 that diffuses through sediments and into the water column is more susceptible to oxidation (e.g., Shelley et al., 2017). Thus, the balance of CH_4 emission pathways can be important in determining the isotopic signature of a particular ecosystem (e.g., Sawakuchi et al., 2016) and how that ecosystem is included in atmospheric mixing models (Fisher et al., 2017; Saunois et al., 2020).

To properly understand the role of stream ecosystems in the global CH_4 cycle, there is a need to not only quantify CH_4 fluxes, but also investigate how CH_4 cycles through these systems, using isotopic and microbial analyses. In this paper, we 1) quantify the relative contributions of diffusive and ebullitive emissions of CH_4 from four headwater streams; 2) examine the $\delta^{13}\text{C}\text{-CH}_4$ and

TABLE 1 | Location, land use, and water quality characteristics of the four streams included in this study. Mean values of runoff, temperature, and stream chemistry are from the period of record.

		F1	F2	D1	D2
Physical descriptors	Latitude (°)	42.77	43.17	42.52	43.13
	Longitude (°)	−70.92	−70.97	−81.18	−70.92
	Area (km ²)	3.9	3.3	4.1	2.3
	Mean observed runoff (cm d ^{−1})	0.77	0.68	0.76	0.96
	Slope (m km ^{−1})	2.1	1.4	1.7	1.9
	Temperature (°C)	9.9	10.1	10.6	9.7
Land cover	Forest (%)	57.0	59.4	13.7	20.8
	Developed (%)	10.7	7.9	72.8	68.7
	Wetland (%)	18.7	17.3	4.3	0.7
Stream chemistry	Nitrate (mg L ^{−1})	0.1	0.2	0.8	0.7
	Chloride (mg L ^{−1})	101	62.5	190	321
	DOC (mg L ^{−1})	7.3	6.0	4.6	4.2

$\delta\text{D-CH}_4$ values of CH_4 associated with these two emission pathways; and 3) explore the CH_4 -associated microbial community in the stream sediments to provide context for CH_4 cycling. Integrating measurements of CH_4 emissions with an investigation of production and oxidation is needed to understand CH_4 cycling at the ecosystem level. By comparing our results to studies of other aquatic ecosystems such as larger rivers and lakes, we explore how CH_4 cycling in streams may differ and what implications this has for aquatic CH_4 budgets.

MATERIALS AND METHODS

Site Description

This study was conducted in four lowland headwater streams in southeastern New Hampshire and northwestern Massachusetts, United States (**Figure 1**). Watersheds were selected to contrast CH_4 dynamics in the two predominant land use conditions in this region (developed versus forested). These streams have been the focus of previous studies that examined nutrient and carbon cycling and the impact of land use (e.g., Wollheim et al., 2005; Wollheim et al., 2015; Wollheim et al., 2017), and were all monitored for CH_4 ebullition from May through October 2019 (Robison et al., 2021). The watersheds of two streams, Sawmill Brook (D1) and College Brook (D2), are characterized by a relatively developed, suburban landscape, while the other two, Cart Creek (F1) and Dube Brook (F2), are predominantly forest-covered. We group these streams into these two watershed land cover classes for comparison, but recognize the limitation of statistical inference with such a small sample size and mixed land use.

The watersheds range in size from 2.3 to 4.1 km², and all streams exhibit relatively shallow slopes (**Table 1**). Mean annual rainfall is 1,280 mm yr^{−1} and mean annual air temperature is 8.9°C (Wollheim et al., 2017). Mean discharge ranges from 25.5 L s^{−1} at D2 to 36.7 L s^{−1} at D1. Stream chemistry is measured as part of the Plum Island Ecosystems LTER project (Morse and Wollheim, 2014; Wollheim et al., 2015). Stream reach characteristics were examined in a concomitant study (Robison et al., 2021). Mean reach depth at baseflow ranged from 7.2 cm at

D2 to 14.5 cm at F1, and mean reach width at baseflow ranged from 1.94 m at D2 to 2.83 m at D1. All reaches exhibit shallow slopes of approximately 2 m km^{−1} or less. Benthic substrates included sand, silt, and fine organic matter, while rocks comprised less than 10% of benthic surface area across all sites. Macrophytes were absent from all reaches in this study, eliminating the possibility of CH_4 efflux via plant-mediated transport.

Dissolved CH_4 Sampling and Diffusive Efflux Estimation

Water samples for dissolved CH_4 analysis were collected at each stream using 60 ml syringes fitted with three-way stopcocks. Samples were collected at variable frequencies ranging from multiple samples a week to roughly weekly between June 1 and October 31, 2019. On average, a sample was collected every 6 days at each site. Syringes were rinsed with stream water prior to sample collection. To collect water samples, syringes were filled with approximately 60 ml of stream water from 5 to 10 cm depth below the stream water surface. Syringes were cleared of air bubbles by inverting and expelling bubbles and water until 30 ml of sample water remained. Samples were stored on ice until returned to the laboratory within 6 h. In the laboratory, 30 ml of ambient air was added to each syringe to achieve a 1:1 ratio of sample water to air. Syringes were then shaken for 2 min to equilibrate gases between water and headspace (Magen et al., 2014). The water was then dispelled from the syringe, and the remaining headspace gas was saved for analysis. If the gas samples were not analyzed immediately, they were stored in evacuated glass vials sealed with a rubber septum until analyzed.

Samples were analyzed for CH_4 concentration in the Trace Gas Biogeochemistry Laboratory at the University of New Hampshire. The CH_4 concentration in parts per million by volume (ppmv) was determined using a Shimadzu Gas Chromatograph Flame Ionization Detector. Concentration was standardized using the average area response of ten injections of a standard CH_4 mixture (Northeast Airgas, 2,006 ppmv or Maine Oxy, 1,000 ppmv) to determine instrument precision (Frolking

and Crill, 1994). If multiple samples were collected from a single site, the mean measured concentration was used.

Diffusive fluxes of CH₄ to the atmosphere (F_{CH_4} , mmol CH₄ m⁻² d⁻¹) were calculated as:

$$F_{CH_4} = k_{CH_4} \times \Delta CH_4, \quad (1)$$

where k_{CH_4} (m d⁻¹) is the gas transfer velocity for CH₄ and ΔCH_4 is the difference in CH₄ concentration (g m⁻³) between the water and the air corrected for Henry's Law. We used a constant air CH₄ concentration of 1.94 ppmv based on measurements at the nearby Global Monitoring Laboratory site at the Isle of Shoals, New Hampshire during the monitoring period in 2019 (Earth System Research Laboratory, G. M. D., 2021). The gas exchange rate for CH₄, k_{CH_4} , was calculated as:

$$k_{CH_4} = \left(\frac{SC_{CH_4}}{600} \right)^{1/2} / k_{600}, \quad (2)$$

where SC_{CH_4} is the Schmidt number for CH₄ at a given water temperature and k_{600} (m d⁻¹) is the gas transfer velocity standardized for a Schmidt number of 600. Without direct measurements of gas transfer velocity across a wide range of flow conditions, we estimated k_{600} for each reach based on the relationship following Raymond et al. (2012):

$$k_{600} = 1162 \pm 192 \times S^{0.77 \pm 0.028} \times V^{0.85 \pm 0.045}, \quad (3)$$

where V is the water velocity at time of sampling (m s⁻¹) and S is the channel slope (unitless). The mean slope of each stream reach was found using StreamStats (Ries et al., 2017). Mean daily discharge (Q ; L s⁻¹) was available from the long-term monitoring projects (Morse and Wollheim, 2014; Wollheim et al., 2017). Predictive relationships for V from Q were made using the equations for scaling stream geometry at a site (Knighton, 1998):

$$V = 0.287 \times Q^{0.4}. \quad (4)$$

Two or fewer direct measurements of the gas exchange velocity were also made at each site, although generally under relatively low flow conditions (< 50 L s⁻¹). These measurements were made using argon as a conservative gas tracer (Hall and Ulseth, 2020). We used these measurements to roughly evaluate the accuracy of our estimates of k_{600} (Supplementary Figure S1).

Ebullitive Sampling and Flux Estimation

Ebullitive CH₄ fluxes were measured during the same time frame as diffusive flux by Robison et al. (2021). Briefly, stationary bubble traps were deployed in triplicate at three to four locations in each stream. Traps were visited at least weekly from June 1 to October 31, 2019, during which the volume of gas collected in each trap was measured and collected for analysis of CH₄ concentration. The flux at each trap was then calculated as the mass of CH₄ emitted over the observation period normalized per unit area under the trap (mmol CH₄ m⁻² d⁻¹). For this study, mean rates of ebullitive CH₄ flux per measurement period at each stream were used, based on the total number of funnels at a stream.

Isotopic Sampling and Analysis

Gas samples were collected for CH₄ isotopic analysis in the first week of August 2019. Additional samples from F1 and D1 from the first week of September 2018 were also included in analysis. Dissolved CH₄ samples from the surface water were collected in the same manner described above. Additional benthic gas samples were collected by physically disturbing the sediment and collecting released bubbles. A minimum of three sediment gas samples and three surface water dissolved gas samples were collected at each site. All samples were stored in evacuated glass vials sealed with a rubber septum for analysis.

Samples were analyzed for $\delta^{13}C$ -CH₄ and δD -CH₄ using an Aerodyne dual tunable infrared laser direct absorption spectrometer (TILDAS; Aerodyne Research Inc, Massachusetts, United States) at the University of New Hampshire. These instruments use high resolution infrared spectrometry to quantify trace gases such as CH₄ (Mcmanus et al., 2011; Nelson and Roscioli, 2015). The TILDAS used here is configured with two 8 μ m quantum cascade lasers (QCLs, Alpes Lasers, Switzerland) and a 200 m multipass absorption cell to simultaneously monitor ¹²CH₄, ¹³CH₄, and CH₃D. The instrument was regularly calibrated with three standard tanks with known isotopic mixing ratios for ¹³CH₄/¹²CH₄ and CH₄/CH₃D. The isotopic composition of the standards was determined using an Aerodyne calibration system. The spectroscopic isotope ratios of four Isometric (now Airgas) CH₄ standards were measured at diluted concentrations ranging from < 1 to 12 ppmv. Keeling plot analysis was used to determine the relationship between the spectroscopic and standard isotope ratios of the Isometric standards. The linear relationship was made to the corresponding measured spectroscopic isotopic ratios of the UNH calibration tanks. The TILDAS is configured with an automated sampling system designed to measure small ($\leq$ 5 ml) injections of high concentration samples by diluting them within the instrument to a target CH₄ mixing ratio of eight ppmv with ultra-zero air. The instrument precision was 0.1‰ for $\delta^{13}C$ -CH₄ and 3‰ for δD -CH₃D at the target CH₄ mixing ratio. Samples with very low concentration (<0.01 mM CH₄) can result in erroneous isotopic measurement, and these datapoints (five of 59 samples) were removed prior to statistical analysis.

