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Received 29 May 2014 | Accepted 22 Sep 2014 | Published 30 Oct 2014

DOI: 10.1038/ncomms6350

Oxic water column methanogenesis as a major component of aquatic CH₄ fluxes

Matthew J. Bogard¹, Paul A. del Giorgio¹, Lennie Boutet¹, Maria Carolina Garcia Chaves¹, Yves T. Prairie¹, Anthony Merante¹ & Alison M. Derry¹

Methanogenesis has traditionally been assumed to occur only in anoxic environments, yet there is mounting, albeit indirect, evidence of methane (CH₄) production in oxic marine and freshwaters. Here we present the first direct, ecosystem-scale demonstration of methanogenesis in oxic lake waters. This methanogenesis appears to be driven by acetoclastic production, and is closely linked to algal dynamics. We show that oxic water methanogenesis is a significant component of the overall CH₄ budget in a small, shallow lake, and provide evidence that this pathway may be the main CH₄ source in large, deep lakes and open oceans. Our results challenge the current global understanding of aquatic CH₄ dynamics, and suggest a hitherto unestablished link between pelagic CH₄ emissions and surface-water primary production. This link may be particularly sensitive to widespread and increasing human influences on aquatic ecosystem primary productivity.

¹ Groupe de Recherche Interuniversitaire en Limnologie, Département des Sciences Biologiques, Université du Québec à Montréal, Case Postale 8888, Succursale Centre-Ville, Montréal, Quebec, Canada H3C 3P8. Correspondence and requests for materials should be addressed to M.J.B. (email: bogard.matt@gmail.com).

ethane (CH₄) emissions currently contribute ~20% to the planet's greenhouse effect, with a large, but poorly defined, fraction derived from freshwater ecosystems 1-4. Accurately placing freshwaters in the global CH₄ budget requires a better understanding of the controls and contributions of CH₄ from different sources²⁻⁵. In this regard, the surface waters of lakes and rivers are systematically supersaturated with CH₄, and it has been traditionally assumed that this CH₄ is derived from anoxic environments, via vertical and lateral transport from profundal and littoral sediments^{3,6–8}. This assumption certainly holds for small, shallow ecosystems where the surface layers are in relatively close contact with sediments. Yet CH₄ supersaturation is also prevalent in large, deep lakes^{3,5,7–12} and oceanic surface waters^{13–17}, where deeper water columns result in significantly reduced surface water exchange with anoxic sediments. Studies of CH4 dynamics in surface waters of oceans and large lakes have indeed concluded that pelagic CH₄ supersaturation cannot be sustained either by lateral inputs from the littoral or from benthic inputs alone^{9,10,13}–15,17–19

A pelagic source of CH₄ would thus be required to sustain the observed CH₄ supersaturation, and multiple lines of evidence suggest that CH₄ is produced in oxic water columns of freshwater and marine systems. In vitro experiments point to CH₄ production in both microanoxic habitats in metazoan guts and on particles 14,17-23, and in particle-free oxic water 15,16,18,19,23-25, via multiple biochemical pathways. The presence 19,22,26-28 and activity^{19,22,27} of hydrogenotrophic and acetoclastic methanogenic archaea have been confirmed at the molecular level in diverse oxic environments. At the same time, marine studies suggest that the microbial decomposition of methylated compounds such as methanethiol¹⁵ and methylphosphonate^{16,23–25} could be an important source of CH₄ in surface waters of the open ocean. Further, metalimnetic peaks of CH4 are a recurrent feature in many ecosystems, which correlate positively with dissolved $\rm O_2$ (DO), algal biomass and production $^{8-10,14,15,18,19}$. However, despite past efforts, the ecological and biogeochemical significance of oxic water column methanogenesis remain speculative, and we have yet to determine the dominant biochemical pathway, its controls, and its contribution to diffusive CH4 emissions from aquatic ecosystems.

In this study, we experimentally assess pelagic CH₄ production in the oxygenated water column of Lac Cromwell, a typical Canadian temperate Shield lake, using floating mesocosms open to the atmosphere, but closed at the bottom and therefore uncoupled from non-pelagic sources of CH₄. The results presented herein represent the first direct, ecosystem-scale demonstration of oxic water CH₄ production, its drivers and source pathway, and the potential widespread importance of this poorly considered process.

Results

CH₄ dynamics in experimental enclosures. The initial CH₄ concentrations in the enclosures were four- to tenfold lower than surrounding lake waters, because of degassing during the initial filtration and filling process, but the enclosures were nevertheless supersaturated relative to the atmosphere at the onset of the experiment (Fig. 1a). Vertical profiles of the enclosures (Supplementary Fig. 1) showed a range in DO between 45.6 and 128.6% saturation, indicating that the entire water column remained oxic in all treatments for the duration of the experiment. Despite oxic conditions, CH₄ concentrations ranged between 0.10 and 0.53 μ M throughout the entire experiment, across all treatments (Fig. 1a), which represents \sim 50- to 265-fold super-saturation relative to the atmosphere.

