

1       **Distribution of branched glycerol dialkyl glycerol tetraethers in**  
2       **surface soils of Qinghai–Tibetan Plateau: implications of brGDGTs-**  
3       **based proxies in cold and dry regions**

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13  
14      **Abstract.** The methylation index of branched tetraethers (MBT) and cyclization ratio of  
15      branched tetraethers (CBT) based on the distribution of branched glycerol dialkyl  
16      glycerol tetraethers (brGDGTs) are useful proxies for the reconstruction of mean annual  
17      air temperature (MAT) and soil pH. Recently, a series of 6-methyl brGDGTs were  
18      identified which were previously co-eluted with 5-methyl brGDGTs. However, little is  
19      known about the distribution of 6-methyl brGDGTs in Qinghai-Tibet Plateau (QTP), a  
20      critical region of the global climate system. Here, we investigated 30 surface soils  
21      covering a large area of QTP, among which 6-methyl brGDGTs were the most abundant  
22      components (average 53±17 % of total brGDGTs). The fractional abundance of 6-methyl  
23      brGDGTs showed a good correlation with soil pH, while the global MBT<sub>5ME</sub> calibration  
24      overestimates MAT in cold regions like QTP. We therefore propose a MBT<sub>5/6</sub> index  
25      including both 5- and 6-methyl brGDGTs, presenting a strong correlation with MAT in  
26      QTP:      $\text{MAT} = -20.14 + 39.51 \times \text{MBT}_{5/6}$  ( $n = 27, r^2 = 0.82; \text{RMSE} = 1.3 \text{ } ^\circ\text{C}$ ) .

27      Another index, namely IBT based on carbon skeleton isomerism of 5-methyl to 6-methyl  
28      brGDGTs, is dependent on soil pH:      $\text{pH} = 6.77 - 1.56 \times \text{IBT}$  ( $n = 27; r^2 =$

29 0.74, RMSE = 0.32). Our study suggests that changing the position of methyl group of  
30 brGDGTs may be another mechanism for some soil bacteria to adapt ambient pH change  
31 besides well-known cyclization.

32

### 33 **1. Introduction**

34 The Qinghai–Tibetan Plateau (QTP), with an area of over 2.5 million km<sup>2</sup> and an  
35 average elevation of over 4000 meters above sea level (a.s.l.), is the world highest and  
36 largest mountain plateau. The uplift of the QTP since early Cenozoic profoundly  
37 influences regional and global climates such as the evolution of Asian monsoon which  
38 affects lives of over two billion people (An et al., 2001; Li, 1991; Lin et al., 2008; Wang  
39 et al., 2008). A number of studies have showed that the QTP is a highly sensitive area  
40 for global climate change (e.g., Kang et al., 2010; Liu & Chen, 2000; Qiu, 2008; Yao et  
41 al., 2007). The record of 97 meteorological stations located over 2000 meters a.s.l. in  
42 China reveals that winter temperature rise is 0.32 °C per decade in the QTP since 1950s,  
43 approximately three times the global warming rate (Liu & Chen, 2000). However, the  
44 history of instrumental measurement is too short to fully record the evolution of the QTP  
45 climate. The reconstruction of the QTP temperature beyond instrumental measurement  
46 is challenging because few quantitative proxies are available. Microfossil assemblages  
47 based on pollen, diatom or chironomid are commonly used paleothermometers, but they  
48 are also influenced by precipitation, salinity, nutrient or other environmental factors (e.g.,  
49 Keatley et al., 2009; Meriläinen et al., 2000; Seppä & Birks, 2001). The  $\delta^{18}\text{O}$  value of  
50 ice core in the QTP shows a good correlation with northern hemisphere temperature  
51 (Thompson et al., 1997; Yao et al., 2002). Unfortunately, ice core with a long term,  
52 continuous record is lacking in most QTP.

53 Over the past decades, two molecular proxies have been developed for estimation  
54 of continental temperature. The first one, namely UK'37, is based on the distribution of  
55 haptophyte-derived long-chain alkenones. This proxy was originally proposed for  
56 paleoceanography (Brassell et al., 1986; Prahl et al., 1988), but was found applicable for  
57 reconstruction of lake surface temperature (e.g., Liu et al., 2006; Zink et al., 2001). A

58 major limitation of UK'37 is that long-chain alkenones are not always present in lakes,  
 59 although they were reported in some QTP lakes (e.g., Chu et al., 2005; Liu et al., 2011;  
 60 Liu et al., 2006). In addition, salinity influences the compositions of long-chain  
 61 alkenones in lakes (Liu et al., 2011). Besides UK'37, the methylation index of branched  
 62 tetraethers (MBT) and cyclization ratio of branched tetraethers (CBT) can be also used  
 63 to infer past continental temperature based on the distribution of branched glycerol  
 64 dialkyl glycerol tetraethers (brGDGTs) (Weijers et al., 2007b):

$$65 \quad \text{MBT} = \frac{\text{Ia}+\text{Ib}+\text{Ic}}{\text{Ia}+\text{Ib}+\text{Ic}+\text{IIa}+\text{IIb}+\text{IIc}+\text{IIIa}+\text{IIIb}+\text{IIIc}+\text{IIa}'+\text{IIb}'+\text{IIc}'+\text{IIIa}'+\text{IIIb}'+\text{IIIc}'} \quad (1)$$

$$66 \quad \text{CBT} = -\log \frac{\text{Ib}+\text{IIb}+\text{IIb}'}{\text{Ia}+\text{IIa}+\text{IIa}'} \quad (2)$$

67 where roman numbers denote relative abundance of compounds in Fig. 1. It should be  
 68 pointed out that the Eq. 1 and 2 are rewritten from original definitions because the peaks  
 69 previously identified as pure 5-methyl brGDGTs (Weijers et al., 2007b) are actually  
 70 mixtures of 5-methyl and 6-methyl isomers (De Jonge et al., 2013).

71 So far, only two species of Acidobacteria were identified to produce brGDGTs  
 72 (Sinninghe Damsté et al., 2011), but the ubiquitous occurrence of brGDGTs in soils/peats,  
 73 lakes and marginal seas suggest that other biological sources are likely (Schouten et al.,  
 74 2013 and references therein). By analyzing globally distributed soils, Weijers et al.  
 75 (2007b) found that the MBT is controlled by mean annual air temperature (MAT) and to  
 76 less extent by soil pH, whereas CBT only relates to soil pH. Such relationship was  
 77 corroborated by the subsequent study of Peterse et al. (2012) who proposed a simplified  
 78 format of MBT (or MBT') based on seven quantifiable brGDGTs.

79 Since the advent, the MBT(MBT')-CBT paleotemperature proxy has been  
 80 increasingly used for lakes (e.g., D'Anjou et al., 2013; Loomis et al., 2012; Sun et al.,  
 81 2011), paleosol-loess sequences (e.g., Peterse et al., 2011; Zech et al., 2012), peat  
 82 (Ballantyne et al., 2010) and marginal seas (e.g., Bendle et al., 2010; Weijers et al., 2007a;  
 83 Zell et al., 2014). However, a relatively large scatter in global MBT/CBT-MAT  
 84 calibrations (about 5 °C for root mean square error; RMSE) suggests that other factors  
 85 besides temperature may influence brGDGTs-based indices (Peterse et al., 2012; Weijers  
 86 et al., 2007b). In arid and semiarid areas such as western United States where

87 precipitation is the ecological limiting factor, mean annual precipitation (MAP) rather  
 88 than MAT is the most important factor that affects brGDGT compositions (Dirghangi et  
 89 al., 2013; Menges et al., 2014). The updated global calibration of MBT'-CBT indices  
 90 (Peterse et al., 2012) also shows a weak correlation with MAT for those soil samples  
 91 from arid regions (MAP < 500 mm). Some studies suggest that regional calibrations are  
 92 needed to improve accuracy of the GDGTs-based proxy (e.g., Loomis et al., 2012;  
 93 Pearson et al., 2011; Shanahan et al., 2013; Zink et al., 2010).

