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Glucose addition increases the magnitude and decreases the age of soil respired carbon in a long-term permafrost incubation study

Elaine Pegoraro, Marguerite Mauritz, Rosvel Bracho, Chris Ebert, Paul Dijkstra, Bruce A. Hungate, Kostas T. Konstantinidis, Yiqi Luo, Christina Schädel, James M. Tiedje, Jizhong Zhou, Edward A.G. Schuur

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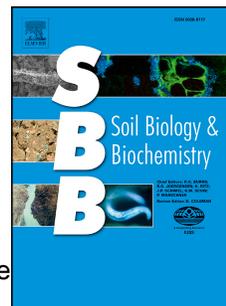
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4 Authors: Elaine Pegoraro^{a,b}, Marguerite Mauritz^{a,b}, Rosvel Bracho^c, Chris Ebert^{a,b}, Paul
5 Dijkstra^{a,b}, Bruce A. Hungate^{a,b}, Kostas T. Konstantinidis^d, Yiqi Luo^{a,b,e,g}, Christina
6 Schädel^{a,b}, James M. Tiedje^f, Jizhong Zhou^g, and Edward A.G. Schuur^{a,b}.

7
8 Affiliations:

9 ^aDepartment of Biological Sciences, Northern Arizona University, Flagstaff, AZ 86011,
10 USA

11 ^bCenter for Ecosystem Science and Society, Northern Arizona University, Flagstaff, AZ
12 86011, USA

13 ^cSchool of Forest Resources & Conservation, University of Florida, Gainesville, FL 32611,
14 USA

15 ^dSchool of Civil and Environmental Engineering and School of Biology, Georgia Institute
16 of Technology, Atlanta, Georgia 30332, USA

17 ^eDepartment of Earth System Science, Tsinghua University, Beijing 100084, China

18 ^fDepartment of Plant, Soil and Microbial Sciences, Center for Microbial Ecology,
19 Michigan State University, East Lansing, Michigan 48824, USA

20 ^gDepartment of Microbiology and Plant Biology, University of Oklahoma, Norman,
21 Oklahoma 73019, USA

22
23
24 Corresponding author: Elaine Pegoraro, (407) 412 8881, efp23@nau.edu

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27 **Abstract**

28 Higher temperatures in northern latitudes will increase permafrost thaw and stimulate
29 above-and belowground plant biomass growth in tundra ecosystems. Higher plant
30 productivity increases the input of easily decomposable carbon (C) to soil, which can
31 stimulate microbial activity and increase soil organic matter decomposition rates. This
32 phenomenon, known as the priming effect, is particularly interesting in permafrost
33 because an increase in C supply to deep, previously frozen soil may accelerate
34 decomposition of C stored for hundreds to thousands of years. The sensitivity of old
35 permafrost C to priming is not well known; most incubation studies last less than one
36 year, and so focus on fast-cycling C pools. Furthermore, the age of respired soil C is
37 rarely measured, even though old C may be vulnerable to labile C inputs. We incubated
38 soil from a moist acidic tundra site in Eight Mile Lake, Alaska for 409 days at 15°C. Soil
39 from surface (0-25 cm), transition (45-55 cm), and permafrost (65-85 cm) layers were
40 amended with three pulses of uniformly ^{13}C labeled glucose or cellulose, every 152 days.
41 Glucose addition resulted in positive priming in the permafrost layer 7 days after each
42 substrate addition, eliciting a two-fold increase in cumulative soil C loss relative to
43 unamended soils with consistent effects across all three pulses. In the transition and
44 permafrost layers, glucose addition significantly decreased the age of soil-respired $\text{CO}_2\text{-C}$
45 with $\Delta^{14}\text{C}$ values that were 115‰ higher. Previous field studies that measured the age
46 of respired C in permafrost regions have attributed younger $\Delta^{14}\text{C}$ ecosystem respiration
47 values to higher plant contributions. However, the results from this study suggest that
48 positive priming, due to an increase in fresh C supply to deeply thawed soil layers, can
49 also explain the respiration of younger C observed at the ecosystem scale. We must
50 consider priming effects to fully understand permafrost C dynamics, or we risk
51 underestimating the contribution of soil C to ecosystem respiration.

52 1. Introduction

53 Temperatures in northern latitudes are currently increasing by 0.6°C each decade, a
54 rate that is two times higher than the global average (IPCC, 2013). Higher temperatures
55 can thaw permafrost (Brown and Romanovsky, 2008; Romanovsky et al., 2010; Harden
56 et al., 2012; Romanovsky et al., 2012) and expose previously protected soil organic
57 matter (SOM) to microbial decomposition (Schuur et al., 2015). Though the permafrost
58 region encompasses only 15% of the total global soil area, it stores two times more
59 carbon (C) than the atmosphere: 1,330–1,580 petagrams C (Pg; 1 Pg = 1 billion metric
60 tons) (Zimov et al., 2006; Tarnocai et al., 2009; Hugelius et al., 2014; Schuur et al., 2015).
61 Microbial mineralization of this stored C can exacerbate the effects of climate change by
62 increasing greenhouse gas concentrations in the atmosphere in the form of carbon
63 dioxide (CO₂) and methane (Schuur et al., 2008).

64 The mineralization rate of permafrost C is partially controlled by its inherent
65 decomposability (Schädel et al., 2014; Schuur et al., 2015). Upon thaw, permafrost C is
66 mobilized (Schuur et al., 2009; Nowinski et al., 2010), and overall soil C losses are high
67 because of the high biolability of organic C in initial stages of decomposition (Dutta et
68 al., 2006; Vonk et al., 2013). However, that initial C is quickly mineralized, and microbes
69 are left with a much larger pool of slowly decomposing C. A previous soil incubation
70 study found that > 85% of the fast decomposing C is depleted after ~ 3 months of
71 incubation at 15°C (Bracho et al., 2016), but it amounts to less than 5% of all C in both
72 organic and mineral soil layers (Schädel et al., 2014). Although soil C comprises several
73 different pools with turnover times spanning less than a year to several hundred years

74 (Trumbore, 2000), the majority of the C stored in permafrost is old (Zimov et al., 2006;
75 Schuur et al., 2009). Therefore, we expect that long-term C losses will largely originate
76 from old and slowly decomposing C pools (Schuur et al., 2008; Knoblauch et al., 2013;
77 Schädel et al., 2013; Schädel et al., 2014).

78 Slowly decomposing C that accumulated over hundreds to thousands of years in
79 permafrost (Czimczik and Welker, 2010; Pries et al., 2012) is thought to be relatively
80 energy-poor. A long-term (~ 1 year) permafrost incubation study revealed that less than
81 3% of the initial soil bulk C is mineralized in the absence of substrate inputs (Dutta et al.,
82 2006). Without an energy source in the form of fresh organic C, microbial activity
83 decreases (Schimel and Weintraub, 2003); however, this energy limitation can
84 potentially be lifted with an increased input of labile C to soil (De Nobili et al., 2001;
85 Fontaine et al., 2007; Blagodatskaya and Kuzyakov, 2008).

86 A warmer climate will also increase shrub expansion in tundra ecosystems, which
87 will increase plant productivity and litter input to soil (Chapin et al., 1995; Shaver et al.,
88 2000; Tape et al., 2006; Natali et al., 2012). Priming theory suggests that a greater
89 supply of fast decomposing C from leaf and root litter, root exudates, and dissolved
90 organic C leachate can increase microbial decomposition of native SOM (Bingeman et
91 al., 1953). Studies show that microbes utilize fresh C as an energy source to produce
92 enzymes that assist in the decomposition of organic molecules that are resistant to
93 microbial degradation (Schimel and Weintraub, 2003; Fontaine et al., 2007; Bernal et al.,
94 2016). Some studies have investigated the implication of tundra shrub expansion on
95 SOM decomposition by adding glucose or low molecular weight C to soil, and found no

96 effects of priming (Rousk et al., 2016; Lynch et al., 2018). However, these studies
97 amended surface and O-horizon soils exclusively, which do not reflect the potential
98 priming effects of deep-rooting tundra species (e.g. *R. chamaemorus* and *E. vaginatum*)
99 whose roots extend to the thaw-front (>45 cm) (Keuper et al., 2017), and thus may
100 increase C input to deep soils and recently thawed permafrost.

101 Priming is a particularly interesting phenomenon in permafrost soil because frozen
102 ground is impermeable to dissolved organic C (DOC) infiltration (Walvoord and Kurylyk,
103 2016). Environmental changes that increase C inputs in deep soil can increase soil C
104 vulnerability to decomposition and loss (Bernal et al., 2016). This means that as
105 permafrost thaws, soil in deep layers may receive fresh C inputs during rain and snow
106 events for the first time in hundreds to thousands of years. Previous studies modeling
107 water movement through soil with underlying permafrost show that DOC input to
108 deeper layers releases more CO₂ (Fan et al., 2013). Fresh C input to soil will also be
109 accompanied by pulses of mineralized N upon permafrost thaw (Rustad et al., 2001;
110 Keuper et al., 2012; Salmon et al., 2018), which could create an ideal environment for
111 higher microbial activity.

112 A more active microbial community, especially in deep soil layers, can alter soil C
113 dynamics by increasing decomposition of old and slowly decomposing organic C, and
114 accelerate C turnover rates (Fontaine et al., 2007; Paterson et al., 2009). While the
115 stability of old soil C can change in the presence of labile C inputs (Fontaine et al., 2007;
116 Bernal et al., 2016), permafrost priming studies have not measured the age of soil
117 respired C. Furthermore, most incubation studies are conducted in short-term

118 laboratory incubations (< 1 year), and thus only focus on fast-cycling soil C pools (Fan et
119 al., 2013; Wild et al., 2014; Wild et al., 2016). To fully understand permafrost C dynamics
120 and priming effects, we measured the magnitude of soil C loss and the age of respired C.

121 We hypothesized that amending permafrost soil with multiple pulses of glucose,
122 the most abundant sugar in rhizodeposits (Derrien et al., 2004), or cellulose, the most
123 common polymer in plant litter (Kögel-Knabner, 2002), would: (1) increase soil C
124 decomposition (positive priming), particularly in permafrost where the majority of the C
125 is slow decomposing C (Schädel et al., 2014), (2) sustain higher rates of soil C
126 decomposition over the long-term because microbes will readily use substrates as an
127 energy source following each pulse, and (3) increase the proportion of old C that is
128 respired as a result of soil priming because it comprises the bulk of the soil C pool in
129 permafrost (Schuur et al., 2009).

130 **2. Materials and Methods**

131 *2.1. Site Description*

132 Soil cores were sampled in 2013 from Eight Mile Lake, Alaska (63° 52' 59"N, 149°
133 13' 32"W), where the mean annual temperature is -1.0 °C, and the mean annual
134 precipitation is 378 mm (Schuur et al., 2009). The site is situated on a well-drained,
135 northeast-facing hillslope (700 m) (Natali et al., 2011) in the discontinuous permafrost
136 zone, but is underlain entirely by permafrost (Osterkamp et al., 2009). Though our site is
137 in the subarctic, it exemplifies a region vulnerable to permafrost degradation that is
138 analog to projected conditions in the Arctic as temperatures increase. The site has
139 cryoturbated mineral soil, comprising glacial till and windblown loess, with dominant

140 amounts of quartz and feldspars. The soil type is Gelisol (Soil Survey Staff, 2014) with an
141 organic horizon ~35 cm thick, and C concentrations greater than 20% (Natali et al., 2011;
142 Pries et al., 2012; Plaza et al., 2017b). In 2013, the maximum thaw depth at the peak of
143 the growing season was < 65 cm (Mauritz et al., 2017). The vegetation is moist acidic
144 tundra, dominated by *Eriophorum vaginatum*. Plant species composition includes
145 *Vaccinium uliginosum*, *Carex bigelowii*, *Betula nana*, *Rubus chamaemorus*, *Empetrum*
146 *nigrum*, *Rhododendron subarcticum*, *V. vitis-idaea*, *Andromeda polifolia* and *Oxycoccus*
147 *microcarpus*. Nonvascular plant cover contains feather moss (primarily *Pleurozium*
148 *schreberi*) and *Sphagnum* species, as well as several lichen species (primarily *Cladonia*
149 spp.) (Schuur et al., 2007; Natali et al., 2011; Deane-Coe et al., 2015).

