1 2 3 4	The influence of near surface sediment hydrothermalism on the TEX ₈₆ tetraether lipid-based proxy and a new correction for ocean bottom lipid overprinting
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14 15	For submission to <i>Biogeosciences</i>
16 17 18 19 20 21	Number of pages: 27 Number of Figures: 6 Number of Tables: 2 Supplementary pages: 6
22	Key Points
23 24 25	• High <i>i</i> GDGTs turnover in shallow sediments is shown to be non-selective and does not impact TEX ₈₆ paleoclimate ratios.
26 27 28	• The proxy can be overprinted by sediment sourced lipids when geothermal temperatures rise above \sim 60–70 °C.
29 30	• A diagenetic correction model is presented to remove overprinting artifacts in the TEX ₈₆ proxy.
31 32	Abstract
33 34 35 36 37 38 39 40 41 42 43 44 45 46 47 48 49	The diversity and relative abundances of tetraether lipids produced by archaea and bacteria in soils and sediments are increasingly used to assess environmental change. For instance, the TetraEther indeX of 86 carbon atoms (TEX ₈₆), based on archaeal isoprenoidal glycerol dialkyl glycerol tetraether (<i>i</i> GDGT) lipids, is frequently applied to reconstruct past sea-surface temperatures (SST). Yet, it is unknown how the ratio fully responds to environmental and/or geochemical variations and if the produced signals are largely the adaptive response by Thaumarchaeota to oceanographic effects associated with climate or seasonal temperature changes in the upper water column. We present the results of a four push-core transect study of surface sediments collected along an environmental gradient at the Cathedral Hill hydrothermal vent system in Guaymas Basin, Gulf of California. The transect crosses a region where advecting hydrothermal fluids reach 155 °C within the upper 21cm below the seafloor (cmbsf) close to the vent center to near ambient conditions at the vent periphery. The recovered <i>i</i> GDGTs closest to the vent center experienced high rates of turnover with up to 94% of the lipid pool being lost within the upper 21 cmbsf. Here, we show that the turnover is nonselective across TEX ₈₆ GDGT lipids and does not affect the ratio independently. However, as evident by TEX ₈₆ ratios being highly correlated to the Cathedral Hill vent sediment porewater temperatures (R ² = 0.84), the ratio can be strongly impacted by the combination of severe lipid loss coupled with the addition of <i>in situ i</i> GDGT production from archaeal communities living in the vent sediments. The resulting overprint produces absolute temperature offsets of up to 4 °C based on the TEX ₈₆ ^R -calibration relative to modern climate records
50 51	of the region. The overprint is also striking given the flux of <i>i</i> GDGTs from the upper water column is estimated to be ~93% of the combined intact polar lipid (IPL) and core GDGT lipid pool initially deposited

52 on the seafloor. A model to correct the overprint signal using IPLs is therefore presented that can similarly be

53 applied to all near-surface marine sediment systems where calibration models or climate reconstructions are

- 54 made based on the TEX_{86} measure.
- 55

56 1. Introduction

57 Archaeal and bacterial tetraether cellular membrane lipids represent a group of common and structurally 58 diverse compounds frequently used to track the presence of living and dead microorganisms as well as 59 geochemical and physical conditions within present-day and paleoenvironments (e.g., Schouten et al., 2002, 60 2004; 2013; Hopmans et al., 2004; Weijers et al., 2007, 2014; Hollis et al., 2012; O'Brien, et al., 2017; Stuart 61 et al., 2017). In this regard, the proportional abundances of these lipids form various prominent proxies for 62 assessing environmental change through time. For example, TEX₈₆ (TetraEther indeX with 86 carbon atoms; 63 Schouten et al., 2002) is a widely used archaeal lipid-based paleotemperature proxy for marine environments. 64 The ratio measures variations in the number of cyclopentyl rings for a select group of archaeal core lipids 65 (CLs) (Supplementary Figure S1) following the assumption that biphytanyl cyclization is an organismal 66 response to changing sea surface temperatures (SSTs). The proxy is therefore used in many regions around the world with TEX₈₆ values typically ranging from 0.2–0.9 in marine settings (e.g. Huguet et al., 2006; Kim 67 68 et al., 2008; McClymont et al., 2012; Tierney, 2014). The utility of TEX₈₆ rests on the premise that *i*GDGTs 69 found in ocean bottom sediments are almost exclusively produced by marine planktonic archaea that inhabit 70 the epipelagic zone (Wakeham et al., 2003; Tierney, 2014; Besseling et al., 2019, 2020). Lipids are therefore 71 required to be efficiently and continually transported from the upper water column to the underlying ocean 72 floor to produce a fossil chemostratigraphic record of microbial response to changing SST conditions with 73 time (Wuchter et al., 2005).

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75 Since its introduction, the reliability of TEX₈₆ to accurately track paleoclimate variations has been questioned. 76 TEX₈₆-based SST estimates have been observed to substantially deviate from other temperature proxies (e.g. 77 Huguet et al., 2006; Rommerskirchen et al., 2011; Seki et al., 2012). For example, over the past decade, 78 considerable effort has been made to reconstruct the early Paleogene greenhouse climate system. However, 79 TEX₈₆ appears to significantly over-estimate reconstructed SSTs (Hollis et al., 2012) relative to other proxies 80 such as Mg/Ca, or clumped isotopic compositions of foraminiferal calcite, as well as various climate models 81 based on partial pressure of carbon dioxide (pCO₂) predictions (Lunt et al., 2012; Naafs et al., 2018). For late 82 Neogene climate reconstructions, TEX₈₆ has been shown to underestimate warming trends relative to the U_{37}^{ky} 83 alkenone-index (Brassell et al., 1986) derived temperatures (Lawrence et al., 2020). The apparent SST offsets 84 have been attributed to how the proxy's associated lipids change in relation to their environment and if these 85 changes are regulated by internal adaptations within the archaeon or by an overarching community 86 succession. In this regard, the debate surrounding these discrepancies largely centers on establishing 87 responses to seasonal biases (e.g. Herford et al., 2006; Wuchter et al., 2006; Huguet et al., 2011); the 88 development of adequate calibration methods (e.g. Kim et al., 2010; Pearson et al., 2013; Tierney et al. 2014); 89 identifying lipid sourcing effects - including subsurface sediments origins for those used with the calculation 90 of TEX₈₆ (e.g. Lipp and Hinrichs, 2009); as well as physical, chemical, and ecological controls for archaeon 91 *i*GDGTs cyclization (e.g. Elling et al., 2015; Qin et al., 2015; Hurley et al., 2016).

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93 For non-thermal influences, the primary concern is what archaeal taxa produce *i*GDGTs and where they are 94 sourced. To this end, most TEX_{86} lipids are thought to be produced by Marine Group I (MGI) planktonic 95 Thaumarchaeota (Brochier-Armanet et al., 2008), which are most abundant below the photic and epipelagic 96 zone (e.g., Karner et al., 2001). Within this context, many regions of the ocean floor may become highly 97 impacted by colder, deeper water column inputs (e.g. Karner et al., 2001; Huguet et al., 2007; Lopes dos 98 Santos et al., 2010; Kim et al., 2012a,b; Pearson et al., 2013; Kim et al., 2015; Ho & Laepple, 2016; Hurley 99 et al., 2016; Lui et al., 2018; Sinninghe Damsté et al., 2018). Other non-thermogenic driving forces impacting 100 the production, cyclization, and relative abundance of TEX₈₆-based lipids include organismal selectivity to 101 specific growth phases and growth rates (Elling et al., 2014; Hurley et al., 2016); redox conditions (Qin et al., 102 2015); and the incorporation of *i*GDGT from archaeal communities living in the ocean floor sediments. With 103 respect to the latter, Lipp and Hinrichs (2009) demonstrated that the production of intact polar lipid GDGTs 104 (IPL-GDGTs) by ocean floor sediment microbial communities collected in the Peru Margin were distinctly 105 different from upper water column sourced CLs and that the conversion of this living pool to fossil lipids 106 would shift TEX_{86} ratios to higher values. However, the overall impact may not be substantial as Umoh et al.

