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1 **Depth related variation of isoprenoid and hydroxylated tetraether lipids in Lake**
2 **Lugu, Southwest China: implications for palaeoenvironmental reconstructions**

3

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18

19 **ABSTRACT**

20 Archaeal glycerol dibiphytanyl glycerol tetraethers (isoGDGTs) and their hydroxylated
21 derivatives (OH-GDGTs) have been increasingly applied to reconstruct past changes in
22 lake temperature and lake-level using down-core sediments. However, a detailed
23 examination of the distribution pattern of iso- and OH-GDGTs in lacustrine sediments
24 is so far limited. To investigate the controls on the sedimentary GDGT distribution in
25 lakes, we examined the archaeal GDGT distribution in surface sediments at different
26 water depths from Lake Lugu, a deep alpine lake in southwest China. Our aim is to
27 determine their distribution, sources and controlling factors. Based on the significant
28 correlations between iso- and OH-GDGTs in deep-water sediments (> 20 m), we
29 suggest that the main biological source of archaeal GDGTs in surface sediments is
30 aquatic Group I.1a Thaumarchaeota (*Nitrosoarchaeum*). The depth-related variation of
31 iso- and OH-GDGTs indicates that water depth is the main factor affecting the
32 distribution of archaeal GDGTs in Lake Lugu, reflecting that Thaumarchaeota prefer to
33 live in the deeper layer above the oxycline. This relationship leads to a positive
34 correlation between %Cren, %OH-GDGTs, and Cren/Cren' with water depth,
35 confirming their potential application for paleo-lake level reconstruction. Our study
36 improves the understanding of the factors that control the archaeal GDGTs in a deep
37 alpine lake and suggest that they might be used as lake-level indicators.

38

39 *Keywords:* isoGDGTs; OH-GDGTs; water depth; biological sources; lake sediments

40

41 **1. Introduction**

42 Isoprenoidal glycerol dibiphytanyl glycerol tetraethers (isoGDGTs) are
43 exclusively produced by archaea and characterized by a varying number of
44 cyclopentane moieties [see [Schouten et al. \(2013\)](#) for a review]. The acyclic GDGT-0
45 is the most common isoGDGT and can be synthesized by all major groups of Archaea,
46 including ammonia-oxidizing Thaumarchaeota, anaerobic methane-oxidizing archaea,
47 methanogens ([Pancost et al., 2001](#); [Blaga et al., 2009](#); [Schouten et al., 2013](#)). IsoGDGT
48 with 1 to 3 cyclopentane moieties (GDGT-1 ~ -3) are commonly found in Euryarchaeota,
49 Crenarchaeota and Thaumarchaeota ([Schouten et al., 2013](#)), and references therein). A
50 structurally unique GDGT containing 4 cyclopentane moieties and 1 cyclohexane
51 moiety is called crenarchaeol ([Sinninghe Damsté et al., 2002](#)), and its isomer
52 (crenarchaeol') is exclusively produced by Thaumarchaeota ([Elling et al., 2017](#);
53 [Sinninghe Damsté et al., 2018](#)). In culture experiments there is a correlation between
54 the production of cyclopentane moieties and temperature ([De Rosa et al., 1980](#); [Elling
55 et al., 2015](#)). Based on this, the degree of cyclization of isoGDGTs with 1–3
56 cyclopentane moieties and crenarchaeol', expressed as TEX₈₆ (TetraEther index of 86
57 carbon atoms), is proposed to reflect surface temperature in marine environments where
58 the predominant source of isoGDGTs is thaumarchaeota ([Schouten et al., 2002](#)). TEX₈₆
59 has also been applied to lacustrine sediments as lake surface temperature proxy ([Powers
60 et al., 2010](#); [Tierney et al., 2010a](#)). However, in lacustrine setting, additional sources of
61 isoGDGTs, for example from methanogens, methanotrophs and other non-
62 thaumarchaeotal sources produced either in the surrounding soil, sediment, or water
63 column, could result in unreliable TEX₈₆-inferred temperature in lakes ([Blaga et al.,
64 2009](#); [Sinninghe Damsté et al., 2012](#); [Li et al., 2019](#); [Yao et al., 2019](#)).

65 Besides the regular isoGDGTs, a structurally similar group of archaeal lipids is
66 called hydroxylated GDGTs (OH-GDGTs). They contain 0-2 cyclopentane moieties
67 (OH-GDGT-0, -1, and -2) and have an additional hydroxyl group located in one of the
68 two biphytanyl chains ([Liu et al., 2012](#)). OH-GDGTs have been found in marine, lake

69 and peat environments (Fietz et al., 2013; Wang et al., 2019; Yang et al., 2019; Kou et
70 al., 2022). Thaumarchaeota Group 1.1a are the likely source organisms of OH-GDGTs,
71 while they are absent from Thaumarchaeota Group I.1b which dominate the archaeal
72 community in most soils (Elling et al., 2015; 2017; Bale et al., 2019). Although the
73 physiological function of the additional hydroxy group remains unknown,
74 hydroxylation may result in the enhanced membrane fluidity to maintain the cell
75 membrane properties based on simulations of molecular dynamics (Huguet et al., 2017).
76 This is consistent with the observation of a significant correlation between the
77 sedimentary distribution of OH-GDGTs and sea water temperature at a global scale
78 (Fietz et al., 2013; Huguet et al., 2013; Lü et al., 2015; Yang et al., 2018; Park et al.,
79 2019). While much previous work on OH-GDGTs has focused on marine settings, the
80 distribution of OH-GDGTs in lake environments is scarcely explored (Wang et al., 2019;
81 Kou et al., 2022).

82 Records of temperature obtained from lacustrine sediment cores based on
83 isoGDGTs have provided promising results within the last decade, these lipids have
84 been increasingly used to reconstruct lake temperatures during the Quaternary (Powers
85 et al., 2011; Blaga et al., 2013; Wang et al., 2015; Morrissey et al., 2018; Sun et al.,
86 2020; Chen et al., 2021). But in addition to temperature, archaeal GDGTs based proxies,
87 such as the fractional abundance of crenarchaeol to total isoGDGTs (%Cren), the ratio
88 of OH-GDGTs to isoGDGTs (%OH-GDGTs), and the ratio of crenarchaeol and
89 crenarchaeol isomer (Cren/Cren') have been proposed as lake-level indicators based on
90 a wide range of lake investigation (Wang et al., 2014a; 2017; 2019; Sun et al., 2020;
91 Kou et al., 2022).

92 However, before any robust paleolimnological interpretation of such palaeo-
93 proxies in lakes is possible, it is important to develop a comprehensive understanding
94 about the origins of GDGTs and their distribution in modern lacustrine sediments, and
95 within the catchment soils. Multiple studies have evaluated the spatial variation of
96 surface sediments with regard to aspects such as organic carbon (Yu et al., 2015), heavy

97 metals (Lin et al., 2016) and lipid biomarkers (Sarkar et al., 2014) within a lake basin.
98 The importance of heterogeneity of organic proxies such as *n*-alkanes has been
99 underlined in two lakes from the Tibetan Plateau (Wang et al., 2012b; Kou et al., 2020).
100 However, the basin-scale spatial distribution of GDGTs and their origin within a single
101 lake has been rarely examined so far, limiting the reliable application of GDGTs from
102 a downcore record to reconstruct past environment change. Thus, evaluating the
103 possible spatial variation of GDGTs in modern lake sediments and catchment on a basin
104 scale could reduce the uncertainties in paleoenvironmental reconstructions.

