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1 **The role of soil characteristics on temperature sensitivity of soil organic matter.**

2

3 **Abstract**

4 The uncertainty associated with how projected climate change will affect global carbon
5 (C) cycling could have a large impact on soil C stocks. The purpose of our study was to
6 determine how various soil decomposition and chemistry characteristics relate to SOM
7 temperature sensitivity. We accomplished this objective using long-term soil incubations
8 at three temperatures (15, 25, and 35°C) and pyrolysis molecular beam mass
9 spectrometry (py-MBMS) on twelve soils from six sites along a mean annual temperature
10 (MAT) gradient (2 to 25.6°C). We calculated Q_{10} values from the CO_2 respired in our
11 long-term incubation using the Q_{10-q} method and found the decomposition of the resistant
12 fraction to be more temperature sensitive with a Q_{10-q} of 1.95 ± 0.08 for the labile fraction
13 and a Q_{10-q} of 3.33 ± 0.04 for the resistant fraction. We used a two pool model (active and
14 slow) with first-order kinetics to fit our soil respiration data, along with a three pool
15 model for comparison purposes. We found that the two and three pool models
16 statistically fit the data equally well. The size of the active pool in these soils, calculated
17 using the two pool model, increased with incubation temperature and ranged from 0.1 to
18 12.8% of initial soil organic C. Sites with an intermediate MAT and lowest C:N ratio had
19 the largest active pool. Pyrolysis Molecular Beam Mass Spectrometry on these soils
20 indicated that there were chemical differences in the SOM between the land use
21 treatments which may have lead to differences found between the total amount of CO_2
22 respired and size of the active pool of the soils.

23

24 **Keywords:** decomposition; temperature sensitivity; soil carbon; long-term incubation;
25 py-MBMS

26

27 **1. Introduction**

28 Temperature effects on chemical reactions have been studied since the late
29 1800's, originating with Arrhenius and van't Hoff (Lloyd and Taylor, 1994); yet, there is
30 still much debate on mechanistically how temperature regulates soil organic matter
31 (SOM) decomposition. Soil organic matter contains two to three times as much C as the
32 atmosphere (Davidson et al., 2000), with respiration of SOM by microbes contributing
33 50-75 Pg of CO₂-C to the atmosphere annually – approximately ten times the current
34 annual emissions from burning fossil fuels (Schimel, 1995). Temperature is an important
35 factor controlling SOM turnover and understanding how temperature affects SOM
36 decomposition will allow us to better predict how global climate change will affect SOM
37 stocks.

38 A Q₁₀ of 1.5-2.0 or an Arrhenius type function, where the effective activation
39 energy for respiration varies inversely with temperature, has commonly been used for the
40 temperature response of SOM decomposition (Friedlingstein et al., 2006), and most
41 terrestrial C models apply the same respiration-temperature relationship to each of the
42 different SOM pools (Melillo et al., 1995; Burke, 2003; Friedlingstein et al., 2006).
43 Contrary to common model formulations in which different types of SOM follow the
44 same temperature decomposition relationship, many recent studies have found that

45 different SOM pools have different temperature responses, although consensus on their
46 apparent temperature sensitivities has not yet been reached (Davidson and Janssens,
47 2006). In addition, although determining the temperature sensitivity of SOM by
48 evaluating the Q_{10} response is commonly calculated using incubations of fixed durations,
49 the approach may generate incorrect values for temperature sensitivity due to
50 comparisons of SOM with different labilities at different temperatures (Leifeld and
51 Fuhrer, 2005; Reichstein et al., 2005; Conant et al., 2008), which further complicates
52 conclusions about SOM temperature sensitivity.

53 Understanding the temperature sensitivity of SOM decomposition is challenging
54 because soil organic matter is composed of many different organic C compounds, with
55 differing inherent kinetic properties (Davidson and Janssens, 2006). To simplify the
56 process of modeling SOM decomposition, this range of compounds is usually classified
57 into a small number of discrete, kinetically-defined pools with some portion of SOM
58 being labile and easily decomposable and the rest comprising one or more other pools
59 being less labile and decomposing slowly. In most decomposition models, temperature
60 effects are modeled as a decomposition rate multiplier for fixed SOM pools (Lloyd and
61 Taylor, 1994; Del Grosso et al., 2005). However, some recent studies have hypothesized
62 that temperature may actually alter the amount of substrate that would be considered
63 easily decomposable or labile (Zogg et al., 1997; Zak et al., 1999; Dalias et al., 2003;
64 Rasmussen et al., 2006). Such a response could be caused by temperature-driven shifts in
65 microbial community composition (Zogg et al., 1997; Zak et al., 1999) or in the selection
66 of compounds being decomposed (Andrews et al., 2000). A change in the microbially-

67 available SOM indicates that reaction rates may not be the only factor controlling
68 decomposition responses to altered temperatures.

69 The objective of this study was to reconcile the differences between the inherent
70 and apparent temperature sensitivity of SOM with differing labilities. To accomplish this
71 we determined the temperature sensitivity of a labile and more resistant soil fraction
72 using a long-term incubation on a set of soils along a mean annual temperature gradient
73 (MAT). At each point along our MAT we had two soils with different land uses as a way
74 to compare similar soils with differing amounts and types of SOM. We used respiration
75 response curves and pyrolysis molecular beam mass spectrometry to characterize the
76 SOM in each of our soils. By utilizing this multi-approach method we wanted to have a
77 better understanding of the apparent temperature sensitivity of SOM.

78

79 **2. Materials and Methods**

80 *2.1. Sample Sites*

81 Surface soil samples were collected from six sites along a mean annual
82 temperature gradient (2 to 25.6°C), each with a native and either cultivated (Indian Head,
83 SK; Mandan, ND; Akron, CO; and Waggoner Ranch, TX) or pasture (Alajuela, Costa
84 Rica and Nova Vida Ranch, Brazil) land use (Table 1). At each site, samples were
85 collected from three locations separated by several meters each (field replicate n = 3)
86 within each land use. Surface litter and above ground vegetation were cleared away prior
87 to sampling. Small pits were dug to a depth of 20 cm, and samples were collected from 0-
88 20 cm. Soils were packaged and transported to the laboratory, where rocks, surface litter

89 and root materials were removed while soil clods were broken by hand and passed
90 through a 2-mm sieve. Soil samples were then air-dried and stored at room temperature
91 until incubations began. Soil organic C (SOC) and total N concentrations were
92 determined with a LECO CHN-1000 autoanalyzer (LECO Corporation, St. Joseph, MI,
93 USA).

94

95 *2.2. Laboratory Incubation*

96 Four laboratory replicates from bulked field samples from each site and treatment
97 combination were incubated at 15, 25, and 35°C for 588 days. For each sample, 80 g of
98 soil were wetted up to 60% water filled pore space to optimize microbial activity (Linn
99 and Doran, 1984). Gravimetric soil moisture was periodically checked throughout the
100 incubation and water was added when water loss was greater than 5% of initial water
101 added. Samples were placed in sealed canning jars fitted with septa, along with
102 scintillation vials containing 20 ml of water to maintain humidity. Incubation starting
103 days were staggered by replicate and each replicate of a soil was measured on a different
104 day to take into account daily measurement variability. Soils were pre-incubated for
105 three days at 25°C and then four days at the respective incubation temperature prior to
106 measurements to allow the soil to equilibrate after wetting up (Paul et al., 2001).
107 Headspace gas samples were analyzed for CO₂ concentration using a Li-Cor LI-6252
108 IRGA (LI-COR Biosciences Lincoln, NE, USA). Jars were flushed with compressed
109 tank air regularly before CO₂ concentrations reached 5% to prevent CO₂ concentration
110 from inhibiting microbial activity. The CO₂ measurements were taken daily during the

111 first two weeks of the incubation, weekly for the next two weeks, and then every four
112 weeks thereafter, generating a total of 36 sampling times over the course of 588 days.