107 (2020) found little effect to the TEX₈₆ paleoclimate ratio when examining surface sediments near 108 hydrothermal vent sites on the Southeast Indian Ridge in the southern Indian Ocean. Lengger et al. (2012, 109 2014) also reported no significant deviation between the TEX_{86} values in sediment cores collected near the 110 oxygen minimum zone from that of the overlying water column in the Arabian Sea with near linear 111 degradation rates of both IPLs and CLs. All together, the *i*GDGT abundances recorded in a TEX₈₆ sediment 112 value may ultimately constitutes a multi-variable datapoint – mixing lipid components that are themselves 113 responses to: temperature, organismal substrate and metabolism dynamics, biozone niche partitions spanning 114 from the ocean surface to the shallow sediment archaeal community, which ultimately become further 115 attenuated by depositional and diagenetic processes.

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117 While not an ideal location to create SST reconstructions, hydrothermal vents of sedimented ocean basins do 118 represent an anomalous endmember to the vast expanse of ambient ocean floor sediment where paleoclimate 119 reconstructions are commonly produced. The sedimented vent systems of Guaymas Basin, Gulf of California 120 (Figure 1) is one such site. The basin experiences high sedimentation rates ranging from 0.4-0.2 cm yr⁻¹ 121 (Curray et al., 1979; Gieskes et al., 1988) due in part to the high productivity of the upper water column. The 122 ocean floor hydrothermally impacted surface sediments are also a location of active and diverse microbial 123 communities with vents that are often covered by Beggiatoa dominated microbial mats (e.g. McKay et al., 124 2012; Meyer et al., 2013; Teske et al., 2016). These sites should in principle, enable a high-resolution archaeal 125 lipid stratigraphic record that provides optimal conditions for studying potential shallow diagenetic and 126 subseafloor interferences to common archaeal lipid-based environmental proxies. The region further offers 127 an ideal setting to compare TEX₈₆ proxy responses to in situ lipid production from thermophilic sedimentary 128 archaea that differ from the pelagic background communities (e.g. Schouten et al., 2003). Recently, Bentley 129 et al. (2022) produced a survey of the source and diagenetic and catagenetic alteration of archaeal lipids from 130 the Cathedral Hill hydrothermal vent complex (Figure 1) in the Guaymas Basin, Gulf of California. Within 131 the investigation, it was observed that most *i*GDGTs are sourced from the overlying water column. Building 132 on the results of Schouten et al. (2003), it was observed that these lipids can become heavily turned over in 133 the hotter portions of the vent site where they rarely survive long enough to become cracked into hydrocarbon 134 biomarkers such as biphytanes and derivatives of biphytanes. For this study, we further examine the *i*GDGT 135 lipid distributions in these near-surface ocean floor sediments to determine if paleoclimate proxy signals can 136 be impacted by the presence of subsurface archaeal populations. The distribution of *i*GDGTs and their 137 corresponding environmental proxy signals were measured within the sediments along a transect at the 138 complex. In this regard, this site offers the unique opportunity to evaluate the response of TEX₈₆ and other 139 tetraether-lipid proxies within a microbially diverse sedimentary environment that is exposed to high 140 temperature vent fluids.





FIGURE 1 A) Location map of Guaymas Basin and the Southern Sill (red outlined box) in the Gulf of
California. Cathedral Hill is marked with a yellow star. B) Photo of Cathedral Hill taken via *Alvin*. C)
Schematic of the push core transect with a color-coding that is consistent for all plots throughout this paper.
Maps modified from Teske et al. (2016), Dalzell et al. (2021), and Bentley et al. (2022).

150 2. Material and methods

151 2.1. Study location and sampling

Four sediment push cores were collected using HOV Alvin (Dive 4462; 10/22/08) at the Cathedral Hill hydrothermal vent site, located at a water depth of 1996 m in the Southern Trough of Guaymas Basin, Gulf of California (27°0.629' N, 111°24.265' W) (Figure 1). The push cores, labeled 1 to 4, were taken along a transect with ~ 2 m spacing extending outwards from microbial mat-covered sediments near the sulfide chimney complex to just outside of the microbial mat area in ambient seafloor sediment. Thermal-probe measurements were sequentially taken beside each core (Table 1). Once the push cores were brought to the surface, the sediments were subsampled into 2-3 cm-thick depth intervals, transferred to combusted glass vials, and immediately stored at -40 °C (onboard the ship) before being shipped under dry ice to the laboratory and later freeze-dried and stored at -80 °C.

0-2 2-4			$(^{\circ}\mathbf{C})^{*}$	temperature (°C) [*]	(g)*	sed ⁻¹)*	<i>i</i> GDGT (µg g ⁻¹) [†]	(μg g ⁻¹) [‡]
2-4	GB4462-5	Black mud with microbial mat filaments	19	19	1.97	11.5	16.7	503.1
2 4	GB4462-5	Brownish-green diatomaceous mud	-	67	2.04	7.65	14.6	461.7
4-6	GB4462-5	Brownish-green diatomaceous mud	85	85	2.03	9.37	6.0	203.3
6-8	GB4462-5	Brownish-green diatomaceous mud	-	105	1.99	2.09	4.3	148.6
8-10	GB4462-5	Brownish-green diatomaceous mud	-	117	2.01	4.38	3.2	59.0
10-12	GB4462-5	Grayish-green mud	121, 124	125	2.01	1.97	1.7	48.8
12-15	GB4402-5	Brownish-green consolidated mud with clay shards	-	135	1.98	1.99	1.4	/8./
13-18	GB4402-5	Brownish-green consolidated clay	142	143	1.90	1.09	0.0	42.0
0-2	GB4462-6	Black mud with microbial mat filaments	9.13	11	2.02	8.48	17.8	591.0
2-4	GB4462-6	Black mud with microbial mat filaments	-	22	1.97	8.65	7.5	266.3
4-6	GB4462-6	Brownish-green diatomaceous mud	20	20	1.95	2.51	2.5	87.4
6-8	GB4462-6	Brownish-green diatomaceous mud	-	47	1.95	3.38	3.4	69.7
8-10	GB4462-6	Brownish-green diatomaceous mud	-	60	1.95	1.48	2.0	48.4
10-12	GB4462-6	Brownish-green diatomaceous mud	69, 77	73	1.94	4.19	2.0	52.1
12-15	GB4462-6	Brownish-green diatomaceous mud	-	87	2.02	1.69	1.0	44.2
15-18	GB4462-6	Brownish-green diatomaceous mud	118	105	1.95	2.01	0.0	22.3
18-21	GB4462-6	Brownish-green diatomaceous mud	109	125	1.94	1.38	0.0	31.2
0-2	GB4462-3	Black mud with microbial mat filaments	3.2	3.2	1.96	7.31	15.3	511.3
2-4	GB4462-3	Brownish-green diatomaceous mud	-	8	1.96	3.91	8.3	308.9
4-6	GB4462-3	Brownish-green diatomaceous mud	15	15	2.00	2.86	7.0	283.5
0-8 8 10	GB4462-3	Brownish-green diatomaceous mud	-	26 24	2.02	5.00	1.5	275.3
8-10 10.12	GB4402-3	Brownish-green diatomaceous mud	54	54 43	2.01	2.02	5.7	231.1
12-15	GB4462-3	Brownish-green diatomaceous mud	-	43 54	1.94	1.30	5.8	184.6
15-18	GB4462-3	Brownish-green diatomaceous mud	61	66	2.01	1.43	12.3	473.1
18-21	GB4462-3	Brownish-green diatomaceous mud	83	80	1.96	1.98	5.2	182.3
0-2	GB4462-8	Black mud	0	0	1.93	3.44	16.7	485.4
2-4	GB4462-8	Brownish-green diatomaceous mud	1.5	8	2.01	3.17	14.6	417.8
4-6	GB4462-8	Brownish-green diatomaceous mud	16	16	1.95	4.00	6.0	480.6
6-8	GB4462-8	Brownish-green diatomaceous mud	-	18	2.02	4.19	4.3	359.7
8-10	GB4462-8	Brownish-green diatomaceous mud	-	21	2.02	4.76	3.2	153.5
10-12	GB4462-8	Brownish-green diatomaceous mud	-	23	1.95	4.84	1.7	459.5
12-15	GB4462-8	Brownish-green diatomaceous mud	-	25	1.95	5.74	1.4	515.2
15-18	GB4462-8	Sample lost during collection	-	-	-	-	0.0	503.1
18-21	GB4462-8	Sample lost during collection	29	-	-	-	0.0	461.7
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173 174	Table 1. Cathedral Hill sample push core, sediment, geochemical, and lipid proxy data.