150 2.2. Soil Core Collection and Processing

151 Four soil cores were collected in June 2013. The seasonally thawed surface soil was
152 cut using a serrated knife, and the underlying frozen soil was cored using a Tanaka drill
153 with a 7.6 cm diameter hollow bit. Soil was sampled to the depth at which the corer hit
154 rocks (~ 85 cm). Cores were wrapped in aluminum foil, kept frozen for shipment, and
155 stored until the start of the experiment.

156 In the lab, the surface vegetation was clipped off, and cores were sectioned to 15
157 cm at the surface and 10 cm increments thereafter (0-15 cm, 15-25 cm, etc.) to the end
158 of the core. The organic/mineral horizon demarcation was determined by %C analysis
159 (mineral < 20% C). Rocks were removed, and we accounted for their mass and volume in
160 bulk soil density calculation. Each depth increment was subsampled for moisture
161 content, and bulk %C, %N, and $\delta^{13}\text{C}$ (‰). Bulk C and N concentrations were determined

162 by dry combustion using a Costech Analytical ECS 4010 elemental analyzer (Valencia, CA,
163 USA) (Pries et al., 2012). Stable C isotope data were determined using a Thermo
164 Finnigan Delta V Advantage continuous flow isotope ratio mass spectrometer
165 (ThermoScientific Inc., Waltham, MA, USA). Soil pH was determined using the slurry
166 method with a 1:1 ratio by proportional weight of soil to deionized water. Soil pH was
167 measured using an Orion 2 Star pH Benchtop (ThermoScientific Inc., Waltham, MA,
168 USA).

169 We measured initial microbial biomass on a subsection of soil that was not
170 incubated. Due to constrains in soil availability, only 3 cores at depths 0-15, 15-25 and
171 45-55 cm, and only 2 cores at 75-85 cm were extracted; we did not have enough soil
172 available for extraction at 65-75 cm. Four analytical replicate samples (4-6 g soil) from
173 each individual core and depth increment were homogenized, and soils were extracted
174 with 25 ml 2M KCl. Two of the replicates were amended with 0.25 ml chloroform to lyse
175 microbial cell membranes. Samples were shaken for 1 hour, then vacuum filtered
176 through pre-leached Whatman GF/A filters. Extracts were sparged with air for 30
177 minutes to volatilize chloroform, and frozen for storage. Dissolved organic C
178 concentrations were measured using a Shimadzu TOC-L analyzer (Kyoto, Japan), and
179 microbial biomass C was calculated as the difference in DOC between extracts
180 performed with and without chloroform (Salmon et al., 2018).

181 2.3. Incubation

182 We incubated soil from five different depths, representative of the surface layer (0-
183 15 and 15-25 cm), the organic/mineral transition layer (45-55 cm), and the permafrost

184 layer (65-75 and 75-85 cm), from 4 replicate soil cores. We weighed triplicates (~ 80 g
185 wet weight) from each depth into 1 L Mason jars and assigned each jar to a treatment:
186 unamended (control), glucose, or cellulose (4 cores x 5 depth x 3 treatments = 60 jars).

187 We pre-incubated all samples for 4 weeks at 15°C so that waterlogged soils from
188 ice-rich sections (65-75 and 75-85 cm) could air dry to less than 60% water holding
189 capacity (WHC). At the end of the 4 weeks, we amended our samples with uniformly ¹³C
190 labeled glucose U-13C6, 24-25 atom% (Cambridge Isotope Laboratories Inc., Andover,
191 MA, USA) or cellulose 97 atom% D from maize (Sigma-Aldrich Co., St. Louis, MO, USA).
192 Soils were amended with 3.5 mg substrate C per gram of initial soil C to account for
193 differences in initial bulk soil %C in each layer. The total concentration added
194 corresponded to 23-84% of the annual net primary productivity (NPP) in our site, scaled
195 to a per gram soil basis (Natali et al., 2012). Labeled substrates were mixed with
196 unlabeled glucose or cellulose (Sigma Life Science, St. Louis, MO, USA) for more diluted
197 $\delta^{13}\text{C}$ values (Supplemental Table 1). Deionized water was added to the mixtures of finely
198 ground glucose or cellulose so that substrate solutions could be injected in each soil
199 section with a syringe to minimize soil disturbance and maximize distribution.
200 Unamended soils received deionized water only. Moisture was adjusted to ~ 60% WHC
201 throughout the incubation in all treatments. Soils were aerobically incubated and
202 maintained at 15°C. A second and third amendment pulse was added at day of
203 incubation (DOI) 153 and 305 (every 152 days). We chose to amend soils every 152 days
204 because Bracho et al. (2016) found that 85-95% of fast-decomposing C is depleted

205 within 100 days of incubation at 15°C; therefore, we assumed most of the added
206 substrate would be consumed before the next C addition.

207 2.4. Flux and isotope measurements

208 Carbon dioxide and $\delta^{13}\text{C}$ measurements were coupled and measured daily for the
209 first week following each pulse, every 2 days for the second week, and once every ~4
210 weeks until the next amendment.

211 Jars were placed in a water bath set to 15°C, connected to an automated soil
212 incubation system (ASIS) that sequentially measured CO_2 concentration in each jar by
213 circulating air through an infrared gas analyzer (IRGA) at 0.9 L min^{-1} (Li-820 Licor,
214 Lincoln, Nebraska). Pressure and CO_2 concentrations in each jar were recorded on a data
215 logger (CR1000, Campbell Scientific, Logan UT) every three seconds for 8 minutes.
216 Carbon dioxide flux, in micrograms of C per gram of initial soil C per day, was calculated
217 as the rate of CO_2 -C increase in the headspace of each jar over 3-4 cycles that lasted 8.5
218 hours each. ASIS is a closed-loop system, but individual jar fluxes were corrected for
219 carry over CO_2 based on the CO_2 concentration in the previous jar and the carry over
220 volume, which included the tubing, IRGA, flow mass controlled, and pump. Details of
221 ASIS are described in Bracho et al. (2016).

222 After each flux measurement, jar headspace was scrubbed for 5 minutes with soda
223 lime and incubated at 15°C for a couple of hours to allow CO_2 to accumulate. We
224 measured $\delta^{13}\text{CO}_2$ -C on a Picarro G2201-*i* Isotopic CO_2 / CH_4 cavity ring-down
225 spectrometer (Picarro Inc., Sunnyvale, California, USA) and recorded the 2-minute
226 $\delta^{13}\text{CO}_2$ -C average for each jar at the end of a 4-minute measurement period. We

227 allowed the Picarro cavity to return to ambient $^{13}\text{CO}_2$ levels before measuring each jar,
228 and we frequently used a reference CO_2 gas before and after measurements for
229 calibration (Airgas, ASG Los Angeles, CA).

230 Because of the high number of jars, the incubation was conducted in 2 rounds,
231 staggered in time to manage the high volume of measurements. Each round was treated
232 the same, and treatments were fully interspersed between rounds.

233 2.5. Priming calculations

234 In the amended treatments, total CO_2 -C flux was a combination of soil-derived C
235 and substrate-derived C, so we applied an isotopic mass balance equation to partition
236 soil-derived C (SOC) from substrate-derived C:

$$237 \quad C_{\text{SOC}} = C_{\text{total}} (\delta_{\text{total}} - \delta_{\text{substrate}}) / (\delta_{\text{SOC}} - \delta_{\text{substrate}}) \quad (1)$$

238 where C_{SOC} is the soil-respired C, C_{total} is total C respired in substrate-amended soils,
239 which includes soil-derived C and substrate-derived C, δ_{total} is the total $\delta^{13}\text{CO}_2$ -C
240 respired, which includes soil-derived $\delta^{13}\text{C}$ and substrate-derived $\delta^{13}\text{C}$, $\delta_{\text{substrate}}$ is the δ
241 $^{13}\text{CO}_2$ -C from the added substrate (end-member), and δ_{SOC} is the soil-respired $\delta^{13}\text{CO}_2$ -C
242 from unamended samples (end-member). Priming was calculated as the difference in
243 soil-respired C in amended samples and soil-respired C in unamended samples (Mau et
244 al., 2015). A positive value indicated that more soil C was mineralized in amended soils
245 relative to control (i.e. positive priming). A negative value indicated that less soil C was
246 mineralized in amended soils relative to control (i.e. negative priming). All values were
247 reported as micrograms of respired C per gram of initial soil C.

248 Though we amended soils with cellulose in Pulse 1, the dilution of the labeled
249 substrate with unlabeled cellulose made the $\delta^{13}\text{C}$ signature too similar to that of the
250 soil; therefore, we were unable to use the substrate end-member to calculate priming.
251 For pulses 2 and 3, we used a cellulose substrate with a higher $\delta^{13}\text{C}$ signature
252 (Supplemental Table 1). We only analyzed and interpreted cellulose data for pulses 2
253 and 3.

254 To calculate substrate use, or the amount of substrate that was respired after each
255 pulse, we implemented a mass balance equation to partition substrate-derived C
256 ($C_{\text{substrate}}$) from the total C respired (C_{total}):

$$257 \quad C_{\text{substrate}} = C_{\text{total}} (\delta_{\text{total}} - \delta_{\text{soil}}) / (\delta_{\text{substrate}} - \delta_{\text{soil}}) \quad (2)$$

258 We calculated the cumulative $C_{\text{substrate}}$ respired after each pulse as a percent of the
259 added substrate ($3.5 \text{ mg } C_{\text{substrate}} \text{ g}^{-1} \text{ C}$) by linearly interpolating the respired $C_{\text{substrate}}$
260 values and integrating the entire sampling period of 105 days.

261 2.6. Radiocarbon

262 To determine the age of respired soil C, we sampled $\Delta^{14}\text{CO}_2$ at DOI 15 and 105 after
263 pulse 1, and DOI 319 (i.e. day 15 after pulse 3). The headspace in each jar was scrubbed
264 free of CO_2 using soda lime to remove any background atmospheric CO_2 contributions.
265 After allowing each headspace to accumulate between 0.5-1.0 mg C, based on the most
266 recent flux rates measured, we collected CO_2 in zeolite molecular sieve traps (Alltech
267 13X; Alltech Associates, Deerfield, IL, USA) for 5 minutes (Hardie et al., 2005). Each
268 molecular sieve trap was baked at 650°C to desorb CO_2 (Bauer et al., 1992). Carbon
269 dioxide was cryogenically purified using liquid nitrogen, and reduced to graphite by H_2

270 reduction with an Fe catalyst on a vacuum line (Vogel et al., 1987). Graphite samples
 271 were sent to the UC Irvine W.M. Keck carbon cycle accelerator mass spectrometry
 272 (AMS) laboratory for $\Delta^{14}\text{C}$ analysis. Radiocarbon samples are analyzed with the standard
 273 oxalic acid II, and the precision of an AMS measurement is $\pm 2\%$, but for many
 274 applications this error is smaller than sampling error (Trumbore et al., 2016b).