The percent of CH₄ oxidized between sediment generation and surface water evasion was estimated using the equation:

$$\% \text{ oxidized} = \frac{\delta^{13}C_{\text{Sample}} - \delta^{13}C_{\text{Source}}}{(\alpha - 1) \times 1000} \times 100, \quad (5)$$

where $\delta^{13}C_{\text{Sample}}$ is the $\delta^{13}C$ -CH₄ value found in the surface water sample, $\delta^{13}C_{\text{Source}}$ is the $\delta^{13}C$ -CH₄ value found in the sediment gas sample, and α is the isotope fractionation factor of CH₄ oxidation. We used two α values to include some uncertainty of isotopic fraction in these stream environments, 1.033 and 1.025 (Tyler et al., 1997; Zhang et al., 2016). This follows previous work estimating the efficiency of CH₄ oxidation in the Amazon River basin (Sawakuchi et al., 2016).

Microbial Sample Collection and Analysis

Sediment cores were collected from the streams for 16S rRNA analysis. A modified Multi Stage Soil Core Sampler (AMS Inc, Idaho, United States), consisting of a stainless steel cylinder and a plastic liner (5 cm diameter), was manually driven into the stream sediments using a sliding weight stand (Wik et al., 2018). Sediment cores were collected the first 2 weeks of July 2019 and processed in the field. Cores were collected at patches in the stream near each set of bubble traps as described in Robison et al. (2021). Four patches at each stream were sampled, except D2 where only three patches were sampled. At each patch, triplicate cores were collected. Sediment collected in the top 2 cm from each core was combined into a composite “surface” sediment sample in a Whirl-Pak sample bag. Similarly, sediment collected between 9 and 11 cm depth was combined into a composite “subsurface” sample. As a result, a total of four composite surface and four composite subsurface samples were collected at each stream except D2, where only three composite samples of each were collected. All tools used to collect cores, including the coring equipment, were sterilized with 70% ethanol between samples and sediment layers to minimize contamination. The depth of 9–11 cm was chosen as the subsurface sample depth because buried rocks or clay layers prevented collection of sediment at depths greater than 12–15 cm in some places. Thus, while 9–11 cm does not represent the maximum or mean depth of the sediment in portions of these streams, it represents a depth layer that could be collected across all sites and is deep enough where complete depletion of oxygen is likely (Crawford et al., 2014). Sediment samples were frozen on-site with liquid nitrogen, stored on ice, and returned to the laboratory within 4 h. Samples were stored in a –80°C freezer until analyzed.

Genomic DNA was extracted from sediment samples using a Qiagen DNeasy PowerSoil kit (Qiagen, Hilden, Germany) with minor changes to the manufacturer’s protocol (see Doherty et al., 2020). DNA was amplified by polymerase chain reaction (PCR) using the primers 515f–806r of the V4 region of the 16S rRNA gene to profile bacterial and archaeal communities (Apprill et al., 2015; Parada et al., 2016). Each reaction contained 6 µl DreamTaq Hot Start Green (Thermo Fisher Scientific, Waltham, MA, United States), 2.6 µl sterile water, 0.7 µl forward primer (5 µM), 0.7 µl reverse primer (5 µM), and 2 µl template DNA (10× diluted). DNA amplification was performed using a T100 Thermal Cycler (Bio-Rad, Hercules, CA, United States). The 16S rRNA conditions were: enzyme activation at 95°C for 3 min; followed by 35 cycles of denaturation at 95 °C for 30 s, annealing at 55°C for 30 s, and extension at 72°C for 60 s; then a final extension at 72°C for 12 min.

Gel electrophoresis was used to confirm the presence of the PCR product, which was then quantified using a Quant-iT dsDNA High-Sensitivity Assay Kit and a Qubit 3.0 fluorometer (Thermo Fisher Scientific, Massachusetts, United States). Resulting concentrations ranged from 0.5 to 28 ng µL^{–1}, with sandier sediment samples typically having lower concentrations. PCR products were sent to the Hubbard Center for Genome Studies (University of New Hampshire, New Hampshire, United States) for sequencing by Illumina HiSeq2500 (250 bp reads) with Rapid Run[®] SBS V2 chemistries (Illumina,

San Diego, CA, United States). Sequence reads were demultiplexed using CASAVA.

Sequences were analyzed using QIIME 2 (version 2020.1; Bolyen et al., 2019) on the Premise high performance computing cluster at the University of New Hampshire. Cutadapt was used to remove primers (Martin, 2011). Using DADA2 (Callahan et al., 2016), sequences were truncated at the length where the median Phred score (quality score) fell below 30 (200 bp for forward read; 225 bp for reverse read) and unique sequences were designated as amplicon sequence variants (ASVs). Each sample was then rarefied to 3,400 sequences per sample. Rarefaction depths were chosen to ensure at least three samples remained for each depth at each stream. Taxonomy was assigned to ASVs using the SILVA database (release SILVA 138 SSU; Quast et al., 2013; Glöckner et al., 2017). The relative abundances of taxa were calculated after removing chloroplasts, mitochondria, and taxa present in less than 5% of samples.

In this study, we focused primarily on taxa associated with CH₄ production and CH₄ oxidation. Following Hough et al. (2020), we identified methanogens based on classification into the archaeal orders: Methanocellales, Methanobacteriales, and Methanomicrobiales, and the families Methanosarcinaceae and Methanosaetaceae, of order Methanosarcinales (Evans et al., 2019). Similarly, we identified methanotrophs as those from the bacterial orders Methylococcales and Methylospirales and the archaeal family Methanoperedenaceae, order Methanosarcinales (Smith and Wrighton, 2019). Taxa from the order Methylococcales are classical aerobic methanotrophs, while those from the order Methylospirales are capable of CH₄ oxidation following reduction of nitrate or nitrite (Wu et al., 2011); that is, they are capable of generating oxygen for CH₄ oxidation when in an anoxic environment. Finally, Methanoperedenaceae are a methanotrophic archaea family (formerly known as ANME-2d) known to be capable of CH₄ oxidation via alternative electron donors like nitrate, iron, or manganese (Ettwig et al., 2016; Leu et al., 2020). Despite the indefinite nature of assigning organismal function using 16S sequencing, traits associated with CH₄ cycling are typically conserved and well defined (Martiny et al., 2015), allowing for confidence in these assignments. The relative abundance of individual samples is calculated, from which a mean and standard deviation from all samples collected at a sediment depth in a stream is determined.

Statistical Analyses

All calculations and statistical analyses were performed in MATLAB and Statistics Toolbox Release 2020a (The MathWorks, Inc, Massachusetts, United States), and the significant level was set at $\alpha = 0.05$. Uncertainty in the dissolved CH₄ concentration for each site was determined by the ratio of the standard error (SE) of the measured concentrations to the median concentration (Hojo and Pearson, 1931):

$$\% \text{ uncertainty} = 1.253 \times \frac{\text{Std Err}}{\text{median CH}_4} \times 100, \quad (6)$$

TABLE 2 | Summary of measured dissolved methane concentration, estimated methane-specific gas exchange rate, calculated diffusive and ebullitive emissions, measured $\delta^{13}\text{C}\text{-CH}_4$ and $\delta\text{D}\text{-CH}_4$ values, estimated percent of the dissolved CH_4 oxidized before emission, and the percent abundance of methanogens and methane oxidizing bacteria in the total detected community for the four streams in this study. SD indicates the standard deviation. *Ebullitive methane flux data is from Robison et al. (2021). Methanotrophs refers to taxa capable of CH_4 oxidation, whether aerobic or anaerobic.

Variable			F1	F2	S1	S2	Overall
Dissolved CH_4 concentration ($\mu\text{mol CH}_4 \text{ L}^{-1}$)	Median		0.9	1.6	1.0	1.6	1.1
	Min		0.3	0.8	0.3	0.2	0.2
	Max		13.6	8.5	2.7	16.1	16.1
k_{CH_4} (m d^{-1})	Median		4.7	3.4	5.9	4.7	4.8
	Min		4.1	2.2	4.5	4.0	2.2
	Max		8.2	8.3	10.3	11.1	11.1
Diffusive CH_4 flux ($\text{mmol CH}_4 \text{ m}^{-2} \text{ d}^{-1}$)	Median		4.1	4.7	5.9	7.8	5.7
	Min		1.4	2.2	1.6	1.3	1.3
	Max		64.9	25.7	20.8	70.8	70.8
Ebullitive CH_4 flux* ($\text{mmol CH}_4 \text{ m}^{-2} \text{ d}^{-1}$)	Median		0.2	1.3	1.0	0.0	0.6
	Min		0.0	0.2	0.01	0.0	0.0
	Max		0.8	4.4	13.5	0.1	13.5
$\delta^{13}\text{CCH}_4$ (‰)	Benthic bubble	Mean	-67.8	-66.5	-63.1	-66.0	-65.9
		SD	0.5	0.4	0.6	0.4	0.5
	Surface dissolved	Mean	-51.5	-52.8	-49.5	-55.0	-52
		SD	1.7	0.7	1.1	2.6	1.5
	Benthic bubble	Mean	-297	-274	-312	-318	-300
		SD	17	16	21	16	18
δDCH_4 (‰)	Surface dissolved	Mean	-231	-257	-290	-265	-261
		SD	21	26	42	28	29
	Benthic bubble	Mean	65.2	54.8	54.4	44.0	55.6
		SD	3.7	2.5	2.6	2.4	3.2
	$\alpha = 1.025$	Mean	49.4	41.5	41.2	33.3	42.1
		SD	2.7	1.8	2.0	1.9	2.1
% CH_4 oxidized	$\alpha = 1.033$	Mean	65.2	54.8	54.4	44.0	55.6
	$\alpha = 1.025$	Mean	49.4	41.5	41.2	33.3	42.1
Methanogens (% abundance)		Mean (0 cm)	0.1	0.2	0.1	0	0.1
		Mean (10 cm)	0.8	0.7	0.3	0.3	0.6
Methanotrophs (% abundance)		Mean (0 cm)	1.0	3.2	5.0	4.5	3.5
		Mean (10 cm)	1.6	1.2	4.7	2.5	2.5

