



Relationships between greenhouse gas production and landscape position during short-term permafrost thaw under anaerobic conditions in the Lena Delta

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Abstract. Soils in the permafrost region have acted as carbon sinks for thousands of years. As a result of global warming, permafrost soils are thawing and will potentially release greenhouse gases (GHGs) such as methane (CH₄) and carbon dioxide (CO₂). However, small-scale spatial heterogeneities of GHG production have been neglected in previous incubation studies. Here, we used an anaerobic incubation experiment to simulate permafrost thaw along a transect from upland Yedoma to the floodplain on Kurungnakh Island. Potential CO₂ and CH₄ production was measured during incubation of the active layer and permafrost soils at 4 and 20 °C, first for 60 d (approximate length of the growing season) and then continuing for 1 year. An assessment of methanogen abundance was performed in parallel for the first 60 d. Yedoma samples from upland and slope cores remained in a lag phase during the growing season simulation, while those located in the floodplain showed high production of CH₄ ($6.5 \times 10^3 \mu\text{g CH}_4\text{-C g}^{-1} \text{C}$) and CO₂ ($6.9 \times 10^3 \mu\text{g CO}_2\text{-C g}^{-1} \text{C}$) at 20 °C. The Yedoma samples from the permafrost layer started producing CH₄ after 6 months of incubation. We conclude that landscape position is a key factor triggering CH₄ production during the growing season time on Kurungnakh Island.

1 Introduction

For the past decades, scientists have warned about the effects of global climate change (IPCC, 2021). The effects of this warming will be pronounced in the polar regions, where the air temperature increase over the last 50 years is already 3 times higher than the increase in the global average for the same period (AMAP, 2021; Rantanen et al., 2022). This particularly affects soils in northern high-latitude permafrost regions, which cover 14.6 % of the Northern Hemisphere (Obu et al., 2019) and contain 1300 Pg of organic carbon (C) (Hugelius et al., 2014a). A majority of this C (822 Pg) is stored in permafrost (Hugelius et al., 2014b), which is defined as ground where the temperature remains at or below 0 °C for more than 2 consecutive years (Washburn, 1973). Due to low temperatures, the organic matter (OM) stored in permafrost soils is characterized by a low decomposition rate (Davidson and Janssens, 2006). However, during summer, the upper part of the permafrost-affected soils thaws (active layer) and allows OM decomposition (Lee et al., 2012). With climate change, permafrost thaw will likely increase and lead to higher OM decomposition rates, releasing greenhouse gases (GHGs) like carbon dioxide (CO₂) and methane (CH₄) (Wagner et al., 2007; Schuur et al., 2015; Knoblauch et al., 2018). This turnover might lead to the transformation of Arctic soils from C sinks to C sources (Koven et al., 2011; Dean et al., 2018; Lara et al., 2019).

Carbon emissions, and mainly CH₄ emissions, greatly vary across the Arctic and especially on small scales (Treat et al., 2018; Lara et al., 2019; Elder et al., 2020). Treat et

al. (2018) showed that C flux variability was strongly associated with specific landscape types due to differences in soil moisture and site drainage in uplands and wetlands that also controlled vegetation communities, which we refer to here as landscape position. Landscape change due to permafrost thaw is also highly affected by landscape position; low-lying ice-rich areas can become waterlogged following permafrost thaw, while higher areas can be drained by water run-off (Osterkamp et al., 2009; Liljedahl et al., 2016). The water-logged areas like thermokarst lakes or wetlands have been identified as CH₄ emission hotspots (Olefeldt et al., 2013; Treat et al., 2018; Kuhn et al., 2021) because of the anaerobic conditions that favor methanogen communities (Conrad, 2002; Yavitt et al., 2006). On the other hand, well-drained sites, such as upland tundra, have the capacity to offset CH₄ emissions by acting as CH₄ sinks due to net oxidation in the surface soil layer (Juncher Jørgensen et al., 2015; Treat et al., 2018). Hence, after permafrost thaw, the redox conditions, determined by the landscape position, lead to different microbial communities and ultimately different CH₄ emissions (McCalley et al., 2014).

To quantify CH₄ and CO₂ production and to understand C turnover from thawing permafrost, numerous incubation studies have been carried out (Lee et al., 2012; Knoblauch et al., 2018; Walz et al., 2018; Holm et al., 2020). Studies have shown that C decomposition depends on several factors, such as organic C quantity, OM quality, temperature, and oxygen availability in the soil (Ganzert et al., 2007; Lee et al., 2012; Schädel et al., 2014; Treat et al., 2015; Knoblauch et al., 2018). In addition, Treat et al. (2015) highlighted that CH₄ production differences were partly explained by the landscape position, with differences between uplands, wetlands, floodplain soils, lowlands, and drained lake basins. For incubation under aerobic conditions, Kuhry et al. (2020) demonstrated that landscape types based on soil type (peaty wetlands, mineral soils) and the origin of the deposits (peat deposits, alluvial deposits) gave a good estimation of soil organic matter (SOM) lability and therefore explained differences in CO₂ production better than using only the %C, which is a commonly used metric for C quality across incubation studies (Treat et al., 2015). However, few studies have specifically focused on how the landscape position affects CO₂ and CH₄ production under anaerobic conditions and whether landscape position is a good indicator of CO₂ and CH₄ production under anaerobic conditions, as might be expected from field observations of CO₂ and CH₄ fluxes (Treat et al., 2018; Elder et al., 2020).

Besides landscape position, climate change affects the environmental factors in the study region. It modifies weather conditions and plays a key role in controlling rain events (frequency and intensity) (Callaghan et al., 2010; Tabari, 2020; Wang et al., 2021; Fewster et al., 2022). During the past 60 years, precipitation in Siberia has increased by 2.6 mm per decade (Wang et al., 2021). This finding likely leads to wetter conditions during the growing season in Siberia, and there-

fore, soil moisture increases. Changes in soil moisture will impact vegetation cover and soil redox conditions. Increasing precipitation and warming will also lead to deepening of the active layer (Zhu et al., 2017; Douglas et al., 2020) and, hence, the release of bioavailable C from the upper part of the permafrost layers. Waldrop et al. (2010) identified more labile C in shallow permafrost than in the active layer, which fueled more CO₂ production in an incubation experiment. On the other hand, other incubation experiments showed higher C turnover and more CO₂ production in the active layer than in the permafrost (Walz et al., 2017; Faucherre et al., 2018). Regarding CH₄ production, incubation studies tend to show higher CH₄ production in the active layer than in shallow permafrost (Treat et al., 2015), but some studies also measured the opposite behavior among their samples (Wagner et al., 2007; Waldrop et al., 2010). Therefore, it is still unclear how much CO₂ and CH₄ can be produced from shallow permafrost. Furthermore, high CH₄ production heterogeneity and long lag times have been measured with samples from Kurungnakh Island, Lena Delta, Russia (Knoblauch et al., 2013, 2018). Hence, the question remains whether the methanogen communities will have the time to become active during the short growing season (60 d) under anaerobic conditions.

The aim of this study was to simulate permafrost thaw under wet growing-season conditions across different landscape units in the Lena Delta and to measure CO₂ and CH₄ production. Here, we incubated upland Yedoma and adjacent lowland floodplain samples under anaerobic conditions and focused on the relationships between GHG production and microbial abundance shifts following short-term (60 d, growing-season length) and longer-term (1 year) permafrost thaw. The objectives of the study were to (1) quantify CH₄ and CO₂ production over 1 year under anaerobic incubation, (2) establish relationships between CH₄ and CO₂ production and methanogen abundances, and (3) characterize the role of the landscape position in gas production in thawed permafrost soils during a growing-season time frame.

