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Similar temperature sensitivity of soil mineral-associated organic carbon regardless of age

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-Temperature sensitivity of mineral-associated SOM was not related to its age

-Temperature sensitivity of POM fraction was similar or slightly higher than MOM

-Within the POM fraction, old C was more sensitive to temperature than recent C

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1 Similar temperature sensitivity of soil mineral-associated organic carbon

2 regardless of age

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30 Abstract

31 Most of the carbon (C) stored in temperate arable soils is present in organic matter (OM) intimately 32 associated with soil minerals and with slow turnover rates. The sensitivity of mineral-associated OM to 33 changes in temperature is crucial for reliable predictions of the response of soil C turnover to global 34 warming and the associated flux of carbon dioxide (CO_2) from the soil to the atmosphere. We studied the 35 temperature sensitivity of C in <63 µm fractions rich in mineral-associated organic matter (MOM) and of 36 C in >63 µm fractions rich in particulate organic matter (POM). The fractions were isolated by physical 37 separation of two light-textured arable soils where the C4-plant silage maize had replaced C3-crops 25 38 years ago. Differences in ¹³C abundance allowed for calculation of the age of C in the soil-size fractions 39 (old C, C3-C > 25 years; recent C, C4-C < 25 years). We incubated bulk soils (< 2 mm) and size 40 fractions sequentially at 6, 18, 26 and 34 °C (ramping up and down the temperature scale) and calculated the temperature sensitivity of old and recent C from ¹²CO₂ and ¹³CO₂ evolution rates. The 41 42 temperature sensitivity was similar or slightly higher for POM than for MOM. Within the POM fraction, old 43 C3-C was more sensitive to changes in temperature than recent C4-C. For the MOM fraction, the 44 temperature sensitivity was unrelated to the age of C. Quantitative PCR analysis indicated that the proportions of bacteria, archaea and fungi did not change during incubation. Our results suggest that 45 46 while OM stabilizing mechanisms affect the temperature sensitivity of soil C, temperature sensitivity 47 appears unrelated to the age of mineral-associated OM.

48

Keywords: Soil organic matter, temperature sensitivity, decomposition, climate change, ¹³C natural
 abundance, Bayesian statistics

51

52 **1. Introduction**

53 Understanding the mechanisms that affect the turnover of organic matter (OM) in soil is crucial in order 54 to predict the response of soil C storage to changes in climate and land use. The effect of changes in 55 temperature on the decomposition of labile soil OM is well known, whereas less is known on the 56 temperature effect on C in soil OM pools with turnover rates counted in decades and centuries 57 (Trumbore, 2000). The soil OM with slow turnover rates is intimately associated with the soil mineral 58 matrix and accounts for most of the C accumulated in the soil profile in temperate arable soils. Thus,

even small changes in turnover rate of stable OM may lead to substantial and long-lasting impacts on
the CO₂ flux from soil to atmosphere (Trumbore, 2000; Schmidt et al., 2011). This spawns a timely need
to link temperature sensitivity to the mechanisms that stabilize or destabilize C in mathematical models
that simulate long-term turnover of soil C.

63

Soil OM persists because either its chemical structure is resistant to enzymes released by decomposers or it has become physically or chemically protected against decomposer activity. The decomposition rate of particulate organic matter (POM) has been linked to its intrinsic chemical complexity (Melillo, 1982; Tuomi et al., 2009), whereas the decomposition rate of stabilized OM relates to its incorporation into organo-mineral complexes and reaction with the extensive surface areas of clay- and silt-sized particles (Christensen, 2001; Kleber et al., 2007).

70

71 The decomposition of stable soil OM is often considered to be more sensitive to changes in temperature 72 than that of labile OM (Conant et al., 2008). This concept is based on the Arrhenius equation (Bosatta 73 and Agren, 1999) according to which the temperature sensitivity of chemical reactions increases with 74 increase in activation energy. Stable OM comprises chemically complex molecules for which decomposition reactions require a high activation energy and thus are highly sensitive to temperature. 75 76 However, OM with long turnover time also includes chemically simple and labile proteins and 77 polysaccharides of microbial origin that have become stabilized in organo-mineral complexes (Kleber et al., 2011). The Arrhenius equation does not necessarily apply when other mechanisms than molecular 78 79 complexity limit the decomposition process (Davidson et al., 2006). Stabilization by organo-mineral 80 interactions is known to reduce the temperature sensitivity of OM (Gillabel et al., 2010; Moinet et al., 81 2018), but the temperature sensitivity of OM with different age residing within a given soil size fraction 82 remains unclear (Leifeld and Fuhrer, 2005; Plante et al., 2010; Poeplau et al., 2017).

