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1 **A diagnostic GDGT signature for the impact of hydrothermal activity on surface deposits**
2 **at the Southwest Indian Ridge**

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16
17 **Abstract**

18 The impact of hydrothermal activity on wider ocean geochemistry and microbial ecology
19 remains a topic of much interest. To explore whether hydrothermal microbial signatures are
20 exported to surrounding marine sediments or if such organisms serve as an important source of
21 sedimentary organic matter, we determined the distributions of glycerol dialkyl glycerol
22 tetraether (GDGT) membrane lipids in surficial normal marine sediments, metalliferous
23 sediments and low-temperature hydrothermal deposits at newly discovered hydrothermal fields
24 and adjacent areas at the Southwest Indian Ridge (SWIR). The GDGTs in those samples varied
25 significantly, evidently representing a variable influence of the hydrothermal activity. GDGT
26 compositions of surficial background sediments in SWIR were similar to those commonly
27 observed in marine sediments, dominated by GDGTs associated with marine planktonic archaea
28 and especially GDGT-0 and crenarchaeol. In contrast, the GDGTs of metalliferous sediments
29 strongly impacted by hydrothermal activity and low-temperature hydrothermal deposits were
30 markedly different, characterized by high relative abundances of isoprenoid GDGTs (*i*GDGTs)

31 bearing multiple rings (yielding a higher ring index), low relative abundances of crenarchaeol,
32 and the presence of glycerol monoalkyl glycerol tetraether lipids (GMGTs; so called ‘H-
33 tetraethers’) that were absent in the normal marine sediments. Sources for these hydrothermal-
34 specific tetraether lipids likely include methanogens and anaerobic methanotrophic archaea
35 (GDGT-0 and GDGT-1-3, respectively), *Thermoprotei* and *Thermoplasmatales* (elevated GDGT-
36 3-4), and other thermophilic archaea including *Methanobacteriales* (GMGTs). Deposits
37 influenced by low-temperature hydrothermal activity also contained higher abundances of
38 branched GDGTs (*brGDGTs*) typically attributed to soil bacteria. The more distal metalliferous
39 sediments influenced by the neutrally buoyant plume did not contain putative hydrothermal
40 GDGTs, having the same GDGT distribution as the background sediments. This suggests that the
41 neutrally buoyant plume has a limited potential to directly influence the organic matter inputs to
42 surrounding sediments, due to a rapidly waning chemosynthetic microbial contribution relative
43 to normal marine contributions as the plume dispersed and was diluted.

44

45 **Keywords:** tetraethers; organic matter; Southwest Indian Ridge; hydrothermal activity;
46 chemosynthetic microbial contribution

47

48 **1. Introduction**

49 In 1977, scientists diving in the submersible *Alvin* made a stunning discovery on the bottom
50 of the Galapagos Rift in the eastern Pacific Ocean, where seafloor hydrothermal activity and a
51 novel ecosystem were observed (Corliss et al., 1979). Since then, these discoveries have changed
52 our understanding of Earth and life, and 532 active and 56 inactive submarine hydrothermal vent
53 fields have been discovered (Beaulieu et al., 2013). The vent deposits are known to harbor high-

54 biomass benthic communities with chemosynthetic primary producers and other microbes
55 serving as the foundation of the food web (Govenar, 2012). They can use chemical energy,
56 derived from mixing of reduced chemicals such as CH₄, H₂S, H₂ and metals in hydrothermal
57 fluids with oxygenated seawater. Black smoker hydrothermal vents exude fluids with μM–mM
58 Fe concentrations (Von Damm et al., 1985; Douville et al., 2002), which can be as much as seven
59 orders of magnitude greater than typical deep ocean dissolved Fe of ~0.2–0.8 nM (Klunder et al.,
60 2011; Noble et al., 2012; Hatta et al., 2015). Low concentrations of the micronutrient iron in
61 seawater are known to limit primary production and nitrogen fixation in large regions of the
62 global ocean, but recent research demonstrates that dissolved iron from hydrothermal vents can
63 be transported thousands of kilometers from the venting site, contributing to the marine dissolved
64 iron inventory, especially in the abyssal ocean (Toner et al., 2012; Fitzsimmons et al., 2014). In
65 this and other areas, it remains vital to examine the impact of hydrothermal activity on the wider
66 ocean geochemistry and microbial ecology.

67 Diverse microbiological investigations, often involving culture-independent molecular
68 studies of 16S rRNA but also enrichment and isolation studies, have been used to examine
69 microbial diversity in deep-sea hydrothermal vent systems (e.g. Takai et al., 2001; Kormas et al.,
70 2006; Sogin et al., 2006; Jaeschke et al., 2012; Reeves et al., 2014). However, these approaches
71 have inherent limitations, such as inconsistent DNA recovery, kinetic biases inherent in
72 polymerase chain reaction, and the need to develop sepecific and appropriate primers and probes
73 (Chowdhury and Dick, 2012). Organic geochemical approaches also have limitations but can
74 offer a complementary view on microbial community structures since they do not require the
75 culturing of microorganisms, are quantitative and reproducible, and can integrate a longer time
76 window than nucleic acid based researches (Mrozik et al., 2014). There has been some research

77 on lipid biomarkers in hydrothermal fluids, sulfides, oxides, hydrothermally heated sediments
78 and organisms from the Mid-Atlantic Ridge, Arctic Mid-Ocean Ridge, East Pacific Rise,
79 Guyamas Basin spreading center and other hydrothermal systems (e.g. Schouten et al., 2003;
80 Phleger et al., 2005; Blumenberg et al., 2012; Hu et al., 2012; Jaeschke et al., 2012; Kellermann
81 et al., 2012; Méhay et al., 2013; Jaeschke et al., 2014; McCollom et al., 2015), including
82 investigations of intact polar lipids (IPLs) (Gibson et al., 2013; Reeves et al., 2014), and these
83 have helped reveal the structure and function of chemosynthetic systems. Since Archaea have
84 high growth temperatures, up to 121 °C (Kashefi and Lovley, 2003), and are widespread in
85 hydrothermal systems, there have been increasing investigations of archaeal membrane lipids in
86 submarine and terrestrial hydrothermal sites in recent years (e.g. Schouten et al., 2003; Pearson
87 et al., 2004, 2008; Pancost et al., 2005, 2006; Boyd et al., 2011; Kaur et al., 2011, 2015; Jaeschke
88 et al., 2012; Kellermann et al., 2012; Boyd et al., 2013; Gibson et al., 2013; Lincoln et al., 2013;
89 Méhay et al., 2013; Jaeschke et al., 2014; Reeves et al., 2014).

90 Here we survey the glycerol dialkyl (and monoalkyl) glycerol tetraether (GDGT and GMGT)
91 membrane lipid distributions (Fig. 1) at the Southwest Indian Ridge, an area where little work
92 has been done using either microbiological or organic geochemical approaches (see below). We
93 examined a combination of hydrothermal deposits and metalliferous (plume) deposits, and used
94 these to obtain a lipid profile and insights into the archaeal community in the hydrothermal field.
95 This allowed us to test whether hydrothermal activity impacted the organic matter (OM)
96 composition of surrounding surface sediments.

97

98 **2. Samples and methods**

99 *2.1. Study area and samples*

100 The SWIR is the ultraslow spreading part of the Indian ridge and the sole modern migration
101 pathway between the diverse vent fauna of the Atlantic and Pacific oceans (German et al., 1998;
102 Zhou and Dick, 2013). It is an area of interest, therefore, with respect to the characterization,
103 distribution and migration of submarine microbes, and potentially for the discovery of new deep
104 sea communities (Rogers et al., 2012; Tao et al., 2012; Amon et al., 2015; Chen et al., 2015a,
105 2015b, 2015c, 2015d). Molecular biological (Peng et al., 2011; Li et al., 2013; Li et al., 2015),
106 element geochemistry and mineralogical (Tao et al., 2011, 2012; Cao et al., 2012) studies have
107 been conducted in the SWIR hydrothermal field, but studies of lipid biomarkers and related
108 biogeochemical processes are rare and focused on hydrocarbons and fatty acids in hydrothermal
109 barnacle shells and sulfides (Huang et al., 2014; Lei et al., 2015).