113

114 *2.3. Temperature Sensitivity*

115 Temperature sensitivity of SOM decomposition/mineralization was determined
116 using the Q_{10-q} method described by Conant et al. (2008a) utilizing the CO_2 respired from
117 our long-term incubations. The method involves determining the amount of time needed
118 for a given amount of C to be respired at a given temperature. The time required to
119 respire a given amount of C at two temperatures is then used to calculate a Q_{10} value.

120 There are two assumptions associated with this method (Conant et al., 2008a). The first
121 is that changes in decomposition rates during the incubation are due to changes in the
122 lability of the SOM being decomposed. The second is that the effect temperature has on
123 the sequence in which SOM compounds are decomposed is small relative to the effect of
124 temperature on decomposition rates. Soil microbial biomass has been found to decline
125 over time in long-term incubations (Follett et al., 2007), but this decline in biomass has
126 not been found to limit the microorganisms capacity to decompose organic matter in
127 long-term incubations (Follett et al., 2007; Steinweg et al., 2008). Because of this we
128 believe that our first assumption that changes in decomposition rates are due to changes
129 in SOM lability is reasonable. Q_{10-q} values were calculated for the initial 0.5% soil C
130 respired, which we considered the labile fraction, and the last 0.5% C respired, which we
131 considered the resistant fraction. Q_{10-q} values were calculated using the 25 and 35°C
132 incubations. The 15°C respiration data was not used due to the limited amount of C

133 respired by the Costa Rica pasture soil. Statistical comparisons of Q_{10-q} values were done
134 by treating the six different sites as random variable replicates (PROC MIXED, SAS
135 Institute, Cary NC).

136

137 2.4. Curve Fitting

138 We used CO₂ respiration results from the 15, 25, and 35°C incubations for curve
139 fitting. The respiration results from the four laboratory replicates of each site-treatment-
140 temperature combination were averaged, and mean respiration rates were used to
141 determine pool size and decomposition rate constants by curve fitting. Pool size and
142 decomposition rate constants were determined using a two-pool model where the two
143 pools, active (C_a) and slow pool (C_s), sum to the total soil C (C_{soil}):

$$144 \quad C_{soil} = C_a + C_s \quad (1)$$

145 We used a two-pool first-order equation (Andren and Paustian, 1987):

$$146 \quad C_{cum}(t) = C_a(1 - e^{-k_a * t}) + C_s(1 - e^{-k_s * t}) \quad (2)$$

147 where $C_{cum}(t)$ is the cumulative soil respiration at time t ($\mu\text{g C (g soil)}^{-1}$), C_a is the size of
148 the active fraction ($\mu\text{g C (g soil)}^{-1}$), and C_s is the size of the slow fraction ($\mu\text{g C (g soil)}^{-1}$).
149 The parameters k_a and k_s are the decomposition rate constants (day^{-1}) for the active and
150 slow pools, respectively. We utilized data from rate curves rather than cumulative
151 respiration because this minimized error accumulation through time (Hess and Schmidt,
152 1995). We used the following rate form of eqn. 2 from Paul et al. (2001) to determine
153 parameter estimates:

154
$$\frac{\Delta C_{cum}}{\Delta t} = C_a * k_a (e^{-k_a * t}) + C_s * k_s (e^{-k_s * t}) \quad (3)$$

155 where $\Delta C_{cum}/\Delta t$ is in units of $\mu\text{g C (g soil)}^{-1} (\text{day})^{-1}$. The size of the active and slow pools
156 adds up to the total amount of C in the soil (C_{soil}) (eqn 1); causing C_s to be determined by
157 the difference of the total soil C and the active pool C. This method of curve fitting also
158 has the assumption that changes in respiration rate over the course of an incubation are
159 due to changes in SOM lability.

160 A three-pool model with first-order kinetics has been found to effectively describe
161 SOC dynamics (Paustian et al., 1992); because of this we also fit our respiration rate
162 curves to a three-pool first order equation. We used the same rate equation as the two-
163 pool model (eqn. 3) and included a resistant pool which was estimated to be fifty percent
164 of the total soil C. The decomposition rate for the resistant pool was estimated using a
165 field mean residence time of 500 years and a Q_{10} adjustment of 2 for the different
166 incubation temperatures. This calculation was only done for comparative purposes and
167 parameter estimates and subsequent temperature sensitivity calculations were done with
168 the two-pool equation.

169 Best fit parameters (C_a , k_a and k_s) for the two-pool model were estimated using
170 non-linear regression of the CO_2 evolved with time in SAS v9.2 PROC NLIN with the
171 Gauss method (SAS Institute, Cary NC). The only restriction imposed on parameters is
172 that the values had to be greater than zero. There were three instances (N. Dakota native
173 grassland 15°C, Colorado native grassland 15°C, and Colorado cultivated 15°C) where
174 the slow pool decomposition rate were not positive and these three samples were not used

175 in subsequent analysis. The errors associated with model parameters are standard error of
176 the model determined in SAS NLIN.

177

178 *2.5 Pyrolysis-molecular beam mass spectrometry (py-MBMS)*

179 Chemical composition of SOM was characterized using py-MBMS. Details of the
180 analytical method are provided in Magrini et al. (2002) and Hoover et al. (2002). Briefly,
181 two subsamples (~0.1 g) from each field replicate sample were weighed in quartz boats
182 and pyrolyzed in a reactor consisting of a quartz tube (2.5 cm inside diameter) with
183 helium flowing through at 5 l min⁻¹ heated and maintained at 500°C. The molecular beam
184 system consisted of an Extrel™ Model TQMS C50 mass spectrometer for both
185 pyrolysis and combustion vapor analysis. Mass spectral data from m/z 20 to 500 were
186 acquired on a Teknivent Vector 2™ data acquisition system using 22 eV electron
187 impact ionization and programmed storage in a personal computer. Repetitive scans
188 (typically one 480 amu scan s⁻¹) were recorded during the evolution of a pyrolysis wave
189 from each soil sample.

190 Overall, 36 samples were collected from the field (6 sites × 3 field replicates × 2
191 land uses). Two aliquots from each of these 36 were analyzed by py-MBMS, resulting in
192 72 total spectra. Two spectra from the analyses of samples from Saskatchewan and one
193 from North Dakota were, however, excluded from further analysis due to data quality
194 issues, leaving a total of 69 spectra.

195 Multivariate analysis has proven to be an important tool for pattern recognition in
196 pyrolysis mass spectrometry (Schulten et al., 1988). Signal intensities from individual

197 spectra (m/z 20-500) were normalized to 100% total ion intensity (TII, the sum of the
198 intensity for each m/z), and reduced data sets (m/z 57-500) were used in the multivariate
199 analyses to omit the small mass units typical of water, CO₂ and other volatiles. We used
200 principal component analysis (PCA) to group samples by similarity with samples having
201 similar chemical compositions being tightly grouped in a PCA score plot, while samples
202 with dissimilar and heterogeneous compositions were more scattered. Eight principal
203 components and full cross validation were used to build the PCA model to determine
204 whether SOM composition could be grouped by site or land use. Partial least squares
205 (PLS) regression analysis was used to predict the dependent variables (i.e., the model
206 estimates for respiration rates during incubation at 15, 25, and 35°C) from our set of
207 independent variables (i.e., the signal intensities from the py-MBMS spectra). Two PLS
208 regressions were performed: one for the respiration rates early in the incubation when the
209 respiration is mostly from the active pool (day 5 of the incubation), and one for
210 respiration rates later in the incubation when the respiration is mostly from the slow pool
211 (day 225 of the incubation). Full cross-validation and Martens' uncertainty test were used
212 to determine statistically significant correlations between py-MBMS spectral data and the
213 biological properties of the samples. PLS analyses were performed iteratively to
214 determine which independent variables were significant based on Martens' uncertainty
215 test, and then subsequent PLS analyses were performed using only the significant
216 variables. This process was repeated until all independent variables were found to be
217 significant. All multivariate analyses were performed using the Unscrambler v.8.0
218 software package (CAMO Process AS, Oslo, Norway).