105 Here we focus on Lake Lugu. It is one of the deepest alpine freshwater lakes in the
106 southeastern margin of the Tibetan Plateau of southwest China (Wang and Dou, 1998).
107 This lake has been considered as an important site for paleolimnological studies due to
108 its sensitivity to changes in the Asian monsoon (Wang et al., 2014b; Sheng et al., 2015;
109 Chang et al., 2018; Zhang et al., 2018). Many studies have showed the significant
110 spatial variability of chironomids (Zhang et al., 2013), diatoms (Wang et al., 2018),
111 trace metals (Lin et al., 2018) and *n*-alkanes (Li et al., 2022) within Lake Lugu.
112 Nevertheless, the spatial distribution pattern of GDGTs and their origin within this lake
113 and catchment are less explored. In order to gain a better understanding of the basin
114 scale heterogeneities and origins of iso- and OH-GDGTs within Lake Lugu, this study
115 mainly focuses on the distribution and origin of archaeal GDGTs from surface
116 sediments with a high spatial resolution. Further, the controlling factor of iso- and OH-
117 GDGTs distribution in Lake Lugu, and the GDGT-based proxy heterogeneities are also
118 discussed for a robust paleolimnological interpretation.

119

120 **2. Materials and methods**

121 *2.1. Study site and sampling*

122 Lake Lugu (27°41'–27°45'N, 100°45'–100°50'E, 2,691 m a.s.l.) is an alpine graben
123 lake that lies at the boundary between Ninglang Yi Autonomous County of Yunnan
124 Province and Yanyuan County of Sichuan Province, the Northwest Yunnan Plateau in

125 southwest China (Fig.1) (NIGLAS, 2019). It is one of the deepest and largest freshwater
126 lakes in China with a maximum water depth of 93.5 m and a mean depth of 40.3 m. The
127 lake has a surface area of 48.45 km² and a catchment area of ca. 171.4 km² (NIGLAS,
128 2019). It lies within the intermontane basin surrounded by mountains, with Maoniuping
129 Peak situated at 3870 m having the highest altitude in this area (Lin et al., 2000). Lake
130 Lugu is featured by a northwest-southeast direction and divided into two interconnected
131 basins, with a peninsula stretching across its center (Zhang et al., 1997). It is a
132 hydrologically semi-closed lake, which is mainly fed by precipitation, ephemeral runoff,
133 and multiple small rivers and streams (NIGLAS, 2019). Two main inflow tributaries
134 enter the lake. In the east the Shankua River and in the south the Sanjiacun River. There
135 is one outflow through the Gaizu River in the southeast that drains the lake via the
136 Caohai wetland into the Yalong River, a tributary of the Yangtze River (Yang, 1984).
137 Lake Lugu is an oligotrophic (TP and TN, 12.7 and 107.9 µg·l⁻¹) and warm temperate
138 monomictic lake, with thermal stratification typically beginning in early April and
139 strong stratification during summer and early autumn. While in early winter the
140 thermocline breaks down and water column mixing occurs in late winter to early spring
141 (Wen, 2017; Wang et al., 2018).

142 The climate of this region is mainly determined by Indian summer monsoon in
143 summer and the southern branch of the westerlies in winter, leading to a temperate
144 climate with distinct dry and wet seasons: approximately 90% of the annual
145 precipitation (~940 mm) occurs between May and October (Wang et al., 2014b; 2018).
146 The mean annual air temperature is ca. 12.8 °C; the mean monthly minimum and
147 maximum temperatures are in December and July (5.6, 19.4 °C, respectively). The
148 terrestrial vegetation in the catchment features a strong altitudinal gradient. Overall, the
149 major plant taxa consist of a mixture of *Acer davidii*, *Populus davidiana* and *Pinus*
150 *yunnanensis* (Li et al., 2013). Submersed macrophytes are mainly distributed in shallow
151 water (<10 m) and are composed of *Ottelia acuminata*, *Potamogeton pectinatus*, and
152 *Potamogeton tepperi* (Tan and Dong, 2011). The main emersed aquatic plants are

153 composed of *Phragmites australis*, *Zizania caduciflora*, *Typha orientalis*, and *Scirpus*
154 *Validus* (Zhao et al., 2016). The predominant soil type in lake catchment consists out of
155 brunisolic soil, red loam, yellow brown soil, purple soil and limestone soil (Zhang et
156 al., 1997).

157 Lake surface sediments at different water depths and mineral soils from the
158 catchment area were obtained during two field campaigns in 2012 and 2015. The
159 locations of the sampling sites are shown in Fig. 1 (For coordinates and water depths,
160 see Supplementary Table S1). 54 lake surface sediments (0–2 cm) were collected using
161 a stainless steel box corer. Nine topsoils (upper 0–2 cm) from within the catchment
162 were also sampled. Surface sediment and soil samples were freeze-dried and ground to
163 powder to homogenize after transport to the laboratory.

164

165 2.2. Lipid extraction and GDGT analysis

166 All samples were ultrasonically extracted as previously described in Li et al.
167 (2022). Briefly, the freeze-dried sample (ca. 5 g dry weight) was extracted four times
168 by ultrasonication using a mixture of dichloromethane and methanol (9:1, v/v). The
169 total lipid extract was subsequently saponified with 6% KOH in methanol. The neutral
170 lipids were extracted with hexane and then separated by column chromatography with
171 activated silica gel as the stationary phase by using hexane and methanol to obtain the
172 apolar and polar fractions, respectively. The polar fraction containing iso- and OH-
173 GDGTs was filtered over a 0.45µm polytetrafluoroethylene (PTFE) filter with
174 hexane/isopropanol (99:1, v/v), and was then dried under gentle stream of N₂ gas prior
175 to further analysis.

176 The archaeal GDGTs were analysed using high performance liquid
177 chromatography-positive ion atmospheric pressure chemical ionization with triple
178 quadrupole mass spectrometry (HPLC-APCI-MS) on an Agilent 1200 series (USA).
179 Analytical separation of GDGTs was achieved using two silica columns in tandem (150
180 mm × 2.1 mm, 1.9 µm; Thermo Finnigan) following the method of Yang et al. (2015).

181 The solvent program started with isocratically eluting for the first 5 min with 84% A
 182 and 16% B, where A = *n*-hexane and B = EtOAc. This was followed by an elution
 183 gradient that consisted out of: 84/16 A/B to 82/18 A/B from 5–65 min and then to 100%
 184 B in 21 min, followed by 100% B for 4 min to wash column and then back to 84/16
 185 A/B to equilibrate the column for 30 min at a constant rate of 0.2 ml/min throughout.
 186 Selected ion monitoring (SIM) mode was used to increase sensitivity and
 187 reproducibility, with *m/z* 1318, 1316, 1314, 1300, 1298 and 1296 as the targeted [M+H–
 188 18]⁺ and [M+H]⁺ ions for detection of OH-GDGTs, while *m/z* 1302, 1300, 1298, 1296,
 189 1292, 1050, 1048, 1046, 1036, 1034, 1032, 1022, 1020 and 1018 for isoGDGTs and
 190 brGDGTs, respectively. Concentrations were determined by integration of peak areas
 191 of the [M+H]⁺ ions of C₄₆ GTGT internal standard (*m/z* 744, concentration of 11.57
 192 ng/μl) (Huguet et al., 2006) and GDGTs of interest, and then normalized to dry weight
 193 (dw). Quantification was semi-quantitative as the relative response factors between
 194 GDGTs and standard were not determined.

195

196 2.3. GDGTs-based proxy calculations

197 Fractional abundances of individual isoGDGT were expressed as the fractional
 198 abundances of the total isoGDGTs, i.e., the sum of crenarchaeol (Cren) and its isomer
 199 (Cren'), GDGT-0, -1, -2, and -3.