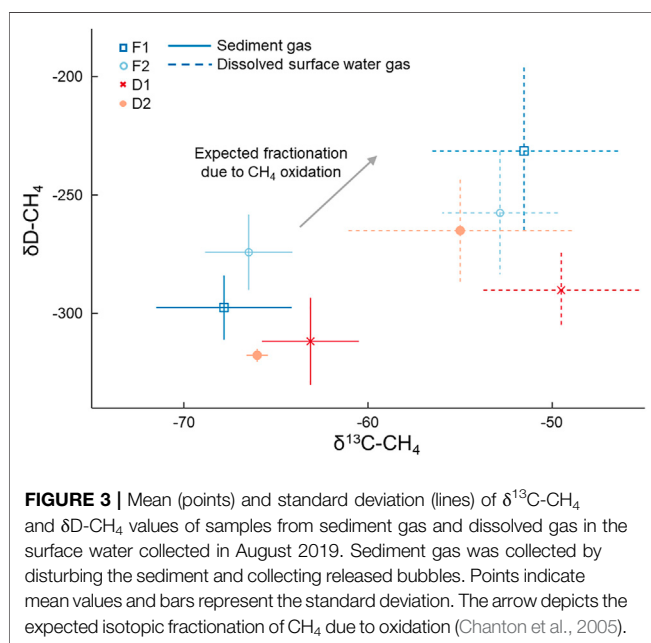
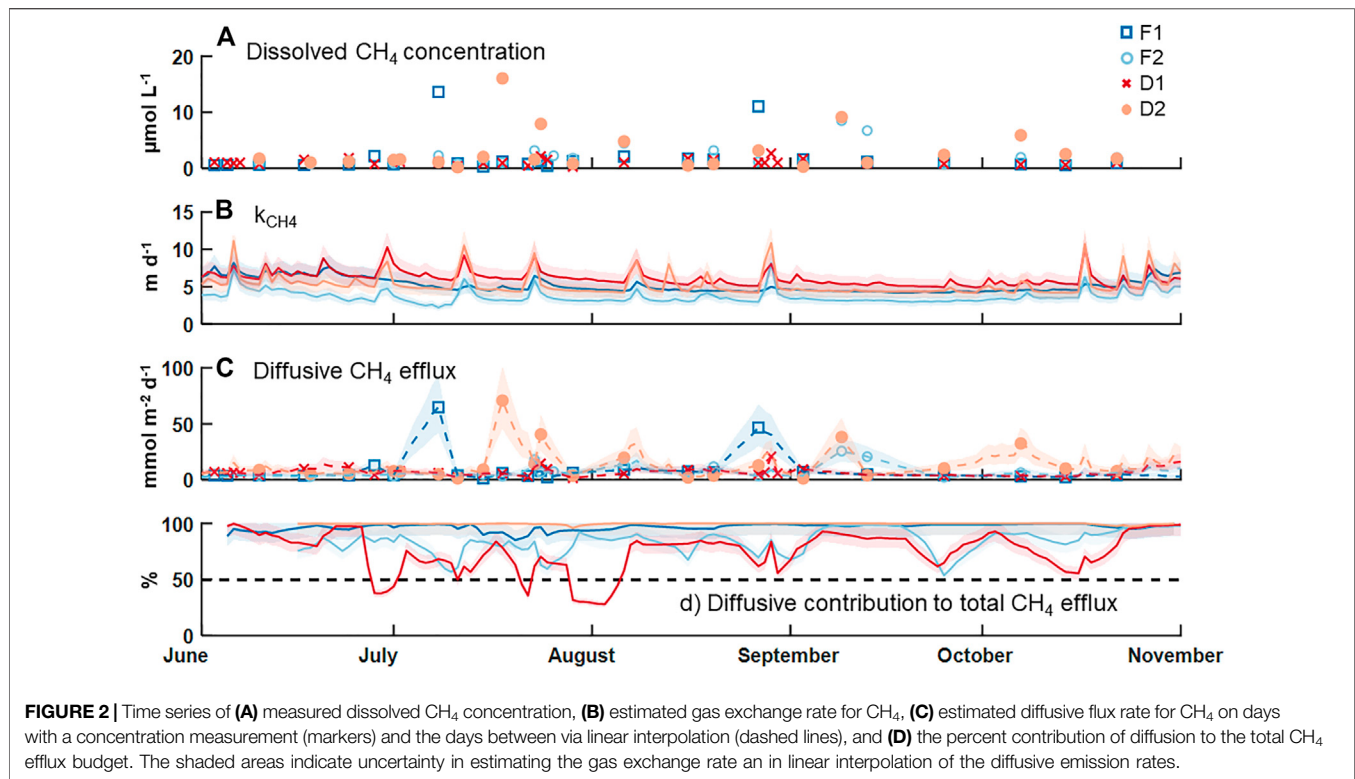
which resulted in a unique percent uncertainty at each site. Uncertainty in the gas exchange rate was determined by the standard deviation included in the factors of Eq. 3. Simple linear interpolation was used to estimate the rate of diffusive efflux on days without a dissolved CH_4 concentration measurement. Uncertainty in the diffusive CH_4 flux was then estimated by propagating the uncertainty from dissolved CH_4 concentration and gas exchange rates. We summarized CH_4 concentration, gas exchange, and rates of diffusion as medians because of the non-normal distribution of the data. Differences in CH_4 concentration, gas exchange, and diffusive emissions between streams were analyzed using the nonparametric Wilcoxon rank sum test. Trends in CH_4 concentration and diffusion across time were analyzed using the nonparametric Kendall rank correlation coefficient. Isotopic and microbial data were summarized as means because of their small sample sizes and unknown distributions. Differences between streams, depths for microbial samples, and type of sample for isotopes (i.e., benthic or surface water) were compared using two-sample *t*-test or one-way ANOVA.

RESULTS

Dissolved CH_4 Concentration and Diffusive Fluxes

Measured concentrations of dissolved CH_4 in surface water were above saturation (roughly $0.003 \mu\text{mol CH}_4 \text{ L}^{-1}$ at 10°C) in all

collected samples, with an overall median $\pm\text{SE}$ of $1.1 \pm 0.4 \mu\text{mol CH}_4 \text{ L}^{-1}$ (Table 2; Figure 2A). F2 and D2 had the highest median concentration of CH_4 ($1.6 \pm 0.9 \mu\text{mol CH}_4 \text{ L}^{-1}$) and F1 had the lowest ($0.9 \pm 0.3 \mu\text{mol CH}_4 \text{ L}^{-1}$). There was no statistical difference in the median dissolved concentration of CH_4 between watershed land cover classes ($p > 0.05$). We also observed no clear pattern in the dissolved concentration across time at any individual stream ($p = 0.54$). The median k_{CH_4} across all sites was $4.8 \pm 0.4 \text{ m d}^{-1}$, ranging from $3.4 \pm 0.3 \text{ m d}^{-1}$ at F2 to $5.9 \pm 0.6 \text{ m d}^{-1}$ at D1 (Table 2; Figure 2B). The watershed land cover classes did differ in the estimated gas exchange rates ($p < 0.01$), where the two streams draining developed landscapes, D1 and D2, exhibited significantly higher k_{CH_4} . Calculated rates of gas exchange were similar to limited direct measurements in each of the streams (Supplementary Figure S1) using short-term continuous injections of a volatile gas tracer (Hall and Ulseth, 2020). Median diffusive CH_4 flux was $5.7 \pm 2.3 \text{ mmol CH}_4 \text{ m}^{-2} \text{ d}^{-1}$ overall (Table 2; Figure 2C). In individual streams, the median diffusive flux ranged from 4.1 ± 2.1 to $7.8 \pm 4.8 \text{ mmol CH}_4 \text{ m}^{-2} \text{ d}^{-1}$ at F1 and D2, respectively, with the developed watershed class exhibiting a higher rate compared to the forested watershed class ($p < 0.01$). We did not observe any clear temporal pattern in diffusive CH_4 emissions across our monitoring period at any individual site ($p > 0.05$). High diffusive CH_4 fluxes were more strongly associated with high measurements of dissolved CH_4 than with high gas exchange



rates (Figure 2). The maximum daily rate of diffusive CH₄ efflux was $70.8 \pm 31.4 \text{ mmol CH}_4 \text{ m}^{-2} \text{ d}^{-1}$ at D2.

Diffusive fluxes of CH₄ were compared to ebullitive fluxes of CH₄ from Robison et al. (2021) during the same period of record (Table 2; Figure 2D). Across all four streams, the median diffusive CH₄ flux ($5.7 \pm 2.3 \text{ mmol CH}_4 \text{ m}^{-2} \text{ d}^{-1}$) was greater

than the median ebullitive CH₄ flux ($0.6 \pm 0.1 \text{ mmol CH}_4 \text{ m}^{-2} \text{ d}^{-1}$). Diffusive emissions of CH₄ comprised approximately 90% of total CH₄ emissions from these four streams over the entire period, ranging from 78% at F2 to nearly 100% at D2. Ebullitive CH₄ flux exceeded diffusive flux only at D1 for short periods (Figure 2D). Diffusion accounted for the majority of CH₄ emissions at the other three sites during the entire monitoring period.

Stable Isotopic Composition of CH₄

Benthic gas and surface water gas samples exhibited clear differences in $\delta^{13}\text{C}-\text{CH}_4$ and $\delta\text{D}-\text{CH}_4$ values (Table 2; Figure 3). Generally, there was more variability between the type of sample (i.e., benthic or surface water) than between streams. Benthic gas $\delta^{13}\text{C}-\text{CH}_4$ and $\delta\text{D}-\text{CH}_4$ values averaged across all streams were $-65.9 \pm 2.0\text{‰}$ and $-300 \pm 18\text{‰}$, respectively. There was some variation in the benthic $\delta^{13}\text{C}-\text{CH}_4$ values between streams, where D1 exhibited a significantly higher mean $\delta^{13}\text{C}-\text{CH}_4$ ($-63.1 \pm 0.6\text{‰}$) than F1 ($-67.8 \pm 0.5\text{‰}$, $p < 0.01$) and F2 ($-66.5 \pm 0.4\text{‰}$, $p = 0.02$). More variation was noted in measured $\delta\text{D}-\text{CH}_4$ values, with F2 having the highest mean value of $-257 \pm 16\text{‰}$ and D2 having the lowest value of $-318 \pm 16\text{‰}$, with only these two sites being significantly different ($p = 0.01$). Dissolved CH₄ $\delta^{13}\text{C}-\text{CH}_4$ and $\delta\text{D}-\text{CH}_4$ values averaged across all four streams were $-52 \pm 1.5\text{‰}$ and $-261 \pm 29\text{‰}$, respectively. These were slightly more varied across streams than the benthic samples. For example, $\delta^{13}\text{C}-\text{CH}_4$ values ranged from $-49.5 \pm 0.6\text{‰}$ at D1 to $-55.0 \pm 2.6\text{‰}$ at D2 in dissolved CH₄ samples, although no site was significantly different from the others. Meanwhile, dissolved $\delta\text{D}-\text{CH}_4$ ranged

