

1 Diploptene $\delta^{13}\text{C}$ values from contemporary thermokarst lake
2 sediments show complex spatial variation

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25 **Abstract**

26 Cryospheric changes in northern high latitudes are linked to significant greenhouse gas flux
27 to the atmosphere, for example, methane that originates from organic matter decomposition
28 in thermokarst lakes. The set of pathways that link methane production in sediments, via
29 oxidation in the lake system, to the flux of residual methane to the atmosphere is complex
30 and exhibits temporal and spatial variation. The isotopic signal of bacterial biomarkers
31 (hopanoids, e.g. diploptene) in sediments has been used to identify contemporary ocean-floor
32 methane seeps and, in the geological record, periods of enhanced methane production (e.g.
33 the PETM). The biomarker approach could potentially be used to assess temporal changes in
34 lake emissions through the Holocene via the sedimentary biomarker record. However, there
35 are no data on the consistency of the signal of isotopic depletion in relation to source or on
36 the amount of noise (unexplained variation) in biomarker values from modern lake
37 sediments. We assessed methane oxidation as represented by the isotopic signal of methane
38 oxidising bacteria (MOB) in multiple surface sediment samples in three distinct areas known
39 to emit varying levels of methane in two shallow Alaskan thermokarst lakes. Diploptene was
40 present and had $\delta^{13}\text{C}$ values lower than -38‰ in all sediments analysed, suggesting methane
41 oxidation was widespread. However, there was considerable variation in $\delta^{13}\text{C}$ values within
42 each area. The most ^{13}C -depleted diploptene was found in an area of high methane ebullition
43 in Ace Lake (diploptene $\delta^{13}\text{C}$ values between -68.2 and -50.1‰). In contrast, significantly
44 less depleted diploptene $\delta^{13}\text{C}$ values (between -42.9 and -38.8‰) were found in an area of
45 methane ebullition in Smith Lake. $\delta^{13}\text{C}$ values of diploptene between -56.8 and -46.9‰ were
46 found in centre of Smith Lake, where ebullition rates are low but diffusive methane efflux
47 occurs. The small-scale heterogeneity of the samples may reflect patchy distribution of
48 substrate and/or MOB within the sediments. The two ebullition areas differ in age and type of
49 organic carbon substrate, which may affect methane production, transport and subsequent
50 oxidation. Given the high amount of variation in surface samples, a more extensive
51 calibration of modern sediment properties, within and among lakes, is required before down-
52 core records of hopanoid isotopic signatures are developed.

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56 **1 Introduction**

57 Arctic lakes are sources of methane within the global carbon cycle (Bastviken 2004). More
58 specifically, thermokarst and thermokarst-affected lakes (those formed and/or influenced by
59 thaw and collapse of ice-rich ground) are recognized as important but variable past and
60 present sources of methane flux to the atmosphere (Shirokova et al., 2012; Walter et al.,
61 2006, 2008; Wik et al., 2013). Predictions of future variation in methane emission rates are
62 largely based on measurements recorded over the last 15 years (e.g. Brosius et al., 2012;
63 Walter Anthony et al., 2014). Long-term (i.e. Holocene) variations in lake-derived methane
64 flux to the atmosphere and changes in emissions during discrete climatic events in the past
65 are generally not well understood (but see Walter Anthony et al., 2014; Walter et al., 2007b).
66 Understanding methane activity in lakes over recent (e.g., decadal/centennial) and longer
67 (millennial) time periods and its relationship with forcing factors (e.g., temperature) could
68 provide useful constraints for the projection of future fluxes with arctic warming.

69 A significant fraction of methane produced in lake sediments may be oxidized and recycled
70 within the lake by methane oxidising bacteria (MOB), a process that offsets methane
71 emissions (Bastviken et al., 2002; Liebner and Wagner, 2007; Reeburgh, 2007; Trotsenko
72 and Khmelenina, 2005). Methane oxidation (MO) is a critical process for tracking past
73 methane production, as the bacteria that carry it out leave a distinctive trace (biomarkers) in
74 the sediments that were their habitat (see below). However, before this proxy can be
75 developed we need to better understand the link between methane production, MO within the
76 lake system and its geochemical representation, and observed fluxes to the atmosphere. Our
77 study contributes towards this goal by assessing the $\delta^{13}\text{C}$ values of bacterial biomarkers
78 obtained from the surficial sediments of two Alaskan lakes to ascertain if i) MO was
79 occurring, and ii) the degree of MO observed in areas characterized by different modes of
80 methane production and transport to the atmosphere.

81 1.1. Methane processing in thermokarst lakes

82 Methane production in thermokarst lakes takes two forms: production occurring in anoxic
83 surface sediments, as is common in most freshwater lakes and reservoirs, and that occurring
84 in deeper sediments, especially along the boundary of the "thaw bulb", which is specific to
85 thermokarst lakes (Figure 1). Commonly, methane production occurs via mineralisation of
86 older organic carbon from sources not found in other types of lake: i) where thermokarst-
87 induced erosion leads to large-scale slumping of banks into the littoral zone; material is

88 typically of Holocene age, but may be older (Figure 1), and ii) the microbial processing of
89 older, labile carbon in the talik, i.e., thawed sediment of the original landscape underlying the
90 lake.

91 Once produced, methane can be transported to the atmosphere through a number of
92 pathways: ebullition (bubbling), turbulent diffusion and plant mediated transport (Bastviken,
93 2004). Walter Anthony et al. (2010) postulate that most thermokarst-specific methane
94 production is transported to the atmosphere via seep ebullition. Thermokarst-specific
95 methane ebullition seeps have been studied using GPS mapping and submerged bubble traps
96 and appear to be persistent, spatially explicit fluxes at the water-air interface (Sepulveda-
97 Jauregui et al., 2015; Walter Anthony and Anthony, 2013; Walter et al., 2006, 2008). Spatial
98 stability is attributed to the development of conduits or 'bubble tubes' (Greinert et al., 2010;
99 Scandella et al., 2011), which form point sources at the sediment-water interface. Typically,
100 such seeps are densest near actively eroding lake margins, which we call the "thermokarst
101 zone". Here, methanogenesis is high due to the thermokarst-specific sources of methane
102 production: thawing of fresh talik and bank collapse (Figure 1; Kessler et al., 2012). Less
103 work has focused on methane production in the surficial sediments of thermokarst lakes, its
104 dissolution and diffusion from the sediments to the water column, and the resultant diffusive
105 emission rates. The diffusive flux component can, however, be relatively high, particularly in
106 older, more stable thermokarst lakes that have accumulated Holocene-aged organic carbon in
107 near-surface sediments (Martinez-Cruz et al., 2015; Walter Anthony et al., 2010).

