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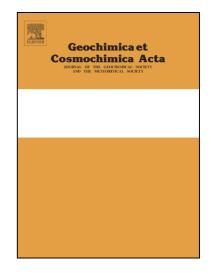


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1	Introducing global peat-specific temperature and pH calibrations based on
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3	
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¹⁶Sorbonne Universités, UPMC, Univ Paris 06, CNRS, EPHE, UMR 7619 METIS. 35 36 Paris, France 37 ¹⁷Department of Forest Sciences, University of Helsinki, Finland ¹⁸Université d'Orléans / CNRS /BRGM, ISTO, UMR 7327, Orléans, France 38 ¹⁹Department of Biology, University of Turku, Finland 39 ²⁰York Institute for Tropical Ecosystems, Environment Department, Wentworth Way 40 University of York, York, UK 41 42 ²¹Department of Geography, Durham University, Durham, UK 43 ²²Departamento de Edafoloxía e Química Agrícola, Universidade de Santiago de 44 Compostela, Santiago de Compostela, Spain 45 ²³Division of Marine Geology & Geophysics, University of Miami – RSMAS, Miami, 46 USA ²⁴Department of Chemistry, Claflin University, Orangeburg, USA 47 ²⁵Peatland Ecology Research Group (PERG), Centre for Northern Studies, Université 48 49 Laval, Quebec City, Canada ²⁶Department of Soil Science, University of São Paulo, Piracicaba, Brazil 50 51 52 *Corresponding author (david.naafs@bristol.ac.uk) 53 54 **Abstract** Glycerol dialkyl glycerol tetraethers (GDGTs) are membrane-spanning lipids from 55 56 Bacteria and Archaea that are ubiquitous in a range of natural archives and especially 57 abundant in peat. Previous work demonstrated that the distribution of bacterial 58 branched GDGTs (brGDGTs) in mineral soils is correlated to environmental factors 59 such as mean annual air temperature (MAAT) and soil pH. However, the influence of 60 these parameters on brGDGT distributions in peat is largely unknown. Here we 61 investigate the distribution of brGDGTs in 470 samples from 96 peatlands around the 62 world with a broad mean annual air temperature (-8 to 27 °C) and pH (3-8) range and 63 present the first peat-specific brGDGT-based temperature and pH calibrations. Our 64 results demonstrate that the degree of cyclisation of brGDGTs in peat is positively correlated with pH, pH = $2.49 \times CBT_{peat} + 8.07$ (n = 51, $R^2 = 0.58$, RMSE = 0.8) and 65 66 the degree of methylation of brGDGTs is positively correlated with MAAT, MAAT_{neat} (°C) = 52.18 x MBT_{5me}' -23.05 (n = 96, R² = 0.76, RMSE = 4.7 °C). 67

68	These peat-specific calibrations are distinct from the available mineral soil
69	calibrations. In light of the error in the temperature calibration (\sim 4.7 $^{\circ}$ C), we urge
70	caution in any application to reconstruct late Holocene climate variability, where the
71	climatic signals are relatively small, and the duration of excursions could be brief.
72	Instead, these proxies are well-suited to reconstruct large amplitude, longer-term
73	shifts in climate such as deglacial transitions. Indeed, when applied to a peat deposit
74	spanning the late glacial period (~15.2 kyr), we demonstrate that MAAT _{peat} yields
75	absolute temperatures and relative temperature changes that are consistent with those
76	from other proxies. In addition, the application of MAAT _{peat} to fossil peat (i.e.
77	lignites) has the potential to reconstruct terrestrial climate during the Cenozoic. We
78	conclude that there is clear potential to use brGDGTs in peats and lignites to
79	reconstruct past terrestrial climate.
80	
81	Keyword: GDGT, biomarker, peatland, calibration, lignite
82	
83	Highlights:
84	- Analysis of brGDGT distributions in global peat dataset
85	- Correlation of brGDGT distributions with peat pH and mean annual air temperature
86	- Development of peat-specific temperature and pH proxies

87	1. Introduction
88	Although reconstructions of terrestrial environments are crucial for the understanding
89	of Earth's climate system, suitable depositional archives (especially longer continuous
90	sequences) are rare on land. Peatlands and lignites (naturally compressed ancient peat)
91	are one exception and offer remarkable preservation of organic matter. Peats can be
92	found in all climate zones where suitable waterlogged conditions exist. Typical peat
93	accumulation rates are on the order of 1-2 mm/year (Gorham et al., 2003) and because
94	they exhibit minimal bioturbation (although roots might be present) they are widely
95	used as climate archives during the late Quaternary, predominantly the Holocene
96	(e.g., Barber, 1993; Chambers and Charman, 2004). Peat-based proxies include those
97	based on plant macrofossils, pollen, and testate amoebae (e.g., Woillard, 1978;
98	Mauquoy et al., 2008; Väliranta et al., 2012), inorganic geochemistry (e.g., Burrows
99	et al., 2014; Chambers et al., 2014; Hansson et al., 2015; Vanneste et al., 2015), (bulk)
00	isotope signatures (e.g., Cristea et al., 2014; Roland et al., 2015) and organic
01	biomarkers (e.g., Nichols et al., 2006; Pancost et al., 2007; Pancost et al., 2011;
02	Huguet et al., 2014; Zocatelli et al., 2014; Schellekens et al., 2015; Zheng et al.,
03	2015). Although these proxies can be used to provide a detailed reconstruction of the
04	environment and biogeochemistry within the peat during deposition, an accurate
05	temperature or pH proxy for peat is currently lacking (Chambers et al., 2012). This is
.06	particularly problematic because temperature and pH are key environmental
07	parameters that directly affect vegetation type, respiration rates, and a range of other
.08	wetland features (e.g., Lafleur et al., 2005; Yvon-Durocher et al., 2014). The aim of
.09	this paper is to develop peat-specific pH and temperature proxies for application to
10	peat cores as well as ancient peats from the geological record preserved as lignites.
11	We focus on using membrane-spanning glycerol dialkyl glycerol tetraether
12	(GDGT) lipids. In general, two types of GDGTs are abundant in natural archives such
13	as peats: 1) isoprenoidal (iso)GDGTs with sn-1 glycerol stereochemistry that are
14	synthesized by a wide range of Archaea, and 2) branched (br)GDGTs with sn-3
15	glycerol stereochemistry that are produced by Bacteria (see review by Schouten et al.,
16	2013 and references therein). A wide range of brGDGTs occur in natural archives
17	such as mineral soils and peat; specifically, tetra-, penta-, and hexamethylated
18	brGDGTs, each of which can contain 0, 1, or 2 cyclopentane rings (Weijers et al.,
19	2006b). In addition, recent studies using peat and mineral soils have demonstrated that
20	the additional methyl group(s) present in penta- and hexamethylated brGDGTs can