219

220 **3. Results**

221 *3.1. Site Comparisons*

222 Soil organic C (SOC) concentrations (0-20cm) ranged from 0.7 to 20.0% among
223 the six sites (Table 2). The Costa Rican soil is allophanic and consequently had much
224 higher SOC concentrations than the other five sites. SOC and total N concentrations
225 differed between land use within each site (except for Texas), with SOC and total N
226 concentrations decreasing after land-use conversion at Saskatchewan, N. Dakota,
227 Colorado, and Costa Rica.

228 Due to the large differences in SOC content among soils, the total amount of C
229 respired was normalized for the initial amount of SOC. After 588 days of incubation, the
230 cumulative amount of SOC respired was greater under warmer incubation temperatures
231 for all soils (Table 2). Overall there was a significant land use difference ($p < 0.001$) for
232 the four sites where the native grassland was converted to wheat cultivation, with the
233 native soils on average respiring significantly more SOC than the cultivated soils, both in
234 absolute terms and after being normalized for total SOC content. Cumulative soil
235 respiration differences were not statistically significant for the native forest to pasture
236 land use conversion in the Costa Rica and Brazil sites ($p = 0.127$).

237

238 *3.3. Temperature Sensitivity*

239 In all but two instances the labile fraction had a smaller Q_{10-q} value than the
240 resistant fraction (Table 2) with N. Dakota cultivated and Brazil native forest being the

241 two exceptions. For the N. Dakota cultivated and the Brazil native forest the Q_{10-q} values
242 for the resistant fraction were much lower than the values for the resistant fraction of the
243 other soils. The Costa Rica pasture site had the highest Q_{10-q} values overall (table 2).
244 There did not appear to be a trend with MAT and Q_{10-q} values for the sites. When sites
245 were treated as replicates and Q_{10-q} values were pooled there was a significant difference
246 in Q_{10-q} values for the labile and resistant fractions ($p = 0.0026$) with the Q_{10-q} for the
247 labile fraction being 1.95 ± 0.08 and the resistant fraction being 3.33 ± 0.04 . There was not
248 significant land use treatment effect ($p = 0.127$) or land use by SOM fraction effect ($p =$
249 0.338)

250

251 3.2. Curve Fitting

252 Respiration rates declined over time for all soils at all three incubation
253 temperatures with the two tropical sites, Costa Rica and Brazil, having a much more rapid
254 decline in respiration rates early in the incubation compared to the other four sites (Fig.
255 1). Respiration rates leveled off for all soils by the end of the incubation, but differences
256 in the respiration rates among the three incubation temperatures were still apparent at the
257 end of the incubation (Fig. 1).

258 Overall the two-pool equation fit the respiration data fairly well; an exception to
259 this was the Costa Rica native forest soil at 35°C where the two-pool equation
260 overestimated the respiration rate later in the incubation (Fig. 1). We also fit the data to a
261 three-pool first order equation, described by Paul et al. (2001), where the third pool was
262 50% of the total SOC and the turnover for the third pool was estimated to be 500 years.

263 We found that both the two and three-pool equations statistically fit the data equally well.
264 Although estimates of the active pool size and decomposition rate were not statistically
265 different between the two-pool and three-pool equation, the three-pool equation
266 consistently had a smaller active pool compared to the two-pool equation by an average
267 of 5% and the three-pool equation consistently had a higher decomposition rate of the
268 active pool compared to the two-pool equation by an average of 6% (Table 3). We used
269 the simpler two-pool model on all subsequent analysis, realizing that the resistant pool
270 contributes only slightly to the respiration of the slow pool because of its high mean
271 residence time.

272 The native grassland Colorado soil had the largest active pool, comprising
273 between 9.4 and 14.0% of total SOC, while the Costa Rica native forest soil had the
274 smallest active pool, comprising between 0.1% and 1.0% of total SOC (Table 4). In the
275 native grassland/cultivated sites (Saskatchewan, N. Dakota, Colorado, and Texas) the
276 native treatment had a significantly larger active pool than the cultivated treatment ($p =$
277 0.003). The size of the active pool at 25°C appeared to be negatively correlated with the
278 C:N ratio of the soil ($r^2 = 0.34$, $p = 0.045$ for all soils) (Fig. 2). The relationship between
279 the active fraction and C:N was improved when the Colorado cultivated and Brazil native
280 forest soils were removed ($r^2=0.84$, $p=0.0002$), which appeared to behave differently than
281 the other soils.

282 Since there were no restrictions placed on the size of the pools, pool sizes were
283 allowed to vary with temperature if it provided the best fit of the data. In all but four
284 instances (N. Dakota cultivated 35°C, Texas cultivated 25°C, Brazil native forest 25°C,

285 Costa Rica pasture 25°C) there was an increase in active pool size with temperature. For
286 the four soils where the active pool size declined with increasing temperature, it was
287 likely due to high initial respiration rates that quickly declined over time. The
288 decomposition rates of the slow pool increased with increasing incubation temperature
289 for all of the soils and the decomposition rate for the active pool increased with
290 increasing temperature in only half the instances (Table 4).

291

292 *3.4. Soil organic matter composition*

293 A semi-empirical quantification of SOM composition can be achieved by
294 assigning the relative intensity of individual mass signals from the py-MBMS to a set of
295 previously identified marker signals associated with several classes of compounds (Sorge
296 et al., 1993; Schulten, 1996). A large proportion of the mass spectra of each sample could
297 be classified, leaving only $6.7 \pm 2.5\%$ of TII unidentified (Table 5). The volatile fraction
298 (m/z 20-56) represented $50.9 \pm 4.5\%$ of the total signal intensity, dominated by m/z 44.
299 The largest classes of identified compounds were various N-containing compounds (11.9
300 $\pm 1.8\%$ of TII) and carbohydrates ($10.6 \pm 2.6\%$ of TII). The various N-containing
301 compounds are identified by Schulten (1996) as heterocyclic N-containing compounds,
302 but the heterocyclic N is thought to be formed by the pyrolysis process (Sharma et al.,
303 2003). Peptides, phenols and lignin monomers, alkyl aromatics, and lipids were also
304 significant contributors (in decreasing order), while sterols and lignin dimers each
305 contributed $< 1\%$ of TII (Table 5). The conversion of native grasslands to agriculture

306 causes a decline in carbohydrates, but the change in carbon compounds with conversion
307 from native forest to pasture was not consistent for the other two sites (Table 5).

308 In many of the sites there were differences in the abundance of chemical
309 compounds with py-MBMS between the native and cultivated/pasture treatments. Figure
310 3 illustrates the differences in chemical composition between the native and cultivated
311 soils for the Saskatchewan site, as an example. The Saskatchewan native grassland had a
312 greater abundance of the lower m/z compounds especially m/z 57 and 96 which are
313 associated with carbohydrates and m/z 67 and 81 which are associated with various
314 nitrogen compounds. The cultivated soil had a greater abundance of higher molecular
315 weight compounds, many of which are associated with lipids, lignin, and alkyl aromatics.

316 Principal components analysis was unable to significantly distinguish SOM
317 composition when all samples were analyzed together (Fig. 4). The first four principal
318 components combined explained only 43% of the variance. Taken together, PCA scores
319 of samples were not tightly grouped on the basis of site or land use, with the exception
320 that native forest and grassland samples appear to be more tightly grouped than cultivated
321 samples. While there was minor separation of samples by site along the PC 1 axis, SOM
322 composition from those sites was comparatively heterogeneous. However, significant
323 clustering was observed on the basis of land-use treatment when sites were analyzed
324 individually (data not shown), although there were no consistent trends across sites. The
325 PCA score results suggest that SOM composition in the Costa Rica samples was highly
326 homogeneous. PCA loadings were dominated by low molecular weight and odd-

327 numbered m/z fragments, which are likely derived from carbohydrate, amino acid and
328 peptide side-chains (data not shown).