The observed surface CH₄ concentrations in the mesocosms represent the net balance between CH₄ production, CH₄ oxidation (MOX) and CH₄ exchange with the atmosphere. In the absence of internal inputs of CH₄, the initial CH₄ supersaturation in the enclosures would have declined to atmospheric equilibrium within approximately 1 week (Fig. 1a, dotted lines), as estimated on the basis of our own empirical measurements of the gas exchange coefficient (see Methods section). Thus, an internal CH₄ source was necessary to sustain the systematic CH₄ supersaturation observed in all mesocosms throughout the experiment. Further, there was an overall significant increase in CH₄ concentrations through time across treatments (Fig. 1a; repeated measures analysis of variance (RM-ANOVA) time effect: P = 0.007), with greater increases in nutrient-amended enclosures (nitrogen and phosphorus (NP), dissolved organic carbon (DOC)-NP), and weaker increases in the DOC and control enclosures (RM-ANOVA treatment \times time effect: P = 0.04). As a result, all enclosures emitted CH₄ to the atmosphere for the duration of the experiment, and the estimated water/air CH₄ fluxes (based on our measurements of gas transfer coefficients in the mesocosms) ranged from 0.07 to 0.36 mmol m⁻² per day (Fig. 1b). Net CH₄ production in the enclosures, calculated as the sum of the observed net increase in concentration and the calculated CH₄ fluxes to the atmosphere, were on average highest in the nutrient-enriched enclosures, whereas the DOC-only addition generated the lowest production rates (Fig. 1b; analysis of variance (ANOVA): P = 0.003). We assessed the robustness of chamber-based calculations of CH₄ emissions by comparing chamber-derived gas exchange coefficients (k_{CH4}) to estimates based on wind speed (detailed in Supplementary Note 1). There was good agreement between the two approaches, and therefore we conclude that chamber-derived results yield reliable estimates of k_{CH4} and CH_4 fluxes for both the enclosures and the lake.

Linking CH₄ production to algal dynamics. The differences among treatments in water column CH₄ concentrations and fluxes were strongly linked to pelagic gross primary production (GPP; Fig. 2a) and net ecosystem production (NEP = GPP-R; Fig. 2b). There was a 20-fold range in GPP across treatments (Fig. 2a), whereas there was only a 5-fold range in ecosystem respiration (R), therefore nutrient additions resulted in a strong shift in NEP (Fig. 2b). The average CH₄ fluxes at day 7 of the experiment were strongly positively related to average GPP rates in the enclosures (Fig. 2a), and to NEP, whereas the DOC addition had no positive influence relative to the control. Increases in CH₄ flux appear to have been driven mostly by increases in GPP rather than heterotrophic metabolism.

Other potential sources of CH₄ in the enclosures were also considered. The enclosures contained very little particulate organic C because of the initial filtration of the surrounding lake water before filling, and there was little or no accumulation of particulate organic C in the bottom of enclosures at the end of the experiment. Moreover, we estimate that zooplankton-derived CH₄, calculated from the measured zooplankton biomass in the enclosures and published production rates (Supplementary Table 1, Supplementary Note 2), contributed <10% to the estimated gross CH₄ production rates in ambient enclosures, a result that is consistent with previous studies 10,18,19,29. Membrane permeability and the resulting exchange with the surrounding lake waters were also considered as a potential cause for CH₄ supersaturation. However, the estimated contributions of CH₄ from the surrounding lake environment were on the order of <1% of gross CH₄ production (Supplementary Methods and Supplementary Table 2). Finally, there was no significant

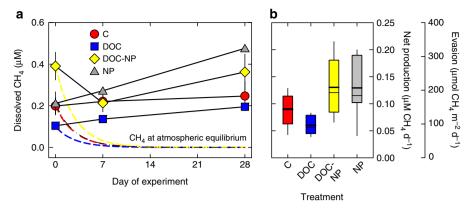


Figure 1 | Surface water CH₄ dynamics in experimental enclosures. Treatments include ambient untreated controls (C), addition of dissolved organic carbon (DOC), nitrogen and phosphorus (NP) or a combination of both (DOC-NP). (a) CH_4 concentrations (error bars ± 1 s.e.m., n=3) generally increased through time, but differed between treatments (RM-ANOVA, treatment: P=0.004, F=10.71; time: P=0.007, F=7.67; treatment* time: P=0.04, F=3.04; Tukey's HSD post-hoc grouping = NP, DOC-NP, C>C, DOC). Dashed lines show the expected declines in dissolved CH_4 because of diffusion at the air-water interface, had no methane been produced in situ. Without internal production of CH_4 , concentrations would have equilibrated with the atmosphere (mean concentration at equilibrium = $0.002 \,\mu$ M) in all treatments within 1-2 weeks. (b) Collectively, net CH_4 production (evasion plus water column accumulation per enclosure) and evasion rates (second y axis) displayed large differences among treatments (ANOVA, P=0.003; post-hoc groups = NP & DOC-NP & C>C>C & DOC). Thick and thin horizontal lines are mean and median, respectively.

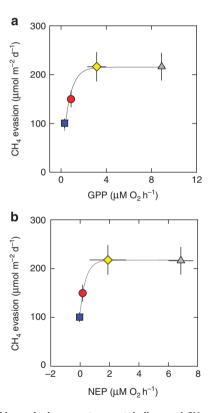


Figure 2 | Linking pelagic ecosystem metabolism and CH₄ **dynamics.** Metabolic estimates from day 7 of the experiment revealed CH₄ evasion was strongly positively correlated to (**a**) ecosystem gross primary production (GPP; P = 0.003; $r^2 = 0.93$; $y = 215.3[1 - e^{-1.6x}]$), and (**b**) shifts to increased net ecosystem production (NEP), which is different between GPP and community respiration (P = 0.02; $r^2 = 0.99$; $y = 111-106[1 - e^{-2.6x}]$). Symbols with error bars (± 1 s.e.m., n = 3) denote the different treatments, including ambient untreated controls (red circles), addition of dissolved organic carbon (blue squares), nitrogen and phosphorus (grey triangles) or a combination of both (yellow diamonds).

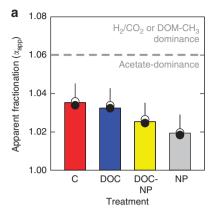
development of biofilm on the walls of the enclosures during the experiment, and this is unlikely to have been a significant source of CH₄.