110 The samples described in this paper were recovered from Dragon Vent Field (49°39' E,
111 37°47' S), a nearby inactive field (50°28' E, 37°39.50' S) and surrounding areas during the
112 DY115-20 and DY115-21 expeditions of R/V Da Yang Yihao in 2009 and 2010 (Fig. 2). Dragon
113 Vent Field, the first active hydrothermal vent to be discovered in the SWIR was found using a
114 remotely operated vehicle from Woods Hole Oceanographic Institution (Tao et al., 2007), at a
115 depth of 2760 m. It harbors many active and inactive sulfide chimneys with mussels, barnacles,
116 sea cucumbers and gastropods (Copley, 2011; Rogers et al., 2012; Tao et al., 2012; Cole et al.,
117 2014). Abundant bivalve and gastropod shells were also observed at the inactive field (~ 200 ×
118 125 m in extent, approximately 73 km away from Dragon Vent Field, at a depth of 1770 m) (Tao
119 et al., 2012).

120 All samples were collected by television grab and divided into three categories (see
121 Supporting Information Table S1) according to the results of mineral and element geochemistry
122 (Pan, 2015): (1) background sediments containing abundant foraminifera detritus, apparently

123 uninfluenced by hydrothermal activity; (2) three metalliferous sediments influenced by various
124 degrees of hydrothermal activity depending on the distance from the hydrothermal vent; and (3)
125 low-temperature hydrothermal deposits enriched in Fe and/or Si, noting that even though these
126 are 'low-temperature' hydrothermal deposits, precipitation temperatures are greater than
127 background sediments of SWIR, ranging between 38.3 to 81.8 °C based on the deduction of
128 oxygen isotopic compositions of amorphous silica in low-temperature hydrothermal deposits
129 from SWIR (Li et al., 2013). Typical samples from the same studied sites (SW35, SW33 and
130 SW36) have been analysed for molecular biology (Peng et al., 2011; Li et al., 2013).

131 The three metalliferous sediments can be classified on the basis of their mineral and element
132 compositions. M-T1 sediments, the furthest from the hydrothermal field, with neutrally buoyant
133 plume fall-outs mixed in, have abundant calcite and slightly higher contents of Fe, Cu and Zn
134 than background sediments. M-T2 sediments, with some oxides mixed in, mainly nontronite and
135 two-line-ferrihydrite, have relatively higher contents of Fe, Mn, Cu and Zn than M-T1, mainly
136 impacting by low-temperature hydrothermal activity. M-T3 sediments have highest Cu and Zn
137 contents, and abundant goethite, representing a direct influence from high-temperature
138 hydrothermal activity (Dias et al., 2008).

139

140 *2.2. Bulk organic parameters*

141 Total carbon (TC) and inorganic carbon (IC) were determined using a Carlo Erba EA1108
142 Elemental Analyzer and a modified Coulomat 702 analyzer, respectively. Total organic carbon
143 (TOC) concentrations were determined by the difference between TC and IC. All reported TOC
144 values were the means of duplicate measurements. Carbon isotopic compositions of TOC
145 ($\delta^{13}\text{C}_{\text{TOC}}$) were obtained after pretreatment with 4 mol/L HCl with a Flash EA 1112 HT-Delta V

146 Advantage (Thermo Company). The $\delta^{13}\text{C}_{\text{TOC}}$ [‰ Vienna Pee Dee Belemnite, VPDB] error was
147 $\pm 0.2\text{‰}$.

148

149 *2.3. Lipid analysis*

150 Two methods were used successively to extract and separate fractions in samples of
151 different types. Metalliferous sediments (M-T2 and M-T3) and low-temperature hydrothermal
152 deposits were processed with method 1, using a modified Bligh-Dyer extraction (Bligh and Dyer,
153 1959) and fractionation protocol based on Dickson et al. (2009). After freeze-drying, about 15 g
154 of each sample were solvent extracted with a culture tube using a single-phase mixture
155 comprised of methanol, chloroform, and aqueous 50 mM phosphate buffer water (pH 7.4) in the
156 volume ratio of 2:1:0.8 (6×). The phases were separated by addition of chloroform and buffer
157 water. The organic phase containing the lipids was collected, and activated copper turnings were
158 added to the extracts for 24 hours to remove elemental sulfur. An aliquot of the total extract was
159 separated into three fractions on a silica column. Fractionation was achieved with
160 chloroform:acetic acid (100:1, v:v), acetone and methanol as eluents to recover simple core
161 lipids (CL), glycolipids (GL) and phospholipids (PL), respectively. The CL fraction was
162 subsequently eluted through a silica column with chloroform saturated with ammonium
163 hydroxide and chloroform:acetic acid (100:1, v:v) to separate neutral components and free fatty
164 acids. GDGTs were not detected in neutral components of the CL fraction based on method 1 and
165 it appears that they were eluted in the GL fraction; previous workers have observed similar
166 behavior and it seems that this method is more appropriate for bacterial membrane lipids (Pitcher
167 et al., 2009).

168 Method 2 was used for the background sediments and M-T1. About 15 g sediments were

169 ultrasonically extracted three times with methanol, dichloromethane:methanol (1:1, v:v) and
170 dichloromethane, respectively (Schouten et al., 2002). The total lipid extract was subsequently
171 separated using a fractionation protocol derived from Oba et al. (2006) and Pitcher et al. (2009),
172 using a silica column and eluting with hexane:ethyl acetate (3:1, v:v), ethyl acetate and methanol
173 to yield CL, GL and PL fractions, respectively.

174 For both methods, to remove polar head groups, 5% HCl in methanol was added to the GL
175 and PL fractions which were then heated at 100 °C for 3 h, after which the organic phase was
176 extracted with double distilled water and chloroform. Because the CL fraction was eluted in the
177 GL fraction in method 1, we have combined GDGT abundances and distributions in the CL and
178 GL fractions for all samples, and discuss only summed CL+GL distributions in order to make an
179 effective comparison among sediments of different types. We also note that silica gel column
180 chromatography separations used in this study are always associated with significant losses of
181 IPL-GDGTs (Lengger et al., 2012); because this likely differs among methodologies, it is
182 inappropriate to compare concentrations even within the constraints of this study – and certainly
183 with other studies. Therefore, this paper focuses solely the distributions of tetraether lipids and
184 how they differ among sediments of different types. This approach should be robust as several
185 studies have shown that although concentrations are methodologically dependent, tetraether lipid
186 distributions are consistent (e.g. Schouten et al., 2009, 2013a). To test this further, we processed
187 one sample with both methods and distributions of GDGTs and GMGTs were similar
188 (see Supporting Information Table S3).

189 Due to the different methods, we do not discuss the potentially fossil vs living signals (core
190 vs intact polar GDGTs). However, we do note that phospholipid distributions (data shown in
191 Supporting Information Table S4) generally show the same relationships among different

192 sediment types as observed for the combined CL+GL fractions discussed below. This is likely a
193 fruitful avenue of future research.

194 Aliquots of all fractions were analysed by high performance liquid chromatography/
195 atmospheric pressure chemical ionisation-MS (HPLC/APCI-MS, Agilent 1100 series) equipped
196 with an autoinjector and Chemstation software (Agilent) in a modification of the procedure of
197 Hopmans et al. (2000) and then Schouten et al. (2007a). Fractions were dissolved in
198 hexane:isopropanol (99:1, v/v) and filtered through 0.45 µm mesh PTFE. Separation of GDGTs
199 was achieved on an Alltech Prevail Cyano column (2.1 mm i.d. × 150 mm, 3 µm) maintained at
200 30 °C with a flow rate of 0.2 ml/min. Injection volume was 20 µl. GDGTs were eluted
201 isocratically with 99% hexane and 1% isopropanol for 7 min, followed by a linear gradient to
202 1.3% isopropanol at 30 min, to 1.6% isopropanol at 35 min, then increasing to 10% isopropanol
203 at 36 min and kept for 8 min, finally equilibrating with 1% isopropanol for 13 min before the
204 next injection. After each two analyses, the column was cleaned by back-flushing
205 hexane:isopropanol 99:1 (v:v) for 7 min and then rinsed by a linear gradient from 90:10 (v:v)
206 hexane:isopropanol to 99:1 (v:v) hexane:isopropanol within 14 min and equilibrated with 1%
207 isopropanol at 30 min. Conditions for APCI-MS were as follows: vaporizer temperature 380 °C,
208 drying gas (N₂) flow 6 l/min and temperature 200 °C, capillary temperature 282 °C, corona
209 discharge current 3 µA. GDGTs were detected in selected ion monitoring (SIM) mode and were
210 semi-quantified by an internal synthetic C₄₆ tetraether standard, based on the procedure of
211 Huguet et al. (2006) and Schouten et al. (2007a).

