- An improved regional branched GDGT-based soil temperature calibration for
- 2 the Tropical Andes of Colombia: Towards a soil calibration for the tropics.

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# **Key Points:**

- New regional air and soil brGDGT calibrations for the Eastern Cordillera of Colombia have root-mean-square-errors of 1.1 °C and 1.2 °C, respectively.
- A soil calibration for the tropics has a root-mean-square-error of 2.2 °C, compared with global calibrations with root-mean-square-errors of 3.8 4.9 °C.

### Abstract

Branched glycerol dialkyl glycerol tetraethers (brGDGTs) are bacterial cell membrane lipids that, when preserved in sedimentary archives, can be used to infer continental paleotemperatures from sedimentary records. Although global calibrations capture a global relationship between the distribution of brGDGTs and temperatures, they underestimate temperatures for tropical regions by ~15 °C. Furthermore, some global calibrations reach saturation at around 24-25 °C, and, in general, they have root-mean-squared errors (RMSEs  $\approx$  ~4 °C) that are too large to resolve small variations in paleoclimate variability in tropical regions. We present a detailed regional calibration of soil brGDGTs along altitudinal transects on both flanks of the Eastern Cordillera of Colombia in the northern tropical Andes that spans ~3200 m in elevation, 17 °C in mean annual soil temperatures, and 19 °C in mean annual air temperatures. These new soil and air regional calibrations yield root-mean-square errors (RMSEs) of 1.2 °C and 1.1 °C, respectively. When combined with existing data from elsewhere in the tropics, the integrated data not only fit a calibration with an RMSE of 2.2 °C, but also fit mean annual air temperatures as high as 27.9 °C, resulting in a calibration that can be used in paleoclimate studies throughout the tropics.

### **Plain Language Summary**

Small variations in temperatures within the tropics can have large effects on climate at mid- and high-latitudes. A common example is how the El Niño Southern Oscillation (ENSO) can alter climate around the world. Quantitative terrestrial temperature records from the tropics, however, are scarce. A new lipid-based proxy, based on the abundances of different bacterial lipids, has enabled quantitative estimates of continental temperature, but the uncertainties in these global estimates remain large (3.8-4.9 °C). Because past temperature changes in the tropics are small

relative to other regions, high-precision temperature estimates are necessary to evaluate small changes that could have bigger effects elsewhere. In this study, we generated new data from the Eastern Cordillera of Colombia to estimate air and soil temperatures using this lipid-based proxy. In order to improve temperature reconstructions within the tropics, we compiled tropical soil data to provide a calibration for this latitudinal range.

### 1. Introduction

Small variations in temperatures within the tropics can have large effects on climate at higher latitudes, as illustrated by teleconnections from the Eastern Tropical Pacific and the El Niño Southern Oscillation (ENSO) (e.g., Halpert and Ropelewski, 1992; Ropelewski and Halpert, 1987, 1989; Sarachik and Cane, 2010; Trenberth et al., 1998). It follows that past mid-latitude climates may also reflect, and perhaps result from, variations in tropical conditions. Most quantitative estimates of past temperature from the tropics, however, come from marine sediment (e.g., Dekens et al., 2007; Groeneveld et al., 2006; Herbert et al., 2010, 2016; Lawrence et al., 2006; Ravelo et al., 2006; Wara et al., 2005; Zachos et al., 2001). Paleoclimate records from continental regions are scarce (e.g., Salzmann et al., 2013) and, in many cases, qualitative (e.g., Hooghiemstra et al., 2006; Van der Hammen et al., 1973). Missing has been a well calibrated proxy that quantifies past temperatures and can be applied widely to past tropical environments.

Recently, lipid-based paleotemperature proxies have evolved as a tool for the quantitative reconstruction of past continental temperatures (e.g., De Jonge et al., 2014; Lu et al., 2016; Peterse et al., 2012, 2014; Thomas et al., 2017; Weijers et al., 2007; Yamammoto et al., 2016). The development of these proxies has been particularly relevant in places where rocks with preserved organic matter may not preserve other materials useful for temperature

reconstructions, such as carbonates and fossil leaves (e.g., Ghosh et al., 2006; Wolfe, 1995). To enable accurate estimates of past temperatures in tropical environments, we calibrated a lipid-based paleotemperature proxy that exploits branched glycerol dialkyl glycerol tetraethers (brGDGTs). We measured both air and soil temperatures and brGDGTs along two altitudinal transect across the Eastern Cordillera of Colombia to construct regional calibrations for each. We then incorporated published data from elsewhere in the tropics to derive a pan-tropical calibration of air temperatures.

### 2. Background on brGDGTs

Branched glycerol dialkyl glycerol tetraethers (brGDGTs) are cell membrane lipids (Figure 1) produced by bacteria in soils, peats, lakes, marine sediment, and anoxic water columns (Liu et al., 2010, 2012, 2014; Russell et al., 2018; Sinninghe Damsté et al., 2000; Weber et al., 2015, 2018; Weijers et al., 2006, 2007). Early studies demonstrated that the distribution of brGDGTs in soils, expressed as the cyclisation of branched tetraethers (CBT, equation 1; Weijers et al., 2007) and the methyl index of branched tetraethers (MBT, equation 2, Weijers et al., 2007), correlated with the pH of the soil and mean annual air temperature (MAAT), respectively (Weijers et al., 2007):

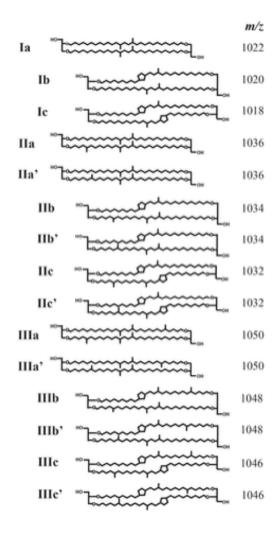
$$CBT = -log \frac{(Ib+IIb+IIb')}{(Ia+IIa+IIa')}, \text{ and}$$
 (1)