Identifying the biochemical source of CH₄. Several methanogenic pathways exist, but as acetoclastic methanogenesis is less isotopically discriminatory than CO_2 reduction or methylotrophy 19,27,28 , apparent fractionation factors (α_{app} ; see Methods for calculation details) can be used to qualitatively distinguish whether CH_4 is produced via methanogenesis from acetate, or CO_2 reduction 30 . Our measured α_{app} values were well within the range indicating dominance of the acetoclastic pathway (Fig. 3a), suggesting that acetate was the dominant source material for pelagic methanogenesis in all treatments. As we only had one estimate of water column MOX per treatment (range = $0.003-0.008 \,\mathrm{mmol}\,\mathrm{m}^{-3}\,\mathrm{h}^{-1}$), we used a scenario analysis with two extreme MOX rates (0 and $0.05 \text{ mmol m}^{-3} \text{ h}^{-1}$; detailed in Methods section) conservatively constrain the possible α_{app} values based on variable MOX rates and associated isotopic fractionation. In all cases, α_{app} values remained well within the acetoclastic range, that is, <1.055 (ref. 30; Fig. 3a). Acetoclastic methanogenesis was likely dominant through time, because $\delta^{13}\text{C-CO}_2$ and $\delta^{13}\text{C-CH}_4$ showed little temporal variability (Fig. 3b).

Rates of oxic water-column methanogenesis. The $\mathrm{CH_4}$ accumulation and outgassing that we measured in the enclosures (Fig. 1) represent the net result of pelagic methanogenesis, gas exchange and MOX. In order to derive a first-order estimate of pelagic methanogenesis, we carried out a detailed $\mathrm{CH_4}$ mass balance (described in the Methods section; summarized in Supplementary Table 2) for day 7 for the control enclosures. Here we combined the measured $\mathrm{CH_4}$ evasion, the change in storage corrected for cross-membrane inputs from surrounding lake water and measured MOX. This exercise yielded rates of apparent pelagic methanogenesis on the order of 0.21–0.24 mmol m $^{-3}$ per day (Table 1), of which $\sim 60\%$ was apparently oxidized and the remainder evaded to the atmosphere.

Oxic water methanogenesis in a whole-lake perspective. Mean pelagic CH₄ evasion from control (that is, non-manipulated) enclosures (0.15 \pm 0.03 mmol m $^{-2}$ per day; Table 1) represented \sim 20% of the average diffusive CH₄ fluxes measured in the lake over the summer (0.78 \pm 0.35 mmol m $^{-2}$ per day), and at the whole-lake scale, were similar to CH₄ ebullitive fluxes

 $(15.2 \pm 32.5 \,\text{mol})$ per day or $0.31 \pm 0.66 \,\text{mmol}\,\text{m}^{-2}$ per day in lake regions $< 3 \,\text{m}$ deep; Table 1). Further, oxic pelagic methanogenesis contributed on the order of 4% to total lake greenhouse



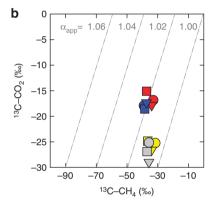


Figure 3 | Assessing the biochemical source of pelagic CH4. (a) Here patterns in the apparent fractionation (α_{app} ; see methods for calculation details) during methanogenesis suggest that the acetoclastic pathway strongly dominated in all treatments, and increased in importance with the addition of nutrients. White and black circles represent values estimated using measured CH₄ oxidation rates (MOX) and a maximum and minimum associated isotopic fractionation factor, respectively. To constrain the effect of error introduced by MOX estimates, α_{app} was estimated from a highly conservative scenario analysis, summarized here by bars and error bars (midpoint and upper potential range, respectively). Values of α_{app} were estimated for day 7 of the experiment, although isotopic signatures of CO2 and ambient CH₄ displayed little change through time for each treatment (b). Red, blue, grey and yellow symbols indicate control, DOC, DOC + NP and NP treatments, respectively, whereas circles, squares and triangles, respectively, indicate days 0, 7 and 28 of the experiment. One enclosure was followed for each treatment, and sample points represent one unreplicated measurement.

gas emissions ($CO_2 + CH_4$, expressed as CO_2 equivalents; Table 1). It was possible to compare fluxes directly between the enclosures and the lake because the physical conditions shaping gas exchange dynamics were quantitatively similar for each. As shown in Supplementary Fig. 2, floating-chamber-derived gas exchange coefficients ($k_{\rm CH4}$) for the lake and the enclosures varied on a diurnal basis, but averaged 0.65 and 0.69 m per day and ranged between 0.10–1.91 and 0.24–1.44 m per day for the enclosures and the lake, respectively.

Discussion

Here we present an ecosystem-scale, experimental demonstration of significant CH₄ production and evasion from the oxic water column of Lac Cromwell that is closely linked to algal dynamics, and which contributes a baseline flux of CH₄ that is likely present in all lakes. The relationship between CH₄ and phytoplankton observed here (Fig. 2) has been hypothesized before to explain both the presence of metabolically active methanogens, and the recurrent metalimnetic and near-surface CH₄ peaks in oxic lake^{9,10,18,19} and marine^{13–17} environments. We confirm this link experimentally, and further show that it generates a significant out flux of CH₄ from the mesocosms to the atmosphere (Fig. 1b). These results in turn imply that factors influencing phytoplankton standing stock and GPP, such as grazing, nutrient availability and the physical structure of the water column, will have a strong bearing on pelagic CH₄ dynamics and resulting CH₄ emissions.

To our knowledge, our results represent the first ecosystemlevel estimates of methanogenesis in oxic freshwaters. For all treatments, intense CH₄ production was needed to sustain the observed patterns in CH₄ concentrations and to offset atmospheric losses. In the absence of methanogenesis, the decreases in enclosure CH₄ concentrations due solely to atmospheric evasion would have rapidly depleted the CH₄ pool (dashed lines, Fig. 1a). These calculations are potentially conservative, as we did not consider MOX. Considering all potential sources and sinks, we estimate an average rate of methanogenesis of $0.23 \, \mathrm{mmol} \, \mathrm{m}^{-3}$ per day (Table 1; Supplementary Table 2). This rate is higher than previous estimates from in vitro incubations for oligotrophic Lake Stechlin, Germany $(0.04-0.09 \, \text{mmol m}^{-3} \, \text{per day})^{18,19}$, but it should be noted that this estimate is very sensitive to the values of MOX used. *In-situ* MOX can vary dramatically across systems³¹, with depth, diurnally and with changes in light exposure 18,19. Therefore, it is possible that our estimates of MOX and methanogenesis may be biased by the fact that water from only one depth was used, and incubations were run in the dark (see Methods section for details). Consequently, we consider our estimates of net CH₄ production (excluding MOX; Fig. 1b) as a lower bound for epilimnetic methanogenesis, as these estimates are analogous to methanogenesis where MOX equals zero. In the