2 Materials and methods

2.1 Site description and sampling

Soil samples were collected in August 2018 on Kurungnakh Island (72.333° N, 126.283° E), Lena Delta, Siberia (Fig. 1). Kurungnakh Island is located in the continuous permafrost zone and is an erosional remnant of Late Pleistocene deposits, characterized by ice- and organic-rich sediments (Grigoriev, 1993; Schwamborn et al., 2002); most of the island is composed of fluvial sandy sediments and Yedoma ice complex (IC) deposits. The IC is made up of ice-saturated sediments (65 % to 90 %) composed of cryoturbated silty sands and peaty deposits of Holocene origin (Schwamborn et al., 2002; Schirrmeister et al., 2011, 2013). Sediments from

the Yedoma IC contain, on average, 3 % total organic carbon (TOC) (Strauss et al., 2013a); however, IC sediments can include organic-rich layers, with TOC content reaching more than 20 % in layers contained within the ice complex sediments (e.g., buried peat horizons; Andreev et al., 2009). Kurunghakh Island is characterized by thermokarst lakes and wetlands due to thermo-erosional activity (Morgenstern et al., 2021). Samples were also collected in the young Kurunghakh Island floodplain area. The young and active floodplains in the Lena River Delta are of Holocene deltaic origin and are composed of stratified middle to fine sands and silts with layers of autochthonous peat and allochthonous OM (Schwamborn et al., 2002; Boike et al., 2013).

The soil sampling was carried out in two stages. First, the active layer was extracted using a spade, and active-layer samples were collected using a fixed-volume cylinder (250 cm³). Then, permafrost soil cores were sampled to a depth of 1 m below the surface by drilling with a modified snow, ice, and permafrost (SPIRE) auger (Jon Holmgren's Machine Shop, Alaska, USA). For this study, three cores were selected; two were from the Yedoma deposits (P15 and P16), and one belonged to a floodplain area (P17). The three coring locations were on a well-drained upland soil (P15; Fig. S1 in the Supplement), on a north-eastern facing slope (P16), and on a floodplain (P17; Fig. 1). The floodplain samples were taken in the highest part of the floodplain, 5 m above the Lena River water level.

These cores were chosen on the basis of geographical proximity to each other, landscape position, moisture gradient, and ice composition. The three cores had an organic layer ranging between 3 to 7 cm (Yedoma and floodplain, respectively). Below this organic surface layer, the soil cores were identified as mineral soil. The permafrost layers from the Yedoma cores were ice-rich, while no visible ice structure was seen for the floodplain core (Table 1).

Cores were described and subsampled in the field. Detailed core descriptions are presented in Table 1 and Table S1 in the Supplement. For the purpose of our study, we chose two samples from each core, one from the active layer and one from the frozen layer (above 1 m depth to simulate shallow permafrost thaw) (Table S1 in the Supplement). Care was taken not to select samples from the top of the active layer in order to avoid the top organic layers. Cores were subsampled in a climate chamber at −4 °C with a hammer and a chisel instead of a saw to limit contamination.

2.2 Sedimentary and geochemical characterization

We characterized the samples for soil texture, C and nitrogen contents, water content, conductivity, and pH. First, samples were thawed at 4 °C overnight; then the pore water was extracted with a Rhizon soil moisture sampler (Rhizon MOM 0.6 µm, Rhizosphere Research Products) (Meijboom and van Noordwijk, 1991). Electrical conductivity and pH were measured from pore water. Prior to further analyses, soil sam-

ples were freeze-dried, and the absolute water content (Eq. 1) was calculated. For TOC weight percent, total carbon (TC) and total nitrogen (TN) subsamples were homogenized and measured with a carbon–nitrogen–sulfur (CNS) analyzer (Elementar Vario EL III). Each subsample was measured in duplicate, and standards and blanks were used to ensure reliable analytical measurements. The bulk density was determined based on a transfer function between absolute water content and bulk density made by Fuchs (2019) (Fig. S2 in the Supplement). Since most of the samples used to establish this correlation came from the same area as our samples, we assumed that the transfer function was applicable to our samples.

$$\Theta = \frac{W_w - W_d}{W_w}, \quad (1)$$

where Θ is the water content, W_w is the wet weight, and W_d is the dry weight.

Carbon storage was calculated by multiplying the TOC contents with the bulk density and then dividing by the sample length.

Another subsample was used for grain size characterization. The grain size analysis was conducted using a laser diffraction particle size analyzer (Mastersizer 3000, Malvern Panalytical, Malvern, UK). Prior to measuring, subsamples were put on a heated shaker for 3 weeks, and H₂O₂ was added daily to remove the organic materials. The samples were measured in a wet dispersion unit, and at least three subsamples from each sample were measured. The average grain size distribution (in vol %) was calculated from the measured replicates.

2.3 Incubation setup and substrate addition

To mimic a wet growing season, the samples were first incubated under anaerobic conditions for 60 d at two different temperatures, 4 and 20 °C. Since 11 samples out of 12 did not produce CH₄ after 2 months of incubation, we extended the incubation time to 363 d to see whether the other cores would produce CH₄. For every sample, three replicates were incubated, resulting in a total of 36 samples. Prior to incubation, the samples were thawed at 4 °C and prepared under oxygen-free conditions using an anoxic glovebox. The samples were homogenized, and 13 g of wet soil was collected and inserted into a 120 mL vial. Sterilized tap water was added only to samples with a gravimetric moisture content of less than 30 % to limit the effect of gas dissolution (Henry's law). The amount of sterilized tap water was calculated to reach 30 % gravimetric moisture based on the original water content and the weight (wet and dry). The flasks were closed with rubber stoppers and aluminum lids. The headspace of the samples was flushed with pure nitrogen for 3 min to remove potential O₂ inside the vials. Then the samples were incubated in the dark.