212 The branched isoprenoid tetraether (BIT) index, a proxy for terrestrial OM input, was used
213 as defined by Hopmans et al. (2004):

$$214 \quad BIT = \frac{[GDGT - I] + [GDGT - II] + [GDGT - III]}{[GDGT - I] + [GDGT - II] + [GDGT - III] + [Cren]} \quad (1)$$

215 , where numbers refer to individual GDGT structures shown in Figure 1. The methylation of
 216 branched tetraether (MBT) and cyclization of branched tetraethers (CBT) ratios were used as
 217 defined by Weijers et al. (2007):

$$218 \quad MBT = \frac{([GDGT - I] + [GDGT - Ia] + [GDGT - Ib])}{\Sigma[\textit{all branched GDGTs}]} \quad (2)$$

$$220 \quad CBT = -\log\left(\frac{[GDGT - Ia] + [GDGT - IIa]}{[GDGT - Ia] + [GDGT - IIa]}\right) \quad (3)$$

221
 222
 223 The ring index (RI) was defined based on Pearson et al. (2004):

$$224 \quad RI = \frac{1 \times [GDGT - 1] + 2 \times [GDGT - 2] + 3 \times [GDGT - 3] + 4 \times [GDGT - 4]}{[GDGT - 1] + [GDGT - 2] + [GDGT - 3] + [GDGT - 4]} \quad (4)$$

225
 226 The methane index (MI) was used as defined by Zhang et al. (2011):

$$227 \quad MI = \frac{[GDGT - 1] + [GDGT - 2] + [GDGT - 3]}{[GDGT - 1] + [GDGT - 2] + [GDGT - 3] + [Cren'] + [Cren'']} \quad (5)$$

228
 229 The tetraether index of tetraethers consisting of 86 carbons (TEX₈₆) was used as defined by
 230 Schouten et al. (2002):

$$231 \quad TEX_{86} = \frac{[GDGT - 2] + [GDGT - 3] + [Cren']}{[GDGT - 1] + [GDGT - 2] + [GDGT - 3] + [Cren']} \quad (6).$$

232
 233 **3. Results**

234 *3.1. Total organic carbon and carbon isotopic composition*

235 The TOC contents and $\delta^{13}C_{TOC}$ values varied among the different sediment types

236 (Supporting Information Table S1 and Fig. 3). Background sediments had the highest TOC
237 contents (1.2% average) and relatively heavy $\delta^{13}\text{C}_{\text{TOC}}$ values (-22.1‰ average). Low-
238 temperature hydrothermal deposits had the lowest TOC contents (0.13% average) and more
239 depleted $\delta^{13}\text{C}_{\text{TOC}}$ values (-24.8‰ average). In the metalliferous sediments, TOC contents and
240 $\delta^{13}\text{C}_{\text{TOC}}$ values in M-T1 were similar to the background sediments, whereas those parameters in
241 M-T2 were close to those of the low-temperature hydrothermal deposits. TOC contents were
242 slightly higher in M-T3 than M-T2 and low-temperature hydrothermal deposits, but similar
243 $\delta^{13}\text{C}_{\text{TOC}}$ values.

244

245 3.2. Tetraether lipid distributions

246 GDGTs observed in the SWIR samples include a range of isoprenoidal GDGTs (*i*GDGTs)
247 bearing 0 to 4 cyclopentyl moieties as well as crenarchaeol and its regioisomer; a suite of the
248 unusual “H-shaped” glycerol monoalkyl glycerol tetraethers (GMGTs, up to four cyclopentyl
249 moieties but mainly GMGT-0); and surprisingly, branched GDGTs, often in high abundances
250 (*br*GDGTs, including GDGT III-IIIb, GDGT II-IIb, GDGT I-Ib). Trace amounts of glycerol
251 trialkyl glycerol tetraether (GTGT-0; i.e. with one biphytanyl and two phytanyl components and
252 no cyclopentyl moieties) were also detected in the low-temperature hydrothermal deposits
253 (Supporting Information Table S2).

254 Among the *i*GDGTs, GDGT-0 and crenarchaeol were dominant in background sediments
255 and M-T1, with crenarchaeol percentages typically being 40-50%. Proportions of GDGTs 1-3
256 were lower and distributions were overall similar to other marine sediments (Schouten et al.,
257 2013b). The proportion of crenarchaeol was markedly lower in M-T2, M-T3 and low-
258 temperature hydrothermal deposits, largely due to higher proportions of GDGTs 0-3 but also

259 GDGT-4, which was not detected in the background sediments. The *i*GDGT distributions of M-
260 T3 was dominated by GDGT-0 and GDGT-4 (Supporting Information Table S2).

261 The isoprenoidal GMGTs (*i*GMGTs) were not detected in background sediments nor in M-
262 T1, but had high relative abundances in the low-temperature hydrothermal deposits and the
263 metalliferous sediments strongly impacted by hydrothermal activity and close to the
264 hydrothermal vent; in fact *i*GMGTs represent >45% of the total tetraether lipids in M-T3, in
265 which GMGT-0 and GMGT-4 were the main components. The proportions of *i*GMGTs in M-T2
266 and low-temperature hydrothermal deposits were lower, but still higher than those of the other
267 sediments, and dominated by GMGT-0.

268 The proportions and abundances of *br*GDGTs were very low in most background samples,
269 M-T1 and M-T3, but markedly higher in M-T2 and low-temperature hydrothermal deposits
270 (>15%). This yielded higher BIT indices for the latter – 0.24 to 0.69 in M-T2 and hydrothermal
271 deposits compared to <0.10 in the other sediment types (Table 1). GDGT-III, GDGT-II and IIa
272 were dominant compounds in most samples. However, GDGT-I was also predominant in M-T2
273 and low-temperature hydrothermal deposits, such that CBT and MBT indices were larger (Table
274 1).

275 In summary, tetraether lipid distributions were dominated by *i*GDGTs in all three SWIR
276 sample categories (Table 1), but the percentages of *i*GDGTs were relatively lower in M-T2, M-
277 T3 and low-temperature hydrothermal deposits, whereas *i*GMGTs were relatively more abundant.
278 Branched GDGTs were also proportionally more abundant in the low-temperature hydrothermal
279 deposits and M-T2. This results in three main groups, shown in the ternary diagram of *i*GDGTs,
280 *i*GMGTs and *br*GDGTs (Fig. 4): Group 1 comprises background sediments and M-T1; Group 2,
281 characterized by relatively higher percentages of *br*GDGTs, comprises some M-T2 and low-

282 temperature hydrothermal deposits; and Group 3, characterized by relatively higher percentages
283 of *i*GMGTs, comprises the other M-T2 and low-temperature hydrothermal deposits as well as M-
284 T3.

285

286 **4. Discussion – Variations in Organic Matter Sources**

287 TOC contents in SWIR background sediments were higher than the global average for deep-
288 sea surficial sediments (0.25~0.50%, Premuzic et al., 1982). This could be associated with
289 elevated concentrations of phytoplankton and zooplankton in the study area, which has been
290 identified as an important carbon sequestration region (Froneman et al., 1998; Llido et al., 2005).
291 $\delta^{13}\text{C}_{\text{TOC}}$ values of most background sediments are in the typical range of marine organic matter
292 (-22‰ to -19‰, Fontugne and Jouanneau, 1987), indicating that organic matter of SWIR
293 background sediments was mainly derived from autochthonous marine organisms. This is also
294 consistent with the presence of lipid biomarkers for phytoplankton (i.e. sterols and alkenones,
295 data not shown). The $\delta^{13}\text{C}$ values of the metalliferous sediments (except M-T1) and most of the
296 low-temperature hydrothermal deposits were lower, consistent with relatively low $\delta^{13}\text{C}$ values
297 for organic matter in other hydrothermal settings (e.g. Southern Mariana Trough, Kato et al.,
298 2010; Loki's Castle, Jaeschke et al., 2012, 2014; PACMANUS, Reeves et al., 2014); this is
299 typically attributed to the production of ^{13}C -depleted OM by chemosynthetic organisms.
300 Intriguingly M-T1 TOC has a $\delta^{13}\text{C}$ value similar to that of background sediments, suggesting that
301 deposition of neutrally buoyant plume material has not imparted an obvious hydrothermal OM
302 signature to the areas far away from the hydrothermal vents.