108 1.2 Determining past methane activity using biomarker proxies

109 Past methane activity can be addressed qualitatively by using indirect proxies, for example,
110 features related to the cycle of methane through the lacustrine food web. Biogenic methane
111 has highly depleted $\delta^{13}\text{C}$ values (usually -80 to -50‰, Whiticar, 1999), depending on the
112 methane production pathway and substrate availability, and this signal can be extracted from
113 various organisms that utilise methane as a food source. Many studies have used the $\delta^{13}\text{C}$
114 values of compounds such as hopanoids from bacteria as indicators of past MO, relating
115 depleted $\delta^{13}\text{C}$ values with increased MO and methane supply (e.g. Boetius et al., 2000;
116 Collister and Wavrek, 1996; Hinrichs et al., 2003; Pancost et al., 2007).

117 The compound diploptene (17 β (H), 21 β (H)-hop-22 (29)-ene), is a hopanoid hydrocarbon
118 derived from a range of bacterial sources including heterotrophs and methanotrophs.

119 Therefore, the $\delta^{13}\text{C}$ values of diploptene represent a mixing relationship, with ^{13}C -depleted

120 MOB at one end and ^{13}C -enriched heterotrophic bacteria (which utilise organic carbon from
121 vegetation) at the other end (Pancost and Sinninghe Damste, 2003 and references therein).

122 In marine sediments, and especially in microbial mats associated with methane seeps,
123 diploptene has been identified as a methanotrophic biomarker via negative $\delta^{13}\text{C}$ values
124 (Elvert et al., 2001b; Pancost et al., 2000a, 2000b). Similarly, it has been argued to have a
125 partial methanotroph source on the basis of low $\delta^{13}\text{C}$ values in Holocene peat (Elvert et al.,
126 2001b; Pancost et al., 2000a, 2000b; van Winden et al., 2010; Zheng et al., 2014). Diploptene
127 and the related diplopterol have been used to infer past patterns of MO from marine sediment
128 records (Jahnke et al., 1999; Pancost et al., 2000a) and lake sediments (Spooner et al., 1994).

129 1.3 Detecting past changes in methane oxidation in thermokarst lakes

130 If MOB are present in the sediments of thermokarst lakes, we would expect to see depleted
131 $\delta^{13}\text{C}$ values of diploptene. To oxidise methane effectively, MOB require access to dissolved
132 methane, and thus it is assumed that MOB-related isotopic depletion indicates oxidation of
133 dissolved methane. However, the strong ebullition observed in some thermokarst lakes
134 complicates the issue, as the relationship between methane that diffuses from sediments (and
135 is either recycled in the lake via MO or released to the atmosphere) and methane that is
136 released to the atmosphere via ebullition remains unclear.

137 Numerous studies suggest that the methane transport pathways diffusion and ebullition co-
138 vary. In deep marine environments a correlation between methane supply in sediments,
139 transported via either diffusive processes or advectively at cold seeps, and MO as indicated
140 by $\delta^{13}\text{C}$ values of specific bacteria and of compounds (Elvert et al., 2001a; Pancost et al.,
141 2001, 2000b). In a shallow (9-m) bight the formation of bubble tubes was linked with
142 increased methane diffusing from the sediments, the proposed explanation being that bubble
143 tubes create an increased surface area that enhances methane diffusion, even though the
144 methane transported via ebullition is taken directly to the atmosphere and is not subject to
145 oxidation (Martens and Klump, 1980).

146 While little work has focused on MO in thermokarst lakes (but see Martinez-Cruz et al.,
147 2015), He et al. (2012) provide evidence for a possible correlation between a methane
148 ebullition seep (in this case, coal-bed sourced) and MO in a thermokarst lake, L. Qalluuraq,
149 Alaska. Using DNA-based stable-isotope probing they calculated the highest MO potentials
150 near the seep, and these were associated with the presence of MOB in the sediments. This
151 suggests a potential link between methane ebullition and increased availability of methane

152 that can be utilised by organisms in the lake sediments and water column. However, He et al.
153 (2012) also observed high variability in MO potentials and methanotroph communities with
154 changing substrates, temperature and sediment depth, indicating the need for further
155 investigation of MO in thermokarst lakes. In contrast, based on $\delta^{13}\text{C}$ and δD stable isotope
156 values and radiocarbon ages of methane in bubbles, Walter et al. (2008) and Walter Anthony
157 et al. (2014) suggest that methane originating in deep thaw-bulb sediments and emitted by
158 ebullition by-passes aerobic MO and that the majority (90%) of deep-sourced methane is
159 transported through ebullition seeps as opposed to via diffusion. Thus there is currently
160 limited and contrasting evidence for a link or otherwise between levels of methane ebullition
161 and methane diffusion in thermokarst lakes.

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163 **2 Regional context & Study sites**

164 Yedoma-like deposits that are similar to those described in Siberia (Schirrmier et al 2011)
165 can be found in Interior Alaska. In Alaska these sediments can have a relatively high organic
166 content (i.e., retransported silt; Péwé, 1975). They are also rich in excess ice (up to 80% in
167 Siberia). Thermokarst lakes that develop in landscapes dominated by such deposits have been
168 categorized as yedoma lakes in previous studies (Walter et al., 2008; Brosius et al., 2012;
169 Sepulveda-Jauregui et al., 2015). Two lakes were sampled in April 2011 and July 2012
170 (Figure 2). Ace Lake represents a yedoma lake (Sepulveda-Jauregui et al., 2015), where the
171 sediments surrounding the lake and eroding into it along its NE margin are predominantly
172 yedoma. Smith Lake is classified as a non-yedoma lake in which Holocene-aged deposits are
173 likely the main source of organic matter fuelling methane production.

174 Smith L. (64°51'55.92"N, 147°52'0.70"W; figure 2) is a shallow (≤ 4 m), productive lake
175 located near the University of Alaska, Fairbanks. It has a gentle bathymetric profile with
176 average water depths between 1-3m. The lake is not subject to a strong fetch or high energy
177 inflow or outflow. It is eutrophic, and observations during ice-free periods suggest high
178 primary productivity, with blue/green algal blooms predominant throughout the summer
179 months. The lake likely originated by thermokarst processes (Alexander and Barsdate, 1971);
180 comparisons of lake shorelines between the 1950s and today suggest that segments of the
181 southern and western margins have been actively thawing and eroding during recent decades,
182 and tilting trees currently lining the margin of a bay on the southeast shore are further

183 evidence of localized thermokarst. Smith Lake's shallow profile reduces the potential of
184 production or storage of methane due to stratification in the ice-free season.