94 Another factor to cause the relatively large scatter of the MBT/CBT-MAT  
 95 calibration is analytical error. By applying advanced analytical techniques, De Jonge et  
 96 al. (2013) identified a series of novel 6-methyl brGDGTs which previously co-eluted  
 97 with 5-methyl GDGTs that were used to calculate the brGDGTs proxies. The successful  
 98 separation of 5- and 6-methyl brGDGTs resulted in a set of new brGDGT proxies, which  
 99 were used to recalibrate traditionally defined MBT-CBT indexes (De Jonge et al., 2014):

$$100 \quad MBT'_{5ME} = \frac{Ia+Ib+Ic}{Ia+Ib+Ic+IIa+IIb+IIc+IIIa} \quad (3)$$

$$101 \quad MAT = -8.57 + 31.45 \times MBT'_{5ME}$$

$$102 \quad (n = 222, r^2 = 0.66; RMSE = 4.8 \text{ } ^\circ\text{C}, P < 0.001) \quad (4)$$

$$103 \quad CBT_{5ME} = -\log \frac{Ib + IIb}{Ia + IIa} \quad (5)$$

$$104 \quad pH = 7.84 - 1.73 \times CBT_{5ME}$$

$$105 \quad (n = 221, r^2 = 0.60; RMSE = 0.84, P < 0.001) \quad (6)$$

$$106 \quad CBT' = \log \frac{Ic + IIa' + IIb' + IIc' + IIIa' + IIIb' + IIIc'}{Ia + IIa + IIIa} \quad (7)$$

$$107 \quad pH = 7.15 + 1.59 \times CBT'$$

$$108 \quad (n = 221, r^2 = 0.85; RMSE = 0.52, P < 0.001) \quad (8)$$

109 For the QTP, several studies have reported GDGTs in lakes, mountains, hot springs  
 110 and paleo-soils (e.g., Günther et al., 2014; He et al., 2012; Liu et al., 2013; Wang et al.,  
 111 2012; Wu et al., 2013; Xie et al., 2012). Wang et al. (2012) analyzed GDGTs in surface  
 112 sediments of the Lake Qinghai and surrounding soils, showing that brGDGTs-inferred  
 113 MAT and soil pH were consistent with measured values. In contrast, Wu et al. (2013)  
 114 found that brGDGTs-derived MAT was higher than instrumentally measured MAT in  
 115 Kusai Lake sediments from the QTP. Based on the distributions of GDGTs in surface

116 sediments of the QTP lakes, Günther et al. (2014) developed the local calibration of  
117 MBT<sup>2</sup>-CBT ( $r^2 = 0.59$ ; RMSE = 1.2 °C). However, there are only 9 lake sediments  
118 included in Günther et al. (2014). For the application of MBT-CBT indices in lakes,  
119 brGDGTs in lake sediments must be exclusively derived from inputs of surrounding soils.  
120 However, in-situ production of brGDGTs occurs in various lakes (e.g., Blaga et al., 2009;  
121 Blaga et al., 2010; Fietz et al., 2012; Pearson et al., 2011; Sinninghe Damsté et al., 2009;  
122 Tierney & Russell, 2009). Furthermore, the 6-methyl brGDGTs were not reported in any  
123 QTP studies, which may explain the relatively low  $r^2$  value of the MBT/CBT-MAT  
124 calibration (e.g., Günther et al., 2014). Given these facts, a direct investigation of soils  
125 with improved chromatography is needed to understand environmental influences on the  
126 brGDGT distributions in the QTP.

127 Here, we analyzed all 5- and 6-methyl brGDGTs in 30 surface soils from the QTP.  
128 Our main objectives are to (1) determine the relative abundance and distribution of 5-  
129 and 6-methyl brGDGTs in the QTP soils; (2) evaluate the effect of recently identified 6-  
130 methyl brGDGTs on soil pH in the QTP; and (3) test whether global brGDGTs-MAT  
131 calibration is applicable in the QTP and thereby understand the influence of temperature,  
132 precipitation and soil pH on 5- and 6-methyl brGDGTs in the QTP.

133

## 134 **2. Materials and methods**

### 135 **2.1. Sampling**

136 A total of 30 surface soil samples (0-10 cm) were collected during two fieldworks  
137 in 2011 and 2012, which cover a large area of the QTP (84.64°~101.20° E; 28.24°~37.45°  
138 N) (Fig. 2). Sampling sites are typical alpine meadow, alpine steppe or alpine meadow  
139 steppe. The extremely dry winter results in the lack of persistent snow cover in most  
140 sampling sites. The soil samples were air-dried and passed through a 2 mm mesh to  
141 remove large gravels. Fine roots (if present) were picked up by steel tweezers. The  
142 detailed information on the sampling sites and environmental variables are listed in the  
143 supplementary material (Table S1).

144

### 145 **2.2 Climate data**

146 There are about 70 meteorological stations in the QTP, mainly distributed in the  
147 eastern part and northern border of the QTP. Thus, direct observation data on temperature  
148 and precipitation at our sampling sites are generally lacking. In this study, we use the  
149 WorldClim dataset (Hijmans et al., 2005) to interpolate annual, seasonal and monthly  
150 mean precipitation and temperature (Table S1). The local climate is dry and cold. The  
151 MAT of our sampling sites ranges from -5.5 to 7.6 °C with a vertical lapse rate of  
152 0.487 °C/100 m to 0.699 °C/100 m (Cheng et al., 2012). The vertical lapse rate of air  
153 temperature decreases from north to south of the QTP. The mean annual precipitation  
154 (MAP) at different altitudes varies from ca. 85 mm to ca. 495 mm. The integrated maps  
155 are derived from monthly temperature and precipitation values gathered from thousands  
156 of weather stations around the world from 1950 to 2000 (47,554 locations for  
157 precipitation and 24,542 locations for temperature). The original point data was splines  
158 interpolated using latitude and longitude at a fine resolution, making it possible to obtain  
159 a reasonable estimation of climatic conditions at individual sites. The WorldClim GIS  
160 data used contain annual average of 6 climate variables at a 30 arc seconds resolution  
161 (~1 km resolution; <http://www.worldclim.org/current.htm>). Besides MAT and MAP,  
162 additional four climate variables were also used to evaluate the relationship between  
163 climate and 5- and 6-methyl brGDGTs indices, including Mean Temperature of Wettest  
164 Quarter (MWQT), Mean Temperature of Driest Quarter (MDQT), Mean Temperature of  
165 Warmest Quarter (MWQT'), Mean Temperature of Coldest Quarter (MCQT). A total of  
166 30 sites from QTP cold and dry regions (Table S1) were extracted by 6 climate variables  
167 using Arcgis 9.3.

168

### 169 **2.3 Soil pH and brGDGT analyses**

170 For pH measurement, soils were mixed with deionized water in a ratio of 1/2.5  
171 (g/ml). The soil pH values were determined by a pH meter with a precision of  $\pm 0.01$  pH.  
172 The pH was reported as an average of three duplicate measurements for each sample  
173 with standard deviation of  $\pm 0.05$ .