329 Sequential application of the Martens' uncertainty test found a small number of
330 m/z values that were significantly correlated to respiration rates at day 5 of the incubation
331 (Table 6). PLS regression using this reduced set of independent variables was able to
332 explain 65% of the variance using the first four components. Respiration rates at day-5 of
333 the incubation were highly correlated with low molecular components associated
334 primarily with carbohydrates, peptides and various N compounds. Regression against
335 respiration rates at day 225 of the incubation resulted in more m/z values that were
336 significantly correlated (Table 6), but each correlation coefficient was smaller than those
337 for day 5. PLS regression of the reduced set of independent variables was able to explain
338 59% of the variance using the first four components. The additional m/z values were
339 generally higher molecular weights. Some were associated with phenols and lignin
340 monomers, lignin dimmers, and lipids, but many were not identified by Schulten (1996),
341 Hempfling & Schulten (1990), or Gillespie et al. (2009) as marker signals for specific
342 compounds classes.

343

344 **4. Discussion**

345 During our incubation the total amount of SOC respired increased with warmer
346 incubation temperatures for all twelve soils. Similar responses across the MAT range of
347 2 to 25.6°C (Table 2) could suggest that increasing global temperatures could affect soils
348 equally. The low respiration rates per unit SOC for the Costa Rica soils were likely due

349 to the unique mineral composition. Protective effects of amorphous aluminum, iron, and
350 allophanic material have been shown to contribute to the much higher SOM
351 concentrations found in these types of soils (Munevar and Wollum, 1977; Boudot et al.,
352 1986; Martin and Haider, 1986; McKeague, 1986). The characteristics that cause SOM
353 accumulation also likely cause their minimal loss of SOC upon incubation (Boudot et al.,
354 1988). However, the relative temperature response of the Costa Rica soil did not differ
355 from the other soils. Respiration rates declined over time in all the soils with the
356 differences in respiration rates between temperatures still apparent at the end of the
357 incubation. Correlations from PLS regression analysis of the py-MBMS products versus
358 the respiration rates could indicate that higher molecular weight compounds, and
359 therefore more resistant SOM, were being utilized in the later stages of the incubation.
360 These correlations support the first assumption of the Q_{10-q} method, that changes in
361 decomposition rates are driven in changes in SOM lability.

362 We found in most instances the resistant SOM was more temperature sensitive
363 than the labile SOM. We utilized the same method of determining temperature
364 sensitivity of SOM decomposition as described by Conant et al. (2008a) along with two
365 of the same sites (Colorado and Texas) and got similar results with an expanded set of
366 sites. The results found by us and Conant et al. (2008a) are consistent with multiple other
367 studies (Bosatta and Agren, 1999; Bol et al., 2003; Fierer et al., 2005; Knorr et al., 2005).
368 However, it is important to point out that there were two soils in our study that did not
369 follow this overall trend and in those two soils the resistant fraction had a lower Q_{10-q}
370 value than the labile fraction and the Q_{10-q} value of the resistant fraction was also much

371 lower than the Q_{10-q} value determined for the resistant fraction of the other soils. Our
372 results illustrate how a study that utilizes a single soil for determining temperature
373 sensitivity of SOM decomposition could possibly come to a different conclusion with
374 regards to SOM temperature sensitivity.

375 The estimated size of the active pool for all the sites ranged from 0.1 to 14.0% of
376 initial SOC (Table 4). This range for the active pool is slightly wider than results found
377 by Rey and Jarvis (2006), which ranged from 0.27% to 11.4%. Many other studies have
378 estimates that fall within these ranges (Collins et al., 2000; Haile-Mariam et al., 2000;
379 Cochran et al., 2007). Our wider range of estimates for the active pool is likely due to the
380 wide range of soils and incubation temperatures utilized. The two central sites, Texas
381 and Colorado, respired the most total SOC by the end of the incubation and had the
382 largest active pools, contrary to the comparison of forested tropical soils and dry forested
383 temperate soils by Trumbore (1993) in which the tropical soils were comprised of more
384 labile C in the upper 22 cm than temperate soils. The larger active pools in our temperate
385 soils and higher cumulative respiration could be due to the greater amount of N in the
386 soils, which indicates a higher proportion of proteinaceous constituents. Our results are
387 consistent with Thomsen et al. (2008), who found the C:N ratio to be an indicator of the
388 decomposability of the SOM in a soil, with soils having lower C:N ratios having greater
389 CO_2 evolution up to a certain threshold.

390 Temperature-induced changes in pool sizes could be an indication that warmer
391 temperatures may enable microbes to quickly decompose a larger portion of SOM.
392 Increased labile pool size with temperature evidenced in this study indicates that at

393 warmer temperatures SOM otherwise unavailable to microbes at lower temperatures
394 becomes available for decomposition. This could be attributed to shifts in microbial
395 community composition at different temperatures (Zogg et al., 1997; Zak et al., 1999),
396 changes in substrate use (Andrews et al., 2000; Schimel and Mikán, 2005), or the
397 overcoming of biochemical resistance of SOM by microbes (Conant et al., 2008b).

398 We had utilized different land use treatments at our sites as a way to compare
399 soils with the same MAT but varying amounts and types of SOM. For the four native
400 grassland/cultivated sites the native grassland treatments respired a greater amount of
401 total soil C and had larger active pools than their cultivated counterparts. The native
402 grassland soils had greater percentages of carbohydrates as found by py-MBMS, which
403 could be the reason for the greater respiration and larger active pool. Plante et al. (2009)
404 found that cultivation resulted in significant decreases in carbohydrates, peptides, and
405 phenols, also utilizing py-MBMS. Surprisingly, the decline in carbohydrates with
406 cultivation did not result in significant trends in Q_{10-q} values with land use. Our MAT
407 gradient did not produce consistent trends with regards to temperature sensitivity, also. It
408 may have been that differences in soil characteristics among sites were too great to
409 elucidate trends with MAT.

410 The size of our estimated active pool did increase with warmer incubation
411 temperatures for most soils indicating that there may be changes in microbial community
412 or function with temperature that should be considered in model incorporation. Our
413 results indicate that although individual soils may have varying apparent temperature
414 sensitivities for labile versus more resistant SOM depending on the inherent

415 characteristics of the soil, overall the more resistant SOM tends to be more temperature
416 sensitive. We were unable to link differences between inherent and apparent temperature
417 sensitivity of SOM to MAT, land use, or the chemical composition of SOM. This is an
418 area that requires more investigation and is necessary to better model temperature effects
419 on SOM decomposition.

420

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596 Table 1. Characteristics of the six locations, sampled in the spring of 2005, used in our long-term incubation

Site	GPS location	MAT/ MAP	Soil Taxonomy	Treatment	% Clay	Vegetation	Year converted to present land use	Citation
Indian Head, Saskatchewan ARGCN	50.533 N -103.517 W	2°C 421 mm	Udic Boroll	Native grassland	50	Grassland containing predominantly cool season grasses	1957	Campbell et al. 1997
				Cultivated	61	Spring-wheat-based rotations		
Mandan, North Dakota NGPRL	46.767 N -100.917 W	5°C 402mm	Typic Argiboroll	Native grassland	28	Warm mixed grass prairie	1984	Black & Tanaka 1997
				Cultivated	33	Continuous spring wheat		
Akron, Colorado CGPRS	40.150 N -103.150 W	9.2°C 420 mm	Aridic Paleustoll	Native grassland	23	Grassland with predominantly C ₄ grasses	1957	Halvorson et al. 1997
				Cultivated	28	Continuous wheat		
Vernon, Texas Waggoner Ranch	33.939 N -99.413 W	17°C 665 mm	Typic Paleustoll	Native grassland	31	Grassland with a mix of C ₄ and C ₃ grasses	N/A	Martin et al. 2003
				Cultivated	44	Continuous wheat with conventional tillage		
Alajuela, Costa Rica Alajuela Research Station	N/A	20°C N/A	Hydric Melanudand	Native forest	9	Tropical forest with predominantly C ₃ species	1979	N/A
				Pasture	12	C ₄ warm season grasses		
Rondônia, Brazil Nova Vida Ranch	10.168 S -62.824 W	25.6°C 2200mm	Paleudult & Kandiudult	Native forest	30	Open humid tropical forest with predominantly C ₃ species	1972	Cerri et al. 2004
				Pasture	25	C ₄ warm season grasses		