83

The effect of OM association with minerals on the temperature sensitivity of recent (<25 years old) C versus older C has not previously been studied. In this study it was determined by measuring the ${}^{12}CO_2$ and ${}^{13}CO_2$ evolved during incubation of OM fractions that were isolated by physical fractionation of soils sampled in an experiment where C3-plants have been replaced by C4-plants 25 years ago.

88

89 The temperature sensitivity of C in soil OM with different mineral association was examined by 90 incubating two particle size fractions (<63 µm, mineral- associated OM (MOM) and >63 µm, particulate 91 OM (POM)). The soils were from an experiment where silage maize (a C4-plant) was grown for 25 years on soil previously under C3-plants exclusively. The difference in δ^{13} C between plants with the C3 and the 92 C4 photosynthetic pathway is typically 15 ‰ (O'Leary, 1988). This allowed us to discriminate between 93 old and recent C (old, C3-C > 25 years; recent, C4-C < 25 years) present in a given fraction and 94 95 between CO₂ originating from old and recent soil C. The objectives were to quantify 1) the temperature 96 sensitivity of C residing in POM and MOM, and 2) the sensitivity of differently aged C associated within 97 each of these size fractions. We hypothesized: 1) that the temperature sensitivity of POM increases with 98 increasing age following Arrhenius equation because chemical recalcitrance controls the decomposition 99 of POM, and 2) that the temperature sensitivity of C in MOM is less affected by age than C in POM 100 because the mineral-association limits the decomposition of MOM. Thus, it follows 3) that the overall 101 decomposition of MOM is less sensitive to temperature than POM.

102

103 The composition of the soil microbial community may change with OM quality and incubation 104 temperature (Biasi et al., 2005). The balance between fungi, bacteria and archaea during the incubations 105 was determined using quantitative PCR, to reveal if respiration rate and changes in isotopic ratio relate 106 to changes in microbial community (Paterson et al., 2009).

107 2. Materials and Methods

108 **2.1. Site and sampling**

109 The study relied on soils sampled in a C3- to C4-vegetation conversion experiment at Askov 110 Experimental Station, Denmark (55°28N, 09°07E). Annual mean temperature and precipitation during 111 1981–2010 were 8.2 °C and 1079 mm, respectively. The experiment was established in 1987 when soil 112 (0-25 cm) was sampled from two agricultural sites. The Askov soil (ASK) has loamy sand texture with 14.1 % clay (< 2 µm), 21.2 % silt (2-20 µm), 30.6 % fine sand (20-200 µm), and 34 % coarse sand (200-113 114 2000 µm). The Lundgaard soil (LUN) has coarse sand texture with 5.5 % clay, 7.7 % silt, 16% fine sand 115 and 70.8% coarse sand. The soils were sieved to <4 cm and placed outdoors in large open-ended 116 cylinders (0.76 m²; 50 cm high) inserted 45 cm into the ground and resting on undisturbed subsoil. Until 117 1987, the ASK and LUN soils had been exclusively under C3-crops (mainly cereals). At the start of the

118 experiment (1988) both soils had a C/N ratio of 13, and ASK soil had pH of 6.4, and LUN soil had pH of 7.6 (H₂O). Soil pH is maintained by occasional addition of Ca(OH)₂. Soil C contents and δ ¹³C values are 119 presented in Table 1. Every year since May 1988, the soils have carried the C4-crop silage maize (Zea 120 121 mays L.). The maize receives NPK mineral fertilizers every spring and is whole-crop harvested for silage 122 in mid-October. Kristiansen et al. (2005) give further details. Soils sampled every two-to-four years since 123 1988 are archived in air-dry condition. The present study received soil that was sampled in 1988 before 124 the first maize crop (year = 0) and in 2013 after 25 years of maize cultivation (year = 25). The 1988 125 samples served as reference for the isotopic composition of C3-derived soil C. The soils were sieved to 126 < 2 mm.

127 **2.**

2.2. Soil fractionation and incubation

The air-dry soil samples from 2013 were divided into four laboratory replicates for physical fractionation, while the amount of soil archived in 1988 did not allow for replication. Samples were soaked overnight in water and fractionated by stirring and wet sieving through a 63- μ m mesh. Then the >63 μ m (POM, including sand) and <63 μ m (MOM) size-fractions and bulk soil (< 2 mm) subsamples were dried at room temperature to constant weight. Soil water holding capacity (WHC) was determined separately for each soil fraction before incubation.