174 The detailed procedure for lipid extraction was described by Wu et al. (2014). About  
175 6 g dry soils were mixed with 600 ng C<sub>46</sub> GDGT (internal standard) (Huguet et al., 2006)

176 and ultrasonically extracted with 20 ml dichloromethane (DCM)/methanol (3:1 v:v) for  
177 15 min (3×). The combined extracts were concentrated to near dryness by a rotary  
178 evaporator and transferred to small vials. The concentrated extracts were completely  
179 dried under a mild stream of N<sub>2</sub> and re-dissolved in DCM. The total extracts were  
180 separated into two fractions by 5 ml hexane/DCM (9:1 v:v) and 5 ml DCM/methanol  
181 (1:1 v:v), respectively, on silica gel columns. The latter fraction containing brGDGTs  
182 was dissolved in 300 µl hexane/EtOAc (84:16,v/v).

183 The GDGTs were analyzed on an Agilent 1200 High Performance Liquid  
184 Chromatography-atmospheric pressure chemical ionization-triple quadruple mass  
185 spectrometry (HPLC-APCI-MS<sup>2</sup>) system (Yang et al., 2015). The injection volume was  
186 10 µl. The separation of 5- and 6-methyl brGDGTs was achieved with two silica columns  
187 in sequence (150 mm × 2.1 mm; 1.9 µm, Thermo Finnigan; USA) at a constant flow of  
188 0.2 ml per min. The solvent gradient was: 84% A (hexane) and 16% B (EtOAc) for 5  
189 min, then increasing the amount of B from 16% at 5 min to 18% at 65 min, and then to  
190 100% B in 21 min. The column was flushed with 100% B for 4 min, and then back to  
191 84/16 A/B to equilibrate it for 30 min. The APCI and MS conditions were: vaporizer  
192 pressure of  $4.2 \times 10^5$  Pa, vaporizer temperature of 400 °C, drying gas flow of 6 L min<sup>-1</sup>,  
193 temperature of 200 °C, capillary voltage of 3500 V, and corona current of 5 µA (3.2  
194 kV). Peak integration was carried out using Agilent MassHunter. Samples were  
195 quantified based on comparisons of the respective protonated-ion peak areas of each  
196 GDGT to the internal standard in selected ion monitoring (SIM) mode. The protonated  
197 ions were m/z 1050, 1048, 1046, 1036, 1034, 1032, 1022, 1020, 1018 and 744 (C<sub>46</sub>  
198 GDGTs). Since we assume same response factors among different brGDGTs and C<sub>46</sub>  
199 GDGTs, our study can be only regarded as semi-quantification.

200

## 201 **2.4 Statistical analyses**

202 In order to assess the relationship of 5- and 6-methyl brGDGT distributions with  
203 environmental variables such as temperature, precipitation and soil pH, we performed  
204 redundancy analysis (RDA) (van den Wollenberg, 1977), a constrained form of the linear  
205 ordination method of principal components analysis (PCA). Species (fractional

206 abundance of 15 brGDGTs) were centered and standardized with zero average and unit  
207 variance before RDA. The significance of the explanatory variances within a 1%  
208 confidence interval was tested with 999 unrestricted Monte Carlo permutations.  
209 Subsequently, a series of partial RDAs (pRDA) were performed to constrain the unique  
210 and independent influence of individual environmental parameter alone, as well as  
211 compared to all other parameters. All statistical analyses were performed with the  
212 CANOCO version 4.5 software (Wageningen UR, USA).

213

### 214 **3. Results and discussion**

#### 215 **3.1 brGDGTs abundance in the QTP soils**

216 All soil samples except for P790, P840 and P855 contain detectable amounts of  
217 brGDGTs. Consequently, 27 soils were used to calibrate brGDGTs' indices in this study.  
218 With the application of two silica LC columns in tandem, 5-methyl and 6-methyl  
219 brGDGT isomers were successfully separated, increasing the number of detectable  
220 brGDGT compounds from 9 (Peterse et al., 2012; Weijers et al., 2006) to 15 (Fig. 1).  
221 There were three tetra-methylated brGDGTs (Ia, Ib and Ic), six penta-methylated  
222 brGDGTs (IIa, IIb, IIc, IIa', IIb', IIc') and six hexa-methylated brGDGTs (IIIa, IIIb, IIIc,  
223 IIIa', IIIb', IIIc'). The mean fractional abundance of 5-methyl brGDGTs ( $f_{5ME}$ ) and 6-  
224 methyl brGDGTs ( $f_{6ME}$ ) was shown in Fig. 3. The 6-methyl brGDGTs accounted for  
225 average 53% of the total amount of brGDGTs, which were dominated by IIa' and IIIa'.  
226 Such composition of brGDGTs is different from that of the global soils (239 soils) that  
227 5-methyl brGDGT (Ia and IIa) are usually the most abundant isomers and 6-methyl  
228 brGDGTs only comprise on average 24% of the total amounts of brGDGTs (De Jonge et  
229 al., 2014), suggesting that the brGDGT-producing bacteria may change their membrane  
230 lipids to adapt environmental conditions. So far, two species of Acidobacteria are only  
231 identified biological sources for brGDGTs, but they only produce tetra-methylated  
232 brGDGTs (Sinninghe Damsté et al., 2011). In our study, the majority of the QTP soils  
233 are weak alkaline (6.2~8.4 pH unit), which may favor thriving of non-Acidobacteria and  
234 thereby lead to the higher proportion of 6-methyl brGDGTs.

235



### 236 **3.2 Environmental control on brGDGT distributions in QTP**

237 A number of studies have demonstrated that temperature, precipitation and pH are  
238 the most important factors that affect the brGDGT distributions in soils (e.g., De Jonge  
239 et al., 2014; Dirghangi et al., 2013; Peterse et al., 2012; Weijers et al., 2006; Weijers et  
240 al., 2007b; Yang et al., 2015; Yang et al., 2014). In order to evaluate the contribution of  
241 these parameters to 5- and 6-methyl brGDGT distributions in the QTP, a RDA was  
242 performed (Fig. 4). The first component explains 65.2% of the variance, mainly  
243 reflecting the variation in soil pH and to less extent MAP. Soil pH presents strong  
244 positive relationships with fractional abundance of brGDGTs IIIa', IIB', IIB, and negative  
245 relationships with that of IIIa, IIa, Ia. The second component of the RDA plot explains  
246 6.1% of total variance, mainly reflecting the variation in MAT and MAP. The brGDGT-  
247 IIIa', IIIa, IIa, IIB, IIC show negative relationships with MAT (in the lower part of RDA),  
248 whereas brGDGT-IIa, Ia, Ib and Ic present positive relationships with MAT (in the upper  
249 part of RDA). These results support a physiological mechanism that soil bacteria change  
250 the number of methyl branches of brGDGTs with temperature in order to maintain  
251 acceptable fluidity of their membranes (Weijers et al., 2007b).

252 Our RDA result shows that MAT and pH have a significant independent effect on  
253 the brGDGT distribution in the QTP soils, however, no significant correlation was  
254 observed between MAP and brGDGTs ( $p > 0.05$ ; Table 1). Soil pH explaining up to 60.1%  
255 of the total variables is the largest contributor to the variance, followed by MAT (up to  
256 16.4%) and MAP (up to 10.8%). The predominant influence of soil pH on brGDGT  
257 distributions was also observed in global soil dataset (De Jonge et al., 2014; Peterse et  
258 al., 2012) and Chinese soils (Yang et al., 2015; Yang et al., 2014). In order to estimate  
259 the independent, marginal effect of MAT, MAP and pH, partial RDA (pRDA) was  
260 performed. The explained variance of pH still remains high (39.9%), indicating that  
261 brGDGT distributions are indeed linked to soil pH, whereas MAT contribute to a smaller  
262 amount (10.6%) of the variance (Table 2). Similar to the result of RDA, pRDA also  
263 showed minor contribution of MAP (2.0%) to brGDGT distributions. The comparison  
264 between RDA and pRDA suggests a decreasing contribution of these three  
265 environmental variables (pH, MAT, MAP) when they are considered as a unique

266 contribution (Table 2). Thus, there is a “synergistic effect” (an “antagonistic action”)  
267 when MAP and pH (MAT and pH) are considered as covariables, resulting in a positive  
268 joint effect of 20.4% for total contribution of pH+MAT+MAP to brGDGT distributions  
269 in the QTP soils.