597 MAT, mean annual temperature; MAP, mean annual precipitation.

598 Table 2. Soil C and N content, C:N ratio, cumulative respiration after 588 days of incubation at 15°C, 25°C, and 35°C, and Q_{10-q}
 599 calculated for the 35/25°C comparison using the cumulative respiration after 588 days for the six sites and land-use types examined in
 600 the cross site comparison. (mean ± 1 standard deviation, n=4)

Site [†]	Land Use [‡]	SOC (%)	Total N (%)	C:N	Cumulative Respiration (% total soil C)			Q _{10-q}	
					15 °C	25 °C	35 °C	labile [§]	resistant [¶]
SK	NG	3.71 ± 0.09	0.358 ± 0.008	10.4	6.52 ± 0.37	10.40 ± 0.60	14.37 ± 0.23	1.6 ± 0.0	2.5 ± 0.2
SK	C	2.29 ± 0.09	0.204 ± 0.021	11.2	4.33 ± 0.14	7.73 ± 0.53	13.05 ± 0.50	2.0 ± 0.1	3.3 ± 1.3
ND	NG	3.24 ± 0.06	0.293 ± 0.004	11.0	3.93 ± 0.30	8.14 ± 0.68	15.03 ± 2.90	1.7 ± 0.1	2.7 ± 0.2
ND	C	2.80 ± 0.24	0.240 ± 0.004	11.7	5.17 ± 0.16	9.17 ± 0.44	11.32 ± 1.06	1.5 ± 0.1	1.3 ± 0.2
CO	NG	1.16 ± 0.08	0.135 ± 0.003	8.6	9.64 ± 0.65	17.93 ± 0.53	30.20 ± 2.39	1.5 ± 0.1	2.8 ± 0.3
CO	C	0.69 ± 0.02	0.100 ± 0.002	6.9	7.45 ± 0.25	13.09 ± 1.58	24.04 ± 2.58	2.2 ± 0.2	4.6 ± 1.4
TX	NG	1.12 ± 0.02	0.129 ± 0.006	8.7	9.60 ± 0.46	14.96 ± 1.10	24.36 ± 0.43	2.0 ± 0.2	3.4 ± 0.5
TX	C	1.02 ± 0.04	0.125 ± 0.004	8.1	8.62 ± 0.20	15.61 ± 0.68	23.22 ± 1.60	1.6 ± 0.1	3.1 ± 0.2
CR	NF	20.02 ± 0.62	1.660 ± 0.045	12.1	2.04 ± 0.10	4.21 ± 0.27	9.70 ± 0.53	1.9 ± 0.2	4.1 ± 0.4
CR	P	14.16 ± 0.19	1.045 ± 0.019	13.5	1.29 ± 0.05	2.68 ± 0.07	5.65 ± 0.17	2.9 ± 0.3	6.5 ± 0.4
BR	NF	1.06 ± 0.02	0.109 ± 0.006	9.7	5.72 ± 0.54	11.81 ± 0.74	19.50 ± 2.28	2.2 ± 0.3	1.4 ± 0.3
BR	P	1.41 ± 0.02	0.133 ± 0.011	10.6	6.76 ± 0.36	10.65 ± 0.45	21.60 ± 0.97	2.4 ± 0.4	4.2 ± 0.6

601 [†] SK, Saskatchewan; ND, North Dakota; CO, Colorado; TX, Texas; CR, Costa Rica; BR, Brazil

602 [‡] NG, native grassland; NF, native forest; C, cultivated; P, pasture

603 [§] Labile was considered the first 0.5% SOC respired in the incubation

604 [¶] Resistant was considered the last 0.5% SOC respired in the incubation

605 Table 3. Parameter estimates averaged across all soils and temperatures and fit estimates for the two-pool and three-pool models for
 606 the six sites and land use types examined in the cross site comparison.

Model	Active Pool ($\mu\text{g C g soil}^{-1}$)	Active Pool Decomp. Rate (day^{-1})	Slow Pool ($\mu\text{g C g soil}^{-1}$)	Slow Pool Decomp. Rate (day^{-1})	Resistant Pool ($\mu\text{g C g soil}^{-1}$)	Resistant Pool Decomp. Rate (day^{-1})	R_a^2
2 pool	1060	2.85E-02	42840	1.36E-04	.	.	0.83
3 pool	1002	3.03E-02	20948	3.10E-04	21950	1.76E-05	0.83
R_a^2 , adjusted r-square							

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610 Table 4. Size of the active pool and active and slow pool decomposition rates at 15°C, 25°C, and 35°C for the two-pool model for the
 611 six sites and land-use types examined in the cross site comparison. Error is model standard error.

Site †	Land use‡	Inc. Temp (°C)	Active Pool (% total soil C)	Active Pool Decomp. Rate (day ⁻¹)	Slow Pool Decomp. Rate (day ⁻¹)
SK	NG	15	3.7 ± 1.3	6.74E-03 ± 1.97E-03	5.60E-05 ± 2.24E-05
		25	5.3 ± 1.4	1.03E-02 ± 2.56E-03	1.01E-04 ± 3.22E-05
		35	6.7 ± 2.1	1.35E-02 ± 4.15E-03	1.72E-04 ± 5.83E-05
SK	C	15	2.1 ± 1.5	6.77E-03 ± 4.03E-03	4.33E-05 ± 2.52E-05
		25	2.6 ± 1.2	1.08E-02 ± 4.73E-03	9.55E-05 ± 2.70E-05
		35	4.2 ± 0.7	1.26E-02 ± 2.12E-03	1.73E-04 ± 1.89E-05
ND	NG	15	ND	ND	ND
		25	3.5 ± 0.7	8.68E-03 ± 1.48E-03	8.82E-05 ± 1.33E-05
		35	7.2 ± 2.5	6.72E-03 ± 1.89E-03	1.67E-04 ± 4.48E-05
ND	C	15	2.0 ± 0.5	1.03E-02 ± 2.35E-03	6.04E-05 ± 1.03E-05
		25	4.0 ± 1.4	7.58E-03 ± 2.26E-03	9.88E-05 ± 2.57E-05
		35	4.0 ± 1.1	1.17E-02 ± 2.94E-03	1.35E-04 ± 2.76E-05
CO	NG	15	ND	ND	ND
		25	9.4 ± 2.4	6.88E-03 ± 1.42E-03	1.85E-04 ± 4.46E-05
		35	14.0 ± 2.7	7.05E-03 ± 1.08E-03	3.31E-04 ± 5.38E-05
CO	C	15	ND	ND	ND
		25	3.1 ± 1.0	1.60E-02 ± 5.11E-03	1.91E-04 ± 2.94E-05
		35	7.3 ± 1.3	1.44E-02 ± 2.48E-03	3.52E-04 ± 4.00E-05
TX	NG	15	7.0 ± 2.7	5.43E-03 ± 1.66E-03	5.93E-05 ± 4.35E-05
		25	8.9 ± 2.0	9.02E-03 ± 1.80E-03	1.25E-04 ± 4.59E-05
		35	10.0 ± 1.2	1.38E-02 ± 1.60E-03	3.14E-04 ± 3.85E-05
TX	C	15	6.8 ± 3.9	5.54E-03 ± 2.51E-03	4.66E-05 ± 6.21E-05
		25	6.7 ± 2.8	1.57E-02 ± 6.53E-03	1.86E-04 ± 8.59E-05
		35	8.8 ± 1.4	1.87E-02 ± 2.99E-03	3.17E-04 ± 5.19E-05
CR	NF	15	0.1 ± 0.0	2.89E-01 ± 6.87E-02	4.63E-05 ± 7.20E-06
		25	0.6 ± 0.1	7.64E-02 ± 8.34E-03	6.57E-05 ± 8.07E-06
		35	1.0 ± 0.1	6.22E-02 ± 8.81E-03	1.63E-04 ± 1.41E-05