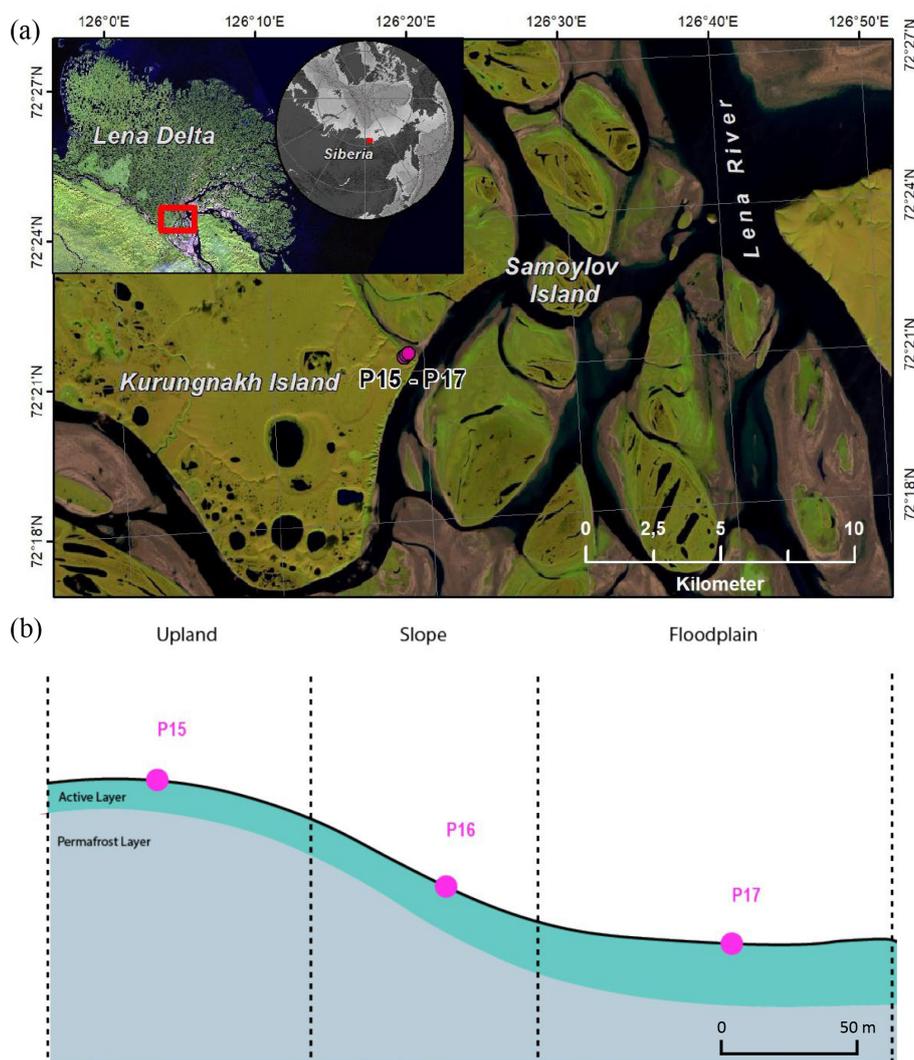


Figure 1. Location of Kurungnakh Island in the Lena Delta (Siberia). The location of the cores used for the study are indicated on the map (a) and along a schematic transect (b). Samples were taken during the Lena summer expedition in 2018.

After 60 d of incubation, $0.7 \text{ mg glucose g}^{-1}$ dry sample weight was added to two of the three replicates to understand the effect of potential substrate limitation in the soil system. The glucose was diluted with Milli-Q water to obtain a 100 g L^{-1} solution. Solutions were injected via syringe to minimize soil disturbance (Pegoraro et al., 2019). The same amount of water as in the glucose solution was added to the third replicate to ensure that differences in gas production were only due to the addition of glucose (Pegoraro et al., 2019; Adamczyk et al., 2021). The glucose addition was also carried out under oxygen-free conditions.

The effects of glucose are usually observed within less than 48 h (Yavitt et al., 1997; Pegoraro et al., 2019). Therefore, after the glucose addition, the gas was measured daily for 1 week (described in the following section). As the first injection had little effect on gas production, a second injection

(day 64) was added with twice the amount of glucose solution ($1.4 \text{ mg glucose g}^{-1}$ dry sample weight).

2.4 Gas analyses

CO_2 and CH_4 in the headspace were measured with a gas chromatograph (GC) (7890A, Agilent Technologies, USA) with flame ionization detection (FID). The temperature in the column was 50°C with a flow of 15 mL min^{-1} and a runtime of 4.5 min. Helium was used as a carrier gas. A Hamilton syringe was used to introduce $250 \mu\text{L}$ of the sampling gas into the GC. For the first week, measurements were made every 2 d, then twice a week for 3 weeks, then once a week until day 60. The incubation vials were flushed when either the CH_4 or CO_2 concentration reached 10^4 ppm to avoid gas saturation inside the flask. The production rate was calculated with the change in concentration of CO_2 and CH_4 over the in-

incubation time. First, the measured CO₂ and CH₄ concentrations were converted from ppmv to μmol L⁻¹ using the ideal gas law, then a linear regression between each measurement point was used to calculate the change in concentration over time. The production rate was calculated using the change in concentration over time from the linear regression, then the rates were normalized using the volume of the soil (for differences in the jar headspace) and the weight of the dry soil samples (Robertson et al., 1999). Then these rates were also normalized by the percentage of C found in each sample to look at substrate quality. For samples with pH > 7, water contents were very low (Table 1); therefore, we assumed that a negligible amount of CO₂ was stored as DIC in the sample water and did not correct the calculation for the pH. However, we are aware that this might underestimate C mineralization.

The impact of glucose on CH₄ and CO₂ production was quantified as a glucose factor, calculated using the cumulative C production at 67 d as follows:

$$Gf = \frac{(P_{gt} - P_t)}{P_t},$$

where Gf is the glucose factor, P_{gt} is the total CH₄ production for samples with glucose, and P_t is the total CH₄ production.

2.5 Quantification of methanotrophic and methanogenic gene copy numbers

Methanogenic archaea were quantified with a quantitative polymerase chain reaction (qPCR) at the following different times during the incubations: when the samples were still frozen (1), after 60 d of incubation (2), and after glucose addition (67 d of incubation) (3). However, due to laboratory restrictions during the Covid-19 pandemic, it was not possible to analyze all the incubated vials. Only one replicate per sample for the first two runs was analyzed. For the last run, we selected the two samples with the highest CH₄ production rates after the glucose addition – the active layers of P16 and P17.

Since methanotrophic bacteria are good indicators of the oxidation level under in situ conditions, they were quantified with qPCR before starting the incubation.

Key genes encoding the enzyme methyl coenzyme M reductase (*mcrA*) (Thauer, 1998) and the enzyme particulate methane mono-oxygenase (*pmoA*) (Theisen and Murrell, 2005) were examined to identify methanogens and methanotrophs, respectively. DNA extractions were performed with a GeneMATRIX Soil DNA purification kit (Roboklon, Germany) according to the manufacturer's protocol. After DNA extraction, the DNA concentration was quantified by fluorescence with the Qubit dsDNA HS Assay Kit (Invitrogen, United States). Gene copy numbers were quantified using a SYBRGreen qPCR assay using the KAPA SYBR FAST qPCR Master Mix (Sigma Aldrich, Germany) on a CFX96 real-time thermal cycler (Bio-Rad Laboratories Inc., United States). All runs were performed in technical triplicates, and

each run was completed through melt-curve analysis in order to check for specificity of the assay (Liebner et al., 2015). Methanogenic archaea were targeted with the primer *mlas-F/mcrA-R* (Microosynth, Balgach, Switzerland) (Hales et al., 1996).

To amplify the methanogenic archaea *mcrA* gene, PCR samples were kept at 95 °C for 5 min to denature the DNA. The amplification process was performed with 40 denaturation cycles at 95 °C for 1 min, annealing at 60 °C for 45 s and elongating at 72 °C for 90 s. To ensure complete amplification, samples were kept at 80 °C for 10 min. In addition, to amplify the methanotrophic *pmoA* gene, using primer *pmoA189-F* and primer *pmoAmb661-R*, two PCR reaction conditions were used. The first PCR comprised initial denaturation at 95 °C for 5 min, 30 cycles with denaturation at 94 °C for 45 s, decreasing annealing temperature from 64 °C to 52 °C for 60 s, elongation at 72 °C for 90s, and final elongation at 80 °C for 90 s. The second PCR comprised an initial denaturation and polymerase activation at 95 °C for 5 min, 22 cycles of denaturation at 94 °C for 45 s, annealing at 56 °C for 60 s, elongation at 72 °C for 90 s, and a final extension at 72 °C for 10 min.

2.6 Statistical analyses

The gas production and microbial data did not show a normal distribution; consequently, it was not possible to test for differences by performing an ANOVA. The differences between cores and depths, and also the impact of temperature on gas production and microbes, were therefore tested using the Kruskal–Wallis test with the R function `kruskal.test`.

