



1 Assessing branched tetraether lipids as tracers of soil organic carbon 2 transport through the Carminowe Creek catchment (southwest 3 England)

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11 **Abstract.** Soils represent the largest reservoir of organic carbon (OC) on land. Upon mobilization, this OC is either returned
12 to the atmosphere as carbon dioxide (CO₂), or transported and ultimately locked into (marine) sediments, where it will act as
13 a long-term sink of atmospheric CO₂. These fluxes of soil OC are, however, poorly quantified, mostly due to the lack of a
14 soil-specific tracer. In this study, a suite of branched glycerol dialkyl glycerol tetraethers (brGDGTs), which are membrane
15 lipids of soil bacteria, is tested as specific tracers for soil OC from source (soils under arable land, ley, grassland and
16 woodland) to sink (Lake Loe Pool sediments) considering a small catchment located in southwest England (i.e. Carminowe
17 Creek draining into Lake Loe Pool). The analysis of brGDGTs in catchment soils reveals that their distribution is not
18 significantly different across different land use types ($p > 0.05$), and thus does not allow tracing land use-specific soil
19 contributions to Lake Loe Pool sediments. Furthermore, the significantly higher contribution of 6-methyl brGDGT isomers
20 in creek sediments (isomerization ratio (IR) = 0.48 ± 0.10 ; mean \pm s.d., standard deviation; $p < 0.05$) compared to that in
21 catchment soils (IR = 0.28 ± 0.11) indicates that the initial soil signal is substantially altered by brGDGT produced *in situ*.
22 Similarly, the riverine brGDGT signal appears to be overwritten by lacustrine brGDGTs in the lake sedimentary record,
23 indicated by remarkably lower Methylation of Branched Tetraethers (MBT'_{SME} = 0.46 ± 0.02 in creek bed sediment and 0.38
24 ± 0.01 in lake core sediment; $p < 0.05$) and higher Degree of Cyclization (DC = 0.23 ± 0.02 in creek bed sediment and $0.32 \pm$
25 0.08 in lake core sediment). Thus, in this small catchment, brGDGTs do not allow us to trace soil OC transport.
26 Nevertheless, the downcore changes in the degree of cyclization and the abundance of isoprenoid GDGTs produced by
27 methanogens in the Lake Loe Pool sediment do reflect local environmental conditions over the past 100 years, and have
28 recorded the eutrophication history of the lake.



29 1 Introduction

30 Globally, around 1500–2000 Pg of carbon is stored in soils in the form of organic matter, which is about two times the
31 amount of carbon in the atmosphere and three times the amount of carbon in vegetation (Janzen, 2004; Smith, 2008). Soil
32 organic carbon (OC) plays an important role in the global carbon cycle, as subtle alterations in the soil OC reservoir may
33 affect the concentration of atmospheric CO₂ and thus influence climate change (Davidson and Janssens, 2006). Atmospheric
34 CO₂ that is fixed by plants through photosynthesis will be stored into soil OC pool, part of which will be transferred to
35 streams and rivers. Upon fluvial discharge, soil OC is buried and locked into the marine or lacustrine sediment, where it will
36 act as a long-term carbon sink. However, instead of a passive pipeline in the carbon cycle, rivers actually represent a
37 dynamic channel, where part of the soil OC is respired back to the atmosphere, and another part may be stored in river bed or
38 lake sediments before reaching the ocean (Cole et al., 2007; Battin et al., 2009; Aufdenkampe et al., 2011). Hence, it is hard
39 to determine the exact amount of soil OC that is transported to the ocean, as the dynamic processes that soil OC undergoes
40 during transport, such as degradation and sequestration, are elusive. This is mostly due to the lack of a specific tracer to
41 distinguish soil OC from the total pool of OC that is also comprised of plant-derived OC, aquatic produced OC, and fossil
42 OC from rock erosion (Blair et al., 2004; Aufdenkampe et al., 2011).

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

52 Branched glycerol dialkyl glycerol tetraethers (brGDGTs; Fig. 1) are membrane spanning tetraether lipids synthesized by
53 heterotrophic bacteria that thrive in soils and peats all over the world (Weijers et al., 2006a, 2007a; Naafs et al., 2017a).
54 Although the exact producers of these lipids are still unknown, after the detection of a brGDGT in an Acidobacterial culture
55 (Sinninghe Damsté et al., 2011, 2014, 2018), it was assumed that Acidobacteria are the source organisms of brGDGTs. The
56 occurrence and relative distribution of brGDGTs in a global set of modern surface soils showed that they can have 4 to 6
57 methyl groups attached to their alkyl backbone, where the degree of branching increases in soils from colder areas.
58 Furthermore, brGDGTs respond to changes in soil pH by forming up to 2 cyclopentane moieties following internal
59 cyclization, where a higher number of cyclopentane moieties corresponds to a higher soil pH (Weijers et al., 2007a).
60 Initially, a combination of two indices, the Methylation of Branched Tetraethers (MBT) index and Cyclization of Branched



61 Tetraethers (CBT) index, was proposed as a proxy to reconstruct the mean air temperature (MAT) and pH of a soil (Weijers
62 et al., 2007a; Peterse et al., 2012). After the identification of novel brGDGT isomers that possess a methyl group at the α
63 and/or ω 6 position rather than at position 5 (Fig. 1) and the improvement of the chromatography method used for brGDGT
64 analysis, a modified temperature proxy, the MBT_{5ME} was developed (De Jonge et al., 2013, 2014b). Furthermore, the
65 relative abundance of 6-methyl brGDGT isomers, quantified as the Isomerization Ratio (IR), appeared to also relate to soil
66 pH (De Jonge et al., 2014a). Indeed, the analysis of brGDGTs in peat profiles and loess-paleosol sequences has resulted in
67 long-term continental paleotemperature records for various areas, e.g. in deglacial central China (Peterse et al., 2011) and
68 northeast China (Zheng et al., 2017), and western Europe during the early Eocene (Inglis et al., 2017).

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

77 Recently, however, brGDGTs have also been found to be produced in aquatic systems such as coastal marine areas (Peterse
78 et al., 2009b; Sinninghe Damsté, 2016), rivers (Zell et al., 2013, 2014a) and lakes (Sinninghe Damsté et al., 2009a; Tierney
79 and Russell, 2009; Loomis et al., 2011, 2014; Schoon et al., 2013; Weber et al., 2015, 2018), which complicates the
80 interpretation of brGDGT-based proxies. Aquatic production became apparent upon comparison of brGDGTs in Svalbard
81 fjord sediments and nearby soils. Whereas the brGDGT signal in the fjord sediments was dominated by compounds
82 containing cyclopentane moieties, soils were characterized by brGDGTs without cyclization (Peterse et al., 2009b). These
83 substantially different brGDGT signatures in combination with the increasing concentration of brGDGTs towards the open
84 ocean then pointed towards a contribution of *in situ* produced brGDGTs to the fjord sediments. Similarly, De Jonge et al.
85 (2014a) found that brGDGTs in suspended particulate matter (SPM) from the Yenisei River matched better with the pH of
86 the river water than that of the soils in the river catchment. Also, the distribution of brGDGTs in SPM remained constant,
87 whereas the Yenisei catchment spans a large latitudinal range with a large range in temperature, which led them to conclude
88 that the majority of the brGDGTs in SPM are produced in the river. Notably, these and subsequent studies proposed ways to
89 recognize *in situ* production of brGDGTs in aquatic environments. For example, the aquatic brGDGTs in the Yenisei River
90 were characterized by a relatively high contribution of 6-methyl brGDGT isomers (De Jonge et al., 2014a), whereas a high
91 degree of cyclization is an indicator of brGDGT production in coastal marine zones (Peterse et al., 2009b; Sinninghe



92 Damsté, 2016), for which Sinninghe Damsté (2016) proposed that a weighed number of rings in tetramethylated brGDGTs,
93 quantified as $\#rings_{tetra} > 0.7$ indicates a purely marine source of brGDGTs in continental margin sediments.