200 The TEX₈₆ reflects the degree of cyclization of particular isoGDGTs and was
 201 calculated according to Schouten et al. (2002):

$$202 \text{TEX}_{86} = \frac{([\text{GDGT} - 2] + [\text{GDGT} - 3] + [\text{Cren}'])}{([\text{GDGT} - 1] + [\text{GDGT} - 2] + [\text{GDGT} - 3] + [\text{Cren}'])} \quad (1)$$

203 The branched versus isoprenoid tetraether (BIT) index was calculated following
 204 Hopmans et al. (2004), reflecting the relative abundance of bacterial brGDGTs vs.
 205 archaeal crenarchaeol.

$$206 \text{BIT} = \frac{([\text{Ia}] + [\text{IIa}] + [\text{IIIa}] + [\text{IIa}'] + [\text{IIIa}'])}{([\text{Ia}] + [\text{IIa}] + [\text{IIIa}] + [\text{IIa}'] + [\text{IIIa}'] + [\text{Cren}])} \quad (2)$$

207 The Methane Index (MI) is expressed as the degree of cyclic GDGT-1, -2 and -3

208 to crenarchaeol and its isomer and was calculated according to [Zhang et al. \(2011\)](#):

$$209 \quad MI = \frac{([GDGT - 1] + [GDGT - 2] + [GDGT - 3])}{([GDGT - 1] + [GDGT - 2] + [GDGT - 3] + [Cren] + [Cren'])} \quad (3)$$

210 The Ring Index (RI) represents the weighted average number of cyclopentane
211 rings and was calculated according to [Zhang et al. \(2016\)](#):

$$212 \quad RI = \frac{(0 \times [GDGT - 0] + 1 \times [GDGT - 1] + 2 \times [GDGT - 2] + 3 \times [GDGT - 3] + 4 \times [Cren] + 4 \times [Cren'])}{([GDGT - 0] + [GDGT - 1] + [GDGT - 2] + [GDGT - 3] + [Cren] + [Cren'])} \quad (4)$$

213 The $R_{i/b}$ ([Xie et al., 2012](#)) represents the relative abundance of total isoGDGTs
214 to total brGDGTs:

$$215 \quad R_{i/b} = \frac{\sum isoGDGTs}{\sum brGDGTs} \quad (5)$$

216 The relative abundance of crenarchaeol and its isomer is calculated according to
217 [Li et al. \(2016\)](#):

$$218 \quad Cren/Cren' = \frac{crenarchaeol}{crenarchaeol'} \quad (6)$$

219 Fractional abundances of individual OH-GDGTs were calculated as the fractional
220 abundances of the total OH-GDGTs, i.e., the sum of OH-GDGT-0, -1 and -2. The %OH-
221 GDGTs is calculated following [Fietz et al. \(2013\)](#):

$$222 \quad \%OH-GDGTs = \frac{\sum OH-GDGTs}{\sum OH-GDGTs + \sum isoGDGTs} \quad (7)$$

223 The ring index of OH-GDGTs (RI-OH) represents the weighted average number
224 of cyclopentane moieties and is calculated according to [Lü et al. \(2015\)](#). The revised
225 ring index of OH-GDGTs (RI-OH') is also calculated according to [Lü et al. \(2015\)](#).

$$226 \quad RI - OH = \frac{([OH-GDGT-1] + 2 \times [OH-GDGT-2])}{([OH-GDGT-1] + [OH-GDGT-2])} \quad (8)$$

$$227 \quad RI - OH' = \frac{([OH-GDGT-1] + 2 \times [OH-GDGT-2])}{([OH-GDGT-0] + [OH-GDGT-1] + [OH-GDGT-2])} \quad (9)$$

228

229 *2.4. Statistical analysis*

230 Principal component analysis (PCA) was performed on the fractional abundances
231 of archaeal GDGTs to provide a view of the variability within the iso- and OH-GDGTs
232 in Lake Lugu using CANOCO version 5. Redundancy analysis (RDA) of archaeal
233 GDGTs and related proxies in surface sediments were performed to analyse their

234 relationship with water depth. The spatial distribution of iso- and OH-GDGTs in surface
235 sediments was performed using ArcGIS 10.6 software, with the kriging method of
236 gridding applied for data interpolation. The independent sample *t*-test was performed
237 using SPSS software to discriminate the significance of differences in concentration of
238 iso- and OH-GDGTs in surface sediments at different water depths.

239

240 **3. RESULTS**

241 *3.1. Distribution of iso- and OH-GDGTs in surface sediments and soils*

242 All isoGDGTs (GDGT-0 to -3, crenarchaeol and its isomer, Cren') were detected
243 in the lake surface sediments. However, concentrations were highly variable with three
244 order of magnitude differences, ranging from 8 to 7370 ng g⁻¹ dw, with an average of
245 1360 ng g⁻¹ dw (Table S1). The summed concentration of isoGDGTs was lowest in
246 the surrounding mineral soils with values from 1 to 120 ng g⁻¹ dw (average of 20 ng
247 g⁻¹ dw) (Table S1). The isoGDGT-0 and crenarchaeol were the two most abundant
248 compounds across all sample types (Fig. 2a). Although the average fractional
249 abundances of GDGT-0 and crenarchaeol in surface sediments were similar (avg. 42
250 and 41%), GDGT-0 had a wider range than crenarchaeol. By contrast, the average
251 fractional abundance of crenarchaeol (41%) was higher than GDGT-0 in soils (36%)
252 (Fig. 2a). The fractional abundance of GDGT-1 to -3 was low in lake surface sediments
253 and soils, with an average of 17% and 18%, respectively. The TEX₈₆ and BIT values
254 of soils (0.89 ± 0.06, 0.94 ± 0.07) were higher than surface sediments (0.81 ± 0.08,
255 0.74 ± 0.15) (Fig. 3 and Table S1). However, the R_{i/b} value of surface sediments was
256 much higher than soils, with an average of 0.7 and 0.1, respectively.

257 OH-GDGTs were detected in 51 of the 54 lake surface sediment samples. The three
258 exceptions were LG-1, -2 and -5, all with a shallow water depth of < 3 m (Table S1).
259 The summed concentration of OH-GDGTs ranged from 2 to 1050 ng g⁻¹ dw, with an
260 average of 160 ng g⁻¹ dw (Table S1). The average fractional abundance of OH-GDGT-
261 0 was higher (40%) than OH-GDGT-1 (27%) and OH-GDGT-2 (34%) (Fig. 2b). The

262 RI-OH value of surface sediments was 1.56 ± 0.09 . The concentration of OH-GDGTs
263 in surrounding soils was below the detection limit (Fig. 2b).

264

265 3.2. Statistical analysis of iso- and OH-GDGT distributions in surface sediments

266 The first two components explain a cumulative 92% of the variance in the PCA
267 (Fig. 2c). On the first principal component (PC1, explaining 84% of the variance) the
268 loading of GDGT-0 is opposite to that of all other isoGDGTs. GDGT-1 and -2 are
269 negatively loaded with the second principal component (PC2, explaining ~8% of the
270 variance), while GDGT-3, crenarchaeol and its isomer are in the same quadrant in the
271 PCA biplot (Fig. 2c). RDA analysis of the fractional abundance of isoGDGTs and
272 proxies was performed to analyse the relationship among TEX₈₆, BIT, RI, MI, and water
273 depth (Fig. 2d). Like the PCA result, GDGT-0 is positively loaded on the first axis of
274 RDA (RDA1, explaining 79% of the variance), and has an opposite loading to the other
275 isoGDGTs (Fig. 2d). The RDA2 explains 6% of the variance. Crenarchaeol is negatively
276 loaded on the RDA2, while GDGT-2, -3, and crenarchaeol' are positively loaded on
277 RDA2 (Fig. 2d). The RDA analysis further highlights the negative correlation between
278 GDGT-0 and water depth, while other isoGDGTs are positively correlated with water
279 depth (Fig. 2d). The average values of GDGT-0/Cren exhibit the following profile:
280 shallow-water sediments (51) > soils (1.7) > deep-water sediments (0.6) (Fig. 3e and
281 Table S1).