270

### 271 **3.3 Evaluation of brGDGT-based proxies in the QTP**

272 Since the identification of 6-methyl brGDGTs (De Jonge et al., 2013), a set of new  
273 brGDGT indices such as MBT'<sub>5ME</sub> and CBT<sub>5ME</sub> have been proposed in order to reduce  
274 uncertainty of reconstructed MAT and soil pH (De Jonge et al., 2014; Weijers et al.,  
275 2007a; Yang et al., 2015). However, even with application of the MBT<sub>5ME</sub>-MAT  
276 recalibration and the multiple regression, relatively large scatter still exists for those  
277 samples from cold regions (De Jonge et al., 2014). Therefore, further calibrations of  
278 brGDGT-derived proxies are needed.

279

#### 280 **3.3.1 MAT calibration in cold and dry regions of the QTP**

281 Consistent with the finding of De Jonge et al. (2014), our result shows that CBT<sub>5ME</sub>  
282 no longer contributes significantly to MAT after the exclusion of 6-methyl brGDGTs ( $p$   
283 = 0.51;  $n = 27$ ). Therefore, we use MBT'<sub>5ME</sub> only to calibrate MAT. Considering that  
284 limited samples from cold regions were included in previous studies (Peterse et al., 2012;  
285 Weijers et al., 2007b), we added our QTP data into the global soil dataset (De Jonge et  
286 al., 2014), resulting in a new calibration of MBT'<sub>5ME</sub>-MAT:

$$287 \text{MAT} = -10.07 + 33.50 \times \text{MBT}'_{5\text{ME}} \\ 288 (n = 249, r^2 = 0.70; \text{RMSE} = 4.7 \text{ }^\circ\text{C}, P < 0.001) \quad (9)$$

289 The correlation coefficient of Eq. 9 ( $r^2 = 0.70$ ) is slightly higher than the previous  
290 global calibration ( $r^2 = 0.66$ ; Eq. 4), while its RMSE (4.7 °C) is similar to the previous  
291 calibration (4.8 °C; Eq. 4) (De Jonge et al., 2014). Furthermore, the comparison of our  
292 estimated MAT and actual MAT ( $\Delta\text{MAT} = \text{MAT}_{\text{est}} - \text{MAT}_{\text{act}}$ ) showed an apparent  
293 overestimation (average 2.8 °C; Fig. 5). Therefore, the simple extension of dataset is not  
294 successful in improving accuracy of the MBT'<sub>5ME</sub>-MAT proxy at the global scale.

295 Alternatively, we conducted a regional calibration of MBT'<sub>5ME</sub> versus MAT based

296 on 27 QTP soils, and a new equation of MBT'<sub>5ME</sub>-MAT was expressed:  
297  $MAT = -10.82 + 28.36 \times MBT'_{5ME}$  ( $n = 27, r^2 = 0.65; RMSE = 1.8 \text{ }^\circ\text{C}, P < 0.001$ ) (10)

298 The slope of Eq. 10 (28.36) is distinct difference from that of global surface soils  
299 (33.50; Eq. 4). Meanwhile, its RMSE value (1.8 °C) is substantially smaller than that of  
300 De Jonge et al. (2014) (4.8 °C). This reduced uncertainty in reconstructed MAT is  
301 attributed to smaller spatial heterogeneity of soils, similar vegetation types (e.g., alpine  
302 meadow) and a narrower MAT range (-5.5~7.6°C) in the QTP. Usually, the regional  
303 calibration has higher  $r^2$  values than the global one due to its smaller size of dataset and  
304 smaller spatial heterogeneity (Loomis et al., 2014; Yang et al., 2014). However, our  
305 calibration for the QTP has a slightly lower  $r^2$  value (0.65) than the global one (0.70; Eq.  
306 9), suggests that the calibration based on MBT'<sub>5ME</sub> alone is not superior to the traditional  
307 MBT calibration. The RDA result reveals that similar to 5-methyl brGDGTs, 6-methyl  
308 brGDGTs also significantly correlate with MAT (Fig. 4). Thus, we propose a new  
309 brGDGT index (MBT<sub>5/6</sub>) including 5-methyl brGDGTs used in the traditional definition  
310 and two dominant 6-methyl brGDGTs (IIa' and IIIa'), expressed as:

$$311 \quad MBT_{5/6} = \frac{Ia + Ib + Ic + IIa'}{Ia + Ib + Ic + IIa + IIb + IIc + IIIa + IIIa'} \quad (11)$$

312 Based on data of the QTP soils, the linear correlation of MAT and MAT<sub>5/6</sub> was  
313 established as (Fig. 6):  $MAT = -20.14 + 39.51 \times MBT_{5/6}$   
314 ( $n = 27, r^2 = 0.82; RMSE = 1.3 \text{ }^\circ\text{C}, P < 0.001$ ) (12)

315 This calibration has substantially higher  $r^2$  (i.e. 0.82) and lower RMSE (i.e. 1.3 °C)  
316 than Eq. 9 (i.e., 0.70 for  $r^2$  and 4.7 °C for RMSE) and Eq. 11 (i.e., 0.65 for  $r^2$  and 1.8 °C  
317 for RMSE), supporting that the inclusion of 5-methyl and 6-methyl brGDGTs is essential  
318 for improved accuracy of MAT reconstruction. However, this result is different from the  
319 finding from the Mount Shennongjia (central China) that 6-methyl brGDGTs are  
320 regarded as the interference, leading to a larger scatter of the MBT'-MAT proxy (Yang  
321 et al., 2015). Nevertheless, these differences highlight the importance of regional  
322 calibrations of brGDGT proxies.

323

324 3.3.2 Effect of soil pH on position of methyl group(s) of brGDGTs

325 It is generally accepted that the proton permeability of the cell membrane plays a  
 326 crucial role in maintaining pH gradient across the membrane of bacteria and archaea  
 327 (Konings et al., 2002). Weijers et al. (2007b) observed a strong correlation between  
 328 relative abundance of cyclopentane moieties of brGDGTs and soil pH, and hypothesized  
 329 that, some soil bacteria can change the methyl groups of brGDGTs into cyclopentyl  
 330 groups with ambient pH rise, which will loosen the packing of the membrane lipids,  
 331 enabling more water molecules to get trapped.

332 Following the approach of De Jonge et al. (2014), we got the following correlation  
 333 between soil pH and CBT' which is a modified format of originally defined CBT  
 334 (Weijers et al., 2007b):

$$335 \text{pH} = 7.01 + 1.49 \times \text{CBT}' \quad (n = 27, r^2 = 0.78, \text{RMSE} = 0.30 \text{ pH unit}) \quad (13)$$

336 The Eq. 12 has slightly lower  $r^2$  and substantially lower RMSE compared with the  
 337 global calibration of pH-CBT' ( $n=221, r^2=0.85, \text{RMSE}=0.52$ ) (De Jonge et al., 2014),  
 338 suggesting that both global and regional calibrations are applicable for soil pH  
 339 reconstruction.