Table 1 G	reenhouse	gas dy	namics in	Lac (Cromwell.
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	CH ₄ and CO ₂ dynamics						
	(mmol m ^{- 3} per day)	(mmol m ^{- 2} per day)	(mol per lake per day)	(mmol CO ₂ eq m ⁻² per day)	(mol CO ₂ eq per lake per day)		
Pelagic production in enclosures	0.23 ± 0.01						
Pelagic diffusive fluxes from enclosures		0.15 ± 0.03	15.5 ± 2.6	1.4 ± 0.2	140 ± 24		
Pelagic diffusive fluxes from lake		0.78 ± 0.35	79.5 ± 31.1	7.1 ± 3.2	722 ± 325		
Ebullition from lake		0.31 ± 0.66	15.2 ± 32.5	2.8 ± 6.0	138 ± 295		
CO ₂ diffusion from lake				33.4 ± 12.4	$3,408 \pm 1,262$		

Both oxic water column production and diffusion of CH_4 in ambient (that is, control) experimental enclosures were estimated. Mean (± 1 s.d.) summertime (June-August inclusive) areal estimates of surface emissions and ebullition were determined for Lac Cromwell alongside the experiment. In all cases, CH_4 emissions were converted to CO_2 equivalents (where 1kg $CH_4 = 25$ kg CO_2 for a 100-year period) to compare with areal and whole-lake CO_2 emissions. Ebullition was not detected at depths >3 m, thereby restricting contributions to whole-lake flux. See Methods section for a detailed description of both enclosure and whole-lake calculations.

ambient enclosures, net production averaged 0.09 and ranged from 0.04 to 0.12 mmol m $^{-3}$ per day, and is in close agreement with estimates of methanogenesis in Lake Stechlin, where MOX was shown to be extremely low throughout the epilimnion^{18,19}. The fact that our approximation of pelagic methanogenesis based on mass balance (Fig. 1b; Table 1) is in good agreement with *in vitro*^{18,19} estimates is very promising, and suggests that both approaches could be incorporated into future detailed studies of CH₄ dynamics for other systems.

Our conclusion that acetate consumption supplied most CH₄ in the enclosures, based on estimates of α_{app} , is particularly sensitive to changes in the rate of MOX. Given the potential variability in MOX discussed above, estimates of α_{app} may be biased by our limited MOX measurement. However, scenario analyses estimating α_{app} from extreme hypothetical MOX rates (see Methods for calculations) were all well within the range of acetoclastic dominance (Fig. 3a). This supports our conclusion that acetate supplied the majority of CH₄ across our treatments, likely for the duration of the experiment (Fig. 3b). Consistent with this conclusion, observations of enriched surface δ^{13} C-CH₄ in numerous lakes^{8,10,18,29,32} have been hypothesized to indicate dominance of acetoclastic methanogenesis in oxic freshwater pelagic zones¹⁸, and both known acetoclastic genera (Methanosaeta and Methanosarcina) have recently been detected in oxic freshwater^{19,26} and terrestrial^{27,28} environments. Furthermore, acetoclastic dominance was enhanced with the addition of nutrients and increased algal production (Fig. 3a). This pattern parallels that from anoxic environments, where increased abundance of biologically young, high-quality algal acetoclastic over material promotes hydrogenotrophic methanogenesis, the latter instead supported by lower quality, aged terrestrial DOC^{30} . Finally, cyanobacteria were near absent in the mixed layer of our enclosures (<3.5% relative abundance, as biomass, estimated from one enclosure per treatment on day 7 in all treatments), ruling out major contributions of diazotroph-derived water-column H₂ for hydrogenotrophic methanogenesis¹⁹, and cyanobacterial methanogenesis during DOM demethylation²³. Taken together, it appears that acetoclastic methanogenesis linked to in-situ, algal DOC production, potentially plays a central role in supporting water column CH₄ production in oxic freshwaters.

The potential importance of acetate for oxic water methanogenesis is intriguing, as acetate concentrations are typically extremely low in oxic surface waters^{18,33–35}, and the presence of oxygen may inhibit acetate consumption²⁷. Yet past reports based on radioactive substrate additions have unequivocally demonstrated extremely rapid turnover of acetate in oxic environments across fresh and marine surface waters^{33–36}, suggesting active production of acetate from multiple metabolic pathways³⁷ in oxic environments, as well as very efficient uptake^{33–36}. Fermentative metabolism by diverse and widespread facultative anaerobic bacteria in particle-associated microanoxic zones could be the main source of water column acetate^{38–40}, and such microhabitats could be directly associated to algae or to particles¹⁹, yet in vitro work as well as our own results further suggest that methanogenesis proceeds even when these potential sites are removed 18,19. One possibility that should be further explored is that fermentative bacteria themselves create the conditions for methanogenesis through syntrophic interactions with acetoclastic methanogens in mixed cell aggregates⁴¹.

