

Hot-s~~H~~ydrothermal spring inputs and climate drive dynamic shifts in microbial-archaeal communities in Lake Magadi, Kenya Rift Valley

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Abstract. The Methane Index (MI) is an organic geochemical index that uses isoprenoid glycerol dialkyl 15 glycerol tetraethers (GDGTs) as a proxy for methane cycling. Here, we report results from sediments in core MAG14-2A that spanning >700 almost <500 ka in Lake Magadi, Kenya. The deposits, which shows abrupt shifts between high and low MI values in through calcareous, tuffaceous and zeolitic silts the core. These shifts coincide with interbedded tuffaceous silt. Where tuffaceous silts are present The MI “switches off” (MI < 0.2); in contrast, where these silts are absent in the core, and on the MI increases (MI > 0.5) 20 through the core with -b- Bulk organic matter is-enriched in ¹³C in Magadi during “MI-off” periods (, with values of ~ -18‰) in the upper part of the core, whereas ¹³C is lower (and -22 to -25‰) in the lower parts of the lower portion sedimentary sequence. Evidence from *n*-alkanes and fatty acid methyl esters (FAMEs) support previous interpretations of an arid environment with a shallower lake where Thermoproteal (formerly Crenarchaeota) archaea thrive in a hot spring spring rich environment over 25 Euryarchaeota. Sediments deposited when the MI switches “on” showed $\delta^{13}\text{C}_{\text{OM}}$ values as low as -89.4 ‰, but most were within the range of -28 to -30‰, which is consistent with contributions from methanogens rather than methanotrophs. Thus, the likely source of these high MI values in Lake Magadi is methanogenic archaea. Our results show that hydrothermal inputs of bicarbonate-rich waters into Lake Magadi combined with further evaporative concentration cause a shift in the dominant archaeal 30 communities, alternating between two stable states.

1 Introduction

Life thrives in East African soda lakes (Schagerl, 2016) and has been the subject of modern studies of both prokaryotic and eukaryotic organisms, but few have studied their sediments over geologic timescales (Schagerl, 2016 and chapters therein). Soda lakes represent ~ 18,500 km² in East Africa (calculated from

35 values in Melack and MacIntyre, 2016). When compared to the three largest African freshwater lakes (lakes Victoria, Tanganyika, and Malawi), these soda lakes account for ~ 13% of the total lake-surface area in East Africa. A survey of microbial isolate diversity in East African lakes found evidence for cyanobacterial and archaeal primary producers with both oxygenic and anoxygenic phototrophs among the microbial population (Grant and Jones, 2016). Unique aerobic and anaerobic heterotrophs that use a
40 variety of electron donors, including sulfur, sulfate, nitrite, carbon dioxide, and methane, were also identified (Grant and Jones, 2016, and sources therein). ~~Near hot spring outflows, many~~ Many thermophilic archaea and bacteria isolates were also ~~classified observed near hot-spring outflows~~ (Grant and Jones, 2016).

45 Saline alkaline (soda) lakes in the East African Rift ~~Valley~~ often become stratified meromictic water bodies ~~of water~~ with a dense monimolimnion below a chemocline (Melack and MacIntyre, 2016). Oxygen rarely penetrates the monimolimnion waters, and as a result, anaerobic bacteria and archaea dominate the bottom waters and sediments. Remineralizing organic matter from the upper water column (mixolimnion) ~~feeds supports the~~ microbes and ~~in~~ generation of anaerobic oxidation of methane (AOM). Methane-
50 oxidizing microbes, specifically archaeal anaerobic methane-oxidizers (ANME), are coupled to sulfate-reducing bacteria in a microbial consortium (Boetius et al., 2000; Hinrichs and Boetius, 2002; Werne et al., 2004). ANME mediate ~~CH₄-methane~~ levels in freshwater and soda lakes and in modern oceanic systems, and account for approximately 90% of methane consumed through AOM (Egger et al., 2018). Rates of methane consumption differ by environment and type of ANME, with global freshwater systems
55 ranging from 1 to 1×10^5 nmol ~~CH₄-methane~~ L⁻¹ day⁻¹ consumed (Martinez-Cruz et al., 2018). Although soda lakes have been less studied, consumption rates as high as 1.6×10^4 nmol ~~CH₄-methane~~ L⁻¹ day⁻¹ have been observed in freshwater Lake Kivu (Roland et al., 2018). Tracking AOM over geologic time periods is important because methane release from tropical wetlands was concomitant with the end of

glacial conditions in Europe and is poorly constrained (DeMenocal et al., 2000; Riddell-Young et al.,
60 2023). Additionally, large methane releases might have been partly responsible for the Permo-Triassic
mass-extinction event (Berner, 2002).