282 The PCA and RDA results can be verified by the distribution pattern of isoGDGTs
283 in surface sediments. The isoGDGTs in shallow lake sediments (water depth < 20 m, n
284 = 14) are dominated by GDGT-0, which constitute ~80% of the total isoGDGTs (Fig. 2a
285 and Table S1). However, crenarchaeol is the main component in deep sediments (water
286 depth > 20 m, n = 40), with an average of 51% (Fig. 2a and Table S1).

287 The PCA and RDA results of OH-GDGTs suggest that OH-GDGTs are less
288 affected by water depth compared to isoGDGTs (Figs. 2e and f). This can be verified by
289 the distribution pattern of OH-GDGTs in surface sediments (Fig. 2b and Table S1). For

290 example, OH-GDGT-0 is dominant in both in shallow (< 20 m) and deep (> 20 m)
291 sediments, with the average of $42 \pm 11\%$ and $39 \pm 3\%$, respectively. Similarly, OH-
292 GDGT-2 (average, $\sim 35\%$) is higher than OH-GDGT-1 (average, $\sim 25\%$) in shallow and
293 deep sediments (Fig. 2b).

294 The different archaeal GDGT distributions between shallow and deep sediments
295 is also seen in their related proxies, such as BIT, RI and Cren /Cren' (Fig. 3). However,
296 OH-GDGT based proxies such as RI-OH and RI-OH' show no substantially difference,
297 as only %OH-GDGTs differ between shallow and deep sediments (Fig. 3h).

298

299 4. DISCUSSION

300 4.1. Potential sources of iso- and OH-GDGTs in surface sediments

301 4.1.1. isoGDGTs

302 The average summed concentration of isoGDGTs in lake surface sediments is
303 much higher than that of soils (Table S1). The substantial discrepancy in concentration
304 between sediments and soils can be explained by *in situ* production of isoGDGTs within
305 the lake. This is further supported by the observation that the different isoGDGT-based
306 proxies are different in lake sediment samples compared to mineral soils (Fig. 3). For
307 example, soils have a higher TEX₈₆ and BIT values than lake sediments (Figs. 3b and
308 c), whereas lake sediments exhibit higher GDGT-2/GDGT-3 and Cren/Cren' ratios
309 compared to soils (Figs. 3f and g), all of which demonstrated by an independent sample
310 *t*-test with $p < 0.05$. Our finding of *in situ* production within this large lake system is
311 consistent with previous studies (Li et al., 2019; Wang et al., 2019; Yao et al., 2019;
312 Baxter et al., 2021; Kou et al., 2022; Sinninghe Damsté et al., 2022). Moreover, the
313 depth related variation in isoGDGTs indicates that there is a deep-water contribution of
314 crenarchaeol. The relative low concentration of oxygen in the deeper and colder
315 (suboxic) part of the lake likely provides a niche for nitrifying thaumarchaea (Könneke
316 et al., 2005; Wuchter et al., 2006; Qin et al., 2017), allowing the proliferation of
317 Thaumarchaeota as seen in other lakes (Baxter et al., 2021; Sinninghe Damsté et al.,

318 2022). More details of depth related variation of isoGDGTs distribution are discussed
319 in Section 4.2.

320 The ratio of isoGDGT-0 over crenarchaeol (GDGT-0/Cren) has been used to infer
321 the relative contribution of methanogenic archaea producing isoGDGTs within lakes
322 with values >2 indicative of a significant methanogen contribution (Blaga et al., 2009;
323 Li et al., 2019). The high values of GDGT-0/Cren in shallow-water sediments (average
324 ~40) thus indicates a dominant contribution of methanogens to the sedimentary
325 isoGDGT pool (Fig. 3e). This methanogen input is unlikely to be derived from soils in
326 the catchment area as the ratio is much lower (average <1) in mineral soils, indicative
327 of a major contribution of Thaumarchaeota (Fig. 3e). Similarly low ratios are found in
328 deep-water sediments, implying a major contribution of Thaumarchaeota and a minor
329 contribution of methanogens in the deeper part of the lake (Fig. 3e). The concentrations
330 of GDGT-0 and crenarchaeol are not correlated in shallow-water sediments ($R^2 = 0.01$,
331 $p < 0.05$, $n = 14$) (Fig. 4f), but do exhibit a significant linear correlation ($R^2 = 0.98$, $p <$
332 0.05 , $n = 40$) in deep-water sediments. This result strongly suggests that contribution to
333 the isoGDGT pool between shallow- and deep-water sediments is different.

334 The ratio of crenarchaeol over its isomer (Cren/Cren') has been used to
335 differentiate between Group I.1a versus I.1b Thaumarchaeota as they produce
336 crenarchaeol and crenarchaeol' in varying proportions. High ratios of Cren/Cren' (>25)
337 are indicative for the Group I.1a Thaumarchaeota, while a lower ratio (<25) is
338 indicative for Group I.1b Thaumarchaeota (Li et al., 2016; Kou et al., 2022). Both the
339 average Cren/Cren' values of shallow- and deep-water sediments (averages of 41 and
340 74) are higher than soils (9) (Fig. 3g). This suggests the dominance of aquatic Group
341 I.1a Thaumarchaeota in lake sediments, while Group I.1b Thaumarchaeota are
342 abundant in soils and further demonstrates that the contribution from isoGDGTs
343 produced in the catchment soils to the lake sediments is small.

344 Recently, analysis of metagenome-assembled genomes (MAGs) of
345 Thaumarchaeota (now referred as the class *Nitrososphaeria* in the phylum

346 *Thermoproteota*, Rinke et al. (2021)) was performed on surface sediments along a water
347 depth gradient in Lake Lugu. It revealed that the dominant thaumarchaeal OTU was
348 *Nitrosoarchaeum* (Ren and Wang, 2022). Additionally, the archaeal 16S rRNA gene
349 amplicon data clarified the depth variation of *Nitrosoarchaeum* (affiliated to Group I.1a)
350 abundance, with a peak ranging from 20 to 50 m depth around the thermocline zone,
351 showing less variations toward shallow layers above 20 m and deep layers below 50 m
352 (Ren and Wang, 2022). This depth variation of *Nitrosoarchaeum* is consistent with the
353 concentration profile of isoGDGTs in surface sediments along our water depth gradient,
354 with the highest concentration at ~ 50 m (Fig. S1a–g). This finding confirms the
355 biological source of isoGDGTs in Lake Lugu to be predominantly Thaumarchaeota
356 (*Nitrososphaeria*). These results consist with previous studies in deep oligotrophic
357 lakes where *Nitrosoarchaeum* is the dominant thaumarchaeal OTU (Hiraoka et al., 2019;
358 Podowski et al., 2022), while in other lakes *Nitrosopumilus* is the main thaumarchaeal
359 OTU (Herber et al., 2020; Klotz et al., 2022; Sinninghe Damsté et al., 2022). This
360 discrepancy may be due to the varying climate and environmental conditions in
361 different regions.

362

363 4.1.2. OH-GDGTs

364 OH-GDGTs have been reported in the cultures of Nitrosopumilales, such as the
365 *Nitrosopumilus* and *Nitrosoarchaeum* clades, which are attributed to Group I.1a
366 Thaumarchaeota (Elling et al., 2017; Bale et al., 2019). However, the Nitrososphaerales
367 (Group I.1b Thaumarchaeota) which are typically the dominant Thaumarchaeota in
368 mineral soils do not produce OH-GDGTs (Elling et al., 2017; Bale et al., 2019). In Lake
369 Lugu, OH-GDGTs are only detected in lake surface sediments and not in the soils from
370 the catchment, indicating that these lipids must be produced *in situ* within the lake. This
371 is consistent with our finding of higher concentrations of summed and individual OH-
372 GDGTs in deep-water sediments compared to shallow-water sediments (Fig. S1h–k).
373 This result is consistent with previous findings of OH-GDGTs reported in mid-latitude

374 Asia, Tibetan Plateau and central European lakes (Wang et al., 2019; Kou et al., 2022;
375 Sinninghe Damsté et al., 2022).