CR	P	15	0.7 ± 0.4	6.15E-03 ± 2.56E-03	9.63E-06 ± 6.10E-06
		25	0.5 ± 0.1	4.94E-02 ± 1.12E-02	3.96E-05 ± 7.75E-06
		35	3.2 ± 0.3	1.17E-02 ± 9.10E-04	4.83E-05 ± 6.15E-06
BR	NF	15	1.1 ± 0.2	5.61E-02 ± 1.14E-02	8.58E-05 ± 1.90E-05
		25	0.6 ± 0.1	1.11E-01 ± 2.15E-02	2.14E-04 ± 2.11E-05
		35	2.1 ± 0.3	4.92E-02 ± 9.52E-03	3.45E-04 ± 3.46E-05
BR	P	15	1.9 ± 0.3	3.59E-02 ± 6.00E-03	9.64E-05 ± 1.75E-05
		25	3.7 ± 0.7	2.23E-02 ± 4.23E-03	1.37E-04 ± 2.55E-05
		35	6.0 ± 0.7	2.91E-02 ± 3.76E-03	3.35E-04 ± 3.94E-05

612 † SK, Saskatchewan; ND, North Dakota; CO, Colorado; TX, Texas; CR, Costa Rica; BR, Brazil

613 ‡ NG, native grassland; NF, native forest; C, cultivated; P, pasture

614 ND values were not determined because the model calculated negative decomposition rates for the slow pool for these samples

615

616 Table 5. Proportions (%) of ion intensity attributable to carbohydrates, peptides, phenols, lignin dimmers, lipids, alkyl-aromatics,
 617 various N-containing compounds (VCN), sterols and the remaining unknown m/z fore each of the soils sampled. (mean \pm 1 standard
 618 deviation, n= 6)

Site ¹	Land use [‡]	m/z 20-56	Carbs	Peptides	Phenols & Lignin Monomers	Lignin dimers	Lipids	Alkyl aromatics	VNC	Sterols	Unknown
SK	NG	52.8	11.0	9.7	4.0	0.3	0.4	2.5	12.9	0.3	4.7
	C	59.0	8.0	8.2	3.0	0.6	0.8	1.9	9.6	0.6	6.6
ND	NG	52.8	11.0	9.1	3.9	0.3	0.5	2.2	13.1	0.3	5.1
	C	54.4	10.1	9.4	4.0	0.3	0.4	2.1	13.1	0.3	4.5
CO	NG	51.8	9.7	8.5	3.9	0.6	0.9	2.5	11.5	0.7	7.5
	C	48.5	8.1	8.4	4.0	1.0	1.8	2.4	9.2	1.1	11.5
TX	NG	53.4	9.1	8.8	3.3	0.6	0.9	2.4	11.0	0.8	7.3
	C	52.1	8.0	7.5	3.5	0.9	1.6	2.2	9.3	1.1	10.7
CR	NF	41.0	15.6	12.3	6.0	0.5	0.2	2.3	14.2	0.4	4.8
	P	47.5	15.4	11.2	4.6	0.3	0.1	2.1	13.3	0.3	3.5
BR	NF	50.0	9.6	9.4	4.2	0.6	0.9	2.1	12.8	0.6	7.5
	P	47.6	11.8	10.3	4.4	0.5	0.7	2.5	13.4	0.6	6.2
Mean		50.9 \pm 4.5	10.6 \pm 2.6	9.4 \pm 1.4	4.1 \pm 0.7	0.5 \pm 0.2	0.8 \pm 0.5	2.3 \pm 0.2	11.9 \pm 1.8	0.6 \pm 0.3	6.7 \pm 2.5

619 [†] SK, Saskatchewan; ND, North Dakota; CO, Colorado; TX, Texas; CR, Costa Rica; BR, Brazil

620 [‡] NG, native grassland; NF, native forest; C, cultivated; P, pasture

621 Table 6. Statistically significant m/z values from Martens' uncertainty test and sequential
 622 PLS regressions of chemical compositions against respiration rate at two different times
 623 of incubation ranked from largest to smallest value of regression coefficient. Where
 624 possible, m/z values marked with a compound class identifier based on classifications by
 625 Schulten (1996).

626

Day 5	97 (Pep), 98 (CH), 60 (CH), 73 (Pep), 111 (VNC), 58, 87 (Pep), 125 (VNC), 126 (CH), 59 (VNC), 74 (Pep), 123 (VNC), 138 (PLM), 101, 139 (VNC), 165, 75 (Pep)
Day 225	97 (Pep), 98 (CH), 83, 60 (CH), 80, 73 (Pep), 111 (VNC), 58, 125 (VNC), 87 (Pep), 126 (CH), 59 (VNC), 74 (Pep), 138 (PLM), 123 (VNC), 86, 72 (CH), 101, 139 (VNC), 168 (PLM), 143, 165, 428, 194 (PLM), 485, 479, 387, 446, 385, 373, 415, 332, 459

627 CH = carbohydrate; Pep = peptide; PLM = phenol and lignin monomer; LD = lignin

628 dimer; LP = lipid, alkane, alkene, fatty acid and n-alkyl ester; AA = alkyl aromatic; VNC

629 = various nitrogen containing compounds; S = sterol

630

631 **Figure Legend**

632 Figure 1. Decomposition rates over time at 15°C, 25°C, and 35°C for the twelve soils. NG
633 is native grassland and NF is native forest. Values represent means \pm SEM (n = 4).

634

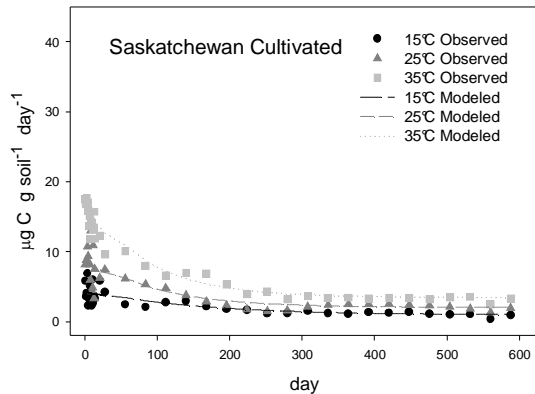
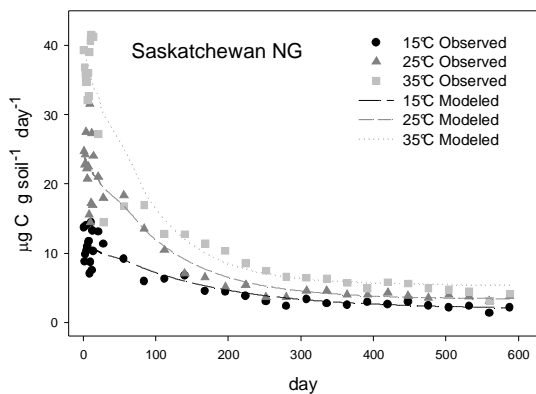
635 Figure 2. Correlation between the C to N ratio of the soil and the size of the active
636 fraction in percent of total SOC for the twelve soils sampled. All sites are represented by
637 closed circles except Colorado cultivated (CO-C), an unfilled square, and Brazil native
638 forest (BR-NF), an unfilled triangle.