Over geologic time, it is possible to gauge periods of increased methane oxidation, as shown by Zhang et
al. (2011) in oceanic systems by using a ratio of archaeal GDGT lipids (de Rosa et al., 1977; Langworthy,
65 1977). The ratio, as described by Zhang et al. (2011), is known as the methane index (MI), which uses
GDGTs produced predominantly by Euryarchaeal ANME. The MI has been used to discern
methanotrophy using the assumption that benthic methanotrophic Euryarchaeota preferentially produce
GDGT-1, -2, and -3, and that GDGTs crenarchaeol (cren) and crenarchaeol^{2'} (cren^{2'}) are thought to come
from Thaumarchaeota and Crenarchaeota, which are part of the TACK superphylum, typically found in
70 the upper water column (Sinninghe Damsté et al., 2002; Pitcher et al., 2009; Zhang et al., 2011).

Currently, the newly suggested names in the Genome Taxonomy Database for Thaumarchaeota and
Crenarchaeota are Nitrososphaerota and Thermoproteota, respectively (Oren and Garrity, 2021; Rinke et
al., 2021), which are used in this paper.
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Moreover, Kim and Zhang (2023) have shown a qualitative and quantitative relationship between the MI
and methanotrophy in deep time, namely from the late Oligocene to the early Miocene. Kim and Zhang
(2023) showed that the MI is applicable to AOM, with other biomarkers co-occurring in high-MI intervals
representative of not only the Group I consortium of anaerobic methanotrophs (ANME) that produce
80 GDGTs, but also of Group 2 and Group 3 consortia (ANME-2 and ANME-3 respectively). Until now, no
studies have directly applied the MI to sediments in African soda lakes despite evidence for AOM in
modern soda lakes. Combined with MI values, other methane-related indices are used here to interpret
methanogenesis and methanotrophy related to AOM. Previous studies have used GDGT-0 and GDGT-2
ratioed to the GDGT crenarchaeol value, which was originally thought to only be produced by mesophilic
85 Thermoproteota (Blaga et al., 2009; Weijers et al., 2012). Blaga et al. (2009) found that methanogens
predominantly produced GDGT-0, whereas Weijers et al. (2011) showed that methanotrophic archaea

predominantly produce GDGT-2. However, the optimum temperature is closer to 40-45 °C (Zhang et al., 2006). Blaga et al. (2009) found that methanogens predominantly produced GDGT-0, whereas Weijers et al. (2011) showed that methanotrophic archaea predominantly produce GDGT-2.

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Lake Magadi (Kenya) is a sulfate-limited lake, and therefore, methanogenesis and methanotrophy may co-occur without suppression of the higher energy yield of sulfate reduction (Nijaguna, 2006; Sorokin et

al., 2007; Deocampo and Renaut, 2016; Lameck et al., 2023). Here, we document evidence of methane cycling in Lake Magadi using archaeal isoprenoid ~~glycerol dialkyl glycerol tetraether~~ (GDGT) lipid

95 biomarkers. Environmental influences on archaeal community composition included precipitation/evaporation fluctuations and variations in hydrothermal activity, the latter often related to contemporary tectonics. This study leverages four methane-related indices: (1) the MI; (2) the %-GDGT-0/crenarchaeol; (3) %-GDGT-2/crenarchaeol; and (4) the ratio of isoprenoid GDGTs ($\frac{[2]}{[3]} / \frac{[2]}{[3]}$) (hereafter, $\frac{[2]}{[3]} / \frac{[2]}{[3]}$) to understand methane cycling in recent and ancient lacustrine sediments. Two

100 distinct communities were found using a combination of the MI and ratios of GDGT-0 and GDGT-2, normalized to crenarchaeol. Intervals of high methanotrophy, as evidenced by MI and %-GDGT-2/crenarchaeol, were related to an equally high proportion of methanogens, while in periods when crenarchaeol was dominant, the methane indices were low.

105 Environmental influences on microbial community composition included precipitation/evaporation fluctuations and variations in hydrothermal activity, the latter often related to contemporary tectonics. Leaf wax data from *n* alkanes was also used to understand changes in the surrounding environment at Lake Magadi. The average chain length of *n* alkanes (ACL₍₂₅₋₃₃₎) and the carbon preference index of both alkanes and fatty acid methyl esters (CPI_{alk} and CPI_{FA}) indicated proportionally higher input of C4
110 vegetation in an arid landscape. Methane indices were typically higher during periods of reduced hydrothermal activity, indicating more Euryarchaeal communities, whereas Thermoproteota communities thrived during periods of higher hydrothermal activity.