376 The biological source of OH-GDGTs in deep-water sediments might be similar to
377 that of the isoGDGTs, as there is a high correlation between the concentrations of both
378 lipid classes ($R^2 = 0.98$, $p < 0.05$, $n = 40$) (Fig. 4f). Moreover, the concentration profiles
379 of the total and individual OH-GDGTs all correlate well with those of GDGT-0 (avg.
380 $R^2 = 0.96$) and crenarchaeol (avg. $R^2 = 0.99$) in deep-water sediments (Figs. 4g, h and
381 S2), strongly suggesting that OH-GDGTs are also sourced from Thaumarchaeota, likely
382 predominantly *Nitrosoarchaeum* (Ren and Wang, 2022). However, there is no strong
383 relationship between OH-GDGTs and isoGDGTs such as GDGT-0 and crenarchaeol in
384 shallow-water sediments (Figs. 4g, h and S2). Only OH-GDGT-0 is moderately
385 correlated ($R^2 = 0.31$, $p < 0.05$, $n = 14$) to GDGT-0, indicating that OH-GDGT-0 might
386 have a (partial) methanogens origin. No correlation between the concentrations of OH-
387 GDGT-0 and crenarchaeol ($R^2 = 0.18$, $p < 0.05$, $n = 14$) also implies that thaumarchaeota
388 are unlikely a major source for OH-GDGT-0 in shallow-water sediments (Fig. S2b).
389 Notably, the concentrations of OH-GDGT-2 and crenarchaeol are strongly correlated
390 ($R^2 = 0.89$, Fig. S2f) in shallow-water sediments, suggesting they may share a similar
391 biological source of *Nitrosoarchaeum* (Ren and Wang, 2022).

392 In summary, the difference in two depth profiles of OH-GDGTs is most likely due
393 to their different sources in shallow- and deep-water sediments. The identical
394 relationship between crenarchaeol and OH-GDGTs in deep-water sediments can be
395 interpreted as their similar biological source, i.e., *Nitrosoarchaeum*, while a
396 contribution of methanogens for OH-GDGT-0 cannot be excluded in shallow-water
397 sediments.

398

399 4.2. Depth-related variation of iso- and OH-GDGTs in surface sediments

400 The isoGDGTs in surface sediments of Lake Lugu are predominantly produced *in*
401 *situ* within the lake and their relative abundances and concentrations are strongly

402 affected by water depth (Figs. S1 and S3). The RDA analysis indicates that water depth
403 is an important factor influencing lake sedimentary isoGDGTs distribution (Fig. 2d).
404 GDGT-0 and crenarchaeol are the two main components of isoGDGTs in sediments
405 with regards to shallow- and deep-waters (Fig. 2), their fractional abundances are both
406 well correlated with water depth ($R^2 = 0.75$ and 0.7 , respectively, $n = 54$) (Figs. 4a and
407 b). Moreover, the relative abundances of other isoGDGTs also exhibit strong/moderate
408 relationships with water depth (GDGT-1, -2, and crenarchaeol, with the R^2 of 0.77 , 0.56
409 and 0.4 , Fig. S3). Thus, our results collectively demonstrate that the isoGDGT
410 distribution in surface sediments of Lake Lugu is primarily controlled by water depth.
411 Although the RDA result indicates that the correlation between OH-GDGTs and water
412 depth is not as strong as that of isoGDGTs (Fig. 2f), their distribution in surface
413 sediments is also associated with water depth. It should be noted that the %OH-GDGTs
414 is significantly correlated with water depth ($R^2 = 0.7$, $p < 0.05$, $n = 54$) (Fig. 4d). As the
415 OH-GDGTs and crenarchaeol are suggested to share similar biological source of
416 *Nitrosoarchaeum* in deep-water sediments (Section 4.1), their strong positive
417 correlation with water depth can be interpreted as a common ecological response of
418 *Nitrosoarchaeum* to water depth in lake environments.

419 Lugu Lake is characterized by substantially enhanced concentrations of iso- and
420 OH-GDGTs in deep-water sediments compared to shallow-water sediments (Fig. S1).
421 The depth-related variations of iso- and OH-GDGTs in surface sediments is confirmed
422 in their interpolated spatial distribution of related proxies (Fig. 5). Crenarchaeol is the
423 most abundant isoGDGTs in deep-water sediments and its average concentration is
424 several orders of magnitude higher ($860 \text{ ng g}^{-1} \text{ dw}$) than that of shallow-water sediments
425 ($20 \text{ ng g}^{-1} \text{ dw}$) (Fig. S1 and Table S1). Thus, low BIT values in deep-water sediments
426 can be explained by the depth-dependent variation of crenarchaeol with increased
427 production in the deeper lake water (Figs. 5d and 4c). Thaumarchaeota are prone to
428 thrive in deep water above the oxycline of deep lakes (Buckles et al., 2013; Baxter et
429 al., 2021; Sinninghe Damsté et al., 2022). In Lake Lugu, the proper niche for

430 *Nitrosoarchaeum* to proliferate is between 20–50 m depth (Ren and Wang, 2022). This
431 control is different from that found for other lipid classes.

432 A depth variation in crenarchaeol abundance in surface sediments has also been
433 observed in lakes from East Africa, midlatitude Asia, and the Tibetan Plateau (Tierney
434 et al., 2010b; Wang et al., 2019; Kou et al., 2022). Based on the significant strong
435 correlation between crenarchaeol and water depth, the ratio of crenarchaeol to total
436 isoGDGTs (%Cren) has been proposed as an indicator for lake level and has been used
437 to reconstruct past changes in lake levels (Wang et al., 2014a; 2017). Recently, %OH-
438 GDGTs (the ratio of total OH-GDGTs to total isoGDGTs) (Wang et al., 2019), and the
439 Cren/Cren' have been applied to reflect the lake level change as well (Sun et al., 2020;
440 Kou et al., 2022). However, the correlations between GDGT-based indices and water
441 depth in these previous studies are weaker compared to our dataset. For example, the
442 correlation coefficient (R^2) between water depth and %Cren in Lake Lugu is 0.77 ($n =$
443 54) (Fig. 4b), however, the correlation coefficient decreases to 0.46 when the dataset (n
444 = 227) of all available midlatitude Asia and Tibetan Plateau lakes was added (Fig. 6a).
445 Similar variations have been observed between the relationship of %OH-GDGTs,
446 Cren/Cren' and water depth. They exhibit moderate ($R^2 = 0.55$) and low ($R^2 = 0.47$)
447 correlations with water depth when previous reported lake datasets are added (Figs. 6b
448 and c). It is likely that on a global scale other potential influencing factors such as
449 temperature, pH, nutrient status, and salinity affect these relationships.

450 The significant strong correlation between BIT and water depth ($R^2 = 0.71$) in Lake
451 Lugu (Fig. 4c), is also consistent with previous findings. Similar relationships have
452 been observed in 82 lakes from a global database (Blaga et al., 2010), as well as in East
453 African lakes (Tierney et al., 2010b), Chilean lakes (Kaiser et al., 2015), Tibetan Plateau
454 lakes (Wang et al., 2012a; 2016; Kou et al., 2022) and midlatitude Asian lakes (Wang
455 et al., 2019). Although the correlation between BIT and water depth ($R^2 = 0.31$, $p < 0.05$,
456 $n = 227$) is not as strong as Lake Lugu when all lakes are integrated, the BIT exhibits a
457 negatively relationship with water depth in most lakes (Fig. 6d). This relationship is

458 probably due to the preferred habitat for Thaumarchaeota in deeper waters.