121	occur on either the α and/or $\omega5$ position (5-methyl brGDGTs) or the α and/or $\omega6$
122	position (6-methyl brGDGTs) (De Jonge et al., 2013; De Jonge et al., 2014).
123	brGDGTs are especially abundant in peat, in fact brGDGTs were first
124	discovered in a Dutch peat (Sinninghe Damsté et al., 2000). The concentration of
125	brGDGTs (as well as isoGDGTs) is much higher in the water saturated and
126	permanently anoxic catotelm of peat compared to the predominantly oxic acrotelm,
127	suggesting that brGDGTs are produced by anaerobic bacteria (Weijers et al., 2004;
128	Weijers et al., 2006a; Weijers et al., 2011), potentially members of the phylum
129	Acidobacteria (Weijers et al., 2009; Sinninghe Damsté et al., 2011; Sinninghe Damsté
130	et al., 2014). Although the exact source organism(s) are/is currently unknown, in
131	mineral soils (and potentially lakes) the distribution of bacterial brGDGTs is
132	correlated with mean annual air temperature (MAAT) and pH (Weijers et al., 2007;
133	Peterse et al., 2012; De Jonge et al., 2014; Loomis et al., 2014; Li et al., 2016). Over
134	the past decade ancient deposits of mineral soils (e.g., Peterse et al., 2014) and peat
135	(e.g., Ballantyne et al., 2010) have been used to reconstruct past terrestrial
136	temperatures.
137	Mineral soils differ from peat as the latter are normally water saturated,
138	consist predominantly of (partially decomposed) organic matter (the organic carbon
139	content of peat is typically> 30 wt.%), are typically acidic (pH 3-6), and have much
140	lower density. The combination of these factors means that peat becomes anoxic at
141	relatively shallow depths, whereas mineral soils are typically oxic. Indeed, Loomis et
142	al. (2011) showed that the brGDGT distribution in waterlogged soils is different from
143	that in dry soils and Dang et al. (2016) recently provided direct evidence of moisture
144	control on brGDGT distributions in soils. These differences suggest that microbial
145	lipids in peat might not reflect environmental variables, i.e. pH and temperature, in
146	the same way as they do in mineral soils.
147	Despite the high concentration of GDGTs in peats relatively few studies have
148	examined the environmental controls on their distribution in such settings (Huguet et
149	al., 2010; Weijers et al., 2011; Huguet et al., 2013; Zheng et al., 2015). Those studies
150	found that the application of soil-based proxies to peats can result in unrealistically
151	high temperature and pH estimates compared to the instrumental record. However,
152	owing to the small number of peats that have been studied to date as well as the lack
153	of peatland diversity sampled (the majority of peats sampled for these studies come
154	from temperate climates in Western Europe), the correlation of temperature and pH

155	with brGDGT distribution in peats is poorly constrained. Notably, the lack of tropical
156	peat brGDGT studies limits interpretations of brGDGT distributions in lignite
157	deposits from past greenhouse climates (Weijers et al., 2011).
158	Here we compare brGDGT distributions in a newly generated global data set
159	of peat with MAAT and (where available) in situ peat pH measurements. Our aim is
160	to gain an understanding of the impact of these environmental factors on the
161	distribution of brGDGTs in peat and develop for the first time peat-specific
162	temperature and pH proxies that can be used to reconstruct past terrestrial climate.
163	
164	2. Material and methods
165	2.1 Peat material
166	We generated a collection of peat comprising a diverse range of samples from around
167	the world (Fig. 1). In total, our database consists of 470 samples from 96 different
168	peatlands. In order to assess the variation in brGDGT distribution within one location,
169	where possible we determined the brGDGT distribution in multiple horizons from
170	within the top 1m of peat (typically representing several centuries of accumulation)
171	and/or analyzed samples taken at slightly different places within the same peatland. A
172	peat deposit typically consists of an acrotelm and catotelm, although marked
173	heterogeneity can exist even over short distances (Baird et al., 2016). The acrotelm is
174	located above the water table for most of the year and characterized by oxic
175	conditions and active decomposition. The acrotelm overlies the catotelm, which is
176	permanently waterlogged and characterized by anoxic conditions and very slow
177	decomposition. Our dataset spans those biogeochemical gradients (e.g. acro/catotelm).
178	Variations in peat accumulation rates differ between sites, implying that the ages of
179	the brGDGT-pool might differ.
180	Our database includes peats from six continents and all major climate zones,
181	ranging from high latitude peats in Siberia, Canada, and Scandinavia to tropical peats
182	in Indonesia, Africa, and Peru (Fig. 2). It covers a broad range in MAAT from -8 to
183	27 °C. Although most samples come from acidic peats with pH <6, the dataset
184	includes several alkaline peats and overall our dataset spans a pH range from 3 to 8.
185	All samples come from freshwater peatlands, except for the one from the Shark River
186	peat (Everglades, USA) that is marine influenced. Unsurprisingly, given their global
187	distribution, the peats are characterized by a wide variety of vegetation, ranging from

188	Sphagnum-dominated ombrotrophic peats that are abundant in high-latitude and
189	temperate climates to (sub)tropical peats dominated by vegetation such as Sagittaria
190	(arrowhead) and Cyperaceae (sedge), and forested tropical peatlands.
191	
192	2.2 Environmental parameters
193	The distribution of brGDGTs was compared to MAAT and in situ pH. MAAT was
194	obtained using the simple bioclimatic model PeatStash, which provides surface air
195	temperatures globally with a 0.5 degree spatial resolution (for details, see Kaplan et
196	al., 2003; Gallego-Sala and Prentice, 2013). The temperature data in PeatStash is
197	obtained by interpolating long-term mean weather station climatology (temperature,
198	precipitation and the fraction of possible sunshine hours) from around the world for
199	the period 1931–1960 (Climate 2.2 data are available online http://www.pik-
200	potsdam.de/~cramer/climate.html). Crucially, mean annual temperatures in peat are
201	similar to MAAT, assuming that the peat is not snow-covered for long periods of time
202	(McKenzie et al., 2007; Weijers et al., 2011). The temperature at the top surface of
203	(high-latitude) peat can differ from the MAAT due to insolation by snow during
204	winter and intense heating during summer. Despite this, the seasonal temperature
205	fluctuations in peat are dampened at depth as temperatures converge to MAAT
206	(Hillel, 1982; Laiho, 2006; McKenzie et al., 2007; Weijers et al., 2011). We assume
207	that all peat horizons experienced MAAT (the only data available on a global basis).
208	This is likely an oversimplification that introduces some additional uncertainty in our
209	calibration.
210	Where available, pH data were obtained from measured values reported in the
211	literature or our measurements during sampling. For peats, pH cannot be determined
212	using dried material, as is normally done for soils (Stanek, 1973). Accurate pH
213	measurements can only be obtained from in situ measurements, especially for
214	groundwater-fed wetlands, and these are not available for all locations.
215	
216	2.3 Lipid extraction
217	For the majority of samples (>430 out of 470), between 0.1 and 0.5 g of dried bulk
218	peat were extracted with an Ethos Ex microwave extraction system with 20 mL of a
219	mixture of dichloromethane (DCM) and methanol (MeOH) (9:1, v/v) at the Organic
220	Geochemistry Unit (OGU) in Bristol. The microwave program consisted of a 10 min
221	ramp to 70 °C (1000 W), 10 min hold at 70 °C (1000 W), and 20 min cool down.