275 Radiocarbon values in unamended samples were corrected for mass-dependent
 276 fractionation to a $\delta^{13}\text{C}$ value of -25% , which is routinely done to natural abundance
 277 samples (Stuiver and Polach, 1977; Trumbore et al., 2016a). For samples amended with
 278 enriched ^{13}C substrates, the increase in ^{13}C atoms relative to ^{12}C is not due to mass-
 279 dependent isotopic fractionation—that is, ^{14}C is not enriched, only ^{13}C is. Therefore,
 280 using $\delta^{13}\text{C} = -25\%$ as a fractionation correction would yield an inaccurate $\Delta^{14}\text{C}$. Thus,
 281 we applied the correction described in Torn and Southon (2001), using the measured
 282 $\delta^{13}\text{C}_{\text{substrate}}$ value as a proxy for the isotopic fractionation correction.

283 Amended samples had a $\Delta^{14}\text{C}$ signature from combined soil and substrate respired
 284 C; therefore, we calculated the soil respired $\Delta^{14}\text{C}$ fraction (f) using $\delta^{13}\text{C}$ signatures of the
 285 substrate and SOC with a similar mass-balance equation implemented in the priming
 286 calculation:

$$287 \quad f = (\delta_{\text{total}} - \delta_{\text{substrate}}) / (\delta_{\text{SOC}} - \delta_{\text{substrate}}) \quad (3)$$

288 where δ_{total} is the total $\delta^{13}\text{CO}_2\text{-C}$ respired, $\delta_{\text{substrate}}$ is the $\delta^{13}\text{CO}_2\text{-C}$ from the added
 289 substrate (end-member), and δ_{SOC} is the soil-respired $\delta^{13}\text{CO}_2\text{-C}$ from control samples
 290 (end-member). We then applied the f of soil respired $\Delta^{14}\text{C}$ to the following equation:

$$291 \quad \Delta^{14}\text{C}_{\text{SOC}} = \Delta^{14}\text{C}_{\text{total}} - ((1-f) * \Delta^{14}\text{C}_{\text{substrate}}) / f \quad (4)$$

292 This soil respired $\Delta^{14}\text{C}$ fraction (f) allows us to calculate the true $\Delta^{14}\text{C}$ value of soil
293 respiration ($\Delta^{14}\text{C}_{\text{SOC}}$) by removing any substrate contribution ($\Delta^{14}\text{C}_{\text{substrate}}$) from the $\Delta^{14}\text{C}$
294 measured by the AMS ($\Delta^{14}\text{C}_{\text{total}}$) (Schuur and Trumbore, 2006). The $\Delta^{14}\text{C}_{\text{substrate}}$ was
295 10.1‰ and 62.3‰ for glucose and cellulose, respectively.

296 2.7. Statistical Analysis

297 Statistical analyses of the main effects and interactions on response variables were
298 performed in R (R Development Core Team, 2015) using linear mixed effects models,
299 lme4 package (Bates et al., 2015). To meet normality assumptions, data were log-
300 transformed when necessary. We removed 6 outlier priming ($\mu\text{g CO}_2\text{-C g}^{-1}\text{C}$) values out
301 of 1347 values (<1% of the data). Normality and homoscedasticity were visually
302 examined using residual plots.

303 Samples from each core ($n=4$) were measured at 5 different depths (0-15, 15-25,
304 45-55, 65-75, and 75-85 cm), but the data were analyzed by pooling depth increments
305 into 3 layers: surface (0-15 and 15-25 cm), transition (45-55 cm), and permafrost (65-75
306 and 75-85 cm) based on statistically similar initial bulk soil %C. Analyses were performed
307 by including soil layer nested in soil core as a random effect so that depth increments
308 would not be treated as independent replicates. Initial soil properties were tested by
309 incorporating soil layer as a fixed effect, and a random effect of soil layer nested in soil
310 core. A post-hoc Tukey HSD test was used to determine significant differences between
311 layers.

312 Priming ($\mu\text{g CO}_2\text{-C g}^{-1}\text{C}$) was tested in response to soil layer, pulse, day(s) after pulse
313 (DAP), and their interactions. Analyses included a random effect of jar nested in soil

314 layer, nested in core, to account for repeated measurements. We accounted for round
315 ($n = 2$) as a random effect in our model, but it had no effect on variance, so we dropped
316 the random effect term. We fit separate models for glucose priming and cellulose
317 priming to avoid an unbalanced design because cellulose data was missing in pulse 1.
318 Changes in cumulative soil C loss in the first week after each pulse were calculated as
319 the response ratio—the ratio between samples amended with glucose and
320 corresponding unamended samples.

321 Soil respired $\Delta^{14}\text{C}$ (‰) was tested in response to treatment, layer, and DOI.
322 Analyses also included a random effect of jar nested in layer, nested in core. We did not
323 have to analyze glucose and cellulose independently because we had radiocarbon data
324 for both treatments for the three different sampling periods. However, to meet
325 conditions of homoscedasticity, we analyzed surface layer samples independently from
326 transition and permafrost layers. Surface soils have C stocks that are more homogenous,
327 and have positive $\Delta^{14}\text{C}$ (‰) values that correspond to modern C that contain elevated
328 ‘bomb’ $\Delta^{14}\text{C}$ values (Turnbull et al., 2016), whereas transition and permafrost layers are
329 more heterogeneous due to cryoturbation, soil subsidence, and soil formation (Harden
330 et al., 2012), and have negative $\Delta^{14}\text{C}$ (‰) values as a result of radioactive decay
331 (Turnbull et al., 2016).

332 We used a Backward step-wise model selection to eliminate fixed effects that
333 increased AIC values by 5 or more. Ninety-five percent confidence intervals (CI) for fixed
334 effects were obtained by bootstrapping parameter estimates (1000 iterations). A fixed

335 effect was considered significant if the 95% CI did not include zero (Pinheiro and Bates,
336 2000).

337 **3. Results**

338 *3.1. Soil properties*

339 Soil properties significantly differed between soil layers (Table 1). Initial bulk soil C
340 and N concentrations significantly decreased with depth; concentrations in the surface
341 were almost 2 times higher than the transition layer, and 4 times higher than the
342 permafrost layer, but there were no significant differences in C:N. Bulk density was
343 significantly lower in the surface, but was not different between the transition and
344 permafrost layers. Carbon concentrations decreased from about 38% in the surface to
345 21% in the transition, and to less than 10% in the permafrost layer. Initial bulk $\delta^{13}\text{C}$
346 values were significantly higher in the surface layer, but did not differ between the
347 transition and permafrost layers. Initial microbial biomass was not significantly different
348 between layers, but was marginally higher in the surface ($1.22 \text{ mg C g}^{-1} \text{ soil C}$) than the
349 transition ($0.36 \text{ mg C g}^{-1} \text{ soil C}$) and permafrost layers ($0.66 \text{ mg C g}^{-1} \text{ soil C}$). Soil pH was
350 in the acidic range, and significantly increased with depth.

351 *3.2. Priming effects*

352 The addition of glucose elicited immediate priming responses in both surface and
353 permafrost layers, though these responses differed in sign and magnitude. Priming in
354 the surface layer was mostly negative in the first week after each pulse. We observed
355 significantly negative priming 48h after each glucose amendment (Surface x 2 DAP: CI -

356 77.36 $\mu\text{g CO}_2\text{-C g}^{-1}\text{C}$ to -9.09 $\mu\text{g CO}_2\text{-C g}^{-1}\text{C}$, Fig. 1, Supplemental Table 2) with consistent
357 effects in all pulses.

358 In the permafrost layer, glucose addition resulted in significant positive priming
359 effects that lasted 7 days after each pulse; this response was similar across all three
360 pulses (Fig. 1, Supplemental Table 2). This strong priming effect in the glucose treatment
361 doubled the cumulative soil C loss relative to the unamended permafrost layer; the
362 cumulative response ratio of soil C loss in glucose treatments relative to control was
363 close to 2 (Figure 2). There were no significant priming responses in the transition layer,
364 with consistent effects across all three pulses and DAP, although mean values generally
365 showed positive or no priming effects (Fig. 1, Supplemental Table 2).

366 Cellulose amendments resulted in positive and negative priming responses
367 depending on the soil layer, and in general had a longer, albeit smaller effect on soil C
368 loss than did glucose. In the surface soil, the 3rd cellulose pulse induced significant
369 negative priming that lasted the entire sampling period (Surface x Pulse 3: CI -49.41 to -
370 34.94 $\mu\text{g CO}_2\text{-C g}^{-1}\text{C}$, Fig. 3, Supplemental Table 2). In the permafrost layer, the 3rd
371 cellulose pulse led to significant positive priming that increased soil C loss (Permafrost x
372 Pulse 3: CI 7.66 to 28.67 $\mu\text{g CO}_2\text{-C g}^{-1}\text{C}$, Fig. 3, Supplemental Table 2). As with glucose
373 addition, there were no significant priming effects in the transition layer, but mean
374 responses trended towards positive or no priming effects (Fig. 3, Supplemental Table 2).

375 3.3. *Age of respired soil C*

376 In unamended soils, the $\Delta^{14}\text{C}$ value of respired C became more negative with depth,
377 indicating that microbes respired older soil C deeper in the soil profile (Fig. 4). Over the

378 course of the incubation, soil respired $\Delta^{14}\text{C}$ averaged 42‰ in the surface layer
379 (Intercept: CI 35‰ to 50‰, Table 2), -201‰ in the transition layer (Intercept: CI -296‰
380 to -110‰, Table 2), and -426‰ in the permafrost layer (Permafrost x DOI 15: CI -516‰
381 to -334‰, Table 2) (Figure 4). We observed the highest variability in soil-respired $\Delta^{14}\text{C}$ in
382 the permafrost horizon; the respired $\Delta^{14}\text{C}$ soil signature in the unamended permafrost
383 layer was significantly more positive on DOI 105 relative to DOI 15, (Permafrost x DOI
384 105: CI -420‰ to -231‰, Table 2), but by the end of the incubation, the soil-respired
385 $\Delta^{14}\text{C}$ was once again similar to DOI 15 (Permafrost x DOI 319: CI -497‰ to -307‰, Table
386 2).

387 Glucose treatments significantly increased the $\Delta^{14}\text{C}$ value of soil respired C by
388 115‰ (Glucose: CI 25‰ to 206‰, Table 2) in the transition and permafrost layers
389 relative to unamended soils in the transition layer (Fig. 4). This means that glucose
390 addition induced soil C mineralization of relatively young C in deeper layers. In the
391 surface layer we saw a similar pattern, soil respired $\Delta^{14}\text{C}$ values in the glucose treatment
392 were closer to 0‰, which indicates soil respired C was younger relative to unamended
393 soils; however, this difference was not statistically significant. Cellulose did not change
394 the $\Delta^{14}\text{C}$ of respired CO_2 relative to unamended soils.

395 3.4. Substrate use

396 Not all the substrate C added was respired by the end of each pulse. The amount
397 of glucose C respired, as a percent of added substrate 105 DAP, was 40-54% in the
398 surface, 37-58% in the transition, and 68-126% in the permafrost layer (Table 3). Out of
399 the total glucose C respired in the permafrost layer 105 DAP, 20-47% was mineralized in

400 the first week, when we observed significant positive priming effects (Table 4), we also
401 observed the highest CO₂ fluxes and $\delta^{13}\text{CO}_2$ in the first 7 DAP that often decreased to
402 match unamended concentrations 15 DAP (Supplemental Figures 1 and 2). The
403 cumulative amount of cellulose loss for pulses 2 and 3 ranged from 20-26% in the
404 surface, 16-25% in the transition, and 9-12% in the permafrost layer (Table 3). The
405 cellulose treatment did not elicit a high peak in CO₂ fluxes in the first 7 DAP
406 (Supplemental Figure 3), and $\delta^{13}\text{CO}_2$ did not decrease to match unamended values 105
407 days after pulses 2 and 3 (Supplemental Figure 4).

