Assessing branched tetraether lipids as tracers of soil organic carbon transport through the Carminowe Creek catchment (southwest England)

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Abstract. Soils represent the largest reservoir of organic carbon (OC) on land. Upon mobilisation, this OC is either returned to the atmosphere as carbon dioxide (CO_2), or transported and ultimately locked into (marine) sediments, where it will act as a long-term sink of atmospheric CO_2. These fluxes of soil OC are, however, difficult to evaluate, mostly due to the lack of a soil-specific tracer. In this study, a suite of branched glycerol dialkyl glycerol tetraethers (brGDGTs), which are membrane lipids of soil bacteria, is tested as specific tracers for soil OC from source (soils under arable land, ley, grassland and woodland) to sink (Lake Loe Pool sediments) considering a small catchment located in southwest England (i.e. Carminowe Creek draining into Lake Loe Pool). The analysis of brGDGTs in catchment soils reveals that their distribution is not significantly different across different land use types ($p > 0.05$), and thus does not allow tracing land use-specific soil contributions to Lake Loe Pool sediments. Furthermore, the significantly higher contribution of 6-methyl brGDGT isomers in creek sediments (isomerization ratio (IR) = 0.48 ± 0.10; mean ± s.d., standard deviation; $p < 0.05$) compared to that in catchment soils (IR = 0.28 ± 0.11) indicates that the initial soil signal is substantially altered by brGDGT produced *in situ*. Similarly, the riverine brGDGT signal appears to be overwritten by lacustrine brGDGTs in the lake sedimentary record, indicated by remarkably lower Methylation of Branched Tetraethers (MBT'_{5ME} = 0.46 ± 0.02 in creek bed sediment and 0.38 ± 0.01 in lake core sediment; $p < 0.05$) and higher Degree of Cyclisation (DC = 0.23 ± 0.02 in creek bed sediment and 0.32 ± 0.08 in lake core sediment). Thus, in this small catchment, brGDGTs do not allow us to trace soil OC transport. Nevertheless, the downcore changes in the degree of cyclisation and the abundance of isoprenoid GDGTs produced by methanogens in the Lake Loe Pool sediment do reflect local environmental conditions over the past 100 years, and have recorded the eutrophication history of the lake.

1 Introduction

Globally, around 1500–2000 Pg of carbon is stored in soils in the form of organic matter, which is about two times the amount of carbon in the atmosphere and three times the amount of carbon in vegetation (Janzen, 2004; Smith, 2008). Soil organic carbon (OC) plays an important role in the global carbon cycle, as subtle alterations in the soil OC reservoir may affect the concentration of atmospheric CO_2 and thus influence climate change (Davidson and Janssens, 2006). Atmospheric CO_2 that is fixed by plants through photosynthesis will be stored into soil OC pool, part of which will be transferred to streams and rivers. Upon fluvial discharge, soil OC is buried and locked into the marine or lacustrine sediment, where it will act as a long-term carbon sink. However, instead of a passive pipeline in the carbon cycle, rivers actually represent a dynamic channel, where part of the soil OC is respired back to the atmosphere, and another part may be stored in river bed or lake sediments before reaching the ocean (Cole et al., 2007; Battin et al., 2009; Aufdenkampe et al., 2011). Hence, it is hard to determine the exact amount of soil OC that is transported to the ocean, as the dynamic processes that soil OC undergoes

during transport, such as degradation and sequestration, are elusive. This is mostly due to the lack of a specific tracer to distinguish soil OC from the total pool of OC that is also comprised of plant-derived OC, aquatic produced OC, and fossil OC from rock erosion (Blair et al., 2004; Aufdenkampe et al., 2011).

To circumvent this problem, lipid biomarkers can be used to trace a specific part of the total OC pool in complex natural environmental systems (Brassell and Eglinton, 1986; Wakeham and Lee, 1993). For example, odd-numbered long chain *n*-alkanes derived from epicuticular plant waxes are widely used to detect the contribution of terrestrial OC to river-dominated marine sediments (Eglinton and Hamilton, 1967; Hedges et al., 1997; Fernandes and Sicre, 2000; Glendell et al., 2018). Similarly, lignin, an abundant biopolymer in vascular plants (Hedges et al., 1997), has been used to trace OC transport along the terrestrial-aquatic continuum by e.g., in the Mississippi River (Goñi et al., 1997; Bianchi et al., 2004), the Amazon River (Hedges et al., 1986, 2000; Feng et al., 2016), and Arctic rivers (Feng et al., 2013). However, these biomarkers are derived from vegetation, which, although land-derived, is not fully representative of soil OC. Thus, in order to specifically trace and quantify the pool of soil OC, another biomarker is needed.

Branched glycerol dialkyl glycerol tetraethers (brGDGTs; Fig. 1) are membrane spanning tetraether lipids synthesized by heterotrophic bacteria that thrive in soils and peats all over the world (Weijers et al., 2006a, 2007a; Naafs et al., 2017a). Although the exact producers of these lipids are still unknown, after the detection of a brGDGT and the presumed brGDGT precursor lipid *iso*-diabolic acid in Acidobacterial cultures (Sinninghe Damsté et al., 2011, 2014, 2018), it was assumed that this members of the phylum are the main source organisms of brGDGTs in soils. However, a biological source outside the phylum of Acidobacteria cannot be excluded (Sinninghe Damsté et al., 2018). The occurrence and relative distribution of brGDGTs in a global set of modern surface soils showed that they can have 4 to 6 methyl groups attached to their alkyl backbone, where the degree of branching increases in soils from colder areas. Furthermore, brGDGTs respond to changes in soil pH by forming up to 2 cyclopentane moieties following internal cyclisation, where a higher number of cyclopentane moieties corresponds to a higher soil pH (Weijers et al., 2007a). Initially, a combination of two proxies, the Methylation of Branched Tetraethers (MBT) index and Cyclisation of Branched Tetraethers (CBT) index, was proposed as a proxy to reconstruct the mean air temperature (MAT) and pH of a soil (Weijers et al., 2007a; Peterse et al., 2012). After the identification of novel brGDGT isomers that possess a methyl group at the α and/or ω 6 position rather than at position 5 (Fig. 1) and the improvement of the chromatography method used for brGDGT analysis, a modified temperature proxy, the MBT'_{5ME} was developed (De Jonge et al., 2013, 2014b). Furthermore, the relative abundance of 6-methyl brGDGT isomers, quantified as the Isomerization Ratio (IR), appeared to also relate to soil pH (De Jonge et al., 2014b). Indeed, the analysis of brGDGTs in peat profiles and loess-paleosol sequences has resulted in long-term continental paleotemperature records for various areas, e.g. in deglacial central China (Peterse et al., 2011) and northeast China (Zheng et al., 2017), and western Europe during the early Eocene (Inglis et al., 2017).

These brGDGTs have not only been found in soils, but also in coastal marine sediments, where they have been used as the terrestrial end-member in the Branched and Isoprenoid Tetraether (BIT) index that determines the relative contribution of fluvially supplied soil organic matter to marine sediments, where the latter is represented by amounts of the isoprenoid GDGT crenarchaeol (Hopmans et al., 2004). For example, the relative abundance of brGDGTs in a marine sediment core from the Bay of Biscay revealed the early re-activation of European rivers after the last deglaciation (Ménot et al., 2006). Furthermore, brGDGTs stored in continental margin sediments are assumed to represent an integrated climate signal of the nearby land, and have been used as such to generate temperature records of deglacial tropical Africa (Weijers et al., 2007b), and Pliocene North-Western Europe (Dearing Crampton-Flood et al., 2018).

Recently, however, brGDGTs have also been found to be produced in aquatic systems such as coastal marine areas (Peterse et al., 2009b; Sinninghe Damsté, 2016), rivers (Kim et al., 2012; Zell et al., 2013, 2014) and lakes (Sinninghe Damsté et al., 2009a; Tierney and Russell, 2009; Loomis et al., 2011, 2014; Schoon et al., 2013; Weber et al., 2015, 2018), which complicates the interpretation of brGDGT-based proxy records. A contribution of *in situ* produced brGDGTs in lakes or on the continental shelf may bias BIT index values towards a more terrestrial signal (e.g. Sinninghe Damsté et al., 2009; De Jonge et al., 2015). Aquatic production in coastal marine areas became first apparent upon comparison of brGDGTs in Svalbard fjord sediments and nearby soils. Whereas the brGDGT signal in the fjord sediments was dominated by compounds containing cyclopentane moieties, soils where characterized by brGDGTs without cyclisation (Peterse et al., 2009b). These substantially different brGDGT signatures in combination with the increasing concentration of brGDGTs towards the open ocean then pointed towards a contribution of *in situ* produced brGDGTs to the fjord sediments. Similarly, brGDGT distributions in lake sediments were found to differ from those in soils surrounding the lake (Sinninghe Damsté et al., 2009; Tierney and Russell, 2009), and generated temperature estimates that severely underestimated actual MAT, mostly due to a high relative abundance of hexamethylated brGDGTs (e.g. Tierney et al., 2010; Loomis et al., 2014; Weber et al., 2015). Finally, the presence of brGDGTs with a polar headgroup still attached in suspended particulate matter (SPM) of several large rivers (Zhang et al., 2012; Zell et al., 2013; De Jonge et al., 2014a) provided strong evidence for aquatic production, as these headgroups are thought to be lost within days after cell death (e.g. Harvey et al., 1986). Notably, these and subsequent studies proposed ways to recognize *in situ* production of brGDGTs in aquatic environments. For example, a high degree of cyclisation is an indicator of brGDGT production in coastal marine zones (Peterse et al., 2009b; Sinninghe Damsté, 2016), for which Sinninghe Damsté (2016) proposed that a weighed number of rings in tetramethylated brGDGTs, quantified as $\#rings_{tetra} > 0.7$ indicates a purely marine source of brGDGTs in continental margin sediments. In rivers, aquatic brGDGTs appear to be characterized by a relatively high contribution of 6-methyl brGDGT isomers, and can be quantified using the IR (De Jonge et al., 2014a).