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$$MBT = \frac{(Ia+Ib+Ic)}{(Ia+Ib+Ic+IIa+IIa'+IIb+IIb'+IIc+IIa'+IIIa+IIIa'+IIIb+IIIb'+IIIc+IIIc')}. (2)$$

Prime numerals (e.g., IIa') indicate 6-methyl isomers, which were originally not separated from 5-methyl isomers (e.g., IIa) by the chromatographic method used in Weijers et al. (2007) and later in Peterse et al. (2012), and thus not used to distinguish these isomers in the

nomenclature, we distinguish between both isomers. Each term in the equations, and illustrated in Figure 1, uses the fractional abundance (FA) of these molecular structures, which is the ratio of the abundance of a particular brGDGT to the total number of brGDGTs in the sample. The MBT index was later refined by Peterse et al. (2012), who excluded brGDGTs-IIIb, -IIIb', -IIIc, and -IIIc' (Figure 1), because, in general, they are found in low abundance and below the analytical limit of detection. Peterse et al. (2012) defined the MBT' as follows:

$$MBT' = \frac{(Ia+Ib+Ic)}{(Ia+Ib+Ic+IIa+IIa'+IIb+IIb'+IIc+IIc'+IIIa+IIIa')}.$$
 (3)



**Figure 1.** Chemical structures of brGDGTs used in this study. Each letter corresponds to different structure from a methyl to a cyclopentyl moiety; 'a' structures have no cyclopentyl moieties, 'b' structures have one cyclopentyl moiety, and 'c' structures have two cyclopentyl moieties. The structures with C6-methylation are referenced with a prime symbol. C5-methylations are highlighted in blue in the corresponding structure. Figure modified from Hanna et al. (2016).

When incorporated in global soil datasets and calibrations, these indices allow for the reconstruction of past soil pH and MAAT using the following relationships (Peterse et al., 2012):

$$pH = 7.90 - 1.97 \text{ x CBT } (n = 176, R^2 = 0.70, RMSE = 0.8),$$
 (4)

MAAT =  $0.81 - 5.67 \times CBT + 31.0 \times MBT'$  ( $n = 176, R^2 = 0.59, RMSE = 5.0 °C$ ), (5)

where n is the number of samples used in the calibration,  $R^2$  is the coefficient of determination between MAAT and the combination of brGDGTs, and RMSE is the root mean square error for each temperature estimate using the respective calibration.

More recently, using an improved chromatographic method capable of separating 5- and 6-methyl isomers, De Jonge et al. (2014) demonstrated that although 5-methyl brGDGTs were mostly influenced by temperature, 6-methyl brGDGTs correlate with pH. By including only 5-methyl brGDGTs from global datasets, De Jonge et al. (2014) produced a new methylation index (MBT'<sub>5me</sub>):

$$MBT'_{5me} = \frac{(Ia+Ib+Ic)}{(Ia+Ib+Ic+IIa+IIb+IIc+IIIa)}.$$
 (6)

Then, they used this index to produce a simple linear equation that estimates MAAT, independently of soil pH:

128 MAAT = 
$$-8.57 + 31.45 \times MBT'_{5me}$$
 ( $n = 231, R^2 = 0.64, RMSE = 4.9 °C$ ). (7)

The MBT' $_{5me}$  index, as all previous MBT indices, is a fraction whose maximum possible value is one. When the MBT' $_{5me}$  index reaches one (MBT' $_{5me}$  = 1), Equation 7 allows for the reconstruction of MAAT only up to 22.7 °C, and therefore, this equation does not allow a reliable reconstruction of MAAT in tropical regions, which exceeds 22.7 °C in most regions. To expand the range of temperature that can be reconstructed, De Jonge et al. (2014) proposed a multiple linear regression using the fractional abundance (FA) of individual 5-methyl brGDGTs (Figure 1):

136 MAAT<sub>mr</sub> = 
$$7.17 + 17.1 \times [Ia] + 25.9 \times [Ib] + 34.4 \times [Ic] - 28.6 \times [IIa]$$
  
137  $(n = 222, R^2 = 0.68, RMSE = 4.6 \, ^{\circ}C)$ . (8)

Their measurements, however, extend only to ~26 °C, and extrapolating to higher temperatures must be done with caution.

More recently, Naafs et al. (2017) refined these global calibrations by filtering samples from the dataset using the relative abundance of 6-methyl brGDGTs (IR<sub>6me</sub> index) proposed by Dang et al. (2016):

$$IR_{6me} = \frac{[IIa'] + [IIb'] + [IIc'] + [IIIa'] + [IIIb'] + [IIIc']}{[IIa'] + [IIb'] + [IIIc'] + [IIIa'] + [IIIb] + [IIIc] + [IIIa] + [IIIb] + [IIIc]}. (9)$$

Dang et al. (2016) found that soils with ratios of 6- over 5-methyl brGDGTs that are lower than 0.5,  $IR_{6me} < 0.5$ , have a higher correlation between MBT' and MAAT. After excluding 173 soils from the global dataset of De Jonge et al. (2014) with  $IR_{6me} > 0.5$ , Naafs et al. (2017) proposed two new air temperature calibrations with higher  $R^2$  and lower RMSEs than those of De Jonge et al. (2014):

MAAT<sub>soil5me</sub> = -14.5 + 39.09 x MBT'<sub>5me</sub> (
$$n = 177, R^2 = 0.76, RMSE = 4.1 \, ^{\circ}C$$
), and (10)

MAAT<sub>mrlsoil5me</sub> = 
$$10.0 + 14.7 \text{ x [Ia]} - 31.7 \text{ x [IIa]}$$
 ( $n = 177, R^2 = 0.77, \text{RMSE} = 3.8 \,^{\circ}\text{C}$ ). (11)

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These recent calibrations, however, cannot be used to estimate temperatures above 24-25 °C, for when the MBT'<sub>5me</sub> = 1 in equation 10, the estimated temperature saturates at 24.6 °C. For equation 11, the maximum temperature that this calibration can estimate is 24.7 °C, when the FA of brGDGT-Ia = 1 and brGDGT-IIa = 0.