2 Materials and methods

2.1 Study locations and sampling

115 Modern Lake Magadi is a seasonally flooded, saline alkaline pan composed of bedded trona ($[\text{Na}_3(\text{CO}_3)(\text{HCO}_3)_2 \cdot 2\text{H}_2\text{O}]$) located in the southern Kenya Rift near the border with Tanzania (Baker, 1958; Eugster, 1980). Its elevation is approximately 600~~to~~⁺605 m above sea level (asl), and it has a maximum depth during the rainy ~~seasons~~ season of a few decimeters ~~up~~ to ~1 m (Fig. 1; Renaut and Owen, 2023).

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The modern lake is fed by ephemeral streams and alkaline hot springs (up to 86°C ~~at adjacent Nasikie Engida~~, distributed along faults around the shoreline (Baker, 1958; Crane, 1981; Allen et al., 1989; Renaut and Owen, 2023). Former high-level shorelines are preserved as coarse clastic sediments and ~~locally as~~ stromatolites around the lake. ~~These vary in age and record lakes of different depths and, with lake sediments of different ages, record deeper paleolakes that occupied the basin~~ during the Quaternary. ~~Other outcrops of eocene sediments relevant to this study are and can be correlated at different ages and are nominally known as the Oloronga Beds and the Green Beds. The chert-bearing Oloronga Beds in outcrop have been dated were deposited from between approximately ~800 to and 300 ka, with cores extending this back to 1000 k Ma (Owen et al., 2019). while the Green Beds outcrops, which include abundant chert and have been, were actively deposited precipitating chert at times between from 220 to and 70 ka variably dated between 191– and 40 ka (Behr and Röhricht, 2000; Owen et al., 2019) with cores suggesting a range off from 380 to 105 ka (Owen et al., 2019). More and more recently, the High Magadi Beds were being deposited between ~25 and 9 ka (Fairhead et al., 1972; Goetz and Hillaire-Marcel, 1992; Williamson et al., 1993; Behr and Röhricht, 2000; Owen et al., 2019; Reinhardt et al., 2019). Calcrete commonly deposits caps areas of the Oloronga Beds (Eugster, 1980), but fluvial where variable erosion locally occurred scourined those sediments, leading to missing facies prior to before the Green Beds being were laid down (Renaut and Owen, 2023).~~

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140 More recently, the stratigraphic terminology has been revised by Owen et al. (2025). They continued to recognise the Oloronga Beds below Lake Magadi under the same name but raised it to formation status

with dates of 1000–380 ka. In contrast, they renamed the Green Beds to the Oloika Formation (380–105 ka) and reassigned the Evaporite Series to the Magadi Formation (105–0 ka) and the High Magadi Beds to the High Magadi Formation. For example, Casanova (1986, 1987) and Casanova and Hillaire-Marcel reported a paleoshoreline at 956 m asl when the High Magadi Beds (Late Pleistocene–Early Holocene) were being deposited, which implies contemporary water depths of >40 m. However, these estimates are approximations due to localized uplift and erosion, introducing uncertainty (Owen et al., 2019; Renaut and Owen, 2023). Various ~~higher~~ lake levels have been suggested based on stromatolites (Casanova and Hillaire-Marcel, 1987) but some details are uncertain (Casanova and Hillaire-Marcel, 1987). Although Lake Magadi is situated near the equator, it lies in a rain shadow and consequently. Consequently, today it has a very large moisture deficit (2400 mm evaporation versus 500 mm precipitation annually; Damnati and Taieb, 1995).

Lake Magadi was cored as part of the *Hominin Sites and Paleolakes Drilling Project* (HSPDP) in June 2014 with the aim of furthering our understanding of the paleoenvironments in the East African Rift Valley and to better contextualize hominin remains and artifacts, and to understand possible environmental influences on hominin evolution and migration (Cohen et al., 2016). A 197.4-meter core (MAG-14-2A) was drilled in the northern end of Lake Magadi in June 2014 (1°51'5.76" S; 36°16'45.84" E; Owen et al., 2019). Approximately In total, 107.7 m of sediments were recovered, with a total core recovery of 55.4% (Cohen et al., 2016). Here, we use the age model from Owen et al. (2019). The core ranges from the modern evaporite–trona surface (0 ka) to the Magadi Trachyte basement, dated at ~1 Ma at the core site (Owen et al., 2019). Cores were sampled in 2016 during the initial core description at the Continental Scientific Drilling Facility (CSD, formerly LacCore) at the University of Minnesota, Minneapolis. Altogether, 61 samples, covering the period from 456 ka to 14.9 ka (Table 1/Table S1), were subsampled and freeze-dried from dark brown to black silty clay intervals in the core. Based on their color, These samples were expected to have a high total organic carbon that would yield the best results/highest quantity of biomarkers from for our study, organic geochemical studies.