All statistics and results analyses were performed with R version 4.0.5 (R Core Team, 2021).

3 Results

3.1 Soil characteristics

All soil samples had a pH between 6.5–7.5, except P15-F. Most electrical conductivities were very low (< 200 μS cm⁻¹), except for two samples, namely P16-F and P17-A (Table 2). Water content was higher in permafrost (54.5%–60.8%) than in the active layer (23.7%–25.8%) for the two yedoma cores, P15 and P16. The water content was higher in the active layer than in the permafrost layer in the floodplain core P17 (36.2% vs. 17.2%) (Table 2).

Sediment TOC ranged from 0.2%–3.8%. Most TOC contents ranged from 2.7%–3.8%, but the TOC content in the permafrost layer of P17 was the lowest of the six samples (0.17%) (Table 2). All the samples had TOC below 6%, and therefore they were considered to be mineral soils (% C < 12%) (Table 2) (Soil Survey Staff, 2014). TN contents were very low for all the samples (< 0.3%) and were below the detection limit of the laser analyzer (below 0.1%) for P17-F. C : N ratios ranged between 12 and 20. The high-

est ratios were measured in P15; the lowest were in P16. The C:N ratio was higher in the permafrost layer of P15 than in the active layer.

The C stock ranged from 2.3 to 38.8 kg m⁻³. The active layers for all the samples were higher than 30 kg m⁻², while the highest C stock in the permafrost layer was 19 kg m⁻³ in the Yedoma core P16. The permafrost layer of the floodplain had the lowest C storage, more than 10 times lower than samples from the active layers (Table 2).

The grain size distribution was similar between P15 and P16. The active layer of P17 contained more clay and the least sand of the other samples, while permafrost in P17-F was the sandiest sample (Table S1 in the Supplement).

3.2 Potential gas production

3.2.1 CH₄ production over one year of incubation

At the end of the 363 d incubation, four of the six samples produced CH₄ at a 20 °C incubation temperature (Fig. 2, Fig. 3). The floodplain active layer (P17-A) was the sample with the highest cumulative CH₄ production over the incubation time (917.2 ± 150 µg CH₄-C g⁻¹ DW). After 6 months of incubation, the CH₄ production rate of P17-A decreased and then plateaued. The floodplain permafrost core (P17-F) produced 1 % of the amount of CH₄ produced by the active layer from the same core (0.5 ± 0.2 µg CH₄-C g⁻¹ DW) at 20 °C. The permafrost layers at 20 °C of both Yedoma cores (P15 and P16) produced similar amounts of CH₄ (20.5 ± 6.1 and 159 ± 104 µg CH₄-C g⁻¹ DW, respectively), while CH₄ production from the active layers of these cores was minimal (P16-A: 3.34 ± 0.25 µg CH₄-C g⁻¹ DW; P15-A: 0.51 ± 0.14 µg CH₄-C g⁻¹ DW). Cumulative CH₄ production at 4 °C was limited to one sample, the active layer of the floodplain core (Figs. 2, 3). Cumulative production of the other cores was less than 1 µg CH₄-C g DW⁻¹ after 1 year (Figs. 2a and b, 3).

The lag time before CH₄ production was observed ranged from 14 d to over 363 d. The active layer of the floodplain core (P17-A-20) was the first to produce CH₄ after 14 d of incubation at 20 °C. In the frozen layers of Yedoma, the cores required at least 6 months of incubation to start producing CH₄ at 20 °C (Fig. 2, Table 2), but in the active layer of the Yedoma cores, CH₄ production took substantially longer, specifically 273 d for P16-A-20, while P15-A did not produce appreciable CH₄ over 363 d in the experiment. At 4 °C, CH₄ production in P17-A started after 333 d but was not observed for the other samples (Fig. 2, Table 3).

3.2.2 CO₂ production over 1 year of incubation

Over the 363 d incubation, cumulative CO₂ production ranged from 90.3 to 701.4 µg CO₂-C g⁻¹ DW. The cumulative CO₂ production of P17-A at 20 °C was the highest among all the samples (701.4 ± 124 µg CO₂-C g⁻¹ DW),

while it was the lowest in permafrost layer of the same core at 20 °C (Figs. 2, 3). The CO₂ production of P15 and P16 was in the same range, between 142 ± 85 µg CO₂-C g⁻¹ DW (P16-A at 4 °C) and 348.3 ± 135 µg CO₂-C g⁻¹ DW (P15-F at 20 °C), except in the case of P16-F at 4 °C (chi squared = 2.80, *df* = 1, *p* = 0.20) (Figs. 2, 3). At 4 °C, the permafrost layers of the Yedoma core P16 and the floodplain core P17 had cumulative production below 60 µg CO₂-C g⁻¹ DW. The results per gram C showed a different pattern for the cumulative CO₂ production of the floodplain core P17. The permafrost layer at 4 °C reached 7.98 mg CO₂-C g⁻¹ C and had higher CO₂ production than the permafrost layer at 20 °C and the active layer at 4 °C (Fig. S3 in the Supplement).

A decrease of CO₂ production at the beginning of the incubation was observed for all the samples (Fig. 2, Fig. S4 in the Supplement). All the active-layer samples (except the active layer of the floodplain P17 at 4 °C), as well as the permafrost layers of the Yedoma cores P15 and P16 at 20 °C, reached the maximum production rates of CO₂ before or at the end of the first 2 months (Fig. 2). The maximum production rate of CO₂ for the active layer of the floodplain at 4 °C was attained after 300 d of incubation, and the other permafrost samples reached the maximum production rate between 2 and 5 months (Fig. 2). Maximum production rates ranged between 57.13 µg C-CO₂ g C⁻¹ d⁻¹ (P16-A) and 754 µg C-CO₂ g C⁻¹ d⁻¹ (P17-F) at 4 °C and between 120.54 and 510.65 µg C-CO₂ g C⁻¹ d⁻¹ for P16-F and P15-A, respectively, at 20 °C (Table S2 in the Supplement). After half a year of incubation, CO₂ production plateaued for all the samples, except for the active layer of the floodplain sample at 4 °C. For all the samples, we noticed an increase in CO₂ production after 60 d of incubation, e.g., after the microbial sampling, followed by a decrease in the CO₂ production. We did not consider those results to be descriptive of the maximum production rates.

After 1 year of incubation, neither the temperature nor the depth impacted the cumulative CO₂ production of the cores (chi squared = 3.86, *df* = 1, *p* = 0.09) (Figs. 2, 3). CO₂ production was higher at 20 °C only for the permafrost layer of the Yedoma core P16 and the active layer of the floodplain P17 (chi squared = 3.86, *df* = 1, *p* < 0.05) (Figs. 2, 3).

The P17-A-20 CO₂:CH₄ ratio decreased rapidly during the first 14 d. The CO₂:CH₄ ratio reached 1 after 40 d and remained stable until the end of incubation (Table 2). In addition, the CO₂:CH₄ ratio of P17-A at 4 °C and of P16-F at 20 °C was low after 363 d of incubation (respectively, 2.7 ± 2.7 and 2.5 ± 2.1). For all the samples, except for P15-F, CO₂:CH₄ ratios were significantly lower at 20 °C than at 4 °C (chi squared = 3.92, *df* = 1, *p* < 0.05) (Table 3).