222	Samples were centrifuged at 1700 rounds per minute for 3 to 5 min and the
223	supernatant was removed and collected. 10 mL of DCM:MeOH (9:1) were added to
224	the remaining peat material and centrifuged again after which the supernatant was
225	removed and combined with the previously obtained supernatant. This process was
226	repeated 3 to 6 times, depending on the amount of extracted material, to ensure that
227	all extractable lipids were retrieved. The total lipid extract (TLE) was then
228	concentrated using rota-evaporation. An aliquot of the TLE (typically 25%) was
229	washed through a short (<2 cm) silica column using DCM:MeOH (9:1) to remove any
230	remaining peat particles. The TLE was dried under a gentle nitrogen flow and then re-
231	dissolved in hexane/iso-propanol (99:1, v/v) and filtered using 0.45 μm PTFE filters.
232	A small number of peats were extracted using different methods and either the
233	TLE or polar fraction was analyzed for GDGTs (see Table S1). Samples from the
234	Kyambangunguru peat in Tanzania were extracted using the Bligh-Dyer protocol.
235	Previous work on peat demonstrated that the brGDGT distribution is similar using
236	Bligh-Dyer extraction as Soxhlet extraction (Chaves Torres and Pancost, 2016). The
237	TLE was cleaned over a short Si column at the OGU in Bristol. Both cleaned TLE
238	and polar fractions were re-dissolved in hexane/iso-propanol (99:1, v/v) and filtered
239	using 0.45 μm PTFE filters.
240	
241	2.4 Analytical methods
242	All samples were analyzed for their core lipid GDGT distribution by high
243	$performance\ liquid\ chromatography/atmospheric\ pressure\ chemical\ ionisation-mass$
244	spectrometry (HPLC/APCI-MS) using a ThermoFisher Scientific Accela Quantum
245	Access triplequadrupole MS. Normal phase separation was achieved using two ultra-
246	high performance liquid chromatography silica columns, following Hopmans et al.
247	(2016). Crucially this method allows for the separation of the 5- and 6-methyl
248	brGDGT isomers. Injection volume was 15 $\mu L,$ typically from 100 $\mu L.$ Analyses were
249	performed using selective ion monitoring mode (SIM) to increase sensitivity and
250	reproducibility (<i>m/z</i> 1302, 1300, 1298, 1296, 1294, 1292, 1050, 1048, 1046, 1036,
251	1034, 1032, 1022, 1020, 1018, 744, and 653). The results were integrated manually
252	using the Xcalibur software. Based on daily measurements of an in-house generated
253	peat standard, analytical precession (σ) over the 12 months during which the data
254	were analyzed is 0.01 for the proxy index we define below (MBT _{5me} ', eq. 2).

- 256 2.5 Proxy calculation
- 257 Guided by previous studies we used a range of proxies to express ratios of different
- 258 GDGTs and the nomenclature of De Jonge et al. (2014) (Fig. 1).

259

eq. (1) MBT

$$=\frac{(Ia+Ib+Ic)}{(Ia+Ib+Ic+IIa+IIa'+IIb+IIb'+IIc+IIIc'+IIIa+IIIa'+IIIb+IIIb'+IIIc+IIIc')}$$

- The original methylation of branched tetraether (MBT) index compared the relative
- abundance of tetramethylated brGDGTs (compounds Ia-Ic) to that of penta-
- 262 (compounds IIa-IIc') and hexamethylated (compounds IIIa-IIIc') brGDGTs that have
- one or two additional methyl groups (Weijers et al., 2007). It was recently discovered
- that the additional methyl groups in penta- and hexamethylated brGDGTs can also
- occur at the C6 position (6-methyl brGDGTs, indicated by a prime symbol; e.g.
- brGDGT-IIa'): the 6-methyl penta- and hexamethylated brGDGTs (De Jonge et al.,
- 267 2013). Excluding the 6-methyl brGDGTs from the MBT index resulted in the
- MBT_{5me}' index. In the global soil database the application of MBT_{5me}' led to an
- improved correlation with temperature (De Jonge et al., 2014).

$$eq. (2) MBT'_{5ME} = \frac{(Ia + Ib + Ic)}{(Ia + Ib + Ic + IIa + IIb + IIc + IIIa)}$$

- 270 In addition to different number of methyl groups, brGDGTs can contain up to two
- 271 cyclopentane moieties (e.g., brGDGT-Ib and -Ic). CBT' is a modified version of the
- original cyclisation of branched tetraether (CBT) index (Weijers et al., 2007) and in
- soils CBT' has the best correlation with pH (De Jonge et al., 2014):

$$eq.(3) \ CBT' = \log \left(\frac{Ic + IIa' + IIb' + IIc' + IIIa' + IIIb' + IIIc'}{Ia + IIa + IIIa} \right)$$

- The isomer ratio of 6-methyl brGDGTs (IR_{6me}) reflects the ratio between 5- and 6-
- 275 methyl brGDGTs (Yang et al., 2015) with low (high) values indicative of a
- dominance of 5-methyl (6-methyl) brGDGTs:

$$eq.(4) IR_{6me}$$

$$=\left(\frac{IIa'+IIb'+IIc'+IIIa'+IIIb'+IIIc'}{IIa+IIa'+IIb+IIIb'+IIIc+IIIc'}\right)$$

- 278 The isomerization of branched tetraethers (IBT) is related to IR_{6me} but reflects the
- isomerization of brGDGT-IIa and -IIIa only (Ding et al., 2015):

$$eq. (5) IBT = -\log\left(\frac{IIa' + IIIa'}{IIa + IIIa}\right)$$

- The branched versus isoprenoidal tetraether (BIT) index (Hopmans et al., 2004)
- reflects the relative abundance of the major bacterial brGDGTs versus a specific
- archaeal isoGDGT, crenarchaeol (Fig. 1), produced by *Thaumarchaeota* (Sinninghe
- 283 Damsté et al., 2002):

$$eq. (6) BIT = \frac{Ia + IIa + IIa' + IIIa + IIIa'}{Ia + IIa + IIa' + IIIa + IIIa' + cren.}$$

- Finally, the isoprenoidal over branched GDGT ratio (R_{i/b}), related to the BIT index,
- 285 records the relative abundance of archaeal isoGDGTs over bacterial brGDGTs (Xie et
- 286 al., 2012).