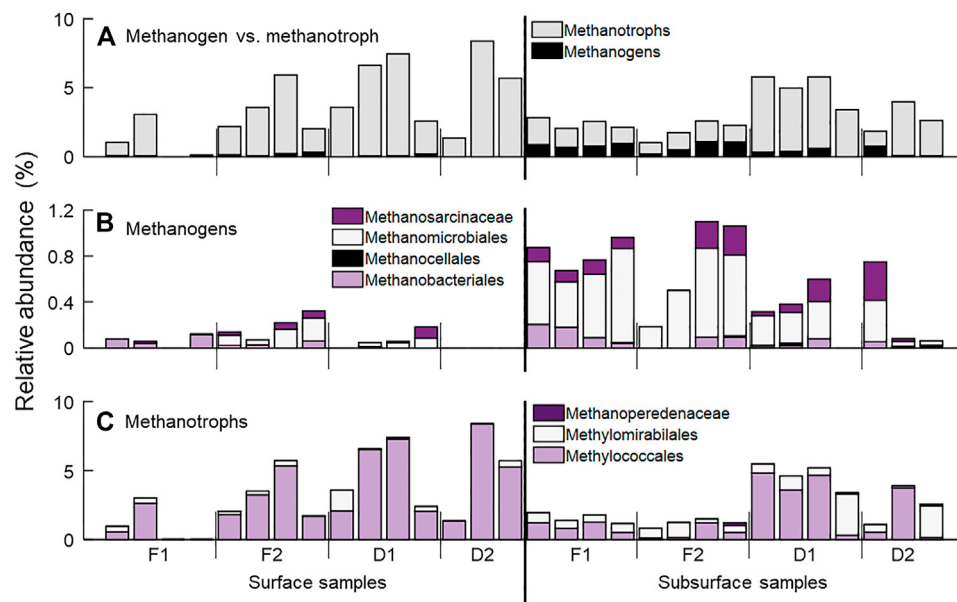


FIGURE 4 | Relative abundance of CH_4 -associated taxa at sampling locations. **(A)** All taxa associated with CH_4 production (methanogens, black) and oxidation (methanotrophs, gray). Here, we use the simplified term methanotrophs to refer to those taxa who are capable of CH_4 oxidation through any of a variety of biochemical pathways. **(B)** Archaeal taxa identified as methanogens detected in sediment samples, including the orders Methanomicrobiales, Methanocellales and Methanobacteriales, and the families Methanosarcinales and Methanosaetaceae, both of the order Methanosarcinales. **(C)** Taxa associated with CH_4 oxidation, separated into categories based on potential biochemical pathways. The bacterial order Methylococcales are canonical, aerobic CH_4 oxidizing bacteria. The bacterial order Methyloirabiales are proposed to be capable of CH_4 oxidation following the reduction of nitrite. Finally, methanotrophic archaea from the order Methanosarcinales, family Methanoperedenaceae (formerly known as ANME-2d) are also capable of CH_4 oxidation following the reduction of nitrate, iron, or manganese.

from $-257 \pm 26\text{‰}$ at F2 to $-290 \pm 42\text{‰}$ at D1, with only these two sites being significantly different ($p = 0.04$). There was no difference in measured isotopic values between watershed land cover classes when comparing either $\delta^{13}\text{C}\text{-CH}_4$ or $\delta\text{D}\text{-CH}_4$ between benthic gas and dissolved samples.

Dissolved CH_4 in surface water was more ^{13}C - and D-enriched relative to the sediment gas overall ($p < 0.01$; **Figure 3**). This amounted to a $14 \pm 1.5\text{‰}$ and $39 \pm 29\text{‰}$ shift towards the heavier isotope for $\delta^{13}\text{C}\text{-CH}_4$ and $\delta\text{D}\text{-CH}_4$, respectively. The $\delta^{13}\text{C}\text{-CH}_4$ values for dissolved CH_4 were all significantly higher than sediment CH_4 at each stream, and the $\delta\text{D}\text{-CH}_4$ values were significantly higher than sediment CH_4 at F1 ($p = 0.01$) and D1 ($p = 0.04$). Given this enrichment in $\delta^{13}\text{C}\text{-CH}_4$ values, we estimated the mean proportion of dissolved CH_4 oxidized was $42.1 \pm 2.1\%$ when using $\alpha = 1.033$, and $55.6 \pm 2.8\%$ when using $\alpha = 1.025$ (**Table 2**). The percentage of dissolved CH_4 oxidized in individual streams ranged from a minimum of $33.3 \pm 3.1\%$ at D2 using $\alpha = 1.025$ to $65.2 \pm 2.6\%$ at D1 given $\alpha = 1.033$.

Methane-Associated Microbial Community

Taxa identified as methanogens and methanotrophs were detected across all four streams (**Figure 4A**, **Supplementary Table S1**). Methanogen taxa were detected in 10 of 15 surface sediment samples, and 14 of 15 subsurface samples (**Figure 4B**). These included the archaeal orders Methanosarcinales (families Methanosaetaceae and Methanosarcinaceae), Methanobacteriales (family Methanobacteriaceae), Methanomicrobiales (families

Methanomicrobiaceae, Methanoregulaceae, and rice cluster II), and Methanocellales (family Methanocellaceae). As a percent of the total community, methanogens comprised roughly 0.1% of the surface sediment community and 0.6% of the subsurface community, on average. Methanogens comprised a larger percentage of the microbial community in the forested watershed streams than in the developed streams ($p = 0.04$).

Taxa classified as methanotrophs were detected in 14 of 15 surface samples and all subsurface samples (**Figure 4C**). Taxa from the bacterial orders Methylococcales (families Methylococcaceae and Methyloirabiales), Methyloirabiales (family Methyloirabillaceae), as well as the archaeal order Methanosarcinales (family Methanoperedenaceae) were detected. Methanotrophs comprised a larger fraction of the total microbial community in all samples compared to methanogens, but this may be a result in part of the bias of our primers for bacteria over archaea (Walters et al., 2016). As a percent of the total community, methanotrophs comprised on average 4.3 and 3.6% in surface and subsurface samples, respectively. Aerobic methanotrophs, i.e., Methylococcales, were significantly more abundant in surface sediment samples ($p = 0.02$), while taxa capable of CH_4 oxidation in anoxic conditions (Methyloirabiales and Methanoperedenaceae) were higher in relative abundance in subsurface samples than in surface samples ($p = 0.02$). Additionally, methanotroph relative abundance was greater in the two streams draining developed watersheds than the two draining forested watersheds ($p < 0.01$).

DISCUSSION

Diffusion via turbulent mixing was the dominant pathway of CH₄ emissions from the stream reaches in this study (**Figure 2D**; **Table 2**). This appears more common in small streams compared to larger rivers or lentic ecosystems, where ebullition tends to be more important. The enrichment of CH₄ isotopic values from the benthos to the surface water is likely a result of oxidation, which is supported by the presence of a diverse microbial community capable of CH₄ oxidation. Emitted CH₄ from these streams thus exhibits isotopic values more characteristic of the dissolved CH₄ pool, i.e., heavier than the benthic gas pool. Together, our results suggest CH₄ emission and isotopic patterns in small streams may be unique amongst aquatic ecosystems, and thus may distinguish how small streams should be included in global CH₄ models.

Diffusive Emissions Dominate CH₄ Efflux Budget

Diffusive emissions were the dominant efflux pathway of CH₄ in these four lowland stream reaches. The median diffusive flux is comparable to a summary of 385 streams and rivers (8.22 ± 25.50 mmol CH₄ m⁻² d⁻¹; Stanley et al., 2016). Total CH₄ emissions were greater than the median of a recent analysis of streams and rivers globally (0.9 mmol CH₄ m⁻² d⁻¹), but within the 90% range (0.0 – 35.8 mmol CH₄ m⁻² d⁻¹; Rosentreter et al., 2021). When compared to lentic waterbodies, the rate of emissions from these four streams is similar on a per area basis (e.g., Holgerson and Raymond, 2016; Sanches et al., 2019). For example, in a variety of lakes and ponds of northern latitudes, total CH₄ emissions ranged from 2.8 to 12.5 mmol CH₄ m⁻² d⁻¹ (Wik et al., 2016b).

The efflux of CH₄ from small streams may be a significant, yet overlooked pathway of carbon fluxes from inland waters. The total aquatic carbon flux for inland waters across North America is estimated to be 24 g C m⁻² yr⁻¹ (Butman et al., 2018), where the flux is normalized for total land area. However, this flux does not include CH₄, as scaling was limited by significant data gaps at the time of estimation. Following scaling methods published previously (Robison et al., 2021), the streams in this study emit roughly 0.08 g CH₄-C m² watershed area yr⁻¹ when both diffusion and ebullition are considered, less than 1% of the total estimated for inland waters. If the greater radiative forcing of CH₄ relative to CO₂ is taken into account (Saunio et al., 2020), the CO₂-equivalent rate increases to 2.2 g C m² watershed area yr⁻¹. Thus, ignoring small stream methane emissions may underestimate watershed carbon emissions by approximately 10%, a significant factor.

Relatively few studies exist with which to compare diffusion and ebullition in lotic systems (**Supplementary Table S2**). Summarizing 26 streams and rivers which considered both pathways, diffusion delivered 68% of total CH₄ emissions on average, but this ranged from 14 to 94% (Stanley et al., 2016). However, it appears smaller streams tend to favor diffusive emissions. For example, diffusion comprised roughly 90% of CH₄ emissions in a small stream in Wisconsin, United States (Crawford et al., 2014). In contrast, studies of larger rivers in the

Amazon (Sawakuchi et al., 2014), on the East Qinghai-Tibet Plateau (Zhang et al., 2020), and in urban China (Wang et al., 2021) found diffusion accounted for 64, 21, and 20% of total CH₄ emissions, respectively. Diffusive fluxes generally account for a smaller fraction of the total CH₄ budget in lentic systems too, e.g., 48% for northern lakes and ponds (Wik et al., 2016b).

Land use may affect diffusive CH₄ emissions, where development promotes higher rates of emission. This pattern follows previous studies which linked low oxygen and high organic carbon delivery to higher rates of CH₄ production in urban streams (Stanley et al., 2016; Wang et al., 2021). With only four streams studied and mixed land use within each watershed, the strength of this comparison is narrow. Individual samples from each stream exhibited relatively high concentrations of dissolved CH₄. The median and maximum of dissolved CH₄ concentrations were similar to those found in a yearlong survey of gas concentrations in streams in New Hampshire (0.63 and 13.5 μmol CH₄ L⁻¹; Herreid et al., 2020). Our dataset precludes analysis of what causes these high concentrations, but we have no reason to discard them from our analysis as they fall within ranges observed in streams and rivers locally and globally (Stanley et al., 2016).

The estimation of diffusive CH₄ efflux is limited by the relatively sparse measurements of dissolved CH₄ concentrations and the estimation of gas exchange. The similarity of direct gas exchange measurements to estimates used in this study suggests this latter uncertainty is modest during low flow conditions (**Supplementary Figure S1**). Similarly, while our results likely suffer in accuracy at fine temporal scales (i.e., daily), it does not necessarily indicate the summarized results are inaccurate. Uncertainty due to sampling frequency on estimating seasonal rates of CH₄ emissions in lakes has been examined previously (Wik et al., 2016a); here, a minimum of 17 measurements of dissolved CH₄ concentration were needed to accurately represent seasonal emission patterns from lakes. The minimum number of measurements of dissolved CH₄ concentration in this study is 25. While the standard for lakes may not be the same for lotic ecosystems, we believe our results reflect the typical seasonal behavior at these streams in which diffusion is the dominant emission pathway for CH₄.