Despite these efforts to improve the global calibration of brGDGT proxies, the scatter in derived temperature estimates remains large (RMSEs  $\approx 3.8 - 4.9$  °C). Moreover, environmental factors besides temperature may also affect the production of brGDGTs through their source organisms and their distribution in the environment. Precipitation, soil moisture, vegetation cover, bacterial community composition, and growing degree days (GDD) of the source organism above freezing are all known sources of variability in the distribution of brGDGTs in

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soils (Dang et al., 2016, De Jonge et al., 2019; Menges et al., 2014; Naafs et al., 2017; Peterse et al., 2014, Wang et al., 2014). These environmental factors can vary regionally, and thus affect the sources and distribution of brGDGTs (Ding et al., 2015; Menges et al., 2014; Wang et al., 2016; Yang et al., 2015). Furthermore, the biological source of brGDGTs remains unknown. Acidobacteria were suggested to represent the most likely source of brGDGTs in a soil profile from the Saxnås Mosse peat bog (Weijers et al., 2009). But, only a small number of acidobacteria grown in culturing studies have been shown to produce brGDGTs, and then only a limited number of them (Sinninghe Damsté et al., 2011, 2018). Furthermore, Weber et al. (2018) showed that stratified lake systems can have a redox-dependent differentiation in the bacterial community in water columns with strong redox gradients, which impact the distribution of brGDGTs at different water depths. The biological sources of brGDGTs not only remain largely unknown, but these lipids have also been found in a wide range of environments. Therefore, given the large diversity of bacteria in soils, the few taxa that account for almost half of this diversity, and the role of environmental factors controlling their distribution (Delgado-Baquerizo et al., 2018), it is reasonable to assume that multiple biological sources of brGDGTs are possible in different types of soils.

Soil bacteria sense the temperature in the soil, not that of the overlying air. Therefore, scatter in inferences of MAATs may also derive from temperature differences between the soil and air. Finally, ignorance of whether bacteria grow in some, but not all, seasons and in how they sense seasonal variations in temperature adds uncertainty to inferences of temperatures, especially in global datasets spanning different latitudes with differing amplitudes of annual temperature variation (Dearing Crampton-Flood et al., 2020; Deng et al., 2016; Wang et al., 2016; Yang et al., 2014). The tropics provide a modest seasonal cycle, allowing temperature

differences between MAAT and, for example, the warmest month, to be close enough to neglect a possible seasonal uncertainty to infer temperature estimates with these calibrations.

### 3. Material and methods

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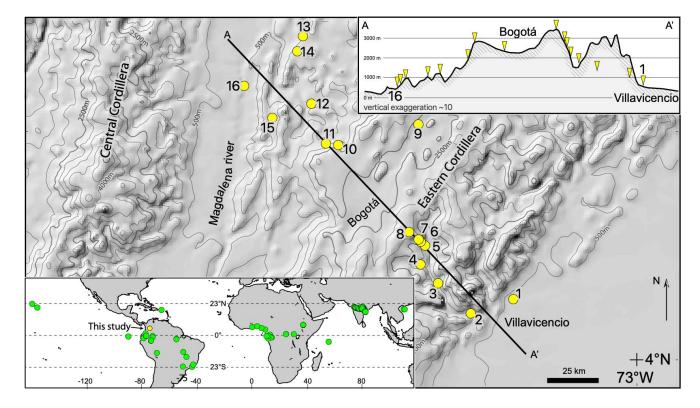
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## 3.1 Sampling in the Eastern Cordillera of Colombia

We collected 32 soil samples from 10 and 50 cm depth at 16 sites along two altitudinal transects spanning ~3200 m of elevation, one across the eastern flank (~400 to ~3400 m) and the other across the western flank (~200 to ~2600 m) of the Eastern Cordillera of Colombia (Figure 2). We also collected four more soil samples from a 30 cm depth at four sites on the western flank (218, 1281, 2185, 2589 m) to evaluate variations of brGDGTs with depth. Temperature data loggers (HOBO UA-001-08 8K) were installed at the same depths of 10 and 50 cm where we collected samples for analysis of brGDGTs, and we ensured that the loggers were fully inserted within the wall of the hole at the appropriate depths. We attached a string to each logger and the holes were covered with the same material that was dug out, making sure that the end of the string was left exposed at the surface. Additionally, we installed air-temperature loggers at each site ~2 meters above the ground from the closest tree and at a distance <~15 m from the soil loggers. Soil and air temperature measurements were recorded every two hours from August 2017 to August 2018 (Table S1 in the supporting information). We recovered all loggers, except for the two soil loggers from the lowest (~ 400 m) site on the eastern flank. Soil pH was measured in the field using an Oaktan waterproof pH 150 meter and a WD-35614-30 probe. We mixed ~2 g of soil with milli-Q water in a 1:2.5 (soil:water) ratio before each measurement and used a new container for each sample. We calibrated the pH-meter at every third station using the buffers with pH 4.00, 7.00, and 10.00.



**Figure 2.** Study area. **a)** Map of the Eastern Cordillera of Colombia for which the regional calibration was generated; yellow symbols indicate the 16 sites with new data presented in this study. The black line indicates the altitudinal transect (from A-A') of the 16 soil sites shown in the insert in the upper right corner. **b)** World map indicating the locations of soils around the tropics used in the tropical compilation.

### 3.2 brGDGTs analysis of soil samples

We freeze dried, homogenized, and sieved (0.3 mm) the soil samples to remove roots, plant debris, and coarse particles. In order to achieve maximum recovery of lipids from each sample, we extracted ~8 g of soil sample twice with dichloromethane:methanol (DCM:MeOH 9:1 v:v) using an accelerated solvent extractor (ASE 200 DIONEX; 100 °C and 2000 psi). We evaporated total lipid extracts (TLEs) under a gentle N<sub>2</sub> stream using a Turbovap and then combined and removed elemental sulfur using copper pellets previously activated with HCl for at least 1 h. Later, we filtered the TLEs through a short Pasteur pipette filled with glass wool and sand:Na<sub>2</sub>SO<sub>4</sub> (8:2) to remove any particles and water. Samples were spiked with 250 ng of the

 $C_{46}$  GDGT internal standard (Huguet et al., 2006). Then, we dissolved the dry samples in 1ml of hexane:isopropanol (hexane:IPA 99:1 v/v) and sonicated, vortexed, and filtered them through a 0.45  $\mu$ m PTFE filter before injection.