3.2.3 Effect of glucose addition

Overall, no effect of glucose injection on CH₄ production was detected after 67 d of incubation (Table 3) (chi

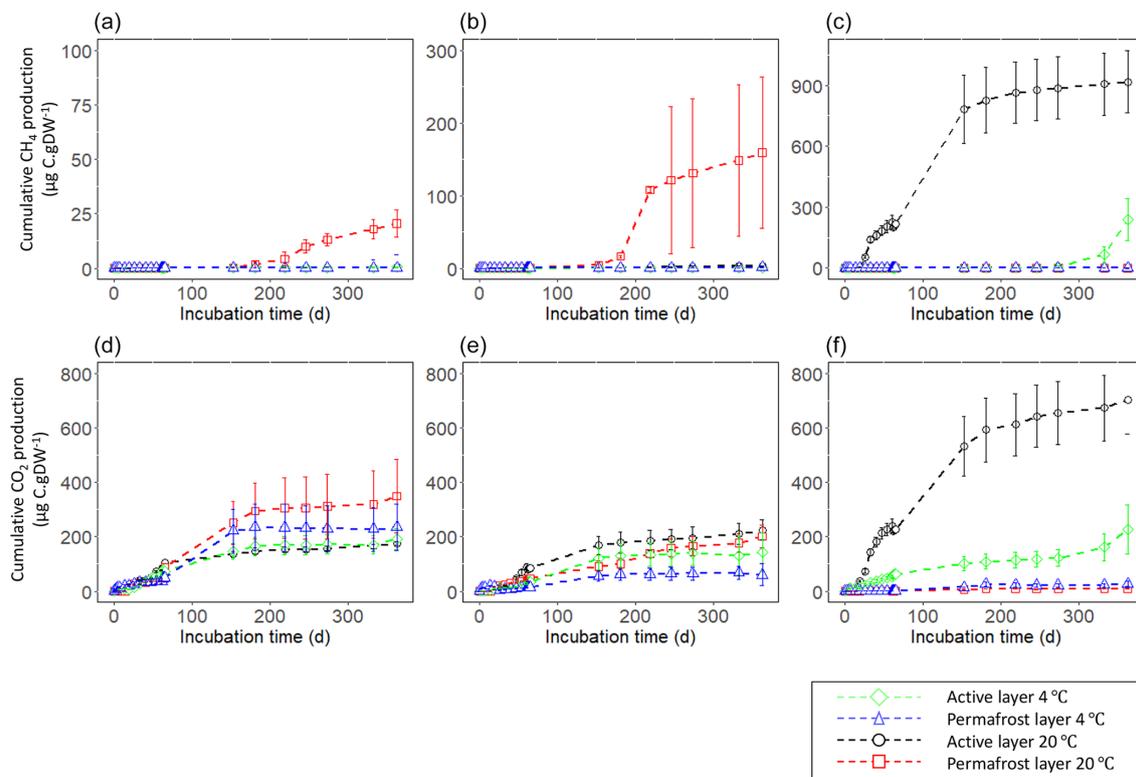


Figure 2. Cumulative gas production per gram dry weight (DW) at 4 and 20 °C for 363 d of incubation. CH_4 production of (a) P15, (b) P16, and (c) P17. CO_2 production of (d) P15, (e) P16, and (f) P17. Error bars show the standard deviation from the means \pm standard error ($n = 3$). Note the differing y-axis scales between cores for CH_4 .

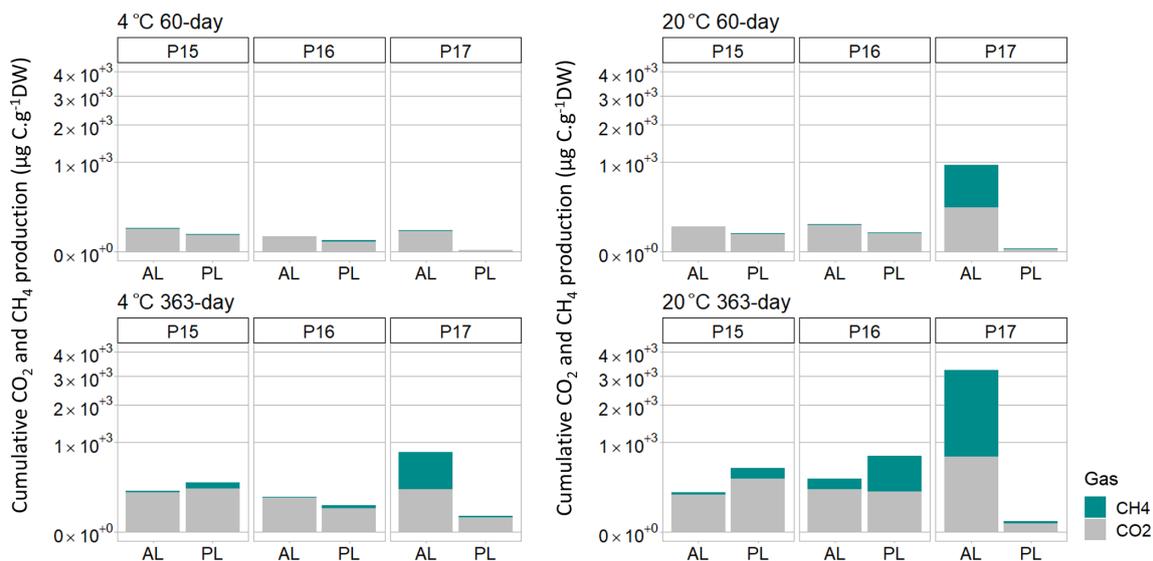


Figure 3. Cumulative production of CO_2 and CH_4 per gram dry weight after 60 d of incubation at 4 and 20 °C and after 363 d. AL stands for active layer, and PL stands for permafrost layer. Scale is expressed as square root for a better representation of the data.

Table 1. Soil description and vertical position of the sampling cores.

Samples	Depth (cm)	Layer	Horizon	Soil description
P15-A	41.5	Active	Mineral	Compacts silt, gray, with brown organic inclusions
P15-F	81.5	Permafrost	Mineral	Ataxilic ice structure, silt, gray
P16-A	38.5	Active	Mineral	Silt, brown, organic rich, slightly sandy
P16-F	102.5	Permafrost	Mineral	Silt, gray-brown, structureless to micro-lenticular
P17-A	31.5	Active	Mineral	Organic rich silt, slightly sandy
P17-F	78.5	Permafrost	Mineral	Sand, no visible ice

Table 2. Chemical and physical properties of the active and frozen layers of the three cores. The conductivity temperature reference was 25 °C. Numbers in brackets are standard deviations.

Samples	pH	Conductivity ($\mu\text{S cm}^{-1}$)	TOC (%)	C:N ratio	TN (%)	C (kg m^{-3})	Water content (wt %)
P15-A	6.75	164.5	3.54	18.13	0.1975	38.8	25.8
P15-F	6.06	150.2	2.70	20.59	0.133	9.4	60.8
P16-A	7.21	98.6	2.70	12.95	0.211	35.2	23.7
P16-F	7.06	479	3.81	12.67	0.3085	18.5	54.5
P17-A	7.22	635	3.48	18.46	0.1935	30.0	36.2
P17-F	7.44	86.4	0.17		<0.10	2.3	17.2

squared = 0.25, $df = 1$, $p = 0.5913$). A CH_4 production peak was detected 1 d after the second glucose addition for P15 and P16. The response factors were low (CH_4 production was between 1.2 and 1.7 times higher with glucose addition compared to without) and appeared only at 20 °C. No impact of the glucose addition was detected on CH_4 production for the samples at 4 °C after neither the first nor the second injection (Fig. S5 in the Supplement).