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

104 2 Methods

105 2.1 Study site and sampling

106 An overview of the study area and sampling sites is given by Glendell et al. (2018). Briefly, the Carminowe Creek catchment
107 is located in Cornwall in southwest England (50°14' N, 5°16' W), covers an area of around 4.8 km² and varies in elevation
108 from 0 to 80 m above sea level (Fig. 2). It is divided into two subcatchments ('north' and 'south'). The two streams converge
109 around 100 m before their joint outlet, and then flow into a natural freshwater lake Loe Pool (50 ha), which is separated from
110 the Atlantic Ocean by a natural shingle barrier. The mean annual temperature (MAT) and mean annual precipitation (MAP)
111 in this area are approximately 11 °C and 1000 mm year⁻¹, respectively. The land use in this studied catchment is dominated
112 by arable land and temporary grasslands (ley), which are under rotation. The steeper hillslopes are under permanent
113 grassland, and riparian woodland covers the areas near the creek. For this study, 74 surface soil samples (0–15 cm) were
114 collected along 14 hillslope transects, including 31 arable land sites, 14 permanent grassland sites, 24 temporary grassland
115 (ley) sites and 5 woodland sites (Fig. 2). Riverbed sediments were collected at three locations along each of the two
116 tributaries (upstream, midstream and downstream), and one more at the joint outlet. A 50 cm long sediment core was taken
117 in the lake, about 150 m away from the joint outlet. The lake core has been dated by the activity of Caesium-137 (¹³⁷Cs), and
118 it covers the last 100 years (Glendell et al., 2018).

119 2.2 Bulk soil properties

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



122 2.3 GDGT extraction and analysis

123 In total, 74 soil samples, 7 creek bed sediment and 25 lake core sediment samples were analysed for GDGTs. First, 5–7 g of
124 the soils or 3–5 g of the sediments were freeze dried and homogenized, after which they were extracted three times with
125 dichloromethane (DCM) : MeOH (9 : 1, v/v) using an accelerated solvent extractor (ASE 350, Dionex™) at 100 °C and 7.7
126 × 10⁶ Pa to obtain a total lipid extract (TLE). After addition of a known amount of C₄₆ GDGT internal standard (Huguet et
127 al., 2006), the TLEs were dried under a N₂ stream, and then separated into apolar and polar fractions by passing them over an
128 activated Al₂O₃ column using hexane : DCM (9 : 1, v/v) and DCM : MeOH (1 : 1, v/v) respectively. The polar fraction,
129 which contains the GDGTs, was evaporated to dryness under a gentle N₂ stream. After this, the samples were prepared for
130 further analysis by re-dissolving them in a hexane : isopropanol (99 : 1, v/v) mixture, and filtration through a 0.45 μm
131 polytetrafluoroethylene (PTFE) filter.

132 The GDGTs were analysed on an Agilent 1260 Infinity ultra high performance liquid chromatography (UHPLC) coupled to
133 an Agilent 6130 single quadrupole mass spectrometer (MS) with settings according to Hopmans et al. (2016). The GDGTs
134 were separated over two silica Waters Acquity UPLC BEH Hilic columns (1.7 μm, 2.1 mm x 150 mm) preceded by a guard
135 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
136 25 min, followed by a linear gradient to 70% A and 30% B for 25 min, where A = hexane and B = hexane : isopropanol (9 :
137 1, v/v). Sample injection volumes were 10 μL. Ionization of the GDGTs was achieved by atmospheric pressure chemical
138 ionization with the following source settings: gas temperature 200 °C, vaporizer temperature 400 °C, N₂ flow 6 L min⁻¹,
139 capillary voltage 3500 V, nebulizer pressure 25 psi and a corona current of 5.0 μA. By scanning the [M+H]⁺ ions (protonated
140 mass) in selected ion monitoring (SIM) mode, the target compounds were detected at *m/z* 1302 (GDGT-0), 1292
141 (crenarchaeol), 1050 (brGDGT-IIIa), 1048 (brGDGT-IIIb), 1046 (brGDGT-IIIc), 1036 (brGDGT-IIa), 1034 (brGDGT-
142 IIb), 1032 (brGDGT-IIc), 1022 (brGDGT-Ia), 1020 (brGDGT-Ib), 1018 (brGDGT-Ic), with *m/z* 744 for the internal
143 standard. Quantitation was achieved by peak area integration of the [M+H]⁺ ions in Chemstation software B.04.03.

144 2.4 GDGT proxy calculations

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

$$148 \text{ BIT} = \frac{[Ia]+[IIa]+[IIIa]+[IIa']+[IIIa']}{[Ia]+[IIa]+[IIIa]+[IIa']+[IIIa']+[crenarchaeol]} \quad (1)$$

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



$$151 \quad MBT'_{5Me} = \frac{[Ia] + [Ib] + [Ic]}{[Ia] + [Ib] + [Ic] + [IIa] + [IIb] + [IIc] + [IIIa]} \quad (2)$$

$$152 \quad \%tetra = \sum[tetramethylated \ brGDGTs] = [Ia] + [Ib] + [Ic] \quad (3)$$

$$153 \quad \%penta = \sum[pentamethylated \ brGDGTs] = [IIa] + [IIb] + [IIc] + [IIa'] + [IIb'] + [IIc'] \quad (4)$$

$$154 \quad \%hexa = \sum[hexamethylated \ brGDGTs] = [IIIa] + [IIIb] + [IIIc] + [IIIa'] + [IIIb'] + [IIIc'] \quad (5)$$

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

$$156 \quad DC = \frac{[Ib] + 2*[Ic] + [IIb] + [IIb']}{[Ia] + [Ib] + [Ic] + [IIa] + [IIa'] + [IIb] + [IIb']} \quad (6)$$

157 The isomerization ratio (IR) is the ratio between penta- and hexamethylated 6-methyl brGDGTs and the total amount of both
158 5- and 6-methyl penta- and hexamethylated brGDGTs (De Jonge et al., 2014a):

$$159 \quad IR = \frac{[IIa'] + [IIb'] + [IIc'] + [IIIa'] + [IIIb'] + [IIIc']}{[Ia] + [IIa'] + [IIb] + [IIb'] + [IIc] + [IIc'] + [IIIa] + [IIIa'] + [IIIb] + [IIIb'] + [IIIc] + [IIIc']} \quad (7)$$

160 2.5 Statistical analysis and data visualization

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

170 3 Results

171 3.1 BrGDGTs in soils

172 Most of the brGDGTs were present in all soils. Only brGDGT-IIIc and brGDGT-IIIc' were always below the detection limit
173 (peak height $> 3x$ baseline), and brGDGT-IIc' was below the detection limit in 13 of the soils (three in arable land, four in
174 grassland and six in ley). The brGDGTs were dominated by pentamethylated ($49.4 \pm 3.0\%$, mean \pm s.d., standard deviation),



175 followed by tetramethylated ($39.7 \pm 4.9\%$) and then hexamethylated brGDGTs ($10.9 \pm 2.6\%$; Table 1). The concentration of
176 brGDGTs ranged between 0.1 and $1.7 \mu\text{g g}^{-1}$ soil, with average of $0.2 \pm 0.1 \mu\text{g g}^{-1}$ soil in arable land, $0.6 \pm 0.4 \mu\text{g g}^{-1}$ soil in
177 grassland, and $0.4 \pm 0.3 \mu\text{g g}^{-1}$ soil in ley (i.e. the temporary grassland). However, the concentration of brGDGTs in
178 woodland was $3.0 \pm 1.0 \mu\text{g g}^{-1}$ soil, which was significantly higher than that in other land use types ($0.4 \pm 0.3 \mu\text{g g}^{-1}$ soil; $p <$
179 0.05 ; Fig. 3a). The C-normalized concentration of brGDGTs in catchment soils ranged between 2.8 to $49.8 \mu\text{g g}^{-1}$ C, $8.1 \pm$
180 $3.6 \mu\text{g g}^{-1}$ C in arable land, $11.2 \pm 6.7 \mu\text{g g}^{-1}$ C in grassland, $10.5 \pm 4.8 \mu\text{g g}^{-1}$ C in ley, and $37.6 \pm 11.0 \mu\text{g g}^{-1}$ C in woodland
181 (Fig. 3a; Table 1). The trend of the concentration of brGDGTs along the soil transects was not obvious.