340 We noted that some non-cyclopentyl brGDGTs such as Ia, IIa and IIIa show  
 341 negative correlations with soil pH, while other brGDGTs show positive correlations with  
 342 soil pH in the RDA (Fig. 4). Based on these facts, we put all positively correlated  
 343 brGDGTs on the numerator and all negatively correlated brGDGTs on the denominator  
 344 to build a new CBT index (or CBT'')

$$345 \text{CBT}'' = \log \frac{\text{Ib}+\text{Ic}+\text{IIb}+\text{IIc}+\text{IIIb}+\text{IIIc}+\text{IIa}' + \text{IIb}' + \text{IIc}' + \text{IIIa}' + \text{IIIb}' + \text{IIIc}'}{\text{Ia}+\text{IIa}+\text{IIIa}} \quad (14)$$

346 A linear correlation between soil pH and CBT'' was established based on 27 QTP soils:

$$347 \text{pH} = 6.93 + 1.49 \times \text{CBT}'' \quad (n = 27, r^2 = 0.80, \text{RMSE} = 0.29 \text{ pH unit}) \quad (15)$$

348 The similar  $r^2$  and RMSE between Eq. 13 and 15 was attributed to minor amounts  
 349 of brGDGTs Ib, IIb, IIc, IIIb and IIIc (average 8% of total brGDGTs; Fig. 3) which were  
 350 excluded from the CBT' index but included in our CBT'' index.

351 The fractional abundance of 6-methyl brGDGTs showed strong positive  
 352 correlations with soil pH in both the QTP ( $r^2 = 0.74$ ; Fig. 8) and global soil dataset ( $0.41$   
 353  $< r^2 < 0.72$ ; De Jonge et al., 2014). This is apparent contrast with the previous assumption

354 that non-cyclopentyl moieties (such as IIa' and IIIa') negatively correlate with soil pH.  
355 Unlike 6-methyl brGDGTs, some 5-methyl brGDGTs did not show positive correlations  
356 with soil pH (de Jonge et al., 2014). Thus, we hypothesize that besides cyclization, the  
357 position of methyl group(s) of brGDGTs also influences cell membrane fluidity. In order  
358 to test this hypothesis, we define a new index about carbon skeleton Isomerism of  
359 Branched Tetraethers (or IBT) as the abundant ratio of non-cyclopentyl 6-methyl to 5-  
360 methyl brGDGTs:

$$361 \quad \text{IBT} = -\log \frac{\text{IIIa}' + \text{IIa}'}{\text{IIa} + \text{IIa}} \quad (16)$$

362 We performed a linear regression of IBT versus soil pH based on 27 QTP soils (Fig.  
363 8), yielding an equation as:

$$364 \quad \text{pH} = 6.77 - 1.56 \times \text{IBT} \quad (\text{n} = 27; r^2 = 0.74, \text{RMSE} = 0.32) \quad (17)$$

365 Meanwhile, the linear correlation of CBT<sub>5ME</sub> and soil pH was also established:

$$366 \quad \text{pH} = 7.98 - 1.12 \times \text{CBT}_{5\text{ME}} \quad (\text{n} = 27; r^2 = 0.66, \text{RMSE} = 0.37) \quad (18)$$

367 For the regional calibration, the IBT index has higher  $r^2$  and lower RMSE than  
368 traditionally defined CBT<sub>5ME</sub> index, supporting that the carbon skeleton isomerism of  
369 brGDGTs (i.e., changing the position of methyl group) is indeed a physiological  
370 mechanism of brGDGTs-producing bacteria to adapt soil pH change.

371

### 372 3.3.3 Seasonality of brGDGTs proxies in the QTP

373 The QTP is under strong influence of Asian Monsoon, characterized by  
374 warm/humid summer (June to August) and dry/cold winter (December to February) (An  
375 et al., 2001; Qiu, 2008). In order to examine if there is a seasonal bias on brGDGT  
376 distributions, we analyze the correlation coefficients of 5- and 6-methyl brGDGTs  
377 proxies (i.e., MBT'<sub>5ME</sub>, MBT<sub>5/6</sub>, CBT<sub>5ME</sub> and IBT) versus annual and seasonal air  
378 temperature (Table S2). Overall, there is no apparent seasonal bias for MBT'<sub>5ME</sub> and  
379 MBT<sub>5/6</sub>. This is likely attributed to significant correlation between seasonal temperature  
380 and MAT in the QTP ( $r^2 > 0.80$ ,  $p < 0.0001$ ). In addition, In addition, no significant  
381 correlation was observed between the CBT indices/IBT and MAT/seasonal temperature  
382 ( $-0.3 < r < 0.3$ ; Table S2), suggesting minor influence of air temperature on these indices.

383 Our results are consistent with that of Weijers et al. (2011) who found no significant  
384 seasonal bias in MBT-CBT indices in mid-latitude soils. Therefore, the reconstruction of  
385 MAT based on the 5- and 6-methyl brGDGTs proxies is doable in the QTP.

386

#### 387 **4. Conclusions**

388 By applying improved chromatography, we successfully separated 5- and 6-methyl  
389 brGDGTs in the surface soils from the Qinghai-Tibet Plateau (QTP), a cold and dry  
390 region. This is the first time to report 6-methyl brGDGTs in the QTP, providing an  
391 opportunity to optimize brGDGTs' proxies in this critical region. Three conclusions were  
392 reached based on brGDGT data in 27 surface soils. Firstly, the 6-methyl brGDGTs are  
393 widely distributed in the QTP soils accounting for average 53% of total amounts of  
394 brGDGTs. Secondly, soil pH is the most important contributor to the variance of  
395 brGDGTs, followed by MAT, while MAP has no significant effect on brGDGTs'  
396 distributions. Thirdly, two new indices including recently identified 6-methyl brGDGTs  
397 were proposed to estimate MAT and soil pH, respectively. The first one, namely MBT<sub>5/6</sub>,  
398 is useful for the MAT reconstruction in cold and dry regions (like QTP) with an improved  
399 RMSE of 1.3 °C. The second one, namely IBT, is allowed to estimate soil pH with an  
400 RMSE of 0.32. Our study demonstrates that besides previously reported cyclization,  
401 isomerization of 5-methyl to 6-methyl brGDGTs (expressed as IBT) is another strategy  
402 for brGDGTs-producing bacteria to adapt ambient pH change.