CO_2 production at 20 °C was, overall, increased by glucose (chi squared = 3.78, $df = 1$, $p < 0.05$). The maximum increase of CO_2 production was seen for the permafrost layer of P16 (4.2 times higher with the glucose addition). No difference in CO_2 production was detected for any of the samples at 4 °C after the glucose addition (Fig. S5 in the Supplement).

3.3 Gene copy numbers of methanogens and methanotrophs

For half of the samples, no methanogenic gene copy numbers were detected when the samples were thawed prior to beginning the incubation. From these samples, only the methanogenic gene copy number for core P17-A was above the detection limit (4.3×10^3). Therefore, it was not possible to compare the methanogenic gene copy numbers before the start of the incubation (Fig. 4).

After 60 d of incubation, the *mcrA* gene copy numbers ranged from 5.35×10^3 to 5.34×10^5 . The qPCR results showed significant differences between cores after 60 d of incubation (chi squared = 3.89, $df = 1$, $p < 0.05$) with the highest copy number per gram soil (Fig. 4c) in P17-A. No

methanogenic gene copy numbers were detected for the permafrost layer of P17 or for the active layer of P16.

P16-F and P17-A had 9 and 36 times higher *mcrA* copy numbers per gram soil, respectively, at 20 °C than at 4 °C (chi squared = 3.9, $df = 1$, $p < 0.05$) (Fig. 4). The temperature response for the active and permafrost layer of P15 was not identified because the results were below the detection limit at both 4 and 20 °C.

Similarly, no comparison was possible between active and permafrost layers for all the samples.

Gene copy numbers of methanotrophic bacteria based on the *pmoA* gene were either below the detection limit or not detected before the incubation; therefore, no interpretation of the oxic conditions under field condition was possible (Fig. 4).

Copy numbers after the addition of glucose did not differ from those without glucose (Fig. 4).

4 Discussion

4.1 CH_4 production in floodplain environment and Yedoma cores across landscape positions

4.1.1 Floodplain core

We mimicked potential CH_4 production during a growing season (60 d) in a floodplain environment of Kurungnakh Island in the Lena River Delta and extended the incubation time to 1 year to capture the CH_4 production behavior. Within the first 2 months, the results showed high rates and a quick onset of CH_4 production, as well as the pres-

Table 3. Summary of lag time of CH₄ offset, CO₂:CH₄ ratios, and glucose factors. Lag time is expressed in days. L.T. stands for samples where the lag time did not end after 1 year of incubation. CO₂:CH₄ ratios represent the means of total production after 363 d of incubation at 20 and 4 °C. Glucose factors were calculated based on cumulative C production 7 d after glucose addition. Positive values indicate the positive impact of glucose on GHG production, while negative values indicate less GHG production after glucose addition.

Samples	Layer	Lag time (days)			Mean CO ₂ :CH ₄		Glucose factor			
		20 °C	4 °C	4 °C	20 °C	4 °C	CH ₄ 4 °C	CH ₄ 20 °C	CO ₂ 4 °C	CO ₂ 20 °C
P15-A	Active	L.T.	L.T.		$522.6 \pm 1.7 \times 10^{+2}$	$409.1 \pm 2.2 \times 10^{+2}$	-0.10	-0.38	0.02	0.18
P15-F	Permafrost	153	L.T.		$1930.1 \pm 2.2 \times 10^{+3}$	$2236.8 \pm 3.9 \times 10^{+3}$	0.02	-0.31	-0.20	-0.44
P16-A	Active	274	L.T.		$1661.4 \pm 1.5 \times 10^{+2}$	$50.1 \pm 4.0 \times 10^{+1}$	-0.41	0.70	-0.02	1.22
P16-F	Permafrost	181	L.T.		$195.5 \pm 1.3 \times 10^{+2}$	2.5 ± 2.1	0.40	-0.93	0.11	3.23
P17-A	Active	14	333		2.7 ± 2.7	0.8 ± 0.1	-0.01	0.24	0.51	0.60
P17-F	Permafrost	L.T.	L.T.		$266.9 \pm 1.8 \times 10^{+2}$	$34.8 \pm 4.2 \times 10^{+1}$	1.18	0.27	-0.11	0.82

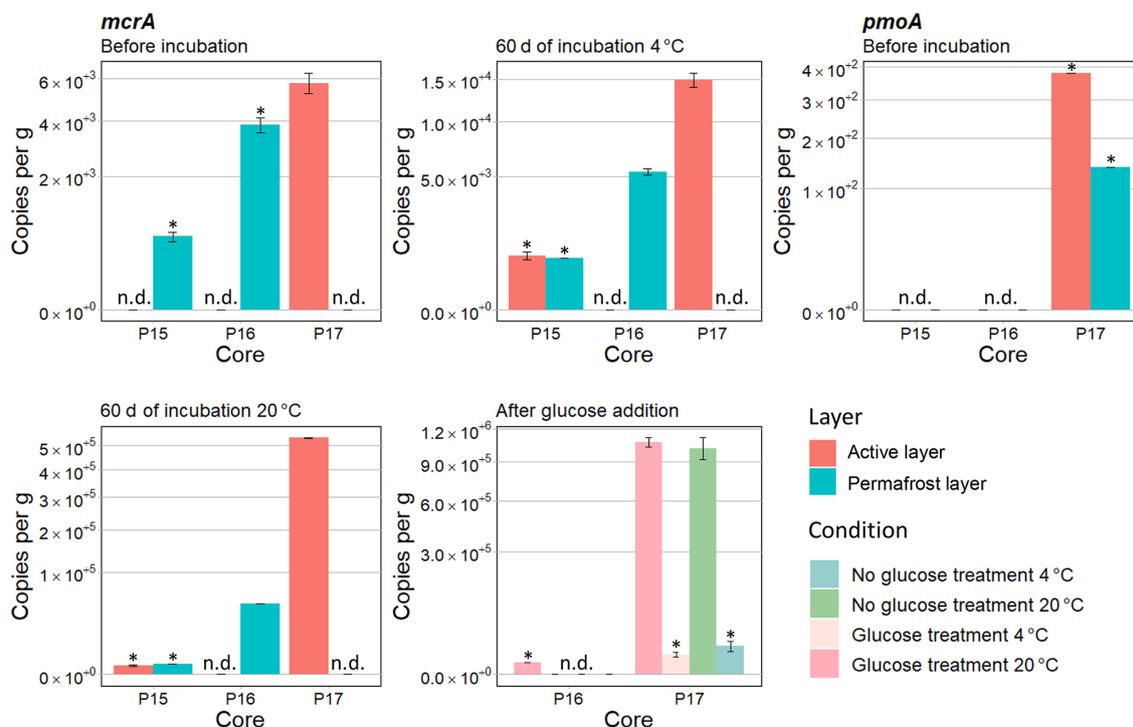


Figure 4. Means of gene copies per gram calculated with qPCR amplification at different times for different conditions – before the incubation, after 60 d of incubation, and at the end. Gene copy numbers of *mcrA* were calculated for P15, P16, and P17. *mcrA* results are shown for the active layers of P16 and P17 with or without glucose treatment after 67 d of incubation. Gene copy numbers of *pmoA* are shown for P15, P16, and P17 before the incubation. Samples below the detection limit are indicated by *, and samples where copies per gram were not detected are indicated by n.d. Scale is expressed as square root in order to have a better display.

ence of methanogen communities (Fig. 4), in the active layer of the floodplain core P17 at 20 °C only. Those findings, as well as the low CO₂:CH₄ ratio, indicated a quick establishment of optimum methanogenic conditions within the growing season time frame of 60 d (Symons and Buswell, 1993) (Figs. 2c, 3; Table 3). Herbst (2022) did a similar incubation study with samples from the active floodplains of Kurungnakh Island and nearby Samoylov Island (Fig. 1). CH₄ was produced from two of the three cores within the first 60 d of incubation (Table S3 in the Supplement). In both this study

and the Herbst study, CH₄ production was triggered quickly after the beginning of the incubation (from 10 to 40 d) within these floodplain samples. Thus, these Arctic floodplain environments may allow the fast establishment of methanogens and therefore rapid CH₄ production under anaerobic conditions.