Here we test brGDGTs as tracers for soil OC in Carminowe Creek catchment, a small catchment in southwest England. Previously, an attempt was made to follow OC transport from soil (source) to Lake Loe Pool, the final sink of this catchment, using a combination of stable isotopes of bulk soil OC and plant leaf wax *n*-alkanes as fingerprints for the different vegetation types present in the catchment (i.e. arable land, grassland, ley and woodland) (Glendell et al., 2018). Although most land use types had a distinct *n*-alkane fingerprint, OC derived from arable land and temporary grassland (ley) could not be distinguished (Glendell et al., 2018). Hence, by assuming a primary soil source of the brGDGTs, their analysis in the same samples may contribute to tracing soil OC from different land use types during transport in Carminowe Creek. Moreover, changes in GDGT distributions in a 50 cm long sediment core from Loe Pool may be used to infer changes in soil OC transport dynamics in the catchment over the past century, and potentially couple them to climate or anthropogenic activity related events in the catchment area.

2 Methods

2.1 Study site and sampling

An overview of the study area and sampling sites is given by Glendell et al. (2018). Briefly, the Carminowe Creek catchment is located in Cornwall in southwest England (50°14' N, 5°16' W), covers an area of around 4.8 km^2 and varies in elevation from 0 to 80 m above sea level (Fig. 2). It is divided into two subcatchments ('north' and 'south'). The two streams converge around 100 m before their joint outlet, and then flow into a natural freshwater lake Loe Pool (50 ha), which is separated from the Atlantic Ocean by a natural shingle barrier. The mean annual temperature (MAT) and mean annual precipitation (MAP) in this area are approximately 11 °C and 1000 mm $year^{-1}$, respectively. The land use in this studied catchment is dominated

by arable land and temporary grasslands (ley), which are under rotation. The steeper hillslopes are under permanent grassland, and riparian woodland covers the areas near the creek. For this study, 74 surface soil samples (0–15 cm) were collected along 14 hillslope transects, including 31 arable land sites, 14 permanent grassland sites, 24 temporary grassland (ley) sites and 5 woodland sites (Fig. 2). Riverbed sediments were collected at three locations along each of the two tributaries (upstream, midstream and downstream), and one more at the joint outlet. A 50 cm long sediment core was taken in the lake, about 150 m away from the joint outlet. The lake core has been dated by the activity of Caesium-137 (^{137}Cs), and it covers the last 100 years (Glendell et al., 2018).

2.2 Bulk soil properties

Total carbon contents were reported by Glendell et al. (2018). Soil pH was measured in this study using a pH meter in a soil to water ratio of 1:5 (w:v) after shaking for two hours.

2.3 GDGT extraction and analysis

In total, 74 soil samples, 7 creek bed sediment and 25 lake core sediment samples were analysed for GDGTs. First, 5–7 g of the soils or 3–5 g of the sediments were freeze dried and homogenized, after which they were extracted three times with dichloromethane (DCM) : MeOH (9 : 1, v/v) using an accelerated solvent extractor (ASE 350, Dionex^{TM}) at 100 °C and 7.7 × 10^6 Pa to obtain a total lipid extract (TLE). After addition of a known amount of C_{46} GDGT internal standard (Huguet et al., 2006), the TLEs were dried under a N_2 stream, and then separated into apolar and polar fractions by passing them over an activated Al_2O_3 column using hexane : DCM (9 : 1, v/v) and DCM : MeOH (1 : 1, v/v) respectively. The polar fraction, which contains the GDGTs, was evaporated to dryness under a gentle N_2 stream. After this, the samples were prepared for further analysis by re-dissolving them in a hexane : isopropanol (99 : 1, v/v) mixture, and filtration through a 0.45 μm polytetrafluoroethylene (PTFE) filter.

The GDGTs were analysed on an Agilent 1260 Infinity ultra high performance liquid chromatography (UHPLC) coupled to an Agilent 6130 single quadrupole mass spectrometer (MS) with settings according to Hopmans et al. (2016). The GDGTs were separated over two silica Waters Acquity UPLC BEH Hilic columns (1.7 μm, 2.1 mm x 150 mm) preceded by a guard column with the same packing. GDGTs were eluted isocratically at a flow rate of 0.2 ml min^{-1} using 82% A and 18% B for 25 min, followed by a linear gradient to 70% A and 30% B for 25 min, where A = hexane and B = hexane : isopropanol (9 : 1, v/v). Sample injection volumes were 10 μL. Ionization of the GDGTs was achieved by atmospheric pressure chemical ionization with the following source settings: gas temperature 200 °C, vaporizer temperature 400 °C, N_2 flow 6 L min^{-1}, capillary voltage 3500 V, nebulizer pressure 25 psi and a corona current of 5.0 μA. By scanning the $[M+H]^+$ ions (protonated mass) in selected ion monitoring (SIM) mode, the target compounds were detected at *m/z* 1302 (GDGT-0), 1292 (crenarchaeol), 1050 (brGDGT–IIIa), 1048 (brGDGT–IIIb), 1046 (brGDGT–IIIc), 1036 (brGDGT–IIa), 1034 (brGDGT–IIb), 1032 (brGDGT–IIc), 1022 (brGDGT–Ia), 1020 (brGDGT–Ib), 1018 (brGDGT–Ic), with *m/z* 744 for the internal standard. Quantitation was achieved by peak area integration of the $[M+H]^+$ ions in Chemstation software B.04.03.

2.4 GDGT proxy calculations

The roman numerals in following equations refer to the molecular structures of GDGTs in Fig.1. The ratios below were calculated based on the fractional abundances (indicated by using square brackets) of GDGTs. The BIT index was calculated according to Hopmans et al. (2004), and modified to also include 6-methyl brGDGTs:

$$BIT = \frac{[Ia]+[IIa]+[IIIa]+[IIa']+[IIIa']}{[Ia]+[IIa]+[IIIa]+[IIa']+[IIIa']+[crenarchaeol]} \quad (1)$$

The degree of methylation (MBT'_{5ME}) and relative abundances of tetra-, penta-, and hexamethylated brGDGTs were calculated following De Jonge et al. (2014b) and Sinninghe Damsté et al. (2016):

$$MBT'_{5Me} = \frac{[Ia]+[Ib]+[Ic]}{[Ia]+[Ib]+[Ic]+[IIa]+[IIb]+[IIc]+[IIIa]} \quad (2)$$

$$\%tetra = \sum[tetramethylated\ brGDGTs] = [Ia] + [Ib] + [Ic] \quad (3)$$

$$\%penta = \sum[pentamethylated\ brGDGTs] = [IIa] + [IIb] + [IIc] + [IIa'] + [IIb'] + [IIc'] \quad (4)$$

$$\%hexa = \sum[hexamethylated\ brGDGTs] = [IIIa] + [IIIb] + [IIIc] + [IIIa'] + [IIIb'] + [IIIc'] \quad (5)$$

Furthermore, the degree of cyclisation (DC) was calculated according to Baxter et al. (2019):

$$DC = \frac{[Ib]+2*[Ic]+[IIb]+[IIb']}{[Ia]+[Ib]+[Ic]+[IIa]+[IIa']+[IIb]+[IIb']} \quad (6)$$

The isomerization ratio (IR) is the ratio between penta- and hexamethylated 6-methyl brGDGTs and the total amount of both 5- and 6-methyl penta- and hexamethylated brGDGTs (De Jonge et al., 2014a):

$$IR = \frac{[IIa']+[IIb']+[IIc']+[IIIa']+[IIIb']+[IIIc']}{[IIa]+[IIa']+[IIb]+[IIb']+[IIc]+[IIc']+[IIIa]+[IIIa']+[IIIb]+[IIIb']+[IIIc]+[IIIc']} \quad (7)$$

2.5 Statistical analysis and data visualization

The statistical analysis and data visualization were undertaken in R programming (version 3.5.2) (R Core Team, 2018). Differences in the concentration of brGDGTs and brGDGT-based proxies between different land use types (i.e. arable land, grassland, ley and woodland), creek bed and lake core sediments were examined by one-way nested ANOVA under generalized linear model (GLM) followed by post-hoc analysis (Tukey HSD (honest significant difference) test), and were performed with package 'car', 'carData' and 'agricolae'. Differences were considered to be significant at level of $p < 0.05$. To show how close our sample mean is to the population mean, standard deviation is used (mean ± s.d.). To examine whether brGDGT signatures could distinguish soil OC derived from different land use types, principal component analysis (PCA) was performed with package 'FactoMineR' and 'factoextra'. The box plot and scatter plots were carried out with package 'ggplot2'.

3 Results

3.1 BrGDGTs in soils

Most of the brGDGTs were present in all soils. Only brGDGT–IIIc and brGDGT–IIIc' were always below the detection limit (peak height > 3x baseline), and brGDGT–IIc' was below the detection limit in 13 of the soils (three in arable land, four in grassland and six in ley). The brGDGTs were dominated by pentamethylated (49.4 ± 3.0%, mean ± s.d., standard deviation), followed by tetramethylated (39.7 ± 4.9%) and then hexamethylated brGDGTs (10.9 ± 2.6%; Table 1). The concentration of brGDGTs ranged between 0.1 and 1.7 μg g^{-1} soil, with average of 0.2 ± 0.1 μg g^{-1} soil in arable land, 0.6 ± 0.4 μg g^{-1} soil in grassland, and 0.4 ± 0.3 μg g^{-1} soil in ley (i.e. the temporary grassland). However, the concentration of brGDGTs in woodland was 3.0 ± 1.0 μg g^{-1} soil, which was significantly higher than that in other land use types (0.4 ± 0.3 μg g^{-1} soil; $p < 0.05$; Fig. 3a). The C-normalized concentration of brGDGTs in catchment soils ranged between 2.8 to 49.8 μg g^{-1} C, 8.1 ± 3.6 μg g^{-1} C in arable land, 11.2 ± 6.7 μg g^{-1} C in grassland, 10.5 ± 4.8 μg g^{-1} C in ley, and 37.6 ± 11.0 μg g^{-1} C in woodland (Fig. 3a; Table 1). The trend of the concentration of brGDGTs along the soil transects was not obvious.