Core ^{*a}	Depth interval (cmbsf)	Alvin dive # and core ID	SUM of TEX ₈₆ cGDGT ^c (µg g ⁻¹)	TEX ₈₆ cGDGT ^c	TEX ^H cGDGT ^d	TEX ^H ₈₆ Reconstructed SSTs (Kim et al., 2010) ^e	RI ^f	MI ^g	TEX ₈₆ _{IPL} GDGT ^c
1	0-2	GB4462-5	110.7	0.56	-0.25	21.2	2.44	0.34	0.58
1	2-4	GB4462-5	117.1	0.58	-0.23	22.6	2.45	0.38	0.58
1	4-6	GB4462-5	47.7	0.58	-0.24	22.3	2.48	0.36	0.55
1	6-8	GB4462-5	33.0	0.58	-0.24	22.2	2.55	0.35	0.57
1	8-10	GB4462-5	13.0	0.59	-0.23	22.9	2.60	0.34	0.72
1	10-12	GB4462-5	10.1	0.57	-0.25	21.8	2.63	0.31	0.70
1	12-15	GB4462-5	17.8	0.61	-0.22	23.8	2.65	0.37	0.69
1	15-18	GB4462-5	9.8	0.61	-0.22	23.9	2.66	0.36	-
1	18-21	GB4462-5	9.3	0.63	-0.20	24.9	2.66	0.38	-
2	0-2	GB4462-6	128.5	0.55	-0.26	20.6	2.52	0.32	0.46
2	2-4	GB4462-6	58.2	0.54	-0.27	20.4	2.52	0.32	0.58
2	4-6	GB4462-6	19.2	0.54	-0.27	20.4	2.53	0.33	0.60
2	6-8	GB4462-6	13.4	0.56	-0.25	21.5	2.68	0.29	0.71
2	8-10	GB4462-6	9.3	0.58	-0.25	21.7	2.70	0.29	0.70
2	10-12	GB4462-6	10.1	0.57	-0.24	21.9	2.71	0.28	0.68
2	12-15	GB4462-6	8.5	0.57	-0.24	21.9	2.73	0.28	0.73
2	15-18	GB4462-6	4.5	0.58	-0.23	22.6	2.68	0.31	-
2	18-21	GB4462-6	6.0	0.59	-0.23	22.8	2.74	0.28	-
3	0-2	GB4462-3	127.0	0.54	-0.27	20.2	2.41	0.37	0.53
3	2-4	GB4462-3	57.7	0.53	-0.27	19.8	2.62	0.27	0.49
3	4-6	GB4462-3	60.0	0.53	-0.27	19.9	2.53	0.31	0.56
3	6-8	GB4462-3	59.8	0.54	-0.27	20.3	2.50	0.33	0.54
3	8-10	GB4462-3	53.0	0.53	-0.27	19.9	2.54	0.31	0.61
3	10-12	GB4462-3	42.1	0.54	-0.27	20.3	2.64	0.27	0.74
3	12-15	GB4462-3	39.2	0.56	-0.25	21.5	2.56	0.30	0.69
3	15-18	GB4462-3	86.8	0.55	-0.26	20.9	2.77	0.26	0.74
3	18-21	GB4462-3	36.4	0.57	-0.25	21.6	2.68	0.29	0.66
4	0-2	GB4462-8	112.9	0.54	-0.27	20.4	2.43	0.35	0.54
4	2-4	GB4462-8	85.3	0.53	-0.27	20.0	2.59	0.30	0.37
4	4-6	GB4462-8	102.7	0.54	-0.27	20.2	2.55	0.31	0.43
4	6-8	GB4462-8	70.8	0.52	-0.28	19.3	2.55	0.29	0.45
4	8-10	GB4462-8	26.6	0.53	-0.27	19.9	2.69	0.26	-
4	10-12	GB4462-8	91.0	0.53	-0.27	19.8	2.54	0.30	-
4	12-15	GB4462-8	73.7	0.53	-0.28	19.7	2.90	0.20	-
4	15-18	GB4462-8	110.7	-	-	-	-	-	-
4	18-21	GB4462-8	117.1	-	-	-	-	-	-

197 Table 1. Cathedral Hill sample push core, sediment, geochemical, and lipid proxy data (continued). 198

199 200 Also reported in Bentley et al. (2022).

[†] Sum of GDGT-1, -2, -3. -4, -5, and -5' (Table S1).

201 [‡]Sum of all detected 1G- and 2G-GDGTs (Table S3).

202 ^{*a} Collected core numbers are relabelled in the sample name to reflect a relative transect position (1-4).

203 ^{*b} Sediment lithology based on freeze-dried sediments.

204 [°] TEX₈₆ = (GDGT-2 + GDGT-3 + GDGT-5')/(GDGT-1 + GDGT-2 + GDGT-3 + GDGT-5'), (Schouten et

205 al., 2002) applied to both core GDGTs and 1-glycosyl-GDGTs (also referred to as MTEX₈₆ in section 3.4).

 d TEX₈₆^H = log ((GDGT-2 + GDGT-3 + GDGT-5')/(GDGT-1 + GDGT-2 + GDGT-3 + GDGT-5')), for 206

207 sediments outside low latitudes (Kim et al., 2010).

208 ^e Following the mean annual sea surface calibration of 0 m water depth (SST = $68.4 \times \text{TEX}_{86}^{\text{H}} + 38.6$) of 209

Kim et al. (2010).

210 ^f Ring index (RI) = $0 \times (GDGT-0) + 1 \times (GDGT-1) + 2 \times (GDGT-2) + 3 \times (GDGT-3) + 4 \times (GDGT-4) + 1 \times (GDGT-4)$

211 $5\times$ (GDGT-5)/ Σ GDGTs, adapted from Pearson et al. (2004) and promoted by Zeng et al. (2016).

212 ^g Methane index (MI) = (GDGT-1 + GDGT-2 + GDGT-3)/(GDGT-1 + GDGT-2 + GDGT-3 + GDGT-5 + GDGT-5)GDGT-5') by Zhang et al. (2011).

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220 2.2. Lipid extraction

221 Lipid extractions followed a modified Bligh and Dyer protocol laid out in Bentley et al. (2022) and following 222 Sturt et al. (2004). A subsample of freeze-dried sediment was added to a Teflon[©] centrifuge tube followed by 223 the addition of 6 ml of mix A solvent solution comprising of 2:1:0.8 v/v/v methanol (MeOH), 224 dichloromethane (DCM), and phosphate buffer (5.5 g L⁻¹ Na₂HPO₄; Avantor Performance Materials, LLC. 225 adjusted to pH of 7.4 with HCl; Anachemia Co.). The solvent sediment mixture was further spiked with 1-226 alkyl-2-acetoyl-sn-glycero-3-phosphocholine (PAF) recovery standard purchased from Avanti Polar Lipids, 227 Inc. The slurry was sonicated for 5 min then centrifuged for 5 min at 1250 rpm. The resulting supernatant 228 was added to a separatory funnel. This procedure was performed twice before being joined by two replicate 229 extractions using mix B, a 2:1:0.8; v/v/v solution of MeOH, DCM, and trichloroacetic acid buffer (50 g L⁻¹ 230 C₂HCl₃O₂: Avantor Performance Materials, LLC. of pH 2) and a final two replicate extractions using mix C, 231 a 5:1 v/v solution of MeOH and DCM. Once complete, the combined A, B, and C. For each step, the organic 232 fraction was collected in a beaker, and the combination of mix A, B, and C were subjected to 10 ml of DCM 233 and H_2O (MilliQ) to achieve separation. The organic phase was drawn off and the water was extracted using 234 3 DCM washes, drawing off the organic phase after each wash. The organic phase was back-extracted with 235 H_2O to ensure purity. The resulting organic phase was then evaporated to dryness at 60 °C under dry nitrogen. 236 The resulting total lipid extract (TLE) was transferred to pre-weighed autosampler vials using DCM:MeOH 237 1:1 v/v, spiked with 1, 2-diheneicosanoyl-sn-glycero-3-phosphocholine (C₂₁-PC; Avanti Polar Lipids, Inc.) 238 and stored at -20 °C.