134

135 Samples of bulk soil, POM and MOM fractions from 2013 were rewetted to 60% WHC and pre-incubated 136 at 20 °C for two weeks to remove any labile C released during the fractionation process (De Nobili et al., 137 2006). Four replicate samples were incubated in 120 mL glass bottles with a rubber septum. The amount 138 of sample incubated ranged between 3 and 31 g. The quantity of sample was determined in a preexperiment in order to allow for sufficient CO₂ for ¹³C analysis when using the same incubation times for 139 140 all fractions. The amounts used for incubation were: 9 g of ASK bulk soil and 19 g of LUN bulk soil, 26 g 141 of ASK > 63 µm fraction and 31 g of LUN > 63 µm fraction, 9 g of ASK < 63 µm fraction and 3 g of LUN < 142 63 µm fraction. Three replicates were used for measurement of CO₂ evolution and one for quantitative 143 PCR (extraction of microbial DNA). After pre-incubation, bottles resided in a water bath for 24 hours to 144 adapt to the first incubation temperature (6 °C). After closing the septum, the bottle headspace was flushed with moist CO₂-free air (80% N₂ and 20% O₂). The soil samples were incubated ramping up and 145 146 down the temperature steps: 6, 18, 26 and 34 °C (sequential method; Hamdi et al., 2013) to minimize

147 potential bias from any change in substrate quality during incubation (Leifeld, 2003). In the sequential method, the CO₂-flux at a given temperature, and in our case also the δ^{13} C content of the respired CO₂, 148 149 is the average of the values measured at a given temperature when ramping up and down the 150 temperature scale. The incubation periods were as short as possible to avoid changes in OM quality 151 while ensuring measurable CO₂ concentrations. Soil samples equilibrated to the next incubation 152 temperature in a water bath for 24 hours before starting the CO_2 collection. Before starting the CO_2 153 collection, the bottle headspace was flushed with CO₂-free air. Then the bottles were incubated in the 154 water bath at each temperature level until the CO₂ concentration in the headspace reached about 1000 155 ppm. At that time point we measured the CO₂ concentration in the headspace and retrieved gas samples for ¹³C analysis. Then the bottle was transferred to the next temperature level and the procedure 156 repeated. The incubation times for CO₂ collection at each temperature step were: Ramping up the 157 158 temperature range: 6 °C: 6 days, 18 °C: 2 days, 24 °C: 1 day, 34 °C: 12 hours, and ramping down the 159 temperature range: 24 °C: 1 day, 18 °C: 2 days, 6 °C: 8 days. The CO₂ concentrations were measured 160 using a Hewlett Packard 6890 gas chromatograph equipped with a TC detector and J&W Scientific 161 Megapore GS-Q column.

162 **2.3. Isotope analysis**

163

For isotope analysis, gas samples were retrieved in a He-flushed glass vial (12 ml Exetainer®, Labco Limited, UK) using a syringe. The δ^{13} C of the CO₂ was determined using a DeltaPlusXL (Thermo Finnigan) continuous flow isotope-ratio mass spectrometer and references NBS-19, NBS-18 and L-SVEC to normalize raw isotope data. The analytical precision was ± 0.15 ‰ (1 σ).

168

Subsamples of size fractions and bulk soil were ball-milled for isotope analysis. The C concentrations were measured using CHN-analyser (CHN-1000, Leco) and ¹³C using NC2500 Carlo-Erba analyser coupled with Delta^{plus}Advantage (Thermo Fisher Scientific) continuous flow isotope-ratio mass spectrometer. The ¹³C results are given as δ^{13} C ‰ based on standard V-PDB and reference materials (IAEA-CH3 and IAEA-CH7). The analytical precision was ± 0.15 ‰ (1 σ). All soil isotope analyses were run in duplicate and results averaged.