BIT index values ranged from 0.57 to 1.00 among land use types (Fig. 3b), with an average value of 0.96 ± 0.03 in woodland, 0.90 ± 0.12 in ley, 0.88 ± 0.14 in grassland and 0.83 ± 0.09 in arable land (without significant differences, $p > 0.05$). However, the BIT values increased from hillslope to downslope along several transects in north catchment, while the BIT values show no clear trends in south catchment (Fig. A1). The MBT'_{5ME} ranged from 0.37 to 0.71 and was mostly similar between all land use types (0.48 ± 0.04; $p > 0.05$; Fig. 3c; Table 1). The degree of cyclisation between land use types was similar (DC = 0.23 ± 0.13; Fig. 3d; Table 1; $p > 0.05$), likewise, the IR ranged from 0.10 to 0.60 (0.28 ± 0.01 on average; Fig. 3e; Table 1; $p > 0.05$), without clear trend along the soil transects. However, four transects in the north catchment have on average significantly higher IR values (> 0.36) than the other transects in the catchment (0.24 ± 0.09; $p < 0.05$; Fig. A1). In general, the IR increases with increasing soil pH in the catchment ($r^2 = 0.36$, $p < 0.001$).

3.2 BrGDGTs in creek bed sediments

All brGDGT compounds were detected in creek bed sediments, except for in the upstream site from north catchment, where brGDGT–IIIc' was below detection limit. The brGDGTs in creek bed sediments were dominated by pentamethylated brGDGTs (45.0 ± 0.7%), followed by tetramethylated brGDGTs (30.1 ± 4.5%), and hexamethylated brGDGTs (24.9 ± 4.7%) (Table 1). The C-normalized concentration of brGDGTs in creek bed sediments was 34.7 ± 17.4 $\mu g\ g^{-1}$ C on average (Fig. 3a; Table 1), where the concentration increased from 32.7 $\mu g\ g^{-1}$ C to 57.0 $\mu g\ g^{-1}$ C downstream in north catchment, and from 14.3 $\mu g\ g^{-1}$ C to 25.2 $\mu g\ g^{-1}$ C downstream in south catchment, reaching a maximum value of 59.3 $\mu g\ g^{-1}$ C at the outlet (Fig. 5a). The concentration of brGDGTs in creek bed sediments was higher than that in soils under any land use types, except for woodland (9.6 ± 4.9 $\mu g\ g^{-1}$ C; Fig. 3a; Table 1).

The BIT values for creek sediments were on average 0.90 ± 0.06 (Fig. 3b; Table 1). The MBT'_{5ME} was relatively constant between 0.44 and 0.49, with an average of 0.46 ± 0.02. The DC ranged from 0.21 to 0.25 in the creek sediments with an average of 0.23 ± 0.02 (Fig. 3e; Table 1). The IR was relatively invariable with an average of 0.48 ± 0.10 (Fig. 3e; Table 1). The brGDGT-based proxies for creek bed sediments were similar to those for soils, except for the IR, which was higher than that in soils under any land use types (0.28 ± 0.11; Fig. 3; Table 1).

3.3 BrGDGTs in Lake Loe Pool sediment core

All brGDGTs were detected in the lake sediment core, except at 20 cm depth, where brGDGT–IIIc' was below the detection limit. The brGDGTs in the lake sediments were mainly dominated by pentamethylated brGDGTs (50.2 ± 1.8%), followed by tetramethylated brGDGTs (28.9 ± 0.7%), and hexamethylated brGDGTs (21.0 ± 1.4%; Table 1). The amount of brGDGTs in lake core sediment ranged from 19.9 to 48.0 $\mu g\ g^{-1}$ C (Fig. 3a; Table 1). The brGDGT concentration in the surface sediment (0–2 cm), of 37.7 $\mu g\ g^{-1}$ C, which was about 1.6 times lower than that in the creek sediment at the outlet (Fig. 5a), increased to a maximum of 48.0 $\mu g\ g^{-1}$ C around 11 cm depth, and then decreased to a minimum of 19.9 $\mu g\ g^{-1}$ C at 23 cm depth (Fig. 6b). The concentration of GDGT-0 ranged between 9.0 $\mu g\ g^{-1}$ C and 27.1 $\mu g\ g^{-1}$ C with an average of 17.4 ± 6.0 $\mu g\ g^{-1}$ C, concentration of crenarchaeol ranged from 0.6 $\mu g\ g^{-1}$ C to 1.4 $\mu g\ g^{-1}$ C with an average of 1.0 ± 0.2 $\mu g\ g^{-1}$ C in the lake sediment core. In general, the concentration of brGDGTs in lake core (34.0 ± 8.7 $\mu g\ g^{-1}$ C; Table 1) was similar with that in river and in woodland, while it was significantly higher than the brGDGTs in soils except for the woodland (9.6 ± 4.9 $\mu g\ g^{-1}$ C; $p < 0.05$; Fig. 3a; Table 1).

The BIT values for the lake sediment core were rather uniform, varying between 0.95 and 0.97 (Fig.3b). Similarly, the values of MBT'_{5ME} along the lake core ranged only between 0.36 and 0.39. The MBT'_{5ME} of 0.37 for the lake surface sediment was significantly lower than that in creek bed sediment (0.46 ± 0.02; $p < 0.05$; Fig 3c; Fig. 5b). Conversely, the DC in the lake surface sediment was 0.39, which was significantly higher than that in creek bed sediment (0.23 ± 0.02; $p < 0.05$;

Fig. 3d; Fig. 5b). The average value of DC for the lake core sediments was 0.32 ± 0.08. The DC increased from the surface to a maximum value (0.44) at around 10 cm depth, and then decreased with slight fluctuations to 0.22 at 43 cm depth (Fig. 6c). The IR was constant downcore (0.32 ± 0.01 on average; Fig. 3e; Table 1) and was significantly lower than that in creek bed sediment ($p < 0.05$; Fig. 3e).

4 Discussion

4.1 Spatial variation of brGDGT signals in catchment soils

Spatial variations in the relative distribution of brGDGTs in all catchment soils were first evaluated by performing principal component analysis (PCA) using the fractional abundances of the 13 major brGDGTs detected. The first two principal components (PCs) explain 65.2% of the variance in the dataset. PC1 describes 49.5% of the variance, and separates acyclic brGDGT–Ia and brGDGT–IIa from all the other brGDGTs (Fig. 4a). In line with this observation, PC1 has a strong positive relationship with the degree of cyclisation of brGDGTs in the soils ($r^2 = 0.97$; Fig. 4c). PC2 describes another 15.7% of the variance, and separates tetramethylated brGDGTs as well as most of the 6-methyl brGDGTs from the majority of the 5-methyl penta- and hexamethylated brGDGTs. As a result, PC2 is negatively correlated with MBT'_{5ME} ($r^2 = 0.49$; Fig. 4d) as well as the IR ($r^2 = 0.58$; Fig. 4e) in soils. Despite the clear relation of the first two PCs with the degree of cyclisation and the degree of methylation, respectively, the position of the soils in the PCA diagram reveals that different land use types are largely overlapping (Fig. 4b). Indeed, the brGDGTs proxies for different land use types are not significantly different ($p > 0.05$; Fig. 3), making it difficult to distinguish the provenance of soil OC solely based on brGDGT signatures.

Indeed, previous work has also shown that brGDGT distributions are not primarily affected by land use. For example, brGDGTs in soils along an altitudinal transect in the Ethiopian highlands revealed that brGDGTs mainly reflect the decrease in temperature with increasing elevation, regardless of drastic changes in land use along the transect (Jaeschke et al., 2018). However, other studies report that vegetation cover does exert a great influence on brGDGT signatures in soils from Minnesota and Ohio, USA (Weijers et al., 2011), around Lake Rotsee, Switzerland (Naeher et al., 2014), in the Tibetan Plateau (Liang et al., 2019), and paddy and upland soils from subtropical (China and Italy) and tropical (Indonesia, Philippines and Vietnam) areas (Mueller-Niggemann et al., 2016). The explanations for the similar distribution of brGDGTs under different land use types in the Carminowe Creek catchment could be the rotation and ploughing in land use in combination with the turnover time of brGDGTs. Although the soil bacterial community composition is generally different across distinct land use types (Fierer and Jackson, 2006; Steenwerth et al., 2003), the regular rotation (generally less than 5 years) of arable land and temporary grassland (ley) in the catchment (Glendell et al., 2018) may create a mixed bacterial community under all vegetation types. Beyond vegetation, regular ploughing as applied across the Carminowe catchment soils (arable land and ley) is recognized to have a more dominant, long-last effect on microbial communities (Drenovsky et al., 2010). Moreover, brGDGTs in terrestrial environments have a relatively long turnover time (ca. 18 years in soils (Weijers et al., 2010), and up to 40 years in peat (Huguet et al., 2017)), especially when compared to the cropland rotation time. Taken together, these factors may contribute to the relatively similar brGDGT signal in all soils in the Carminowe catchment, further limiting the variation in brGDGT signals in catchment soils.