182 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
183 woodland, 0.90 ± 0.12 in ley, 0.88 ± 0.14 in grassland and 0.83 ± 0.09 in arable land (without significant differences, $p >$
184 0.05). However, the BIT values increased from hillslope to downslope along several transects in north catchment, while the
185 BIT values show no clear trends in south catchment (Fig. A1). The MBT'_{SME} ranged from 0.37 to 0.71 and was mostly
186 similar between all land use types (0.48 ± 0.04 ; $p > 0.05$; Fig. 3c; Table 1). The degree of cyclization between land use types
187 was similar ($\text{DC} = 0.23 \pm 0.13$; Fig. 3d; Table 1; $p > 0.05$), likewise, the IR ranged from 0.10 to 0.60 (0.28 ± 0.01 on
188 average; Fig. 3e; Table 1; $p > 0.05$), without clear trend along the soil transects. However, four transects in the north
189 catchment have on average significantly higher IR values (> 0.36) than the other transects in the catchment (0.24 ± 0.09 ; $p <$
190 0.05 ; Fig. A1). In general, the IR increases with increasing soil pH in the catchment ($r^2 = 0.36$, $p < 0.001$).

191 3.2 BrGDGTs in creek bed sediments

192 All brGDGT compounds were detected in creek bed sediments, except for in the upstream site from north catchment, where
193 brGDGT-IIIc' was below detection limit. The brGDGTs in creek bed sediments were dominated by pentamethylated
194 brGDGTs ($45.0 \pm 0.7\%$), followed by tetramethylated brGDGTs ($30.1 \pm 4.5\%$), and hexamethylated brGDGTs ($24.9 \pm$
195 4.7%) (Table 1). The C-normalized concentration of brGDGTs in creek bed sediments was $34.7 \pm 17.4 \mu\text{g g}^{-1}$ C on average
196 (Fig. 3a; Table 1), where the concentration increased from $32.7 \mu\text{g g}^{-1}$ C to $57.0 \mu\text{g g}^{-1}$ C downstream in north catchment,
197 and from $14.3 \mu\text{g g}^{-1}$ C to $25.2 \mu\text{g g}^{-1}$ C downstream in south catchment, reaching a maximum value of $59.3 \mu\text{g g}^{-1}$ C at the
198 outlet (Fig. 5a). The concentration of brGDGTs in creek bed sediments was higher than that in soils under any land use types
199 except for woodland ($9.6 \pm 4.9 \mu\text{g g}^{-1}$ C; Fig. 3a; Table 1).

200 The BIT values for creek sediments were on average 0.90 ± 0.06 (Fig. 3b; Table 1). The MBT'_{SME} was relatively constant
201 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
202 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).
203 The brGDGT-based indices in creek bed sediments were similar with the indices in soils, except for IR, which was higher
204 than that in soils under any land use types (0.28 ± 0.11 ; Fig. 3; Table 1).



205 3.3 BrGDGTs in Lake Loe Pool sediment core

206 All brGDGTs were detected in the lake sediment core, except at 20 cm depth, where brGDGT-IIIc' was below the detection
207 limit. The brGDGTs in the lake sediments were mainly dominated by pentamethylated brGDGTs ($50.2 \pm 1.8\%$), followed by
208 tetramethylated brGDGTs ($28.9 \pm 0.7\%$), and hexamethylated brGDGTs ($21.0 \pm 1.4\%$; Table 1). The amount of brGDGTs in
209 lake core sediment ranged from 19.9 to $48.0 \mu\text{g g}^{-1} \text{C}$ (Fig. 3a; Table 1). The brGDGT concentration in the surface sediment
210 (0–2 cm), of $37.7 \mu\text{g g}^{-1} \text{C}$, which was about 1.6 times lower than that in the creek sediment at the outlet (Fig. 5a), increased
211 to a maximum of $48.0 \mu\text{g g}^{-1} \text{C}$ around 11 cm depth, and then decreased to a minimum of $19.9 \mu\text{g g}^{-1} \text{C}$ at 23 cm depth (Fig.
212 6b). The concentration of GDGT-0 ranged between $9.0 \mu\text{g g}^{-1} \text{C}$ and $27.1 \mu\text{g g}^{-1} \text{C}$ with an average of $17.4 \pm 6.0 \mu\text{g g}^{-1} \text{C}$,
213 concentration of crenarchaeol ranged from $0.6 \mu\text{g g}^{-1} \text{C}$ to $1.4 \mu\text{g g}^{-1} \text{C}$ with an average of $1.0 \pm 0.2 \mu\text{g g}^{-1} \text{C}$ in the lake
214 sediment core. In general, the concentration of brGDGTs in lake core ($34.0 \pm 8.7 \mu\text{g g}^{-1} \text{C}$; Table 1) was similar with that in
215 river and in woodland, while it was significantly higher than the brGDGTs in soils except for the woodland ($9.6 \pm 4.9 \mu\text{g g}^{-1}$
216 C; $p < 0.05$; Fig. 3a; Table 1).

217 The BIT values for the lake sediment core were rather uniform, varying between 0.95 and 0.97 (Fig.3b). Similarly, the
218 values of $\text{MBT}'_{5\text{ME}}$ along the lake core ranged only between 0.36 to 0.39. The $\text{MBT}'_{5\text{ME}}$ of 0.37 for the lake surface sediment
219 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
220 lake surface sediment was 0.39, which was significantly higher than that in creek bed sediment (0.23 ± 0.02 ; $p < 0.05$; Fig.
221 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
222 maximum value (0.44) at around 10 cm depth, and then decreased with slight fluctuations to 0.22 at 43 cm depth (Fig. 6c).
223 The IR was constant downcore (0.32 ± 0.01 on average; Fig. 3e; Table 1) and was significantly lower than that in creek bed
224 sediment ($p < 0.05$; Fig. 3e).

225 4 Discussion

226 4.1 Spatial variation of brGDGT signals in catchment soils

227 Spatial variations in the relative distribution of brGDGTs in all catchment soils were first evaluated by performing principle
228 component analysis (PCA) using the fractional abundances of the 13 major brGDGTs detected. The first two principal
229 components (PCs) explain 65.2% of the variance in the dataset. PC1 describes 49.5% of the variance, and separates acyclic
230 brGDGT-Ia and brGDGT-IIa from all the other brGDGTs (Fig. 4a). In line with this observation, PC1 has a strong positive
231 relationship with the degree of cyclization of brGDGTs in the soils ($r^2 = 0.97$; Fig. 4c). PC2 describes another 15.7% of the
232 variance, and separates tetramethylated brGDGTs as well as most of the 6-methyl brGDGTs from the majority of the 5-
233 methyl penta- and hexamethylated brGDGTs. As a result, PC2 is negatively correlated with $\text{MBT}'_{5\text{ME}}$ ($r^2 = 0.49$; Fig. 4d) as
234 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 cyclization and



235 the degree of methylation, respectively, the position of the soils in the PCA diagram reveals that different land use types are
236 largely overlapping (Fig. 4b). Indeed, the brGDGTs indices for different land use types are not significantly different ($p >$
237 0.05; Fig. 3), making it difficult to distinguish the provenance of soil OC solely based on brGDGT signatures.

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

254 Some spatial trends are visible in spite of the overall comparable brGDGT signals across the catchment (Fig. A1), which
255 may be explained by variations in other environmental factors than land use or vegetation. Mean air temperature and soil pH
256 have been shown to be the main factors controlling the distribution of brGDGTs in soils worldwide (Weijers et al., 2007a;
257 Peterse et al., 2012; De Jonge et al., 2014b). However, in the small (ca. 4.8 km²) Carminowe Creek catchment, the annual
258 mean air temperature is practically the same for all soils. Similarly, the range in soil pH is relatively small among different
259 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
260 signals based on these parameters. Additionally, the soil water content (SWC) has been shown to affect the distribution and
261 abundance of brGDGTs in soils, either directly by changing the microbial community, or indirectly by altering soil
262 temperature, soil pH, or soil oxygen content (Dirghangi et al., 2013; Menges et al., 2014; Dang et al., 2016). The SWC
263 positively correlates with the abundance of brGDGTs in soils from Qinghai-Tibetan Plateau (Wang et al., 2013), as well as
264 in soils along an aridity transect in the USA (Dirghangi et al., 2013). Moreover, the degree of methylation of 6-methylated
265 brGDGTs is sensitive to the SWC, especially in semi-arid and arid regions (Dang et al., 2016). Although MAP is also the
266 same for the whole catchment, the subtle altitudinal differences in this small creek catchment (i.e. 0-80 m above sea level)



267 may result in an increase in SWC from hilltop to downslope. This would introduce just enough variability in SWC to explain
268 some of the trends in brGDGT signals along hillslope transects. In the north of the catchment, the BIT index values gradually
269 increase > 0.4 from the presumably better aerated soils at the hilltops towards the wetter soils closer to the creek for two of
270 the transects (Fig. A1). The change in BIT index values is driven by both an increase in the amount of brGDGTs and a slight
271 decrease in crenarchaeol concentrations, similar to previous findings (Dirghangi et al., 2013; Wang et al., 2013; Menges et
272 al., 2014). The trend in BIT is likely enhanced by the (minor) change in soil pH along these two transects (from 6.2 to 6.1
273 and from 6.6 to 5.7), which may influence the BIT index as a result of the generally positive relation of crenarchaeol
274 concentrations and a negative relation of brGDGT concentrations with increasing soil pH (Weijers et al., 2006b; Peterse et
275 al., 2010). Nevertheless, these trends in the BIT index are visible in only two of the hilltop transects in the north part of the
276 catchment.