Over the past million years, Lake Magadi has varied from ~~a swampy fresh water bodies to a large fresh~~
170 ~~to mildly saline lakes~~~~humid that was continuously lake, when the freshwater lake was~~ fed by rivers and groundwater~~continuously~~, to ~~the a-~~~~smaller hypersaline lakes bounded by the Magadi grabens that dried~~
~~to trona, tectonically restricted, saline alkaline pan pans~~ partly fed by hot springs (Owen et al., 2019;
Renaut and Owen, 2023). From 545 to 380 ka ~~the~~ Magadi ~~catchment progressively changed transitioned~~
175 to a more ~~pronounced~~ arid ~~system condition with the palaeolake~~ marked by abundant ~~calcareous~~, organic-
rich sediments (Owen et al., 2019). Periodic freshwater inundation occurred from 380 to 105 ka ~~into a~~
~~highly saline, alkaline lake that accumulated minor as marked by interspersed~~ calcite and magnesium-
rich calcite at ~~the~~ lake margins. ~~Ash that fell into this waterbody reacted to form e a variety~~ ~~vidence~~ of
zeolites ~~formed by trachytic volcanic glass reinteracting with alkaline water with anoxic, and~~ sulfate-
rich bottom~~-~~water brines ~~that were~~ subjected to microbial sulfate reduction (Owen et al., 2019; Deocampo
180 et al., 2022). The most recent phases of the lake -(105 to 0 ka) were more evaporatively enriched, with
abundant trona ~~and and~~ minor nahcolite. ~~Well-preserved diatoms in sediments deposited after ~500 ka~~
~~suggest very high aqueous silica in the palaeolake aqueous silica in order to explain the preservation of~~
~~their frustules under highly alkaline conditions, which may reflect strong evaporative concentration of~~
~~silica-rich hydrothermal inflows as well as, and~~ metals (i.e., Br, Pb, Zn, Sb, Ag), indicating increased~~that~~
185 ~~might be related to periods of increased hydrothermal inflow, especially from ca. 20 to 10 ka~~ (Owen et
al., 2019).

2.2 Leaf wax and bulk organic preparation and analysis

2.2.1 Lipid extraction

190 To obtain a total lipid extract (TLE), 61 samples from Lake Magadi were freeze dried and homogenized
and ca. 5–10 g of sediment were ultrasonically extracted with 2:1 DCM:MeOH. The TLE for each sample
was treated with activated copper shot to remove elemental sulfur. The TLEs were then separated into
three fractions (apolar (AP), polar one (P1), and polar two (P2)) using activated alumina via ~~a~~ short
195 column chromatography. The AP fraction was eluted with 4 mL of 9:1 Hexane (Hex):DCM (v/v), the P1
fraction with 4 ml 1:1 DCM:MeOH, and the P2 fraction with 4 ml MeOH. The P1 fractions were dried

down and re-dissolved in 99:1 Hex:Isopropanol (IPA) (v/v) and filtered through a 0.45 μm 4 mm diameter PTFE filter prior to GDGT analysis.

2.2.2 Leaf wax preparation and analysis

~~Leaf waxes (i.e., free fatty acids and *n* alkanes) were identified and quantified in the same intervals as 200 GDGTs in the Lake Magadi core. The TLE from the ultrasonic extraction contained free fatty acids (FFAs), which were methylated using BF_3 in a methanol solution, converting the FFAs to fatty acid methyl esters (FAMEs). The apolar *n* alkane fraction did not require methylation to be run. Compound concentrations were determined via gas chromatography mass spectrometry (GC-MS) using a Thermo Scientific Trace 1310 GC, equipped with an Agilent DB-5 column (30 m x 0.320 mm, 0.25 μm film) in 205 tandem with a FID and a Thermo Scientific ISQ-QD Single Quadrupole Mass Spectrometer. Samples were run separately to avoid loss from the FID. The inlet was operated in splitless mode for the FAMEs at a temperature of 250 $^{\circ}\text{C}$. Column flow was set to 1.5 ml min^{-1} with an initial oven temperature of 70 $^{\circ}\text{C}$, which was held for 1 minute and then ramped to 130 $^{\circ}\text{C}$ over 6 min at a rate of 10 $^{\circ}\text{C min}^{-1}$, and then ramped to 320 $^{\circ}\text{C}$ over 57.5 min at a step of 4 $^{\circ}\text{C min}^{-1}$ and held at 320 $^{\circ}\text{C}$ for 3 min. The column carrier 210 gas was He and the gas used in the FID was a mixture of air (350 ml min^{-1}), He (20 ml min^{-1}), and H₂ (35 ml min^{-1}). Quantification of compounds was completed using 5 α -androstane as an internal standard and compounds were identified on the FID by comparing to relative retention times with a FAME standard.~~