Some spatial trends are visible in spite of the overall comparable brGDGT signals across the catchment (Fig. A1), which may be explained by variations in other environmental factors than land use or vegetation. Mean air temperature and soil pH have been shown to be the main factors controlling the distribution of brGDGTs in soils worldwide (Weijers et al., 2007a; Peterse et al., 2012; De Jonge et al., 2014b). However, in the small (ca. 4.8 km^2) Carminowe Creek catchment, the annual mean air temperature is practically the same for all soils. Similarly, the range in soil pH is relatively small among different land use types (from 5.4 ± 0.3 in woodland to 6.6 ± 0.1 in arable land; Table 1), which makes it difficult to separate brGDGT

signals based on these parameters. Additionally, the soil water content (SWC) has been shown to affect the distribution and abundance of brGDGTs in soils, either directly by changing the microbial community, or indirectly by altering soil temperature, soil pH, or soil oxygen content (Dirghangi et al., 2013; Menges et al., 2014; Dang et al., 2016). The SWC positively correlates with the abundance of brGDGTs in soils from Qinghai-Tibetan Plateau (Wang et al., 2013), as well as in soils along an aridity transect in the USA (Dirghangi et al., 2013). Moreover, the degree of methylation of 6-methylated brGDGTs is sensitive to the SWC, especially in semi-arid and arid regions (Dang et al., 2016). Although MAP is also the same for the whole catchment, the subtle altitudinal differences in this small creek catchment (i.e. 0-80 m above sea level) may result in an increase in SWC from hilltop to downslope. This would introduce just enough variability in SWC to explain some of the trends in brGDGT signals along hillslope transects. In the north catchment, the BIT index values gradually increase from the presumably better aerated soils at the hilltops towards the wetter soils closer to the creek (Fig. A1). The increase is > 0.3 for Transects 1 and 8, but also Transects 2, 3, and 7 show an increase in BIT values downslope, albeit to a smaller degree (0.17, 0.19, and 0.04, respectively; Table A1). The change in BIT index values is driven by both an increase in the amount of brGDGTs and a slight decrease in crenarchaeol concentrations with the presumed increase in SWC downslope, similar to previous findings (Dirghangi et al., 2013; Wang et al., 2013; Menges et al., 2014). The trend in BIT is likely enhanced by the (minor) change in soil pH along Transects 1 and 8 (from 6.2 to 6.1 along Transect-1 and from 6.6 to 5.7 along Transect-8), which may influence the BIT index as a result of the generally positive relation of crenarchaeol concentrations and a negative relation of brGDGT concentrations with increasing soil pH (Weijers et al., 2006b; Peterse et al., 2010). Nevertheless, these trends in the BIT index are visible in five of the transects and only occur in the north part of the catchment.

Interestingly, the IR is also significantly higher in soils along four transects in north catchment (all > 0.36 averagely for Transects 1, 2, 7, and 8) compared to the average IR value for the rest of the transects in the entire catchment (0.24 ± 0.09; $p < 0.05$). The majority of the sites with higher IR are in cropland, except for those in the Transect-1, which is under grassland (Fig. A2). Although a relative increase in 6-methyl brGDGTs has been linked to higher soil pH in the global soil dataset (De Jonge et al., 2014b), this relation is not so strong in the soils from the Carminowe creek catchment ($r^2 = 0.36$, $p < 0.001$), likely due to the relatively minor range and variation in soil pH (from 5.4 ± 0.3 to 6.6 ± 0.1). Nevertheless, the soils with high IR values in the north catchment also have pH values > 6.0 with an average value of 6.6 ± 0.1.

4.2 Tracing brGDGTs from soils to creek bed sediments

Based on the similar brGDGT signatures for soils under different land use types, these compounds cannot be used to trace back the exact source of the soil OC after mobilisation and transport throughout the catchment. However, the concentration and general soil signature of the brGDGTs can be compared with those in creek bed sediments to trace the transfer of OC from the soils into the creeks. The C-normalized concentration of brGDGTs in the creek sediments is higher than that in most of the soils (34.7 ± 17.4 μg g^{-1} C and 9.6 ± 4.9 μg g^{-1} C respectively), except for those in the woodland soils at the riverbanks (37.6 ± 11.0 μg g^{-1} C; Table 1). Thus, purely based on the concentration, this suggests that brGDGTs in the creek would be primarily derived from the woodland, which also appeared to be the main source of *n*-alkanes in creek bed sediment (Glendell et al., 2018). However, when looking at the relative distribution of the brGDGTs, the percentage of hexamethylated brGDGTs in creek sediments is higher than that in soils (24.9 ± 1.8% and 10.9 ± 0.3%, respectively), whereas the percentage of tetramethylated brGDGTs is lower than in soils (30.1 ± 1.7% and 39.7 ± 0.6%, respectively; Table 1). Furthermore, brGDGTs in creek sediments have a significantly higher IR (i.e. 0.48 ± 0.04) than soils under any of the land use types (0.28 ± 0.01 on average in the catchment; $p < 0.05$; Fig. 3e; Table 1). This is clearly reflected in the PCA, which separates the creek sediments from both the soils and lake sediments on PC2 that is associated with the IR (Fig. 4e). The higher IR in the creek bed sediments can be explained by a contribution of aquatically (i.e. *in situ*) produced 6-methyl

brGDGTs. Similar contributions of 6-methyl brGDGTs, and thus higher IR, were also observed in suspended particulate matters from the Yenisei River (De Jonge et al., 2014a), and upstream of the Iron Gates in the Danube River, where the higher IR was coupled to in-river production facilitated by the lower flow velocity and decreased turbidity of the river water (Freymond et al., 2017). Hence, the significantly higher IR in combination with the higher C-normalized concentrations of brGDGTs in the Carminowe creek sediments suggests that the brGDGT signal is mainly aquatic.

In attempt to further prove the riverine *in situ* production of brGDGTs, we roughly estimate the minimum amount of 6-methyl brGDGTs that needs to be produced in the creek in order to reach the higher IR. We hereby assume that the brGDGTs derived from woodland soils are completely transferred into creek without any degradation. Thus, the concentration of 6-methyl brGDGTs in the creek sediments [6-me $_{creek}$] resembles the sum of the average concentration of 6-methyl brGDGTs in woodland soils [6-me $_{woodland}$] and those produced *in situ* [6-me $_{in\ situ}$]. The minimum amount of 6-methyl brGDGTs produced *in situ* can then be calculated using the brGDGT-concentration-weighed IR for creek sediments (IR_{creek} = 0.47) and the following equation (Eq. 8).

$$IR_{creek} = \frac{[6\text{-}me_{creek}]}{[5\text{-}me_{creek}] + [6\text{-}me_{creek}]} = \frac{[6\text{-}me_{woodland}] + [\mathbf{6\text{-}me_{in\ situ}}]}{[5\text{-}me_{creek}] + [6\text{-}me_{woodland}] + [\mathbf{6\text{-}me_{in\ situ}}]} \quad (8)$$

Solving this equation results in a minimum amount of 7.4 μg g^{-1} C 6-methyl brGDGTs that needs to be additionally produced in the creek to reach the higher IR. This accounts for 65% of the total amount of 6-methyl brGDGTs in the creek bed sediment that we measured. Considering a mixture of all soils rather than only woodland as source for soil-derived brGDGTs in the creek results in the *in situ* production of 9.3 μg g^{-1} C 6-methyl brGDGTs, corresponding to 81% of the 6-methyl brGDGT pool in the creek bed sediments. This implies that the initial soil brGDGT signal is rapidly overprinted by a riverine *in situ* signal upon entering the creek. Only the IR for the downstream site in the northern creek approaches that of the adjacent soil (IR = 0.30 in the creek bed sediment and 0.38 ± 0.07 for Transect-7; Fig. A2), and may be explained by its use as arable land (Fig. 5a), which involves regular ploughing and subsequent soil mobilisation and implies a temporary, local overprint.

The absence of a clearly recognizable soil brGDGT signal in the creek bed sediments may be further explained by the relatively limited input of soil material into the creek. So far, river systems that have shown to transport a soil-derived brGDGT signal are either characterized by a distinct rainy season (e.g. the Congo River (Weijers et al., 2007b; Hemingway et al., 2017) or the Amazon River (Kim et al., 2012)), or have experienced a recent episode of extreme rainfall (e.g. the Danube River, >100 mm in 3 days causing a 100-year flood event, (Freymond et al., 2017) or the Rhône River, with heavy rainfall during sampling (Kim et al., 2015)). The Carminowe creek area does not have a clear rainy season, and is further characterized by its limited relief. Hence, the relatively minor input of soil-derived brGDGTs seems to be easily overprinted by riverine *in situ* production. Alternatively, the soil-derived brGDGTs could be preferentially degraded in an aquatic environment as a result of priming effect (Bianchi, 2011), which would lead to a signature that is dominated by brGDGTs that are produced *in situ*.

4.3 Sources of brGDGTs in the sediments of Lake Loe Pool

In theory, rivers would transport soil-derived OC together with any aquatic OC produced along the way. Once discharged, in this case into a lake, the OC would settle and then be buried into the sediments where it would act as a long-term sink of OC. However, the soil brGDGT signal cannot be recognized in the sediments from Loe Pool since it is already lost upon entering the Carminowe creek. Indeed, the PCA of the relative distributions of brGDGTs indicates that lake sediments plot completely separated from both the soils and creek sediments, mostly due to a higher relative abundance of GDGT–IIIa (Fig. 4a, b). As a result, the MBT'$_{5ME}$ is significantly lower in Loe Pool sediments (0.38 ± 0.00) compared to in the creek bed

sediments (0.46 ± 0.01; $p < 0.05$) and soils (0.48 ± 0.01; $p < 0.05$; Fig. 5b; Table 1). Furthermore, the DC is significantly higher in lake sediments than in both soil and creek bed sediments (0.32 ± 0.02, 0.23 ± 0.01 and 0.23 ± 0.01, respectively; $p < 0.05$; Fig. 3d; Table 1). The distinct brGDGT signature of the lake sediments suggests that brGDGTs in the lake again are significantly altered compared to those in the soils and creek sediments. This implies that the riverine brGDGT signal is either replaced or overwritten in the lake.