Evidence for In-Stream CH₄ Production and Oxidation From Isotopic and Microbial Data

The isotopic and microbial results provide evidence for active CH₄ processing in these streams, with production and oxidation ubiquitous. For example, the presence of ebullition and the detection of methanogens suggests active CH₄ production in the sediment of all four streams. The δ¹³C-CH₄ and δD-CH₄ values of the sediment-derived gas suggest a combination of acetoclastic (-60% to -50% δ¹³C-CH₄ and -400% to -250% δD-CH₄) and hydrogenotrophic (-110% to -60% δ¹³C-CH₄ and -250% to -170% δD-CH₄) methanogenesis (Whiticar, 1999; Chanton et al., 2005), which is common in aquatic environments (Shelley et al., 2015; Wik et al., 2020). The presence of methanotrophs and an enrichment in both δ¹³C-CH₄ and δD-CH₄ values between benthic and surface samples suggests

CH₄ oxidation was prevalent across the four streams as well, representing a potentially significant control on emissions. The estimated oxidation proportions are similar, but generally lower than the 57–82% range estimated for large rivers in the Amazon River basin (Sawakuchi et al., 2016) or the 57–100% estimated for lakes in Sweden (Bastviken et al., 2002). The lower oxidation proportion in the streams could be a result of relatively faster turnover of the water column. That is, while the efficiency of CH₄ oxidation could be similar across all aquatic ecosystems, relative fast mixing in small streams may limit the time for CH₄ oxidation to occur. Regarding microbial evidence for CH₄ production and oxidation, detection of microbial taxa via 16S rRNA sequencing can recover dormant or dead microbes (Oliver, 2005), thus the detections do not directly indicate activity. However, combined with the magnitude of CH₄ fluxes and isotopic values, we can confidently infer the presence of microbial CH₄ production and oxidation in the streams.

Alternative explanations for the change in isotopic values are possible. The difference could result from dissolved CH₄ inputs from a different source in the watershed, e.g., the riparian zone (Crawford et al., 2013). If the balance of isotopic values of this other source is distinct from that in the stream sediments, the isotopic values of the dissolved CH₄ in the streams may be confounded by the mixed sources. Additionally, diffusion exerts a fractionation effect ($< 3\text{‰ } \delta^{13}\text{C-CH}_4$) on CH₄, where lighter CH₄ is preferentially emitted (Knox et al., 1992). This fractionation is much smaller than that of oxidation and the time for diffusion is relative fast in these streams, limiting the impact of this effect. Because isotopic and microbial sampling occurred in August and July, respectively, the results of our oxidation analysis may reflect the conditions of this warmer period of the year. For example, CH₄ isotopes in aquatic ecosystems may be heaviest in the summer (Atkins et al., 2017), possibly as a result of more active microbial oxidation (Shelley et al., 2015). If so, this would indicate we may have measured conditions under which oxidation is most efficient. However, considering the isotopic results in the context of the detection of methane oxidizing microbes, we are confident CH₄ oxidation is prevalent in these streams with the greatest uncertainty around the temporal dynamics of oxidation efficiency.

Mechanisms Driving CH₄ Emission Pathway and Oxidation Potential in Headwater Streams

The dominance of diffusive CH₄ emissions and the relatively lower proportion of CH₄ oxidation appear to distinguish small streams from other aquatic ecosystems. Identifying the factors that drive this difference is thus key to understanding how CH₄ dynamics may differ in small streams. We propose two characteristics that are fundamental in differentiating small streams from larger rivers and from lentic systems: flow and depth. We hypothesize limited bubble formation and CH₄ oxidation results from the unique physical and biogeochemical conditions of small streams caused by increased water exchange between the water column and sediments (hyporheic flow) and short residence time of CH₄ in the water column (gas exchange).

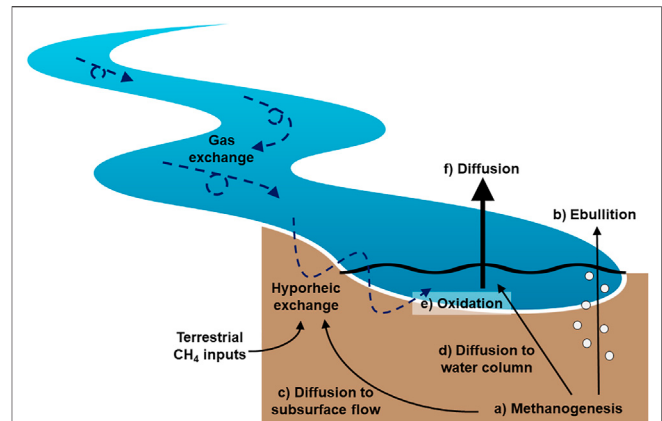


FIGURE 5 | Proposed conceptual model of CH₄ production, transport, oxidation, and emissions in stream ecosystems. The inherent flow and shallow nature of streams promotes hyporheic exchange and high gas exchange rates. This serves to limit bubble formation and promote the transport of dissolved CH₄ out of the sediments *via* diffusion and subsurface flow. While the dissolved CH₄ pool is exposed to oxidation, the proportion of CH₄ that is oxidized is limited by the relatively rapid exchange of water and gases in stream ecosystems. Most CH₄ is emitted *via* diffusion across the water-air interface in streams, and the limited exposure of this CH₄ pool results in relatively heavy dissolved CH₄ isotopic values. Because diffusive CH₄ emissions dominate the overall emission budget, the mean isotopic value of emitted CH₄ reflects this heavier signature and CH₄ isotopes in streams are enriched relative to other aquatic ecosystems.

Flow provides lotic ecosystems with consistent turbulent mixing, and the relatively shallow nature of small streams propagates this turbulence to the sediment–water interface. These likely limit bubble formation broadly *via* changes to sediment quality and depth. Because bubble formation is greatly affected by the grain size and physical structure of aquatic sediments (Liu et al., 2016), processes like turbulence which affect the sediment will in turn affect ebullition. For example, the erosional nature of streams (Knighton, 1998) may inhibit ebullition by increasing mean grain size or reducing deposition of organic matter, both of which have been shown to limit ebullition in streams (Crawford et al., 2014; Bodmer et al., 2020). Turbulence also enhances exchange at both the sediment–water and air–water interface within streams (Boano et al., 2014), affecting both bubble formation and oxidation rates. Advective water inputs into and through stream sediments *via* hyporheic flow (Packman and Salehin, 2003) create patches of oxic and anoxic conditions in close proximity (MahmoodPoor Dehkordy et al., 2019; Nelson et al., 2019). The CH₄ produced in anoxic zones can be transported through sediments by this advective flow (Sobczak and Findlay, 2002), limiting bubble formation further, exposing CH₄ to oxidizing conditions, and decreasing the residence time of CH₄ within the sediment. This advective exchange may ultimately limit the overall potential for CH₄ oxidation. For example, in Amazonian rivers, the extent of CH₄ oxidation was positively related to residence time in surface sediments (Sawakuchi et al., 2016). Oxidation within the water column is also limited by higher rates of gas exchange in streams and rivers

compared to lentic ecosystems (Looman et al., 2021), as does the shallower depth of streams compared to larger rivers (Raymond et al., 2012). For example, 51–100% of the CH_4 in lakes originating in sediments in deeper portions of lakes was oxidized in the water column, while only 24–40% of the CH_4 from shallower areas was oxidized due to the relatively shorter transit time (Bastviken et al., 2008).

We propose a conceptual model of CH_4 production, transport, oxidation, and emission unique to small streams, where CH_4 is emitted primarily through diffusive fluxes and oxidation is relatively limited (Figure 5). In this model, 1) CH_4 is primarily produced by methanogens in anoxic sediments. While bubble formation is generally limited in small streams, 2) some CH_4 is emitted *via* ebullition. The bulk of CH_4 production, plus inputs from the upslope catchment via groundwater flow, remains dissolved. In small streams, dissolved CH_4 3) diffuses through the sediments entering hyporheic flow paths, 4) although small amounts of vertical diffusion through the sediments may also occur. Hyporheic flow, which remains primarily anoxic in many systems, moves dissolved CH_4 back to the water column. We propose 5) oxidation primarily occurs near the sediment–water interface, where the redox gradient is greatest and the highest abundance of methanotrophs are detected. In the surface water, CH_4 oxidation is likely limited due to relatively high gas exchange rates. After oxidation, 6) the remaining dissolved CH_4 emitted via diffusion is enriched isotopically relative to that emitted via ebullition. The relative importance of each of these processes will depend on stream characteristics such as slope, water velocity, water depth, and sediment characteristics (i.e., depth, grain size) that influence the degree of hyporheic exchange, hyporheic residence time, redox conditions, and microbial activity. Some of the proposed mechanisms described in this conceptual model were not directly measured (e.g., hyporheic exchange) and require further exploration and testing. We specifically urge the examination of CH_4 cycling and transport within the benthic environment in relation to subsurface flow and redox conditions.

Lotic Ecosystems May Be Isotopically Enriched Relative to Lentic Systems

One important implication of this conceptual model is how streams are considered in global CH_4 models (Dean et al., 2018), particularly with regards to the isotopic values of emitted CH_4 . For example, given both the imbalance of emission pathways in this study, as well as the isotopic values of ebullitive and diffusive gases, we calculate mean isotopic values of $-55.7 \pm 2.1\text{‰}$ $\delta^{13}\text{C}\text{-CH}_4$ and $-260 \pm 36\text{‰}$ $\delta\text{D}\text{-CH}_4$ for the total CH_4 emitted. The $\delta^{13}\text{C}\text{-CH}_4$ reflects relatively heavy values compared to global averages of approximately -62‰ $\delta^{13}\text{C}\text{-CH}_4$ used for aquatic systems (Schwietzke et al., 2016). Considering streams and rivers account for roughly 8% of the global aquatic CH_4 budget (Rosentreter et al., 2021), this difference could be enough to alter the global isotopic signature used for aquatic CH_4 sources.