403

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407

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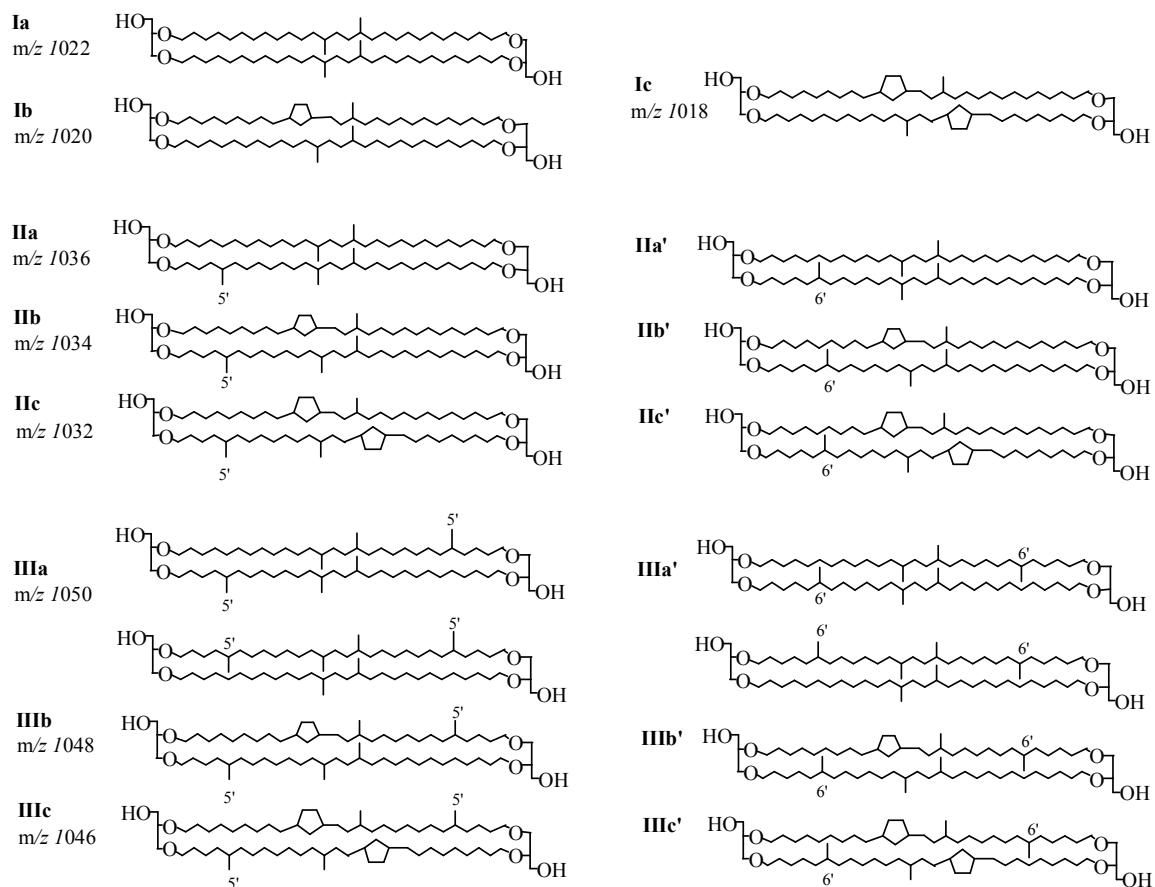
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611

612 **Fig. 1.** Molecular structures of 5- and 6-methyl branched GDGTs used in this study. The  
 613 compounds that have one or two methyl groups at the  $\alpha_6$  or  $\omega_6$  position are defined as  
 614 6-methyl brGDGTs, while the compounds that have one or two methyl groups at the  $\alpha_5$   
 615 or  $\omega_5$  position are defined as 5-methyl brGDGTs.

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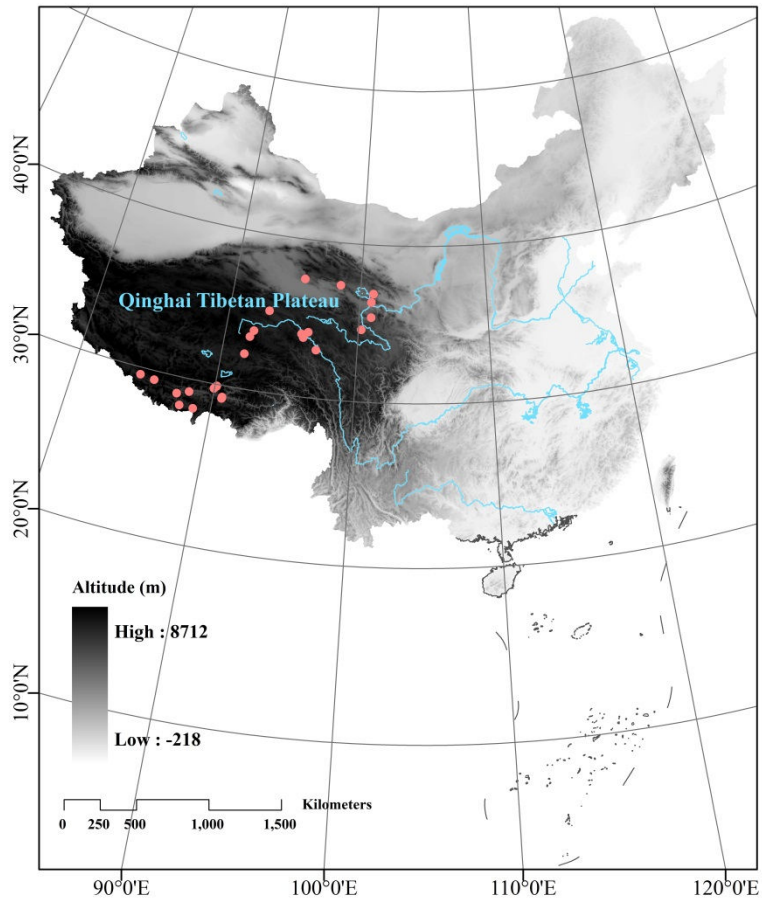
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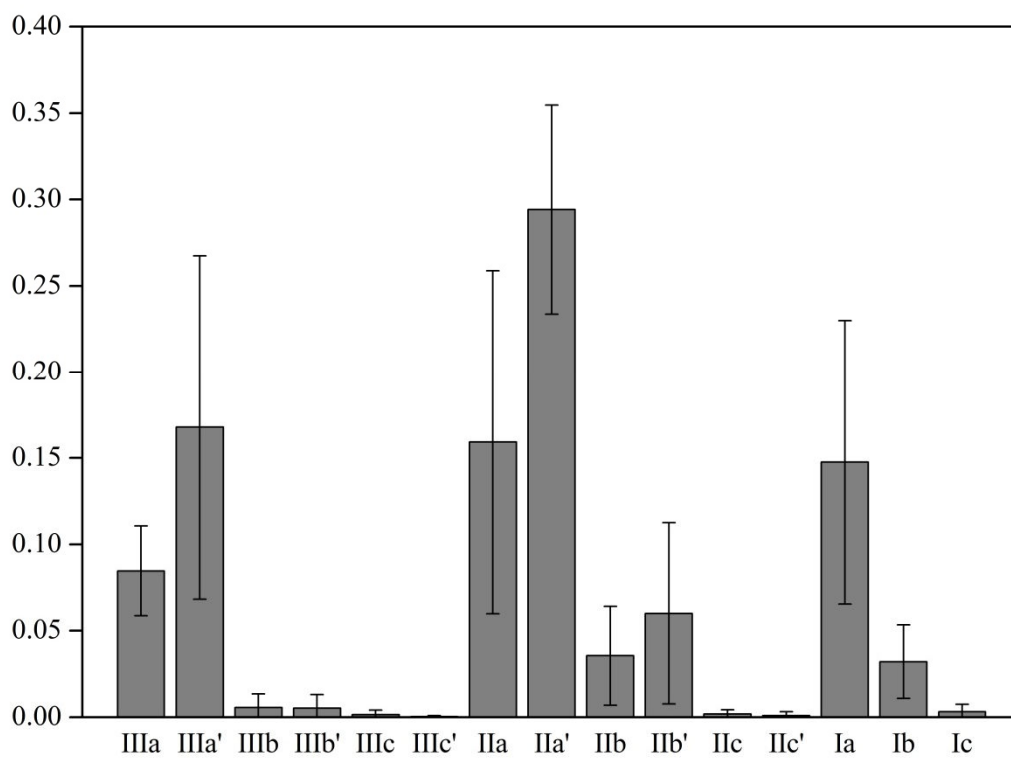
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**Fig. 2.** Locations of soil sampling sites (n = 30) in the QTP (Pink solid circles).