408 **4. Discussion**

409 Models forecast that 5-15% of the terrestrial permafrost C pool (~130-160 Pg C) is
410 vulnerable to decomposition in this century under current warming trajectories (Schuur
411 et al., 2015). Higher temperatures will also increase shrub expansion in tundra regions
412 (Chapin et al., 1995; Tape et al., 2006), and total belowground plant biomass, root
413 production, and rooting depth (Sullivan et al., 2007; Zamin et al., 2014; Keuper et al.,
414 2017), thus increasing fresh C inputs to soil. Our findings show that adding glucose to
415 permafrost soils can change the magnitude and age of respired soil C.

416 *4.1. Permafrost soil C loss increases with glucose addition*

417 Glucose additions caused a two-fold increase in permafrost soil C loss relative to
418 unamended samples in the first week after each pulse. The additional soil C mineralized
419 in the permafrost layer 7 days after each pulse was 0.71 to 1.20 mg C g⁻¹C, which is 1 to
420 almost 2 times greater than the initial microbial biomass in that layer (0.66 mg C g⁻¹ C).
421 This suggests that the increase in C loss resulted primarily from an increase in SOM

422 decomposition (real priming effect), rather than an accelerated turnover of the
423 microbial C pool (apparent priming effect) (Blagodatskaya and Kuzyakov, 2008;
424 Blagodatskaya et al., 2011). A previous priming incubation study in Siberian permafrost
425 conducted over 6 days also reported a two-fold increase in SOM decomposition in deep
426 mineral soils amended with glucose (Wild et al., 2014). De Baets et al. (2016) similarly
427 reported significant positive priming effects on a 10-day glucose addition study
428 conducted in tussock tundra mineral soil and permafrost. Our results corroborate the
429 hypothesis that the addition of an easily decomposable C substrate alleviates energy
430 constraints in deeper soil layers that have higher concentrations of slowly decomposing
431 C.

432 We hypothesized that amending deeper soil layers with a simple substrate would
433 increase soil C decomposition because slowly decomposing C constitutes most of the C
434 at depth. Our initial soil parameters show that the fraction of slowly-decomposing C
435 increased in the permafrost layer. We observed lower bulk soil $\delta^{13}\text{C}$ values at depth,
436 which are common for soils that are waterlogged, and can indicate the accumulation of
437 recalcitrant materials, like lignin (Alewell et al., 2011). The increased fraction of slowly-
438 decomposing C in deeper layers was also reflected in the nuclear magnetic resonance
439 spectra (NMR) analyses performed on subsamples of our soil cores. We found that the
440 proportion of alkyl C relative to *O*-alkyl C increased in the permafrost layer (Plaza et al.,
441 2017a). This pattern in the alkyl/*O*-alkyl ratio represents progressive organic matter
442 degradation, and decreased plant-derived fresh organic matter relative to microbial
443 biomolecules (Baldock and Skjemstad, 2000).

444 The negative priming effects in the surface layer that we observed after each
445 glucose pulse can be attributed to a switch from microbial decomposition of SOM to
446 decomposition of the added, more accessible substrate (Cheng, 1999; Kuzyakov, 2002).
447 These results are consistent with the theory that C availability is not limiting in highly
448 organic and less decomposed soils. Priming studies conducted in boreal, subarctic, and
449 permafrost regions that amended organic surface soils with glucose or low-molecular
450 weight C reported no net priming effects (Hartley et al., 2010; Lindén et al., 2014; Wild
451 et al., 2014; De Baets et al., 2016; Karhu et al., 2016; Rousk et al., 2016; Lynch et al.,
452 2018). Labile C addition to highly organic layers seem to shift microbial substrate
453 utilization to more N-rich compounds, which can reduce total SOM decomposition
454 (Rousk et al., 2016). Considering the entire soil profile, the 7-day cumulative response
455 ratio of respired soil C in the surface layer is very close to 1. The glucose effect on the
456 magnitude of soil C loss in the glucose treatment was negligible, and closely matched
457 rates in unamended surface samples. Meanwhile, positive priming in the permafrost
458 layer doubled the amount of C respired relative to permafrost unamended samples.

459 The strong negative priming response in surface soils during the 3rd cellulose pulse
460 may reflect the increasing C limitation of microorganisms during long-term incubations
461 (Bracho et al., 2016). Since 74-92% of the added cellulose remained in the soil after each
462 pulse, and $\delta^{13}\text{CO}_2\text{-C}$ was still high 105 DAP, it is possible that microbes relied on the
463 accumulated cellulose as a source of C instead of SOM after almost one year of
464 incubation.

465 In the permafrost layer; the stimulation of daily CO₂-C production (7.66 to 28.67 µg
466 CO₂-C g⁻¹C) caused by the 3rd cellulose pulse was small compared to the initial microbial
467 biomass in the permafrost layer (660 ± 120 µg microbial C g⁻¹C). Thus, we cannot fully
468 attribute this positive priming effect in the cellulose treatment to higher SOM
469 decomposition, but potentially to an increase in microbial turnover (i.e. apparent
470 priming effect) (Jenkinson et al., 1985; Blagodatskaya and Kuzyakov, 2008). If we
471 consider the effect of cellulose addition on the entire soil profile, negative priming
472 effects in the surface were 2-5 times higher than positive priming effects in the
473 permafrost.

474 4.2. *Persistent SOM decomposition response to multiple glucose additions*

475 Our results show that over the course of the incubation, each pulse of glucose
476 elicited a priming response. In the permafrost layer, glucose addition consistently
477 doubled soil C loss (Figure 2). Priming studies that repeatedly amended soils with
478 multiple pulses of substrate C have reported higher soil C losses than single additions at
479 the beginning of an incubation (De Nobili et al., 2001; Hamer and Marschner, 2005).

480 Glucose was readily used by microbes in the permafrost layer, and 68-126% of the
481 added substrate was completely respired 152 days following each pulse. Out of the
482 substrate respired, 20-47% was respired just in the first week, which is when we
483 observed positive priming effects. Glucose often produces a rapid response in microbial
484 activity (Bernal et al., 2016), and leads to rapid metabolic changes in a wide variety of
485 fast-growing bacteria that utilize it as a substrate (Stotzky and Norman, 1961; Hungate
486 et al., 2015). This fast turnover of glucose in permafrost soil, which can be seen in the

487 rapid decline in CO₂ flux and $\delta^{13}\text{CO}_2\text{-C}$ (Supplemental Figures 1 and 2), and the sustained
488 positive priming effects over the course of the incubation suggest that multiple pulses
489 will continue to stimulate soil C loss, and single-pulse incubation studies might
490 underestimate soil C losses.

491 4.3. *Age of respired soil C decreases with glucose input*

492 Carbon in deep soil layers is mostly old, and its longer mean residence time is due
493 to different biological, physical, and chemical stabilization processes; in permafrost, soil
494 C persistence may also be promoted by low rates of DOC inputs (Schimel et al., 1994;
495 Schuur et al., 2009; Czimczik and Welker, 2010; Walvoord and Kurylyk, 2016). Though a
496 previous study found that soluble C inputs increased the proportion of older CO₂-C
497 fluxes in boreal soils (Lindén et al., 2014), our results indicate that an input of glucose
498 decreased the age of soil respired C in transition and permafrost layers. Glucose
499 amendment induced soil C mineralization that was overall 115% higher (younger)
500 relative to soil C respired in unamended and cellulose treatments. Though other
501 permafrost priming studies do not report radiocarbon age of respired soil C, two priming
502 studies in other ecosystems also reported younger C losses. Blagodatskaya et al. (2011)
503 found that primed SOM decomposition originated mainly from younger C, even though
504 the proportion of young and old C equally contributed to SOM. Sullivan and Hart (2013)
505 conducted a priming experiment at four sites along a substrate age gradient from 0.93-
506 3000 ky, and found the greatest cumulative priming in younger sites, while the oldest
507 site experienced either negative or no net priming .

508 Recent arguments have been raised against the long-standing theory that
509 persistent soil C is found in large humic substances, but rather in a continuum of
510 progressively more decomposed compounds that transition from polymers to
511 monomers as OM is oxidized (Lehmann and Kleber, 2015). Older C fractions are not
512 always composed of complex or recalcitrant compounds (Kleber et al., 2011; Nunan et
513 al., 2015). Observations in a long-term bare-fallow soil that was depleted of plant litter
514 inputs showed that microbial communities became adapted to metabolizing simple and
515 small compounds. Microbes tended to use molecules or derivatives from the Krebs cycle
516 (i.e. α -ketoglutaric acid, citric acid), and were less able to decompose polymers, and
517 although old C molecules can be readily mineralized, the energetic payoff is low (Nunan
518 et al., 2015). The microbial mining hypothesis suggests that SOM decomposition
519 increases when microbes use extracellular enzymes to release nutrients locked in
520 polymers (Craine et al., 2007; Fontaine et al., 2011; Dijkstra et al., 2013). When a fresh C
521 substrate is introduced that increases microbial activity and extracellular enzyme
522 production, microbes can access C and nutrients in more complex polymeric structures.
523 If these complex C structures are younger, then positive priming may increase the
524 proportion of younger C that is respired. Soil C stabilization mechanisms in permafrost
525 are not completely understood, and in general, permafrost C is considered stable
526 primarily due to low temperature constraints on decomposition. Once temperature
527 constraints are lifted, mechanisms like waterlogging, microaggregation, and association
528 with silt and clay particles may also contribute to C stabilization (Six et al., 2002). Future

529 priming studies in permafrost should focus on physical stabilization processes that
530 influence C turnover rates.

531 A previous study at Eight Mile Lake found that the proportion of old C respired from
532 deeply thawed, experimentally warmed plots, was lower relative to less thawed areas.
533 They attributed this phenomenon to higher plant respiration, which increased the
534 contribution of younger C to ecosystem respiration and diluted the old C signature
535 (Schuur et al., 2009; Hicks Pries et al., 2015). Our results suggest that positive priming in
536 deep soil layers can also contribute to a younger signature of ecosystem respiration. The
537 average $\Delta^{14}\text{C}$ in unamended samples was -201‰ (CI: -296 to -110‰) in the transition
538 layer and -426‰ (CI: -516 to -334‰) in the permafrost layer. Priming increased the
539 respired $\Delta^{14}\text{C}$ values to -86‰ in the transition layer and -311‰ in the permafrost layer.
540 Though these values still reflect relatively older C, the change corresponds to potential
541 $\Delta^{14}\text{CO}_2$ that could be measured in layers that are 10-30 cm shallower than the depths
542 they originated from. Applying a partitioning model without considering priming effects
543 could cause the model to underestimate deep soil C contribution to ecosystem
544 respiration by attributing them to decomposition of shallower soils that have a greater
545 proportion of younger C.

546 4.4. *Implications for permafrost-C feedback*

547 Our findings indicate that priming effects can have implications for the magnitude
548 and turnover of soil C in permafrost. Priming doubled permafrost soil C losses in the first
549 week after each pulse, and decreased the overall age of respired soil C. Field ecosystem
550 respiration measurements show that moist acidic tundra sites at Eight Mile Lake are a

551 net annual CO₂-C source to the atmosphere, despite increases in plant biomass (Mauritz
552 et al., 2017). This effect could be exacerbated by increased inputs of easily
553 decomposable C to mineral and permafrost soils. We extrapolated our incubation
554 results to field conditions at Eight Mile Lake by using field soil temperatures to
555 temperature-correct soil C fluxes, and applied a site specific Q₁₀ of 2.6 based on the
556 temperature sensitivity analysis of CO₂ fluxes from soils collected in our site in 2010
557 (Bracho et al., 2016). Our estimates suggest that priming-induced soil C losses
558 correspond to 4-12% of the cumulative C released during a growing season (235 - 408 g
559 CO₂-C m⁻², reported in Mauritz et al. 2017). Though we must be cautious when relating
560 incubation results to the field, as these estimates are likely high because soils can
561 remain waterlogged after thaw, and our sample size is small, our study suggests that
562 priming may be an important mechanism that exacerbates soil C losses to the
563 atmosphere. Future priming studies should focus on field studies, and measure the
564 radiocarbon age of primed soil C. We must consider priming effects to fully understand
565 permafrost C dynamics, or we may potentially underestimate the contribution of soil C
566 to ecosystem respiration rates.