303 To explore these differences in OM source and microbial ecology further we have examined
304 the tetraether lipids of surface sediments from SWIR. Tetraether lipid distributions differ among

305 the normal marine sediments, metalliferous sediments and the low-temperature hydrothermal
306 deposits. Based on the relative distributions of *i*GDGTs, *i*GMGTs and *br*GDGTs, background
307 sediments but also M-T1 have OM sources typical of deep marine sediments, primarily GDGTs
308 exported from overlying waters (Group 1). M-T2 and M-T3 distributions are similar to low-
309 temperature hydrothermal deposits but can still be divided into two sub-groups: Group 2
310 with >11% *br*GDGTs and Group 3 with >10% *i*GMGTs.

311 Combined with the previous molecular biological analyses of samples from the same
312 studied sites at SWIR (Peng et al., 2011; Li et al., 2013), likely sources of different tetraether
313 lipid classes can be assigned. The GDGT-0/crenarchaeol, MIs and RIs all indicate that the
314 hydrothermal deposits have additional archaeal sources compared with background sediments.
315 Moreover, the percentages of *br*GDGTs and BIT indices, combined with unusual MBT/CBT
316 ratios, appear to reflect in situ bacteria production.

317

318 *4.1. Isoprenoidal GDGTs*

319 The *i*GDGTs in background sediments and M-T1 of SWIR were dominated by GDGT-0 and
320 crenarchaeol and lower contents of *i*GDGTs 1-3. This is similar to distributions in the surficial
321 sediments from other oceans and indicates a major contribution to tetraether membrane lipids
322 from non-thermophilic Thaumarchaeota in the marine environment (Schouten et al., 2002).
323 TEX₈₆ values in these samples were also consistent with SWIR sea surface temperatures (SSTs,
324 <http://www.ospo.noaa.gov/data/sst/contour/global.c.gif>) and global calibrations (Fig. 5a; Kim et
325 al., 2010), suggesting a predominantly allochthonous source of water-column GDGTs, which has
326 been confirmed by recent 16s rRNA analysis (mainly Thaumarchaeota, unpublished data).

327 The compositions of *i*GDGTs in most metalliferous sediments and low-temperature

328 hydrothermal deposits, with higher contents of total *i*GDGTs 0-4 and lower contents of
329 crenarchaeol, were different from background sediments. Some of this could be attributed to the
330 presence of crenarchaeota *Thermoprotei* and euryarchaeota *Thermoplasmatales*, previously
331 documented for some of these samples (Peng et al., 2011; Li et al., 2013) and known sources of
332 *i*GDGTs 1-4 (reviewed in Pearson and Ingalls, 2013; Schouten et al., 2013b).

333 Additional contributions of non-pelagic archaea to the isoprenoidal GDGT pool can be
334 ascertained by testing expected TEX₈₆ indices against the RI. Group 2 and 3 sediments are both
335 associated with high RIs, but RIs are higher for the latter and Group 2 TEX₈₆ values are similar
336 (albeit at the high end) of the background sediment range. Therefore, it appears that *i*GDGTs in
337 Group 3 sediments – and perhaps Group 2 sediments – derive from sources additional to those
338 that dominate normal marine sediments, presumably hydrothermal organisms.

339 Additionally, the *i*GDGT distributions could primarily reflect the different environmental
340 conditions under which Group 2 and 3 sediments formed. TEX₈₆ indices in marine sediments
341 have a positive correlation with sea surface temperatures in overlying waters but generally not
342 with pH (e.g. Schouten et al., 2002; Kim et al., 2008, 2010; Boyd et al., 2011). Other studies
343 have also shown that temperature is an important control on the distribution of archaeal tetraether
344 membrane lipids, with RIs increasing with growth temperature (Pearson et al., 2004; Uda et al.,
345 2004; Elling et al., 2015; Kaur et al., 2015). Similar studies, however, have shown that pH also
346 governs GDGT distributions in thermophilic archaea, with RI increasing as pH decreases in
347 diverse settings (e.g. Boyd et al., 2011, 2013; Wu et al., 2013; Kaur et al., 2015).

348 To date, the highest temperature vent fluid observed in the Dragon vent field is 379 °C
349 (unpublished data). Previous studies showed the pH_(in situ) of the highest temperature vent fluid (>
350 380 °C) measured in situ with solid-state electrochemical sensors, is slightly acidic (5.1–5.4).

351 However, mixing of seawater with vent fluid results in seawater dominated conditions with
352 attendant pH increases (Ding et al., 2005), such that below 121 °C, the upper temperature limit
353 for life (Kashefi and Lovley, 2003), $\text{pH}_{(\text{in situ})}$ is usually greater than 6.0, approaching neutrality
354 (Ding et al., 2005). It is unclear if a pH range from about 6 to 8 can explain the large variations
355 in GDGT distributions observed here due to the lack of in situ pH information. Pure culture
356 study of marine planktonic thaumarchaeal isolates demonstrated that pH variations over a range
357 of 7.3 to 7.9 exerted a minor influence on GDGT cyclization, however, pH might influence
358 environmental GDGT distribution indirectly by selecting for specific thaumarchaeal lineages
359 with distinct lipid compositions (Elling et al., 2015). Moreover, pH has been shown to be a
360 control on RIs in other settings, albeit over a larger range. Therefore, the unusual *i*GDGT
361 distributions in Group 2 and especially Group 3 likely reflect a range of ecological but also
362 environmental factors, primarily dictated by temperature but possibly also related to pH
363 variations.

364 Both GDGT-0 and crenarchaeol occur in marine group 1 Crenarchaeota, but only GDGT-0
365 appears to be produced by methanogens (Schouten et al., 2007b). The GDGT-0/crenarchaeol
366 ratio in marine group 1 Crenarchaeota typically varies between 0.2 and 2 (Schouten et al., 2002)
367 and ratios >2 suggest an additional source for GDGT-0 (Blaga et al., 2009). The GDGT-
368 0/crenarchaeol ratios were <2 in background sediments and M-T1 (Group 1), whereas the ratios
369 in some metalliferous sediments and low-temperature hydrothermal deposits (Group 3) were > 2
370 (Fig. 5b). Unlike RIs, greater abundances of GDGT-0 are difficult to ascribe to higher
371 temperatures or lower pH and we instead suggest that this is evidence for an additional source,
372 possibly methanogenic archaea. Abundant methanogens (mainly Methanosarcinaceae or
373 Methanobacteriales) have been detected in some hydrothermal deposits at the same SWIR sites

374 (Peng et al., 2011; Li et al., 2013). However, other archaeal species cannot be excluded as
375 contributors (e.g. Pearson et al., 2013; Schouten et al., 2013b; Villanueva et al., 2014) and we
376 note that Archaeoglobales was also detected in these settings (Peng et al., 2011).

377 The GDGT-0/crenarchaeol ratios were <2 in Group 2, which could suggest a hydrothermal
378 origin for both compounds, and crenarchaeol has been found in both terrigenous (e.g. Pearson et
379 al., 2004; Schouten et al., 2007b; Pitcher et al., 2011) and marine hydrothermal systems (Méhay
380 et al., 2013); alternatively, it is consistent with a smaller hydrothermal overprint of Group 2
381 samples (compared to Group 3), consistent with the lower RIs and TEX_{86} values.

382 GDGT MIs below 0.3 to 0.5 are typical of normal marine sediments, whereas MIs of
383 sediments impacted by additional microbial inputs, including anaerobic methanotrophs, are
384 typically >0.5 (Pancost et al., 2001a; Zhang et al., 2011). The MI of most samples of Groups 1
385 and 2 were <0.3 , with several background sediments in the range of 0.3~0.5; in contrast, the MIs
386 of most Group 3 samples were >0.6 (Fig. 5b), indicating an additional contribution. As
387 hydrothermal plume samples collected from Dragon Vent Field have higher CH_4 contents than
388 background water, by at least one order of magnitude (Wang et al., 2015), it seems likely that
389 archaea involved in methane production and consumption could have contributed to the *i*GDGT
390 signature of Group 3, affecting both MIs and %GDGT-0. However, the elevated MIs can also be
391 explained by environmental impacts on GDGT distributions, as discussed above.