Lacustrine *in situ* production of brGDGTs has been reported in other studies (Sinninghe Damsté et al., 2009; Tierney and Russell, 2009; Buckles et al., 2014; Loomis et al., 2011, 2014a; Weber et al., 2015, 2018; Miller et al., 2018). However, there are no generally recognized indicators (yet) to identify lacustrine brGDGT production, although several studies reported a "cold bias" while attempting to reconstruct the mean air temperature (MAT) based on brGDGTs in lake sediments using a soil-based transfer function (Tierney et al., 2010). In a study on East African lakes, this cold bias was linked to a large *in situ* contribution of brGDGT–IIIa (Tierney et al., 2010), similar to in Loe Pool. However, the East African lake dataset was generated using the 'old' chromatography method that does not separate 5-methyl and 6-methyl brGDGTs. A recent study that has re-analysed the East African Lake dataset indicates that the presumed contribution of GDGT–IIIa mainly consists of brGDGT–IIIa' (Russell et al., 2018), which is less prominent in lake Loe Pool. Although the identity of brGDGT-producer(s) in lakes still remain(s) elusive, a recent study from the stratified Lake Lugano (Switzerland) showed that the majority of the brGDGTs are produced in the lower, anoxic part of the water column rather than in the sediment (Weber et al., 2018). Furthermore, the combination of brGDGT analysis with molecular biological methods revealed that brGDGTs appeared to be produced by multiple groups of bacteria thriving under different redox regimes in this stratified lake. Specifically, brGDGT–IIIa occurred in the entire water column and continuously increased with depth, whereas brGDGT–IIIa' was mainly produced in the upper, oxygenated part of water column (Weber et al., 2018). Extrapolating the ecological niches of brGDGT production in Lake Lugano to Loe Pool we can speculate that brGDGT-IIIa, which is dominating the brGDGT signal in the Loe Pool sediments, is mostly produced in the lake during summer, when the eutrophic state of the lake may seasonally cause the anoxic conditions favourable for its (i.e. brGDGT-IIIa) production. However, our dataset does not allow to further pinpoint the time and depth of lacustrine brGDGT production, or whether brGDGTs are solely produced in the water column of Loe Pool or also in the lake sediment.

4.4 Reconstructing local environmental changes based on GDGTs in Loe Pool lake sediments

Downcore variations in the brGDGT distribution of Lake Loe Pool sediments may provide information on past environmental changes in the catchment, in spite of the lacustrine *in situ* production in Lake Loe Pool. The 50 cm deep sediment core covers about the last 100 years based on ^{137}Cs activity (Glendell et al., 2018). The peak activity correlated with bomb testing in the 1960s was detected at 26 cm depth (Fig. 6a), which can thus be linked to 1963 (Glendell et al., 2018).

The C-normalized concentration of brGDGTs starts to increase around 23 cm, reaching a maximum concentration of 48.0 μg g^{-1} C at 11 cm depth (Fig. 6b). The increased brGDGT concentrations coincide with an increase in the degree of cyclisation (Fig. 6c), which generally responds to a change in pH, where more cyclopentane moieties correspond to a higher pH (Weijers et al., 2007a; Schoon et al., 2013). According to historical records, agriculture and anthropogenic perturbations such as mining and urban pollution intensified in the 1960s (~ 26 cm depth), which increased the input of soil and nutrients into Lake Loe Pool (Coard et al., 1983), and resulted in eutrophication (i.e. blooms of cyanobacteria and algae) since at least 1986 (~ 23 cm depth) (O'Sullivan, 1992; Flory and Hawley, 1994). Earlier studies have also recognized an increased use of farmyard manures and septic tanks at this time in the nitrogen isotopic composition of the lake sediments, and have detected higher inputs of terrestrial organic material resulting from intensified farming practices and a higher erosion rate during the 1960s to 1980s based on ratios of aquatic- and terrestrial-derived plant waxes (Glendell et al., 2018). Thus, the high brGDGT

concentrations and DC in the sediments likely reflect the eutrophic conditions of the lake resulting from the increased nutrient input to the lake (Coard et al., 1983). The DC has then recorded the increase in lake water pH associated with eutrophication, whereas brGDGT concentrations express increased aquatic production. Due to remediation measures taken by the local government in 1996 (~ 12 cm depth), the eutrophication has reduced over the past twenty years (Glendell et al., 2018). The partial recovery of the lake has likely resulted in a return to lower lake water pH, as manifested in the decrease in the DC from ~ 10 cm depth upwards (Fig. 6c).

The process of eutrophication and subsequent recovery can also be recognized in the ratio between GDGT–0 and crenarchaeol, which are isoprenoidal GDGTs produced by Archaea. Crenarchaeol is produced by ammonia oxidizing Thaumarchaeota (Sinninghe Damsté et al., 2002) in aquatic environments (Schouten et al., 2000; Powers et al., 2004) and to a lesser extent also in soils (Weijers et al., 2006a), whereas GDGT–0 is a membrane lipid that occurs in all major groups of Archaea, but is indicative of methanogens and thus anaerobic conditions, with a typical ratio of GDGT-0 and crenarchaeol > 2 (Blaga et al., 2009). The ratio of GDGT-0/crenarchaeol in the sediments of Loe Pool is > 2 throughout the entire core, and ranges between 10.9 and 24.3, indicating that at least the bottom waters of the lake have been (seasonally) anoxic over the past 100 years (Fig. 6d), although the isoGDGTs may potentially be produced in deeper sediments. The ratio reaches its maximum at 16 cm depth, suggesting that eutrophic conditions and bottom water anoxia were most severe around this time. The recovery of the lake after the remediation measures is again reflected in the return to pre-1960 values at ~ 10 cm depth (Fig. 6d).

5 Conclusions

In this study, brGDGTs were tested as a tracer for the transport of soil OC from different vegetation and land use types from source (soil) to sink (lake Loe Pool) in the Carminowe Creek catchment with the aim to reconstruct the provenance of the soil OC in lake Loe Pool sediments over time. Unfortunately, brGDGT signatures in the catchment soils are not distinct for land use types, indicating that other environmental parameters have a larger influence on the distribution of brGDGTs in these soils. Although temperature and precipitation can be considered equal for all soils due to the small size of the catchment, changes in BIT index values and the relative contribution of 6-methyl brGDGTs along a part of the hilltop transects indicate that soil water content (SWC) may exert a control on brGDGT signals, assuming that SWC increases downslope. The regular rotation of cropland in this catchment and the relative long turnover time of brGDGTs in soils could be another reason to explain the limited spatial variation in brGDGT signals.

Comparison of the soil-derived brGDGT signals to that of creek bed sediments reveals that the soil brGDGT signal is almost completely overprinted by aquatically produced brGDGTs, indicated by a substantially higher fractional abundance of 6-methyl brGDGTs in the creek. Upon discharge into the lake, the creek brGDGT signal is replaced by and/or mixed with a lacustrine *in situ* produced brGDGT signal, which is characterized by a relatively higher DC and lower MBT'_{5ME}, as well as a specifically high fractional abundance brGDGT-IIIa. Despite regular ploughing of the land, the absence of a profound rainy season and limited relief likely limits the degree of soil mobilisation necessary to transfer the soil-derived brGDGT signal to the lake sediments in the modern system. Still, downcore variations in GDGT distributions in the sediments of Loe Pool do reflect local environmental conditions over the past 100 years. The degree of cyclisation of brGDGTs as well as the ratio of isoprenoidal GDGT-0 and crenarchaeol produced by Archaea trace the historical record of lake eutrophication induced by increased nutrient input from intensified agricultural activity in the catchment during the 1960s to 1980s, and its recovery after measures taken by the owner since 1996. Our study shows that GDGTs in sedimentary archives are good recorders of past environmental and land management (e.g. agricultural intensification, increased fertilizer use) change, although the ability of brGDGTs to trace soil OC along a soil-aquatic continuum requires a higher degree of soil mobilisation.

Data availability

All data are available in the Supplementary Information.

Author Contribution

J.M., F.K., and F.P designed the study, M.G. and J.M. collected the sample material. J.G. conducted the biomarker analysis and interpreted the data under supervision of F.P. and J.J.M, J.G. and F.P wrote the paper with input from all co-authors.

Competing interests

The authors declare that they have no conflict of interest.