175 **2.4. Quantitative PCR (qPCR)**

176 Quantitative PCR (qPCR) was used for examining the proportions of fungi (ITS), bacteria (B16S) and archaea (A16S) in soil at each incubation temperature. Gene copy numbers were determined on the 177 additional set of bottles incubated along with bottles used for CO₂-collection. Samples for DNA extraction 178 were taken from the same bottle after incubation at 6, 18, 26 and 34°C, when ramping up the 179 temperature scale. Microbial DNA was extracted from freeze-dried subsamples using the NucleoSpin 180 181 Soil DNA extraction kit (Macherey-Nagel). Quantitative polymerase chain reaction (gPCR) assays were 182 conducted in thin-walled PCR tubes (Qiagen) on a Rotor-Gene 6000 real time PCR machine (Corbett 183 Life Science). We amplified gPCR products obtained with fungal ITS1F and ITS2 (Gardes and Bruns, 184 1993; White et al., 1990), bacterial 16S rRNA 1055F and 1392R (Lee et al., 1993; Olsen et al., 1986; 185 Stahl et al., 1988) and archaeal 16S rRNA Arch967f and Arch1060R primer pairs (Amann et al., 1990; Reysenbach and Pace, 1995; Riley-Buckley, 2001). The assays were run with Maxima[™] SYBR Green 186 187 qPCR Master Mix (2X) (Thermo Scientific) in a 20 μl reaction volume containing 1μl template, 0.375 μM 188 of each primer, and 1x qPCR master mix. For qPCR conditions see Peltoniemi et al. (2015). Fluorescence was measured at the end of each extension step. The standard curves were constructed 189 190 with plasmids containing corresponding inserts, taking into account the concentration and molecular 191 mass of the plasmid, including the insert. The samples and standards were run in duplicate. All samples 192 were replicated and absence of PCR inhibition was verified through 1:10 dilution. The copy numbers in 193 samples were calculated based on comparison with threshold cycle values of the standard curve; the 194 numbers are given per gram of soil C.

195 **2.5. Calculations**

Soil C and respired CO₂ was divided into C3-C (pre-1988) and C4-C (post-1988) using the isotope massbalance equation (Balesdent and Mariotti, 1996). The fraction of the total soil C originating from C4 plants, F_{C4-C} , was estimated as:

199

200
$$F_{C4-C} = \frac{\delta^{13} c_{t=25} - \delta^{13} c_{t=0}}{\delta^{13} c_{c4-C} - \delta^{13} c_{t=0}},$$
 (1)

201

where $\delta^{13}C_{t=25}$ is the value of soils retrieved in 2013 and $\delta^{13}C_{t=0}$ is the value of the corresponding samples from 1988. The calculation of $\delta^{13}C_{c4-c}$ accounts for the small changes in isotopic composition of maize biomass recorded over the experimental period (Christensen et al., 2011). This was estimated for *year*205 2013 as follows:

207
$$\delta^{13}C_{C4-C} = -0.043 \ year + 74.15.$$
 (2)

208

The CO₂ respiration rates at a given temperature (up and down the temperature range) were averaged.
The Gaussian temperature function is as follows:

211

212
$$R(t) = R_0 e^{aT + bT^2},$$
 (3)

213

where *T* is the temperature (°C) and the parameters are $R_0 > 0, a > 0, b < 0$ (Tuomi et al., 2008). These were fitted to measurements using Bayesian framework (Gelman et al., 2013). Measurement errors in the Gaussian mode were assumed to be independent and follow a log-normal distribution. This error distribution was a natural choice for exponential model to account heteroscedasticity. The prior information of the parameter R_0 was modelled with a log-normal distribution with mean 100 and variance 100^2 . Uninformative priors were used for parameters *a* and *b*, that is $p(a, b) \propto 1$. In this way, the following posterior was obtained:

221

222
$$p(\log (R_0), a, b | y) \propto p(\log (R_0))p(a, b)p(y | \log (R_0), a, b) \propto Normal((V^{-1} + H^T H/\sigma^2)^{-1}(V^{-1}m + HTy/\sigma^2), (V-1+HTH/\sigma^2)-1,$$
 (4)

224

where each row in matrix *H* is $[1, T_i, T_i^2]$ with measurement temperatures T_i and data vector *y* contains the corresponding CO₂ measurements y_i . The matrix *V* and vector *m* relate to prior values. From the above posterior, analytical equations were derived for marginal densities.

228

The temperature coefficient Q_{10} describes the change in respiration rate when temperature increases by 10 °C and was calculated from *a* and *b* as follows:

232
$$Q_{10}(T) = \frac{R(T+10)}{R(T)} = e^{10a+20bT+100b}$$
(5)

233

resulting in a log-normal posterior distribution for $Q_{10}(T)$.

235

We analysed the C age using the isotopic composition of soil converted from C3 to C4 vegetation 25 years ago. The soil C pool was previously observed to be in steady-state equilibrium (Thomsen and Christensen, 2010). This allows a simple calculation of mean residence time (MRT) of C in bulk soil and both POM and MOM fractions using the δ^{13} C values according to Amelung et al. (2008):

240

241
$$MRT = -\frac{t}{\ln(1 - F_{C4-C})},$$
 (6)

242

where F_{C4-C} is the C4-C fraction of the soil C pool (Eq. 1) and *t* is time in years since the vegetation change (*t*=25). The significance of trends in temperature series was tested with a *t-test* for trend. The calculations for the Eqs. 1-3. and 6 were done in MS Excel. Differences in the qPCR results (logtransformed gene copy numbers) between different SOC fractions were tested with one-way ANOVA combined with Tukey test (IBM SPSS Statistics 22). Test for data normality used the Shapiro-Wilk test, and Levene test was used for testing of homogeneity of variances.