459

460 4.3. Implications for paleolimnological studies

461 Iso- and OH-GDGT based proxies, i.e., %Cren, %OH-GDGTs and Cren/Cren'
462 have been used to reconstruct lake water level from ancient and modern lake sediments
463 (Wang et al., 2017; 2019; Sun et al., 2020; Wu et al., 2020; Chen et al., 2021). In Lake
464 Lugu, these archaeal GDGTs also correlate well with water depth and can serve as the
465 indicator of lake level. However, a prerequisite for applying these archaeal GDGT
466 indicators is to ascertain that nitrifying Thaumarchaeota could thrive under certain lake
467 condition, such as the larger and deeper lakes. Thus, the oligotrophic/mesotrophic lakes
468 with sufficient oxygen for ammonia oxidation do offer a proper niche for the nitrifying
469 thaumarchaea such as *Nitrosopumilale* and *Nitrosoarchaeum*, and subsequently leading
470 to the high production of crenarchaeol. In contrast, small and shallow lakes that are
471 prone to be affected by eutrophication, which favor high production of GDGT-0 but
472 minor crenarchaeol or OH-GDGTs, limiting the application of archaeal GDGTs as lake-
473 level indicator.

474 The spatial distribution of iso- and OH-GDGTs in Lake Lugu have showed that
475 water depth is the main factor for their distribution, which is different from the spatial
476 heterogeneity of *n*-alkanes that are mainly affected by soil erosion in this lake (Li et al.,
477 2022). The depth variation of archaeal GDGTs in Lake Lugu revealed that the fractional
478 abundances of iso- and OH-GDGTs remain constant between the depth range of 40–60
479 m (Figs. 4 and S3). For example, %Cren reached the maximum (~ 60) at the water depth
480 of 40 m and show less variability at the depth range < 60 m (Fig. 6a). Similar
481 distribution pattern of %Cren is also observed in mid-altitude Asian and Tibetan Plateau
482 lakes (Wang et al., 2019; Kou et al., 2022), the %Cren all remain unchanged at the water
483 depth > 40 m (Fig. 6a). This likely reflects the physiology of Thaumarchaeota in lake
484 environments, which prefer to live in the deep layer of lake with oxygen.

485

486 **5. CONCLUSIONS**

487 We examined the distribution of iso- and OH-GDGTs in surface sediments from
488 different water depths of Lake Lugu as well as mineral soils from the catchment to
489 determine their sources and factors that impact their distributions. Different isoGDGTs
490 distribution between mineral soils and lake surface sediments indicates that isoGDGTs
491 in lake sediments are mainly produced within the lake, while the terrestrial contribution
492 is minor. The biological source of crenarchaeol in lake sediments is probably
493 *Nitrosoarchaeum*, the dominant thaumarchaeal OTU based on a previous molecular
494 biological study. Our data show that water depth is the main factor affecting the archaeal
495 GDGT distribution in Lake Lugu. Both the %Cren and %OH-GDGTs of surface
496 sediments are correlated with water depth, suggesting that they could be used as proxies
497 for past lake-level changes.

498

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708

Figure Captions

Fig. 1. Bathymetry of Lake Lugu and sampling sites. Surface sediments ($n = 54$) from different water depths and topsoils ($n = 9$) from the catchment of Lake Lugu were collected. The light blue dots indicate samples collected at the water depth < 20 m ($n = 14$). The dark blue dots represent samples collected at the water depth > 20 m ($n = 40$). The yellow triangles indicate soils collected in the catchment of Lake Lugu.

Fig. 2. (a) Average distribution of isoGDGTs in surface sediments and surrounding soils within the catchment of Lake Lugu. (b) Average distribution of OH-GDGTs in surface sediments. (c) Principal component analysis (PCA) on the distribution of isoGDGTs from surface sediments and surrounding soils. (d) Redundancy analysis (RDA) on the distribution of isoGDGTs from surface sediments and surrounding soils. (e) PCA biplot of OH-GDGTs in surface sediments. (f) RDA triplot of OH-GDGTs in surface sediments. The light blue dots indicate samples collected at the water depth < 20 m ($n = 14$). The dark blue dots represent samples collected at the water depth > 20 m ($n = 40$). The yellow triangles indicate soils collected in the catchment of Lake Lugu.

Fig. 3. Boxplots of the concentration of crenarchaeol (a); TEX₈₆ (b); BIT(c); RI(d); GDGT-0/Cren(e); GDGT-2/GDGT-3(f); Cren/Cren'(g) in the surface sediments and surrounding soils. (h) Boxplot of %OH-GDGTs in the surface sediments. Shallow sediments indicate samples collected at the water depth < 20 m ($n = 14$). Deep sediments represent samples collected at the water depth > 20 m ($n = 40$).

Fig. 4. Cross plots of %GDGT-0 (a), %Cren (b), BIT (c), and %OH-GDGTs (d) in surface sediments versus water depth. Cross plots of the concentrations of GDGT-0 and crenarchaeol (e), summed OH-GDGTs and isoGDGTs (f), summed OH-GDGTs and GDGT-0 (g), and summed OH-GDGTs and crenarchaeol (h) in the surface sediments. The light blue triangles indicate samples collected at the water depth < 20 m. The dark blue dots represent samples collected at the water depth > 20 m. The linear regression of shallow sediment (< 20 m) is indicated in light blue. The linear regression of deep sediment (> 20 m) is indicated in dark blue. The black linear regression represents for all sediment samples ($n = 54$).

Fig. 5. Spatial distribution of %Cren (a), %OH-GDGTs (b), Cren/Cren' (c), and BIT

(d) in the surface sediments of Lake Lugu.

Fig. 6. Cross plots of %Cren (a), %OH-GDGTs (b), Cren/Cren' (c), and BIT (d) in surface sediments versus water depth. The dark grey dots indicate samples in this study. The light grey dots include lake samples from this study, midlatitude Asia ([Wang et al., 2019](#)) and Tibetan Plateau ([Kou et al., 2022](#)).

Fig. 1.

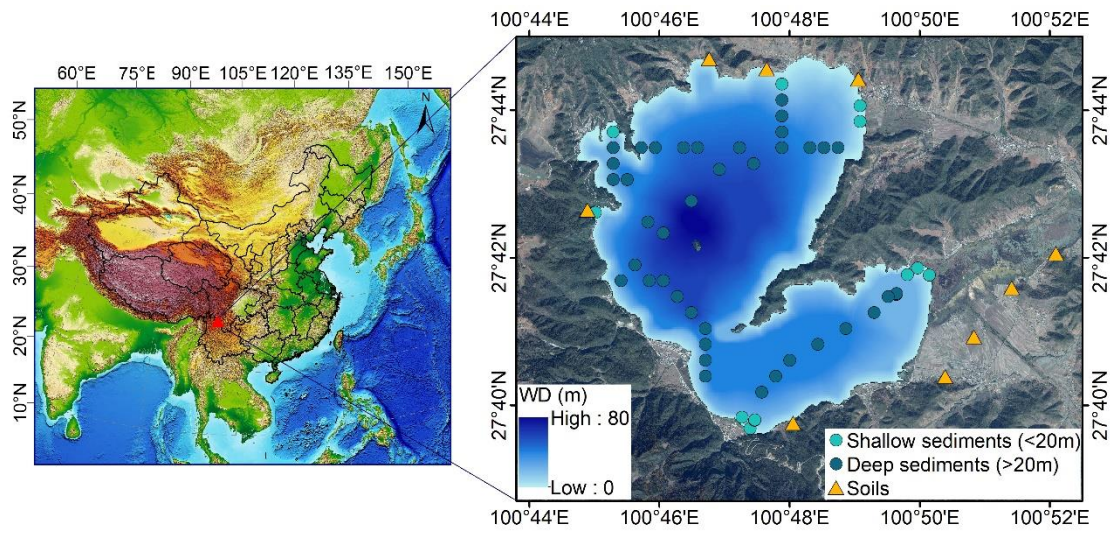


Fig. 2.