185 Ace L. (64°51'45.49N, 147°56'05.69W) is part of the Ace-Deuce system (Alexander and
186 Barsdate, 1974) situated within an area covered by the Pleistocene Gold Hill Loess and
187 Goldstream Formation (Péwé, 1975). Ace L. is thermokarst in origin and formed through the
188 thawing of ice bodies in the loess. The Ace-Deuce Lake system has high nutrient levels and
189 can be described as a eutrophic lake with a strong seasonal nutrient cycle (Alexander and
190 Barsdate, 1974). As with Smith L., blue/green algal blooms are common throughout the
191 summer months.

192

193 **3 Methods**

194 **3.1 Establishing the thermokarst zones**

195 Walter Anthony and Anthony (2013) defined the 'thermokarst' zone for a number of lakes,
196 and we continue to use this definition here, i.e., the region of active thermokarst margin
197 expansion observed using historical aerial photographs obtained during the past 60 years. In
198 most lakes, the density of ebullition seeps is higher in thermokarst zones compared to
199 elsewhere (Walter Anthony and Anthony, 2013). In Ace and Smith Lakes, ebullition
200 emission rates have been quantitatively monitored through a combination of early-winter ice-
201 bubble surveys and bubble-trap flux measurements in previous studies (see Sepulveda-
202 Jauregui et al., 2015 for methods). We obtained surface sediment cores, from both the ice and
203 open water well within the zone boundaries and as close to observed ebullition seep locations
204 as possible (figure 2). The deepest part of Ace L. (the central area) was not sampled. The
205 development of a thermocline and anoxic bottom waters in deeper sections of Ace L. would
206 likely have an effect on both the rate of production and oxidation of methane that occurs in
207 the surface sediments. Eliminating such factors reduces the number of variables which might
208 explain the $\delta^{13}\text{C}$ values derived in this study.

209 **3.2 Methane monitoring**

210 Ebullition gas samples were collected from seep locations (October 2009 at Smith L and
211 April, 2011 at Ace L.) in the thermokarst zone (n1 and n5 for Smith L, and Ace L.
212 respectively) in the manner described in Walter Anthony et al. (2012) for determination of

213 bubble methane concentration and stable isotope analyses. Gases were collected from
214 submerged bubble traps into 60-ml glass serum vials following Walter et al. (2008), sealed
215 with butyl rubber stoppers, and stored under refrigeration in the dark until analysis in the
216 laboratory. We measured methane concentration using a Shimadzu 2014 equipped with an
217 FID at the Water and Environmental Research Centre at University of Alaska Fairbanks
218 (UAF). We determined $\delta^{13}\text{C}_{\text{CH}_4}$, using a Finnegan Mat Delta V at Florida State University.
219 Subsamples of gas were combusted to CO_2 , purified, and catalytically reduced to graphite
220 (Stuiver and Polach, 1977), and the $^{13}\text{C}/^{12}\text{C}$ isotopic ratios were measured by accelerator
221 mass spectrometry at the Woods Hole Oceanographic Institution's National Ocean Sciences
222 AMS Facility. Stable isotope compositions are expressed in δ (‰) = $103 \left(\frac{R_{\text{sample}}}{R_{\text{standard}}} - 1 \right)$,
223 where R is $^{13}\text{C}/^{12}\text{C}$ standard refers to the Vienna Pee Dee Belemnite (VPDB). The analytical
224 error of the stable isotopic analysis was ± 0.1 ‰ $\delta^{13}\text{C}$. We express radiocarbon data as
225 percent modern carbon pmC (%) = $\left(\frac{(^{14}\text{C}/^{12}\text{C})_{\text{sample}}}{(^{14}\text{C}/^{12}\text{C})_{\text{standard}}} \right) \times 100$, which is the
226 percentage of $^{14}\text{C}/^{12}\text{C}$ ratio normalized to $\delta^{13}\text{C} = -25$ ‰ and decay corrected relative to that of
227 an oxalic standard in 1950 (Stuiver and Polach, 1977).

228 **3.3 Biomarker analysis**

229 Surface sediment samples were retrieved using a gravity corer and the 0-5cm sequence was
230 extruded at 1-cm resolution and retained for analysis; the 1-2 cm slice was subsampled for
231 biomarker analysis and not the top as the sediment-water interface was often difficult to
232 sample cleanly due to unconsolidated sediments. The 1-2 cm slice integrates a number of
233 years of sediment accumulation (>10years) which reflects samples from a
234 palaeoenvironment. Two sequential sediment extractions were performed to obtain the total
235 lipid extract. The first step was a modified Bligh and Dyer extraction (Bligh and Dyer, 1959).
236 Briefly, buffered water was prepared adjusting a solution of 0.05M KH_2PO_4 in water to pH
237 7.2 through the addition of NaOH pellets. Subsequently, a monophasic solvent mixture was
238 made up with buffered water, CHCl_3 and MeOH (4:5:10 v/v). Samples were sonicated in
239 Bligh-Dyer solvent mixture for 15 minutes and then centrifuged at 3000 rpm for 5 minutes.
240 Supernatant was collected in a round bottom flask. This step was repeated twice and all
241 supernatants were combined and dried to obtain the total lipid extraction (TLE) labelled
242 TLE1. Post-extraction sediment residues were air-dried. The Bligh and Dyer post-extraction
243 residues were sonicated in DCM for 15 minutes and then centrifuged at 3000 rpm for 5
244 minutes. This step was repeated first with DCM:MeOH (1:1, v/v) and then with MeOH.

245 Supernatants were combined after every step of sonication-centrifugation to obtain TLE2.
246 Both TLE1 and TLE2 were then combined to yield the final TLE.
247 The TLE was split into three fractions of increasing polarity using silica flash column
248 chromatography (Oba et al., 2006; Pitcher et al., 2009). Silica gel columns (0.5 g, 60 Å
249 particle size) were prepared and conditioned with 4 ml of *n*-hexane:ethyl acetate (3:1, v/v).
250 Fractions were eluted with 3 ml of *n*-hexane:ethyl acetate (3:1, v/v) to obtain the simple lipid
251 fraction, 3 ml of ethyl acetate to obtain glycolipids and 10ml of MeOH to obtain
252 phospholipids. The simple lipid fraction was further split into neutral lipid and the fatty acid
253 fractions. The organic phase was then collected into a round bottom flask and Na₂SO₄
254 anhydrous was added until complete removal of water. Silica gel columns (again, 0.5 g, 60 Å
255 particle size) were prepared and conditioned with 4 ml of the recently prepared CHCl₃ sat
256 solution. The simple lipid fraction was then loaded onto the column and subsequently, the
257 neutral lipid fraction was eluted with 9 ml of CHCl₃ sat. Finally, the neutral lipids were
258 separated into apolar and polar lipid fractions. Columns were prepared with approximately
259 0.5 g of activated alumina (Al₂O₃) and compounds eluted with 4 ml of *n*-hexane:DCM (9:1,
260 v/v) and 3 ml of DCM:MeOH (1:2, v/v) to yield the two fractions, respectively. Here, we
261 focus on analyses of the neutral lipid apolar fraction as this is the fraction where diploptene
262 will elute.