Is oxic water methanogenesis a significant component of the CH₄ budget of Lac Cromwell? We have combined our estimated rates of oxic water methanogenesis and CH₄ emissions in the ambient enclosures with measurements of diffusive and ebullitive CH₄ fluxes from the surrounding lake waters (summarized in Table 1), and conclude that methanogenesis in the oxic pelagic

zone may potentially contribute $\sim\!20\%$ of mean summertime $\rm CH_4$ diffusive fluxes, approximately the same magnitude as ebullitive fluxes measured for the entire lake. Careful extrapolation of our absolute emissions rates (Fig. 1b, Table 1) to other ecosystems should be made, as previously overlooked, system-specific differences in gas exchange coefficients $(k_{\rm CH_4})$ and non-Fickian fluxes will introduce error to conventional cross-system flux comparisons (see Supplementary Note 3 for extended coverage of this potential problem). Although oxic water methanogenesis contributed $\sim\!4\%$ to Lac Cromwell's summertime greenhouse gas footprint, it is conceivable that this pathway could have greater relative importance in more nutrient-rich, productive ecosystems, and in hardwater environments where low to negative (that is, net $\rm CO_2$ uptake) $\rm CO_2$ emissions are common $^{42-44}$.

Within Lac Cromwell, it is interesting to note that oxic water methanogenesis contributed a significant component of wholelake emissions, despite the fact that it is a small (0.1 km²), relatively shallow (~ 3.5 m mean depth) lake with an extensive littoral area, and a CH₄-rich anoxic hypolimnion, all of which are typically major sources of CH₄. A corollary is that in large, deep lakes and open ocean sites where surface waters are increasingly decoupled from benthic or littoral sources of CH₄, surface CH₄ concentrations should be mostly driven by pelagic methanogenesis and therefore a function of algal biomass and metabolism. In this regard, whereas chlorophyll a does not explain the large-scale patterns in surface CH₄ across lakes in general⁴⁵, the relationship we observed between surface water CH₄ and chlorophyll a in our isolated enclosures $(\log_{10}[CH_4] = 0.46\log_{10}[chl \ a] + 0.68,$ $r^2 = 0.68$, n = 4) generally agrees with observations specifically from large lakes, extending all the way to ultraoligotrophic open ocean regions (Fig. 4; $\log_{10}[CH_4] = 0.99\log_{10}[chl \ a] - 1.63$, $r^2 = 0.73$). This continuity suggests that algal-linked pelagic CH₄ production may explain much of the ambient CH₄ dynamics observed in large and deep aquatic ecosystems, regardless of the fact that different mechanisms have been identified as potentially important in marine (that is, methylphosphonate or methanethiol catabolism)^{15,16,23–25} versus freshwater (hydrogenotrophic or acetoclastic methanogenesis; Fig. 3)^{18,19}. There are clearly a number of potential sources of CH₄ in oxic pelagic waters, and although it remains unknown how each pathway differs in relative importance across these diverse aquatic environments, our results highlight a consistent relationship between open-water phytoplankton dynamics and the regulation of oxic pelagic CH₄.

Taken together, our findings suggest that pelagic CH₄ production in oxic environments may be widespread, and likely supports a baseline evasion of CH4 from all oxic water columns. In small and shallow ecosystems, such as Lac Cromwell, this pathway will be quantitatively significant but secondary relative to anoxic littoral and benthic sources. In large and deep aquatic environments, however, this pathway could become the single dominant source fueling CH₄ supersaturation and fluxes to the atmosphere. It should be emphasized that consideration of oxic water column methanogenesis does not necessarily lead to higher flux estimates for lakes, as the CH₄ derived from this process is already included in routine measurements of surface water pCH₄ and CH₄ surface fluxes²-7,32,46. Incorporating the origin of the CH₄ that outfluxes to the atmosphere from these lakes, however, is critical to our understanding of the regulation of these fluxes and our capacity to predict their future change. In this regard, there are potentially large global implications for algal-driven oxic-water methanogenesis. CH₄ emissions from surface waters, particularly from freshwaters^{2–4}, are a major element of the global atmospheric CH4 budget, and we suggest here that the algal-mediated baseline flux is not only a major contributor to these overall aquatic CH₄ emissions, but also one that

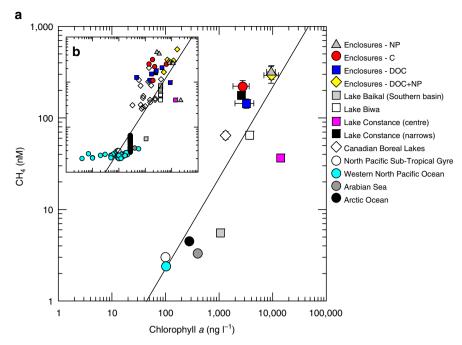


Figure 4 | Linking dissolved CH₄ and algal biomass across diverse open-water aquatic ecosystems. Here we combine experimental data from this study with literature-derived measurements from large (>10 km²), deep (>10 m maximum depth) lakes and marine environments where both CH₄ and chlorophyll a samples were simultaneously measured. A strong positive relationship between CH₄ and chlorophyll a exists across systems: (**a**) averaged data ($[\log_{10}x] = 0.993[\log_{10}y] - 1.630$, $r^2 = 0.73$, P < 0.0001), where error bars represent ± 1 s.e.m.; (**b**) raw data ($[\log_{10}x] = 0.855[\log_{10}y] - 1.166$, $r^2 = 0.72$, P < 0.0001). Guidelines for meta-analysis are discussed in the Methods section, and raw data are available in Supplementary Table 3.

is particularly sensitive to environmental change. As such, widespread and intensifying human- and climate-driven changes in pelagic nutrient availability^{47–49}, terrestrial DOC inputs^{50,51} and physical structure of the water column^{47,48,51}, which strongly shape aquatic algal dynamics^{47,51–54}, may have major, but previously unconsidered, consequences for oxic water methanogenesis and aquatic CH₄ emissions.

Methods

Study site. Lac Cromwell is located at the Station de Biologie des Laurentides, a field research facility of the Université de Montréal. The lake turns over in spring and fall, is relatively shallow and small (mean depth = 3.5 m, maximum depth = 9.5 m, surface area of 0.11 km², volume of 3×10^5 m³), and is oligomesotrophic, with little phytoplankton biomass (~ 6.0 µg of chl a per litre)⁵⁵.