277 Interestingly, also the IR is significantly higher in soils along four transects in north catchment (> 0.36) compared to the
278 average IR value for the rest of the catchment (0.24 ± 0.09 ; $p < 0.05$). The majority of the sites with higher IR are in
279 cropland except for those in the Transect-1, which is under grassland (Fig. A2). Although a relative increase in 6-methyl
280 brGDGTs has been linked to higher soil pH in the global soil dataset (De Jonge et al., 2014a), this relation is not so strong in
281 the catchment soils ($r^2 = 0.36$, $p < 0.001$), likely due to the relatively minor range and variation in soil pH (from 5.4 ± 0.3 to
282 6.6 ± 0.1). Nevertheless, the soils with high IR values in the north catchment have pH values > 6.0 with an average value of
283 6.6 ± 0.1 .

284 4.2 Tracing brGDGTs from soils to creek bed sediments

285 Based on the similar brGDGT signatures for soils under different land use types, these compounds cannot be used to trace
286 back the exact source of the soil OC after mobilization and transport throughout the catchment. However, the concentration
287 and general soil signature of the brGDGTs can be compared with those in creek bed sediments to trace the transfer of OC
288 from the soils into the creeks. The C-normalized concentration of brGDGTs in the creek sediments is higher than that in
289 most of the soils ($34.7 \pm 17.4 \mu\text{g g}^{-1} \text{C}$ and $9.6 \pm 4.9 \mu\text{g g}^{-1} \text{C}$ respectively), except for those in the woodland soils at the
290 riverbanks ($37.6 \pm 11.0 \mu\text{g g}^{-1} \text{C}$; Table 1). Thus, purely based on the concentration, this suggests that brGDGTs in the creek
291 would be primarily derived from the woodland, which also appeared to be the main source of *n*-alkanes in creek bed
292 sediment (Glendell et al., 2018). However, when looking at the relative distribution of the brGDGTs, the percentage of
293 hexamethylated brGDGTs in creek sediments is higher than that in soils ($24.9 \pm 1.8\%$ and $10.9 \pm 0.3\%$, respectively),
294 whereas the percentage of tetramethylated brGDGTs is lower than in soils ($30.1 \pm 1.7\%$ and $39.7 \pm 0.6\%$, respectively; Table
295 1). Furthermore, brGDGTs in creek sediments have a significantly higher IR (i.e. 0.48 ± 0.04) than soils under any of the
296 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,
297 which separates the creek sediments from both the soils and lake sediments on PC2 that is associated with the IR (Fig. 4e).
298 The higher IR in the creek bed sediments can be explained by a contribution of aquatically (i.e. *in situ*) produced 6-methyl



299 brGDGTs. Similar contributions of 6-methyl brGDGTs, and thus higher IR, were also observed in suspended particulate
300 matter from the Yenisei River (De Jonge et al., 2014a), and upstream of the Iron Gates in the Danube River, where the
301 higher IR was coupled to in-river production facilitated by the lower flow velocity and decreased turbidity of the river water
302 (Freymond et al., 2017). Hence, the significantly higher IR in combination with the higher C-normalized concentrations of
303 brGDGTs in the Carminowe creek sediments suggests that the brGDGT signal is mainly aquatic. This implies that the initial
304 soil brGDGT signal is rapidly overprinted by a riverine *in situ* signal upon entering the creek. Only the IR for the
305 downstream site in the northern creek approaches that of the adjacent soil (IR = 0.30 and 0.38 ± 0.02 , respectively; Fig. A2),
306 and may be explained by its use as arable land (Fig. 5a), which involves regular ploughing and subsequent soil mobilization
307 and implies a temporary, local overprint.

308 In attempt to further prove the riverine *in situ* production of brGDGTs and explain the higher IR in creek bed sediment than
309 that in soils, we roughly estimate the minimum amount of 6-methyl brGDGTs produced in creek to reach the higher IR. We
310 assume that the brGDGTs derived from woodland are completely transferred into creek without any degradation, thus, the 6-
311 methyl brGDGTs in creek is composed by 6-methyl brGDGTs produced *in situ* ($6'_{Me \text{ in situ}}$), which we are going to estimate
312 the exact values, and 6-methyl brGDGTs derived from woodland ($6'_{Me \text{ woodland}}$). The minimum amount of 6-methyl
313 brGDGTs produced *in situ* was calculated based on the IR equation using absolute concentration of brGDGTs (Eq. 8).

$$314 \quad IR_{concentration} = \frac{6'_{Me \text{ creek}}}{5'_{Me \text{ creek}} + 6'_{Me \text{ creek}}} = \frac{6'_{Me \text{ woodland}} + 6'_{Me \text{ in situ}}}{5'_{Me \text{ creek}} + 6'_{Me \text{ woodland}} + 6'_{Me \text{ in situ}}} \quad (8)$$

315 The minimum amount of riverine *in situ* 6-methyl brGDGTs production was $10.67 \mu\text{g g}^{-1} \text{ C}$, occupied 93.95% of the total
316 amount of 6-methyl brGDGTs in creek bed sediment we measured. Similarly, when considering the total soil transfer other
317 than only woodland, the minimum amount of riverine *in situ* 6-methyl brGDGTs was $10.66 \mu\text{g g}^{-1} \text{ C}$, accounting for 93.86%
318 of the total 6-methyl brGDGTs in creek bed sediment. It proves that the riverine *in situ* production of 6-methyl brGDGTs is
319 thriving.

320 The absence of a clearly recognizable soil brGDGT signal in the creek bed sediments may be explained by the relatively
321 limited input of soil material into the creek. So far, river systems that have shown to transport a soil-derived brGDGT signal
322 are either characterized by a rainy season (e.g. the Congo River; Weijers et al., 2007b; Hemingway et al., 2017), or have
323 experienced a recent episode of extreme rainfall (e.g. the Danube River, >100 mm in 3 days causing a 100-year flood event;
324 Freymond et al., 2017). The Carminowe creek area does not have a distinct rainy season, and is further characterized by its
325 limited relief. Hence, the relatively minor input of soil-derived brGDGTs seems to be easily overprinted by riverine *in situ*
326 production. Alternatively, the soil-derived brGDGTs are preferentially degraded in an aquatic environment, resulting in a
327 signature that is dominated by brGDGTs that are produced *in situ*.



328 4.3 Sources of brGDGTs in the sediments of Lake Loe Pool

329 In theory, rivers would transport soil-derived OC together with any aquatic OC produced along the way. Once discharged, in
330 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.
331 However, the soil brGDGT signal cannot be recognized in the sediments from Loe Pool since it is already lost upon entering
332 the Carminowe creek. Indeed, the PCA of the relative distributions of brGDGTs indicates that lake sediments plot
333 completely separated from both the soils and creek sediments, mostly due to a higher relative abundance of GDGT-IIIa (Fig.
334 4a, b). As a result, the MBT_{SME} is significantly lower in Loe Pool sediments (0.38 ± 0.00) compared to in the creek bed
335 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
336 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
337 < 0.05 ; Fig. 3d; Table 1). The distinct brGDGT signature of the lake sediments suggests that brGDGTs in the lake again are
338 significantly altered compared to those in the soils and creek sediments. This implies that the riverine brGDGT signal is
339 either replaced or overwritten in the lake.