263 **3.4 Compound identification and Compound-specific δ¹³C isotope analysis**

264 GC-MS analyses were performed using a Thermoquest Finnigan Trace GC and MS. The GC
265 was fitted with an on-column injector and the stationary phase was CP Sil5-CB. Detection
266 was achieved with electron ionization (source at 70 eV, scanning range 50-580 Daltons). The
267 temperature program consisted of three stages: 70-130 °C at 20 °C/min rate; 130-300 °C at 4
268 °C/min; and 300 °C, temperature held for 10 min.

269 Gas chromatography combustion isotope ratio mass spectrometry (GC-IRMS) was performed
270 using a ThermoScientific Trace GC Ultra coupled to a Conflo IV interface and DeltaV mass
271 Spectrometer. The GC conditions and program were the same as for GC-MS analyses.

272 Calibration was achieved using CO₂ reference gas of known isotopic composition and sample
273 δ¹³C values were expressed against the standard VPDB. All measurements were performed in
274 duplicate.

275 3.5 Mass Balance equation

276 A carbon isotopic mass balance equation (Equation 1), or two-part mixing model, was
277 developed to evaluate the contribution of MOB to the total bacterial biomass, and, therefore,
278 the relative amount of oxidation occurring at each sample location. By developing this
279 mixing model and considering in more detail the potential end member values for the $\delta^{13}\text{C}$
280 values of diploptene derived from different sources (MOB and other heterotrophic bacteria)
281 we can obtain a semi-quantitative estimation of the distribution patterns of MOB across the
282 samples.

283 The resulting end member values are given in table 1. The equation is as follows:

$$284 f_{\text{mob}} = \frac{\delta^{13}\text{C}_{\text{dip_sample}} - \delta^{13}\text{C}_{\text{hetero_dip}}}{\delta^{13}\text{C}_{\text{mob_dip}} - \delta^{13}\text{C}_{\text{hetero_dip}}} \quad (1)$$

285 Where f_{mob} is the fraction of diploptene generated by MOB and $\delta^{13}\text{C}_{\text{dip_sample}}$ is the stable
286 carbon isotopic composition of diploptene in a given sample. $\delta^{13}\text{C}_{\text{hetero_dip}}$ and $\delta^{13}\text{C}_{\text{mob_dip}}$ are
287 the inferred $\delta^{13}\text{C}$ values of diploptene if it were derived solely from heterotrophic bacteria
288 and methanotrophic bacteria, respectively. Both are expressed as the $\delta^{13}\text{C}_{\text{bacterial_biomass}} -$
289 $\Delta^{13}\text{C}_{\text{biosynthesis}}$, the latter term reflecting fractionation during biosynthesis of diploptene.

290 For heterotrophs, it is likely that $\delta^{13}\text{C}_{\text{bacterial_biomass}}$ is similar to that of the substrate organic
291 carbon and is calculated from the $\delta^{13}\text{C}_{\text{bulk_sediment}}$ taken in each zone of the lake. For
292 heterotrophic bacteria, $\Delta^{13}\text{C}_{\text{biosynthesis}}$ can vary from ~2 to 8‰ or more (Pancost and Sinninghe
293 Damsté 2003, and references therein) and a representative value of 4‰ is used here. Given
294 the small range in $\Delta^{13}\text{C}_{\text{biosynthesis}}$ and $\delta^{13}\text{C}_{\text{bulk_sediment}}$ values, the minimum and maximum
295 values for $\delta^{13}\text{C}_{\text{hetero_dip}}$ are similar.

296 For MOB, $\delta^{13}\text{C}_{\text{bacterial_biomass}}$ is calculated from the $\delta^{13}\text{C}_{\text{methane}}$ minus the fractionation that
297 occurs during carbon uptake by methanotrophs (0-30‰; Jahnke et al., 1999). The $\delta^{13}\text{C}_{\text{methane}}$
298 is the measured value of methane captured at seep locations in the thermokarst zones at each
299 lake. As the value is based on a limited number of data points (n=1 and n=5 for Smith L. and
300 Ace L. respectively), it is likely there will be more variation than is seen in the model. In
301 order to incorporate the large range for fractionation that occurs during carbon uptake by
302 methanotrophs (Jahnke et al., 1999), we used both the minimum and maximum value of
303 fractionation (0 and 30‰) to show different scenarios rather than assuming a single value.
304 This is likely larger than variation due to differing $\delta^{13}\text{C}_{\text{methane}}$. With little information
305 available on the fractionation of hopanoids during their biosynthesis by MOB, we assumed a

306 conservative value of 10‰ for our study. This is larger than value assigned for heterotrophic
307 bacteria but still remains a realistic estimate. We calculated mass balances based on both the
308 maximum and minimum end member $\delta^{13}\text{C}$ values for heterotroph- and methanotroph-derived
309 diploptene.

310

311 **4 Results**

312 **4.1 Methane signatures**

313 Early-winter ice-bubble surveys, combined with bubble-trap measurements of ebullition flux
314 and bubble methane concentration, revealed that ebullition seeps occur with high density in
315 the thermokarst zones of both lakes (2.27 seeps m^2 and 4.2 seeps m^2 for Smith L. and Ace L.,
316 respectively as estimated from ice-bubble surveys) compared to the rest of the lake (0.35
317 seeps m^2 and 0.67 seeps m^2 for Smith L. and Ace L., respectively).

318 Seep ebullition rates in the thermokarst zones were 85 and 151 $\text{mg CH}_4 \text{m}^{-2} \text{d}^{-1}$ for Smith L.
319 and Ace L., respectively (Figure 2). In the rest of each lake (lake centre and non-thermokarst
320 margins) seep ebullition rates were 6 and 20 $\text{mg CH}_4 \text{m}^{-2} \text{d}^{-1}$ for Smith L. and Ace L.,
321 respectively. The $\delta^{13}\text{C}$ values for methane in bubbles collected from seeps in the thermokarst
322 zones were -60.9‰ and -64.6‰ for Smith Lake and Ace L., respectively.

323 **4.2 Diploptene $\delta^{13}\text{C}$ values**

324 Diploptene was detected in all but one of the samples analysed (Table 2; figure 3). The
325 isotopic values ranged from -68.2 to -38.8‰ and had an overall standard deviation of 7.8‰.