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241 2.3. High performance liquid chromatography – mass spectrometry (HPLC-MS) 242

243 Mass spectrometric analyses were performed on an Agilent Technologies 1260 Infinity II HPLC coupled to 244 an Agilent Technologies 6530 quadruple time-of-flight mass spectrometer (qToF-MS) operated in positive 245 mode. Chromatographic separation used a reverse-phase method outlined by Zhu et al. (2013). The HPLC 246 was fitted with an Agilent Technologies ZORBAX RRHD Eclipse Plus C_{18} (2.1 mm × 150 mm × 1.8 µm) 247 reverse phase column and guard column maintained at 45 °C. The sample injection solvent was methanol. An 248 aliquot of each sample representing 1% of the TLE was analyzed. A 0.25 mL min⁻¹ flow rate was established 249 with mobile phase A consisting of methanol/formic acid/ammonium hydroxide (100:0.04:0.10 v/v/v) held at 250 100% for 10 min, thereafter mixed following a linear gradient with mobile phase B (propan-2-ol/formic 251 acid/ammonium hydroxide (100:0.04:0.10 v/v/v) to 24%, 65%, and 70% over 5-, 75-, and 15-min intervals, 252 respectively. Each sample run was finished by re-equilibrating the system with 100% mobile phase A for 15 253 min The effluent was ionized by an electrospray ionization source with a gas temperature of 300 °C, a 3 L 254 min⁻¹ drying gas flow, and a 5.33 μ A source current. The mass spectrometer was set to a 100–3000 m/z scan 255 range in positive mode in an untargeted method with 10 ppb resolution to simultaneously resolve both 256 archaeal IPLs and CLs.

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258 Analyte identification was achieved by accurate mass resolution, mass spectral analysis using Agilent 259 Technology's MassHunter software, and comparison of fragmentation patterns with the literature (e.g., 260 Knappy et al., 2009; Liu et al., 2010; Yoshinaga et al., 2011 – see Bentley et al., 2022 for further details). 261 Mass fragments consistent with the loss of a biphytane (m/z 743.7) were screened for all archaeal lipids. 262 Quantification was achieved by summing the integration peak areas of [M+H]⁺, [M+NH₄]⁺, and [M+Na]⁻ 263 adducts for the respective IPLs and CLs of interest. Concentration values were obtained relative to the internal 264 C_{21} -PC standard and reported in $\mu g g^{-1}$ dry sediment weight. Response factors were determined by a series of 265 injections of a standard solution containing: PAF, C₂₁-PC, 1,2-diacyl-3-O-(α-D-galactosyl1-6)-β-D-266 galactosyl-sn-glycerol (DGDG), 1,2-diacyl-3-O-β-D-galactosyl-sn-glycerol (MGDG), 1,2-di-O-phytanyl-sn-267 glycerol (archaeol), 1',3'-bis[1,2-dimyristoyl-sn-glycero-3-phospho]-glycerol (14:0 Cardiolipin) from Avanti 268 Polar Lipids, Inc., USA, and 2,2'-di-O-decyl-3,3'-di-O-(1'',\u00f6''-eicosanyl)-1,1'-di-(rac-glycerol) (C₄₆-269 GTGT) from Pandion Laboratories, LLC in amounts ranging from 100 pg to 30 ng. Response factors were 270 calculated relative to the C₂₁-PC, and the appropriate correction factor was then applied to the lipid class of 271 interest.

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A series of samples were re-run to identify or confirm deviations in the data set. The variations between the concentrations of GDGTs in the re-run and the initial runs yielded a maximum difference of $\sim \pm 4 \ \mu g \ g^{-1}$ per GDGT compound, providing confidence in the initial results and confirming the presence of two outliers in
the data set (Bentley et al., 2022). These outliers are Core 4 at 8-10 cm, with abnormally low concentrations
of all compounds that are likely ion suppression from a sample heavily impregnated with oil, and Core 3 at
15–18 cm, which contains relatively high lipid concentrations that are yet to be explained.

279 280

281 **3. Results and Discussion**

282 **3.1.** Archaeal lipid diversity and turnover

283 The Cathedral Hill transect sediments have *i*GDGTs containing 0-4 cyclopentyl (GDGT 0-4) as well as 284 crenarchaeol (Cren) and the isomer of crenarchaeol (Cren') that contains five rings (four cyclopentyl and one 285 cyclohexyl moiety) (Table S1). Branched GDGTs (brGDGTs) including Ia-c, IIa-c, and IIIa were found to 286 have discontinuous and/or low absolute abundances, with some compound classes not being detected (i.e. 287 brGDGT-IIIb; Table S2). The brGDGTs are therefore not further examined in this study. For cores 1 to 3 the 288 concentrations of nearly all *i*GDGT compounds systematically decrease with depth (Figure 2). Bentley et al. 289 (2022) established the sedimentation of archaeal lipids from the upper water column as being uniform both 290 in terms of spatial loading across the length of the transect as well as over an inferred 52.5–105 yrs of 291 sedimentation as penetrated by the length of the push core (based on sedimentation rates). From this, it is 292 estimated that $\sim 70.6 \pm 23.5 \ \mu g \ iGDGTs \ g^{-1}$ sed yr⁻¹ is being deposited on the seafloor from the overlying 293 water column. However, for cores closest to the vent site, lipid abundances exhibited a much sharper decrease 294 with depth, which Bentley et al. (2022) attribute to the turnover of archaeal lipids coupled to, but not directly 295 caused by, hydrothermalism. For cores 1 and 2, losses reach as high as 94% within the upper 21 cmbsf (cm 296 below seafloor). The lipid loss is less severe for core 3 at ~60%. For the ambient core 4, *i*GDGTs have similar 297 down core stratigraphic trends with a near-consistent average of 400 µg g⁻¹ sediment concentration and no 298 systematic loss of lipids.

299

300 Due to the high temperature conditions of the vent fluids at Cathedral Hill, the identified archaeal *i*GDGT-301 based IPLs within the sediments most likely represent the composition of cellular membrane material from 302 archaeal communities living in the sediments. These lipids have exclusively monoglycosyl (1G) or diglycosyl 303 (2G) head groups linked to a 2,3-sn-glycerol. Within the pyrolytic environment, the transformation of IPL 304 iGDGTs could hypothetically add to the core iGDGT lipid pool. Similar to CLs, the 1G-GDGTs contain 0-4 305 cyclopentyl moieties and include Cren and Cren'. Surface concentrations of these lipids are ~15 µg g-1 sed. in 306 cores 1 to 3 (residing within the microbial mat) and 11 μ g g⁻¹ sed. for core 4 (Table S2). Also similar to the 307 CLs, the archaeal IPL concentrations decrease down core and are closely coupled to increasing porewater 308 temperatures (Table S2). For cores 1 and 2, the maximum depths for detectable 1G-GDGTs are 15-18 and 309 12-15 cmbsf, corresponding to vent porewater temperatures of 145 and 87 °C, respectively. In core 3, 1G-310 GDGTs persist down core with a consistent lipid depletion that reaches its lowest concentration of 5.2 μ g g⁻¹ 311 sed. in the bottom of the core at 18–21 cmbsf sediment depth where porewater temperatures rise to 80 °C. In 312 core 4, which is most similar to the ambient ocean bottom conditions and falls outside of the area covered by 313 the microbial mat, the lipid concentrations average is $\sim 8 \ \mu g \ g^{-1}$ sed. across the depth of the core. The 2G-314 GDGTs have 0 to 2 cyclopentyl rings that for cores 1 and 2 are restricted to the upper 4 to 6 cmbsf. These 315 lipids are not further investigated in this study as 2G-GDGTs are of limited abundance (max summed 316 concentrations $<2 \mu g g^{-1}$ sed.) and their structural diversities negligibly affect isoprenoid-based proxies.

317

318 Lipid-based proxies for the calibration or reconstruction of paleoclimate records such as TEX₈₆, are based on 319 environmentally scaled contributions of select GDGT compounds. These proxies could be negatively 320 impacted should other ocean floor sediment systems experience high rates of lipid turnover (Lengger et al., 321 2014). To evaluate whether down-core depletions of lipid concentrations impacted tetraether-based proxies, 322 the concentrations of the highly abundant GDGT-0 was plotted relative to the TEX₈₆ ratio lipids (*i*GDGT-1, -323 2, -3, and Cren') (Figure 3A). For figure 3A, straight lines in the logarithmic plot indicate near-equal depletion 324 rates between the paired x- and y-axis lipid classes. Similarly, parallel slopes for the various lipid pairs also 325 indicates near-equal depletion rates, with vertical offsets between pairs marking different initial starting 326 abundances of the compared lipid. In this regard, iGDGT-0, -1, -2, and Cren' have undergone the same rate 327 of turnover. However, the depletion rate of iGDGT-3's is lower than that of other lipid classes for cores 1 and 328 2. Although, this may represent a distinct resilience to turnover, we suggest it instead results from overprinting
 329 by the subsurface hyperthermophilic archaeal community (see below).