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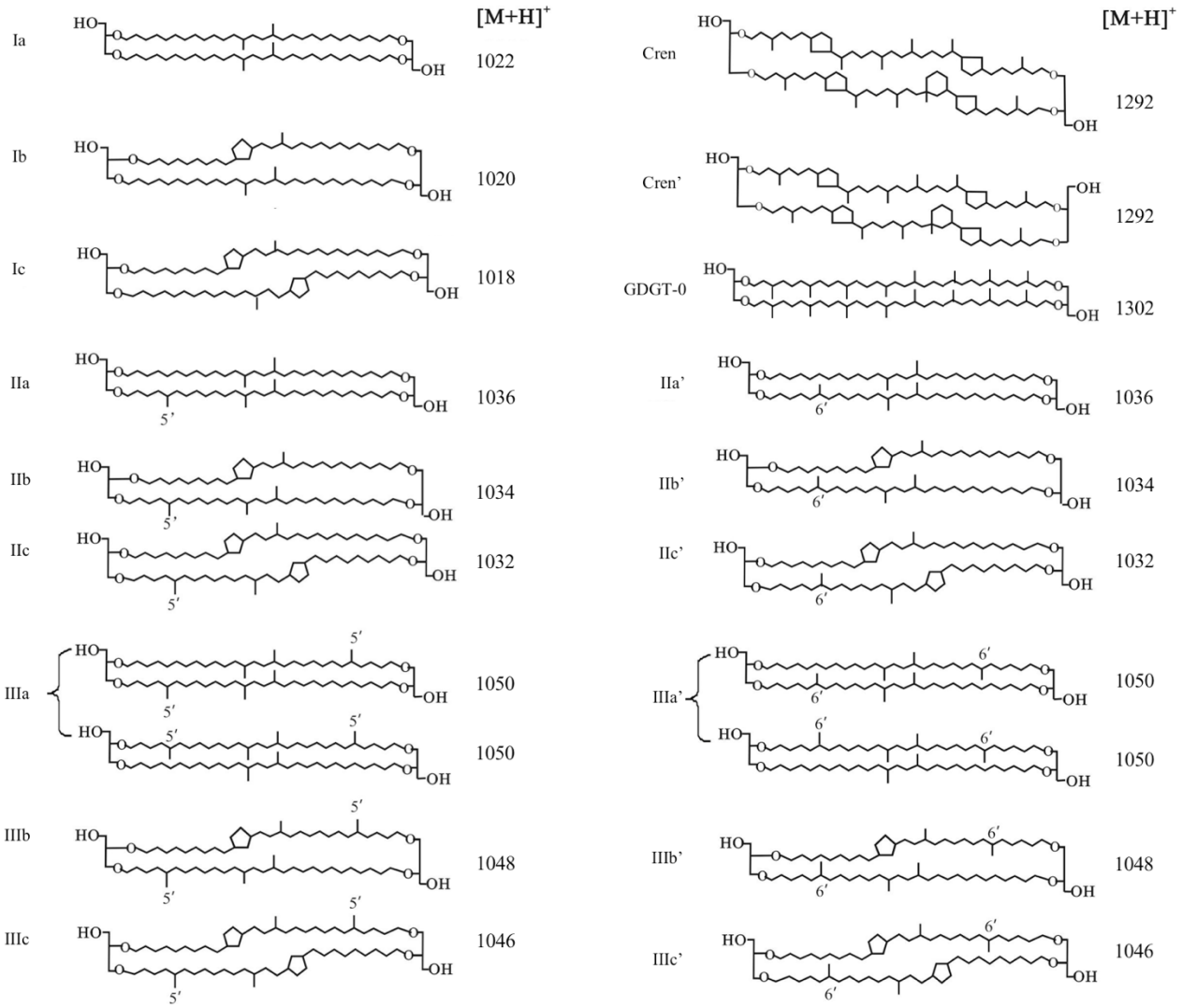
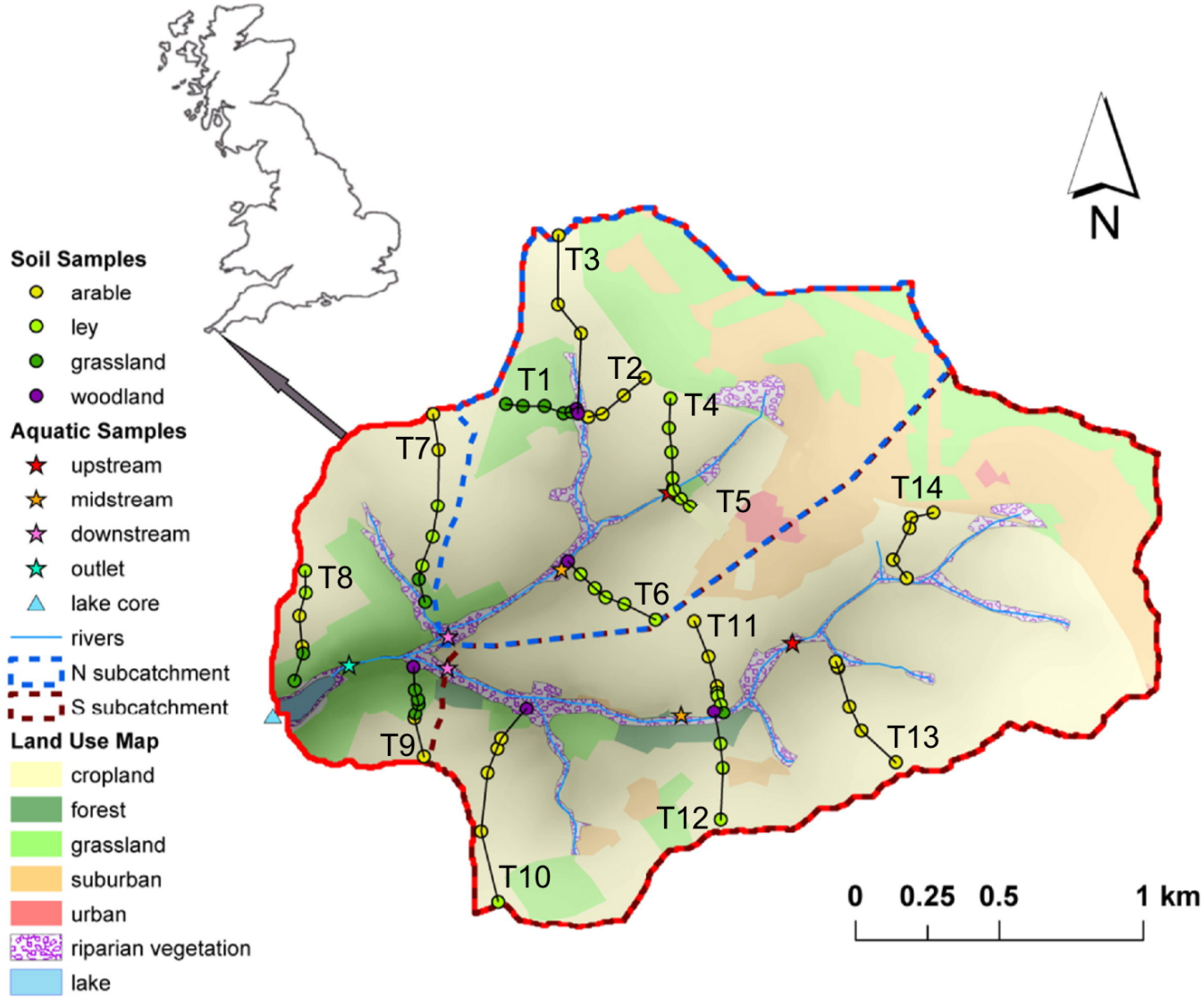


682

683 **Figure 1: Molecular structures of 5-methyl and 6-methyl branched GDGTs, GDGT-0 and crenarchaeol. The 6-methyl brGDGTs**
684 **are represented by apostrophe. The structures of penta- and hexamethylated brGDGTs with cyclopentane moiety(ies) IIb', IIc',**
685 **IIIb', IIIc' are tentative.**



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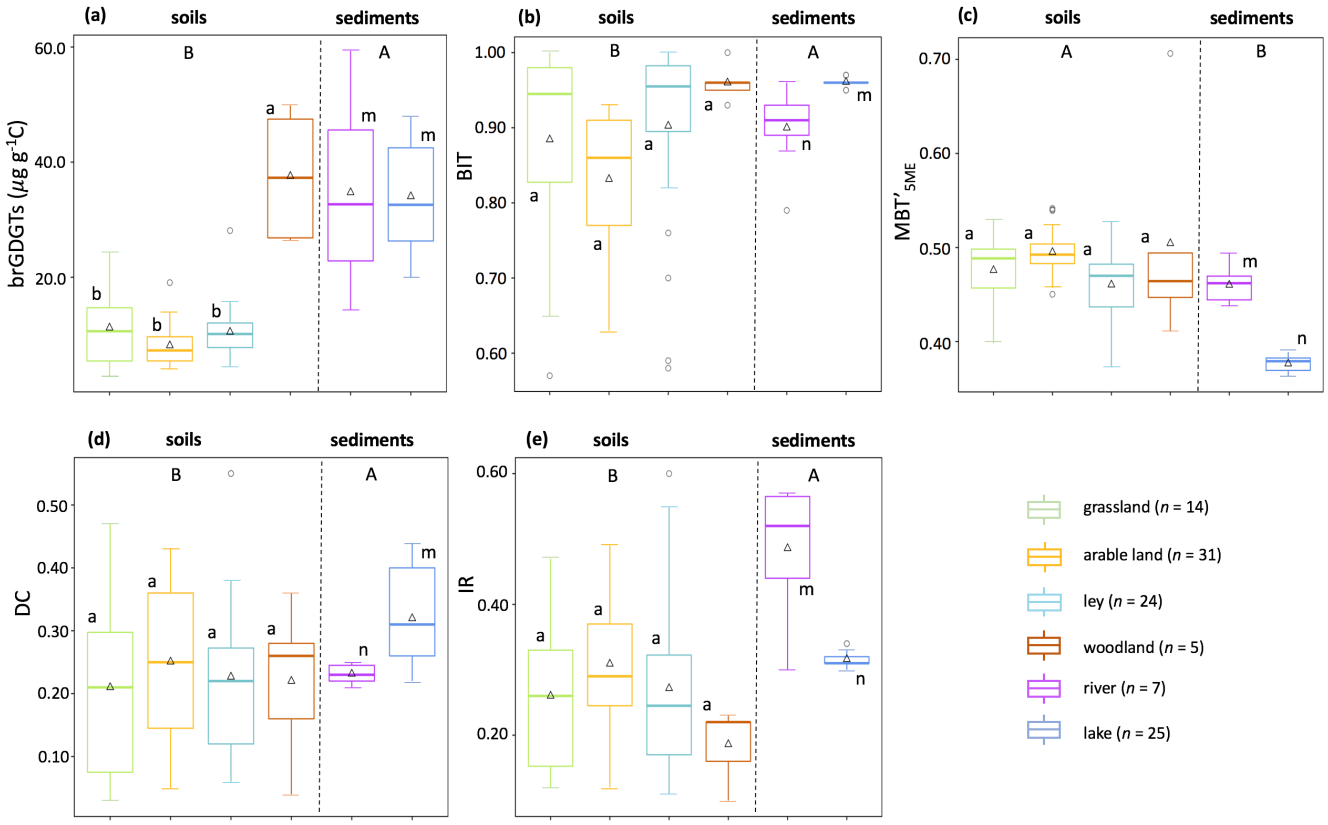
687 **Figure 2: Map of the Carminowe Creek catchment in southwest England showing land use types, 14 soil transects (labelled T1-14),**
688 **creek bed and lake core sediment sampling locations. The coloured circles and stars indicate soil samples under different land use**
689 **types and creek bed sediments along the streams, respectively. Adjusted from Glendell et al. (2018).**