326 In the Ace L. thermokarst zone, diploptene values ranged from the lowest value for the whole
327 dataset of -68.2 to -50.1‰. Both the most negative and least negative values were found at
328 the greatest water depth (3.2m) in samples located very close to one another, suggesting high
329 variability across small spatial scales.

330 In Smith L., diploptene $\delta^{13}\text{C}$ values ranged from -56.8 to -38.8‰. The most negative value
331 was found in the centre of the lake, and the difference between this and the least negative
332 value (-46.9‰) in the centre of the lake is almost 10‰. In the Smith L. thermokarst zone
333 there was less variability in diploptene $\delta^{13}\text{C}$ (-42.9 to -38.8‰); however, there is still a
334 difference in values of 4.1‰. Samples from the centre of the lake and the thermoakrst zone
335 (n=6, n=3 respectively) were compared using a Mann-Whiney U test (H_0 : diploptene $\delta^{13}\text{C}$

336 values are not different). The test suggested a significant difference between samples from
337 the centre and the thermokarst zone suggesting a difference in bacterial community
338 composition.

339 A comparison of both thermokarst zones shows that diploptene $\delta^{13}\text{C}$ values at Ace L. were
340 more negative than those at Smith L. by at least 10‰. The samples in the thermokarst zone
341 of Ace L. and the centre of Smith L. (n=4, n=6 respectively) were not significantly different
342 according to a Mann Whitney U test.

343 **4.3 Mixing model predictions**

344 The potential contributions of MOB to the diploptene signal under different end-member
345 assumptions are shown in Table 3. The minimum and maximum contributions range from 19
346 to 85%, 7 to 27% and 19 to 63% for Ace L. thermokarst zone, Smith L. thermokarst zone and
347 Smith centre, respectively.

348 Ace L. thermokarst zone had the highest overall potential contributions but also the largest
349 range of predicted values. Smith L. centre had the second highest contribution of MOB to the
350 diploptene signal, and, apart from one sample, suggested a more consistent contribution
351 across the zone. Smith L. thermokarst zone had the lowest potential contribution of the total
352 dataset; even when choosing end member values that yield the greatest MOB contribution,
353 values only reached 27%.

354

355 **5 Discussion**

356 Ace L. thermokarst zone had the highest observed ebullition emission rates, the most
357 depleted $\delta^{13}\text{C}$ diploptene values and the highest potential MOB contribution (according to the
358 mixing model results). The highest ebullition emission rates in Smith L. were in the
359 thermokarst zone, which had the lowest MOB contributions and least depleted $\delta^{13}\text{C}$
360 diploptene values. The centre of Smith L. had very low ebullition rates but depleted $\delta^{13}\text{C}$
361 diploptene values and high predictions of MOB contributions.

362 The $\delta^{13}\text{C}$ diploptene signatures are similar to those that have been previously highlighted as
363 evidence for methanotrophy in lacustrine sediments (-64‰ to -55‰; Spooner et al., 1994;
364 Naeher et al., 2014), marine sediments (-62‰ to -35‰; Freeman et al., 1994; Thiel et al.,
365 2003) and wetlands (-40‰ to -30‰ to; van Winden et al., 2010; Zheng 2014). Therefore, we
366 conclude that diploptene $\delta^{13}\text{C}$ values are documenting the presence of at least some MOB

367 bacteria in lake sediments. The lowest values in Ace L. are among the lowest reported for
368 lacustrine (or other terrestrial) systems, suggesting a relatively high degree of methanotrophy
369 at those sampling sites. Moreover, although the diploptene values were highly variable, the
370 highest values yielded MOB fractions >10%, even when using the most conservative
371 assumptions (Table 3).

372 The results of the mixing model suggest that MOB can contribute anywhere between 7-83%
373 of the diploptene production across all sampled areas (Table 3). These estimates have a large
374 degree of uncertainty and we note that there are some important caveats to using this mixing
375 model. Crucially, diploptene is not derived from all bacteria nor even all methanotrophic
376 bacteria (Rohmer et al., 1987). Nor is it likely to occur in constant biomass-to-lipid ratios in
377 those organisms from which it can derive. Thus, using a diploptene mass balance to infer
378 bacterial biomass distributions should be done cautiously, and the data should be considered
379 semi-quantitative. Nonetheless, a MOB contribution to total biomass of ~10 to 80% is similar
380 to that derived from other studies (11-80%; Bastviken et al. 2003; Sundh et al. 2005;
381 Kankaala et al. 2006). Regardless of absolute MOB estimates, our data show that the centre
382 of Smith L. and the thermokarst zone at Ace L. likely have the highest proportion of MOB in
383 the total bacterial biomass.

384 At Ace L., MOB biomass was high relative to other samples collected in this study and in the
385 context of previous studies. Ace L. has been classified as a 'yedoma-type' lake in previous
386 studies (Walter et al., 2008; Sepulveda-Jauregui, et al., 2015; see above). Walter Anthony and
387 Anthony (2013) suggest that yedoma thermokarst lakes typically produce more methane than
388 non-yedoma thermokarst lakes owing to a higher availability of labile carbon in thick, thawed
389 yedoma sequences. Given the coincidence of high ebullition emission rates, depleted $\delta^{13}\text{C}$
390 diploptene signatures and high estimated MOB biomass, it is likely that the supply of
391 dissolved methane is high in the thermokarst zone and that this methane might be derived
392 from thermokarst-specific sources. Alternatively, lake-edge thermokarst erosion of yedoma-
393 type sediments is also known to supply nitrogen and phosphorus to lakes (Walter Anthony et
394 al. 2014), enhancing primary production, which in turn can fuel methanogenesis and MO
395 from contemporary (atmospheric) carbon (Martinez-Cruz et al., 2015).

396 Within the thermokarst zone at Smith L. the $\delta^{13}\text{C}$ values of diploptene were less variable than
397 in the Ace L. thermokarst zone, and the $\delta^{13}\text{C}$ values were more enriched. In fact, the
398 thermokarst zone in Smith L. had the lowest proportion of MOB for the entire dataset, with a
399 MOB contribution to diploptene being near-equivocal for most of these samples, with values

400 at or below 10% according to the mixing model. On the other hand, despite evidence for
401 much lower methane efflux, samples from the centre of Smith L. had diploptene $\delta^{13}\text{C}$ values
402 that were similar to those of the Ace L. thermokarst zone. The differences between the centre
403 and the thermokarst zone in Smith L. could be explained by several processes. They could
404 arise from variation in the microbial community that is manifest as different MOB
405 expressions of hopanoids. For example, the thermokarst zone MOB might not be
406 biosynthesising diploptene or its precursor. Alternatively, there may be differences in the
407 balance of MO contributing to energy versus biomass production in the bacterial community.
408 Another explanation, which could be validated through further investigation, relates to
409 potential differences in methane production pathways, as highlighted by Walter et al. (2008).
410 In this case, the higher $\delta^{13}\text{C}$ values of diploptene in the thermokarst zone could be due to
411 more enriched methane formed through acetate fermentation. However, the most direct
412 interpretation of the data is that MOB are more abundant in the centre of the lake than at the
413 thermokarst margin and, by extension, more MO is taking place in the lake centre.