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641 **Fig. 3.** Average ( $n = 27$ ) fractional abundance of brGDGTs in surface soils of the QTP.

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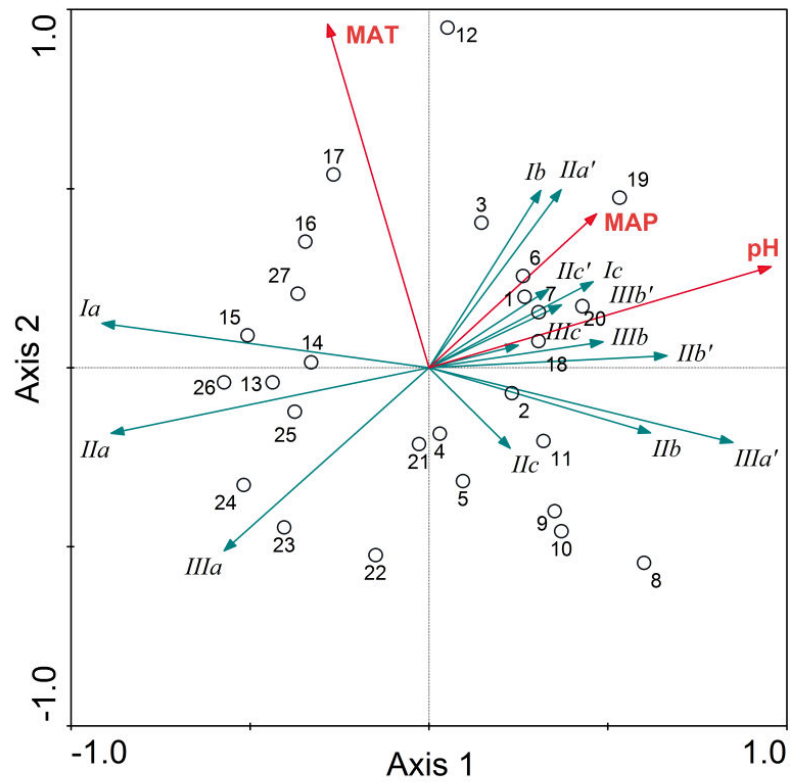
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658 **Fig. 4.** RDA triplot showing the relationship between 5- and 6-methyl brGDGTs%, MAT,  
 659 MAP and soil pH from the QTP. Numbers in the plot correspond to the soils in  
 660 supplementary material (Table S1). The first and second axis explained 65.2% and 6.1%  
 661 of the variance, respectively.

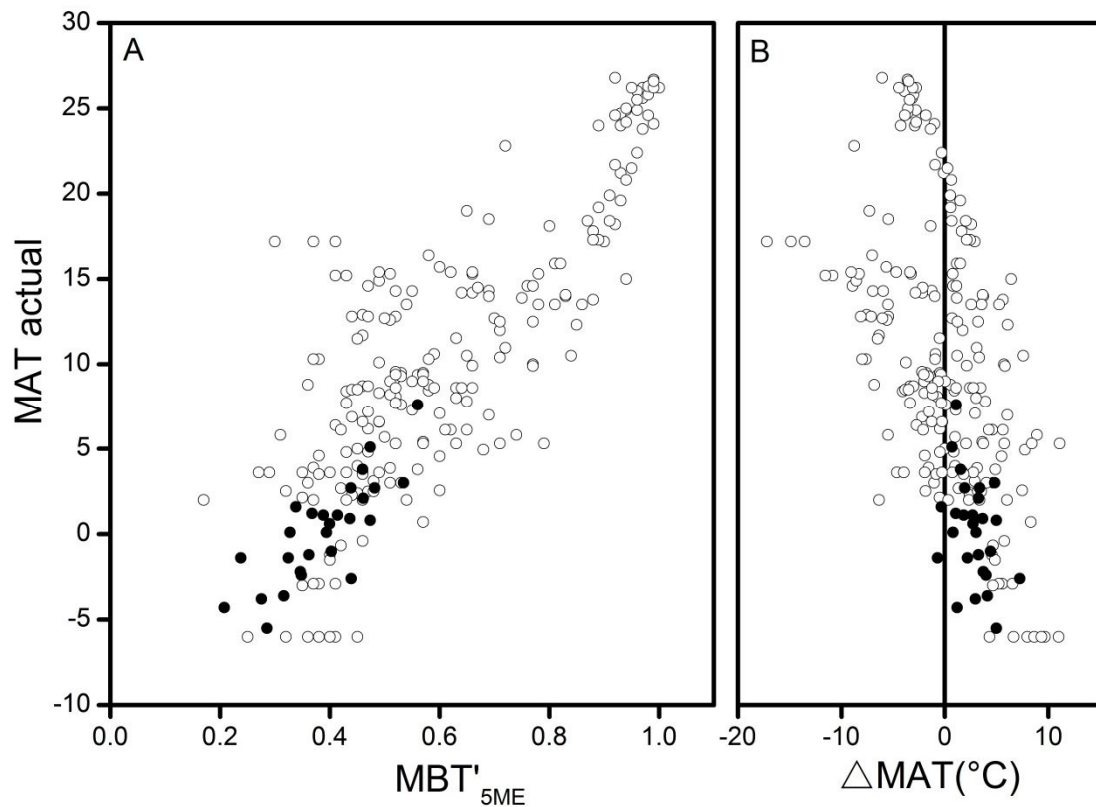
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668 **Fig. 5.** A) Scatterplot of  $MBT'_{5ME}$  with actual MAT; B: difference between estimated  
 669 MAT and actual MAT ( $\Delta MAT$ ). Solid and empty circles represented soils in this study  
 670 and global soils (de Jonge et al., 2014), respectively.

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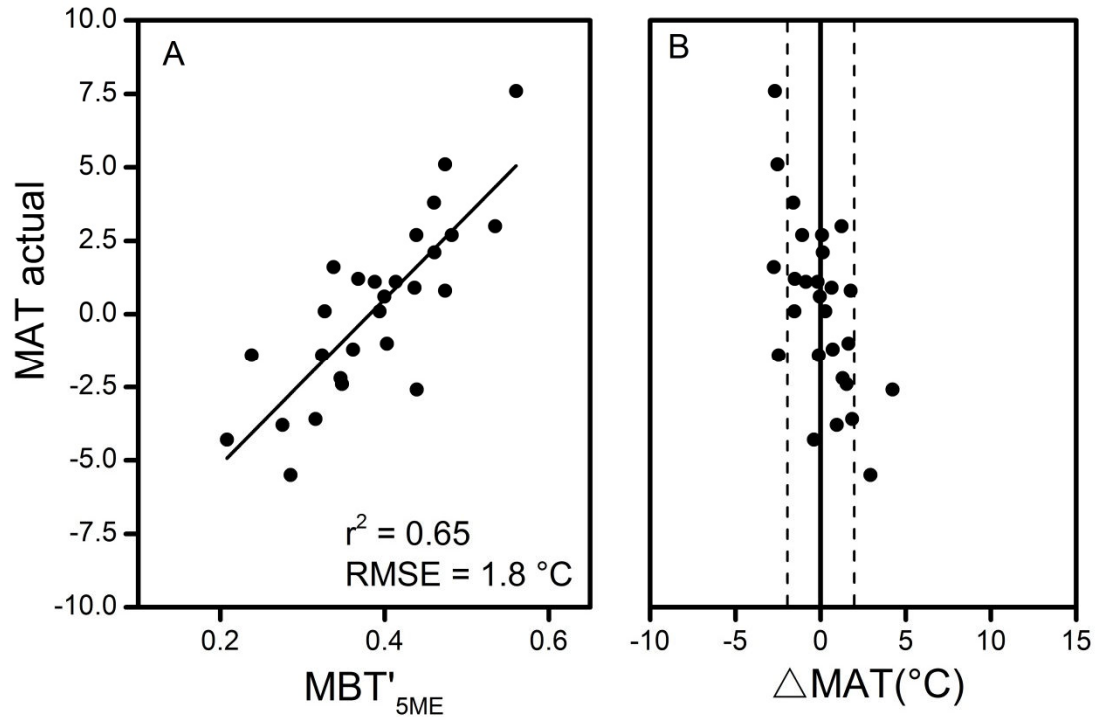
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678 **Fig. 6.** A) Linear regression of MBT'5ME with actual MAT; B) difference between

679 estimated MAT and actual MAT ( $\Delta\text{MAT}$ ). Data are from this study for 27 surface soils

680 of the QTP.