639

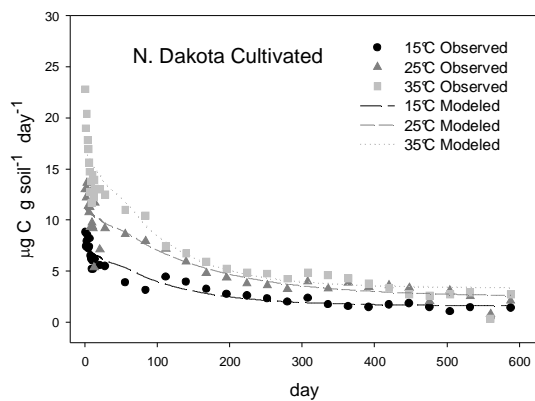
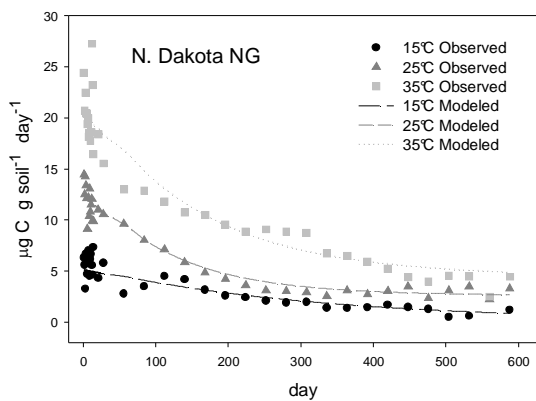
640 Figure 4. Normalized mass spectrum for spectral range $m/z=57-500$ from the pyrolysis
641 molecular beam mass spectrometry analysis of Saskatchewan native grassland (NG), N.
642 Dakota cultivated, and the difference between the two spectra. In the difference graph
643 m/z that are more abundant in the native grassland are positive and m/z that are more
644 abundant in the cultivated are negative. Spectra are means of 6 samples.

645

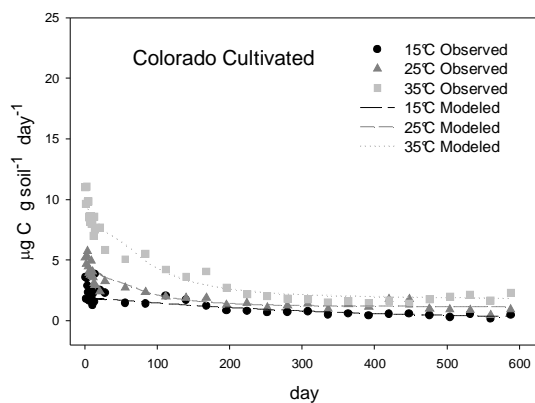
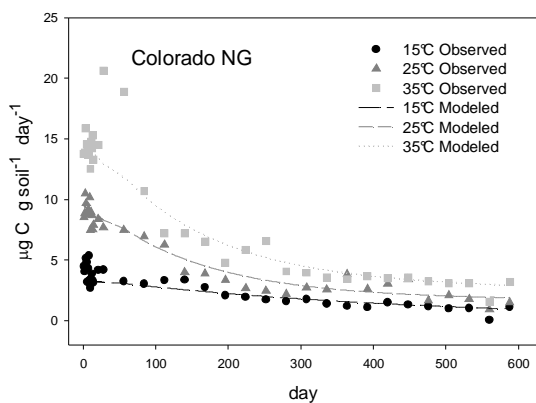
646 Figure 3. PCA scores of whole-soil samples analyzed by py-MBMS for the twelve soils
647 sampled. Closed symbols are native treatments, open symbols are cultivated or pasture
648 treatments.



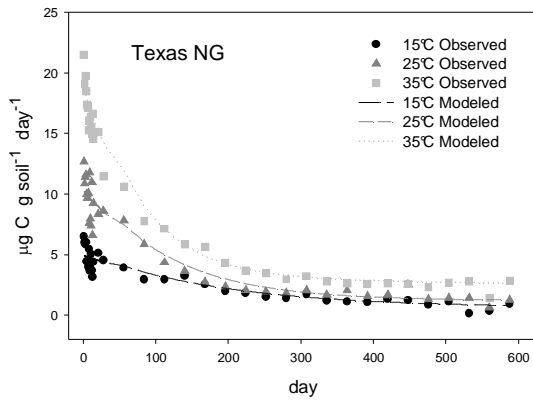
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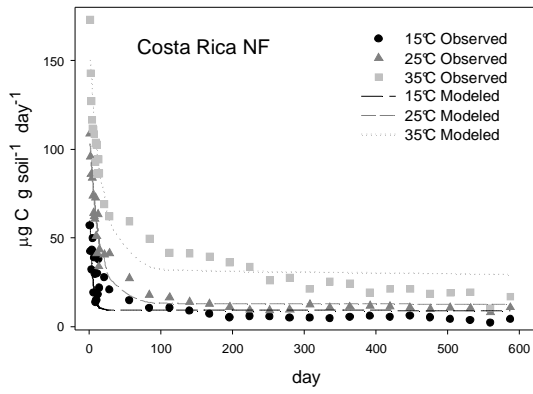
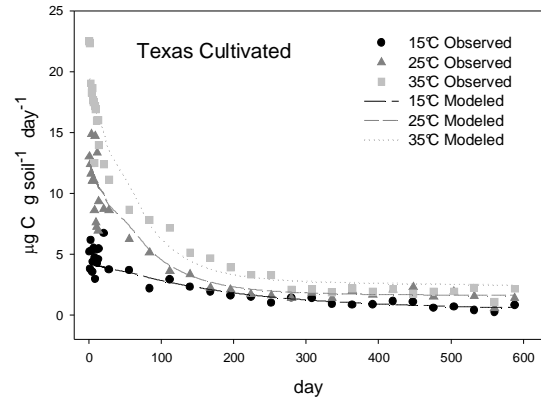
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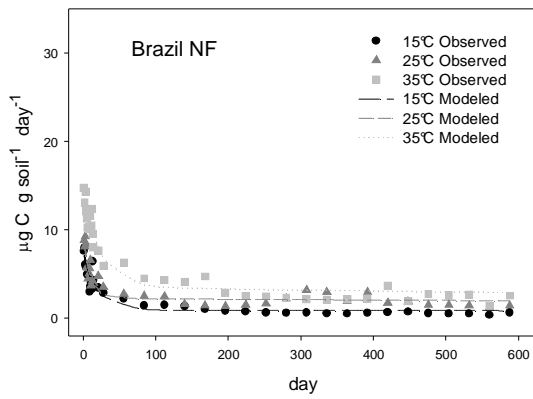
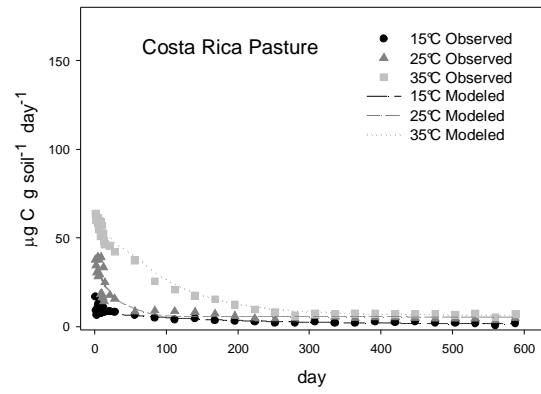
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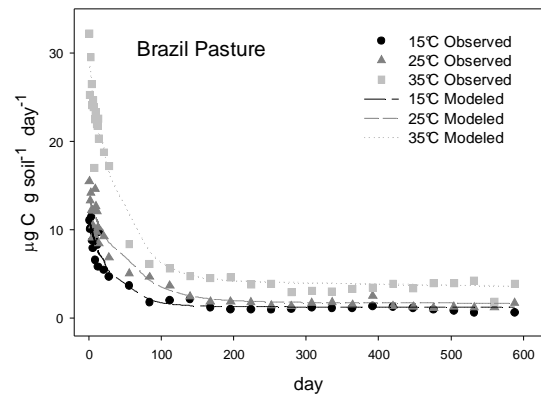
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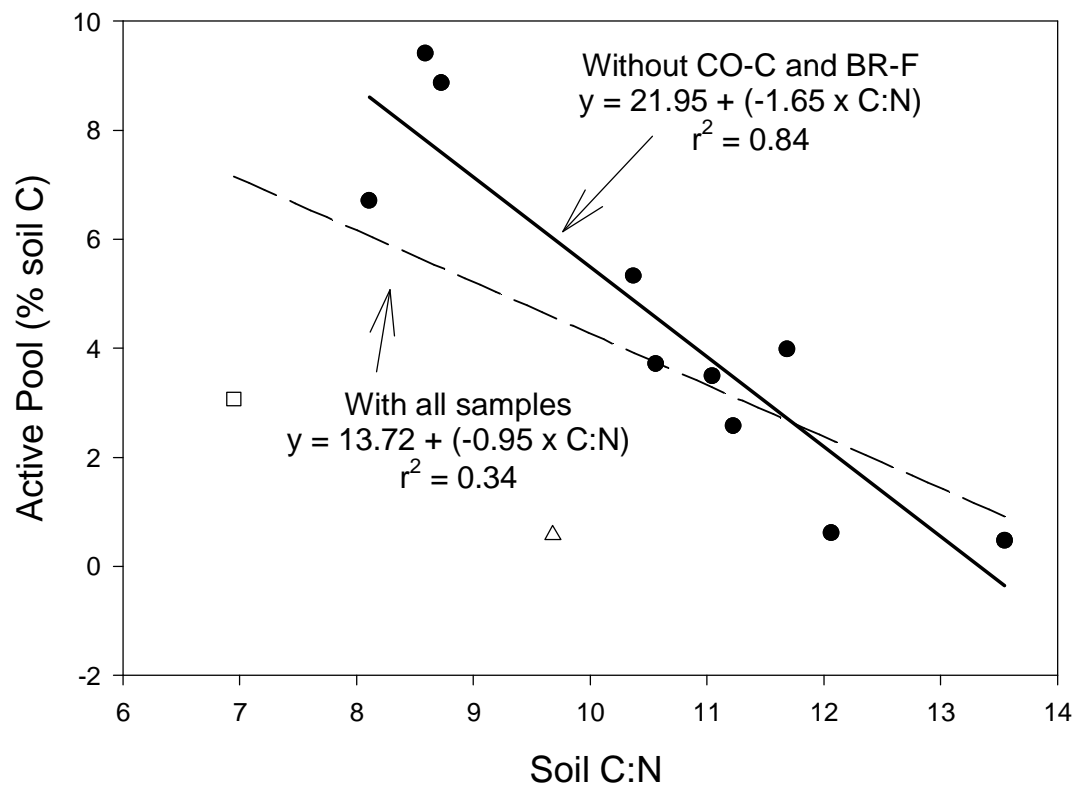
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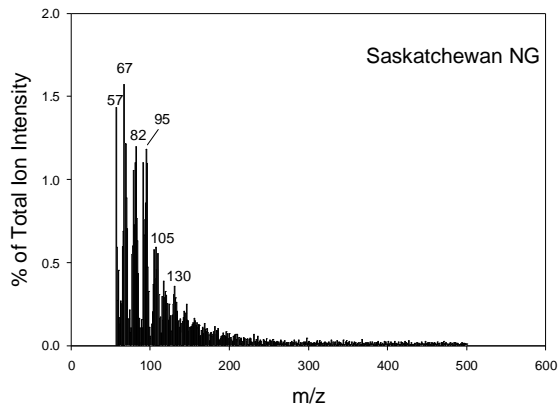
655 Figure 1.