To better track changes across each core, the degradation rate constants (k') of TEX₈₆ lipid classes were calculated for each push core (Figure S2; Table S3) using a first-order kinetic model:

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$$C_t = C_{i'} e^{-k't} \tag{1}$$

in which C_t and C_i are concentration at time (*t*) and the initial concentration, respectively (e.g. Schouten et al., 2010). Rearranging Eq. 1, the k' were calculated as

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 $k' = (-\ln[C_t/C_i])/t$ (2)

From these data, it is evident that the down core concentrations of each lipid decrease at equivalent rates (i.e. they have the same slopes for their rates of decay $s^2=0.2$). the exception to this is core 2, which independent of two outliers has different decay paths for GDGT-3 and GDGT-5. This is consistent with the TEX₈₆ *i*GDGT lipid classes largely being removed from the sediment lipid pool in a non-selective manner.

344 Based on these results, the TEX₈₆, ring index (RI), and methane index (MI) values were plotted against their 345 respective summed *i*GDGTs lipid concentrations (Fig 3B–D). For samples located within the habitable zone 346 (having porewaters ranging from 0–123 °C; Kashefi and Lovley, 2003), no correlation is observed between the lipid abundances and proxy ratios of TEX₈₆, RI, or MI (Figure 3B–D). This further suggests these proxies 347 348 are not affected by turnover in the habitable zone. However, once sediment burial reaches beyond the 349 habitable zone, TEX₈₆ ratios trend to higher values (similarly also reflected in GDGT-3 concentration trends 350 of Figure 3A). Collectively, these data strongly indicate that archaeal lipid turnover is largely nonselective of 351 the TEX₈₆ lipid classes and will therefore theoretically not in and of themselves significantly impact archaeal 352 lipid paleoclimate proxy reconstructions.

353

354 Apart from paleoclimate reconstructions, archaeal lipid CLs are sometimes used to resolve aspects of 355 localized biogeochemical cycles within sediments. To this end, the location and degree of anaerobic oxidation 356 of methane (AOM) is determined by methane and archaeal lipid carbon isotope measures (e.g. Boetius et al., 357 2000; Schouten et al., 2003; Stadnitskaia et al., 2008; Biddle et al., 2012) as well as by the proportional 358 abundances of core GDGTs (cGDGTs) in the form of the MI (Zhang et al., 2011; Carr et al., 2018; Petrick et 359 al., 2019). With respect to the latter, the MI proxy is used to differentiate regions of normal marine (with 360 values between 0-0.3) and active AOM conditions in and around cold seeps (where values >0.5-1 are 361 reported for gas hydrate impacted sediments and subsurface environments with high AOM levels). To our 362 knowledge, the use of this proxy for hydrothermal vent systems has not been thoroughly investigated even 363 though this microbial process has been well documented at Guaymas Basin. For example, highly ¹³C-depleted 364 CLs reaching up to -70% in hydrothermal vent sediments with porewater temperatures as high as 95 °C 365 indicates thermophilic archaea actively engaging in AOM (Schouten et al., 2003). Biddle et al. (2012) through 366 the detection of relevant archaeal communities by 16S RNA in conjunction with highly depleted methane 367 carbon isotope values determined active AOM spanning 35 to 90 °C porewater conditions. AOM is not likely 368 to be the dominant form of carbon and sulfur metabolism as it generally accounts for less than 5% of sulfate 369 reduction (Kallmeyer and Boetius, 2004). When applying the MI to the Cathedral Hill push core transect 370 survey low values (ranging from 0.2–0.38; Table 1) are recorded with no correspondence to thermal controls 371 across the vent transect (Figure 4). Although, it could be considered that the low values arise from a lack of 372 AOM within these sediments the low MI values are consistent with a high upper water column *i*GDGTs 373 loading as estimated by Bentley et al. (2022).





FIGURE 2. Down core profiles of the Cathedral Hill core *i*GDGTs absolute and relative lipid abundances
and their generated *i*GDGT proxies: TEX₈₆, RI, and MI. The pink background indicates transect intervals
within zones of active GDGT lipid heterotrophy (Bentley et al., 2022). The gray background are transect
regions where porewater temperatures exceeded 123 °C, marking the known upper thermal limit of life
(Kashefi and Lovley, 2003). Yellow fields are zones where oil generation and hydrocarbon degradation occur
(Dalzell et al., 2021).





●Core 1 ●Core 2 ●Core 3 ●Core 4



394 3.2. TEX₈₆ and reconstructed SSTs

McClymont et al. (2012) reported a GDGT-based reconstructed annual SSTs of 16–18 °C from particulate
 organic matter collected in ambient sediment traps in the Guaymas Basin during an annual cycle from 1996–
 1997. The reconstructed temperatures followed the calibration model for sediments outside of polar regions

proposed by Kim et al. (2010). These authors demonstrated the temperatures derived from the TEX₈₆ reconstruction were significantly lower than those produced by the closely co-varying $U_{37}^{k\prime}$ paleoclimate proxy, and satellite measured estimates that jointly estimated a mean annual sea surface temperature (MASST) of 23 °C. The longer 21-year (1982–2004) satellite-derived MASST is also reported to be higher at 24 °C (Herrera-Cervantes et al., 2007). Although, the sites and time frames of these surveys do not match that of the Cathedral Hill survey, they do provide context to what our reconstructed TEX₈₆ values should record.

406

407 The high sedimentation rate at Cathedral Hill has resulted in near homogenous inputs of organic matter from 408 the upper water column across the transect area (Dalzell et al., 2021; Bentley et al., 2022). Therefore, TEX₈₆ 409 reconstructions should produce equivalent cross-transect trends with sediment depth. Nonetheless, as with 410 changes in the archaeal lipid concentrations, the profiles of iGDGT proxies TEX₈₆ and RI of the transect 411 similarly have down core trends (Figure 2; Bentley et al., 2022). For core 4, TEX₈₆ span a narrow range of 412 values (n=7; 0.52-0.54, avg. 0.53 ± 0.01 ; Figure 4A) across a period of ~ 37.5 to 75 yrs. corresponding to the 413 depth of the cores. To a slightly lesser degree, the core top (0-2 cmbsf) across the transect also display near-414 equal values to core 4 (n=4; 0.56–0.54; avg. 0.55 \pm 0.01). These values mark a TEX^H₈₆ reconstructed mean 415 annual SST of 19.3–20.4 °C following the Kim et al. (2010) calibration model (Table 1). However, the TEX₈₆ 416 values recorded in cores 1 to 3 at Cathedral Hill have considerably larger ranges with values spanning from 417 0.53 to 0.63 (Table 1) that systematically increase with rising porewater temperatures ($R^2 = 0.83$; Table 1; 418 Figure 2 and 4A). This increase is most noticeable in core 1 where the highest TEX₈₆ values are obtained 419 from the bottom core sediments (10-21 cmbsf; marking the non-habitable zone) where TEX₈₆ values span 420 0.57-0.63 (Table 1; Fig 4A) corresponding to a TEX^H₈₆ reconstructed SST change of 3.1 °C marking a range 421 from 21.8 to 24.9 °C (Table 1). The fundamental driver for the proxy's is likely influenced by the archaeal 422 community composition that is responding to their exposure to *in situ* vent fluid temperatures (Figure 4).

423

424 Two mechanisms are considered for the observed proxy variations. The first is that progressive ring-loss due 425 to carbon-carbon bond cleavage of pentacyclic rings moleties by exposure to the sharp geothermal gradient 426 acts to systematically attenuate the iGDGT lipid pool. Hydrous pyrolysis experiments conducted by Schouten 427 et al. (2004) demonstrated that at extreme temperatures (ca. >160 °C), TEX₈₆ values become negatively 428 impacted by the preferential destruction of polycyclic GDGTs. Such losses produce progressively lower ratio 429 values. Although, the transect sediment porewaters do not reach the pyrolytic temperatures of the Schouten 430 et al. (2004) experiment, they are high enough to generate hydrocarbons (Dalzell et al., 2021) and 431 thermochemically degrade iGDGTs in the hottest regions of the transect they are also more long-lived than 432 what is produced from a laboratory experiment. However, the observed stratigraphic TEX_{86} trends do not 433 match those of predicted ring loss as the values increase rather than decrease in relation to elevated porewater 434 condition. Nonetheless, the thermochemical oxidative loss of GDGTs and its effect on the TEX₈₆ ratio is 435 further explored below (section 3.4).