$$eq. (7) R_{i/b} = \frac{\sum isoGDGTs}{\sum brGDGTs}$$

- 288 2.6 Statistical methods
- 289 Temperature and pH calibrations were obtained using the average proxy value for
- 290 each peat and Deming regressions. The software we used was RStudio
- 291 (RStudio Team, 2015) and Method Comparison Regression (MCR) package
- 292 (Manuilova et al., 2014), which are freely available to download¹. The Rscript and
- 293 data are available in the appendices.
- Deming regressions differ from simple linear regression, which so far have
- been used in brGDGT proxy calibrations, as they account for error in the data on both
- the x- (e.g., proxy) and y-axis (e.g., environmental variable) (Adcock, 1878).
- We used the average proxy value for each peat to calculate Deming regressions,
- 298 calibration errors (RMSE, see below), and calibration coefficients of determination
- 299 (R^2). The errors associated with proxy measurements (e.g. MBT_{5me}') and
- and environmental parameters (MAAT/pH) are independent and assumed to be normally
- distributed. To calculate a Deming regression, the ratio of variances (δ) must be
- calculated. For MAAT we took a standard deviation (σ) of 1.5 °C based on the
- estimated mean predictive error of up to 1.4 °C for mean temperature in a similar
- dataset (New et al., 1999). For pH we took a standard deviation of 0.5 based on the
- 305 average reported heterogeneity in pH for the peatlands used in the database (see
- Supplementary Table 1). For MBT_{5me}', CBT', and CBT_{peat} we calculated the average

¹ https://www.rstudio.com and https://cran.r-project.org/web/packages/mcr/index.html

307	standard deviation of each proxy from the entire peat data set (0.05, 0.25, and 0.2,
308	respectively). This results in a ratio of variances of 0.0011 for the MBT _{5me} '/MAAT
309	calibration and 0.25 and 0.16 for the pH calibration based on CBT' and CBT _{peat} ,
310	respectively. Residuals were calculated for the full dataset and using
	eq. (8) $Residual_y = y_{observed} - y_{predicted}$
311	The root mean square error (RMSE) for y, the predictive error for the
312	environmental parameter of interest (MAAT or pH), was calculated for the average
313	proxy value of each peat and using
	eq. (9) $RSME_y = \sqrt{\frac{\sum_{x=1}^{n} (y_{x,observed} - y_{x,predicted})^2}{n}} \times \frac{n}{df}$
314	Where df stands for degrees of freedom, which in this case is n-1.
315	
316	3. Results
317	Although we did not calculate concentrations, based on changes in signal intensity the
318	relative abundance of GDGTs was always higher at depth compared to the top (\sim 0 $-$
319	20) cm of peat. BIT indices (eq. 6) range between 0.75 and 1, but 99% of the samples
320	have a BIT value \geq 0.95. Similarly, $R_{i/b}$ ratios are typically <0.5. Only three samples
321	from the São João da Chapada peat in Brazil have a $R_{i/b}$ ratio >1 .
322	The majority of brGDGTs are tetramethylated and 5-methyl penta- and
323	hexamethylated brGDGTs. The most abundant brGDGTs in peat are brGDGT-Ia and
324	IIa. By extension, the IR _{6me} ratio (eq. 4) is low. brGDGTs containing cyclopentane
325	moieties are much less abundant than acyclic brGDGTs and brGDGT-IIIb(') and -
326	IIIc(') are either below detection limit or present at trace abundances ($\leq 1\%$ of total
327	brGDGTs). Indeed, three brGDGTs dominate the entire global dataset: tropical peats
328	contain almost exclusively brGDGT-Ia (up to 99% of total brGDGTs), whereas in
329	high-latitude peats brGDGT-IIa and -IIIa are dominant (Fig. 3).
330	
331	4. Discussion
332	The observation that $R_{\text{i/b}}$ ratios are low in most peats is consistent with previous
333	observations that bacterial brGDGTs dominate over archaeal isoprenoidal GDGTs in
334	peat (Schouten et al., 2000; Sinninghe Damsté et al., 2000; Pancost et al., 2003) and
335	mineral soils (Hopmans et al., 2004).
336	

337	4.1 Shallow vs deep GDGT distributions
338	The apparent increase in GDGT abundance with depth is consistent with previous
339	observations in peatlands (Weijers et al., 2004; Peterse et al., 2011) and reflects the
340	combined effects of preferential GDGT production in anaerobic settings and the
341	accumulation of fossil GDGTs over time at depth (Liu et al., 2010; Weijers et al.,
342	2011).
343	In one high-latitude peat (Saxnäs Mosse, Sweden) the distribution of both
344	intact polar lipids (compounds still containing a polar head groups) and core
345	brGDGTs (compounds having lost their polar head group) differed between the acro-
346	and catotelm and brGDGT abundances were much higher in the latter (Weijers et al.,
347	2009; Peterse et al., 2011). Based on these results Peterse et al. (2011) speculated that
348	microbial communities differed between the oxic acrotelm and anoxic catotelm. As
349	oxygen content can influence cellular lipid composition of bacteria, Huguet et al.
350	(2010) speculated that oxygen availability could be one of the factors directly
351	influencing the brGDGT synthesis by bacteria in peat, as opposed to influencing the
352	type of source organism(s). Studies from lakes also suggested that changes in lake
353	oxygenation state can influence the brGDGT distribution (Tierney et al., 2012;
354	Loomis et al., 2014).
355	Our dataset consists of a mixture of surface (0-15 cm) and deeper samples
356	that extend through the top one meter of peat. For the majority of peats there is no
357	detailed information available on water table depths and location of the acro/catotelm
358	boundary. Nonetheless, to provide a first order assessment on whether there is a
359	systematic and significant difference in core brGDGT distribution between the upper
360	(assumed to be generally oxic) and underlying anoxic peat, we compared the relative
361	abundance of the three most abundant brGDGTs (Ia, IIa, and IIIa) in the shallow
362	surface peat (top 15 cm) with that of the deep peat below 15 cm (Fig. 3), although we
363	acknowledge that this is likely an oversimplification.
364	There are some differences. In general the relative abundance of brGDGT-Ia
365	is slightly higher in the top 15 cm of a peat compared to the peat below 15 cm,
366	especially when its abundance is < 60%. Overall, however, the distributions plot
367	along the 1:1 line, indicating that there is no systematic difference in brGDGT
368	distribution between the (assumed) oxic surface and the peat below 15 cm (likely
369	anoxic). This does not preclude differences in brGDGT production between oxic and
370	anoxic conditions, but this appears to be primarily expressed via greater production of