Experimental design. Here, 12, 1 m diameter, 6 m deep, 4712-L polyethylene enclosures were attached to floating wooden frames anchored in the middle of the lake (\sim 8 m depth). On 5 June 2012, bags were filled by pumping epilimnetic water sequentially across two mesh screens of 110 and 45 μm to remove zooplankton, and three mesocosms each received N+P, DOC, N+P+DOC additions, or no addition (control). Both KH₂PO₄ and NaNO₃ additions were made to reach target concentrations of 50 and 700 μg P and N per litre, respectively. For DOC, Superhume brand humic slurry was added to reach a target concentration of 15 mg of DOC per litre. For a full review of Superhume properties and applicability to freshwater experimentation, see Lennon *et al.*⁵⁶. The mesocosms were allowed to stabilize for 1 week before initial sampling and re-stocking zooplankton at ambient lake concentrations. Zooplanktons were collected from Lac Cromwell by vertical tows using a 54-μm Nitex net, and immediately released into mesocosms after each haul.

Limnological measurements. All limnological samples were gathered at $\sim\!1200$ hours on days 0, 7 and 28 of the experiment (13, 19 June 2012 and 10 July 2012, respectively; Supplementary Fig. 3). Dissolved oxygen (DO) and temperature were measured at 0.5 m depth in all bags, whereas water column profiles were taken from one enclosure per treatment using an Yellow Springs Instruments (YSI) Pro Plus multiparameter sonde (Supplementary Fig. 1). On all three dates, water was collected at 0.5 m depth, filtered to 0.45 μ m and chlorophyll a was measured spectrophotometrically in ethanol filter extracts. Dissolved CH₄ concentrations at 25 cm depth were measured in triplicate for each mesocosm following Prairie and del Giorgio 5 . Partial pressures were converted to concentrations using Henry's

constant corrected for temperature. On day 7, dark incubations to estimate MOX using unfiltered surface water were conducted in the laboratory at room temperature, lasting 96 h (Supplementary Fig. 4). Incubations began at approximately noon on 18 June, day 6 of the experiment; samples for measurement of MOX were collected from one mesocosm per treatment, at 25 cm depth, into 12 ml precombusted vials. Samples were capped to exclude any air with a gastight, 3-mm-thick, butyl rubber-lined plastic cap, then incubated in the dark at 20 °C. Ambient CH₄ measured in the mesocosms was used as a time 0 concentration, and samples were subsequently killed every 24 h by injection of HgCl₂, and stored at 4 °C in the dark until analysis. Upon analysis, 5 ml of water was displaced with ambient air, and the remaining sample was equilibrated with the air by physically shaking the vials. Headspace equilibration and gas chromatography analysis were performed as for ambient CH4. Respiration rates were determined by dark in situ, 24 h incubations on experiment days 6 and 7 in all enclosures, using unfiltered water in 4-l cubitainers. Initial and final samples of DO were collected and measured by membrane inlet mass spectrometry⁵⁷.

CH₄ **chamber flux estimates.** Air–water CH₄ fluxes in the mesocosms and the surrounding lake were calculated by combining the measured surface water pCH₄ and the CH₄ gas exchange coefficient ($k_{\rm CH4}$), the latter derived from the measured gas exchange coefficient for CO₂ ($k_{\rm CO2}$) in mesocosms supersaturated in CO₂. $k_{\rm CO2}$ was measured in one mesocosm per treatment, 3–5 times daily during days 6 and 27–28 of the experiment, using the floating chamber method following Vachon et al. ⁵⁸ Briefly, chambers were connected via a closed loop system to an EGM-4 infrared gas monitor (PP Systems). Fluxes were calculated as the rate of change of chamber pCO₂ per min during a 15-min interval. Diel sampling periods were spaced to obtain flux estimates from the morning, afternoon and nighttime periods, and mean values were used for daily flux estimates. Dissolved CO₂ samples were taken before all chamber measurements using the same headspace technique as for CH₄, but with direct injection of the equilibrated gas into the EGM-4 in the field. The gas exchange coefficient for CH₄ was then calculated from kCO₂ using equation (1), following Jahne et aL. ⁵⁹:

$$k_{\text{CH}_4} = k_{\text{CO}_2} * \left(Sc_{\text{CH}_4} * Sc_{\text{CO}_2}^{-1} \right)^{-2/3}$$
 (1)

where Sc is the Schmidt number of CO₂ and CH₄, respectively⁶⁰. To calculate CH₄ fluxes, we used these average k_{CH_4} values, the measured $p\text{CH}_4$ and the temperature-dependent solubility of the gas following equation (2):

$$F = k_{\text{gas}} * K_{\text{h}} * \Delta_{\text{gas}} \tag{2}$$

where F is flux of CH₄ (mmol m⁻² per day), K_h is the temperature-corrected Henry's constant and Δ_{gas} is the difference in partial pressures between the air and

the water partial pressures of $\mathrm{CH_4}$. We assumed an atmospheric partial pressure of $\mathrm{CH_4}$ of 1.75 p.p.m.

Isotopic analyses. Isotope compositions are reported in delta notation following equation (3):

$$\delta^{13} C \text{ or } \delta^{18} O = ((R_{\text{sample}}/R_{\text{standard}}) - 1) * 1,000$$
 (3)

where R is the ratio of heavy to light isotope, $\delta^{13}C$ represents $\delta^{13}C$ –CH₄ or ^{13}C of dissolved inorganic carbon (DIC), $\delta^{18}O$ represents $\delta^{18}O$ –O₂ or $\delta^{18}O$ –H₂O and $R_{\rm standard}$ is the $\delta^{13}C$ or $\delta^{18}O$ signature of Vienna Pee Dee Belemnite or standard mean ocean water, respectively. All samples were taken at \sim 25 cm depth. Samples for $\delta^{18}O$ –O₂ and $\delta^{13}C$ –CH₄ were stored in 12-ml precombusted borosilicate vials, preserved with HgCl₂, and air-free samples capped with a gastight rubber-lined plastic cap. $\delta^{18}O$ –O₂ samples were analysed at the University of Ottawa Stable Isotopes Laboratory, and $\delta^{13}C$ –CH₄ samples were measured at the University of Waterloo, both using standard laboratory methods. A single sample for $\delta^{18}O$ –H₂O was collected from the lake before filling enclosures, analysed with a Picarro L230i isotopic water analyser, and assumed representative for all enclosures for the duration of the experiment. $\delta^{13}C$ –DIC samples were collected in acidwashed, 40 ml vials with both Teflon and rubber-lined gastight plastic caps, and measured at the Colorado Plateau Stable Isotopes Laboratory using standard methods. Day 7 respiration (R), $\delta^{18}O$ –O₂ and $\delta^{18}O$ –H₂O were used to estimate GPP and NEP following Quiñones-Rivera *et al.*61