2.2.24 Bulk organic $\delta^{13}\text{C}$ analysis

~~Samples were subsampled from the same intervals as organic biomarkers for bulk organic carbon isotope 215 analysis. Powdered sediment samples were weighed ~~out~~ in silver capsules and carbonates were removed by adding 5% HCl in four-hour increments. Samples were analyzed on a Costech Elemental Analyzer coupled to a ThermoFinnigan Delta V Plus isotope ratio monitoring mass spectrometer (IRMS). Samples are reported as per mil (‰) deviations from the Vienna Pee Dee Belemnite (VPDB) standard in conventional delta notation.~~

220 **2.3 GDGT preparation and analysis**

2.3.1 GDGT analysis

Polar samples from Lake Magadi were analyzed for core lipid isoprenoid glycerol dialkyl glycerol tetraethers (iso-GDGTs) at the University of Massachusetts Amherst on an Agilent 1260 series high performance liquid chromatograph (HPLC; [Fig. S1](#)) in tandem with an Agilent 6120 series single quadrupole mass selective detector (MSD). Compounds were ionized using atmospheric pressure chemical ionization (APCI). The columns used for GDGT separation were a pre-column guard followed by two ultra-high performance liquid chromatography (UHPLC) silica columns (BEH HILIC, 2.1x150 mm, 1.7 μ m, Waters) connected in series and kept at 30 °C. Elution solvents followed Hopmans et al. (2016) using a flow rate of 0.2 mL min⁻¹. Two solvent mixtures, hexane (A) and 9:1 Hex:IPA (B), [and](#) were eluted isocratically for 25 minutes with 18% B, a linear gradient to 35% B in 25 minutes, a second linear gradient to 100% B in 30 minutes.

2.3.2 GDGT indices

Several different ratios based on the relative abundance of different isoprenoid GDGTs have been developed to determine their source(s). The methane index (MI) is defined by Zhang et al. (2011) and is calculated as in Eq. (1):

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

MI values range between 0 and 1 with values > 0.5 considered to be derived from methanotrophic communities and values < 0.3 considered normal sedimentary conditions (Zhang et al., 2011). These proposed ranges from Zhang et al. (2011) were derived from GDGTs found in marine sediments, so the cutoff values for methanotrophy may differ [from](#) [in](#) lacustrine sediments, [particularly those in saline, alkaline environments, even those that are saline and alkaline.](#)

245 The ratio of GDGT-2 / crenarchaeol (%-GDGT-2-/cren) also indicates methanotrophy (values > 0.2), specifically methanotrophy associated with sulfate-methane transition zones (Weijers et al., 2011). These values were normalized and converted to percentages so that the numbers produced could be contextualized with the other indices used (Eq. 2). As a result, %-GDGT-2-/cren contributions greater than 33% will be considered methanotrophic signals.

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$$\%GDGT-\frac{2}{cren} = \frac{[GDGT-2]}{[GDGT-2]+[cren]} * 100\% \quad (2)$$

Methanogenic inputs are calculated similarly to Eq. 2 above using GDGT-0 in place of GDGT-2. Blaga 255 et al. (2009) found that values of GDGT-0 / (GDGT-0 + cren) > 2 are associated with methanogenic archaeal communities in a study of freshwater lakes. Similarly, in a study of Eocene-~~aged~~ marine sediments, Inglis et al. (2015) normalized the equation and converted it to a percentage, a convention we follow (Eq. 3). They found that contributions from methanogens were indicated by values greater than 67%.

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$$\%GDGT-\frac{0}{cren} = \frac{[GDGT-0]}{[GDGT-0]+[cren]} * 100\% \quad (3)$$

The GDGT-2 / GDGT-3 ([\[2\]](#)/[\[3\]](#)[\[2\]](#)/[\[3\]](#)) index was used here to describe both mesophilic environments 265 as well as environments with high MI values. [Rattanasriampaipong et al. \(2022\) found that differences in \[2\]/\[3\] are linked to distinct archaeal communities whereby low values of \[2\]/\[3\] \(ca. 0.55\) are observed in thermophilic cultures while elevated values are indicative of hot spring mats \(ca. 1.00\), shallow aerobic ammonia-oxidizing archaea \(AOA; ca. 1.16\), or archaea in suspended particulate matter \(ca. 2.52\)](#). This is the same version described in Rattanasriampaipong et al. (2022) (Eq. 4).