371	brGDGTs under anoxic conditions as demonstrated by the higher abundance of
372	GDGTs across the acro/catotelm boundary (Weijers et al., 2006a). These results
373	provide indirect evidence that oxygen availability does not significantly impact the
374	degree of methylation of (core) brGDGTs. One possible explanation for why oxygen
375	availability does not affect distributions is that brGDGTs could be predominantly
376	produced by anaerobes throughout the peat, in low abundance in anaerobic
377	microenvironments in shallow peat and in high abundance in the anaerobic catotelm.
378	Several (high-latitude) peats, however, do appear to exhibit strong variations
379	between deep and shallow sections of the peat. The down core records from Stordalen
380	(Sweden) and Andorra (S. Patagonia), for example, are characterized by a large and
381	abrupt shift in brGDGT distribution at depth (Fig. 4). The MBT _{5me} ' indices recorded
382	at the very top of these high-latitude peats are between 0.8 and 0.6, as high as those
383	found in mid-latitude and subtropical peats, but decrease to values between 0.2 and
384	0.4 below ~30 cm. Peats from temperate climates (e.g. Walton moss, UK) and the
385	tropics (e.g. Sebangau, Indonesia) display much smaller or no change in brGDGT
386	distribution with depth (Fig. 4 and 5). It appears that this offset in brGDGT
387	distribution with depth is amplified in high-latitude peats. This is consistent with
388	previous studies that indicated a difference in brGDGT-distribution between the acro-
389	and catotelm in a high-latitude peat from southern Sweden (Weijers et al., 2009;
390	Peterse et al., 2011).
391	We argue that the high MBT_{5me} ' values at the top of these high-latitude peats
392	are heavily biased towards summer temperatures. At these settings winter
393	temperatures are often below freezing for a prolonged period, likely causing bacterial
394	growth and GDGT production to slow down significantly. Summer temperatures are
395	much higher (e.g. mean warmest month temperature at Stordalen is around 13 °C), in-
396	line with the observed relatively high MBT _{5me} ' values (e.g., 0.6-0.7 at Stordalen, see
397	Figure 4). Deeper in the peat, seasonal temperature fluctuations are much less
398	pronounced and temperatures rapidly converge to the MAAT (Vitt et al., 1995; Laiho,
399	2006; McKenzie et al., 2007; Weijers et al., 2011), likely accounting for the lower
400	MBT _{5me} ' values in the deeper peat horizons. Moreover, the greater production of
401	GDGTs in the anaerobic part of the peat will cause GDGT-based temperatures to
402	rapidly converge on the deep peat growth temperature, overprinting the seasonal
403	summer bias of fossil GDGTs synthesized at the surface.

404	This effect is diminished in temperate and especially tropical peatlands from
405	around sea level, which we attribute to the lack of a preferred growing season in
406	settings with smaller seasonal temperature ranges. In such settings temperatures are
407	less frequently (or never) below freezing and brGDGT production in the top of the
408	peat likely occurs for all or most of the year, such that GDGTs produced in both the
409	shallow and deeper part of the peat record MAAT. This hypothesis needs further
410	testing but indicates that 1) brGDGT production may be biased towards the warm
411	season in the upper part of high-latitude/altitude peats; 2) care has to be taken when
412	interpreting brGDGT-based trends in the top of such peats; and 3) the temperature
413	signal in such peats is imparted at depth, such that downcore GDGT variations in
414	ancient peat archives could potentially be temporally offset (precede) the climate
415	events that caused them. However, as brGDGTs in long peat cores, and by extension
416	ancient lignites (fossilized peats), are dominated by production at depth where
417	temperature equals MAAT (see section 2.2) it is very unlikely that temperatures
418	obtained from these archives are seasonally biased.
419	In the remainder of this work, for high-latitude peats that show a clear offset
420	between the top and deeper part of the peat we use only the average GDGT
421	distribution from below 20 cm, as the majority of change appears to occur in the top
422	20 cm. For the other peats we retain all data from the upper 1 m, not differentiating
423	between the acro- and catotelm. To generate the temperature and pH calibrations we
424	use the average brGDGT distribution for each peatland. For peats where multiple
425	samples were analyzed, error bars indicate the deviation (1 σ) from the average.
426	
427	4.2 Influence of temperature and pH on brGDGTs in peats
428	It is well established that in soils and lakes, environmental conditions such as
429	temperature and pH are highly correlated with the brGDGT distribution (e.g., Weijers
430	et al., 2007; Peterse et al., 2012; Schoon et al., 2013; De Jonge et al., 2014; Loomis et
431	al., 2014; Xiao et al., 2015; Li et al., 2016). In the following sections we investigate
432	the influence of these parameters on the brGDGT distribution in peat using the
433	average proxy value (e.g. MBT _{5me} ') for each peatland.
434	
435	4.2.1 Influence of peat pH on brGDGT distribution
436	Weijers et al. (2007) demonstrated that in a global mineral soil database the degree of
437	cyclisation of brGDGTs is correlated to pH, with a higher fractional abundance of

438	brGDGTs that contain cyclopentane moieties in soils with a higher pH. Following the
439	discovery of 6-methyl brGDGTs (De Jonge et al., 2013), it was shown that the degree
440	of isomerization of brGDGTs, the ratio of 6-methyl versus 5-methyl brGDGTs, is also
441	correlated to soil pH, with a higher fractional abundance of 6-methyl brGDGTs in
442	soils with a higher pH (De Jonge et al., 2014; Xiao et al., 2015). Owing to the limited
443	pH range of the few peats used to study brGDGTs so far and because all of these
444	studies pre-date the recent analytical advances that allow for the separation of 5- and
445	6-methyl brGDGTs, it is unknown whether pH has an influence on brGDGTs in peats
446	or whether the dependence is similar to that found in soils. Our peat database spans a
447	pH range from 3 to 8, similar to that of the soil database, allowing us to assess the
448	influence of pH on the brGDGT distribution in such settings.
449	Although pH measurements are only available in 51 out of 96 peats, our
450	results indicate that 6-methyl brGDGTs are present at either only trace abundances
451	(IR _{6me} $<$ 0.1) or are absent in acidic peats with pH $<$ 5.4 (Fig 6). Higher ratios occur in
452	peats with higher pH. The highest ratio (0.58) occurs in the marine-influenced
453	alkaline peat from the Everglades. Not surprisingly, the fractional abundances of the
454	three most common 6-methyl brGDGTs (brGDGT-IIa', -IIb', -IIIa') are significantly
455	correlated with pH with R-values between 0.4 and 0.6 (p<0.01) (Fig. 7). These results
456	are consistent with observations from soils that indicate a positive correlation between
457	the fractional abundance of 6-methyl brGDGTs and pH (De Jonge et al., 2014; Xiao
458	et al., 2015).
459	As a result, the IR_{6me} as well as the related IBT index, both of which have
460	been used to reconstruct pH in soils (Ding et al., 2015; Xiao et al., 2015), are
461	correlated with pH in the peats (not shown). However, this comparison is complicated
462	by the fact that 6-methyl brGDGTs are absent in many of the peats. For IR_{6me} the
463	absence of 6-methyl brGDGTs results in values that are 0, whereas IBT cannot be
464	calculated for samples that lack 6-methyl brGDGTs as the logarithm of zero is
465	undefined.
466	The abundance of 6-methyl brGDGTs is generally lower in peats than in
467	mineral soils with comparable pH. Indeed, 6-methyl brGDGTs are present in 99% of
468	all soils in the global soil database, including soils with pH ${<}5$ where IR_{6me} ratios can
469	be as high as 0.4 (Fig. 6). Recent work has shown that in addition to pH the fractional
470	abundance of 6-methyl brGDGTs is negatively correlated with soil water content,
471	with fewer 6-methyl brGDGTs versus 5-methyl brGDGTs in soils with 60% water