We analyzed brGDGTs using a Thermo Scientific Ultimate 3000 high performance liquid chromatograph (HPLC) coupled to a Q Exactive Focus Orbitrap-Quadrupole high resolution mass spectrometer via an atmospheric pressure chemical ionization (APCI) source. We achieved the chromatographic separation, identification, and quantification of brGDGTs by using a slighly modified version of the protocol described by Hopmans et al. (2016). Rather than starting at 18% hexane:isopropanol (9:1, v/v) (Hopmans et al., 2016), we began the eluent gradient with 30% hexane:isopropanol (9:1, v/v) to shorten overall run times without compromising the chromotographic separation of brGDGTs (Crump et al., 2019; Harning et al., 2019). We reequilibrated the HPLC column for 20 minutes between runs. The injection volume was 10µl instead of the 5µl in Hopmans et al. (2016). The positive ion APCI settings are in Table S2 in the supporting information. We analyzed samples on full-scan mode with a mass range of 500-1500 m/z at 70,000 mass resolution. We calibrated the Q Exactive within a mass accuracy range of 3 ppm using the Pierce LTQ Velos ESI Positive Ion Calibration Solution (Thermo Scientific). We identified brGDGTs based on their characteristic molecular ions and elution patterns and performed manual integration of the area under each peak using the Thermo Scientific Xcalibur software.

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### 3.3 Tropical soil dataset compilation

We compiled existing brGDGTs data from sites within 23.5° of the equator from De Jonge et al. (2014), Dearing Crampton-Flood et al. (2020), Jaeschke et al. (2018), and Naafs et al. (2017),

and combined them with the results from our study to create a soil brGDGT calibration that can be used for tropical paleoclimate reconstructions (Table S3 in the supporting information). We included only sites for which 5- and 6-methyl isomers were separated, their fractional abundance (FA) was available, and their modern MAAT was measured. We excluded four soils reported by De Jonge et al. (2014), also excluded by Naafs et al. (2017), because the elevation of those sites was unknown. De Jonge et al. (2014) and Naafs et al. (2017) inferred different values of MAAT, as large as 13 °C for the same site. Naafs et al. (2017) interpolated temperatures from a gridded database with 0.5° of resolution in both latitude and longitude: When the elevation of a soil site fell inside a cell grid but differed by more than 250 m from the average elevation of that grid cell, Naafs et al. (2017) used the temperature value of the nearest grid cell, whose elevation differed from that of site by less than 250 m. We consider that this approach is inappropriate for the tropics, where the biggest temperature differences derive from differences in elevation that can vary across short distances. Therefore, we decided to use the temperatures reported by De Jonge et al. (2014) because they relied on air temperatures from the closest meteorological station.

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### 3.4 Statistical analyses

We chose not to perform simple linear regression using the MBT'<sub>5me</sub> index, as this index can estimate temperatures only up to 22.7 °C (De Jonge et al., 2014) and 24.6 °C (Naafs et al., 2017). Instead, we carried out multiple linear regressions to derive equations to estimate temperatures, first for the Eastern Cordillera of Colombia and then for the tropics as a whole. The original dataset from De Jonge et al. (2014) excluded all 6-methyl brGDGTs because of their lack of correlation with temperature. We also excluded 5-methyl brGDGTs-IIIb and -IIIc, as they have

been shown to be present in only 26% of global soils (Peterse et al., 2012). We computed the Akaike Information Criterion (AIC, equation 12, Akaike, 1974) for various candidate relationship that relate brGDGTs to MAAT, each with different combinations of predictors. In this case, the predictors were FAs of individual brGDGTs, and the equation with minimum AIC value was selected to be the best fit to the data. The AIC is computed as follows:

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$$AIC = -2L + k*log(n),$$
 (12)

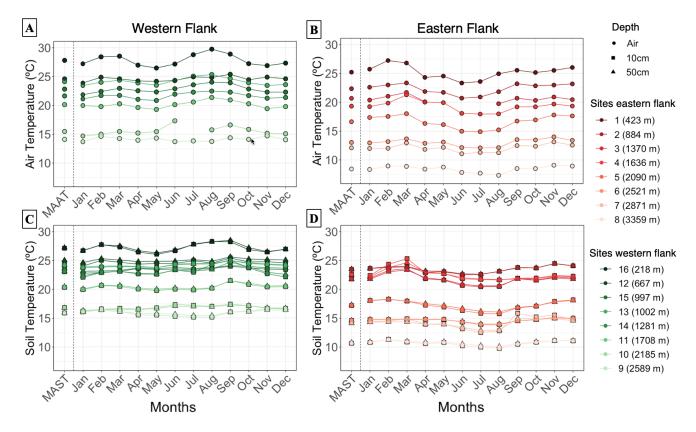
where k is the number of parameters estimated by the equation, n is the number of observations, and L is the log-likelihood function. The first term corresponds to the mean squared error (i.e. goodness of fit), and the second term is a penalty for complexity of the equation (i.e. the number of parameters). When comparing two different equations, we used the analysis of variance (ANOVA) to determine whether the difference between their means was statistically significant.