436

437 The second mechanism is that subsurface microbial communities donate enough core GDGTs to overprint 438 the detrital signal source. The RI (Figure 4B) values were similarly compared to recorded porewater 439 temperatures to better interpret the TEX_{86} trends and to ensure that the Cathedral Hill reconstructed 440 temperatures are influenced by the subsurface microbial community. In this regard, RI is used to monitor the 441 adaptive response of an archaeal community at the hydrothermal vent site. Lipid cyclization is an adaptive 442 response to changing environmental temperature or acidity in which an archaeon increases its rigidity by 443 decreasing the fluidity and permeability of its cellular membrane that, therefore, also further regulates the 444 flow of solutes and nutrients in and out of the cell (Gliozzi et al., 1983; De Rosa and Gambacorta, 1988; Uda 445 et al., 2001; Schouten et al., 2002; Macalady et al., 2004; Boyd et al., 2013). Both cores 1 and 2 have RI 446 values highly correlated to temperature ($R^2 = 0.87$ and 0.75, respectively) consistent with heat stress adaption. 447 This same was also observed in the Guaymas Basin by Schouten et al. (2003) who reported an increase in the 448 RI of core lipid GDGTs with in situ temperature. As such, a significant proportion of the measured *i*GDGTs 449 likely emanates from archaeal communities living in the shallow sediments of Cathedral Hill. As such, the 450 lipid cyclization pattern may reflect stratigraphically discrete thermophilic to hyperthermophilic communities 451 that are selectively adapted to more extreme temperature conditions (see Bentley et al., 2022 for further 452 discussion on the lipid-based taxonomic make-up of the vent site).



FIGURE 4. Cross plots of A) TEX₈₆, B) RI, and C) MI, *i*GDGT proxies versus porewater temperature. TEX^{*H*}₈₆ reconstructed MASSTs are based on Kim et al. (2010). Blue field indicates MI values for normal marine conditions (Zang et al., 2011).

455 **3.3.** Lipid signal sourcing

456 To evaluate the sources of measured archaeal lipids, CL and $_{IPL}TEX_{86}$ (the ratio applied to IPLs that contain 457 equivalent core lipids) indices were compared as signal responses from their respective pools of living and 458 dead cellular debris (Figure 5). For cores 1, 2, and 3 the 1G-*i*GDGT $_{IPL}TEX_{86}$ measures are positively

459 correlated with temperature ($R^2 = 0.46$, 0.74, and 0.66, respectively; Figure 5A). In this regard, 1G-*i*GDGT

460 IPLTEX₈₆ ratio appears to be largely influenced by in situ porewater temperatures as well as may by the 461 archaeal community ecology of the vent system. Factors such as community composition and adaptation 462 may further impact the $_{IPL}TEX_{86}$ ratio as the rates of changes between cores 1–3 are not the same. Similar to 463 the _{CL}TEX₈₆ values, the _{IPL}TEX₈₆ is not correlated to their summed TEX₈₆ lipid abundances (Figure 5B). Such 464 a condition is largely consistent with the living lipid pool being modified by the archaeal community's 465 response to thermal stress and not by subsequent thermal-oxidative transformation occurring shortly after cell 466 death. 467 The IPL and CL lipids of transect samples can be further grouped into three clusters (A, B, C), suggesting a 468 mixed signal for the sourcing of archaeal GDGTs from both the living and dead pools of archaea (Figure 5C) 469 closely tracking temperature. In this plot, we assume that clusters falling on the 1:1 line indicate the living 470 biota can equally contribute to the dead pool of total recovered GDGTs. Those off-axis contribute either less 471 or more to one or the other lipid pool. The three clusters mark unique thermal zones within the transect area 472 with cluster A being composed of the ambient core 2 to 4 seafloor surface samples; cluster B marking a mix 473 of intermediate temperature samples from all cores; and cluster C being composed of high temperatures 474 samples. The lipid groups likely mark distinct archaeal communities. As cluster B resides on the 1:1 line, the 475 TEX₈₆ core lipids likely have a mix of detrital and *in situ* inputs. Cluster C, however, appears likely dominated 476 by *in situ* lipid production. The thermal zonation and equivalent directionality of the resulting ratios (i.e., both 477 CL and IPLTEX₈₆ ratios increase with porewater temperature) further supports overprinting of the original

478 _{CL}TEX₈₆ sea surface signal by the ocean bottom sediment archaeal community as a mechanism for the observed _{CL}TEX₈₆ trends.

480

481 Collectively, these results suggest the source of the archaeal CLs measured in the TEX₈₆ and RI indices 482 progressively become more dominated by subsurface microbial communities adapted to the hotter 483 hydrothermal vent fluids. Our results also indicate that in select natural environments, such as hydrothermal 484 vent complexes, the TEX₈₆ SST-proxy may entirely record ocean bottom sediment porewater temperatures. 485 To our knowledge, a clear case of overprinting to this level has not yet been demonstrated.



●Core 1 ●Core 2 ●Core 3 ●Core 4

FIGURE 5. Cross plots of 1G-*i*GDGTs $_{IPL}TEX_{86}$ versus (A) porewater temperatures and (B) the concentration of 1G-*i*GDGTs in the sediments. C) TEX_{86} proxy of core GDGTs vs 1G-GDGTs. Clusters A–C may represent different archeal communities that are providing varying inputs of *i*GDGT to the core GDGT lipid pool. The dotted trendline is the partial least square regression of the complete core lipid TEX_{86} data set. The solid line marks the 1:1 CL to IPL proxy correspondence indicating both allochthonous and autochthonous sources contribute equally to the core GDGT lipid pool.

488 **3.4. TEX₈₆ overprint corrections**

The measured TEX_{86} ($_M\text{TEX}_{86}$) value of the Cathedral Hill sediments is herein considered to be a weighted sum of a sea surface TEX_{86} ($_{SS}\text{TEX}_{86}$) value acquired from lipids sourced in the upper water column that is further modified by a component of the deeper water column sourced core lipids ($_{WC}\text{TEX}_{86}$) as well as by additions of archaeal lipids from the benthic and subsurface microbial communities ($_{Sed}\text{TEX}_{86}$). These ratio 493 loadings are collectively also potentially further modified by diagenetic influences in the ocean bottom 494 sediments. Over the cumulative sediment burial period and in consideration of the measured porewater 495 temperatures of the Cathedral Hill push core sediments, these influences include the selective loss of lipids 496 by their binding into protokerogen (*K*) and by potential changes due to the loss of lipid by turnover (φ ; section 497 3.1). Additional catagenetic effects from thermochemical alteration of lipids (θ) may also attenuate the sum 498 of sedimentary core lipids by their exposure to high temperature vent fluids. Collectively, these effects are 499 considered to form the following relationship:

$${}_{M}\text{TEX}_{86} = \frac{a_{SS}\text{TEX}_{86} + b_{WC}\text{TEX}_{86} + c(d_{0-n})_{Sed}\text{TEX}_{86}}{\varphi + K + \theta}$$
(3)

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503 where *a*, *b*, and *c*, are measured scaling parameters for lipid loading and φ , *K*, and θ are diagenetic and catagenetic alteration parameters. Solving for _{SS}TEX₈₆: 505

$$_{ss}\text{TEX}_{s6} = \frac{_{M}\text{TEX}_{s6}(\varphi + K + \theta)}{a} - \frac{b_{wc}\text{TEX}_{s6} + c(d_{0-n})_{Sed}\text{TEX}_{s6}}{a}$$
(4)

506 507

508 In this regard, a portion of the archaeal community from the upper water column, presumably initially sourced 509 of IPLs, and an additional community inhabiting the ocean floor sediments were assumed to eventually die 510 with their respective IPLs gradually hydrolyze, joining the CL pool where they further contribute to the 511 observed $_{M}TEX_{86}$ value. For this study, no data was collected to calculate $b_{WC}TEX_{86}$ and its potential impact 512 on $_{M}$ TEX₈₆ cannot be further considered in this study. However, it is highly likely, given the longer residence 513 times for glycosidic-based headgroups of the identified archaeal IPLs and their relatively short settling time 514 through the water column (Lengger et al., 2012; Xie et al., 2013) that a component of this lipid source was 515 already mixed with the SedTEX₈₆ contribution. For this study, SedTEX₈₆ is an IPLTEX₈₆ ratio based on detected 516 1G-GDGT-1, -2, -3, Cren' and 2G-GDGT-1, -2, as present in the original paleoclimate proxy (Table 1; Figure 517 6). Testing the removal of 2G-GDGTs lipids, which have a low absolute concentration ($\leq 2 \mu g g^{-1}$ sed.) and 518 shallow stratigraphic zones of occurrence (section 3.1; Table S2), yielded a negligible <1 °C change in the 519 summed average reconstructed SST.