392 To explore these two options, we have examined other biomarker classes. Archaeol has
393 always been found in association with anaerobic methane-oxidising Archaea (Blumenberg et al.,
394 2004) and it was found here (1.6–120 ng/g sediment) and was indeed more abundant in the
395 Group 3 sediments (Pan, 2015). Intriguingly, the samples with high MIs, RIs and GDGT-
396 0/crenarchaeol ratios also contained abundant non-isoprenoidal dialkylglycerol ethers (DAGEs)

397 and *iso/anteiso*C_{15:0} and *iso/anteiso*C_{17:0} fatty acids (Pan, 2015), potentially derived from
398 sulphate-reducing bacteria (Hinrichs et al., 2000; Pancost et al., 2001b). These could reflect
399 independent bacterial inputs, or organisms syntrophically associated with archaeal
400 methanotrophs. However, many thermophilic bacteria synthesize abundant DAGEs and *iso*- and
401 *anteiso*-branched fatty acids (e.g. Huber et al., 1992; Sturt et al., 2004; Yang et al., 2006;
402 Schubotz et al., 2013; Reeves et al., 2014). Moreover, many Archaea (including methanogens,
403 halophiles, Marine Benthic Group B and Miscellaneous Crenarchaeotic Group) produce archaeol
404 (e.g. Koga and Morii, 2005; Lipp and Hinrichs, 2009). In fact, many of these compounds have
405 been detected in hydrothermal deposits that appear to have no AOM influence (e.g. Bradley et al.,
406 2009; Kaur et al., 2011, 2015). As such, the co-occurrence of these biomarkers with high MIs
407 could be evidence for an AOM influence, but that evidence is weak and other explanations
408 remain possible. Compound-specific stable carbon isotope analysis could resolve these
409 competing hypotheses (e.g. Pancost et al., 2001a; Elvert et al., 2005; Niemann and Elvert, 2008),
410 but that was not possible due to their low abundances.

411

412 4.2. *Isoprenoidal GMGTs*

413 Isoprenoidal GMGTs were only found in some metalliferous sediments and low-
414 temperature hydrothermal deposits from the SWIR, being absent in Group 1 sediments and
415 occurring in relatively low abundances in Group 2 (mainly GMGT-0). Abundances were much
416 higher in Group 3 (especially SW40 in M-T3, enriched in copper and zinc, and strongly
417 influenced by high-temperature hydrothermal activity), and dominated by both GMGT-0 and
418 GMGT-4.

419 GMGTs have been found in many marine sediments, albeit at generally low abundances

420 (Schouten et al., 2008); they appear to be particularly abundant in hydrothermal settings,
421 including Loki's Castle (Jaeschke et al., 2012, 2014), Lost City (Lincoln et al., 2013; Méhay et
422 al., 2013) and PACMANUS deep-sea hydrothermal fields (Reeves et al., 2014). They have been
423 reported in *Methanobacteriales* (*Methanothermus fervidus*, Morii et al., 1998), *Thermococcales*
424 (*T. celer*, *P. horikoshii*, Sugai et al., 2004; Jaeschke et al., 2012), DHVE2-cluster
425 (*Aciduliprofundum boonei*, Reysenbach et al., 2006) and *Desulforococcales* (*Ignisphaera*
426 *aggregans*, Knappy et al., 2011). Because a thermophilic crenarchaeon and *Methanobacteriales*
427 were previously detected in some of these samples (Peng et al., 2011; Li et al., 2013), we
428 speculate that GMGTs (and maybe GTGT-0) in the metalliferous sediments and low-temperature
429 hydrothermal deposits mainly originated from these archaea (Schouten et al., 2013b).

430 Additionally, there is a strong positive correlation between RI and percentages of GMGTs
431 (of total tetraethers) in the metalliferous sediments strongly influenced by hydrothermal activity
432 and low-temperature hydrothermal deposits from the SWIR (Fig. 6). A global synthesis shows
433 that this correlation is generally widespread and somewhat consistent across a range of
434 hydrothermal settings (Fig. 6; data from Jaeschke et al., 2012; Lincoln et al., 2013; Jaeschke et
435 al., 2014; Reeves et al., 2014). Given the strong positive linear relationship between RI and
436 temperature (Elling et al., 2015; Kaur et al., 2015), it seems likely temperature also governs the
437 relative abundances of GMGTs – although again, we cannot entirely preclude a pH control.
438 Regardless of the direct control, it seems clear that %GMGTs also are indicative of hydrothermal
439 input in the SWIR, similar to what has been suggested for Lost City Hydrothermal Field (Lincoln
440 et al., 2013).

441

442 4.3. Branched GDGTs of Putative Bacterial Origin

443 At the SWIR, proportions of *br*GDGTs are low (<10%) for most background sediments, M-
444 T1, M-T3 and some low-temperature hydrothermal deposits (mainly Group 1 and 3); similarly,
445 BIT indices were generally less than 0.10. These observations are consistent with those from
446 previous studies of open marine sediments, wherein *br*GDGT generally comprise less than 10%
447 of total GDGTs (Schouten et al., 2013b). Moreover, the low percentages of *br*GDGTs in M-T3,
448 which has been influenced by high-temperature hydrothermal activity, are similar to high-
449 temperature hydrothermal sulfides from other areas (Jaeschke et al., 2012; Reeves et al., 2014).
450 However, higher relative amounts of *br*GDGTs occurred in some background marine sediments
451 (e.g. SW12, SW21) of Group 1, the hydrothermally impacted metalliferous sediments (e.g.
452 SW32) and low-temperature hydrothermal deposits (e.g. SW33, SW36, SW41) of Group 2.
453 Distributions also differed, with GDGT-I being dominant over GDGT-III, GDGT-II and GDGT-
454 IIa in Group 1 and 3 but less so in Group 2; similarly, CBT and MBT indices were relatively
455 higher in Group 2 than in Group 1 and 3.

456 Although *br*GDGTs are mainly considered to be products of heterotrophic anaerobic
457 bacteria in terrigenous soil (Weijers et al., 2009), a thermophile source has been inferred for
458 some terrestrial hot springs (Hedlund et al., 2013; Zhang et al., 2013). Furthermore, it seems
459 likely that the terrigenous contribution to the organic matter in the study area, located at an ocean
460 ridge >2000 km away from the nearest mainland, was minor; this is consistent with minor inputs
461 of Al and Ti (terrigenous indicators) and very low abundances of leaf wax biomarkers (high-
462 molecular-weight fatty acids and alkanols, Pan, 2015). Instead, we suggest that the *br*GDGTs in
463 all samples but especially where proportions exceed 10% derive from in situ bacterial production;
464 this could include Acidobacteria which are abundant in some Group 2 samples (SW33 and SW36;
465 Li et al., 2013). Recent studies have shown that *br*GDGTs are synthesized by bacteria in marine

466 sediments (Peterse et al., 2009; Zhu et al., 2011), hydrothermal systems (Hu et al., 2012; Lincoln
467 et al., 2013) and shelf systems (Sinninghe Damsté, 2016). However, it remains unclear why
468 particularly high proportions are largely restricted to Group 2 sediments in this setting.

469

470 *5. Synthesis*

471 At submarine hydrothermal vents, microorganisms thrive on inorganic energy sources, such
472 as methane, reduced iron and manganese that are abundant in hydrothermal vent fluids
473 (Tagliabue et al., 2010; Breier et al., 2012). These inorganic elements are dispersed more widely
474 by hydrothermal plumes rising hundreds of metres off the seafloor and traveling thousands of
475 kilometres from the vents (Dick and Tebo, 2010; Toner et al., 2012; Fitzsimmons et al., 2014).
476 The abundance of chemosynthetic microorganisms within hydrothermal plumes makes such
477 plumes an important dispersal mechanism and a significant source of organic matter to the deep
478 ocean (McCollum, 2000; Lam et al., 2004, 2008). Moreover, these microorganisms appear to be
479 active in plumes and partially determine the geochemical fate of these hydrothermal inputs
480 (Lilley et al., 1995).

481 Previous work has confirmed that species richness and phylogenetic diversity is typically
482 highest near the vent orifice, with the abundance of chemosynthetic microorganisms decreasing
483 with increasing distance from the vent (Sheik et al., 2015). This is consistent with our analyses.
484 The M-T2 and M-T3 sites, with high Fe, Mn, Cu and Zn contents and in close proximity to the
485 vent, have GDGT distributions similar to those of hydrothermal deposits and distinct from
486 background sediments. Overall, the hydrothermal GDGT signature was consistent with previous
487 work and the expected influence of higher growth temperature, including high Ring Indices,
488 TEX₈₆ values and %GMGT. Other features, including high %GDGT-0, appear to be consistent

489 with an active methane cycle in these sites. Crucially, the more distal M-T1 sediments have
490 GDGT distributions largely indistinguishable from background sediments, suggesting a rapidly
491 waning chemosynthetic contribution relative to normal marine contributions as the plume
492 dispersed and was diluted.