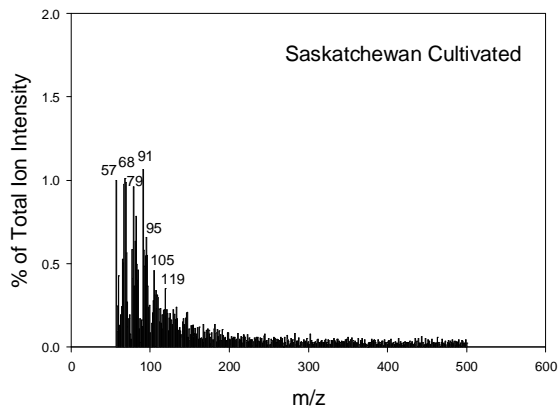
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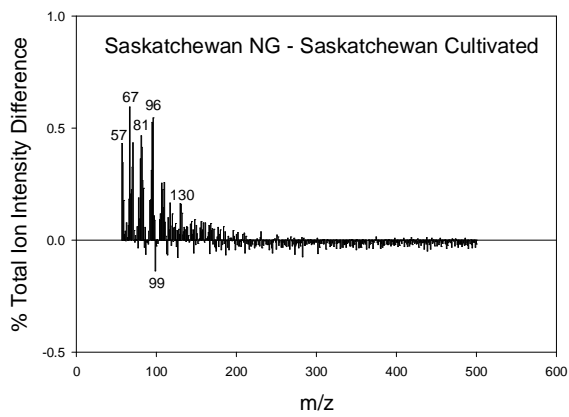
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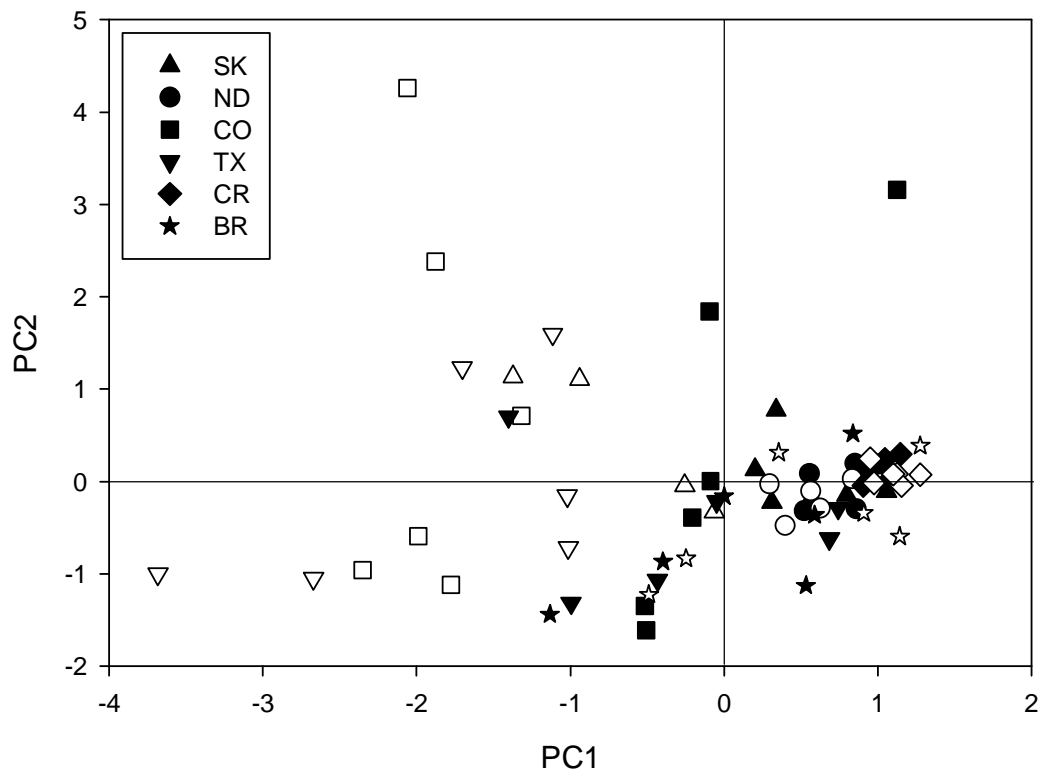


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661

662 Figure 3.



663

664 Figure 4.