340 Lacustrine *in situ* production of brGDGTs has been reported in other studies (Sinninghe Damsté et al., 2009a; Tierney and
341 Russell, 2009; Loomis et al., 2011, 2014; Schoon et al., 2013; Weber et al., 2015, 2018). However, there are no generally
342 recognized indicators (yet) to identify lacustrine brGDGT production, although several studies reported a “cold bias” while
343 attempting to reconstruct the mean air temperature (MAT) based on brGDGTs in lake sediments using a soil-based transfer
344 function (Tierney et al., 2010a). In a study on East African lakes, this cold bias was linked to a large *in situ* contribution of
345 brGDGT-IIIa (Tierney et al., 2010a), similar to in Loe Pool. However, the East African lake dataset was generated using the
346 ‘old’ chromatography method that does not separate 5-methyl and 6-methyl brGDGTs. A recent study that has re-analysed
347 the East African Lake dataset indicates that the presumed contribution of GDGT-IIIa mainly consists of brGDGT-IIIa'
348 (Russell et al., 2018), which is less prominent in lake Loe Pool. Although the identity of brGDGT-producer(s) in lakes still
349 remain(s) elusive, a recent study from the stratified Lake Lugano (Switzerland) showed that brGDGTs are mostly produced
350 in the lower, anoxic part of the water column rather than in the sediment (Weber et al., 2018). Furthermore, the combination
351 of brGDGT analysis with molecular biological methods revealed that brGDGTs appeared to be produced by multiple groups
352 of bacteria thriving under different redox regimes in this stratified lake. Specifically, brGDGT-IIIa occurred in the entire
353 water column and continuously increased with depth, whereas brGDGT-IIIa' was mainly produced in the upper, oxygenated
354 part of water column (Weber et al., 2018). Extrapolating the ecological niches of brGDGT production in Lake Lugano to Loe
355 Pool we can speculate that brGDGT-IIIa, which is dominating the brGDGT signal in the Loe Pool sediments, is mostly
356 produced in summer, when the eutrophic state of the lake may seasonally cause the anoxic conditions favourable for its (i.e.
357 brGDGT-IIIa) production. However, our dataset does not allow to further pinpoint the time and depth of lacustrine brGDGT
358 production, or whether brGDGTs are solely produced in the water column of Loe Pool or also in the lake sediment.



359 4.4 Reconstructing local environmental changes based on GDGTs in Loe Pool lake sediments

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

364 The C-normalized concentration of brGDGTs starts to increase around 23 cm, reaching a maximum concentration of $48.0 \mu\text{g}$
365 $\text{g}^{-1} \text{C}$ at 11 cm depth (Fig. 6b). The increased brGDGT concentrations coincide with an increase in the degree of cyclization
366 (Fig. 6c), which generally responds to a change in pH, where more cyclopentane moieties correspond to a higher pH
367 (Weijers et al., 2007a; Schoon et al., 2013). According to historical records, agriculture and anthropogenic perturbations such
368 as mining and urban pollution intensified in the 1960s (~ 26 cm depth), which increased the input of soil and nutrients into
369 Lake Loe Pool (Coard et al., 1983), and resulted in eutrophication (i.e. blooms of cyanobacteria and algae) since at least
370 1986 (~ 23 cm depth) (O'Sullivan, 1992; Flory and Hawley, 1994). Earlier studies have also recognized an increased use of
371 farmyard manures and septic tanks at this time in the nitrogen isotopic composition of the lake sediments, and have detected
372 higher inputs of terrestrial organic material resulting from intensified farming practices and a higher erosion rate during the
373 1960s to 1980s based on ratios of aquatic- and terrestrial-derived plant waxes (Glendell et al., 2018). Thus, the high brGDGT
374 concentrations and DC in the sediments likely reflect the eutrophic conditions of the lake resulting from the increased
375 nutrient input to the lake (Coard et al., 1983). The DC has then recorded the increase in lake water pH associated with
376 eutrophication, whereas brGDGT concentrations express increased aquatic production. Due to remediation measures taken
377 by the local government in 1996 (~ 12 cm depth), the eutrophication has reduced over the past twenty years (Flory and
378 Hawley, 1994; Glendell et al., 2018). The partial recovery of the lake has likely resulted in a return to lower lake water pH,
379 as manifested in the decrease in the DC from ~ 10 cm depth upwards (Fig. 6c).

380 The process of eutrophication and subsequent recovery can also be recognized in the ratio between GDGT-0 and
381 crenarchaeol, which are isoprenoidal GDGTs produced by Archaea. Crenarchaeol is produced by ammonia oxidizing
382 Thaumarchaeota (Sinninghe Damsté et al., 2002) in aquatic environments (Schouten et al., 2000; Powers et al., 2004) and to
383 a lesser extent also in soils (Weijers et al., 2006a), whereas GDGT-0 is a membrane lipid that occurs in all major groups of
384 Archaea, but is indicative of methanogens and thus anaerobic conditions, with a typical ratio of GDGT-0 and crenarchaeol >
385 2 (Blaga et al., 2009). The ratio of GDGT-0/crenarchaeol in the sediments of Loe Pool is > 2 throughout the entire core, and
386 ranges between 10.9 and 24.3, indicating that at least the bottom waters of the lake have been (seasonally) anoxic over the
387 past 100 years (Fig. 6d). The ratio reaches its maximum at 16 cm depth, suggesting that eutrophic conditions and bottom
388 water anoxia were most severe around this time. The recovery of the lake after the remediation measures is again reflected in
389 the return to pre-1960 values (Fig. 6d).



390 **5 Conclusions**

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

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

413 **Data availability**

414 All data are available in the Supplementary Information.

415 **Author Contribution**

416 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
417 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.