### 4. Results and Discussion

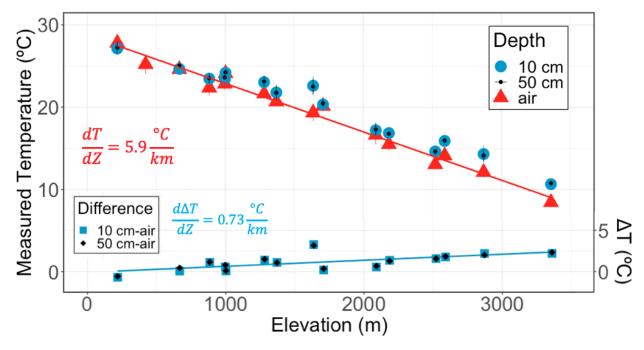
## 4.1 Logger temperature data from the Eastern Cordillera of Colombia

Mean monthly air temperatures (MMATs) and mean monthly soil temperatures (MMSTs) recorded by the loggers show a small temperature seasonality in this region, with the greatest difference between the mean warmest month and the mean coldest month for the same site of only 3.27 °C (Figure 3 and Table S1 in the supporting information). Average temperature gradients with elevation across the entire Eastern Cordillera of Colombia are 5.9 °C/km for air ~2 m above the ground, 5.2 °C/km for soil at a 10 cm depth, and 5.3 °C/km for soil at a 50 cm depth (Figure 4). The difference between soil (for both 50 cm and 10 cm) and air temperatures increases with elevation from ~ -0.5 °C to ~2.5 °C (indicating an elevation gradient of the

difference of 0.73 °C/km; Figure 4). Since the mean annual soil temperatures (MASTs) measured at both 10 and 50 cm at each station of the Eastern Cordillera of Colombia were almost identical, with an average absolute difference of 0.1 °C (Figure 4), we included temperatures from both depths in the following calibration.



**Figure 3.** Mean monthly temperatures recorded by data loggers from August 2017 to August 2018 along the two elevation transects in the Eastern Cordillera of Colombia. Panels (a) and (b) show the Mean Monthly Air Temperature (MMAT) for each flank. Panels (c) and (d) show the Mean Monthly Soil Temperature (MMST) at 10 and 50 cm depths for each flank. Each plot shows the mean annual air temperature (MAAT) and mean annual soil temperature (MAST) for each site on the left-hand side adjacent to the y-axis.

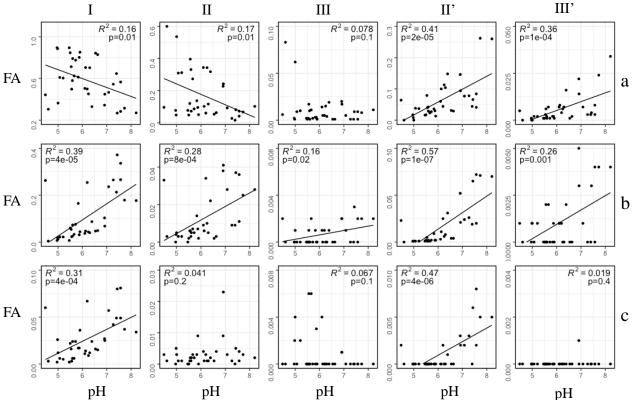


**Figure 4.** Mean *in situ* soil and air temperatures measured from August 2017 to August 2018 versus elevation. Top: points show mean temperatures at 10cm (blue circles) and 50cm (black circles), and overlying air temperatures (red triangles), with their respective monthly standard deviations. The red line shows a gradient of air temperature relative to elevation in the Eastern Cordillera of 5.9°C/km. Bottom: points show differences between mean temperatures of soil (at 10 cm: blue squares and at 50 black diamonds) and air. The blue line shows a gradient of the difference in temperature relative to elevation of 0.73°C/km.

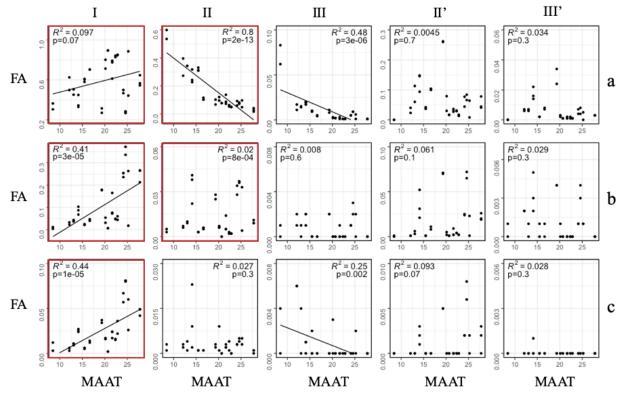
### 4.2 Regional soil brGDGTs calibration for the Eastern Cordillera of Colombia

We compare the new measured soil brGDGTs data from the Eastern Cordillera of Colombia with the measured pH and MAAT (Table S3 in the supporting information) in Figures 5 and 6, respectively. Figure 5 shows regressions between the fractional abundance (FA) of each individual brGDGT with pH. Most 6-methyl brGDGTs (those in columns II' and III' of Figure 5) show a higher correlation with pH than 5-methyl brGDGTs (those in columns II and III of Figure 5). The highest correlation with pH comes from the brGDGT-IIb' ( $R^2 = 0.57$ ). Consistent with previous work (De Jonge et al., 2014), 6-methyl brGDGTs correlate better with pH than with

MAAT, demonstrating that these isomers are not needed for the establishment of a temperature calibration and therefore can be excluded from it.



**Figure 5.** Regressions between fractional abundances (FA) of brGDGTs and measured soil pH from the Eastern Cordillera of Colombia. Roman numerals and letters on the right refer to the different brGDGT structures, II' and III' correspond to the structures with C6-methylation position. Linear regressions are shown for those brGDGTs with an  $R^2 \ge 0.1$ , with the corresponding values of p.



**Figure 6.** Regressions between fractional abundances (FA) of brGDGTs and measured mean annual air temperatures (MAAT) of each soil in the Eastern Cordillera of Colombia. Roman numerals and letters on the right refer to the different brGDGT structures, II and III correspond to the structures with C5-methylation position. Linear regressions are shown for those brGDGTs with  $R^2 \ge 0.1$ , with the corresponding values of p. Red boxes show the brGDGTs used in the regional air calibration (equation 14).