Very few studies which combine measurements of CH_4 isotopes and emission pathways are available in lotic systems,

limiting comparison of our results (Supplementary Table S2). The most extensive data in rivers comes from the Amazon, where consideration of both ebullitive and diffusive emissions provides mean $\delta^{13}\text{C}\text{-CH}_4$ values for total emitted CH_4 of -49.6‰ (Sawakuchi et al., 2016) and -38.4‰ (Sawakuchi et al., 2021). In these large tropical rivers, diffusion comprises a smaller fraction of the total emission budget, but a greater proportion of dissolved CH_4 is oxidized and thus the emitted CH_4 is more isotopically enriched. Our proposed conceptual model allows for these differences, where the higher efficiency of oxidation in larger rivers is enough to counter the higher prevalence of ebullition (Sawakuchi et al., 2021). Further, hyporheic flow is relatively less important in large rivers (Battin et al., 2008), potentially resulting in a greater proportion of CH_4 accumulating in bubbles in river sediments compared to streams. Even fewer studies exist with which to compare $\delta\text{D}\text{-CH}_4$ values. The streams in this study (-260‰) are $\delta\text{D}\text{-CH}_4$ enriched relative to Swedish lentic ecosystems (-310‰ ; Wik et al., 2020), which further supports the preferred emissions of oxidized CH_4 from streams. More closely comparing lotic $\delta^{13}\text{C}\text{-CH}_4$ signatures to wetlands of similar latitudes further supports relative isotopic enrichment in lotic systems. Latitudinal patterns have been observed in aquatic $\delta^{13}\text{C}\text{-CH}_4$ signatures, with enriched values typically characteristic of more tropical environments (Ganesan et al., 2018). The values measured in the present study are enriched relative to wetlands at a similar latitude (approx. -67‰ ; Fisher et al., 2017). Similarly, $\delta^{13}\text{C}\text{-CH}_4$ values measured in fluvial portions of the Amazon River basin (Sawakuchi et al., 2016; Sawakuchi et al., 2021) are enriched relative to other tropical wetlands (approx. -60‰ ; Brownlow et al., 2017). There remains uncertainty in how much aquatic CH_4 emissions are contributing to variation in atmospheric $\delta^{13}\text{C}\text{-CH}_4$ values (Turner et al., 2019), thus expanding measurement of aquatic CH_4 isotopes remains a valuable endeavor. While many additional measurements across ecosystems will be needed to confirm our proposed model, the possibility of a unique isotopic signature for streams and rivers is significant.

CONCLUSION

This study demonstrates how small streams can emit CH_4 at similar rates to other aquatic ecosystems on a per area basis, but with distinct isotopic signatures compared to lentic systems. The dominance of diffusive emissions to the total CH_4 efflux appears to be characteristic of small streams. Still, the presence of methanogens and methanotrophs highlight the active production and oxidation of CH_4 within stream ecosystems. While oxidation of dissolved CH_4 is limited by the relatively rapid transport of dissolved CH_4 out of the ecosystem, the dominance of diffusive emissions to the overall CH_4 emissions budget ensures the isotopic signature of emitted CH_4 is enriched relative to lentic waterbodies but similar to larger rivers. Consideration of a distinct lotic CH_4 isotopic signature may be critical for accurately incorporating streams and rivers in

regional and global atmospheric mixing CH_4 models. Our proposed conceptual model provides a framework on which to further examine CH_4 production and emissions from streams and how the relative importance of each process varies as a function of stream characteristics. At a global scale, concurrent examination of CH_4 production, oxidation, isotopic values, emissions, and associated microbial communities is needed to determine the generality of our observed patterns in streams. The resultant mechanistic perspectives are required to robustly understand the contribution of lotic ecosystems to the global CH_4 cycle.

DATA AVAILABILITY STATEMENT

The datasets presented in this study can be found in online repositories. The names of the repository/repositories and accession number(s) can be found below: genetic data: <https://www.ncbi.nlm.nih.gov/>, PRJNA757371. All other data: <https://pie-liter.ecosystems.mbl.edu/data>

AUTHOR CONTRIBUTIONS

All authors contributed significantly to this manuscript. AR, WW, AC, JM, RV, and JE designed the project. AR, AC, and PC led field sampling and laboratory analyses. CP and RV led isotopic analyses. AC, JM, and JE led microbial laboratory preparation. AR, CP, and AC led data analyses. AR led manuscript development with input from all authors.

FUNDING

Funding for this project comes from the Plum Island Ecosystems LTER NSF Award OCE-1637630, the National Aeronautics and Space Administration Interdisciplinary Science award NNX17AK10G, and the University of New Hampshire College

of Life Sciences and Agriculture's Paine Fund. Partial funding was provided by the New Hampshire Agricultural Experiment Station. This is Scientific Contribution Number 2923. This work was supported by the USDA National Institute of Food and Agriculture Hatch Projects NH00659 and NH00667.

ACKNOWLEDGMENTS

We thank the members of the Water Systems Analysis Group and the Trace Gas Biogeochemistry Laboratory at the University of New Hampshire for their assistance in project and manuscript development. We thank Eliza Balch, Sarah Bower, and Christopher Whitney for their support in field work, and Apryl Perry for her assistance in laboratory analyses. We also thank Dr. Joanne Shorter from Aerodyne Research Inc. for access to the dual tunable infrared laser direct absorption spectrometer (TILDAS).

SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at: <https://www.frontiersin.org/articles/10.3389/fenvs.2021.791305/full#supplementary-material>

Supplementary Figure S1 | Comparison of measured and estimated gas exchange rates (k_{600}) in the four streams. Gas exchange is measured directly by addition of Argon as a conservative gas tracer. Gas exchange is estimated using Equations 3 and 4 as described in the text. The 1:1 line is displayed as the dashed line.

Supplementary Table S1 | Mean relative abundance (%) of methanogen and methanotroph archaeal and bacterial orders in sediments of the four streams as classified using the SILVA database. Samples are from the surface sediments (0–2 cm depth) or subsurface (9–11 cm depth). nd, no detection. In the order Methanosarcinales the Methanosarcinaceae and Methanosaetaceae families are classified as methanogens, and the Methanoperedenaceae family are classified as methanotrophs.

Supplementary Table S2 | Rates of diffusive and total CH_4 emissions from a variety of aquatic ecosystems, and $\delta^{13}\text{C}-\text{CH}_4$ and $\delta\text{D}-\text{CH}_4$ values of emitted CH_4 . ^Emitted isotope values are inferred from source signatures and both diffusive and ebullitive emission pathway balances.

REFERENCES

- Apprill, A., McNally, S., Parsons, R., and Weber, L. (2015). Minor Revision to V4 Region SSU rRNA 806R Gene Primer Greatly Increases Detection of SAR11 Bacterioplankton. *Aquat. Microb. Ecol.* 75, 129–137. doi:10.3354/ame01753
- Atkins, M. L., Santos, I. R., and Maher, D. T. (2017). Seasonal Exports and Drivers of Dissolved Inorganic and Organic Carbon, Carbon Dioxide, Methane and $\delta^{13}\text{C}$ Signatures in a Subtropical River Network. *Sci. Total Environ.* 575, 545–563. doi:10.1016/j.scitotenv.2016.09.020
- Bastviken, D., Cole, J. J., Pace, M. L., and Van de Bogert, M. C. (2008). Fates of Methane from Different lake Habitats: Connecting Whole-lake Budgets and CH_4 emissions. *J. Geophys. Res.* 113, 1–13. doi:10.1029/2007JG000608
- Bastviken, D., Ejlertsson, J., and Tranvik, L. (2002). Measurement of Methane Oxidation in Lakes: A Comparison of Methods. *Environ. Sci. Technol.* 36, 3354–3361. doi:10.1021/es010311p
- Battin, T. J., Kaplan, L. A., Findlay, S., Hopkinson, C. S., Marti, E., Packman, A. I., et al. (2008). Biophysical Controls on Organic Carbon Fluxes in Fluvial Networks. *Nat. Geosci.* 1, 95–100. doi:10.1038/ngeo101
- Bishop, K., Buffam, I., Erlandsson, M., Fölster, J., Laudon, H., Seibert, J., et al. (2008). Aqua Incognita: the Unknown Headwaters. *Hydrol. Process.* 22, 1239–1242. doi:10.1002/hyp.7049
- Boano, F., Harvey, J. W., Marion, A., Packman, A. I., Revelli, R., Ridolfi, R., et al. (2014). Hyporheic Flow and Transport Processes: Mechanisms, Models, and Biogeochemical Implications. *Rev. Geophys.* 52, 1–77. doi:10.1002/2012RG000417
- Bodmer, P., Wilkinson, J., and Lorke, A. (2020). Sediment Properties Drive Spatial Variability of Potential Methane Production and Oxidation in Small Streams. *J. Geophys. Res. Biogeosci.* 125, 1–15. doi:10.1029/2019JG005213
- Bolyen, E., Rideout, J. R., Dillon, M. R., Bokulich, N. A., Abnet, C., Al-Ghalith, G. A., et al. (2018). QIIME 2: reproducible, interactive, scalable, and extensible microbiome data science. *PeerJ* 6, e27295v2–354.
- Brownlow, R., Lowry, D., Fisher, R. E., France, J. L., Lanoisellé, M., White, B., et al. (2017). Isotopic Ratios of Tropical Methane Emissions by Atmospheric Measurement. *Glob. Biogeochem. Cycles* 31, 1408–1419. doi:10.1002/2017GB005689
- Butman, D., Striegl, R. G., Stackpoole, S. M., Del Giorgio, P. A., Prairie, Y. T., Pilcher, D. J., et al. (2018). Chapter 14: Inland Waters, in *Second State of the Carbon Cycle Report (SOCCR2): A Sustained Assessment Report*. Washington D. C.: U.S. Global Change Research Program. doi:10.7930/SOCCR2.2018.Ch14