418 **Competing interests**

419 The authors declare that they have no conflict of interest.

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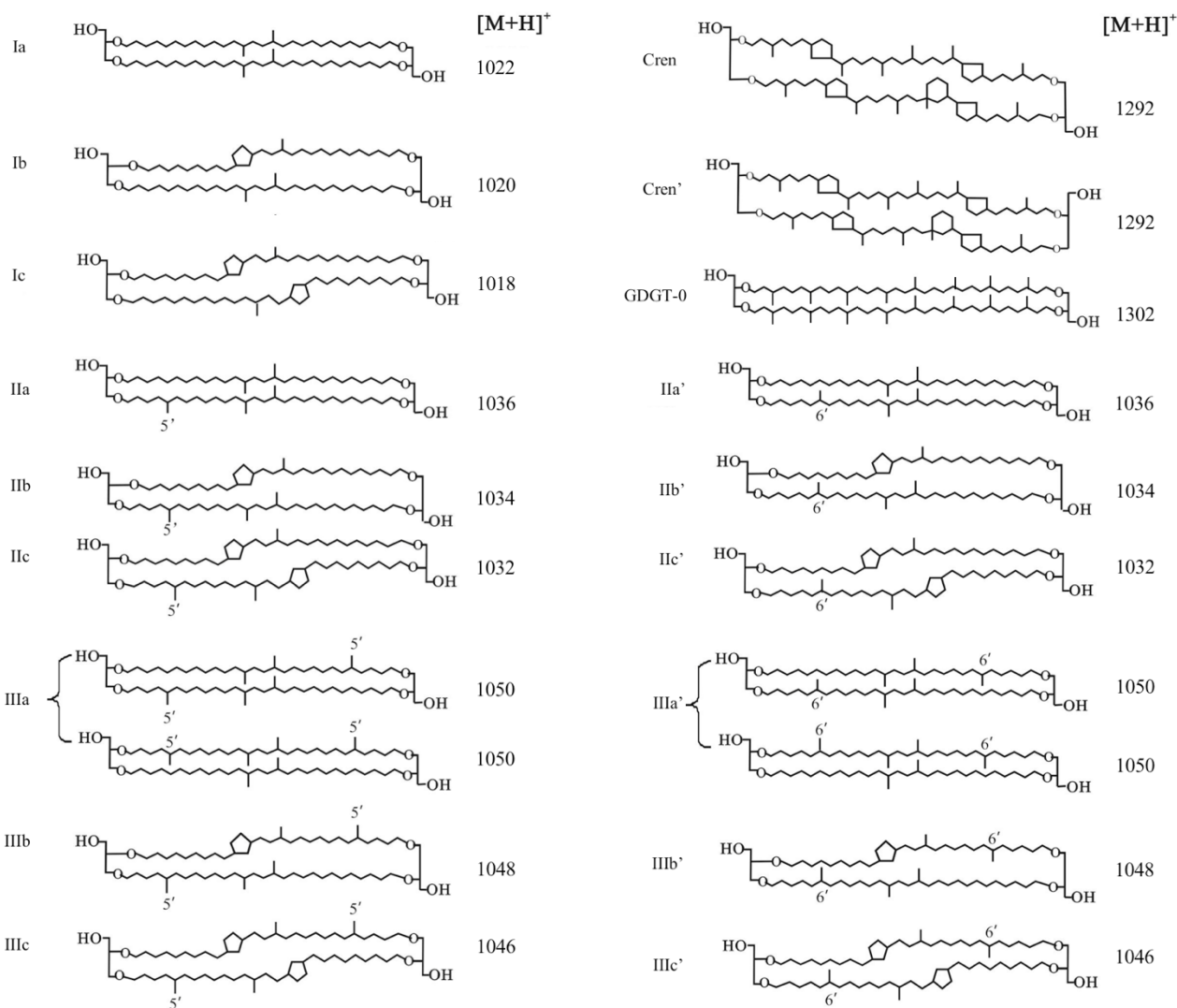
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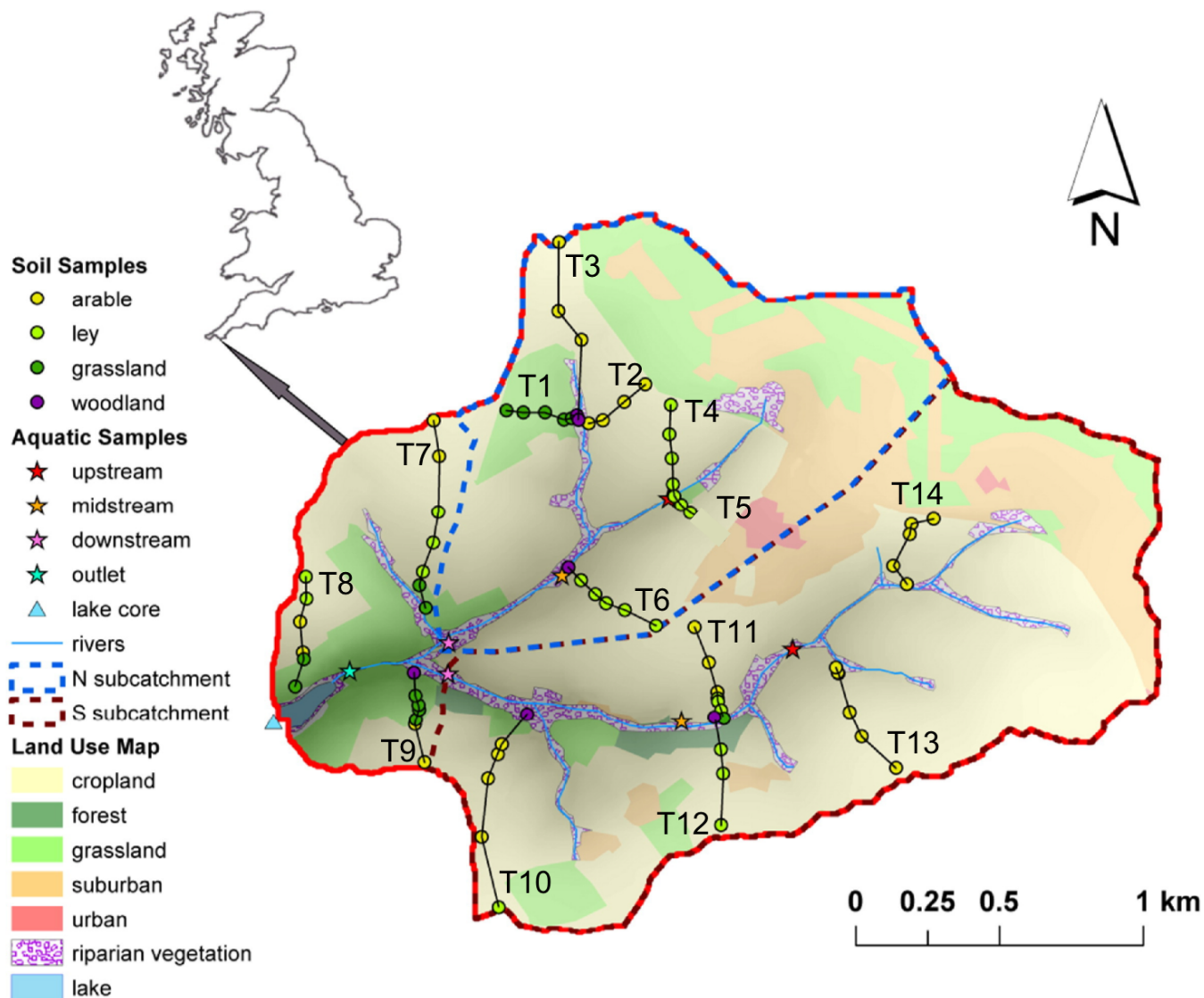


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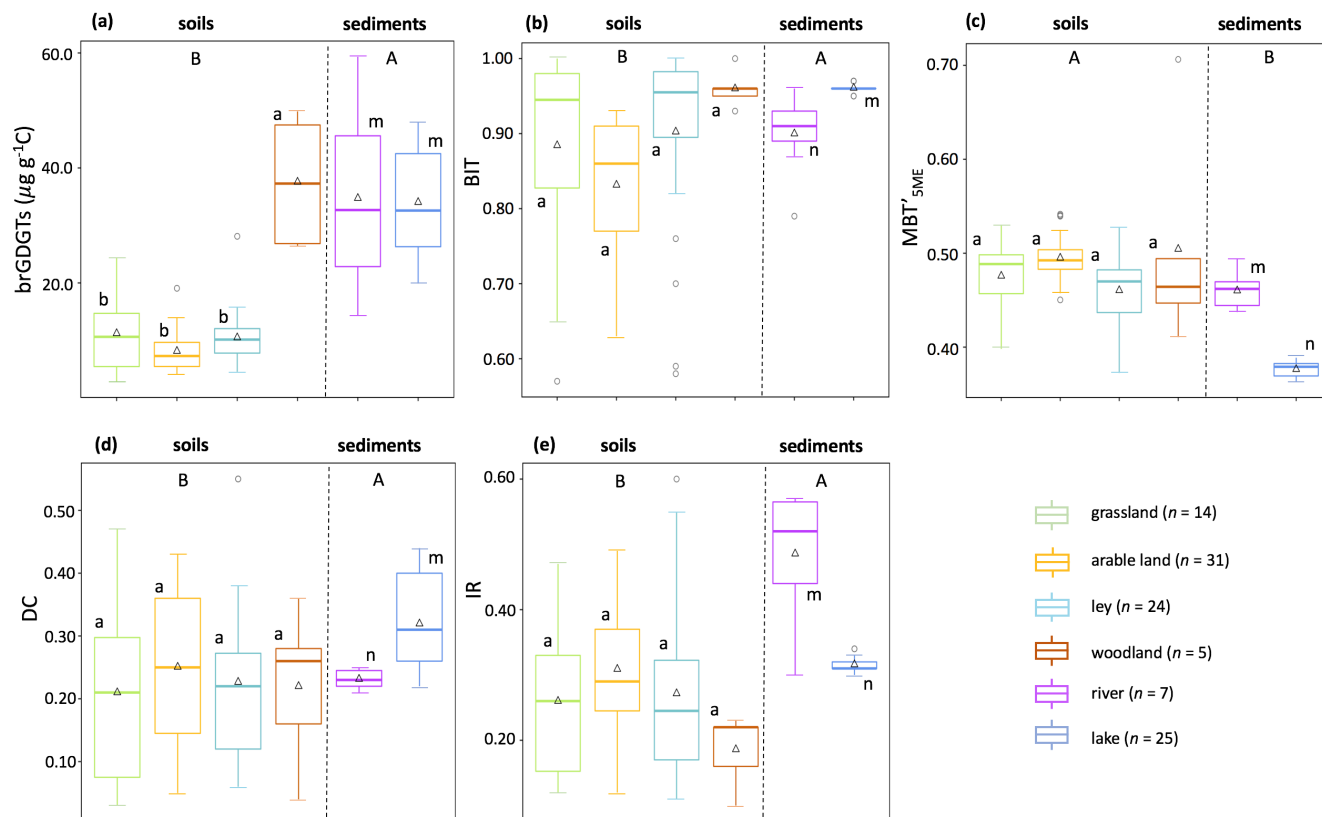
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644 **Figure 1: Molecular structures of 5-methyl and 6-methyl branched GDGTs, GDGT-0 and crenarchaeol. The 6-methyl brGDGTs**
645 **are represented by apostrophe. The structures of penta- and hexamethylated brGDGTs with cyclopentane moiety(ies) IIb',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