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865

866 Table 1. Initial soil properties (mean \pm SE) grouped by soil layers. A linear mixed effects model tested the effect of soil layer on bulk density,
 867 initial bulk soil organic carbon and nitrogen (%), C:N, initial $\delta^{13}\text{C}$ (‰), microbial biomass, and pH. Values not sharing the same letter indicate a
 868 significant difference.

Layer	Bulk Density (g/cm ³)	Initial C (%)	Initial N (%)	C:N	Initial $\delta^{13}\text{C}$ (‰)	Microbial biomass (mg C g ⁻¹ soil C)	pH
Surface	0.17 \pm 0.04 ^a	37.96 \pm 2.52 ^a	1.51 \pm 0.11 ^a	25.52 \pm 2.59 ^a	-26.22 \pm 0.17 ^a	1.22 \pm 0.76 ^a	3.96 \pm 0.09 ^a
Transition	0.53 \pm 0.15 ^b	20.99 \pm 3.92 ^b	0.86 \pm 0.20 ^b	25.08 \pm 1.88 ^a	-27.22 \pm 0.19 ^b	0.36 \pm 0.13 ^a	4.77 \pm 0.13 ^b
Permafrost	0.46 \pm 0.03 ^b	9.69 \pm 3.68 ^c	0.37 \pm 0.13 ^c	24.80 \pm 1.50 ^a	-27.36 \pm 0.23 ^b	0.66 \pm 0.12 ^a	5.12 \pm 0.15 ^c

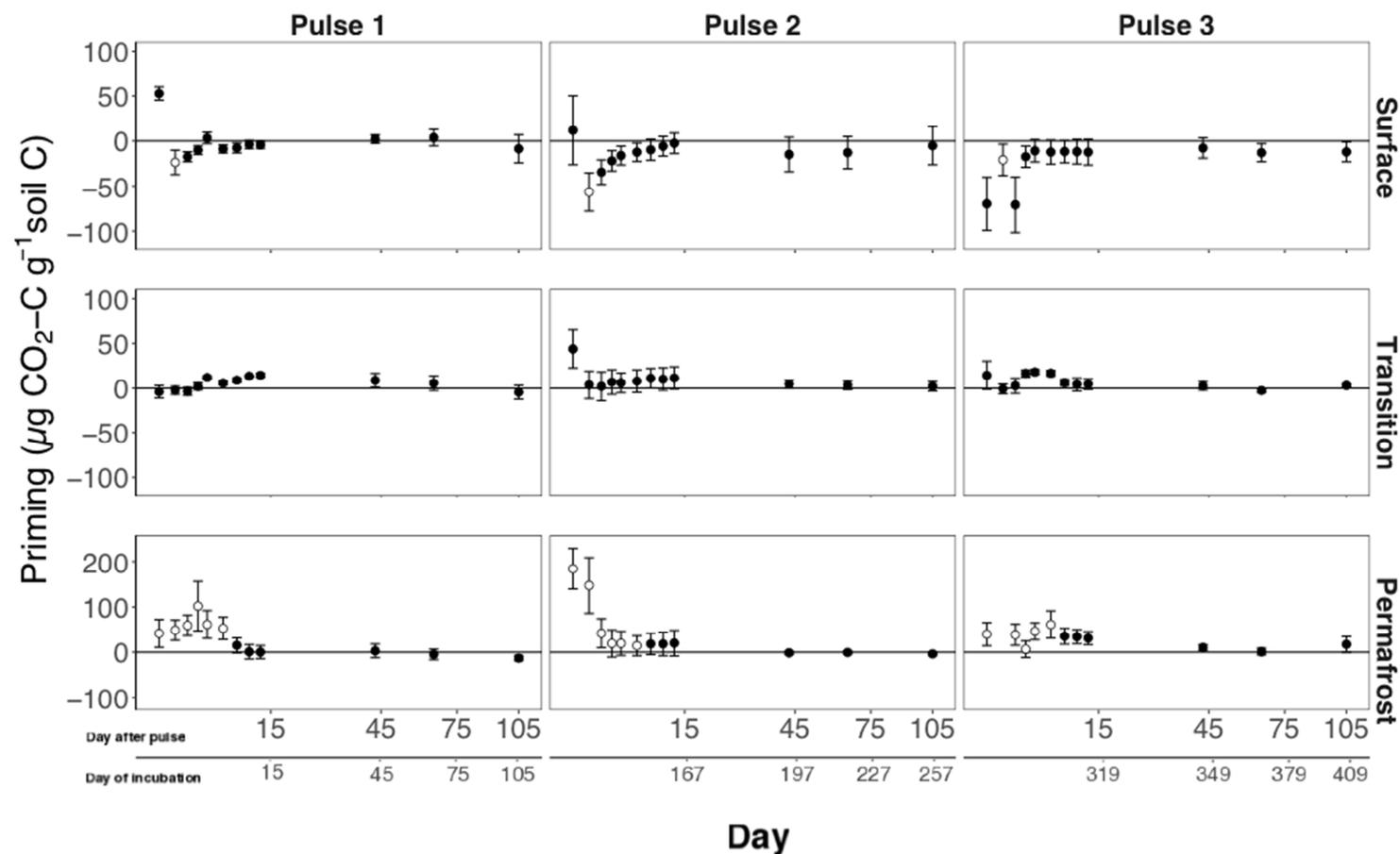
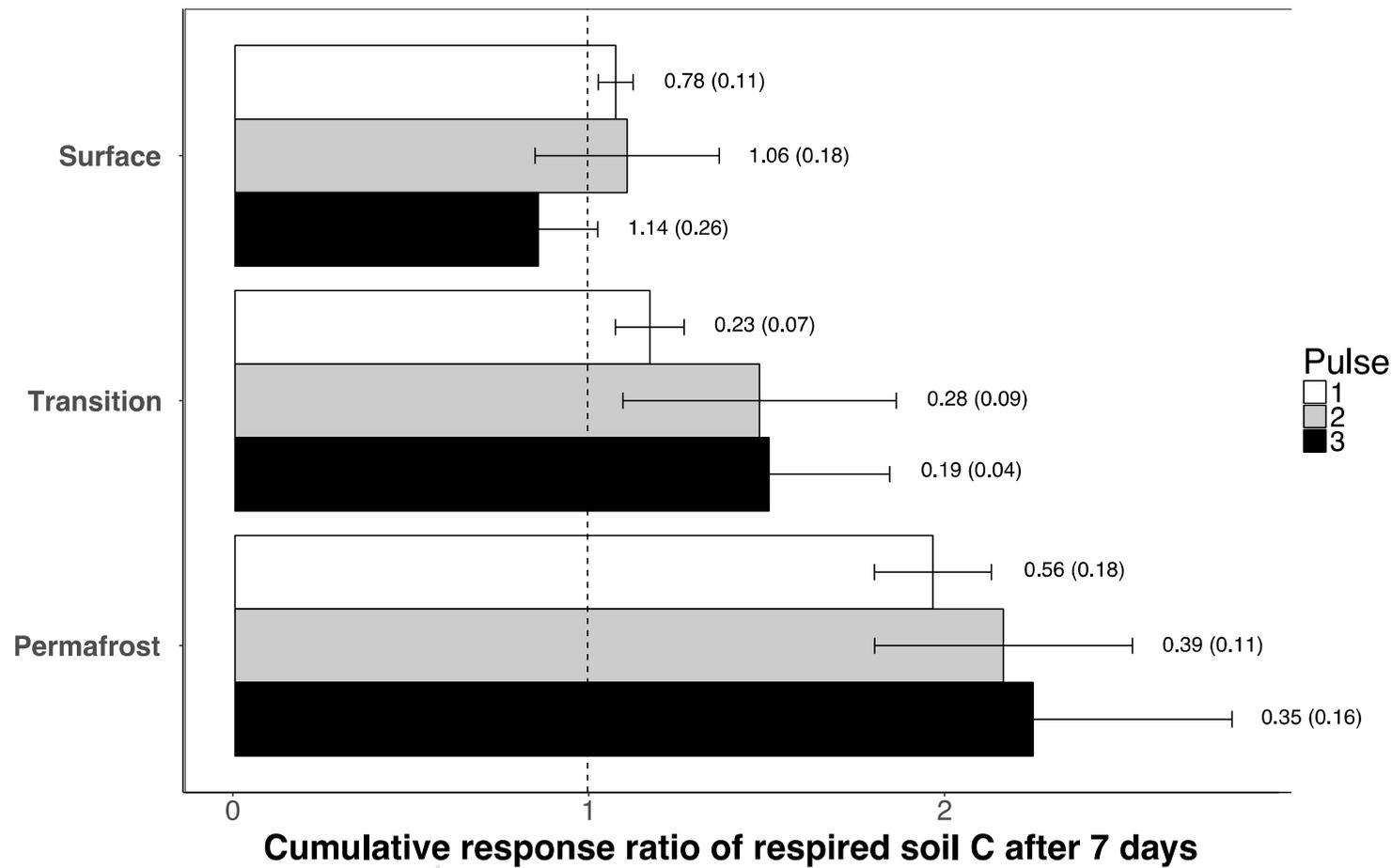