414 Overall, the Smith L. thermokarst zone had lower methane ebullition rates and less negative
415 $\delta^{13}\text{C}$ of methane as measured from ebullition flux than Ace L. Therefore, compared with Ace
416 L., the availability of methane produced in this the Smith L. thermokarst zone may be lower
417 due to physical differences in substrate organization. At Smith L. it is likely that methane is
418 not produced in the talik but in near-surface sediments related to peat slumping at the margin.
419 The large size of the sediment blocks and the early stage of decomposition of the slumped
420 organic material may mean there is less exposed substrate surface area for methane
421 production, as compared with yedoma-lake production from fine-grained and more labile
422 talik sediments. Also, methane production in near-surface sediments (often linked to shallow
423 water depths) is subject to reduced partial pressure and faster release of bubbles from the
424 sediment. Bubble tubes initiated in sediments shallower than the talik bulb are likely to be
425 reduced in overall number and size.

426 An important outcome of this study is the large degree of variation seen in the $\delta^{13}\text{C}$ values of
427 diploptene across small spatial distances. The variation does not clearly correspond to
428 patterns in methane production (e.g. high and low ebullition areas). In fact, in Smith L,
429 diploptene $\delta^{13}\text{C}$ values are lower in the low methane flux lake centre than in the high flux
430 thermokarst zone. Despite the caveats associated with interpreting diploptene $\delta^{13}\text{C}$ values, the
431 difference is so large that it likely does indeed reflect more MO in the lake centre. We
432 suggest that this is due to more efficient MO in some diffusive settings than in some

433 thermokarst settings, where ebullition may effectively bypass the MO community. Regardless
434 of mechanism, this variability is a significant finding, as often whole-lake dynamics are
435 interpreted from a single sediment core in palaeoenvironmental studies. Such large variation
436 in $\delta^{13}\text{C}$ values in surface sediments taken from the same zone within a lake – as well as the
437 complex relationships between inferred MO and methane flux – highlight the need for
438 caution when interpreting shifts in $\delta^{13}\text{C}$ values through time using down-core values.

439 Interestingly, a previous study of MOB in lake sediments also reported considerable variation
440 in bacterial communities at small spatial scales (Kankaala et al., 2006). High spatial and/or
441 temporal variability in MOB and other elements of the bacterial biomass could also affect the
442 isotopic composition of heterotrophs higher in the food web, if they consume MOB (e.g.,
443 chironomid larvae). This could have implications for interpretation of not only biomarkers
444 but also other geochemical records. For example, investigations of the biological and
445 geochemical connections between MOB and isotopic signatures of organisms at higher
446 trophic levels are needed, if such organisms are used to interpret past methane emissions.

447 While the results of this study show the potential of diploptene $\delta^{13}\text{C}$ signatures to highlight
448 MO in lakes, further work is needed to understand what this signature is reflecting in terms of
449 methane production and flux. Whether there is a positive correlation between ebullition flux
450 and high diffusion in the thermokarst zone is still to be determined. The current data show no
451 clear link. While this is one of very few studies to use within-lake replicates, and differences
452 are statistically significant, the sample number is small and the system could usefully be
453 tested further prior to developing down-core studies.

454

455 **6 Conclusions**

456 Our primary goal was to contribute towards the understanding of the sedimentary signature of
457 methane production and oxidation in thermokarst lakes using diploptene $\delta^{13}\text{C}$ values as a
458 proxy for the occurrence of MOB. Diploptene was present in almost all samples, and the
459 $\delta^{13}\text{C}$ values were depleted, suggesting the presence of MOB in three zones with differing
460 levels of methane ebullition emissions rates. A two-part mixing model highlighted the
461 potential variation in total MOB biomass, with almost no MOB contributing to bacterial
462 biomass in some samples but forming over half the total bacterial population in others.
463 Critically, these $\delta^{13}\text{C}$ values were highly variable within zones, suggesting small-scale spatial
464 heterogeneity in MOB abundance and thus methane oxidation. The data do not show a

465 consistent relationship between MOB abundance and methane emission rates at the lake
466 surface; in fact, in Smith L, it appears that high MOB abundance occurs where methane
467 emissions are low, suggesting that pathways of carbon flow are as or more important than
468 total flux. Therefore, further investigation of the different types of methane ebullition
469 observed in thermokarst lakes, the relationship between these and diffusion and the different
470 expression of these pathways and MOB biomass are critical. There is also a need to examine
471 localized spatial variability of MO within lakes and how any spatial variation is integrated
472 temporally, as this may critically affect observed down-core patterns of biomarkers and their
473 isotopic signals.

474

475 **Acknowledgements**

476 This research was supported by a NERC grant (NE/K000233/1) to M. Edwards and P.
477 Langdon, a QRA new researcher's award to K. Davies and a PhD Scholarship to K. Davies
478 from Geography and Environment, University of Southampton. We gratefully acknowledge
479 field and equipment assistance from Nancy Bigelow, Charlotte Clarke, Rob Collier and Ben
480 Gaglioti, and permission from the owners to work at Ace Lake. Mark Dover (Cartography
481 Unit, G&E) made valuable improvements to the figures.

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654 Table 1. Mixing model end member values and $\delta^{13}\text{C}$ values of the primary variables used to
655 calculate the proportion of MOB at each sample point. $\delta^{13}\text{C}_{\text{bulk}}$ is the average bulk sediment

656 value from each lake, \pm indicates the standard deviation of the $\delta^{13}\text{C}_{\text{bulk}}$. MOB and
657 heterotrophic bacteria have been assumed to have maximum levels of lipid biosynthesis
658 occurring (10 and 4‰ respectively). $\delta^{13}\text{C}_{\text{mob_dip_min}}$ is the estimated minimum stable isotope
659 value given the $\delta^{13}\text{C}$ value of methane at each lake and the maximum potential fractionation
660 of carbon by MOB. $\delta^{13}\text{C}_{\text{mob_dip_max}}$ is the estimated value of MOB with no fractionation
661 during assimilation. $\delta^{13}\text{C}_{\text{hetero_dip_max}}$ is the maximum estimated stable isotope value of
662 heterotrophic bacteria if no fractionation is occurring during assimilation and the bulk
663 sediment is +1.0 standard deviation (S.D.) from the mean at each lake. $\delta^{13}\text{C}_{\text{hetero-hopane_min}}$
664 represents the minimum value for heterotrophic hopanes given maximum possible
665 fractionation during assimilation and if bulk sediment is -1.0 S.D from the mean.