249 **3. Results**

250 **3.1. Soil characteristics**

The <63 μ m fraction (MOM) had a higher C concentration and contained more C than the >63 μ m fraction (POM) (Table 1). The MOM represented 67 to 69 % of the total soil C in ASK both in 1988 and 2013. For LUN, MOM accounted for 74 % in 1988 and 57 % in 2013, mainly because of a higher C concentration in POM in 2013. The C concentration in MOM was about 5 % while that in POM ranged from 0.3 to 1.3 %.

256

The MOM and POM isolated from LUN soil had very similar δ^{13} C values before maize growing started in 1988, whereas for ASK soil MOM was somewhat depleted in ¹³C compared to POM (Table 1). Maize cultivation increased the δ^{13} C values of each soil fraction, and the values of the POM increased most. In 2013, maize-derived C represented 37 % of the C in POM in ASK and 39 % in LUN, while maize-C accounted for 14 % of the C in MOM at both sites (see Fig. 4). For LUN as well as ASK soil, the MRT of

262 C in POM was 50 years and about 170 years for C in MOM (Table 1).

263 **3.2. Temperature sensitivity of SOC fractions**

The ASK bulk soil, POM and MOM fractions all showed similar respiration rates during incubation (Fig. 1a). In LUN, the average respiration rate increased from the bulk soil to the MOM and POM. The respiration rates were higher than deducted from the MRTs of the fractions (seen Table 1). For example, the respiration rate measured at 18 °C equalled 80 μ g CO₂ g⁻¹ C hour⁻¹, corresponding to about 0.2 g C g⁻¹ C per year.

269

270 The Gaussian equation was used to describe the temperature dependence of the respiration. When fitted to measured respiration from MOM and POM isolated from ASK soil, the 95 % probability 271 272 distributions of the two parameter values of this equation overlapped only little (Fig. 2a). This indicates a 273 different temperature dependence as illustrated by the Q₁₀ value in Figs. 2c, e-g. For example, in the 274 temperature range 5 to 15 °C, the most probable Q₁₀ value of the POM equalled 4.8 and that of the MOM 275 3.6. Because of the Gaussian equation, the Q_{10} values and the temperature sensitivity of the MOM and 276 POM fractions from ASK soil differed for different temperatures. The effect of temperature on the 277 respiration did not differ between size fractions from LUN soil (Fig. 2b, d-g).

278

The δ^{13} C value of respired CO₂ was some 4 ‰ higher than that of C in the corresponding soil fraction (Fig. 3, Table 1). This indicates that decomposers favoured the younger maize-derived C. Maize-derived C represented 63 % (LUN) and 70 % (ASK) of the C respired from POM, 52 % (ASK) and 60 % (LUN) of the C respired from bulk soil, and 39 % (ASK and LUN) of the C respired from MOM (Fig. 4). The δ^{13} C values did not vary significantly with incubation temperature, except for C respired from POM isolated from the LUN soil (Fig. 3). Here the δ^{13} C decreased with an increasing temperature indicating that an increasing proportion of the older C3-C was respired at higher temperatures.

286

3.3. Microbial community composition measured by quantitative PCR

The ITS, B16S and A16S copy numbers did not change systematically with incubation temperature. In the ASK soil, the POM fraction had lower ITS, A16S and B16S copy numbers per mg soil C than bulk soil and MOM, but these differences were not statistically significant at all temperatures (Fig. 5). In the

LUN soil, the MOM fraction had higher ITS and B16S gene copy numbers compared to bulk soil andPOM fraction (Fig. 5).

292 4. Discussion

293 We studied the temperature dependence of C in bulk soil and their particle size fractions from two 294 differently textured arable soils converted from C3-plants to C4-plants 25 years ago. The >63 µm fraction 295 with a MRT of decades is dominated by particulate organic matter (POM), whereas the <63 µm fraction 296 is dominated by OM intimately associated with soil minerals (MOM) and with a MRT of more than a 297 century. Our estimates of old C3-derived C and recent C4-derived C depend on $\delta^{13}C_{v=0}$ determined for 298 the soils before maize cultivation was initiated in 1988. Accounting for the annual increase in $\delta^{13}C_{v=0}$ of 299 0.0083‰ (Christensen et al., 2011) changed the percentages of C3-C and C4-C in soil particle size 300 fractions by only 1%. Menichetti et al. (2015) showed that in experiments lasting for more than three to 301 four decades, isotopic fractionation could lead to larger changes in soil δ^{13} C.