- Callahan, B. J., McMurdie, P. J., Rosen, M. J., Han, A. W., Johnson, A. J. A., and Holmes, S. P. (2016). DADA2: High-Resolution Sample Inference from Illumina Amplicon Data. *Nat. Methods* 13, 581–583. doi:10.1038/nmeth.3869
- Chanton, J., Chaser, L., Glasser, P., and Siegel, D. (2005). "Carbon and Hydrogen Isotopic Effects in Microbial, Methane from Terrestrial Environments," in *Stable Isotopes and Biosphere - Atmosphere Interactions* (San Diego, CA, USA: Elsevier Inc.), 85–105. doi:10.1016/B978-012088447-6/50006-4
- Chanton, J. P., Fields, D., and Hines, M. E. (2006). Controls on the Hydrogen Isotopic Composition of Biogenic Methane from High-Latitude Terrestrial Wetlands. *J. Geophys. Res.* 111, 1–9. doi:10.1029/2005JG000134
- Chanton, J. P., Martens, C. S., and Kelley, C. A. (1989). Gas Transport from Methane-Saturated, Tidal Freshwater and Wetland Sediments. *Limnol. Oceanogr.* 34, 807–819. doi:10.4319/lo.1989.34.5.0807
- Chanton, J. P. (2005). The Effect of Gas Transport on the Isotope Signature of Methane in Wetlands. *Org. Geochem.* 36, 753–768. doi:10.1016/j.orggeochem.2004.10.007
- Chowdhury, T. R., and Dick, R. P. (2013). Ecology of Aerobic Methanotrophs in Controlling Methane Fluxes from Wetlands. *Appl. Soil Ecol.* 65, 8–22. doi:10.1016/j.apsoil.2012.12.014
- Conrad, R. (2005). Quantification of Methanogenic Pathways Using Stable Carbon Isotopic Signatures: A Review and a Proposal. *Org. Geochem.* 36, 739–752. doi:10.1016/j.orggeochem.2004.09.006
- Conrad, R. (2009). The Global Methane Cycle: Recent Advances in Understanding the Microbial Processes Involved. *Environ. Microbiol. Rep.* 1, 285–292. doi:10.1111/j.1758-2229.2009.00038.x
- Crawford, J. T., Stanley, E. H., Spawn, S. A., Finlay, J. C., Loken, L. C., and Striegl, R. G. (2014). Ebullitive Methane Emissions from Oxygenated Wetland Streams. *Glob. Change Biol.* 20, 3408–3422. doi:10.1111/gcb.12614
- Crawford, J. T., Striegl, R. G., Wickland, K. P., Dornblaser, M. M., and Stanley, E. H. (2013). Emissions of Carbon Dioxide and Methane from a Headwater Stream Network of interior Alaska. *J. Geophys. Res. Biogeosci.* 118, 482–494. doi:10.1002/jgrg.20034
- Dean, J. F., Middelburg, J. J., Röckmann, T., Aerts, R., Blauw, L. G., Egger, M., et al. (2018). Methane Feedbacks to the Global Climate System in a Warmer World. *Rev. Geophys.* 56, 207–250. doi:10.1002/2017RG000559
- Doherty, S. J., Barbato, R. A., Grandy, A. S., Thomas, W. K., Monteux, S., Dorrepaal, E., et al. (2020). The Transition from Stochastic to Deterministic Bacterial Community Assembly during Permafrost Thaw Succession. *Front. Microbiol.* 11, 2923. doi:10.3389/fmicb.2020.596589
- Earth System Research Laboratory, G. M. D. (2021). Carbon Cycle Gases Offshore Portsmouth, New Hampshire (Isles of Shoals), United States. Available at: <https://gml.noaa.gov/dv/iadv/index.php?code=nha> (Accessed July 1, 2021).
- Ettwig, K. F., Zhu, B., Speth, D., Keljens, J. T., Jetten, M. S. M., and Kartal, B. (2016). Archaea Catalyze Iron-dependent Anaerobic Oxidation of Methane. *Proc. Natl. Acad. Sci. USA* 113, 12792–12796. doi:10.1073/pnas.1609534113
- Evans, P. N., Boyd, J. A., Leu, A. O., Woodcroft, B. J., Parks, D. H., Hugenholtz, P., et al. (2019). An Evolving View of Methane Metabolism in the Archaea. *Nat. Rev. Microbiol.* 17, 219–232. doi:10.1038/s41579-018-0136-7
- Fisher, R. E., France, J. L., Lowry, D., Lanoisellé, M., Brownlow, R., Pyle, J. A., et al. (2017). Measurement of the $\delta^{13}\text{C}$ Isotopic Signature of Methane Emissions from Northern European Wetlands. *Glob. Biogeochem. Cycles* 31, 605–623. doi:10.1002/2016GB005504
- Frolking, S., and Crill, P. (1994). Climate Controls on Temporal Variability of Methane Flux from a Poor Fen in southeastern New Hampshire: Measurement and Modeling. *Glob. Biogeochem. Cycles* 8, 385–397. doi:10.1029/94gb01839
- Ganesan, A. L., Stell, A. C., Gedney, N., Comyn-Platt, E., Hayman, G., Rigby, M., et al. (2018). Spatially Resolved Isotopic Source Signatures of Wetland Methane Emissions. *Geophys. Res. Lett.* 45, 3737–3745. doi:10.1002/2018GL077536
- Glöckner, F. O., Yilmaz, P., Quast, C., Gerken, J., Beccati, A., Ciuprina, A., et al. (2017). 25 Years of Serving the Community with Ribosomal RNA Gene Reference Databases and Tools. *J. Biotechnol.* 261, 169–176. doi:10.1016/j.jbiotec.2017.06.1198
- Hall, R. O., and Ulseth, A. J. (2020). Gas Exchange in Streams and Rivers. *WIREs Water* 7, 1–19. doi:10.1002/wat2.1391
- Hampton, T. B., Zarnetske, J. P., Briggs, M. A., MahmoodPoor Dehkordy, F., Singha, K., Day-Lewis, F. D., et al. (2020). Experimental Shifts of Hydrologic Residence Time in a sandy Urban Stream Sediment-Water Interface Alter Nitrate Removal and Nitrous Oxide Fluxes. *Biogeochemistry* 149, 195–219. doi:10.1007/s10533-020-00674-7
- Herreid, A. M., Wymore, A. S., Varner, R. K., Potter, J. D., and McDowell, W. H. (2020). Divergent Controls on Stream Greenhouse Gas Concentrations across a Land-Use Gradient. *Ecosystems* 24, 1299–1316. doi:10.1007/s10021-020-00584-7
- Hojo, T., and Pearson, K. (1931). Distribution of the Median, Quartiles and Interquartile Distance in Samples from a Normal Population. *Biometrika* 23, 315–363. doi:10.2307/2332422
- Holgersen, M. A., and Raymond, P. A. (2016). Large Contribution to Inland Water CO₂ and CH₄ Emissions from Very Small Ponds. *Nat. Geosci.* 9, 222–226. doi:10.1038/ngeo2654
- Hotchkiss, E. R., Hall Jr, R. O., Jr, Sponseller, R. A., Butman, D., Klaminder, J., Laudon, H., et al. (2015). Sources of and Processes Controlling CO₂ Emissions Change with the Size of Streams and Rivers. *Nat. Geosci.* 8, 696–699. doi:10.1038/ngeo2507
- Hough, M., McClure, A., Bolduc, B., Dorrepaal, E., Saleska, S., Klepac-Ceraj, V., et al. (2020). Biotic and Environmental Drivers of Plant Microbiomes across a Permafrost Thaw Gradient. *Front. Microbiol.* 11, 1–18. doi:10.3389/fmicb.2020.00796
- Knighton, D. (1998). *Fluvial Forms and Processes: A New Perspective*. First. New York, New York: Hodder Arnold.
- Knox, M., Quay, P. D., and Wilbur, D. (1992). Kinetic Isotopic Fractionation during Air-Water Gas Transfer of O₂, N₂, CH₄, and H₂. *J. Geophys. Res.* 97, 20335. doi:10.1029/92jc00949
- Leu, A. O., Cai, C., McIlroy, S. J., Southam, G., Orphan, V. J., Yuan, Z., et al. (2020). Anaerobic Methane Oxidation Coupled to Manganese Reduction by Members of the Methanoperedenaceae. *ISME J.* 14, 1030–1041. doi:10.1038/s41396-020-0590-x
- Liu, L., Wilkinson, J., Koca, K., Buchmann, C., and Lorke, A. (2016). The Role of Sediment Structure in Gas Bubble Storage and Release. *J. Geophys. Res. Biogeosci.* 121, 1992–2005. doi:10.1002/2016JG003456
- Looman, A., Maher, D. T., and Santos, I. R. (2021). Carbon Dioxide Hydrodynamics along a wetland-lake-stream-waterfall Continuum (Blue Mountains, Australia). *Sci. Total Environ.* 777, 146124. doi:10.1016/j.scitotenv.2021.146124
- Magen, C., Lapham, L. L., Pohlman, J. W., Marshall, K., Bosman, S., Casso, M., et al. (2014). A Simple Headspace Equilibration Method for Measuring Dissolved Methane. *Limnol. Oceanogr.* 12, 637–650. doi:10.4319/lom.2014.12.637
- MahmoodPoor Dehkordy, F., Briggs, M. A., Day-Lewis, F. D., Singha, K., Krajnovich, A., Hampton, T. B., et al. (2019). Multi-scale Preferential Flow Processes in an Urban Streambed under Variable Hydraulic Conditions. *J. Hydrol.* 573, 168–179. doi:10.1016/j.jhydrol.2019.03.022
- Martin, M. (2011). Cutadapt Removes Adapter Sequences from High-Throughput Sequencing Reads. *EMBnet j.* 17, 10–2809. doi:10.14806/embj.17.1.200
- Martiny, J. B. H., Jones, S. E., Lennon, J. T., and Martiny, A. C. (2015). Microbiomes in Light of Traits: A Phylogenetic Perspective. *Science* 350, 350. doi:10.1126/science.aac9323
- Marx, A., Dusek, J., Jankovec, J., Sanda, M., Vogel, T., van Geldern, R., et al. (2017). A Review of CO₂ and Associated Carbon Dynamics in Headwater Streams: A Global Perspective. *Rev. Geophys.* 55, 560–585. doi:10.1002/2016RG000547
- Mcmanus, J. B., Zahniser, M. S., and Nelson, D. D. (2011). Dual Quantum cascade Laser Trace Gas Instrument with Astigmatic Herriott Cell at High Pass Number. *Appl. Opt.* 50, A74–A85. doi:10.1364/AO.50.000A74
- McManus, J. B., Zahniser, M. S., Nelson, D. D., Shorter, J. H., Herndon, S. C., Jervis, D., et al. (2015). Recent Progress in Laser-Based Trace Gas Instruments: Performance and Noise Analysis. *Appl. Phys. B* 119, 203–218. doi:10.1007/s00340-015-6033-0
- Morse, N. B., and Wollheim, W. M. (2014). Climate Variability Masks the Impacts of Land Use Change on Nutrient export in a Suburbanizing Watershed. *Biogeochemistry* 121, 45–59. doi:10.1007/s10533-014-9998-6
- Nelson, A. R., Sawyer, A. H., Gabor, R. S., Saup, C. M., Bryant, S. R., Harris, K. D., et al. (2019). Heterogeneity in Hyporheic Flow, Pore Water Chemistry, and Microbial Community Composition in an alpine Streambed. *J. Geophys. Res. Biogeosci.* 124, 3465–3478. doi:10.1029/2019JG005226
- Oliver, J. D. (2005). The Viable but Nonculturable State in Bacteria. *J. Microbiol. Spec No.* 93–100.
- Packman, A. I., and Salehin, M. (2003). Relative Roles of Stream Flow and Sedimentary Conditions in Controlling Hyporheic Exchange. *Hydrobiologia* 494, 291–297. doi:10.1023/A:1025403424063