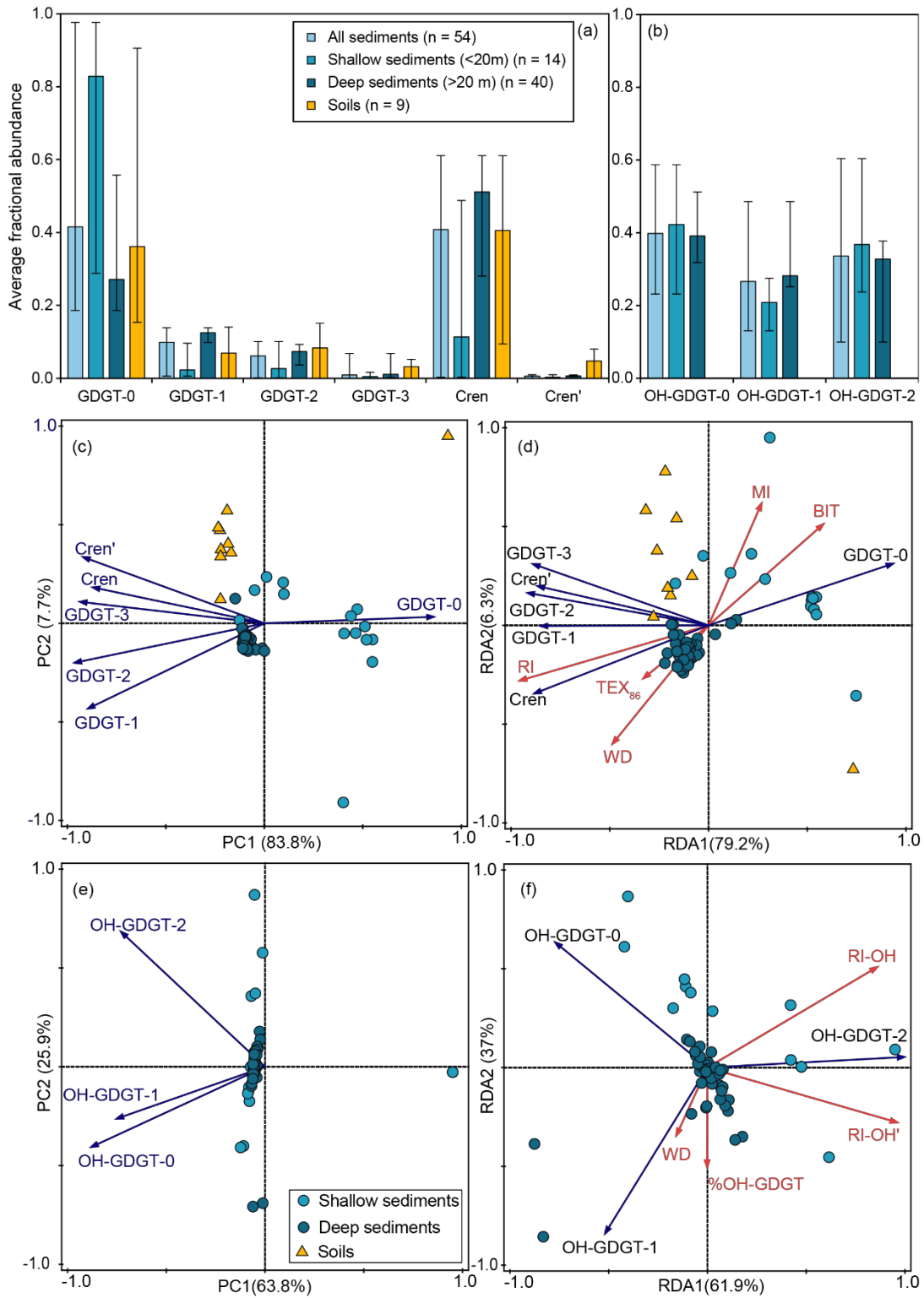


Fig. 3.