302

303 Our study relied on air-dried and sieved soil samples and a soil fractionation procedure that eliminates 304 soil structure. The potential effect of the rewetting on the observed temperature sensitivity of the soil OM 305 was minimized by pre-incubating samples for two weeks. The pre-incubation showed a flush of CO₂ for 306 bulk soil as well as POM from the LUN soil (Fig. S1), suggesting that this flush of CO₂ was not only due 307 to fractionation procedure. Based on the decreasing respiration rates during the pre-incubation it was 308 estimated that the C released from bulk soil accounts for less than 30 % of C in the >63 µm fraction and 309 more than 70% of C in the <63 µm fraction. If the respiration rate of the >63 µm fraction did not change 310 due to fractionation, then the unexplained increase in the respiration rate of <63 µm fraction after 311 fractionation corresponds to 11 % and 42 % of the total respiration in ASK and LUN, respectively. Thus, 312 part of the respiration from POM as well as MOM may originate from C released during re-wetting and 313 fractionation. For both soils, however, old C-3 derived C made up a larger proportion of the C respired 314 from MOM than from POM. For POM, a larger part of the soil C and respired C originated from recent C-315 4 derived C (Fig. 4). This suggests that we did manage to separate two soil fractions with OM of different 316 age, allowing us to compare the temperature sensitivity of soil C older and younger than 25 years.

317

We estimate that on average 0.9% of the soil C was lost during the incubation. With this incubation setup, it was also possible to observe if the isotopic signature changed to the same direction when moving up and down of the temperature scale (e.g. for POM in LUN, the δ^{13} C value exhibits slopes of -0.046 and -0.060 with increasing temperature when measured up and down the temperature scale). Therefore, we do not consider changes in substrate quality during the incubation to have affected our results.

323 4.1. Temperature sensitivity and age of C

In the LUN soil, the relative contribution of C3-derived C to respiration from POM increased with temperature (Fig. 3), reflecting a higher temperature sensitivity of the old C3-C than the recent maizederived C4-C. This is ascribed to a chemically more complex structure of the part of POM that has survived for > 25 years. The surviving old POM demands higher activation energy and thus shows higher temperature sensitivity.

329

330 The high temperature sensitivity of old C in the POM fraction supports our first hypothesis that C release 331 from this fraction follows Arrhenius kinetics. We hypothesised that mineralisation of C in this fraction 332 reflects the intrinsic temperature sensitivity of C in POM as this fraction is freely available to microbes. 333 However, this effect of age was less pronounced in ASK loamy sand, and the trend was not statistically 334 significant. This may be due to less complete soil dispersion in the more clayey ASK soil. Even though 335 the fractionation procedure aimed at breaking all soil aggregates, some microaggregates (63-250 µm) in 336 the ASK soil may have survived dispersion. This may explain why the ASK size fractions and bulk soil 337 had rather similar respiration rates. For the coarser LUN soil with a more complete soil dispersion, POM 338 showed higher respiration rate. There was slightly more old C3-C respired from the POM fraction of the 339 LUN soil compared with that of the ASK soil, even though the proportions of C3-C and C4-C in POM 340 were almost identical for both soils. This supports the previous notion that physical protection of C within 341 microaggregates that survived dispersion may have affected POM availability in the ASK soil.

342

There was no change with temperature in the isotopic signal of CO_2 from the MOM fraction, suggesting that mineral-associated recent C4-C and old C3-C in MOM exhibit similar temperature sensitivity. This supports our second hypothesis stating that temperature sensitivity of MOM is less affected by age than that of POM. Our results indicate that the most temperature sensitive OM pool in soil is the free, but chemically recalcitrant C with intermediate turnover time, aligning with recent studies, which have also

found that C fractions with decadal-scale turnover times show the highest temperature sensitivity (Gillabel et al., 2010; Hopkins et al., 2012).

350 4.2. Temperature sensitivity of soil C fractions

351 Our third hypothesis stated that organo-mineral (MOM) association of C reduces its intrinsic temperature 352 sensitivity compared to C in POM. This hypothesis was only partially supported by the lower temperature 353 sensitivity of MOM and higher sensitivity of POM in ASK soil (Fig. 2a) as illustrated by different Q₁₀ values, especially in the temperature range of 5-15 °C (Fig. 2e). This is in line with previous studies 354 355 showing that stabilization of C in organo-mineral complexes reduces its temperature sensitivity (Gillabel 356 et al., 2010; Moinet et al., 2018). However, differences in Q_{10} s of the incubated size fractions were less 357 clear at higher temperatures and not statistically significant for LUN soil. We speculate that the stabilising 358 effect of organo-mineral interaction was stronger in MOM from the finer textured ASK soil, whereby 359 stabilisation with minerals reduced the Q_{10} more compared to the Q_{10} of C in POM, which was similar for 360 both soils.