493

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504

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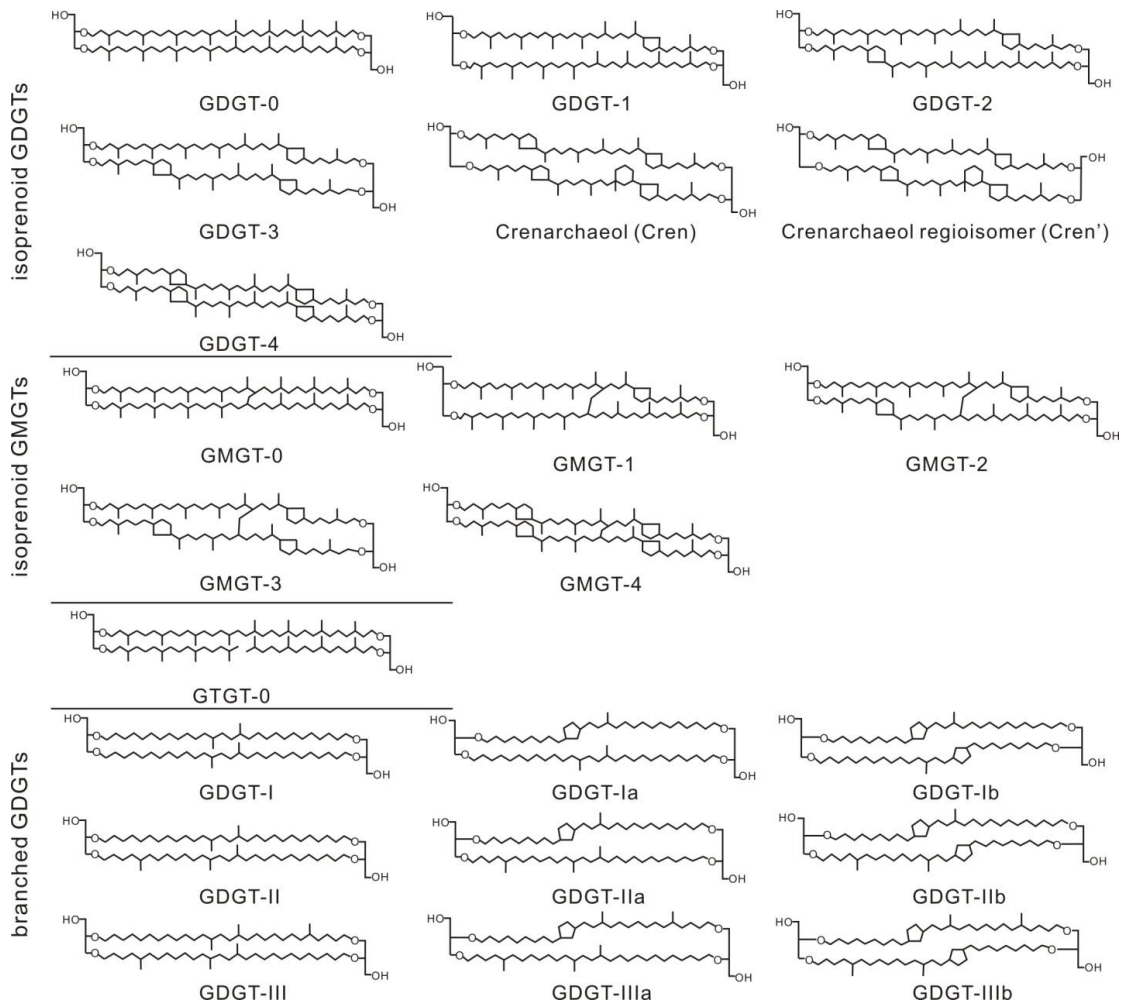


Fig. 1. Structures of tetraether lipids detected in SWIR hydrothermal field.

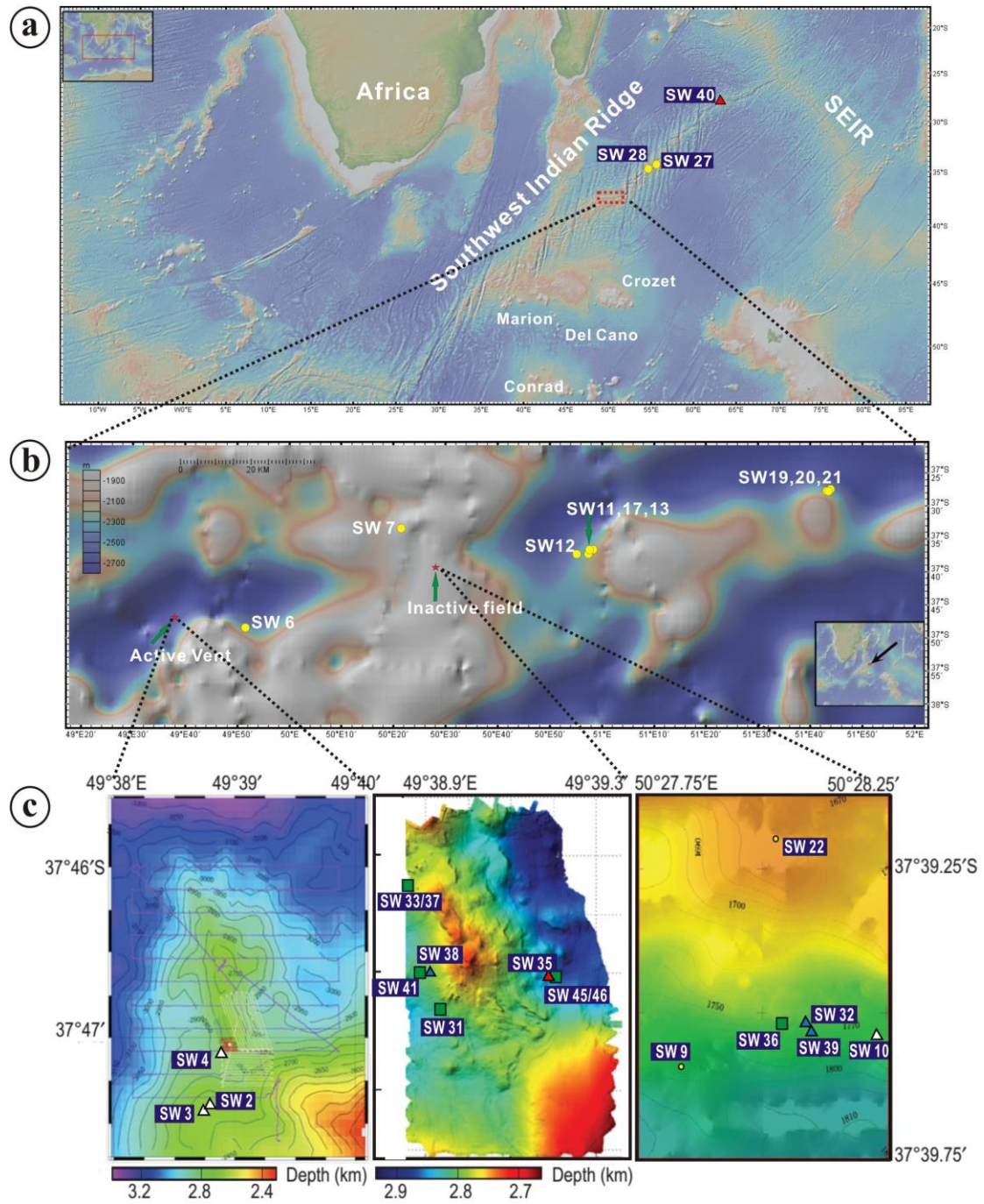


Fig. 2. Location of the samples collected from SWIR. Yellow circles mark background sediments; white, blue and red triangles mark M-T1, M-T2 and M-T3, respectively; green squares mark low-temperature hydrothermal deposits. Panel c modified after (Tao et al., 2012).