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$$[2]/[3] = [GDGT-2]/[GDGT-3] \quad (4)$$

(5)

275 Carbon preference indices (CPI) were calculated for both the free fatty acid methyl esters (CPI_{FAs}; Eq. 6) and *n* alkanes (CPI_{alk}; Eq. 7) to determine the source of leaf waxes and possible degradation of each wax type. The equation for the CPI_{alk} is based on the work of Marzi et al. (1993) with an adaptation for the CPI_{FAs} from Naraoka and Ishiwatari (1999). Average chain length (ACL₍₂₅₋₃₃₎; Eq. 8) for *n* alkane leaf waxes with chain lengths of 25–33 carbons was also calculated based on the equation from Freeman and
280 Paneost (2014).

$$CPI_{FAs} = 2 * \frac{\sum \text{even } C_{26} \text{ to } C_{28}}{(\sum \text{odd } C_{19} \text{ to } C_{27}) + (\sum \text{odd } C_{24} \text{ to } C_{29})} \quad (6)$$

$$CPI_{alk} = \frac{1}{2} * \frac{(\sum \text{odd } C_{23} \text{ to } C_{31}) + (\sum \text{odd } C_{25} \text{ to } C_{33})}{\sum \text{even } C_{24} \text{ to } C_{32}} \quad (7)$$

$$ACL_{(25-33)} = \frac{1}{2} * \frac{(25 * [C_{25}] + 27 * [C_{27}] + 29 * [C_{29}] + 31 * [C_{31}] + 33 * [C_{33}])}{([C_{25}] + [C_{27}] + [C_{29}] + [C_{31}] + [C_{33}])} \quad (8)$$

290 **2.4 Bulk geochemistry**

Bulk geochemical data and core descriptions from both Owen et al. (2019, 2024) and Owen et al. (2024) were also used to interpret hot spring influences in the intervals of focus (i.e., Intervals 1, 3, and 5). They attributed REE anomalies to increased lake alkalinity, that which reflected increased evaporation and the development of highly saline, alkaline lakes and possibly increased hydrothermal/fluvial inflow ratios.
295 All statistical analyses were performed using the GraphPad Prism 10[®] software (<https://www.graphpad.com/>). Only the necessary data to determine relationships between the bulk

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geochemistry of rare earth elements (REEs; La, Ce, Nd, Sm, Eu, Yb, Lu) and methane indices (MI, %GDGT-0/cren, %GDGT-2/cren, and [2]/[3]) were imported. Prior to performing a principal component analysis (PCA) or correlation matrix, the data needed to be tested for normality. To determine whether 300 the data were normally distributed, the built-in “Normality and Lognormality Tests” function in GraphPad[®] which yielded lognormal distributions of each dataset. The data were found to be non-normally distributed.

2.4.1 Principal component analysis (PCA)

For the PCA, the imported data were analyzed using the built-in PCA function in GraphPad[®]. The data 305 were standardized, which scaled the data to have a mean of 0 and a standard deviation of 1. The principal components were selected based on their eigenvalues using the Kaiser Rule, which selects eigenvalues greater than 1.0. Principal components 1 and 2 explained 58.0% of the variance in the data.

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2.4.2 Correlation matrix

The correlation matrix was performed using the built-in function in GraphPad[®]. Because the data were 310 non-normally distributed, the nonparametric Spearman correlation was chosen over the Pearson correlation. An r value was computed for every pair of Y datasets without using the default two-tailed option at a confidence interval 95% confidence 95% interval.

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3 Results

3.1 GDGT lipid variability

315 Samples are split into six intervals (1-6) based partly on their fractional abundances of GDGT-0 and cren as well as their MI values: (1) 35.6~~768~~ to 32.61 m; ca. 17.7 to 14.9 ka, (2) 67.81~~5~~ to 43.51 m; ca. 129 to 38.9 ka (3) 86.06~~5~~ to 70.78 m; 197 to 149 ka, (4) 96.38 to 94.91~~5~~ m; ca. 318 to 315 ka, (5) 104.10 to 103.16 m; ca. 324 to 323 ka, and (6) 130.21 to 119.64 m; ca. 456 to 391 ka (Table 4Table S1).

320 In each of the intervals of the core, Fractional abundances of GDGTs along with sample depth and age are shown in Table 4Table S1 for each interval of the MAG 14.2A core demarcated by the respective

sampling interval. GDGT 0 to 3 are present in addition to crenarchaeol and its regioisomer (hereafter, eren and cren'). Downcore GDGT ratios (MI, %GDGT 2 / cren, % GDGT 0 / cren and [2] / [3]) are summarized along with values of $\delta^{13}\text{C}_{\text{OM}}$ in Fig. 3 and are plotted against age (Fig. 3 and Table 1).