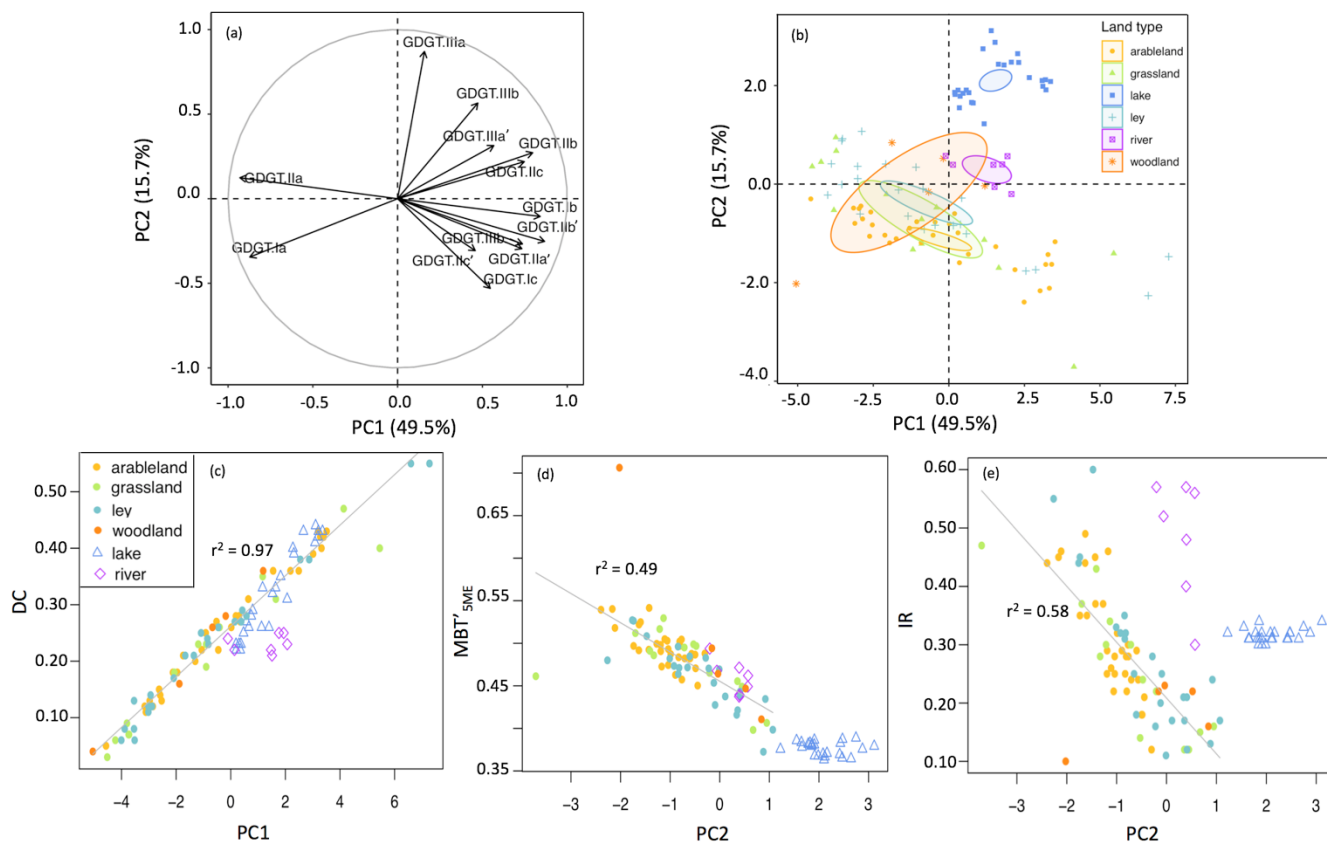
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648 **Figure 2:** Map of the Carminowe Creek catchment in southwest England showing land use types, 14 soil transects (labelled T1-14),
649 creek bed and lake core sediment sampling locations. The coloured circles and stars indicate soil samples under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



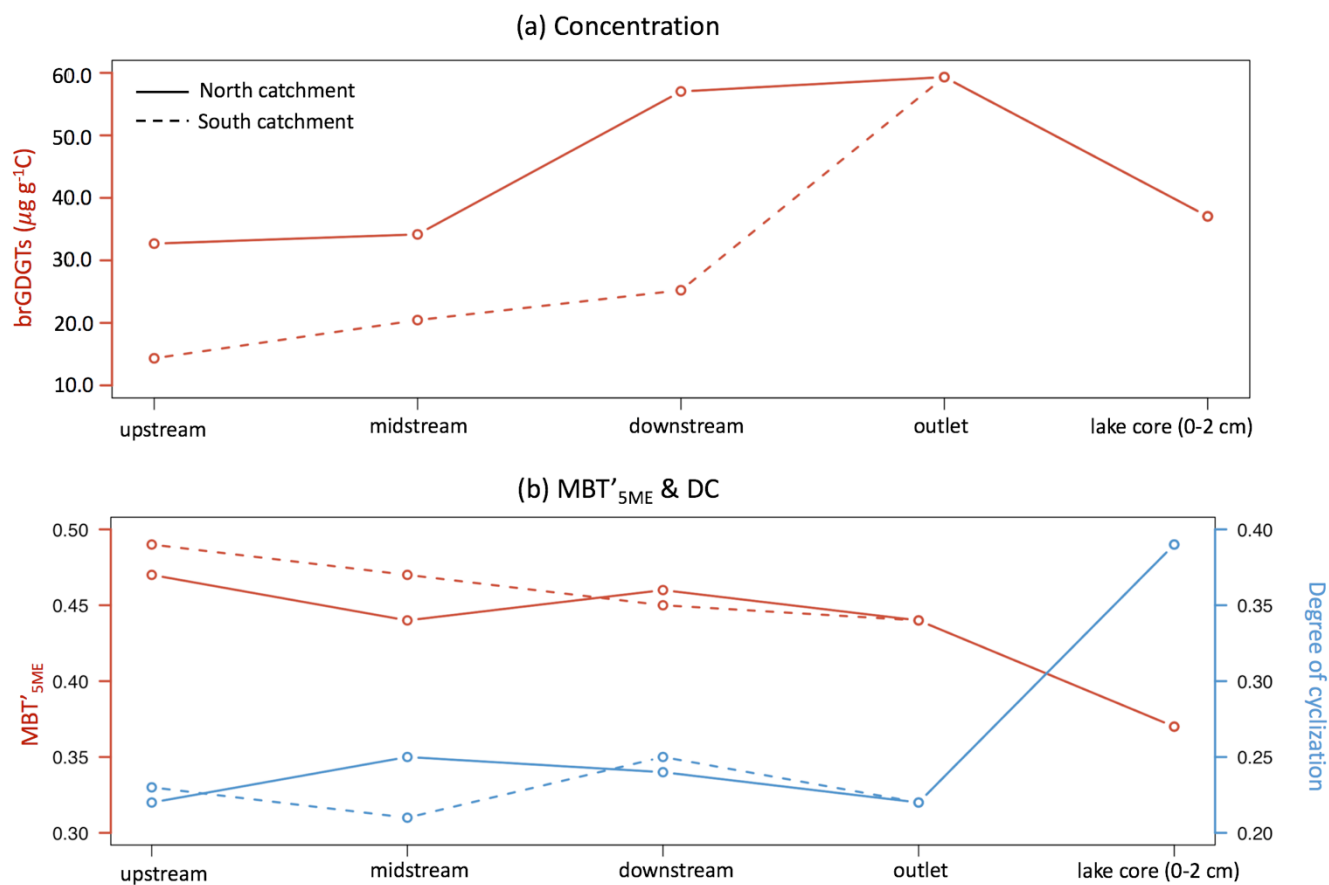
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652 **Figure 3: Box plots displaying (a) the C-normalized concentration of brGDGTs, and brGDGT-based indices: (b) BIT index**
 653 **(branched and isoprenoid tetraether ratio), (c) $\text{MBT}'_{5\text{ME}}$ (methylation of 5-methyl branched tetraethers),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