Numerical methods. We used RM-ANOVA to compare the effects of treatments, and treatment by time interactions, on ambient CH4. One-way ANOVA was used to assess differences between treatments for grouped data, and $log_{10}(x+1)$ transformations applied to maintain the homogeneity of variances, followed by Bonferroni post-hoc pairwise comparisons of treatment means. All regression analyses were performed using Sigmaplot version 12, and ANOVAs were computed with SPSS version 16. We used ordinary least squares regression to quantify the link between GPP or NEP and CH₄ evasion. To estimate rates of MOX, the loss of CH4 through time was fitted with a polynomial function, the derivative was taken and the slope at time 0 was calculated (Supplementary Fig. 4). This approach was deemed more accurate in quantifying the slope at time 0, as traditional methods, such as log transformations followed by visual selection of the linear portion of the data set³¹, can introduce large errors in the slope of the regression line when deciding which points to fit and which to exclude. Although MOX was estimated only once, our estimates are within the range of pelagic MOX $(0.170-0.250~\mathrm{mmol\,m^{-3}}$ per day) previously reported \$\frac{31,32,62}{3}\$, particularly for boreal lakes of similar trophic condition⁶².

Apparent isotopic fractionation of methanogenesis. We estimated the apparent fractionation factor (α_{app}) during methanogenesis following Conrad³⁰. The apparent fractionation factor is defined using equation (4) as

$$\alpha_{app} = (\delta^{13}\text{C-CO}_2 + 10^3)/(\delta^{13}\text{C-CH}_{4\text{source}} + 10^3)$$
 (4)

where $\delta^{13}\text{C-CH}_{4\text{source}}$ is the isotopic signature of source CH₄, and $\delta^{13}\text{C-CO}_2$ was calculated from measured $\delta^{13}\text{C-DIC}$ following the study by Stumm and Morgan⁶³, using fractionation factors from the study by Mook *et al.*⁶⁴. As we measured ambient $\delta^{13}\text{C-CH}_4$ ($\delta^{13}\text{C-CH}_{4\text{ambient}}$), we estimated $\delta^{13}\text{C-CH}_{4\text{source}}$ by correcting for the isotopic fractionation effects of MOX and evasion to the atmosphere using an open-system, steady-state model for isotope fractionation⁶⁵, as defined in equation (5):

$$\delta^{13}\text{C-CH}_{4\text{source}} = f\left(\delta^{13}\text{C-CH}_{4\text{ambient}} - \Delta_{\text{evasion}}\right) + (1 - f)\left(\delta^{13}\text{C-CH}_{4\text{ambient}} - \Delta_{\text{MOX}}\right)$$
(5)

where f and 1-f are the fractions of CH₄ loss, standardized to daily rates for the entire enclosure, estimated, respectively, as evasion and MOX rates divided by the sum of each. Δ is the isotopic effect of each loss pathway. In cases such as ours where $\Delta < 100\%$, Δ can be approximated using equation (6):

$$\Delta = (1 - \alpha)10^3 \tag{6}$$

where α is the fractionation factor of a given reaction. Literature-derived values of α were used for both loss pathways. For evasion, an α value of 0.9992 was used⁶⁶, and for MOX, the range reported by Bastviken *et al.*⁶² (0.9816–0.9792) was used.

To better constrain the potential range of α_{app} estimated on day 7 of our experiment, we performed a scenario analysis by varying combinations of MOX rates and isotopic fractionation values, to determine the sensitivity of our estimates of α_{app} to potential variability in MOX. For this analysis, we chose two very conservative, extreme (for oxic pelagic freshwaters) values in MOX rates (0 and 0.05 mmol m $^{-3}$ h $^{-1}$) and paired each rate with both minimum and maximum values reported for isotopic fractionation during MOX⁶² (18.4 and 20.8‰). Because the MOX rate of zero did not have associated isotopic fractionation, we calculated three different combinations of extreme parameters: no oxidation with no fractionation, high oxidation with low fractionation and high oxidation rates with high associated fractionation. We then took the maximum and minimum

estimates of α_{app} , based on this analysis, as upper and lower bounds for α_{app} estimated using our actual measurements of MOX.

Mesocosm mass balances. To quantify methanogenesis in the oxic mixed layer of ambient enclosures, mass balances from the ambient control treatment enclosures were constructed (Supplementary Table 2) for comparison with other lake-based CH₄ fluxes using equation (7):

$$CH_{4gross} = \Delta_{CH4} + E + O - M \tag{7}$$

Here rates of methanogenesis ($\mathrm{CH_{4gross}}$) are estimated by summing the rate of change in storage (Δ_{CH4}), defined as the rate of increase in the ambient $\mathrm{CH_{4}}$ pool, with the rates of $\mathrm{CH_{4}}$ evasion (E) and oxidation (O), minus the horizontal influx of external $\mathrm{CH_{4}}$ via diffusion across the walls of the enclosures (M; detailed in the Supplementary Methods). In all cases, mass balance terms were first standardized to the entire mixed layer or surface of the enclosure (as mmol per enclosure per day) in order to combine aerial (E and M) and volumetric (Δ_{CH4} and O) processes on each sampling date.