472 content compared to soils with < 10% water content (Dang et al., 2016). It is likely 473 that the negative correlation between soil water content and fractional abundance of 6-474 methyl brGDGTs can explain the overall lower IR_{6me} in peats as these are generally 475 water saturated. 476 In addition to 6-methyl brGDGTs, the fractional abundances of brGDGTs 477 containing cyclopentane moieties (brGDGT–Ib and –IIb) are also significantly 478 correlated to pH (R = 0.73 and 0.56, p<0.01, respectively) (Fig. 7a and 7c). The other 479 brGDGTs are not significantly correlated to pH. These observations are consistent 480 with those from soils, where both 5- and 6-methyl brGDGTs containing cyclopentane 481 moieties are more abundant at higher pH (Weijers et al., 2007; Peterse et al., 2012; De 482 Jonge et al., 2014). Consequently, and similar to soils (De Jonge et al., 2014; Xiao et 483 al., 2015), CBT' (eq. 3) in peat can be modeled as a function of pH (Fig. 8): eq. (10) $pH = 2.69 \times CBT' + 9.19$ (n = 50, $R^2 = 0.44$, RMSE = 1.0) 484 The slope of this calibration is different (higher) from that found in soils (see 485 supplementary information), but the coefficient of determination is lower, and the 486 RMSE is higher. A stronger correlation is obtained by using only compounds that are 487 significantly correlated to pH in the numerator, CBT_{peat}: $eq. (11) CBT_{peat} = \log \left(\frac{Ib + IIa' + IIb + IIb' + IIIa'}{Ia + IIa + IIIa} \right)$ $eq. (12) pH = 2.49 \times CBT_{peat} + 8.07 (n = 51, R^2 = 0.58, RMSE = 0.8)$ 488 Although the coefficient of determination increases and RMSE decreases using 489 CBT_{peat}, the calibration uncertainties are still larger than those reported for soils (see 490 supplementary information). 491 It is noteworthy that in peats the correlation between brGDGT distributions and pH is much weaker than that with MAAT (see below). This contrasts with 492 mineral soils, for which the correlation of CBT' with pH ($R^2 = 0.85$), is stronger than 493 that of MAT_{mr} with MAAT ($R^2 = 0.68$) (De Jonge et al., 2014). The weaker 494 495 correlation can partly be explained by the smaller sample set used for the peat 496 calibration (n = 51) versus soil calibration (n = 221), However, taking 51 random 497 mineral soils from the latter still yields a stronger correlation between CBT' and pH 498 than we obtain for the peat data set. In addition, the coefficient of determination of a 499 calibration based only on peats with pH \geq 5 is \sim 0.5 for CBT_{peat}, similar to that of the 500 complete data set. We argue that the difference could be related to the observation 501 that in mineral soils water content also influences the brGDGT distribution, especially

502	that of 6-methyl brGDGTs (e.g., Menges et al., 2014). Recently Dang et al. (2016)
503	showed that CBT _(5me) is higher in dry soils compared to wet soils. Because alkaline
504	soils are often also dry whereas acidic soils are often wet, this effect could enhance
505	the correlation between CBT' and pH in soils. As peats are typically water saturated,
506	the additional effect of soil water content is lacking, which may explain the weaker
507	correlation between CBT' and pH in peats compared to mineral soils.
508	
509	4.2.2 Influence of MAAT on brGDGTs in peats
510	In mineral soils the distribution of brGDGTs is influenced by MAAT, with the degree
511	of methylation decreasing as temperature increases (Weijers et al., 2007; De Jonge et
512	al., 2014). A temperature effect on the brGDGT distribution was recently also found
513	in one peatland (Huguet et al., 2013). Although the producers of brGDGTs are
514	currently unknown, such a response is consistent with homeoviscous adaptation
515	(Weijers et al., 2007). Here we investigate whether temperature has a significant
516	correlation with brGDGTs in peats on a global scale.
517	When plotted against MAAT, only 5-methyl brGDGTs lacking cyclopentane
518	moieties (brGDGT-Ia, -IIa, and -IIIa) have significant correlations with MAAT (Fig.
519	9). brGDGT-Ia is positively correlated with MAAT ($R = 0.72$, p<0.01), whereas
520	brGDGT-IIa ($R = 0.82$, $p < 0.01$), and -IIIa ($R = 0.63$, $p < 0.01$) are negatively correlated
521	with MAAT. These correlations are significantly higher than those found in the global
522	soil data set (De Jonge et al., 2014). The degree of methylation of 5-methyl brGDGTs
523	is captured in the MBT_{5me} ' index (eq. 2). As such we use the MBT_{5me} ' index to
524	construct a peat-specific temperature proxy (Fig. 10):
	$eq. (13) MAAT_{peat} (^{\circ}C) = 52.18 \times MBT'_{5me} - 23.05 (n = 96, R^2 = 0.76, RMSE)$
	= 4.7 °C)
525	Crucially, no correlation is observed between MBT _{5me} ' and pH ($R^2 = 0$ and $p > 0.8$)
526	and we observe no trend in the residuals. The coefficient of determination (R ²) of
527	MAAT _{peat} is higher compared to a Deming regression of the expanded soil dataset (R ²
528	= 0.60, see supplementary information) as well as that of the linear MBT_{5me} '
529	calibration ($R^2 = 0.66$) suggested by De Jonge et al. (2014). Crucially, because the
530	slope of the MAAT _{peat} calibration is steeper, it could have greater utility for the
531	reconstruction of tropical temperatures (MAAT _{peat} reaches saturation at 29.1 °C),
532	although these maximum temperatures are higher than the maximum MAAT in the