520

521 The $c(d_{0-n})$ measured scaling parameter was calculated as

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$$c(d_{0-n}) = \sum_{i=0}^{n} \left(\frac{[GDGTs_{IPL-TEX_{ss} lipids}]_n}{[GDGTs_{CL-TEX_{ss} lipids}]_{0-2cm}} \right)$$
5)

523 524

525 using the summed concentrations of 1G- and 2G-GDGTs that have the potential to become converted to 526 cGDGTs by progressive burial diagenesis and $d_{0,n}$ marking the range of sampled sediment depths, with 0 527 being the 0-2 cmbsf core top and n the deepest point of sediment burial. These intervals are divided by the 528 water column input of TEX₈₆ lipids ([GDGTs $_{CL-TEX86 \text{ lipids}}]_{0-2cm}$) estimated to be 120 µg g⁻¹ sed. based on their 529 average measured concentration across the four-core transect. The function assumes the surface sediment does not hydrolyze its IPL-GDGTs to CLs (Table 2). When applied to Eq. 4 and further excluding φ , *K*, and θ , the *ss*+*wc*TEX^H₈₆ reconstructed SSTs average 19.68 ±0.79 °C (Table 2; Figure 6A) with the total samples 530 531 532 having an unchanging depth profile that mirrors the range of values measured in the ambient sediments of 533 core 4 (Figure 2).

534

535 The selective lipid removal by diagenetic and catagenetic processes theoretically may also affect the TEX_{86} 536 value; however, their perspective impact on the directionality and magnitude of the ratio are difficult to predict 537 and equally hard to discretely measure. Although the loss of GDGTs to protokerogen formation could 538 potentially impact the ratio, it was shown to be a negligible sink for the lipids (Bentley et al., 2022). As such, 539 the K parameter in Eqs. 3 and 4 was therefore assigned a 0 value. Due to the high geothermal gradient at 540 Cathedral Hill, some of the transect push core sediments resided within zones of active catagenesis (Fig. 2; 541 Dalzell et al., 2021). The degradation rates of each TEX_{86} lipid were independently measured for the four 542 push cores (Eq. 2; Fig. S2). As the abundance of both CLs and IPLs differentially decreases through the

544 changes (section 3.1), the degradation rates must also record the effects of thermochemical oxidative 545 weathering (Fig. 3B). In this case, φ and θ are treated as grouped parameters. To determine if individual lipid 546 classes were selectively removed during degradation, the variance (s^2) of the rate change as measured from 547 its respective regression slope (i.e. $m_{logk'}$) from the TEX₈₆ lipids (Figure S2; Table S4 from Eq. 2) were 548 calculated. For the Cathedral Hill transect, the calculated mlogk' s² is 0.20, which is due to accelerated 549 degradation rates for higher ring lipids, GDGT-3 and Cren', in samples from cores 1 and 2, where high vent 550 temperatures resulted in hydrocarbon generation of the sediments (Dalzell et al., 2021). A weighing function 551 for the degree of lipid class selectivity during turnover is proposed:

$$\varphi + \theta = 1/_M \text{TEX}_{86}^{0.2} \tag{6}$$

555 When applied to Eq. 4, the corrected data series produces an average transect SS+WCTEX^H₈₆ reconstructed SST 556 of 23.66 ±0.59 °C with a near-zero partial least squares regression slope (Table 2; Figure 6B). As these 557 modeled values are within the 23-24 °C obtained for the 21-year (1982-2004) satellite-derived MASST data 558 for the Guaymas Basin region (Herrera-Cervantes et al., 2007). Based on these calculations, nearly all MTEX₈₆ 559 attenuation can be attributed to sediment microbial overprinting coupled to diagenetic and catagenetic loss of 560 lipids consistent with prior observations at Guaymas Basin (Schouten et al., 2003; Zhang et al., 2011). The 561 high degree of influence this has on the TEX₈₆ proxy is striking given that the upper water flux of GDGTs at 562 Cathedral Hill is estimated to represents up to 93% of the total intact polar and core GDGT lipid pool within 563 these sediments. Although, this study demonstrates the benthic microbial community can influence TEX_{86} 564 values in anomalous, end-member environments; the above model has not yet been tested across conventional 565 ocean shelf environments.

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FIGURE 6. Reconstructed combined $_{SS}TEX_{86}$ and SSTs $_{WC}TEX_{86}$ from Eq. 4 (A) with and (B) without φ , *K*, and θ scaling parameters compared to measured porewater temperatures. Colored circles indicate recorded values from the four push cores. $_{M}TEX_{86}$ values are plotted for reference (open black diamonds). Blue field is the 23–24 °C range observed for the 21-year (1982–2004) satellite-derived MASST data (Herrera-Cervantes et al., 2007).

580				Ĩ					
Sample	Depth (cmbsf)	Porewater Temp. (°C)	t Time (yrs.)	MTEX ₈₆ (Measured <i>i</i> GDGT TEX ₈₆)	TEX ^H 86 Reconstructed SST (°C)	TEX ₈₆ 1G- & 2G- GDGT IPLs (μg g ⁻¹)	Cumulative 1G- & 2G- GDGTs Loading with Depth (µg g ⁻¹)	SedTEX ₈₆ (i.e. 1G- & 2G-GDGT 1PLTEX ₈₆)	c(do-n) Cumulative Weighted IPL Loading (Eq. 5)
Core 1 (0-2cm)	1	19	10	0.56	21.2	4.80	0	0.58	0.00
Core 1 (2-4cm)	3	67	20	0.58	22.6	3.41	4.80	0.58	0.04
Core 1 (4-6cm)	5	85	30	0.58	22.3	1.29	8.21	0.55	0.07
Core 1 (6-8cm)	7	105	40	0.58	22.2	1.14	9.50	0.57	0.08
Core 1 (8-10cm)	9	117	50	0.59	22.9	1.41	10.64	0.72	0.09
Core 1 (10-12cm)	11	125	60	0.57	21.8	0.76	12.05	0.70	0.10
Core 1 (12-15cm)	13	135	70	0.61	23.8	0.72	12.81	0.69	0.11
Core 1 (15-18cm)	17	145	80	0.61	23.9	0.00	13.53	0.69*	0.11*
Core 1 (18-21cm)	20	153	90	0.63	24.9	0.00	13.53	0.69*	0.11*
Avg.				0.59	22.84				
Std. Dev.				0.02	1.16				
Core 2 (0-2cm)	1	11	10	0.55	20.6	4.33	0	0.49	0.00
Core 2 (2-4cm)	3	22	20	0.54	20.4	1.80	4.33	0.57	0.04
Core 2 (4-6cm)	5	20	30	0.54	20.5	0.76	6.13	0.60	0.05
Core 2 (6-8cm)	7	47	40	0.56	21.5	1.31	6.89	0.73	0.06
Core 2 (8-10cm)	9	60	50	0.58	22.3	0.88	8.20	0.70	0.07
Core 2 (10-12cm)	11	73	60	0.57	22.0	0.92	9.08	0.68	0.08
Core 2 (12-15cm)	13	87	70	0.57	21.8	0.40	10.00	0.73	0.08
Core 2 (15-18cm)	17	105	80	0.58	22.6	0.00	10.40	0.73*	0.09
ore 2 (18-21cm)	20	125	90	0.59	22.7	0.00	10.40	0.73*	0.09*
Avg.				0.56	21.61				
Std. Dev.				0.02	0.91				
Core 3 (0-2cm)	1	3.2	10	0.54	20.2	3.51	0	0.56	0.03
Core 3 (2-4cm)	3	8	20	0.53	19.9	1.79	3.51	0.51	0.01
Core 3 (4-6cm)	5	15	30	0.53	19.9	1.45	5.30	0.57	0.01
Core 3 (6-8cm)	7	26	40	0.54	20.3	1.77	6.74	0.55	0.01
Core 3 (8-10cm)	9	34	50	0.53	19.9	1.70	8.51	0.61	0.01
Core 3 (10-12cm)	11	43	60	0.54	20.3	2.16	10.21	0.71	0.02
Core 3 (12-15cm)	13	54	70	0.56	21.4	2.52	12.37	0.69	0.02
Core 3 (15-18cm)	17	66	80	0.55	20.9	4.72	14.89	0.73	0.04
Core3 (18-21cm)	20	80	90	0.57	21.6	2.10	19.61	0.65	0.02
Avg.				0.54	20.50				
Std. Dev.				0.01	0.67				
Core 4 (0-2cm)	1	2	10	0.54	20.4	2.43	0	0.54	0.02
Core 4 (2-4cm)	3	8	20	0.53	20.0	1.75	2.43	0.44	0.01
Core 4 (4-6cm)	5	16	30	0.54	20.2	2.15	4.18	0.49	0.02

579	Table 2. F	Reconstructed	sea surface	temperatures.
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Core 4 (4-6cm)

Core 4 (6-8cm)	7	18	40	0.52	19.3	1.76	6.34	0.47	0.01
Core 4 (8-10cm)	9	21	50	0.53	19.9	0.44	8.09	-	-
Core 4 (10-12cm)	11	23	60	0.53	19.8	2.20	8.54	-	-
Core 4 (12-15cm)	13	25	70	0.53	19.7	0.00	10.74	-	-
Avg.				0.53	19.90				
Std. Dev.				0.01	0.34				
Cumulative Avg.					19.68				
Cumulative Std. Dev.					0.79				

* Marks inherited values from the above sediment horizon.