361

362 Studies using incubations of physically or chemically isolated fractions to study their temperature 363 sensitivity are scarce and results are partly contradictory. Plante et al. (2010) found the Q₁₀ to increase in 364 the order: hydrolysis residue<POM<whole soil. In their study, POM was fractionated with wet sieving 365 size-separation method comparable to our study, whereas Q₁₀ was determined from long-term (126-366 days) parallel incubations. This could have decreased the Q₁₀ of labile POM fraction more relative to bulk soil Q₁₀ values. Contrary to Plante et al. (2010), in our study there were no clear differences between the 367 368 Q₁₀ values of POM and bulk soil. Benbi et al. (2014) found higher temperature sensitivity for POM 369 compared to MOM, which aligns with our results from the ASK soil. Their study relied on a size-370 fractionation procedure similar to that of our study, but divided POM into two size classes (53-250 µm and 250-2000 µm). Ding et al. (2014) reached opposite results, and found that the Q₁₀ for the different 371 372 size-fractions increased in the order: sand<silt<clay. In their study, however, soil dispersion by 373 ultrasonication and the crushing of the size-fractions after drying prior to incubation could have increased 374 the availability of mineral associated OM. Flocculation of clay with CaCl₂ could have also affected C 375 availability from that fraction. Leifeld and Fuhrer (2005) found little difference between size fractions (<63 376 μ m and >63 μ m) whereas C in residues from acid hydrolysis of finer sized soil fractions showed a high 377 sensitivity to temperature. The residue obtained after acid hydrolysis is considered less suitable for

378 studying stable C in incubations studies. The isolated soil OM fractions are impacted by chemical 379 changes that may make mineral associated C either more available (Leifeld and Fuhrer, 2005) or less available due to cation bridging with Ca²⁺ added to neutralise pH after hydrolysis (Plante et al., 2010). 380 Our fractionation procedure avoided these pitfalls. However, laboratory incubations with optimal 381 382 conditions for microbial decomposition always provide shorter turnover time estimates than methods based on changes in the abundance of ¹³C after C3- to C4-vegetation change (Feng et al., 2016). This 383 384 was also true in our study. Although decomposition rates of MOM in incubation studies may not refer 385 directly to field conditions, they are still valid when comparing the temperature sensitivity of older versus 386 younger C in a given fraction.

387 4.3. Microbial balance

388 The composition of the microbial community may affect mineralization rates and the isotopic composition 389 of respired CO₂ (Paterson et al., 2009). We did not directly analyse the microbial community, but 390 estimated the balance of the community targeting the three domains fungi, bacteria and archaea of the 391 microbial community by qPCR. No significant changes in the balance were observed with temperature 392 although the relative abundance of archaeal 16S rRNA copy numbers tended to increase with 393 temperature. Archaea are the least abundant microbial group in soil (Tamez-Hidalgo et al., 2016), and 394 their functional importance is less understood than that of bacteria and fungi. The small changes in this 395 microbial group are unlikely to explain observed changes in CO₂ production or isotope signatures of 396 respired CO₂. The rather similar balance between the three domains in soil-size fractions and bulk soils 397 suggests that trends in isotopic composition of respired CO₂ were not derived from differences in the 398 composition of the microbial community. The qPCR analysis showed higher gene copy numbers in the 399 MOM than in the POM fraction isolated from LUN soil. Most of the decomposer community live attached 400 to the surfaces of clay and silt particles (Mills, 2003), whereby these particles may become enriched in 401 gene copies. Because qPCR records genes associated with live as well as dead microbial biomass, a 402 higher gene copy number in the MOM fraction may indicate a larger contribution of microbial necromass 403 to the stabilised C in this fraction (Liang et al., 2017).

404 **5. Conclusions**

The temperature sensitivity of C in the <63 μ m particle size fraction (MOM), dominated by mineral associated C, was smaller or similar to the sensitivity of C in the >63 μ m size fraction (POM). The

407 temperature sensitivity of respiration increased with age of C in the POM fraction while the sensitivity 408 was unaffected by age of C in the MOM fraction where OM is stabilised by association with minerals. No 409 evidence was found for highly temperature sensitive C being a significant component of the MOM 410 fraction. Our results imply similar temperature sensitivity of older and younger C, when residing in 411 organo-mineral associations. This aligns with recent evidence suggesting that the most temperature 412 sensitive C is in soil OM pools cycling on intermediate time scales. Our results suggest that the temperature sensitivity of most of the C residing in arable topsoils relates to the mechanisms that 413 414 stabilize C in soil rather than to the age of the OM stabilized in organo-mineral complexes.