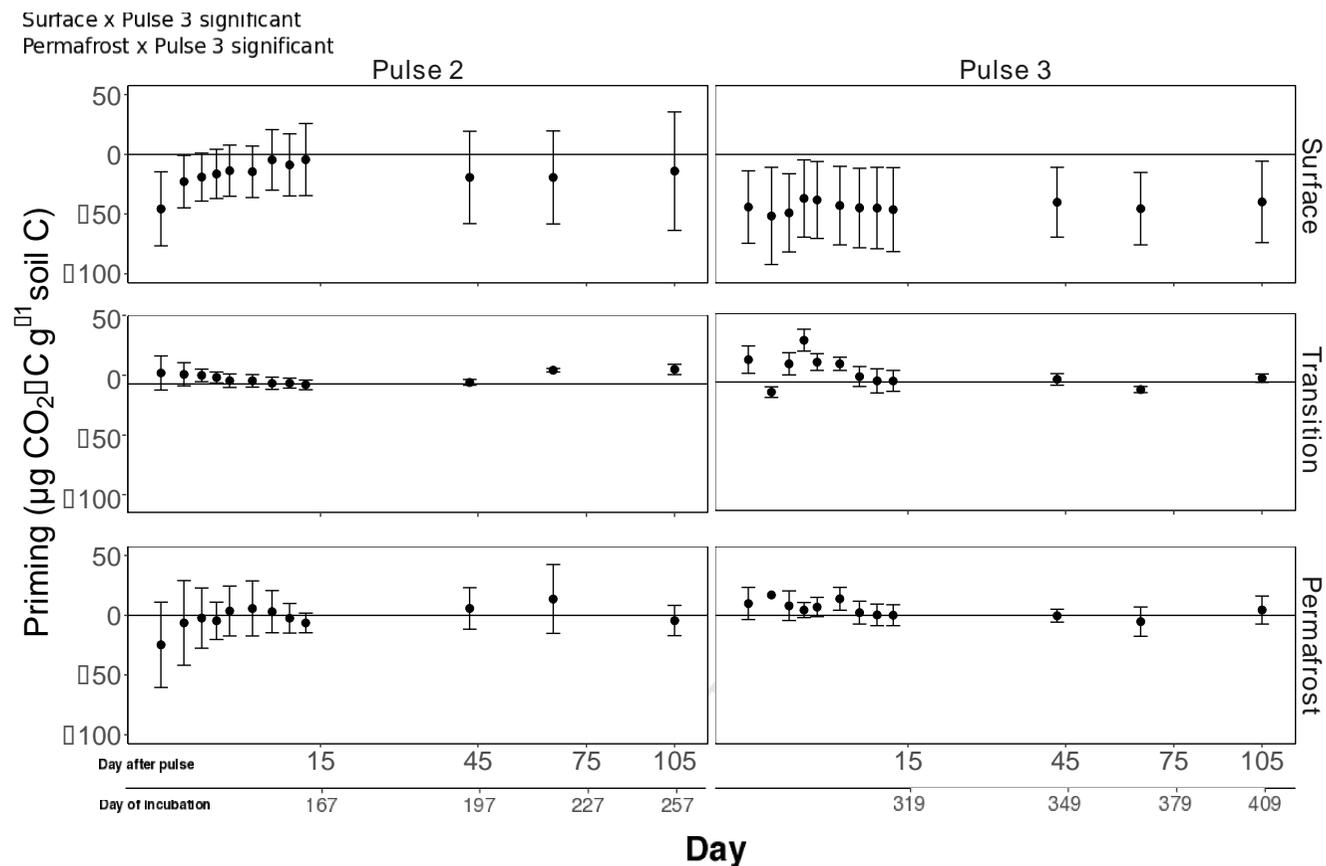
Figure 1. Priming

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 870 ($\mu\text{g CO}_2\text{-C g}^{-1}\text{C}$) in surface, transition, and permafrost layers over time (in days) after glucose pulse amendment (DAP) (primary x-axis). The
 871 secondary x-axis indicates day since the start of incubation (DOI). The x-axis was square-root transformed to make it easier to see values in the
 872 first 15 DAP. Values above zero indicate higher soil C losses relative to control (positive priming), and values below zero indicate lower soil C
 873 losses relative to control (negative priming). Bars are one standard error from the mean, and open circles indicate a significant priming effect.
 874 Note that the scale for the y-axis in the permafrost layer is two times higher.



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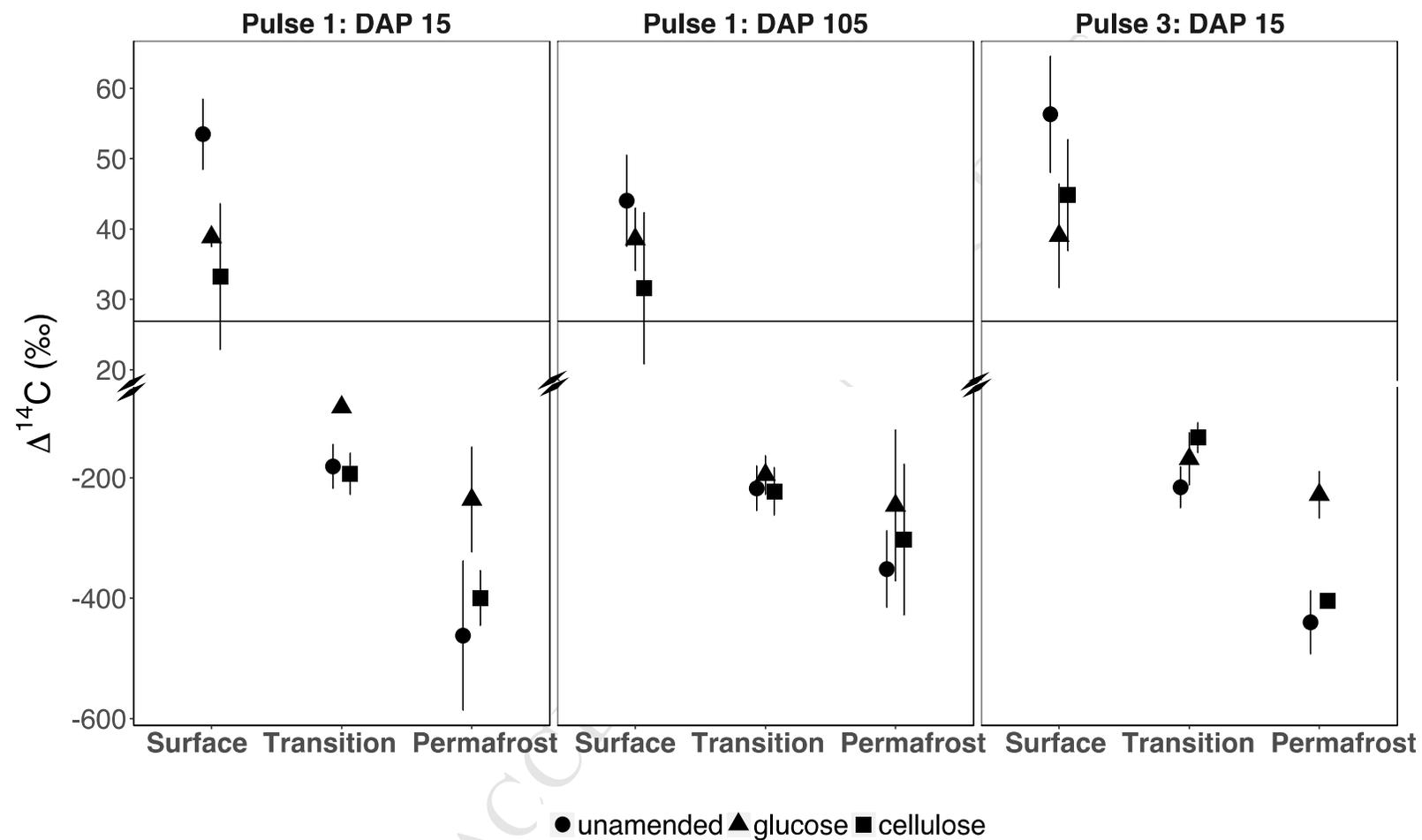
Figure 2. Cumulative response ratio of respired soil C in glucose treatments relative to respired soil C in unamended samples over 7 DAP. Glucose to unamended ratios that are less than 1 indicate negative priming, equal to 1 indicate no priming effect, greater than 1 indicate positive priming. Numbers in the figure are the unamended cumulative soil C losses ($\text{mg soil C g}^{-1}\text{C} \pm \text{SE}$) 7 days after each pulse. Bars are one standard error from the mean.



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Figure 3. Priming ($\mu\text{g CO}_2\text{-C g}^{-1}\text{C}$) in surface, transition, and permafrost layers over time (in days) after cellulose pulse amendment (DAP) (primary x-axis). The secondary x-axis indicates day since the start of incubation (DOI). The x-axis was square-root transformed to make it easier to see values in the first 15 DAP. Labelling was not high enough in Pulse 1 to calculate priming effects (data not shown). Values above zero indicate higher soil C losses relative to control (positive priming), and values below zero indicate lower soil C losses relative to control (negative priming). Bars are one standard error from the mean. Surface and permafrost layers had significantly negative and positive priming, respectively, in Pulse 3.

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Figure 4. Mean soil respired $\Delta^{14}\text{C}$ (‰) in surface, transition, and permafrost layers by treatment 15 and 105 days after pulse 1, and 15 days after pulse 3 (DOI 319). Bars are one standard error from the mean. Horizontal line indicates atmospheric $\Delta^{14}\text{C}$ concentration in 2013. Note the break in the y-axis.

893 Table 2. Mixed linear effects model for soil respired $\Delta^{14}\text{C}$ at 15 and 105 days after the 1st amendment pulse, and DOI 319 (15 days after the 3rd
 894 amendment pulse). The intercept for the surface model represents soil respired $\Delta^{14}\text{C}$ (‰) in unamended surface layer, 15 days after pulse 1. The
 895 intercept for the deep layers (transition and permafrost) model represents soil respired $\Delta^{14}\text{C}$ (‰) in unamended transition layer, 15 days after
 896 pulse 1. Coefficients represent effect size on intercept, and significant effects are bolded.
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Response variable	Full model	Final Model	Variable	Coefficient	Min CI	Max CI
Soil respired $\Delta^{14}\text{C}$ (‰)	Treatment x DOI	Intercept	Intercept (Unamended, Surface, DOI 15)	42.4	34.8	50.1
Soil respired $\Delta^{14}\text{C}$ (‰)	Treatment x Layer x DOI	Treatment	Intercept (Unamended, Transition, DOI 15)	-200.9	-295.9	-109.7
		Layer	Glucose	115.4	25.1	206.3
		DOI	Cellulose	30.4	-55.5	118.2
		Layer x DOI	DOI 105	-59.5	-121.0	1.4
			DOI 319	-20.1	-76.8	36.8
			Permafrost x DOI 15	-224.8	-314.8	-133.3
			Permafrost x DOI 105	158.5	65.3	253.9
			Permafrost x DOI 319	44.4	-51.7	138.5

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899 Table 3. Cumulative substrate C loss as a percent ($\% \pm \text{SE}$) of added substrate (3.5 mg substrate C per g initial C) 105 days after each amendment
 900 pulse.

	<i>Pulse 1</i>		<i>Pulse 2</i>		<i>Pulse 3</i>	
	Glucose	Cellulose	Glucose	Cellulose	Glucose	Cellulose
Surface	54.4 \pm 3.9	NA	39.9 \pm 2.4	20.0 \pm 2.4	45.9 \pm 6.8	25.9 \pm 2.0
Transition	58.3 \pm 15.0	NA	37.8 \pm 7.6	15.6 \pm 6.8	36.9 \pm 11.7	24.5 \pm 5.5
Permafrost	125.9 \pm 49.0	NA	67.5 \pm 32.2	9.4 \pm 2.3	80.2 \pm 39.8	11.8 \pm 5.8

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903 Table 4. Seven-day cumulative substrate C loss as a percent ($\% \pm \text{SE}$) of respired substrate 105 days after each amendment pulse

	<i>Pulse 1</i>		<i>Pulse 2</i>		<i>Pulse 3</i>	
	Glucose	Cellulose	Glucose	Cellulose	Glucose	Cellulose
Surface	37.9 \pm 2.1	NA	40.7 \pm 1.8	5.6 \pm 0.7	43.4 \pm 4.2	6.9 \pm 0.5
Transition	13.8 \pm 2.4	NA	29.0 \pm 5.6	5.6 \pm 1.1	24.2 \pm 1.9	4.8 \pm 0.5
Permafrost	46.6 \pm 3.9	NA	47.4 \pm 3.7	6.0 \pm 1.7	20.0 \pm 1.7	5.2 \pm 0.9