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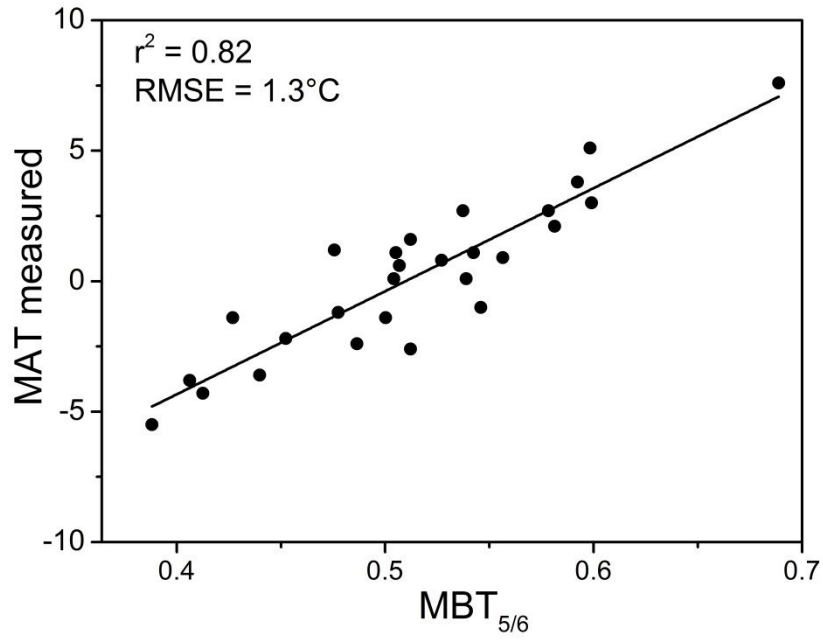
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694 **Fig. 7.** Linear regression plot of MBT<sub>5/6</sub> versus MAT in the QTP.

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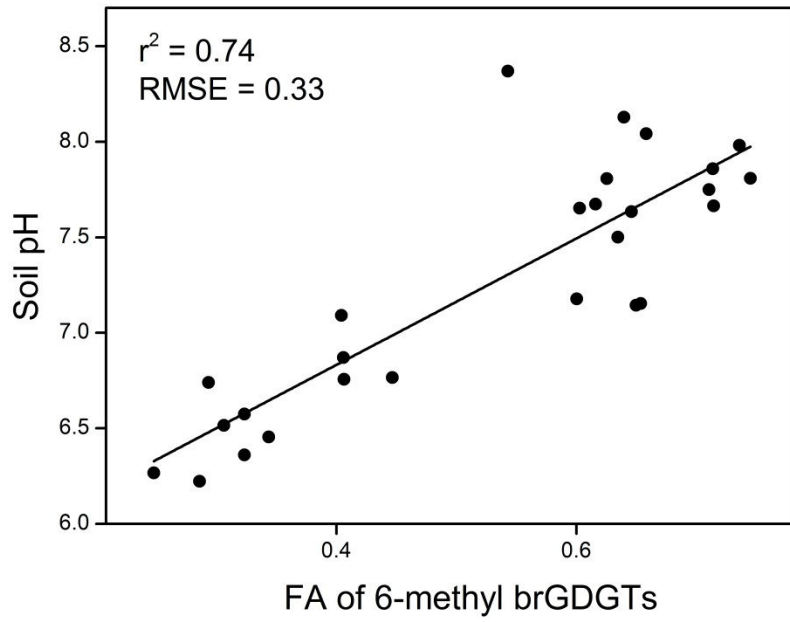
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709 **Fig. 8.** Plots of fractional abundance of 6-methyl brGDGTs of the total amount of  
 710 brGDGTs ( $f_{6ME}$ ) versus soil pH in the QTP.

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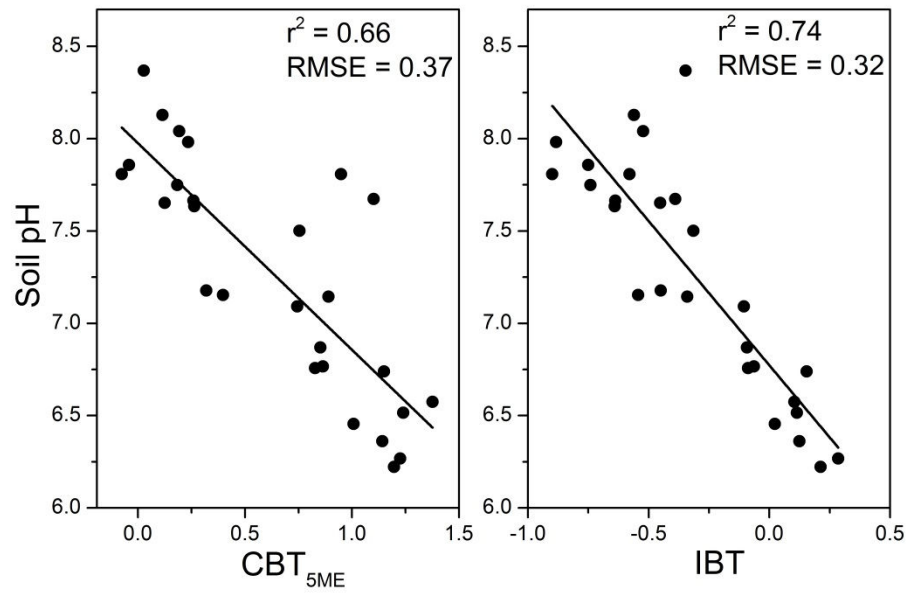
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726 **Fig. 9.** Scatterplots of A) soil pH versus  $CBT_{5ME}$  and B) soil pH versus IBT based on 27  
 727 soil samples in the QTP.

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746 **Table 1:** Results of RDA and partial RDA (pRDA) showing the total and unique  
 747 contributions of soil pH, MAT and MAP to the variance in brGDGT distributions in the  
 748 QTP soils.

Variables	Total contribution (%)		Unique contribution (%)	
	<i>p</i> Value	Max eigenvalues*	<i>p</i> Value	Eigenvalues
pH	0.001	60.1	0.001	39.9
MAT	0.001	16.4	0.001	10.6
MAP	0.179	10.8	0.172	2.0
All variables	0.001	72.9		
Joint effects				20.4
MAT + pH	0.001	56.4		
MAT + MAP	0.001	12.8		
MAP + pH	0.001	62.1		

749 \*The first environmental variable which has been selected into the analysis has the maximum eigenvalues (explained  
 750 variances), there are 6 sequences with different arrangement of pH, MAT and MAP. However, no matter which  
 751 sequence has been selected for RDA, the total variables contribution is invariant.

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