Putting together the modern soil brGDGTs data presented in Figure 6 and the MAAT and MAST derived from loggers (Figure 3), we developed air and soil temperature calibrations for the Eastern Cordillera of Colombia. Although we obtained similar AIC values (21.25 and 21.65) from the two best equations for the soil temperature calibration, we selected the one with the second lowest AIC value (equation 13):

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$$MAST = 18.79 + 4.07 \text{ x [Ia]} + 28.73 \text{ x [Ib]} - 17.3 \text{ x [IIa]} - 81.22 \text{ x [IIb]}$$

$$(n = 30, R^2 = 0.93, RMSE = 1.2 \text{ °C}),$$
(13)

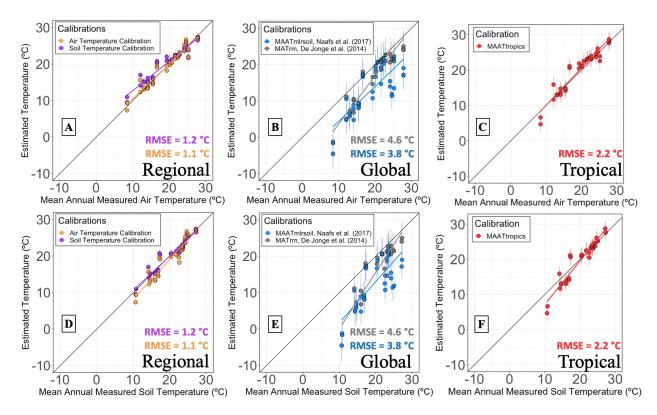
The selected equation includes brGDGT-Ia, which is a common and abundant brGDGT in warm environments. The two equations show no statistically significant difference in the variances captured. It is not uncommon to have equations with AIC values close to each other around the minimum value. Although a single best equation is identified, it is good to consider other close equations, such as the case here (e.g., Mendoza et al., 2014; Regonda et al., 2006, and references therein).

For the air temperature calibration, equation 14 yields the smallest AIC value:

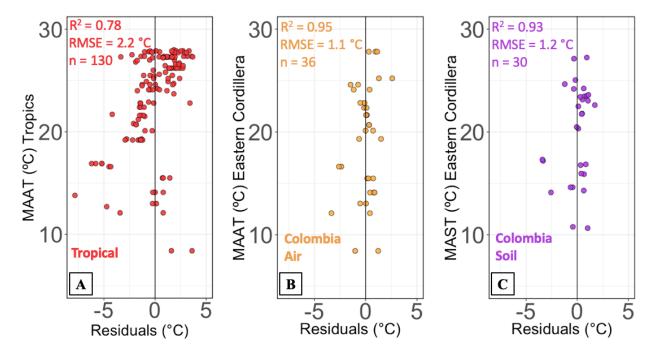
$$MAAT = 10.93 + 11.29 \text{ x [Ia]} + 32.39 \text{ x [Ib]} + 53.22 \text{ x [Ic]} - 11.84 \text{ x [IIa]} - 79.94 \text{ x [IIb]} (14)$$

$$(n = 36, R^2 = 0.95, RMSE = 1.1 \, ^{\circ}C).$$

Both regional MAAT and MAST calibrations reduce the RMSE substantially, from 4.6 °C (equation 8, De Jonge et al., 2014) and 3.8 °C (equation 11, Naafs et al., 2017), to 1.1 °C for MAAT and 1.2 °C for MAST (Figure 7a, b, d, e). Moreover, all air and soil temperatures implied by the regional calibrations differ from those measured with temperature loggers by less than 5 °C (Figure 8). By contrast, global calibrations underestimate air temperatures for the Eastern Cordillera of Colombia by up to 13.2 °C. Equations 13 and 14 provide a more precise reconstruction of temperatures in this region than the global calibrations (Figure 7). These regional calibrations not only allow for a larger range of temperature reconstruction than previous global calibrations (De Jonge et al., 2014: Naafs et al., 2017; Peterse et al., 2012), but also extend the temperature measurements within the calibration to higher values (27.8 °C for air and 27.5 °C for soil). MAAT and MAST estimates outside these calibration ranges, however, should be made with caution, as they rely on extrapolations.



**Figure 7.** Regressions of estimated temperatures against measured Mean Annual Air Temperatures (a, b, and c) and Mean Annual Soil Temperatures (d, e, f) from August 2017 to August 2018 in the Eastern Cordillera, using temperatures estimated with different brGDGT calibrations. In figures **a** and **d**, temperatures inferred from brGDGTs are compared using the regional calibrations for air temperatures (orange) (equation 14) and soil temperatures (purple) (equation 13). In figures **b** and **e**, temperature values are compared with global calibrations (equations 8 and 11) from De Jonge et al. (2014) (gray) and Naafs et al. (2017) (light blue). In panels **c** and **f**, temperature values are compared with the tropical calibration (equation 16). Black lines represent 1:1 relationships.



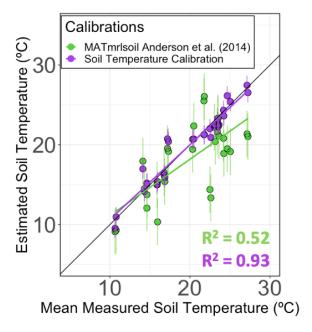
**Figure 8.** Residuals for the three multiple linear regressions calculated in this study. (a) the MAAT tropical calibration, (b) the air temperature calibration of the Eastern Cordillera of Colombia, and (c) the soil temperature calibration of the Eastern Cordillera of Colombia.