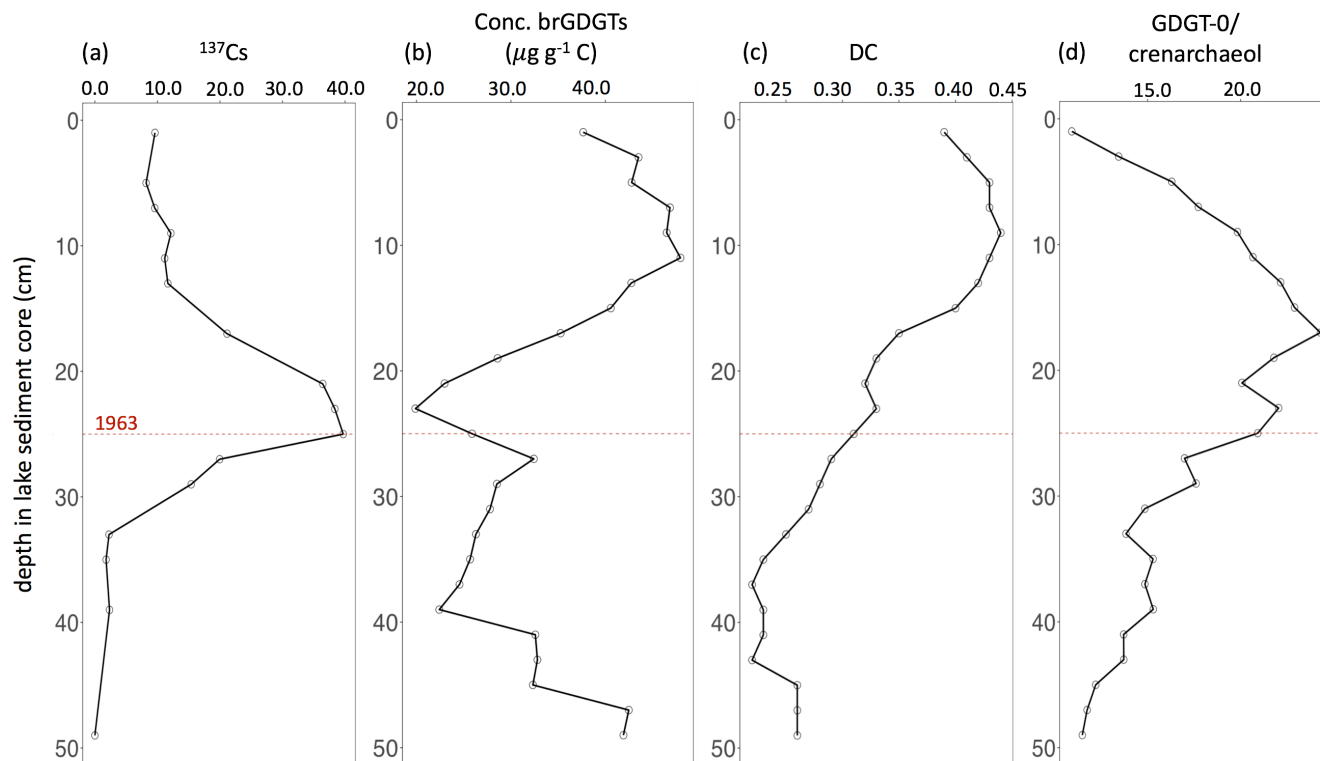
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661 **Figure 4: PCA based on the relative abundances of 13 major brGDGTs. Figure (a) shows the distribution of 13 brGDGTs**
662 **(brGDGT-IIIc and brGDGT-IIIc' are excluded as they are below the detection limit) along the first two PCs, roman numerals and**
663 **English alphabet represent the compounds shown in the Fig. 2. Figure (b) shows sampling sites loading scores on the first two PCs**
664 **and 95% confidence ellipses around the categories of land use: arable land ($n = 31$), grassland ($n = 14$), ley ($n = 24$) and woodland**
665 **($n = 5$), and creek ($n = 7$) and lake ($n = 25$). Figure (c) shows cross plots between PC1 and DC (degree of cyclization). Figure (d)**
666 **and (e) show cross plots of PC2 with MBT'_{5ME} (methylation of 5-methyl branched tetraethers) and IR (isomerization ratio)**
667 **respectively. The linear correlation was calculated excluding creek and lake sediment.**



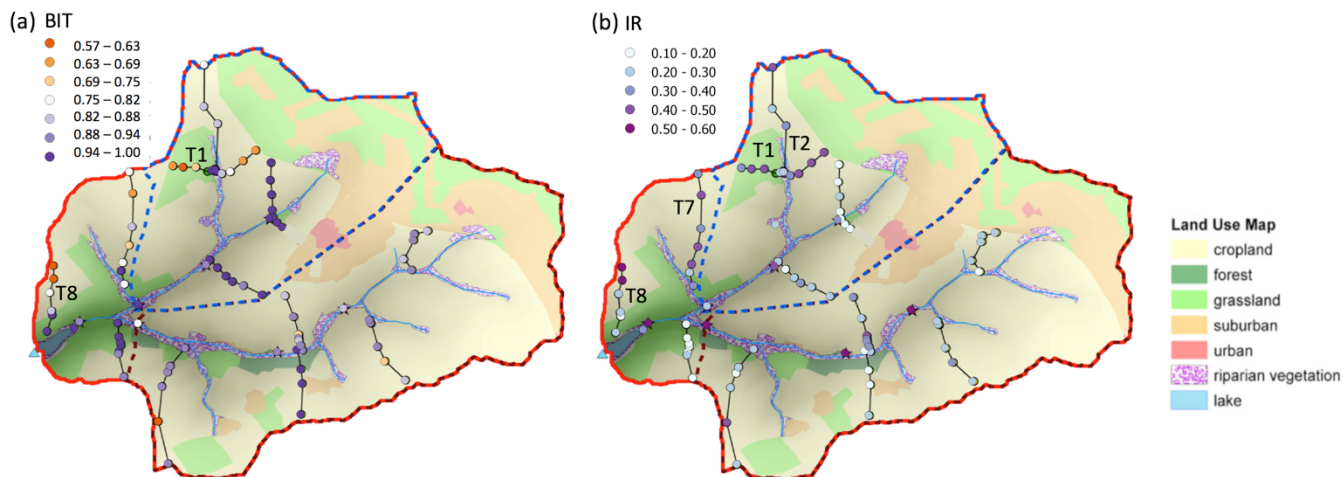
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669 **Figure 5: Spatial variability of (a) C-normalized concentration of brGDGTs and (b) MBT'_{SME} (methylation of 5-methyl branched**
670 **tetraethers) and DC (degree of cyclization) in downstream direction of both substreams in the Carminowe Creek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



671

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



674

675 **Appendix: Spatial variability of the (a) BIT (branched and isoprenoid tetraether ratio) and (b) IR (isomerization ratio) along 14**
676 **soil transects in the Carminowe Creek catchment. The coloured circles show the concentrations and proxy values. Tx indicates soil**
677 **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 indices 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 cyclization), IR (isomerization ratio) (mean ± standard deviation, s.d.).

Land use (n)	C% *	pH	Conc. ($\mu\text{g g}^{-1}$ soil)	Conc. ($\mu\text{g g}^{-1}$ 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

678 *From Glendell et al. (2018)