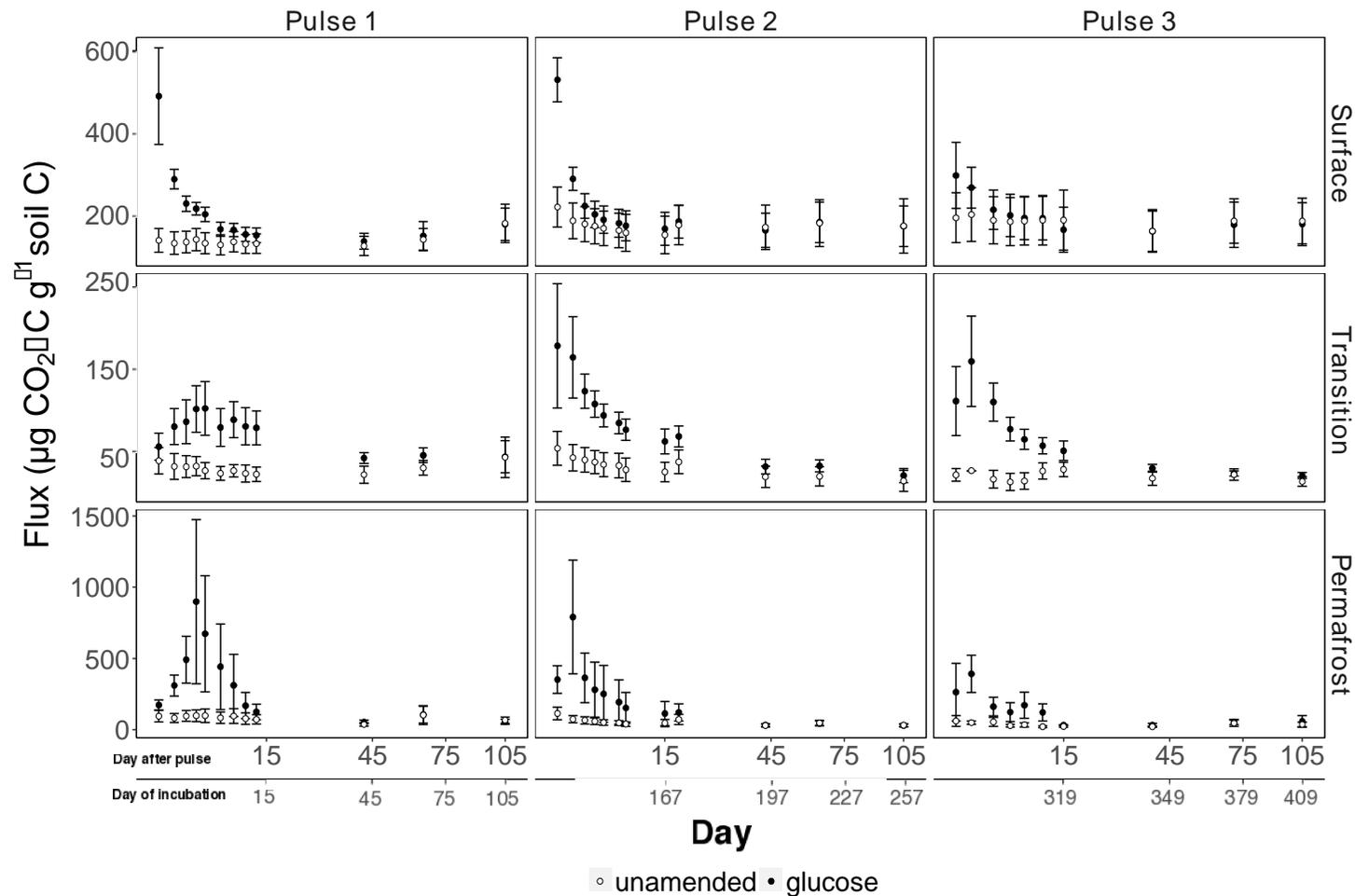
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906 Supplementary Materials

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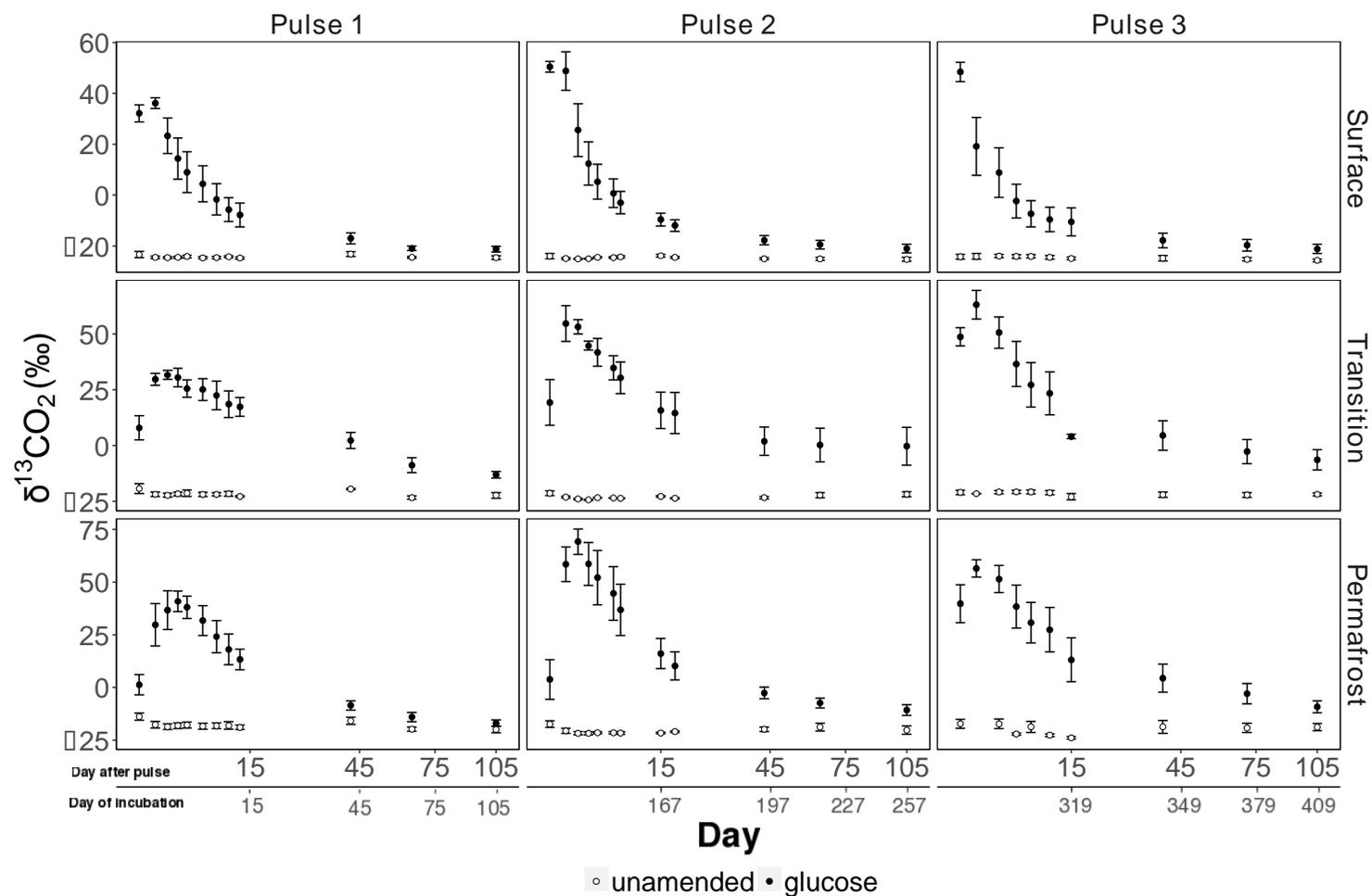
908 Table 1. Glucose and cellulose $\delta^{13}\text{C}$ values (‰) by pulse and round. These $\delta^{13}\text{C}_{\text{substrate}}$ values were used as end-members in the isotopic mass
909 balance equation to calculate priming (Equation 1).

	<i>Pulse 1</i>		<i>Pulse 2</i>		<i>Pulse 3</i>	
	Round 1	Round 2	Round 1	Round 2	Round 1	Round 2
Glucose	73.4	68.9	115.2	98.6	92.0	113.5
Cellulose	8.8	-21.2	76.4	234.1	494.9	494.9



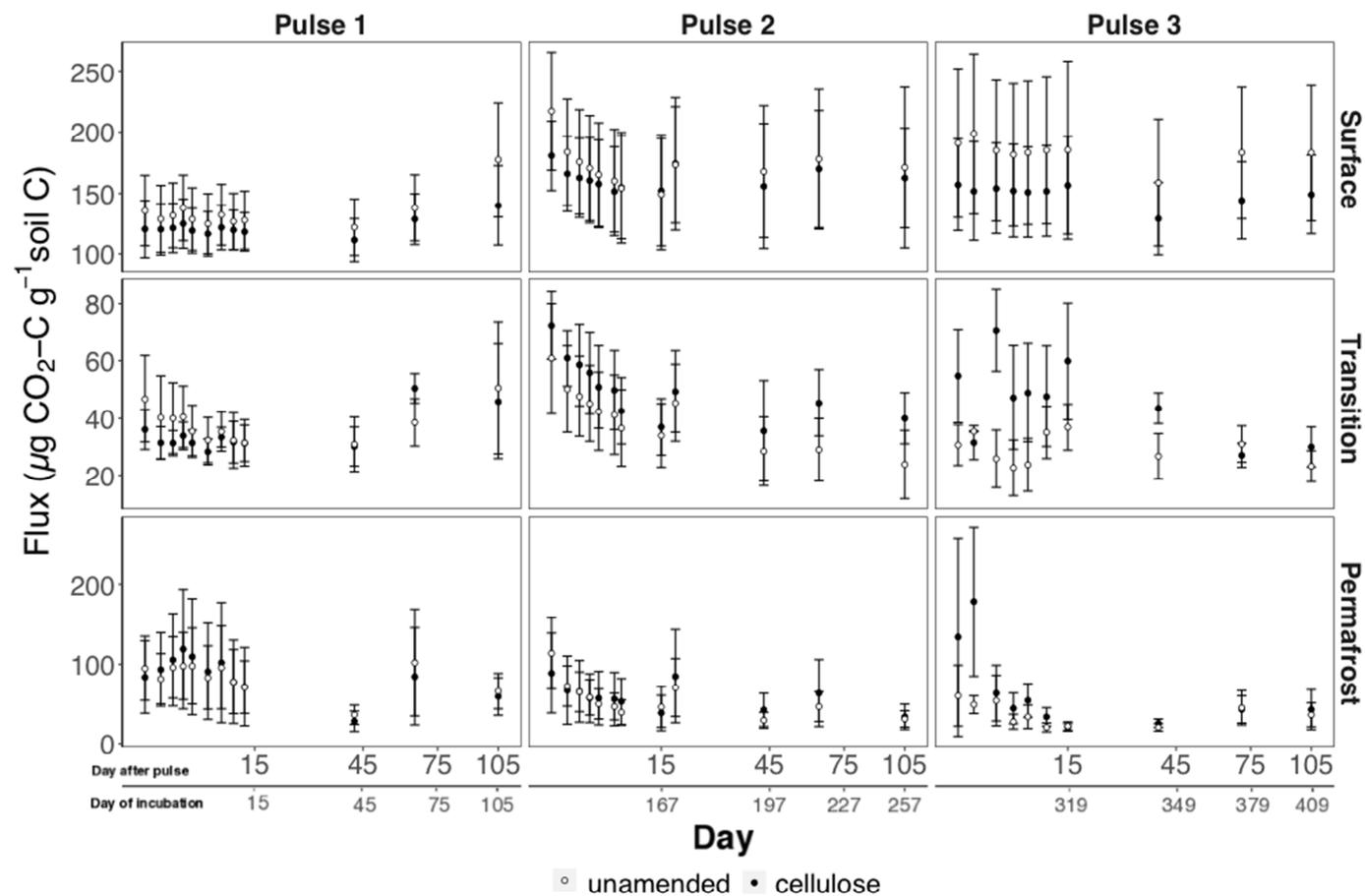
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Figure 1. Total respired C ($\mu\text{g CO}_2\text{-C g}^{-1}\text{C}$) in unamended and glucose treatments in surface, transition, and permafrost layers over time (in days) after glucose pulse amendment (DAP) (primary x-axis). The secondary x-axis indicates day since the start of incubation (DOI). The x-axis was square-root transformed to make it easier to see values in the first 15 DAP. Bars are one standard error from the mean. Note that the scale for the y-axis is different for each layer.