Previous work on the Eastern Cordillera of Colombia by Anderson et al. (2014), which used a chromatographic method that did not separate 5- and 6- methyl brGDGTs, yielded a local soil calibration with an RMSE of 2.9 °C (equation 15):

379 MASTsoil = 29.1 - 0.017 x (Ia) - 0.61 x log(Ib) - 3.34 x log(Ic) - 0.34 x (IIa+IIa') - 0.11 x log(IIb + IIb') + 0.44 x log(IIc + IIc') - 0.067 x (IIIa + IIIa') (15) 
$$(n = 24, R^2 = 0.77, \text{RMSE} = 2.9 \text{ °C}),$$

Although this equation resulted in an improvement over the RMSE of 5 °C from the global air temperature calibration (equation 5, Peterse et al., 2012), this value is still higher than what can be obtained when 6-methyl brGDGTs are chromatographically separated and excluded (equation 13, Figure 9). The influence of 6-methyl brGDGTs is particularly noticeable at the warm sites (>15 °C), where the calibration from Anderson et al. (2014) underestimates temperatures by as

much as ~9 °C. Furthermore, for sites where we measured MAST, we obtained an  $R^2 = 0.93$  with the soil calibration in equation 13 compared to the  $R^2 = 0.52$  for the calibration of Anderson et al. (2014). This improvement confirms the utility of separating 5- and 6-methyl brGDGTs when estimating temperatures.

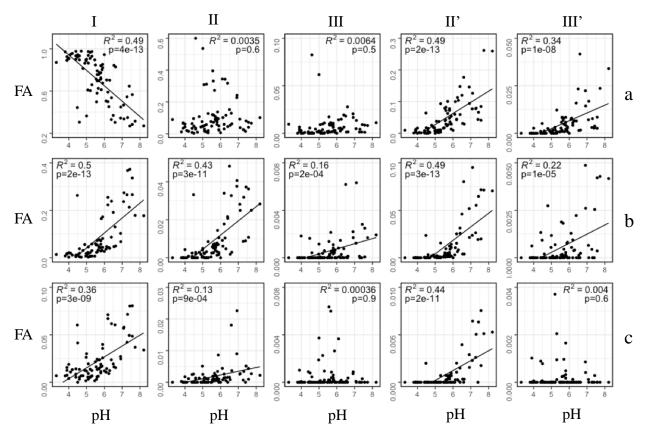


**Figure 9.** Comparison between the soil calibration from Anderson et al. (2014) (green, equation 15) and the regional soil calibration (purple, equation 13) with soil temperature measurements from the Eastern Cordillera of Colombia.

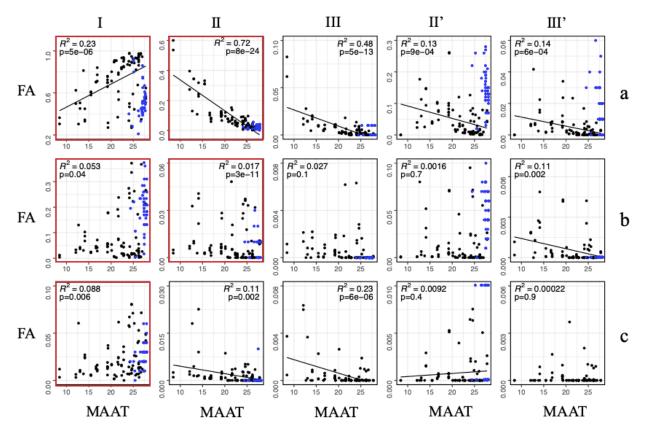
### 4.3 Integrating brGDGTs for a pan-tropical soil calibration

In order to integrate the new data obtained from the Eastern Cordillera of Colombia into datasets from elsewhere in the tropics, we compared the available measured pH and MAAT for each soil in the tropical compilation (Table S3 in the supporting information) with the FA of each individual brGDGT (Figures 10 and 11, respectively). From all soil data available, the regressions in Figures 10 and 11 only include 83 samples, as Dearing Crampton-Flood et al. (2020) reported only two decimal values for the FA of each brGDGTs and no pH measurement.

The 6-pentamethyl brGDGTs exhibit the highest correlation with pH in the 83 soil samples from the tropical compilation (Figure 10), as reported for our data (Figure 5) and as previously suggested by De Jonge et al. (2014). These isomers also exhibit an insignificant correlation with MAAT (e.g.,  $R^2 = 0.12$ ; p-value = 0.002 for IIa', Figure 11). Furthermore, only 2 out of 130 soil samples from this tropical compilation have a FA higher than 0.006 for 5-methyl brGDGT-IIIb and -IIIc (Figure 11). Since these samples represent only 1.5% of the tropical dataset, we excluded brGDGT-IIIb and brGDGT-IIIc from temperature calibrations.



**Figure 10.** Regressions between fractional abundances (FA) of brGDGTs and measured soil pH from the compilation of brGDGT measurements from the tropics. Only 83 out of 130 soil samples have reported pH measurements. Roman numerals and letters on the right refer to the different brGDGT structures, II' and III' correspond to the structures with C6-methylation position. Linear regressions are shown for those brGDGTs with an  $R^2 \ge 0.1$ , with the corresponding values of p.

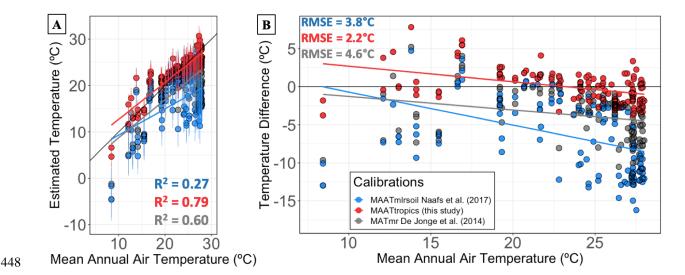


**Figure 11.** Regressions between fractional abundances (FA) of brGDGTs and measured mean annual air temperatures (MAAT) of each soil in the tropical compilation. Roman numerals and letters on the right refer to the different brGDGT structures, II' and III' correspond to the structures with C6-methylation position. Linear regressions are shown for those brGDGTs with  $R^2 \ge 0.1$ , with the corresponding values of p, and are done only with 83 soil samples. Data reported from India (Dearing Crampton-Flood et al., 2020) are shown in blue but were not used in the linear regressions for each brGDGT. Red boxes show brGDGTs used in the tropical calibration (equation 16).