	Eq.	4 excluding φ+θ+	K	Eq. 4 including φ+θ+K			
Sample	ss+wc TEX₈₆ (M TEX₈₆ - c(do-n)*sed TEX₈₆)	ss+wcTEX ^H (after Kim et al., 2010)	ss+wcTEX ^H Reconstructed SST (°C)	$\phi+\theta$ (Eq. 6) (where s ² = 0.20; Table S4)	SS+WCTEX86	ss+wcTEX ^H ₈₆ Reconstructed SST (°C) (after Kim et al., 2010)	
Core 1 (0-2cm)	0.56	-0.25	21.2	1.12	0.63	24.7	
Core 1 (2-4cm)	0.56	-0.25	21.4	1.12	0.63	24.9	
Core 1 (4-6cm)	0.54	-0.27	20.3	1.13	0.62	24.2	
Core 1 (6-8cm)	0.53	-0.27	19.8	1.13	0.61	23.9	
Core 1 (8-10cm)	0.52	-0.28	19.5	1.14	0.61	23.7	
Core 1 (10-12cm)	0.50	-0.30	17.9	1.15	0.58	22.6	
Core 1 (12-15cm)	0.53	-0.27	20.0	1.13	0.62	24.2	
Core 1 (15-18cm)	0.53	-0.27	19.8	1.13	0.61	24.1	
Core 1 (18-21cm)	0.55	-0.26	21.0	1.13	0.63	25.0	
Avg.	0.54	-0.27	20.10	1.13	0.61	24.14	
Std. Dev.	0.02	0.02	1.08	0.01	0.02	0.75	
Core 2 (0-2cm)	0.55	-0.26	20.6	1.13	0.62	24.2	
Core 2 (2-4cm)	0.52	-0.28	19.2	1.14	0.60	23.3	
Core 2 (4-6cm)	0.51	-0.29	18.7	1.14	0.59	22.9	
Core 2 (6-8cm)	0.52	-0.28	19.3	1.14	0.60	23.4	
Core 2 (8-10cm)	0.53	-0.28	19.7	1.14	0.61	23.8	
Core 2 (10-12cm)	0.52	-0.28	19.1	1.14	0.60	23.4	
Core 2 (12-15cm)	0.51	-0.29	18.5	1.14	0.59	23.0	
Core 2 (15-18cm)	0.52	-0.28	19.2	1.14	0.60	23.5	
Core 2 (18-21cm)	0.52	-0.28	19.3	1.14	0.60	23.6	
Avg.	0.52	-0.28	19.32	1.14	0.60	23.47	
Std. Dev.	0.01	0.01	0.60	0.00	0.01	0.40	
Core 3 (0-2cm)	0.52	-0.28	19.4	1.14	0.60	23.3	
Core 3 (2-4cm)	0.52	-0.28	19.4	1.14	0.60	23.3	
Core 3 (4-6cm)	0.53	-0.28	19.5	1.14	0.60	23.4	
Core 3 (6-8cm)	0.53	-0.27	19.9	1.13	0.60	23.6	
Core 3 (8-10cm)	0.52	-0.28	19.4	1.14	0.60	23.3	
Core 3 (10-12cm)	0.53	-0.28	19.6	1.14	0.60	23.5	
Core 3 (12-15cm)	0.55	-0.26	20.7	1.13	0.62	24.3	
Core 3 (15-18cm)	0.52	-0.28	19.3	1.14	0.60	23.4	
Core3 (18-21cm)	0.55	-0.26	21.0	1.13	0.62	24.6	
Avg.	0.53	-0.27	19.79	1.14	0.60	23.64	
Std. Dev.	0.01	0.01	0.62	0.00	0.01	0.49	
Core 4 (0-2cm)	0.53	-0.27	19.8	1.13	0.60	23.6	
Core 4 (2-4cm)	0.53	-0.28	19.7	1.14	0.60	23.4	

Table 2. Reconstructed sea surface temperatures (continued).

Core 4 (4-6cm)	0.53	-0.28	19.8	1.14	0.60	23.5
Core 4 (6-8cm)	0.52	-0.29	19.0	1.14	0.59	22.9
Core 4 (8-10cm)	-	-	-	-	-	-
Core 4 (10-12cm)	-	-	-	-	-	-
Core 4 (12-15cm)	-	-	-	-	-	-
Avg.	0.53	-0.28	19.51	1.07	0.60	23.38
Std. Dev.	0.01	0.01	0.38	0.00	0.01	0.31
Cumulative Avg.			19.68			23.66
Cumulative Std. Dev.			0.79			0.59

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584 4. Conclusions

585 In this study, we demonstrate a pronounce overprint of cGDGTs sourced from the ocean floor sedimentary 586 archaeal community at the Cathedral Hill vent site in Guaymas Basin. The overprint is marked by lipids with 587 more cyclized ring moieties marking an adaptive response by archaea to rigidify the cellular membranes 588 against localized heat stress. This in turn has resulted in the commonly used TEX₈₆ paleoclimate proxy to 589 partially record advecting porewaters temperatures. As the vast majority of cGDGTs in these sediments is 590 sourced from the overlying water column, the impact on the TEX₈₆ ratio is further the product of rapid lipid 591 turnover rates and diagenetic and catagenetic alteration processes potentially unique to the hydrothermal system. Together, these factors resulted in absolute TEX^H₈₆ temperature offsets of up to 4 °C based on 592 593 calibrations closely suited to the latitudinal position of Guaymas Basin. To untangle the impact of these 594 coupled drivers on the TEX_{86} proxy, we further present a method to correct the overprints by both the water 595 column and subsurface archaeal community using IPLs extracted from both of these sources. Although, we 596 have not been able to test this model with lipid inputs from the overlying water column, we have demonstrated 597 its effectiveness at removing sediment sourced overprints, which may not be unique to hydrothermal systems. 598 This approach should be capable of being extended to all near-surface marine sediment systems and may 599 improve the quality of calibration models or climate reconstructions that are based on modern TEX₈₆ 600 measures.

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604 Acknowledgments

605 A special thank you is extended to Associate Editor Jack Middelburg and the reviewers of Biogeosciences 606 who provided highly constructive feedback. We are grateful to Carl Peters, formally at Saint Mary's 607 University, who provided considerable feedback and advice during the course of this study. We further thank 608 the officers, crew, and pilots of the R/V Atlantis and HOV Alvin for their expert help at sea and their 609 outstanding efforts acquiring the samples for this study. Julius Lipp, Florence Schubotz, and Kai-Uwe 610 Hinrichs of MARUM, assisted our lab in the development of lipidomic analytical techniques. Special thanks 611 is extended to Clarissa Sit for the use of her HPLC-qToFMS. Sean Sylvia assisted with the preparation of 612 push cores used in sampling. Funding: Funding for this study through NSERC Canadian Research Chair, 613 Canada Foundation for Innovation (CFI) JELF-CRC, NSERC Discovery Grant (Application Number: 614 RGPIN-2017-05822), WHOI Deep Ocean Exploration Initiative 2008, and NSF grant MCB-0702677 (to JSS 615 and SMS).

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618 **Conflicts of Interest**

619 The authors declare no conflict of interest.

Supplementary information Supplementary material related to this article can be found on-line at https://doi.org/.....

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