However, not all floodplain soils showed fast establishment of methane communities and high rates of potential methane production. Unlike the active layer, the permafrost layers of floodplain did not produce appreciable CH₄ after 1

year of incubation at either 4 or 20 °C and were still considered to be in the lag phase. The absence of detectable *mcrA* copy numbers per gram soil after 60 d of incubation indicated the absence of methanogen communities in the permafrost samples (Fig. 4). Similarly, low rates of CO₂ production, low C content, and high sand content in this permafrost sample indicate non-suitable conditions for many types of soil microbes (Figs. 2, 4; Table 1) (Eskelinen et al., 2009).

As expected, our results showed significant differences between CH₄ production rates at 4 and 20 °C. At 4 °C, almost 300 d of incubation were needed to trigger CH₄ production in the active layer of the floodplain (versus 14 d at 20 °C), with a total cumulative CH₄ production 4 times lower than at 20 °C (Fig. 3). Other studies showed similar patterns; e.g., CH₄ production increases with temperature and has shorter lag times (Ganzert et al., 2007; Treat et al., 2015). This is explained by a strong temperature sensitivity of methanogen communities (Westermann, 1993; Li et al., 2015). At the end of the growing-season simulation, our results showed *mcrA* copy numbers 36 times lower at 4 °C than at 20 °C (Fig. 4), again indicating that the methanogen community required both time and high temperatures to establish.

4.1.2 Yedoma cores

This study highlights a different CH₄ production behavior between the floodplain and the Yedoma cores. The permafrost layers from the Yedoma cores only started producing CH₄ after 6 months of incubation at 20 °C, whereas the floodplain core produced CH₄ earlier (Fig. 2). The CO₂:CH₄ ratios remained high after 1 year of incubation (Table 3), meaning that the methanogenic conditions were not yet optimal (Symons and Buswell, 1993). The low *mcrA* copies after the 60 d of incubation compared to the active layer of the floodplain, as well as the long lag times, showed that the methanogen communities took more time to become active in the permafrost Yedoma cores (Figs. 2, 4).

Our results indicated higher CH₄ production rates in the permafrost layer than in the active layer, while others generally show the opposite (Yavitt et al., 2006; Treat et al., 2015, p.201). However, most of the studies on CH₄ production from Yedoma cores showed high variability in the cumulative CH₄ production. As explained above, lag times measured from former studies, as well as CH₄ production rates, differed (Lee et al., 2012; Knoblauch et al., 2013; Walz et al., 2018; Jongejans et al., 2021). It is therefore difficult to estimate the potential production of CH₄ after thawing from Yedoma soils due to this high variability. Methanogens are highly constrained microbial communities, and the community size varied strongly between the sites in this study (Fig. 4), which partly explains the discrepancies among the studies on Yedoma soils due to the ecological and phylogenetic narrowness of the methanogen communities (Ernakovich et al., 2022).

The active layers at 4 and 20 °C and the permafrost layers at 4 °C were still in the lag phase without appreciable CH₄ production after 1 year of incubation (Figs. 2, 3; Table 3), which agrees with the absence of detected methanogen community (Fig. 4). Several multiannual studies also observed long and heterogeneous lag times at 4 °C for Yedoma soils (from 53 ± 23 up to 2500 d; Knoblauch et al., 2018; Walz et al., 2018). Knoblauch et al. (2018) explained the long lag time as being the result of a lack of methanogens or a lack of active methanogenic communities in soil samples. We added glucose to test whether the absence of CH₄ production was due to a lack of labile C or a lack of established methanogenic communities. If the methanogen community was small but established, we would expect to have community growth after the glucose addition. Since glucose had no effect on either CH₄ production rates or P15 and P16 methanogen community growth, we concluded that the absence of CH₄ production for those samples was because the methanogens were not active (or not active enough to detect). It has been shown that the establishment of a microbial community after thaw is correlated to the community characteristics and the thaw disturbance (Deng et al., 2015; Ernakovich et al., 2022). For ecologically and phylogenetically narrow microbial communities, like methanogens, random environmental processes like microtopography (stochastic processes) strongly influence the abundances and activation of the microbial communities. After an abrupt thaw, like we simulated in our incubation study, the role played by stochastic processes on constrained microbial communities is even stronger (Deng et al., 2015; Ernakovich et al., 2022). Therefore, although the incubation was carried out under anaerobic conditions, the quantity and the establishment of an active methanogen community in the samples after thaw was strongly controlled by stochastic processes.

4.1.3 Controls on CH₄ production across landscape positions

CH₄ production over the incubation time was not correlated with the TOC and TN % (Fig. 3, Table 2). The landscape position rather than soil characteristics played a key role in the establishment of microbial activity and, consequently, explained much of the variation in GHG production among the samples. Periodic water saturation in core P17 was indicated by oxidation marks at several depths in the field. These redox features indicate periodically anoxic conditions that likely favored the establishment or persistence of methanogen communities and reduced the lag times prior to CH₄ production under anaerobic conditions (Chasar et al., 2000; Paul et al., 2006; Jaatinen et al., 2007; Keller and Bridgham, 2007) (Fig. 2, Table 3). On the other hand, well-drained conditions were found in the field for the active layers of both the upland and the slope cores (P15 and P16), which did not produce appreciable CH₄ after 1 year of incubation (Figs. 2, 3). The aerobic conditions due to the dry environment likely in-

hibited methanogenesis (Megonigal and Schlesinger, 2002). Unlike the active layers, the permafrost layers of Yedoma showed low but existing *mcrA* results from the Yedoma permafrost layers at 20 °C (Fig. 4) and started producing CH₄ after 6 months. The methanogen community was likely established prior to or during the deposit of the Yedoma sediments, and the microbial community survived despite being freeze-locked (Holm et al., 2020). In addition, we quantified methanotroph communities to include more information about the potential for methane oxidation under field conditions, but the results showed amounts below the detection limit before the incubation (Fig. 4). These results support our hypothesis concerning the impact of landscape position on CH₄ production, namely that aerobic conditions in the landscape coincide with poor establishment of methanogenesis even when incubation conditions become favorable for methanogens.

4.2 Controls on potential CO₂ production

The rates of CO₂ production per gram C were on the same order of magnitude as other Yedoma incubation studies from Kurungnakh Island (Knoblauch et al., 2013, 2018) and the nearby Lena River Delta (Walz et al., 2018). These similar results suggest that C in these Yedoma soils is easily available due to the organic-rich characteristics (Strauss et al., 2013a). On the other hand, the adjacent samples from the permafrost layers of the floodplain showed CO₂ production per gram similar to the Yedoma cores, while it had the lowest CO₂ cumulative production per gram dry weight of soil. Although floodplain environments in the Lena Delta are considered to be low-C-content pools (Siewert et al., 2016), our results showed that the C in this soil was highly labile and comparable to the lability of Yedoma soils.