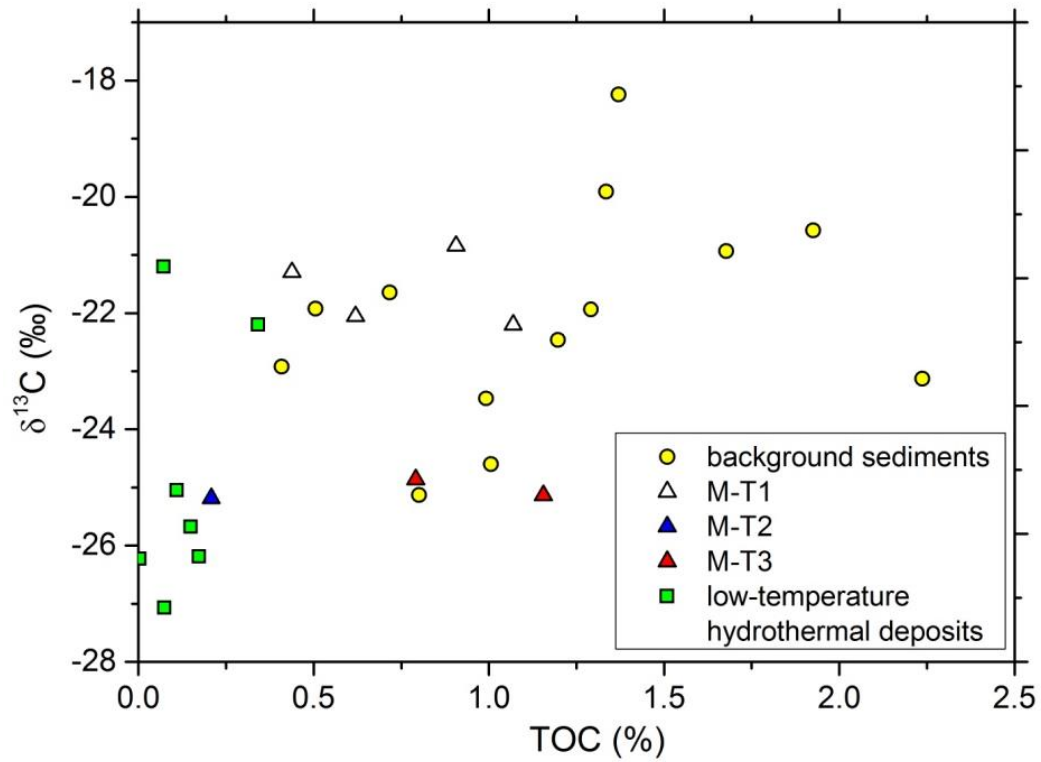


Fig. 3. Crossplot showing the overall positive correlation between TOC content and $\delta^{13}\text{C}_{\text{TOC}}$ values in background and metalliferous sediments and hydrothermal deposits from the SWIR.

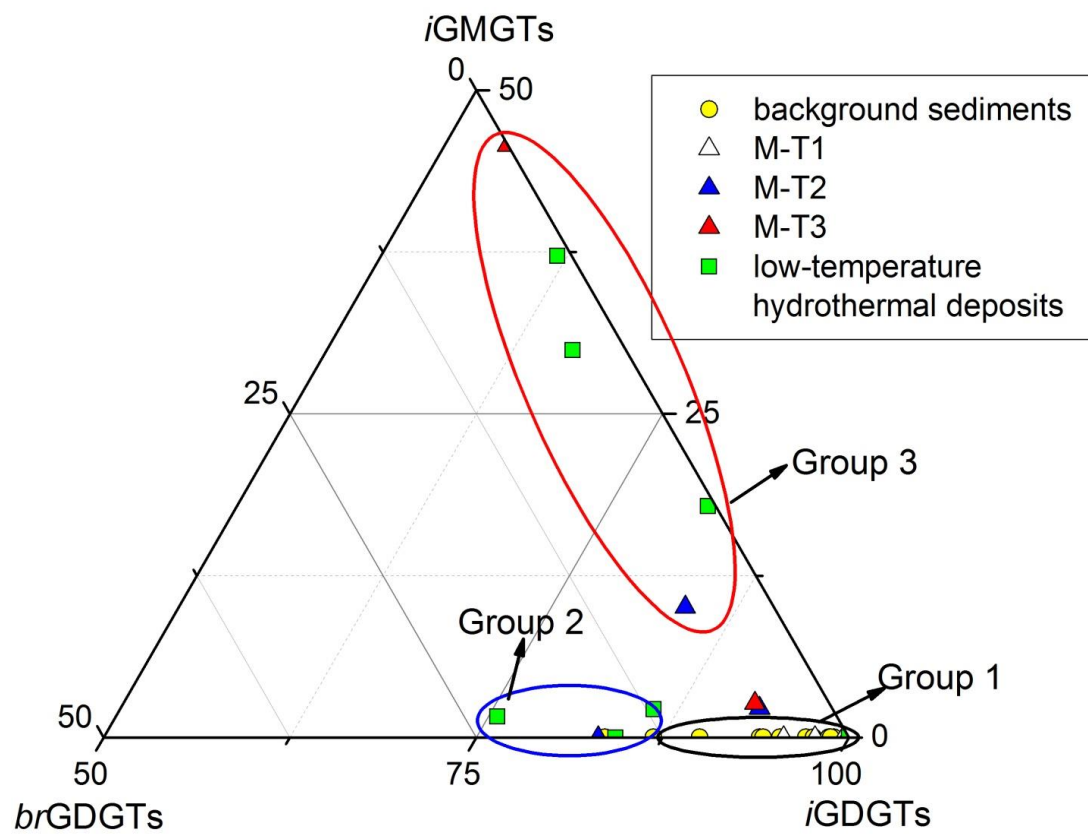


Fig. 4. Ternary diagram of *iGDGTs*, *iGMGTs* and *brGDGTs* in background and metalliferous sediments and hydrothermal deposits from the SWIR. Groups 1 to 3 were divided according to relatively higher contents of *iGDGTs*, *brGDGTs* and *iGMGTs*, respectively.

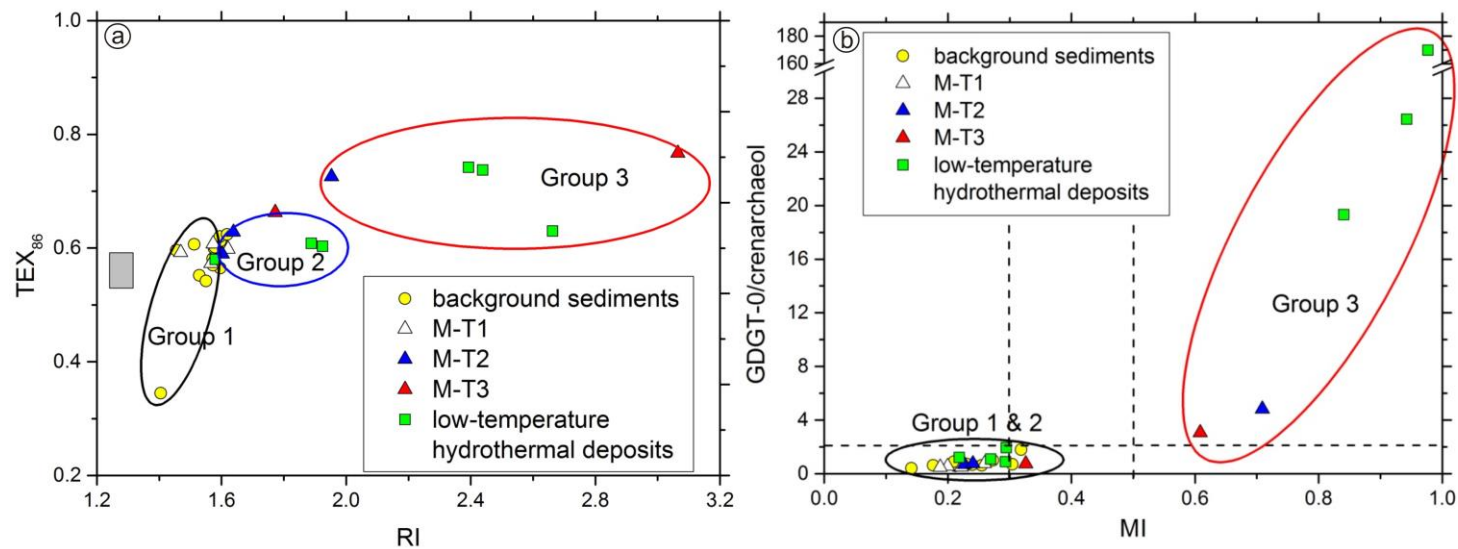


Fig. 5. Crossplots of TEX₈₆ versus Ring Index (RI) (Panel a) and MI versus GDGT-0/ crenarchaeol (Panel b) in background and metalliferous sediments and hydrothermal deposits from the SWIR. In Panel a, the bar shows expected TEX₈₆ values for the overlying water sea surface temperature (SST) of background sediments in the SWIR (SST in the range of 19 to 23 °C according to <http://www.ospo.noaa.gov/data/sst/contour/global.c.gif>, and using the TEX₈₆^H calibration of Kim et al., 2010, $SST=68.4 \times \log \text{TEX}_{86} + 38.6$); note that Group 1 but also Group 2 sediments are consistent with this, whereas Group 3 are characterized by higher than expected TEX₈₆ values. Both Group 2 and Group 3 exhibit elevated RIs.