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691 **Figure 3: Box plots displaying (a) the C-normalized concentration of brGDGTs, and brGDGT-based proxies: (b) BIT index**
692 **(branched and isoprenoid tetraether ratio), (c) MBT'$_{5ME}$ (methylation of 5-methyl branched tetraethers), (d) DC (degree of**
693 **cyclisation) and (e) IR (isomerization ratio). The triangles represent the average values, the bold line indicates the median (50th**
694 **percentile), bottom and top of the box indicate first quartile (25th percentile) and third quartile (75th percentile) respectively,**
695 **whiskers cover the smallest and largest value within 1.5 times of the interquartile range (i.e. the distance between the top and**
696 **bottom of the box). Any data points outside the whiskers are considered as outliers. Different letters indicate differences between**
697 **samples: A and B for differences between catchment soils and aquatic sediments, a and b for soils under different vegetation types,**
698 **and m and n for creek bed and lake core sediments ($p < 0.05$).**

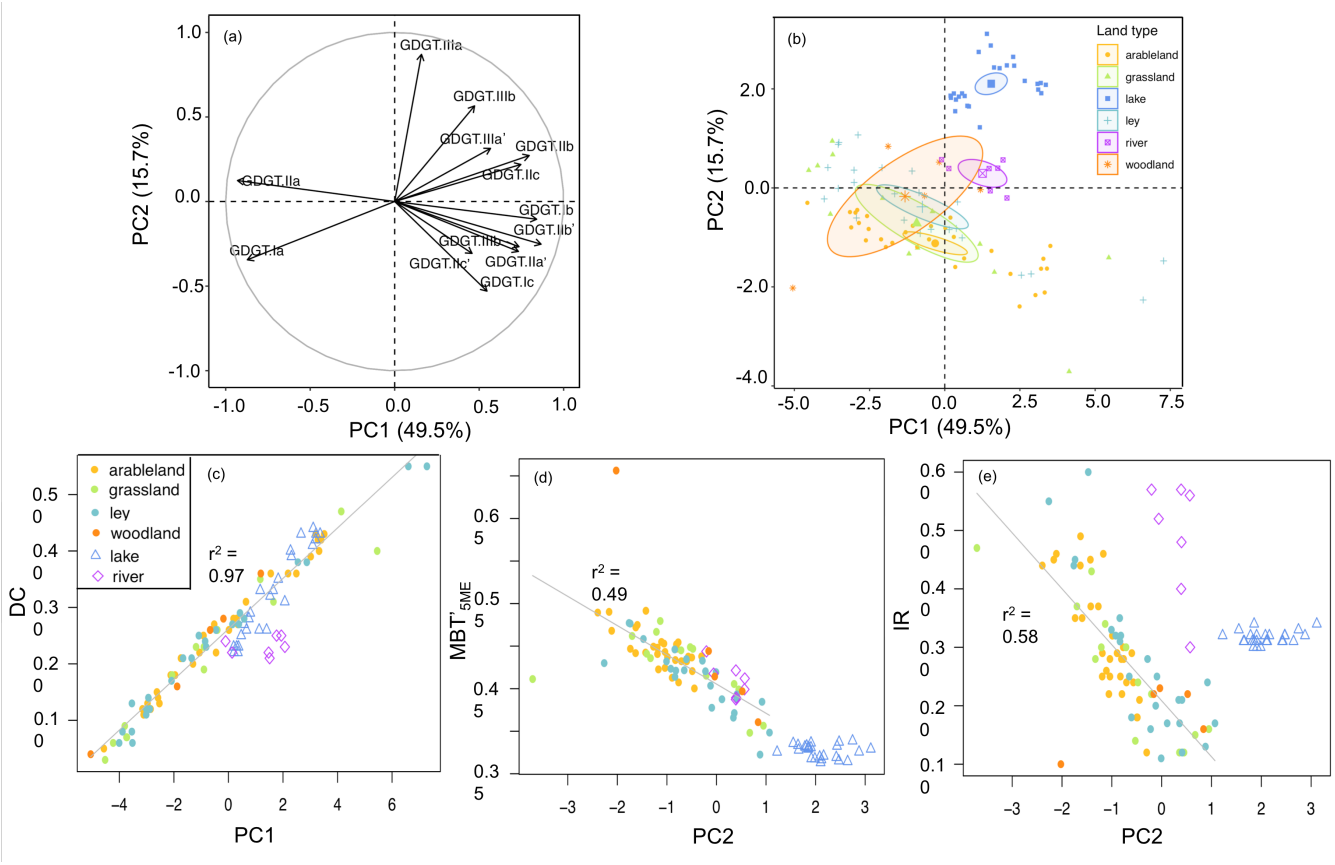


Figure 4: PCA based on the relative abundances of 13 major brGDGTs. Figure (a) shows the distribution of 13 brGDGTs (brGDGT-IIIc and brGDGT-IIIc' are excluded as they are below the detection limit) along the first two PCs, roman numerals and English alphabet represent the compounds shown in Fig. 1. Figure (b) shows sampling sites loading scores on the first two PCs and 95% confidence interval ellipses surrounding the mean point of different groups of land use: arable land (*n* = 31), grassland (*n* = 14), ley (*n* = 24) and woodland (*n* = 5), and creek (*n* = 7) and lake (*n* = 25). Figure (c) shows cross plots between PC1 and DC (degree of cyclisation). Figure (d) and (e) show cross plots of PC2 with MBT'_{5ME} (methylation of 5-methyl branched tetraethers) and IR (isomerization ratio) respectively. The linear correlation was calculated excluding creek and lake sediment.

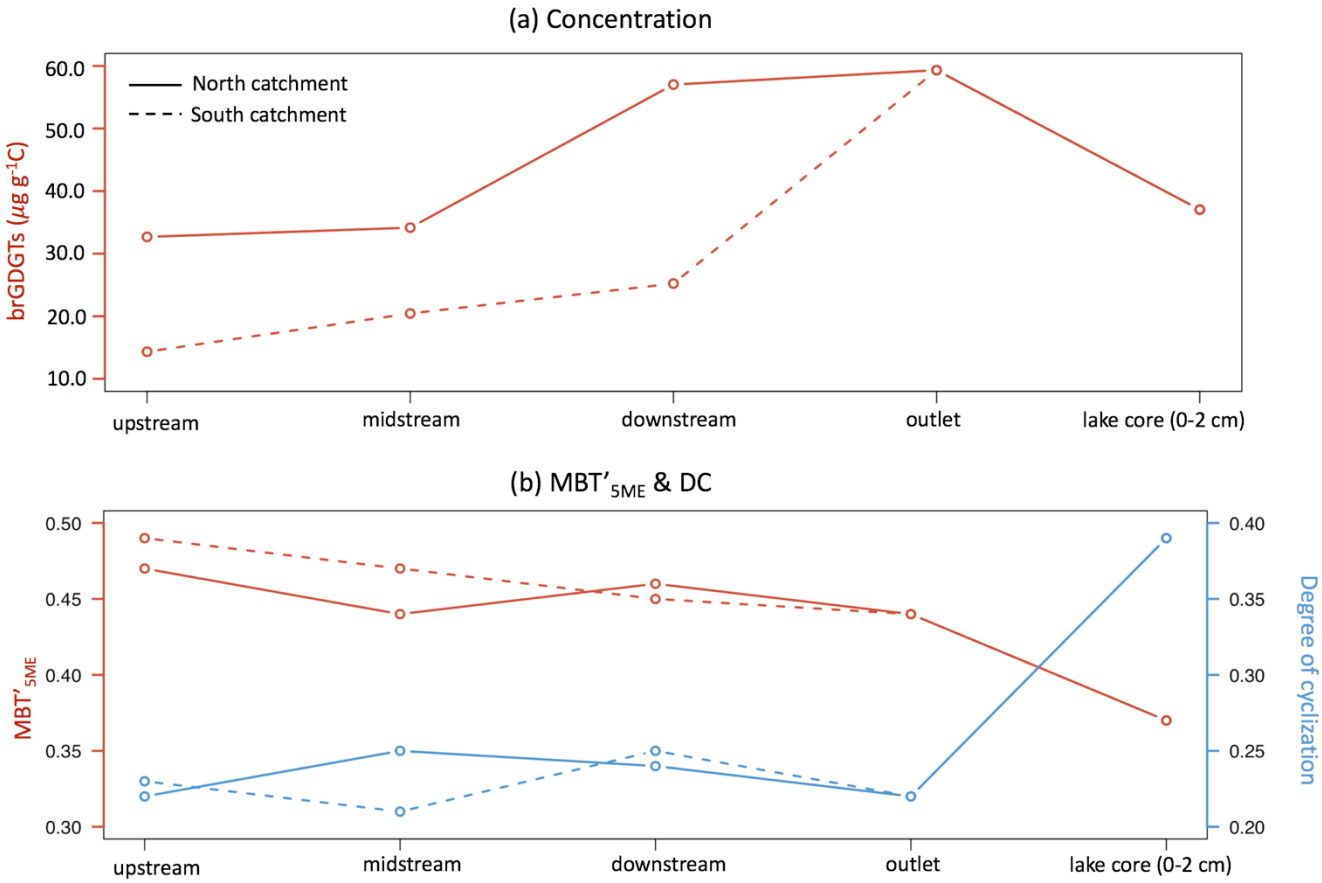
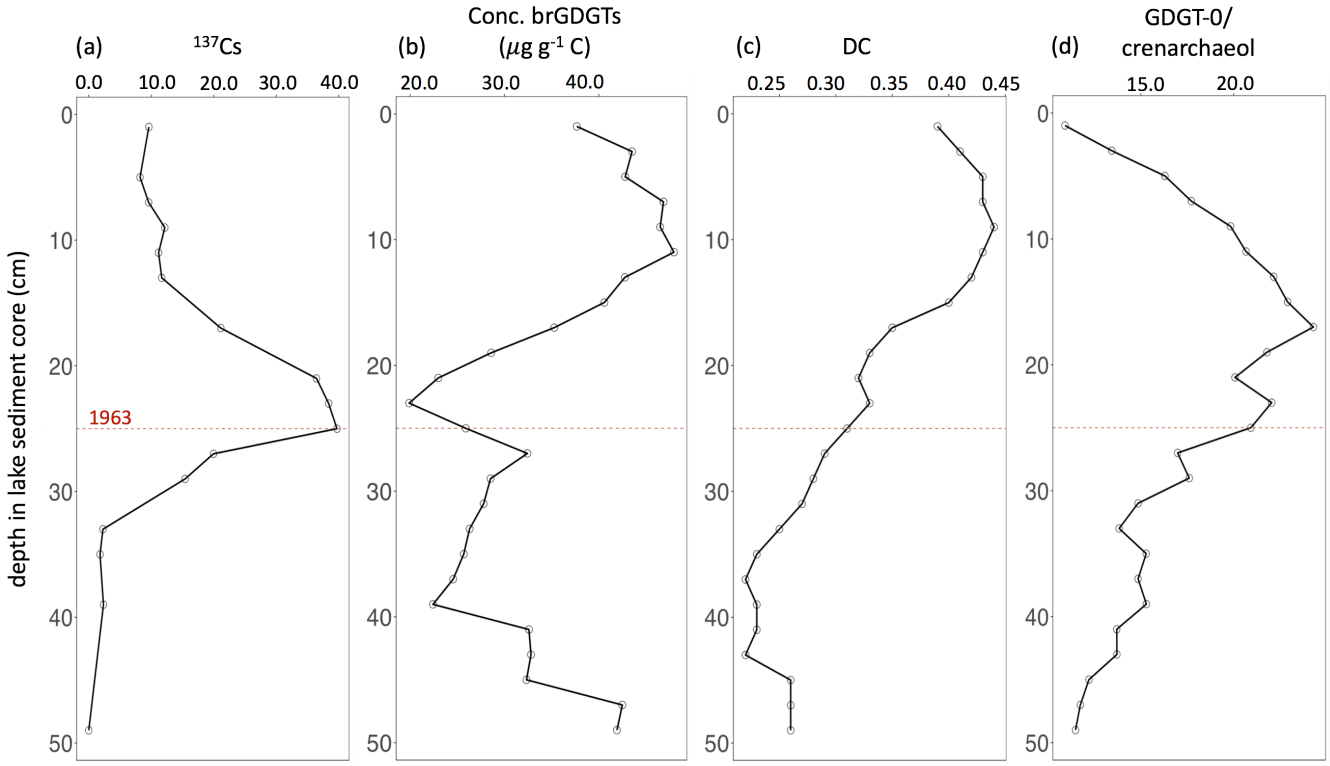