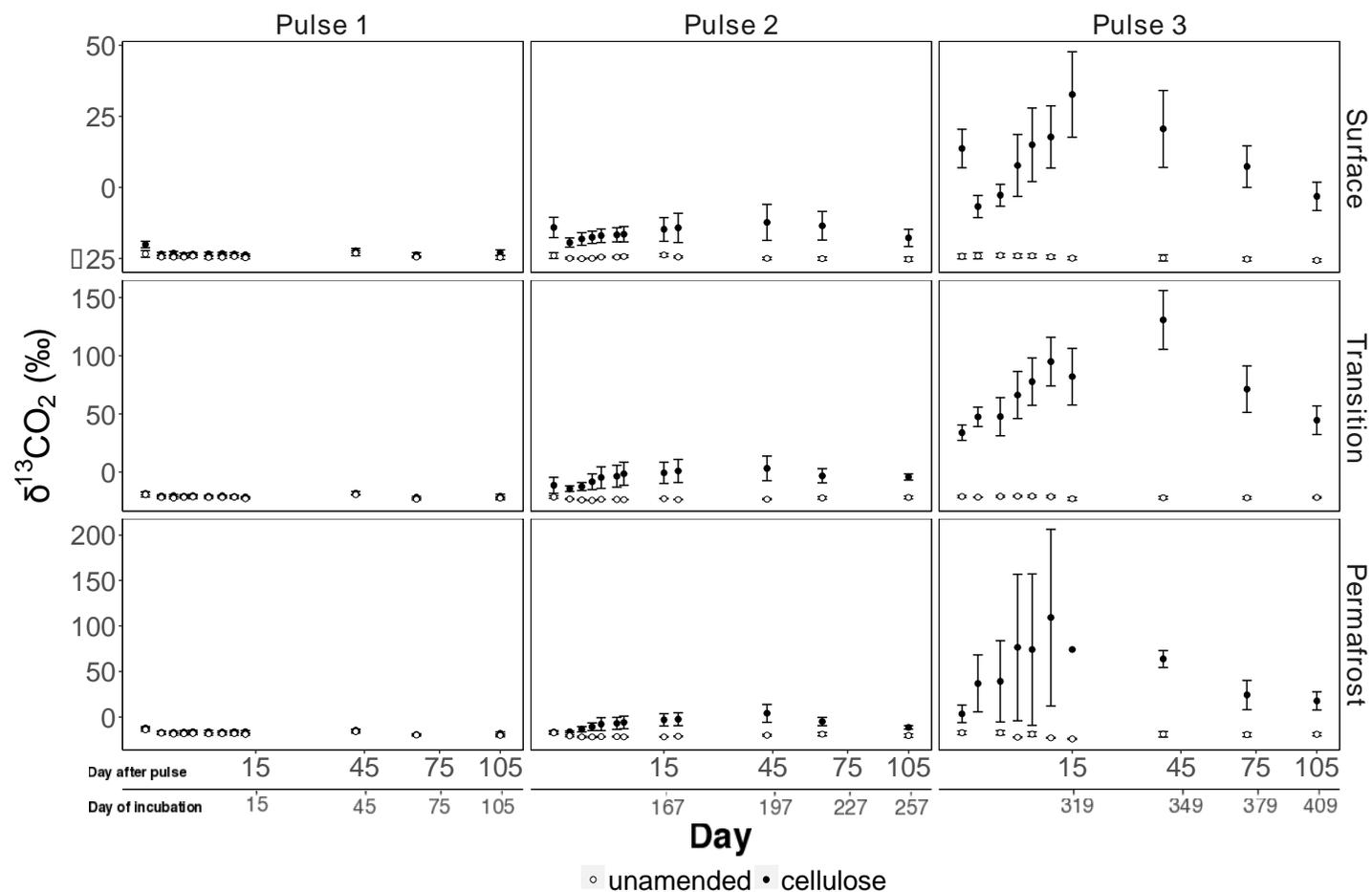
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Figure 2. Total respired $\delta^{13}\text{CO}_2\text{-C}$ (‰) in unamended and glucose treatments in surface, transition, and permafrost layers over time (in days) after each glucose pulse amendment (DAP) (primary x-axis). The secondary x-axis indicates day since the start of incubation (DOI). The x-axis was square-root transformed to make it easier to see values in the first 15 DAP. Bars are one standard error from the mean. Note that the scale for the y-axis is different for each layer.



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 921 Figure 3. Total respired C ($\mu\text{g CO}_2\text{-C g}^{-1}\text{C}$) in unamended and cellulose treatments in surface, transition, and permafrost layers over time (in days)
 922 after each cellulose pulse amendment (DAP) (primary x-axis). The secondary x-axis indicates day since the start of incubation (DOI). The x-axis
 923 was square-root transformed to make it easier to see values in the first 15 DAP. Bars are one standard error from the mean. Note that the scale
 924 for the y-axis is different for each layer.

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Figure 4. Total respired $\delta^{13}\text{CO}_2\text{-C}$ (%) in unamended and cellulose treatments in surface, transition, and permafrost layers over time (in days) after each cellulose pulse amendment (DAP) (primary x-axis). The secondary x-axis indicates day since the start of incubation (DOI). The x-axis was square-root transformed to make it easier to see values in the first 15 DAP. Bars are one standard error from the mean. Note that the scale for the y-axis is different for each layer.

931 Table 2. Mixed linear effects model parameters for glucose and cellulose priming effects following each amendment pulse. Coefficients
 932 represent daily priming values in micrograms CO₂-C per gram soil C, calculated from the effect size on the intercept. The intercept for the
 933 glucose model is: unamended surface soil, at 1 DAP, Pulse 1. The intercept for the cellulose is: unamended surface soil, at 1 DAP, Pulse 2. If the
 934 Min and Max CI do not overlap zero, it indicated a significant priming response (bolded).

Response variable	Full model	Final Model	Variable	Coefficient	Min CI	Max CI
Glucose priming ($\mu\text{g CO}_2\text{-C g}^{-1}\text{C}$)	Pulse x DAP x Layer	Pulse	Intercept (Surface, 1 DAP, Pulse 1)	5.72	-22.81	35.11
		DAP	Surface x 2 DAP	-42.55	-77.36	-9.09
		Layer	Surface x 3 DAP	-17.06	-52.17	17.95
		Layer x DAP	Surface x 4 DAP	14.65	-19.95	49.98
		DAP x Pulse	Surface x 5 DAP	13.08	-21.63	46.59
		Surface x 7 DAP	1.43	-35.14	37.75	
		Surface x 9 DAP	-2.47	-36.48	31.33	
		Surface x 11 DAP	-1.58	-37.7	34.03	
		Surface x 13 DAP	-1.38	-37.29	33.13	
		Surface x 43 DAP	7.04	-27.38	40.73	
		Surface x 65 DAP	5.73	-29.03	41.46	
		Surface x 105 DAP	-5.28	-40.28	29.68	
		Transition x 1 DAP	21.97	-19.71	63.46	
		Transition x 2 DAP	-9.57	-55.91	37.76	
		Transition x 3 DAP	16.71	-30.07	63.26	
		Transition x 4 DAP	36.23	-8.14	81.07	
		Transition x 5 DAP	29.37	-14.72	73.26	
		Transition x 7 DAP	19.01	-25.73	64.79	
		Transition x 9 DAP	11.7	-32.26	55.02	
		Transition x 11 DAP	9.54	-34.46	52.97	
Transition x 13 DAP	10	-33.56	56.6			
Transition x 43 DAP	14.03	-30.81	59.23			

Transition x 65 DAP	10.11	-34.3	54.93
Transition x 105 DAP	-1.91	-44.87	43.74
Permafrost x 1 DAP	91.32	58.46	123.17
Permafrost x 2 DAP	86.82	47.35	126.75
Permafrost x 3 DAP	60.71	24.09	97.09
Permafrost x 4 DAP	67.59	31.23	102.93
Permafrost x 5 DAP	58.44	22.32	95.47
Permafrost x 7 DAP	64.78	29.39	101.69
Permafrost x 9 DAP	24.3	-12.62	61.34
Permafrost x 11 DAP	35.9	-0.6	73.03
Permafrost x 13 DAP	15.3	-21.07	54.35
Permafrost x 43 DAP	11.36	-25.44	48.13
Permafrost x 65 DAP	6.53	-29.06	42
Permafrost x 105 DAP	-1.99	-38.56	34.79
1 DAP x Pulse 2	55.88	26.56	84.08
1 DAP x Pulse 3	-41.04	-69.06	-12.55
2 DAP x Pulse 2	-18.44	-58.19	21.76
2 DAP x Pulse 3	-22.35	-75.37	28.72
3 DAP x Pulse 2	-28.33	-68.56	12.56
3 DAP x Pulse 3	-40.58	-83.53	0.74
4 DAP x Pulse 2	-18.96	-59.58	22.15
4 DAP x Pulse 3	-19.86	-61.09	20.33
5 DAP x Pulse 2	-12.65	-51.59	28.02
5 DAP x Pulse 3	1.96	-38.03	42.87
7 DAP x Pulse 2	-14.95	-53.87	24.24
7 DAP x Pulse 3	6.44	-36.06	46.97
9 DAP x Pulse 2	0.08	-39.62	40.83

			9 DAP x Pulse 3	1.36	-39.49	41.22
			11 DAP x Pulse 2	5.8	-34.07	47.4
			11 DAP x Pulse 3	5.47	-35.19	45.21
			13 DAP x Pulse 2	8.39	-30.88	48.36
			13 DAP x Pulse 3	4.57	-35.24	43.76
			43 DAP x Pulse 2	-1.06	-39.76	40.23
			43 DAP x Pulse 3	4.07	-36.41	45.1
			65 DAP x Pulse 2	1.81	-38.27	41.99
			65 DAP x Pulse 3	0.79	-40.75	40.81
			105 DAP x Pulse 2	2.72	-38.06	42.82
			105 DAP x Pulse 3	8.25	-33.1	49.37
Cellulose priming ($\mu\text{g CO}_2\text{-C g}^{-1}\text{C}$)	Pulse x DAP x Layer	Pulse	Intercept (Surface, Pulse 2)	-16.87	-46.95	12.78
		Layer	Surface x Pulse 3	-42.54	-49.41	-34.94
		Pulse x Layer	Transition x Pulse 2	4.97	-43.03	51.4
			Transition x Pulse 3	8.85	-0.02	24.77
			Permafrost x Pulse 2	-1.7	-38.22	35.93
			Permafrost x Pulse 3	8.29	7.66	28.67

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937 Table 3. Average respired $\delta^{13}\text{C}$ (\pm SE) in unamended soils for the entire duration of the incubation experiment. Values not sharing the same
938 letter indicate a significant difference.

	$\delta^{13}\text{CO}_2$ (‰)	
Surface	-24.47 ± 0.07^a	
Transition	-21.99 ± 0.17^b	
Permafrost	-18.95 ± 0.20^c	

Title: Glucose addition increases the magnitude and decreases the age of soil respired carbon in a long-term permafrost incubation study

Authors: Elaine Pegoraro^{a,b}, Marguerite Mauritz^{a,b}, Rosvel Bracho^c, Chris Ebert^{a,b}, Paul Dijkstra^{a,b}, Bruce A. Hungate^{a,b}, Kostas T. Konstantinidis^d, Yiqi Luo^{a,b,e,g}, Christina Schädel^{a,b}, James M. Tiedje^f, Jizhong Zhou^g, Edward A.G. Schuur^{a,b}.

Highlights:

Glucose addition increased permafrost soil C loss two-fold.

Glucose addition to surface soils did not elicit a priming response.

Cellulose addition to surface soils elicited a negative priming response.

Glucose addition decreased the age of soil-respired carbon in deep layers.