brGDGT-IIa has the highest correlation with MAAT ( $R^2 = 0.71$  in 83 soil samples; Figure 11). In general, brGDGT-Ia has the highest FA in warm and tropical regions, but our results suggest that variations in temperature in these regions are better captured by brGDGT-IIa. Both of these brGDGTs, along with brGDGT-Ib, -Ic, and -IIb, correlate best with temperature in tropical areas where the seasonal cycle of temperature is small. When all tropical data are used, equation 16 fits data best, according to the AIC criterion:

 $MAAT_{tropics} = 17.55 + 7.45 \text{ x [Ia]} + 36.16 \text{ x [Ib]} - 28.89 \text{ x [Ic]} - 25.02 \text{ x [IIa]} - 69.22 \text{ x [IIb]}$  (16) 431  $(n = 130, R^2 = 0.78, RMSE = 2.2 \, ^{\circ}C).$ 432 Although equation 16 exhibits a larger RMSE (2.2 °C) than the regional air temperature 433 calibration (1.1 °C, equation 14), it is roughly half that of global calibrations (RMSE = 3.8-4.6 434 °C; Figure 7). 435 Global calibrations (equation 8, De Jonge et al., 2014; equation 11, Naafs et al., 2017) 436 underestimate temperature in tropical sites by as much as ~ -16 °C. The global calibration by De 437 Jonge et al. (2014) has the same  $R^2$ , 0.78, as our tropical calibration, but the RMSE of the 438 tropical calibration is less than half (2.2 °C) that of the global calibration (4.6 °C). When 439 considering sites with lower temperatures, which come from higher sites in the case of the 440 tropics, the tropical calibration slightly overestimates the temperatures at these sites (Figure 12). 441 A possible explanation for this overestimation is the limited number of sites (11 out of 130) with 442 a MAAT below 15 °C in the tropical compilation, which may lead to biases towards warmer 443 estimates. Another possible contribution to this underestimation could be the observed increasing 444 difference between soil and air temperatures with increasing elevation (Figure 4), where bacteria 445 living in high-elevation soils (>3000 m) register the warmer temperature of the soil compared to 446

the colder overlying air.

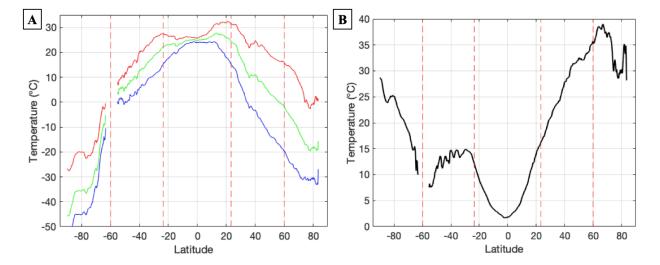


**Figure 12.** a) Comparison between tropical (equation 16) and global (equations 8 and 11) calibrations for air temperatures using all tropical sites. b) Differences between the estimated and the measured temperatures for each tropical soil versus measured MAAT (n = 130).

Global calibrations of MAAT rely on soil bacteria that live at different latitudes and that can grow at different times within a year. Whether bacteria that live in the soil grow all year round, or solely on the days above freezing, or during a specific season is still unknown. Therefore, temperature estimates that are calibrated with global MAAT could misrepresent the temperatures that bacteria record. If bacterial growth were biased towards warmer than colder days, global calibrations would underestimate temperature. In order to quantify the typical temperature variability in an annual cycle at different latitudes, we obtained terrestrial air temperature data from around the world for the 2000-2018 period from the European Centre for Medium Range Weather Forecasts (ECMWF) Reanalysis version 5 (ERA5) with a resolution of 0.25° in both latitude and longitude (Hersbach et al., 2019). Figure 13a shows the average MAAT (green), mean warmest month (red), and mean coldest month (blue) of terrestrial temperatures. Figure 13b shows the mean annual range of temperature by latitude, calculated as the difference between the warmest month and the coldest month for terrestrial temperatures. The tropics are

the only place where all of regressions of brGDGTs abundances against MAAT, mean annual temperature above freezing, and warmest months are similar enough that they can all represent the same value. Thus, lipid-based proxy calibrations might be better constrained by regional calibrations limited to restricted latitudinal ranges rather than global generalizations.





**Figure 13. a)** Zonally averaged mean air temperatures at 2 m above land areas (obtained from climate copernicus ERA5 data). Mean annual air temperature (green), mean warmest month (red), and mean coldest month (blue). **b)** Zonally averaged mean annual range of air temperature calculated as the difference between the warmest month and the coldest month for each site.

### 5. Conclusions

We measured temperatures in soils at depths of 10 cm and 50 cm, and in the air 2 m above the ground along profiles spanning 3200 m on the eastern and western sides of the Eastern Cordillera of Colombia. We also took soil samples from the same depths and measured the fractional abundances of the different brGDGTs in them with the goal of improving calibrations of temperatures inferred from brGDGTs to *in situ* temperatures. We obtained linear equations 13 and 14 relating the fractional abundances of certain brGDGTs to soil and overlying air

temperatures with RMSE of 1.2 °C and 1.1 °C, respectively. Combining these data with published measurements from other sites in the tropics, with latitudes < 23.5°, we obtained a calibration of air temperature (equation 16) with an RMSE of 2.2 °C for use in other tropical locations. These calibrations contrast with, and improve upon, global calibrations with RSME or 3.8 – 4.9 °C. Tropical calibrations are not subjected to possible seasonal biases as mean monthly temperatures differ little from the mean annual values. Moreover, these calibrations extend the temperature range accessible by brGDGTs to temperatures typical of low-latitudes, which is necessary for brGDGTs to be used to infer past temperatures from tropical environments, especially during intervals of greenhouse conditions, such as the Early Pliocene, Eocene, and Cretaceous.

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## **Data Availability Statement**

- Soil dataset is currently being archived in the Pangaea data repository and will be made available
- at acceptance.
- Global climate data used in this study are freely and publicly available online, and may be
- accessed directly as:
- 514 ECMWF Reanalysis version 5:
- 515 https://cds.climate.copernicus.eu/#!/search?text=ERA5&type=dataset

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