	$\delta^{13}\text{C}_{\text{bulk}}$			$\delta^{13}\text{C}_{\text{mob_dip_min}}$	$\delta^{13}\text{C}_{\text{mob_dip_max}}$	$\delta^{13}\text{C}_{\text{hetero_dip_min}}$	$\delta^{13}\text{C}_{\text{hetero_dip_max}}$
	(‰)	n	\pm	(‰)	(‰)	(‰)	(‰)
Ace	-30.8	10	2.1	-104.6	-74.6	-36.9	-32.7
Smith	-29.3	10	0.8	-100.9	-70.9	-34.1	-32.5

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681 Table 2 $\delta^{13}\text{C}$ values of diploptene at the study sites. The values are an average of three
 682 replicates. The standard deviation of these replicates and of each zone and across all samples
 683 is also given.

	Sample Number	$\delta^{13}\text{C}_{\text{dip}}$ (‰)	Sample replicate standard Deviation (SD)		Standard Deviation (SD)
Ace					
TK zone	a1	-50.1	1.5		
	a2	-58.5	2.0		
	a3	-53.1	0.4		
	a4	-68.2	0.1	TK zone	8.0
Smith					
Centre TK zone	1	-51.4	2.7		
	2	-48.3	0.0		
	3	-56.8	N/A		
	4	-49.2	1.0		
	5	-46.9	1.8		
	6	-48.0	0.1	Centre	3.6
	7	-38.8	0.3		
	8	-40.9	0.2		
	9	-42.9	0.1		
	10	N/A	N/A	TK zone	2.0
			Total	7.8	

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688 Table 3. Estimated contribution of MOB to the diploptene signal. Calculations assume
 689 fractionation due to biosynthesis of 10‰ for MOB and 4‰ for heterotrophic bacteria. $f_{\text{mob_min}}$
 690 was calculated assuming the highest fractionation for both MOB and heterotrophs (30 and
 691 4‰ respectively). $f_{\text{mob_max}}$ assumes no fractionation during assimilation. $f_{\text{mob_average}}$ was
 692 calculated using average $\delta^{13}\text{C}$ values for $\delta^{13}\text{C}_{\text{mob-hopane}}$ and $\delta^{13}\text{C}_{\text{hetero-hopane}}$.

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	Sample Number	$f_{\text{mob_min}}$	$f_{\text{mob_max}}$	$f_{\text{mob_average}}$	
Ace					
TK zone	a1	0.19	0.42	0.28	
	a2	0.32	0.62	0.43	
	a3	0.24	0.49	0.33	
	a4	0.46	0.85	0.61	
Smith					
Centre	1	0.26	0.49	0.34	
	2	0.21	0.41	0.28	
	3	0.34	0.63	0.45	
	4	0.23	0.44	0.30	
	5	0.19	0.37	0.26	
	6	0.21	0.40	0.28	
	TK zone	7	0.07	0.17	0.11
		8	0.10	0.22	0.14
		9	0.13	0.27	0.18

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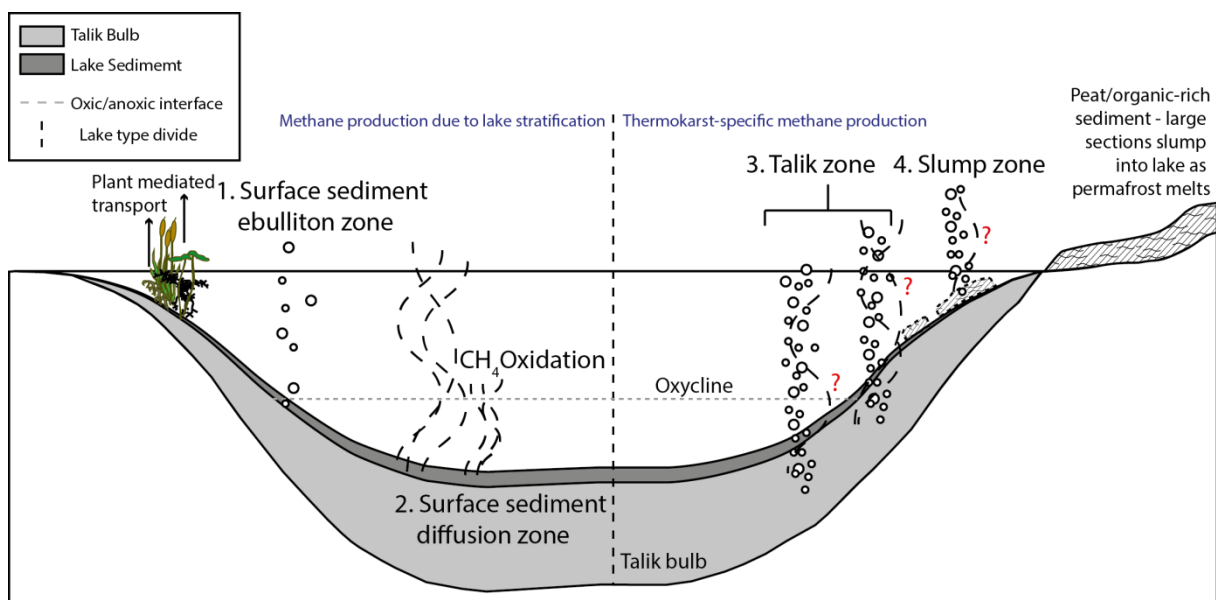
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701 Figure 1. Illustration of methane production zones and emission pathways in lakes alongside
 702 thermokarst-specific zones and pathways. 1) Surface sediment ebullition zone (background
 703 methane production). Methane that is produced in the anoxic surface sediments is released
 704 via ebullition, usually near the margins (Bastviken et al., 2004). (2) Surface sediment
 705 diffusion zone. Methane is produced in the anoxic surface sediments and diffuses in the
 706 sediments above and into the water column. Some of this methane will reach the water
 707 surface-air interface but a large amount is likely to be oxidised by MOB (Kankaala et al.,
 708 2006). This process is common in many lakes also. (3) Talik zone. Methane is produced in
 709 the deeper talik sediments underneath the lake and is released via ebullition seeps (Walter et
 710 al. 2008). Often this is a higher flux and is more constant than surface sediment ebullition.
 711 This production zone and pathway is a thermokarst-specific process. (4) Slump zone.
 712 Methane production in the surface sediments is increased due to the introduction of large
 713 volumes of slumped sediments. This methane is also released via ebullition seeps. Often, the
 714 flux from these ebullition seeps is higher than surface sediment ebullition but not as high as
 715 talik ebullition. This process might occur in any lakes that have dynamic margins and high
 716 erosion rates; however, it is likely that this process is most common in thermokarst lakes due
 717 to the melting of permafrost, so it is termed thermokarst-specific. Red question marks
 718 indicate where methane diffusion from the sediments has not been studied in detail.