The enclosure mixed layer average depth of 1.68 m (Supplementary Fig. 1) was used to calculate both mixed layer volume, and the surface area of the enclosure wall that contributed to CH₄ influx from the surrounding lake environment. The change in storage ($\Delta_{\rm CH_4}$) was calculated as the rate of change in ambient CH₄ concentrations in each bag as a function of time using the slope of the regression line as determined by least-squares regression analysis, then averaged and adjusted volumetrically to the enclosure scale (mmol per enclosure per day; Supplementary Table 2). The rate of MOX (O) for day 7 of the experiment was used for all calculations. Finally, estimates of CH_{4gross} were converted to units of mmol m⁻³ per day for comparison with other lake-based measurements (Table 1).

Whole-lake CH4 dynamics. Enclosure measurements and mass balance results were compared with the surrounding Lac Cromwell environment. To quantify summertime rates of ebullition in Lac Cromwell (Table 1), bubble traps were fixed at five permanent sampling sites along a transect from the littoral to pelagic at \sim 1, 2, 3, 5 and 7 m depths. Bubble traps were left in place over the entire sampling period and collected monthly in June, July and August 2012. Traps consisted of an inverted funnel (63.5 cm diameter) suspended at 0.5 m below the surface of the water (0.25 m at the shallowest sample site). A graduated 1-l glass bottle was attached to the funnel by gluing the sides of the bottle cap to the neck of the funnel to make a gastight seal. The bottles were covered with aluminum foil to prevent overheating or light exposure of the collected gas. Bottles were filled with water upon deployment, so the gas accumulated by displacing water in the bottle. Bottles were collected once a month or until a volume greater than 250 ml of gas was detected. All gas was carefully removed by displacement with ambient water using a stopcock cap fitted with two syringes. A subset of the gas was injected in 30 ml saline vials, and analysed upon return to the laboratory by gas chromatographic analysis as detailed for enclosure samples. As the majority of samples settled in the bottles for up to a month before collection, we assumed that the composition of CH₄ in bubbles was 0.6 atm or 60%, based on a subset of samples analysed that were collected within days of sedimentary release. This assumption is consistent with literature estimates of bubble $\mathrm{CH_4}$ concentration³².

Diffusive evasion of lake CH₄ was estimated by first measuring concentrations of CH₄ on a monthly basis near each bubble trap site along the transect. Concentrations ranged between 0.63 and 2.09 with an average of 1.21 μ M, and did not vary across sites (ANOVA, $P=0.98,\,F=0.10$). As detailed above for the enclosure mass balance, atmospheric fluxes were estimated by applying the average 24 h gas exchange coefficient measured at 7 m depth along the transect (mean $k_{\rm CH_4}=0.65\,\rm m$ per day, Supplementary Fig. 2).

Whole-lake CH_4 and CO_2 diffusive fluxes were calculated by applying average areal estimates of diffusion to the entire lake surface area (102,000 m²). As ebullition was only detected at depths <3 m, average rates of ebullition (0.31 \pm 0.66 mmol m $^{-2}$ per day) were applied to the lake surface <3 m in depth (49,166 m²). This area was estimated by determining the area of the lake <3 m in depth from a hypsographic curve for Lac Cromwell⁶⁷, and existing bathymetric information for Lac Cromwell⁶⁸.

Data compilation for meta-analysis. To assess the relationship between algal dynamics and $\mathrm{CH_4}$ in open water environments, we compared the results from our experimental enclosures with directly obtained results from large (> $10\,\mathrm{km^2}$), deep (> $10\,\mathrm{m}$ maximum depth) boreal lakes 45 and other freshwater and marine data from the literature (raw data and selection methods tabulated in Supplementary Table 3). Literature data include only studies where $\mathrm{CH_4}$ and chlorophyll a were measured simultaneously, except for Lake Baikal, where detailed studies of $\mathrm{CH_4}$ and chlorophyll a were measured independently, but overlapped temporally and thus were included. To remain consistent with the sampling approach from our study, and sampling of boreal lakes, only near-surface, open-water results from the centre of the lake or furthest from the coastal environment (in marine studies) were used. Summertime results were taken if seasonal data were presented. If surface water results were given as a range, the midpoint of that range was chosen. If multiple samples were taken at one site, then results were averaged.

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Acknowledgements

Simon Gauthier-Fautaux, Juan Pablo Nino Garcia, Cynthia Soued, Marilyne Robidoux and Ryan Hutchins provided field and laboratory assistance. Jean-Francois Lapierre, Dominic Vachon, Adam Heathcote and David Bastviken provided conceptual and methodological advice. Annick St. Pierre, Alice Parks and the employees of the Station de

biologie des Laurentides de l'Université de Montréal provided logistical support. Terhi Rasilo and Tonya Del Sontro revised an early version of the manuscript. M.J.B. was supported by a doctoral scholarship from the National Science and Engineering Research Council of Canada (NSERC). This project is part of the programme of the NSERC/HQ Industrial Research Chair in Carbon Biogeochemistry in Boreal Aquatic Systems (CarBBAS), and was co-funded by grants from NSERC (to P.A.d.G. and A.M.D.), Hydro-Québec (to P.A.d.G.) and a Fonds de recherche Québec-Nature et Technologie award from the government of Québec (to A.M.D.).

Author Contributions

M.J.B, P.A.d.G. and A.M.D. designed the study; M.J.B., L.B., M.C.G.C. and A.M. performed field and lab work; M.J.B. and Y.T.P. analysed the data; M.J.B. and P.A.d.G. wrote the paper; all authors revised the paper; A.M.D. and P.A.d.G. contributed materials.

Additional information

Supplementary Information accompanies this paper at http://www.nature.com/naturecommunications

Competing financial interests: The authors declare no competing financial interests

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How to cite this article: Bogard, M. J. *et al.* Oxic water column methanogenesis as a major component of aquatic CH₄ fluxes. *Nat. Commun.* 5:5350 doi: 10.1038/ncomms6350 (2014).