Figure 5: Spatial variability of (a) C-normalized concentration of brGDGTs and (b) MBT'$_{5ME}$ (methylation of 5-methyl branched tetraethers) and DC (degree of cyclisation) in downstream direction of both substreams in the Carminowe Creek catchment.



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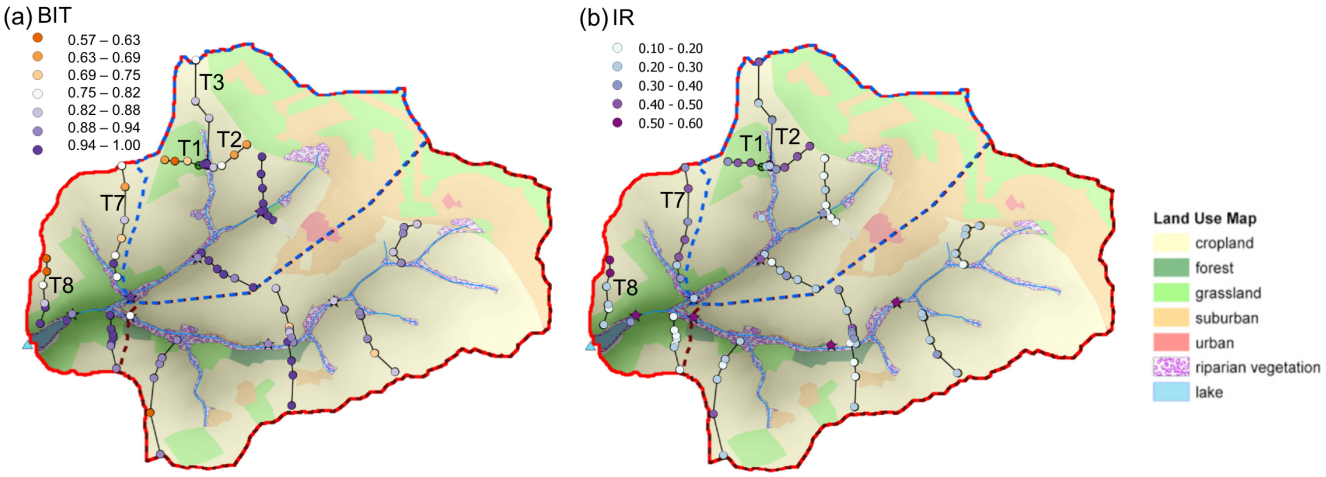
711 **Figure 6: Lake sediment core profiles of (a) ^{137}Cs to date, (b) C-normalized concentration of brGDGTs, (c) DC (degree of**
712 **cyclisation) and (d) ratio between GDGT-0 and crenarchaeol. The red dashed line indicates the year of 1963.**



Appendix Fig.1: Spatial variability of the (a) BIT (branched and isoprenoid tetraether ratio) and (b) IR (isomerization ratio) along 14 soil transects in the Carminowe Creek catchment. The coloured circles show the concentrations and proxy values. Tx indicates soil transects discussed in the text. The background colours indicate different land use types. Adjusted from Glendell et al. (2018).

Table 1. **C% (carbon content), pH values, average concentrations of brGDGTs and brGDGT-based proxies under different land use types. BIT (branched and isoprenoid tetraether ratio), MBT'_{5ME} (methylation of 5-methyl branched tetraethers), %tetra (percentage of tetramethylated brGDGTs), %penta (percentage of pentamethylated brGDGTs), %hexa (percentage of hexamethylated brGDGTs), DC (degree of cyclisation), IR (isomerization ratio) (mean ± standard deviation, s.d.).**

Land use (n)	C% *	pH	Conc. ($\mu g\ g^{-1}$ soil)	Conc. ($\mu g\ g^{-1}$ C)	BIT	MBT'_{5ME}	%tetra	%penta	%hexa	DC	IR
arable (31)	2.9 ± 0.5	6.6 ± 0.4	0.2 ± 0.1	8.1 ± 3.6	0.83 ± 0.09	0.50 ± 0.02	40.1 ± 3.1	49.7 ± 1.5	10.2 ± 1.8	0.25 ± 0.11	0.31 ± 0.10
grass (14)	5.6 ± 1.2	6.0 ± 0.5	0.6 ± 0.4	11.2 ± 6.7	0.88 ± 0.14	0.48 ± 0.04	39.8 ± 4.0	49.4 ± 2.3	10.8 ± 2.1	0.21 ± 0.14	0.26 ± 0.12
ley (24)	3.6 ± 0.9	6.0 ± 0.3	0.4 ± 0.3	10.5 ± 4.8	0.90 ± 0.12	0.46 ± 0.04	37.8 ± 3.7	50.2 ± 2.0	12.0 ± 2.9	0.23 ± 0.14	0.27 ± 0.13
woodland (5)	8.2 ± 2.1	5.4 ± 0.7	3.0 ± 1.0	37.6 ± 11.0	0.96 ± 0.03	0.50 ± 0.12	45.4 ± 13.0	44.4 ± 8.3	10.3 ± 4.8	0.22 ± 0.12	0.19 ± 0.05
all soils (74)	4.0 ± 1.8	6.2 ± 0.5	0.6 ± 0.8	11.5 ± 8.9	0.87 ± 0.12	0.48 ± 0.04	39.7 ± 4.9	49.4 ± 3.0	10.9 ± 2.6	0.23 ± 0.13	0.28 ± 0.11
creek (7)	2.3 ± 0.8	7.1 ± 0.2	0.8 ± 0.4	34.7 ± 17.4	0.90 ± 0.06	0.46 ± 0.02	30.1 ± 4.5	45.0 ± 0.7	24.9 ± 4.7	0.23 ± 0.02	0.48 ± 0.10
Lake (25)	7.5 ± 1.0	5.7 ± 0.2	2.6 ± 0.7	34.0 ± 8.7	0.96 ± 0.01	0.38 ± 0.01	28.9 ± 0.7	50.2 ± 1.8	21.0 ± 1.4	0.32 ± 0.08	0.32 ± 0.01

717 *From Glendell et al. (2018)

Appendix Table 1. BIT values along 14 transects (Tx indicates the transect number, and Sx indicates the sample point, where 1 represents the hilltop and subsequent numbers are further downslope).

		North catchment								South catchment					
	BIT	**T1**	*T2*	*T3*	T4	T5	T6	*T7*	**T8**	T9	T10	T11	T12	T13	T14
hilltop	S1	**0.65**	*0.66*	*0.77*	0.97	0.99	0.97	*0.77*	**0.58**	0.92	0.92	0.84	0.95	0.84	0.86
	S2	**0.57**	*0.66*	*0.86*	0.99	0.99	0.93	*0.65*	**0.59**	0.91	0.63	-	-	0.73	0.88
	S3	**0.73**	*0.77*	*0.87*	1.00	1.00	0.94	*0.82*	**0.80**	0.98	0.90	0.92	0.98	0.91	0.88
	S4	**0.88**	*0.83*	*0.96*	0.99	-	0.96	*0.70*	**0.85**	1.00	0.92	0.72	0.98	0.92	0.86
	S5	**0.95**	-	-	-	-	0.97	*0.76*	**0.98**	1.00	0.91	0.91	1.00	0.91	0.93
	S6	-	-	-	-	-	0.96	*0.94*	**0.97**	1.00	0.93	0.85	-	0.90	-
	S7	-	-	-	-	-	-	*0.81*	-	0.95	-	0.92	-	-	-
downslope	S8	-	-	-	-	-	-	-	-	-	-	0.92	-	-	-

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