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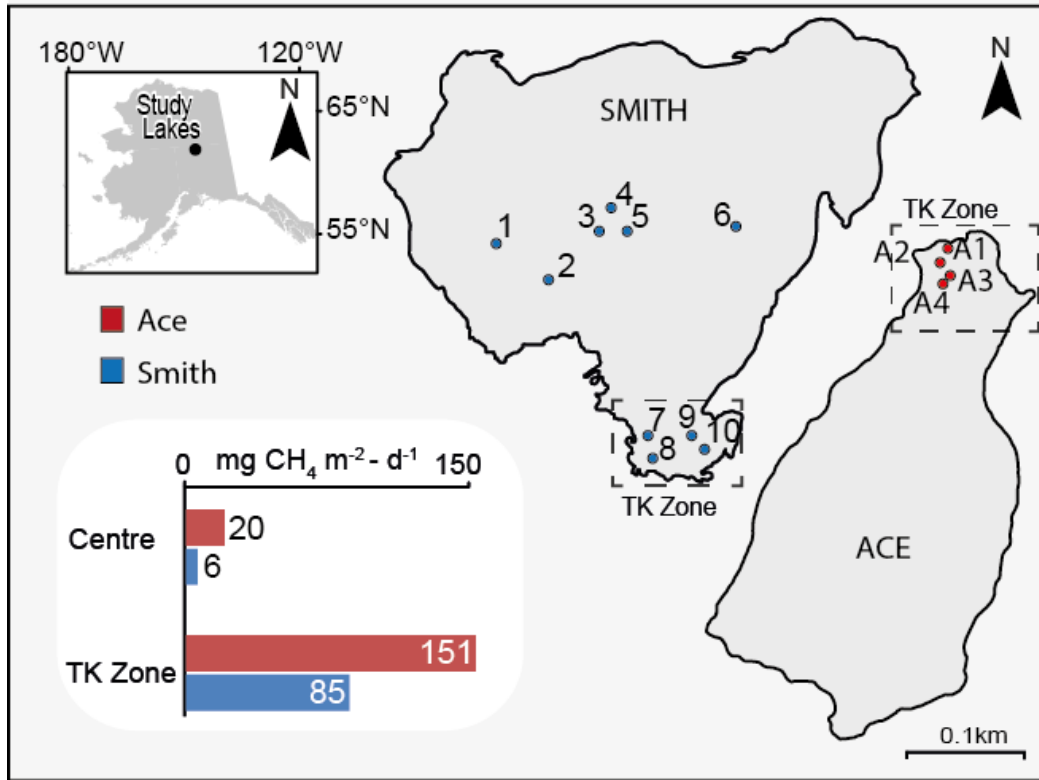


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723 Figure 2. Locations of the study lakes in Alaska and the sediment sample points within each
724 lake. The red (Ace L.) and blue (Smith L.) bars indicate the flux values as averaged within a
725 given area of the lake. Flux measurements were taken on October 2009 at Smith L and April,
726 2011 at Ace L.



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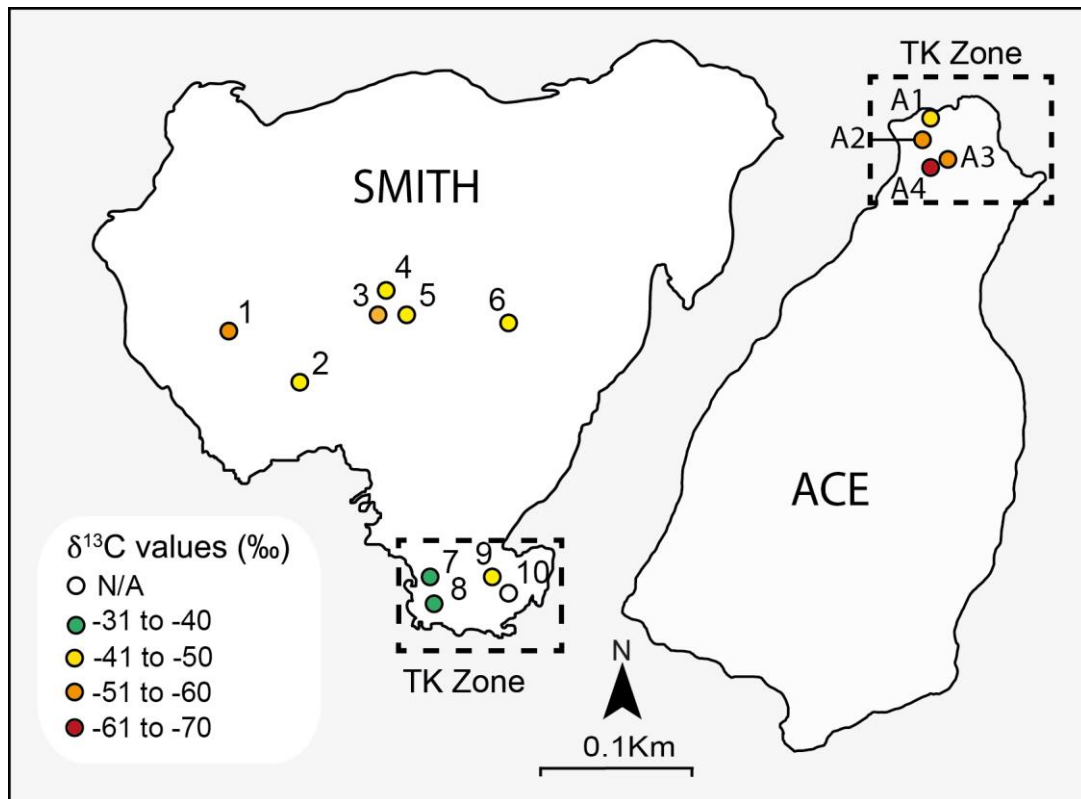
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739 Figure 3. Diploptene $\delta^{13}\text{C}$ values at Smith Lake and Ace Lake. In general the most depleted
740 values are found in Ace and in the centre of Smith. The Thermokarst zone at Smith L. has the
741 least depleted values for the whole dataset



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