- Parada, A. E., Needham, D. M., and Fuhrman, J. A. (2016). Every Base Matters: Assessing Small Subunit rRNA Primers for marine Microbiomes with Mock Communities, Time Series and Global Field Samples. *Environ. Microbiol.* 18, 1403–1414. doi:10.1111/1462-2920.13023
- Quast, C., Pruesse, E., Yilmaz, P., Gerken, J., Schweer, T., Yarza, P., et al. (2013). The SILVA Ribosomal RNA Gene Database Project: Improved Data Processing and Web-Based Tools. *Nucleic Acids Res.* 41, D590–D596. doi:10.1093/nar/gks1219
- Raymond, P. A., Zappa, C. J., Butman, D., Bott, T. L., Potter, J., Mulholland, P., et al. (2012). Scaling the Gas Transfer Velocity and Hydraulic Geometry in Streams and Small Rivers. *Limnol. Oceanogr.* 2, 41–53. doi:10.1215/21573689-1597669
- Ries, K. G., III, Newson, J. K., Smith, M. J., Guthrie, J. D., Steeves, P. A., Haluska, T. L., et al. (2017). StreamStats, Version 4. In U.S. Geological Survey Fact Sheet 2017–3046. doi:10.3133/fs20173046
- Robison, A. L., Wollheim, W. M., Turek, B., Bova, C., Snay, C., and Varner, R. K. (2021). Spatial and Temporal Heterogeneity of Methane Ebullition in lowland Headwater Streams and the Impact on Sampling Design. *Limnol. Oceanogr.* 66, 4063–4076. doi:10.1002/lno.11943
- Rosentreter, J. A., Borges, A. V., Deemer, B. R., Holgersson, M. A., Liu, S., Song, C., et al. (2021). Half of Global Methane Emissions Come from Highly Variable Aquatic Ecosystem Sources. *Nat. Geosci.* 14, 225–230. doi:10.1038/s41561-021-00715-2
- Sanchez, L. F., Guenet, B., Marinho, C. C., Barros, N., and de Assis Esteves, F. (2019). Global Regulation of Methane Emission from Natural Lakes. *Sci. Rep.* 9, 1–10. doi:10.1038/s41598-018-36519-5
- Sanders, I. A., Heppell, C. M., Cotton, J. A., Wharton, G., Hildrew, A. G., Flowers, E. J., et al. (2007). Emission of Methane from Chalk Streams Has Potential Implications for Agricultural Practices. *Freshw. Biol.* 52, 1176–1186. doi:10.1111/j.1365-2427.2007.01745.x
- Saunio, M., Stavert, A., Poulter, B., Bousquet, P., Canadell, J., Jackson, R., et al. (2020). The Global Methane Budget 2000–2017. *Earth Syst. Sci. Data* 12, 1561–1623. doi:10.5194/essd-12-1561-2020
- Sawakuchi, H. O., Bastviken, D., Enrich-prast, A., Ward, N. D., Camargo, P. B., and Richey, J. E. (2021). Low Diffusive Methane Emissions from the Main Channel of a Large Amazonian Run-Of-The-River Reservoir Attributed to High Methane Oxidation. *Front. Environ. Sci.* 9, 1–13. doi:10.3389/fenvs.2021.655455
- Sawakuchi, H. O., Bastviken, D., Sawakuchi, A. O., Krusche, A. V., Ballester, M. V. R., and Richey, J. E. (2014). Methane Emissions from Amazonian Rivers and Their Contribution to the Global Methane Budget. *Glob. Change Biol.* 20, 2829–2840. doi:10.1111/gcb.12646
- Sawakuchi, H. O., Bastviken, D., Sawakuchi, A. O., Ward, N. D., Borges, C. D., Tsai, S. M., et al. (2016). Oxidative Mitigation of Aquatic Methane Emissions in Large Amazonian Rivers. *Glob. Change Biol.* 22, 1075–1085. doi:10.1111/gcb.13169
- Schwietke, S., Sherwood, O. A., Bruhwiler, L. M. P., Miller, J. B., Etiope, G., Dlugokencky, E. J., et al. (2016). Upward Revision of Global Fossil Fuel Methane Emissions Based on Isotope Database. *Nature* 538, 88–91. doi:10.1038/nature19797
- Shelley, F., Abdullahi, F., Grey, J., and Trimmer, M. (2015). Microbial Methane Cycling in the Bed of a Chalk River: Oxidation Has the Potential to Match Methanogenesis Enhanced by Warming. *Freshw. Biol.* 60, 150–160. doi:10.1111/fwb.12480
- Shelley, F., Ings, N., Hildrew, A. G., Trimmer, M., and Grey, J. (2017). Bringing Methanotrophy in Rivers Out of the Shadows. *Limnol. Oceanogr.* 62, 2345–2359. doi:10.1002/lno.10569
- Smith, G. J., and Wrighton, K. C. (2019). Metagenomic Approaches Unearth Methanotroph Phylogenetic and Metabolic Diversity. *Curr. Issues Mol. Biol.* 33, 57–84. doi:10.21775/cimb.033.057
- Sobczak, W. V., and Findlay, S. (2002). Variation in Bioavailability of Dissolved Organic Carbon Among Stream Hyporheic Flowpaths. *Ecology* 83, 3194–3209. doi:10.1890/0012-9658(2002)083[3194:vibodo]2.0.co;2
- Stanley, E. H., Casson, N. J., Christel, S. T., Crawford, J. T., Loken, L. C., and Oliver, S. K. (2016). The Ecology of Methane in Streams and Rivers: Patterns, Controls, and Global Significance. *Ecol. Monogr.* 86, 146–171. doi:10.1890/15-1027.1
- Trimmer, M., Maanoja, S., Hildrew, A. G., Pretty, J. L., and Grey, J. (2010). Potential Carbon Fixation via Methane Oxidation in Well-Oxygenated River Bed Gravels. *Limnol. Oceanogr.* 55, 560–568. doi:10.4319/lno.2009.55.2.056010.4319/lno.2010.55.2.0560
- Turner, A. J., Frankenberg, C., and Kort, E. A. (2019). Interpreting Contemporary Trends in Atmospheric Methane. *Proc. Natl. Acad. Sci. USA* 116, 2805–2813. doi:10.1073/pnas.1814297116
- Tyler, S. C., Bilek, R. S., Sass, R. L., and Fisher, F. M. (1997). Methane Oxidation and Pathways of Production in a Texas Paddy Field Deduced from Measurements of Flux, $\delta^{13}\text{C}$, and δD of CH_4 . *Glob. Biogeochem. Cycles* 11, 323–348. doi:10.1029/97gb01624
- Walters, W., Hyde, E. R., Berg-Lyons, D., Ackermann, G., Humphrey, G., Parada, A., et al. (2016). Improved Bacterial 16S rRNA Gene (V4 and V4-5) and Fungal Internal Transcribed Spacer Marker Gene Primers for Microbial Community Surveys. *mSystems* 1, 1–10. doi:10.1128/msystems.00009-15
- Wang, D. T., Welander, P. V., and Ono, S. (2016). Fractionation of the Methane Isotopologues $^{13}\text{CH}_4$, $^{12}\text{CH}_3\text{D}$, and $^{13}\text{CH}_3\text{D}$ during Aerobic Oxidation of Methane by *Methylococcus Capsulatus* (Bath). *Geochimica et Cosmochimica Acta* 192, 186–202. doi:10.1016/j.gca.2016.07.031
- Wang, G., Xia, X., Liu, S., Zhang, L., Zhang, S., Wang, J., et al. (2021). Intense Methane Ebullition from Urban Inland Waters and its Significant Contribution to Greenhouse Gas Emissions. *Water Res.* 189, 116654. doi:10.1016/j.watres.2020.116654
- Whiticar, M. J. (1999). Carbon and Hydrogen Isotope Systematics of Bacterial Formation and Oxidation of Methane. *Chem. Geology* 161, 291–314. doi:10.1016/S0009-2541(99)00092-3
- Wik, M., Johnson, J. E., Crill, P. M., DeStasio, J. P., Erickson, L., Halloran, M. J., et al. (2018). Sediment Characteristics and Methane Ebullition in Three Subarctic Lakes. *J. Geophys. Res. Biogeosci.* 123, 2399–2411. doi:10.1029/2017JG004298
- Wik, M., Thornton, B. F., Bastviken, D., Uhlbäck, J., and Crill, P. M. (2016a). Biased Sampling of Methane Release from Northern Lakes: A Problem for Extrapolation. *Geophys. Res. Lett.* 43, 1256–1262. doi:10.1002/2015GL066501.Received
- Wik, M., Thornton, B. F., Varner, R. K., McCalley, C., and Crill, P. M. (2020). Stable Methane Isotopologues from Northern Lakes Suggest that Ebullition Is Dominated by Sub-Lake Scale Processes. *J. Geophys. Res. Biogeosci.* 125, 1–13. doi:10.1029/2019JG005601
- Wik, M., Varner, R. K., Anthony, K. W., MacIntyre, S., and Bastviken, D. (2016b). Climate-sensitive Northern Lakes and Ponds Are Critical Components of Methane Release. *Nat. Geosci.* 9, 99–105. doi:10.1038/ngeo2578
- Wollheim, W. M., Mulukutla, G. K., Cook, C., and Carey, R. O. (2017). Aquatic Nitrate Retention at River Network Scales across Flow Conditions Determined Using Nested *In Situ* Sensors. *Water Resour. Res.* 53, 9740–9756. doi:10.1002/2017WR020644
- Wollheim, W. M., Pellerin, B. A., Vörösmarty, C. J., and Hopkinson, C. S. (2005). N Retention in Urbanizing Headwater Catchments. *Ecosystems* 8, 871–884. doi:10.1007/s10021-005-0178-3
- Wollheim, W. M., Stewart, R. J., Aiken, G. R., Butler, K. D., Morse, N. B., and Salisbury, J. (2015). Removal of Terrestrial DOC in Aquatic Ecosystems of a Temperate River Network. *Geophys. Res. Lett.* 42, 6671–6679. doi:10.1002/2015GL064647
- Wu, M. L., Ettwig, K. F., Jetten, M. S. M., Strous, M., Keltjens, J. T., and Niftrik, L. v. (2011). A New Intra-aerobic Metabolism in the Nitrite-dependent Anaerobic Methane-Oxidizing Bacterium *Candidatus 'Methyloirabilis Oxyfera'*. *Biochem. Soc. Trans.* 39, 243–248. doi:10.1042/BST0390243
- Zhang, G., Yu, H., Fan, X., Ma, J., and Xu, H. (2016). Carbon Isotope Fractionation Reveals Distinct Process of CH_4 Emission from Different Compartments of Paddy Ecosystem. *Sci. Rep.* 6, 1–10. doi:10.1038/srep27065
- Zhang, L., Xia, X., Liu, S., Zhang, S., Li, S., Wang, J., et al. (2020). Significant Methane Ebullition from alpine Permafrost Rivers on the East Qinghai-Tibet Plateau. *Nat. Geosci.* 13, 349–354. doi:10.1038/s41561-020-0571-8

Conflict of Interest: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

Publisher's Note: All claims expressed in this article are solely those of the authors and do not necessarily represent those of their affiliated organizations, or those of the publisher, the editors, and the reviewers. Any product that may be evaluated in this article, or claim that may be made by its manufacturer, is not guaranteed or endorsed by the publisher.

Copyright © 2022 Robison, Wollheim, Perryman, Cotter, Mackay, Varner, Clarizia and Ernakovich. This is an open-access article distributed under the terms of the Creative Commons Attribution License (CC BY). The use, distribution or reproduction in other forums is permitted, provided the original author(s) and the copyright owner(s) are credited and that the original publication in this journal is cited, in accordance with accepted academic practice. No use, distribution or reproduction is permitted which does not comply with these terms.