533	modern calibration data set which is 26.7 °C. In contrast, the Deming MBT _{5me} ' soil				
534	calibration reaches saturation (i.e. MBT_{5me} = 1) at a temperature of 24.8 °C (see				
535	supplementary information), while the linear MBT _{5me} ' calibration suggested by De				
536	Jonge et al. (2014) has a maximum of 22.9 °C.				
537					
538	4.3 Implications for paleoclimate reconstructions and future work				
539	Compared to the natural archives previously used to reconstruct past terrestrial				
540	temperature change (e.g., riverine, lacustrine, and marine sediments), peats have a				
541	major advantage. For example, the brGDGTs in peat are mainly derived from in situ				
542	production. Mixing of brGDGT source areas, which complicates the application of				
543	GDGTs in sediments that represent a large catchment area (e.g., Zell et al., 2014; De				
544	Jonge et al., 2015; Sinninghe Damsté, 2016), is unlikely to be a problem. In addition,				
545	peats are overall characterized by anoxic conditions and the preservation potential of				
546	organic compounds such as brGDGTs is high. Finally, as peats are water saturated,				
547	especially the catotelm where the majority of brGDGT production occurs, the				
548	additional influence of changes in moisture content (Menges et al., 2014; Dang et al.,				
549	2016) is also negligible. Nevertheless, there are limitations to this proxy that need to				
550	be considered when evaluating suitable palaeoclimate applications, and we explore				
551	those below.				
552					
553	4.3.1 Late Holocene climate				
554	Here we provide peat-specific temperature and pH proxies that could be used to				
555	reconstruct terrestrial climate over a broad range of time scales, including the late				
556	Holocene. However, the estimated variation in terrestrial temperature of most places				
557	on earth during the last millennium is typically less than 1°C (Mann et al., 2009;				
558	Pages 2k Consortium, 2013), although there could be local exceptions. Such				
559	temperature change is much smaller than the calibration error (RMSE of \sim 4.7 $^{\circ}$ C).				
560	Although based on different lipids produced by different organisms, GDGT proxies				
561	can potentially record temperature changes smaller than the calibration errors when				
562	utilized within a highly constrained site-specific study (Tierney et al., 2010), although				
563	this interpretation was recently contested (Kraemer et al., 2015).				
564	Regardless of calibration issues, application of the MAAT _{peat} calibration to				
565	late Holocene palaeoclimate remains problematic. A potential seasonal bias in the top				
566	of some high-latitude peats, as well as a potential difference between oxic and anoxic				

567	production, appear to prevent application of this proxy to shallow peat sediments.				
568	Indeed, our downcore profiles spanning the top 1 meter of peat exhibit changes in				
569	brGDGT distributions equivalent to temperature variations of up to several degrees				
570	Celsius, larger than the expected climate variations. Moreover, as discussed above,				
571	GDGTs appear to be predominantly generated at depth, and although this evidently				
572	ensures they record MAAT it does mean that their reconstructed temperature signal				
573	start in deeper peat horizons, i.e. stratigraphically preceding the climate changes that				
574	caused them.				
575	Future work should determine whether these peat-specific proxies can be used				
576	to reconstruct small amplitude and/or short-lived temperature variation. However we				
577	currently urge caution in applying the peat-specific proxies to shallow peat cores to				
578	reconstruct late Holocene climate (e.g., Little Ice Age or Medieval Warm anomaly).				
579					
580	4.3.2 Application to the last glacial				
581	We envision these proxies are well-suited to reconstruct large amplitude and more				
582	long-term temperature excursions such as those associated with the last glacial				
583	termination and early Holocene. Such transitions are recorded in some particularly				
584	long peat cores at several places around the world (e.g., McGlone et al., 2010;				
585	Vanneste et al., 2015; Zheng et al., 2015; Baker et al., 2016). To test whether the				
586	novel peat-specific temperature calibration can be used to reconstruct				
587	glacial/interglacial temperature variability, we applied this proxy to samples from the				
588	Hani peat sequence (Fig. 2). Hani peat is located in northeastern China and in places				
589	is up to 10 meters thick, spanning \sim 16,000 cal yrs (Zhou et al., 2010). We analyzed				
590	two samples from ~840 cm depth (corresponding to the late glacial at around 15.3				
591	kyr), and compared $MAAT_{peat}$ with that of two samples from around 100 cm depth				
592	(corresponding to the late Holocene with an age of 700-1000 yrs). Using $MAAT_{peat}$				
593	we obtained an average temperature of around -0.8 °C for the late glacial (15.3 kyr).				
594	For the late Holocene (0.7-1 kyr) we obtained an average temperature of around 4.6				
595	°C (Table 1).				
596	Taking the calibration error of ~4.7 °C into account the reconstructed late				
597	Holocene temperatures (4.6 $^{\circ}$ C) are close to the observed modern-day MAAT of				
598	around 4 $^{\circ}\mathrm{C}$ at this locality (Zhou et al., 2010). In contrast, applying soil calibrations				
599	to reconstruct MAAT at this site results in significantly higher values (up to 11 °C;				
500	Table 1) MAAT (as well as the soil MBT ₅ ; calibration) indicates that				

601	temperatures increased from the late glacial to the late Holocene by around 5 °C. In					
602	contrast the MAT_{mr} mineral soil calibration indicates a smaller increase of around 3					
603	$^{\circ}\text{C.}$ A ~5 $^{\circ}\text{C}$ increase is similar to that observed in east Asian loess-paleosol					
604	sequences (Peterse et al., 2014), although that is based on the MBT(')/CBT method.					
605	In addition a 5 °C deglacial temperature increase is similar to those of several sea					
606	surface temperature records available from similar latitudes in the Sea of Japan (Lee,					
607	2007). The next step should be multiproxy temperature reconstructions in a variety of					
608	locations to test the new calibration and to determine whether absolute temperatures					
609	obtained using MAAT _{peat} are reliable. Nonetheless, this initial analysis indicates that					
610	MAAT _{peat} yields temperature estimates that are consistent with both modern day					
611	observations and other proxy estimates for the last glacial.					
612						
613	4.3.3 Deep time application					
614	We see considerable scope for future work with this proxy to reconstruct terrestrial					
615	temperature during past greenhouse periods and across hyperthermals (e.g					
616	Paleocene/Eocene Thermal Maximum; PETM). These events are recorded in lignite					
617	deposits. For example the PETM is documented in lignites from the UK (Collinson et					
618	al., 2003; Pancost et al., 2007). Importantly, lignites are the lowest (maturity) rank of					
619	coal and have not experienced significant burial and associated temperature and					
620	pressure that leads to the loss of GDGTs (Schouten et al., 2004, 2013). Due to their					
621	low thermal maturity, lignites are thought to retain their original brGDGT distribution					
622	over geological timescales. For example, brGDGTs have been reported in an					
623	immature late Paleocene lignite from the USA (Weijers et al., 2011), early Eocene					
624	lignites from Germany (Inglis et al., 2017), as well as Miocene lignite from Germany					
625	(Stock et al., 2016). Although analyzed using the classical analytical method that did					
626	not separate 5 and 6-methyl brGDGTs, the brGDGT distribution in a late Paleocene					
627	lignite from North America is dominated by brGDGT-Ia (Weijers et al., 2011),					
628	similar to that seen in modern peats from the tropics and suggesting high terrestrial					
629	temperatures. This is consistent with our overall understanding of terrestrial climate					
630	during the greenhouse world of the late Paleocene and early Eocene (Huber and					
631	Caballero, 2011).					
632	As the brGDGT distribution in peat deposits is dominated by production in the					
633	anoxic catotelm below the water table where the seasonal temperature cycle is muted					
634	(see section 4.1) brGDGT-based temperatures obtained from lignite deposits can be					