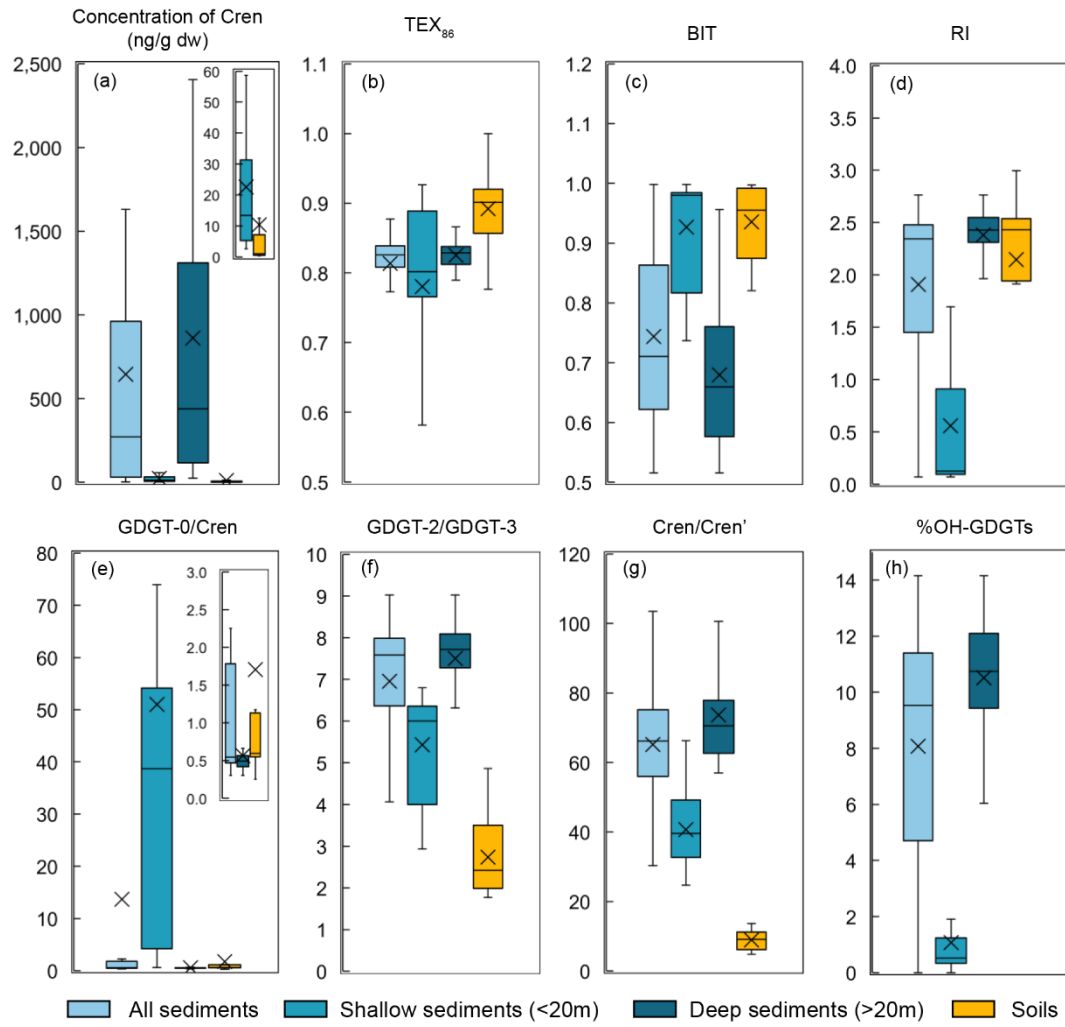


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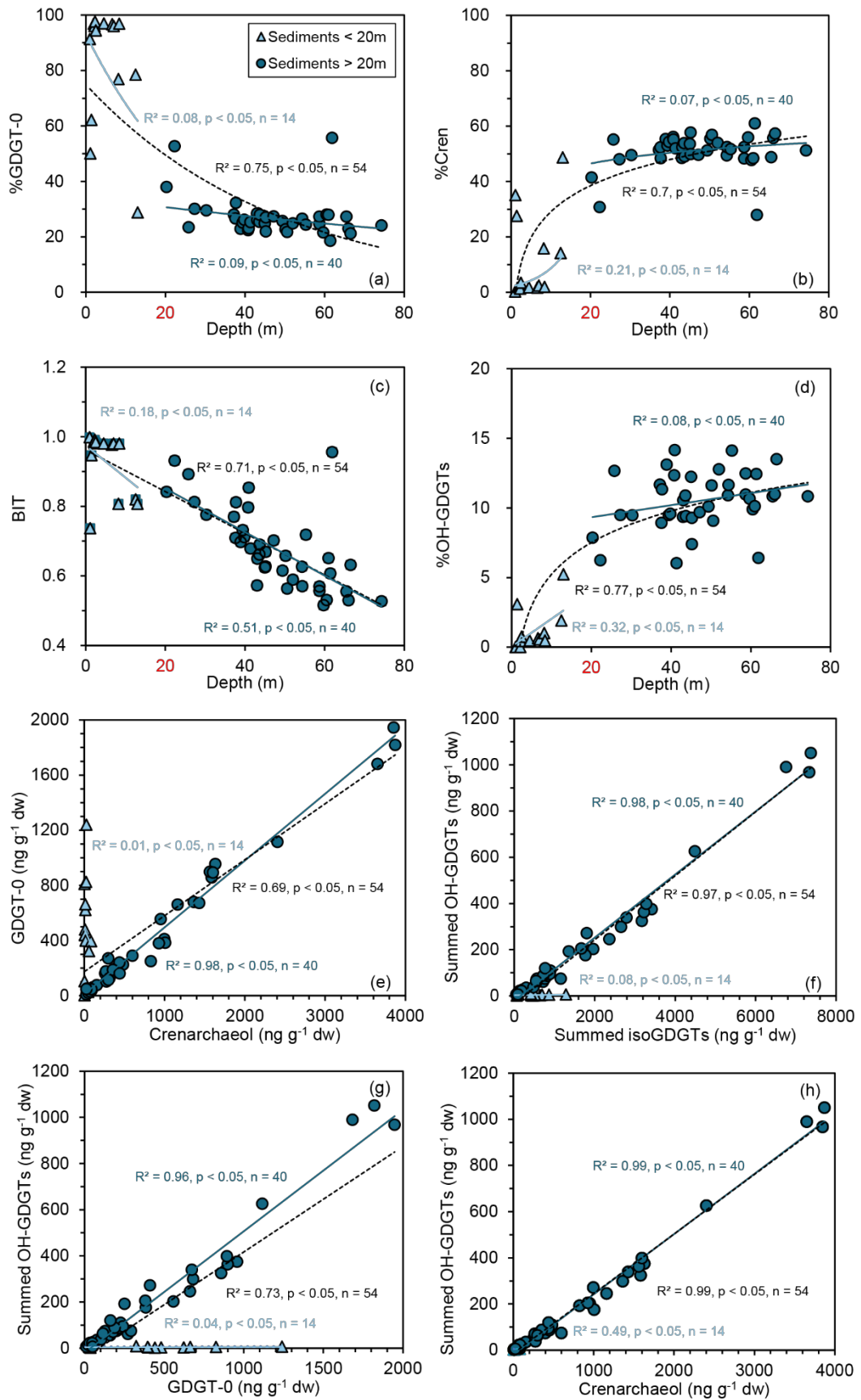


Fig. 5

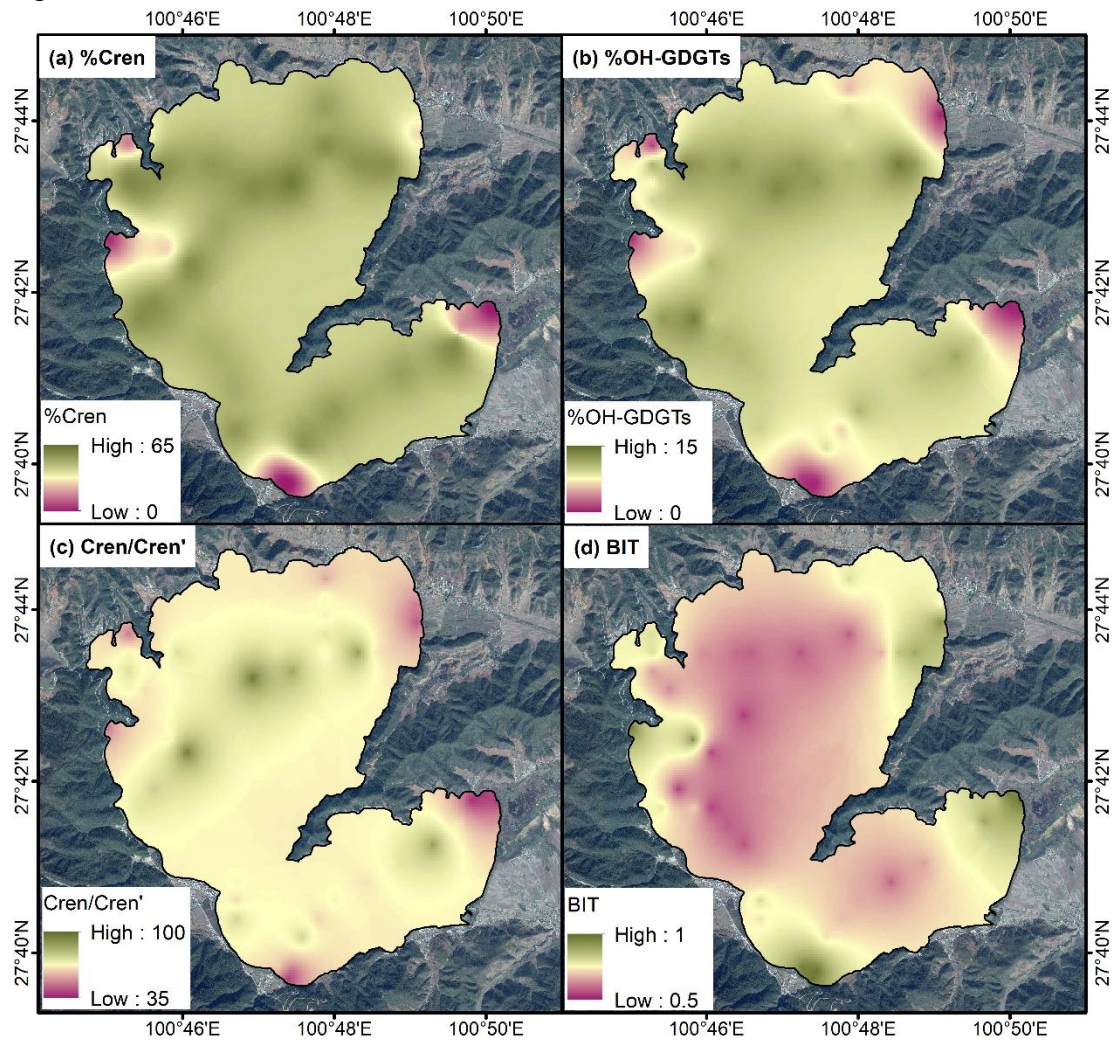


Fig. 6