325 MI, %-GDGT-0/~~+~~cren, and %-GDGT-2/~~+~~cren values oscillate between high and low values, changing abruptly ~~throughout the core from one interval to the next~~. The methanotrophic (%-GDGT-2/~~+~~cren) and the methanogenic (%-GDGT-0/~~+~~cren) indices track similarly to MI values; that is, when values of MI are high, so are the other two indices. It should be noted that there are some large gaps in sampling between intervals in the core ~~as a result of due to~~ our sampling 330 regime (i.e. targeting intervals with high apparent organic matter based on darker silty matrix).

Interval 1 is characterized by a higher proportion of cren and lower overall index values. The %-GDGT-0/~~+~~cren index averages 36.3 % ($\pm 0.09\%$) in this interval while the %-GDGT-2/~~+~~cren index averages 10.6 % ($\pm 0.05\%$; Fig. 3|Fig. 2). The MI in this interval is correspondingly low with an average of 0.25

335 (± 0.07), well below the MI = 0.5 cutoff range for methanotroph-impacted communities. As such, this interval could be used for the [2] / [3]|[2] / [3] index; values averaged 2.1 (± 1.02). Interval 2 has much higher values for each of these indices, where the average %-GDGT-0/~~+~~cren = 99.3 % ($\pm 0\%$) and the average %-GDGT-2/~~+~~cren = 93.6 % ($\pm 0.04\%$). MI values in Interval 2 are also high with an average of 0.96 (± 0.02). Of note, there is a large gap where no measurements were taken from 43.55 to 46.6815 m 340 (~9.7 kyr) as well as from 50.36 to 58.74 m (~32.5 kyr). Interval 3 averages for %-GDGT-0 / cren and % GDGT-2/~~+~~cren are 54.3 ($\pm 0.27\%$) and 20.4 % ($\pm 0.26\%$), respectively. However, there is one anomalously high value at 77.32 m with %-GDGT-0/~~+~~cren and %-GDGT-2/~~+~~cren values at 99.6 and 93.8 % and an MI = 0.96. Excluding the high index value, the averages were lowered to 48.6 ($\pm 0.22\%$) and 11.2 % ($\pm 0.05\%$) for the %-GDGT-0/~~+~~cren and %-GDGT-2/~~+~~cren values and the MI average was 345 lowered from 0.33 (± 0.23) to 0.26 (± 0.06). With the exclusion of 77.32 m, the [2] / [3]|[2] / [3] index averaged 1.5 (± 0.80) in this interval, lower than Interval 1. Interval 4 is characterized by high index values, with a similarly abrupt shift from low values. Averages of the %-GDGT-0/~~+~~cren and %-GDGT-2/~~+~~cren are 98.1 ($\pm 0.04\%$) and 88.2 % ($\pm 7.41\%$) and an average MI of 0.92 (± 0.04); these average index values are similarly high as compared to Intervals 2 and 6.2. Interval 5 is a shift to lower overall index values

350 with averages of %-GDGT-0-+-cren and %-GDGT-2-+-cren at 40.1 ($\pm 0.17\%$) and 9.2 % ($\pm 0.03\%$) and an average MI of 0.22 (± 0.05). Finally, Interval 6 shows a period in the core with high index values throughout. Averages of %-GDGT-0-+-cren, %-GDGT-2-+-cren, and MI are 97.6 % ($\pm 0.03\%$), 89.4 % ($\pm 0.08\%$), and 0.95 (± 0.05), respectively.

3.2 Bulk $\delta^{13}\text{C}_{\text{OM}}$ values

355 ~~Values of bulk $\delta^{13}\text{C}_{\text{OM}}$ values are shown in Fig. 3 and Table 1~~—Table S1—Bulk $\delta^{13}\text{C}_{\text{OM}}$ values follow a similar pattern to the indices described in section 3.1, that is the values oscillate between high and low values between intervals. Samples in Interval 1 ranged from -21.9 to -16.8 ‰ and had an average $\delta^{13}\text{C}_{\text{OM}}$ value of -18.4 ‰ with respect to VPDB. Interval 2 samples had the most ^{13}C -depleted values in all sampled intervals, ranging from -89.4 to -24.7 ‰ with an average of -35.1 ‰ and excluding the three outlier values (-48.1, -64.2, and -89.4 ‰), the Interval 2 average was -28.2 ‰. In Interval 3, the $\delta^{13}\text{C}_{\text{OM}}$ had a narrower range from -24.4 to -21.4 ‰ and an average of -22.5 ‰. A lighter signal from Interval 4 yielded a narrow range of values from -27.0 to -25.4 ‰ averaging -26.0 ‰. Interval 5 had slightly heavier values ranging from -25.0 to -18.1 ‰ with an average of -22.1 ‰. Lastly, Interval 6 had depleted $\delta^{13}\text{C}_{\text{OM}}$ values similar to intervals 2 and 4, with a range of -28.2 to -22.1 ‰ and an average of -25.2 ‰. Analytical reproducibility of duplicate runs was better than $\pm 0.15\%$ VPDB.