635	considered to reflect MAAT. We envision that future studies applying our new peat-					
636	specific calibrations to immature lignites will provide valuable new insights into					
637	terrestrial climate during the geological past. In addition, the GDGT concentrations in					
638	peats are generally much higher than those found in soils. We therefore propose that					
639	for studies of brGDGT distributions in (marine) sediments with a peat-dominated					
640	catchment area (e.g. Siberia (Frey and Smith, 2005)) or that contain independent					
641	evidence for the input of peat-derived material (e.g. high concentration of C_{31} $\alpha\beta$ -					
642	2 hopanes or palynologic evidence for the input of typical peatland vegetation), the					
643	majority of GDGTs is likely derived from peatlands. In such settings it is more					
644	appropriate to use a peat-specific calibration rather than a mineral soil calibration.					
645	.69					
646	5. Conclusions					
647	Using 470 samples from 96 peatlands from around the world we explored the					
648	environmental controls on the bacterial brGDGT distribution in peats. We					
649	demonstrate that brGDGT distributions are correlated with peat pH and especially					
650	mean annual air temperature (MAAT). We develop for the first time peat-specific					
651	brGDGT-derived pH and temperature calibrations. In addition to their application in					
652	ancient peat-forming environments, we also suggest that these calibrations could be					
653	preferable to the available mineral soil calibration in marginal marine settings when it					
654	is clear that brGDGTs are predominantly derived from peats. We suggest caution in					
655	applying this proxy to late Holocene peat (e.g., covering the Medieval Climatic					
656	Anomaly and/or Little Ice Age) as both the calibration error and downcore variation					
657	appears to be larger than expected climate signals during this period. Taken together					
658	our results demonstrate that there is clear potential to use GDGTs in peatlands and					
659	lignites to reconstruct past terrestrial climate, opening up a new set of sedimentary					
660	archives that will help to improve understanding of the climate system during the					
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1056	
1057	Fig. 2: Map with the location of all peats used in this study. The star indicates the
1058	location of the Hani peat sequence in NE China.
1059	
1060	Fig. 3: Fractional abundances of the three main brGDGTs in the top 15 cm of each
1061	peat (assumed to be representative of the oxic acrotelm) versus the fractional
1062	abundance of these brGDGTs between 15 and 100 cm in the peat (assumed to be
1063	representative for the anoxic catotelm). For peats where multiple samples were
1064	analyzed, error bars represent 1 σ from the average fractional abundance.
1065	
1066	Fig. 4: Down core record of MBT _{5me} ' in four peats: a high-latitude peat from Sweden
1067	(blue squares), high-latitude peat from Patagonia (orange squares), temperate peat
1068	from the UK (green triangles), and tropical peat from Indonesia (purple diamonds).
1069	(For interpretation of the references to color in this figure legend, the reader is
1070	referred to the web version of this article.)
1071	
1072	Fig. 5: Standard deviation of MBT _{5me} ' for each low-altitude (< 1000 m) peat versus
1073	latitude. The four peats used in figure 4 are highlighted.
1074	
1075	Figure 6: Ratio of 6-methyl over 5-methyl brGDGTs (IR _{6me}) versus pH for peat
1076	samples (black squares) together with the IR_{6me} in the top 10 cm of mineral soils
1077	(orange circles) (De Jonge et al., 2014; Ding et al., 2015; Xiao et al., 2015; Yang et
1078	al., 2015; Lei et al., 2016). Vertical error bars on the peat data represent 1σ and are
1079	based on the analysis of multiple horizons from the same peat. Horizontal error bars
1080	represent the spread in pH reported for each peat. (For interpretation of the references
1081	to color in this figure legend, the reader is referred to the web version of this article.)
1082	
1083	Figure 7: Fractional abundance of brGDGT versus pH for those compounds with a r-
1084	value greater than 0.45 A) brGDGT-Ib, B) brGDGT-IIa', C) brGDGT-IIb, D)
1085	brGDGT-IIb', and E) brGDGT-IIIa' (p $<$ 0.01 for all compounds). Samples with
1086	fractional abundances ≤ 0.001 are not included. Vertical error bars represent 1σ and
1087	are based on the analysis of multiple horizons from the same peat. Horizontal error
1088	bars represent the spread in pH reported for each peat.
1089	

1090	Fig. 8: A) Average CBT' for each peat versus pH (black circles) and C) average
1091	CBT _{peat} for each peat versus pH (black circles). Solid blue lines in A and C represent
1092	the Deming regression used to obtain the calibrations, while dashed black lines reflect
1093	simple linear regressions. Horizontal error bars represent 1σ and are based on the
1094	analysis of horizons samples from the same peat. Vertical error bars represent the
1095	spread in pH reported for each peat. Also shown is the residual pH for all analyzed
1096	peat samples (yellow circles), obtained by subtracting the estimated pH using the
1097	CBT' (B) and CBT _{peat} (D) deming calibrations from the observed pH. (For
1098	interpretation of the references to color in this figure legend, the reader is referred to
1099	the web version of this article.)
1100	
1101	Fig. 9: Fractional abundance of the three main brGDGT versus MAAT A) brGDGT-
1102	Ia, B) brGDGT-IIa, and C) brGDGT-IIIa (p \leq 0.01 for all compounds). Samples with
1103	fractional abundances $<\!\!0.001$ were not included. Vertical error bars represent 1σ and
1104	are based on the analysis of multiple horizons from the same peat.
1105	
1106	Fig. 10: Average MBT _{5me} ' for each peat versus MAAT (black circles). The solid blue
1107	line represents the Deming regression, whereas dashed lines reflect the simple linear
1108	regression. Horizontal error bars represent 1σ and are based on the analysis of
1109	multiple horizons from the same peat. Also shown is the residual MAAT of all
1110	analyzed peat samples (yellow circles) obtained by subtracting the estimated MAAT
1111	using the MBT_{5me} ' Deming calibration from the observed MAAT. (For interpretation
1112	of the references to color in this figure legend, the reader is referred to the web
1113	version of this article).

Depth	Age		MAT _{mr} soil (RMSE 4.6 °C)	MAT _{5me} ' soil (RMSE 4.8 °C)	MAAT _{peat} (RMSE 4.7°C)
(cm)	(yr)	MBT _{5ME} '	De Jonge, 2014	De Jonge, 2014	This study
86	~700	0.53	6.6	10.9	4.5
102	~1000	0.53	6.6	11.3	4.8
838	~15,100	0.46	4.4	6.7	1.2
846	~15.400	0.39	2.8	5.4	-2.7
		Δ ΜΑΑΤ	3.0 °C	5.0 °C	5.4 °C