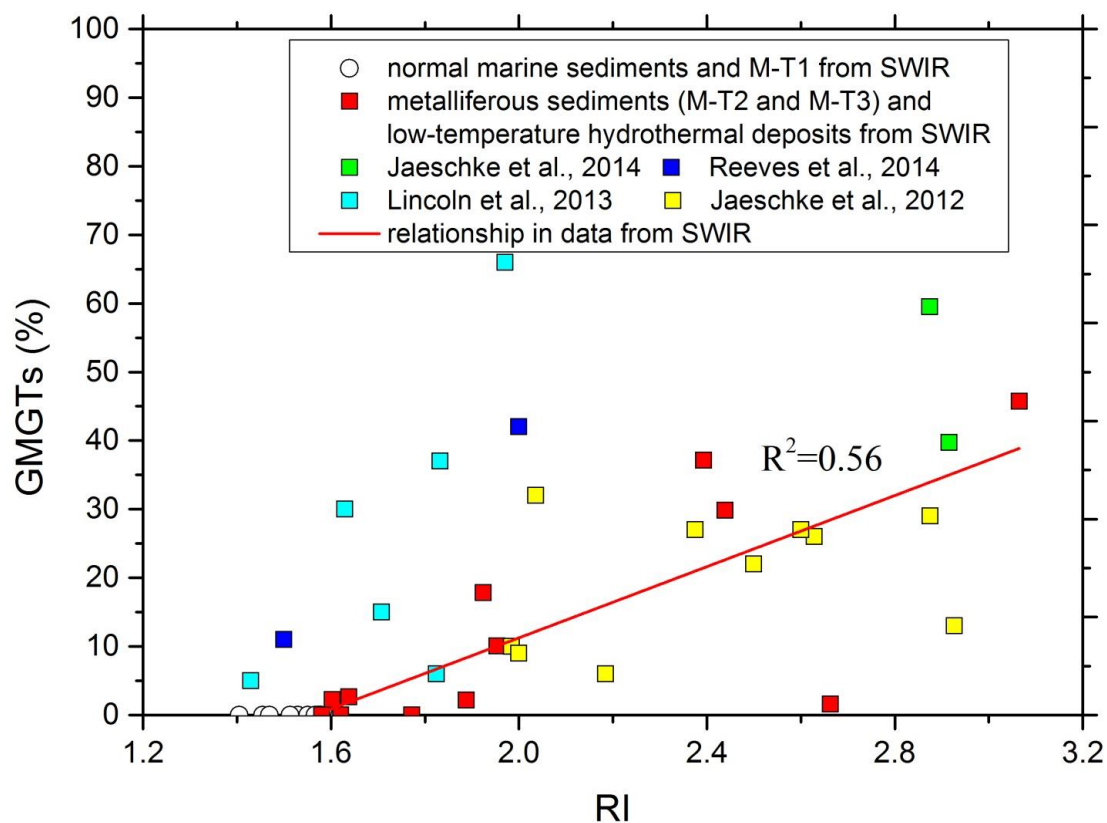


Fig. 6. Crossplot showing %GMGTs (as percentage of total tetraethers) versus RI (ring index) of background and metalliferous sediments and hydrothermal deposits from the SWIR and other hydrothermal systems. The line shows the positive relationship between GMGTs and RI in metalliferous sediments (M-T2 and M-T3) and hydrothermal deposits from SWIR, which is broadly consistent with observations from other sites. Note the several Group 1 and 2 samples with no or low GMGTs and correspondingly low RIs.

Table 1 Various tetraether lipid-based parameters for background sediments, metalliferous sediments and low-temperature hydrothermal deposits from the SWIR.

Type	No.	<i>i</i> GDGTs (%)	<i>i</i> GMGTs (%)	<i>br</i> GDGTs (%)	MBT	CBT	BIT	GDGT-0/ Crenarchaeol	RI	TEX ₈₆	MI	<i>i</i> GMGTs/ <i>i</i> GDGTs	Group
background sediments	SW6	100	0.00	0.00	—	—	0.00	0.75	1.5	0.55	0.22	0.00	1
	SW7	95	0.00	5.0	0.16	-0.11	0.05	0.63	1.6	0.60	0.26	0.00	
	SW9	97	0.00	3.0	0.11	-0.11	0.05	0.71	1.6	0.54	0.30	0.00	
	SW11	99	0.00	1.0	0.00	—	0.02	0.42	1.5	0.61	0.14	0.00	
	SW12	87	0.00	13	0.11	-0.12	0.19	1.0	1.6	0.58	0.27	0.00	
	SW13	98	0.00	2.0	0.09	-0.19	0.03	0.70	1.6	0.61	0.23	0.00	
	SW17	90	0.00	10	0.17	0.17	0.12	0.62	1.5	0.60	0.18	0.00	
	SW19	99	0.00	1.0	0.13	-0.49	0.01	0.71	1.6	0.56	0.22	0.00	
	SW20	94	0.00	6.0	0.11	0.35	0.10	0.67	1.6	0.61	0.24	0.00	
	SW21	84	0.00	16	0.14	0.14	0.31	1.8	1.4	0.35	0.32	0.00	
	SW22	94	0.00	6.0	0.00	-0.42	0.06	0.59	1.6	0.60	0.21	0.00	
	SW27	99	0.00	1.0	0.27	—	0.03	0.90	1.6	0.57	0.21	0.00	
	SW28	99	0.00	1.0	0.04	0.10	0.02	0.72	1.6	0.62	0.23	0.00	
M-T1	SW2	100	0.00	0.00	0.14	0.44	0.01	0.52	1.6	0.62	0.22	0.00	1
	SW3	96	0.00	4.0	0.00	0.12	0.06	0.56	1.5	0.59	0.20	0.00	
	SW4	98	0.00	2.0	0.00	0.28	0.03	0.50	1.6	0.61	0.19	0.00	
	SW10	100	0.00	0.00	0.00	—	0.00	0.76	1.6	0.57	0.26	0.00	
M-T2	SW32	83	0.00	17	0.44	1.06	0.28	0.71	1.6	0.60	0.23	0.00	2
	SW38	84	10	6.0	0.30	0.12	0.26	4.8	2.0	0.73	0.71	0.12	3
	SW39	93	2.2	4.8	0.28	0.25	0.07	0.76	1.6	0.59	0.24	0.02	n.d.
M-T3	SW35	92	2.7	5.0	0.15	-0.35	0.05	0.75	1.6	0.63	0.33	0.03	n.d.
	SW40	54	46	0.00	—	—	0.00	3.1	3.1	0.77	0.61	0.84	3
low-temperature hydrothermal deposits	SW31	100	0.00	0.00	—	—	0.00	2.0	1.8	0.66	0.29	0.00	n.d.
	SW33	76	1.6	23	0.69	0.04	0.26	0.89	2.7	0.63	0.29	0.02	2
	SW36	84	0.00	16	0.06	-0.28	0.26	1.2	1.6	0.58	0.22	0.00	2
	SW37	82	18	0.52	1.0	—	0.24	26	1.9	0.60	0.94	0.22	3
	SW41	86	2.2	12	0.32	0.83	0.24	1.1	1.9	0.61	0.27	0.03	2
	SW45	66	30	3.6	0.22	0.05	0.42	19	2.4	0.74	0.84	0.45	3
	SW46	62	37	0.94	0.30	0.34	0.69	170	2.4	0.74	0.98	0.60	3

— Some of the components involved in the index were not detected, precluding its calculation.

n.d., not determined; these samples were distinct from Group 1, but not characterized by the defining features of either Group 2 or Group 3.