415

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422

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544 **Figure captions**

- 545 Figure 1. Measured respiration rates at different temperatures and the modelled Gaussian temperature 546 curve for soil particle size fractions and bulk soil: 1 a) ASK and 1 b) LUN.
- 547
- Figure 2. 95% probability contours of parameter values *a* and *b* in the Gaussian model for the bulk soil and particle size fractions in ASK (a) and in LUN (b), Q_{10} as a function of temperature in ASK (c) and in

550 LUN (d). Dashed lines represents 95% credible intervals (Bayesian confidence intervals). Posterior 551 distributions of Q_{10} calculated for temperature range of 5-15 °C (e), 15-25 °C (f) and 25-35 °C (g).

552

Figure 3. Mean \pm standard deviation (N=6) of isotopic composition of CO₂ respired from the bulk soils and soil size fractions at different temperatures in ASK (a) and in LUN (b). Lines represent linear regressions fitted to the means: a) ASK: MOM (y = 0.03x - 22.44, R²=0.70, p=0.16), POM (y = -0.03x -16.59, R²=0.63, p=0.21), Bulk soil (y = 0.01x - 19.94, R²=0.08, p=0.72) b) LUN: MOM (y = -0.00x - 21.24, R²=0.01, p=0.89), POM (y = -0.05x - 16.80, R²=0.92, p=0.04), bulk soil (y = -0.05x - 17.49, R²=0.68, p=0.18). The trend in POM fraction (>63 µm) in LUN is statistically significant.

559

Figure 4. The proportions of C4-C and C3-C in soil and in respired CO_2 calculated from their isotopic composition in ASK (a) and LUN (b). The δ^{13} C values for CO_2 are averages four replicates and 4 temperatures measured twice.

563

Figure 5. Mean \pm standard deviation of the gene copy numbers/ mg soil C for ITS gene in a) ASK and b) LUN, for B16S in c) ASK and d) LUN, and A16S in e) ASK and f) LUN. The Anova and Tukey's test for differences in gene copy numbers between different SOC fractions (bulk, MOM <63 µm and POM <63 µm) were tested separately in each temperature. Different letters indicate statistically significant differences between fractions. The figure presents non-transformed data, but the y-axis in the figure is in log-scale. For the statistical analysis log-transformed data was used.

571 Tables

572 Table 1. Mean \pm standard deviation of C contents, mass proportions and isotopic composition of ASK 573 and LUN bulk soils, POM and MOM fractions in 1988 and 2013 as well as the mean residence time 574 (MRT) of C. In 2013, N=2 for δ^{13} C except for the POM fraction where N=8.

| | 1988 (t=0) | | | 2013 (t=25) | | | |
|-----------|---------------|----------|-----------------------|-------------|----------|-----------------------|-----------|
| | C (%) | Mass (%) | δ ¹³ C (‰) | C (%) | Mass (%) | δ ¹³ C (‰) | MRT years |
| N | 2 | 1 | 4 | 2 | 4 | 2/8 | |
| ASK | | | | | | | |
| Bulk soil | 2.5 ± 0.1 | 100 | -27.9 ± 0.2 | 2.4 ± 0.1 | 100 | -24.5 ± 0 | 100 ± 4 |
| MOM | 5.4 ± 0.3 | 33 | -28.2 ± 0.1 | 5.7 ± 0.3 | 32 ± 3.9 | -26.1 ± 0.1 | 170 ± 8 |
| POM | 1.3 ± 0.3 | 67 | -27.4 ± 0.4 | 1.2 ± 0.1 | 68 ± 3.9 | -22.1 ± 0.8 | 54 ± 9 |
| LUN | | | | | | | |
| Bulk soil | 1 ± 0.1 | 100 | -26.9 ± 0 | 0.9 ± 0 | 100 | -22.6 ± 0.3 | 71 ± 7 |
| MOM | 4.8 ± 0.1 | 15 | -26.7 ± 0.2 | 4.9 ± 0.1 | 12 ± 1.7 | -24.8 ± 0.2 | 170 ± 15 |
| POM | 0.3 ± 0.2 | 85 | -26.7 ± 0.5 | 0.5 ± 0.2 | 88 ± 1.7 | -21.4 ± 1 | 51 ± 12 |







Figure3





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