The CO₂ production followed trends in total C and N contents. The samples with similar C and N contents produced comparable ranges of CO₂, whereas the sample (P17-F) with the lowest TOC and N content showed low CO₂ production (per gram DW) during the incubation (Fig. 2, Table 2, Table S2 in the Supplement). As shown by Schädel et al. (2014), the C : N ratio was positively correlated to the cumulative CO₂ production. However, the correlation was stronger at 4 than at 20 °C (Fig. S7 in the Supplement). Therefore, consistent with other studies (Schädel et al., 2014; Knoblauch et al., 2018), the quality (N), quantity (C), and the bioavailability (C : N) of the OM is a key factor for the mineralization into CO₂ production.

Our CO₂ and CH₄ production results combined with microbial analysis indicated that CO₂ production pathways might change according to the landscape position. The CO₂ : CH₄ production ratio of 1 : 1, as well as the presence of a high number of methanogenic archaea, indicated that the CO₂ production in the active-layer floodplain could have originated from methanogenesis (Figs. 3, 4) (Symons and Buswell, 1993; Knoblauch et al., 2018; Holm et al.,

2020). In drier environments, like the P15 and P16 cores, the high CO₂ : CH₄ production rates resulted from other, under-terminated anaerobic decomposition pathways. Anaerobic respiration is a common function, and diverse microbial communities are able to decompose the OM to CO₂; therefore, the anaerobic microbial community is not a limiting factor for C mineralization to CO₂ (Elderfield and Schlesinger, 1998). Based on the positive correlation between C : N and the cumulative CO₂ (Fig. S7 in the Supplement), as well as the broad microbial community that is able to produce CO₂, our CO₂ production is rather controlled by the quality (N) and the quantity (TOC) of the OM rather than by the microbial communities (Knoblauch et al., 2018; Holm et al., 2020).

4.3 Implication for C feedback on Kurungnakh Island during the growing season

With climate change, Arctic environments will be subject to changes in moisture conditions, vegetation shifts, increased active-layer depth, and abrupt permafrost thaw (Serreze et al., 2000; Hinzman et al., 2005; Myers-Smith et al., 2011; Turetsky et al., 2019). Our permafrost thaw simulation under wet summer conditions for Kurungnakh Island soil showed that the CO₂ production for the Yedoma cores was similar in magnitude to other studies including Yedoma soil. In the incubation experiment, all the Yedoma samples reached the maximum production rates within the first 2 months of incubation (Fig. 2). Schädel et al. (2014) attributed the decrease in CO₂ production rates after some time in incubation to a rapid C turnover that relied mainly on the decomposition of the labile C pool (Schädel et al., 2014; Walz et al., 2018; Schädel et al., 2020). However, several studies pointed out the small size of the labile C pool of Yedoma deposits (Knoblauch et al., 2013; Strauss et al., 2013b). Here, the Yedoma soils from Kurungnakh Island showed labile-C-pool depletion after 6 months of incubation (e.g., the CO₂ production rates decreased, and the cumulative CO₂ production plateaued after 6 months of incubation; Fig. 2). Therefore, under wet summer conditions, it is likely that there will be rapid C turnover and CO₂ production during the growing season.

The active layer of the floodplain at 20 °C produced up to 300 µg CH₄-C g⁻¹ DW at the end of the simulated growing season. The low-lying position of the floodplain allows regular flooding from the river. The hydrological conditions of this area provide favorable redox conditions, i.e., anaerobic conditions, for the establishment of active methanogen communities if the temperature is high enough (Figs. 3, 4). Therefore, we expect CH₄ production from the floodplain site relatively soon after the beginning of the growing season. Long-term in situ measurements in the Lena Delta have shown the highest CH₄ emission rates for moist to dry dwarf-shrub-dominated tundra, located mainly in lower floodplain environments (5048.5 mg m⁻² yr⁻¹; Schneider et al., 2009). In this area of the Lena Delta, CH₄ emissions have been measured from June to September, with the highest emis-

sion rates in July (Rößger et al., 2022). Our results showed a high CH₄ production potential from the floodplain; however, floodplain environments are periodically flooded, meaning there might be periods where the floodplain would be too dry to allow CH₄ production (Huissteden et al., 2005; Obligov et al., 2020). A long-term study like that by Rößger et al. (2022) in floodplain environments would help to quantify CH₄ emission from floodplains, how often it occurs during the growing season, and how the CH₄ flux would respond to changes in soil moisture over time.

While CH₄ production did occur in the year-long anaerobic incubation of Yedoma samples, other factors might result in these tundra regions still being a net CH₄ sink barring abrupt permafrost thaw (Schneider et al., 2009; Juncher Jørgensen et al., 2015). In these dry Yedoma sites, the net CH₄ flux is the balance between CH₄ production in anoxic soil layers and CH₄ oxidation in overlying aerobic layers. Here, we showed that CH₄ production was possible from these soils given high-enough temperatures (Fig. 3), but this required a long lag time for the establishment of the methanogen communities (Fig. 4). Therefore, it is unlikely that the active methanogen communities have enough time (>60 d; Fig. 4) or warm-enough temperatures (Fig. 3) to establish during the growing season in the upland and slope areas (P15, P16) on Kurungnakh Island even with deeper active layers, soil moisture increases, and highly bioavailable C in Yedoma sediments (Anthony et al., 2014; Mann et al., 2014; Spencer et al., 2015). These conditions may constrain the potential CH₄ emissions from some Yedoma soils even with warmer climates, wetter soils, and permafrost thaw but will require additional field-based observations to account for plant transport and methane oxidation processes occurring in situ.

5 Conclusions

In this study, we provide new information regarding the importance of the landscape position for CH₄ production during the growing season on Kurungnakh Island in the Lena River Delta. High CH₄ production was measured in waterlogged (floodplain) areas within the 60 d simulation of the growing season at 20 °C thanks to a fast establishment of the methanogen community (14 d). In contrast, the well-drained Yedoma active-layer samples were still in the lag phase at the end of the growing-season simulation and did not produce appreciable CH₄ emissions, and C turnover resulted in CO₂ production. CH₄ was produced by the Yedoma permafrost layers after a lag phase of 6 months at 20 °C. Although the permafrost layer of the floodplain had low TOC, we identified similar C lability for the three cores as for other studies with samples from Siberia and therefore high potential C gas production throughout this region but mainly as CO₂. As a result, the data presented in this case study contribute to quantifying and understanding C turnover in permafrost ar-

reas. Questions remain regarding how to upscale results from laboratory incubation to in situ conditions, and our results highlighted the need to better understand changes in redox conditions across the landscape position to improve upscaling.

Data availability. The data sets used in this paper are available at <https://doi.org/10.1594/PANGAEA.945685> (Laurent et al., 2023).

Supplement. The supplement related to this article is available online at: <https://doi.org/10.5194/bg-20-2049-2023-supplement>.

Author contributions. ML, CCT and SL designed the study. ML conducted all the experiments (soil analyses, incubations, and microbe quantification). MF and AR collected the soil samples and field notes during the expedition in 2018 and created the map. SL furnished the laboratory materials to perform microbe analyses and gas measurements. TH provided data from her incubation experiments. ML wrote the paper with contributions from all the co-authors.

Competing interests. The contact author has declared that none of the authors has any competing interests.

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