3.3 Leaf wax distributions

365 ~~Only 15 samples were analyzed for FAMEs as they were part of pilot sampling and were not further expanded due to time constraints. Analyzed FAMEs exhibit an even over odd predominance, which is diagnostic of a primarily terrestrial source. $\text{C}_{16:0}$ to $\text{C}_{34:0}$ FAMEs were present in the 15 samples analyzed with $\text{C}_{34:0}$ and $\text{C}_{24:0}$ averaging the highest percent abundance at 16.5 and 11.5 %, respectively, followed by $\text{C}_{17:0}$ and $\text{C}_{16:0}$ with averages of 10.8 and 10.3 %, respectively (Table 2). The CPI_F ranged from a minimum of 1.1 to a max of 8.5 with an average of 3.82 in the 15 samples, indicating an overall terrestrial source of FAMEs.~~

375 The *n* alkanes represent a more robust sampling interval (n=56) that closely matches the total number of
GDGT sampling intervals (n=58). The *n* alkanes analyzed showed an odd over even predominance, with
higher overall $ACL_{(25-33)}$ values indicating primarily terrestrial sources and no evidence for *n* alkanes C_{17}
to C_{22} . Overall, C_{31} and C_{33} *n* alkanes accounted for 54.4 % of all *n* alkanes in every interval and were
particularly dominant in Intervals 2 through 6, while In contrast, *n* alkanes in Interval 1 were more evenly
380 distributed, though C_{31} is still more dominant, averaging 14.5%. The average value for the CPI_{alk} in
Interval 1 was 2.0 and the $ACL_{(25-33)}$ was 29.2. Intervals 2, 3, and 4 were much higher than Interval 1,
with average values of the CPI_{alk} at 6.7, 7.0, and 6.1, respectively, while the values for the $ACL_{(25-33)}$ were
30.7, 30.9, and 30.3, respectively. Intervals 5 and 6 had similar values with the CPI_{alk} averaging 4.6 and
4.2, respectively, and the $ACL_{(25-33)}$ averaging 29.7 and 30.1 respectively.

385 In order To understand proportions of sulfate reducing bacteria (SRB), we focus here on the short chain
length FAMEs. The short chain $C_{15:0}$ FAMEs were not observed in Magadi sediments, possibly due to
degradation of the FAMEs smaller than $C_{16:0}$. These short chain FAMEs are diagnostic of bacteria with
different respiratory pathways and are useful in understanding bacterial contributions in sedimentary
390 environments (Cho and Salton, 1966; Parkes and Taylor, 1983). Among bacterially diagnostic short chain
FAMEs, $C_{15:0}$, $C_{15:0}\text{ iso}$, $C_{17:0}$, and $C_{17:0}\text{ iso}$ are used to identify SRB in sediments (Boon et al., 1977; Ueki
and Suto, 1979; Parkes and Taylor, 1983). In Magadi, only 15 samples were analyzed for FAMEs, and of
those 15 samples, only 22.8% of samples in Magadi contained $C_{17:0}$ fatty acids indicative of SRB. In
Interval 1, only one sample contained $C_{17:0}$ at ca. 14.9 ka with a concentration of 6.2 $\text{ng g}^{-1}\text{ sed}^{-1}$. Interval
395 2 had higher concentrations of $C_{17:0}$ with a range from 8.8 to 469.0 $\text{ng g}^{-1}\text{ sed}^{-1}$ and an average of 97.1 $\text{ng g}^{-1}\text{ sed}^{-1}$. Similar to interval 1, interval 3 had only one sample with $C_{17:0}$ present at 390.8 $\text{ng g}^{-1}\text{ sed}^{-1}$, coinciding
with the high MI value in that interval at ca. 185 ka. Interval 4 $C_{17:0}$ values were similarly sparse, only
having two values at 8.3 and 73.9 $\text{ng g}^{-1}\text{ sediment extracted}$. In Interval 6, three values of $C_{17:0}$ were
recorded with values of 5.9, 31.5, and 359.6 $\text{ng g}^{-1}\text{ sediment extracted}$.

400 **3.34 Bulk geochemistry**

~~Bulk geochemical data and core descriptions from both Owen et al. (2019) and Owen et al. (2024) were also used to interpret hot spring influences in the intervals of focus (i.e., Intervals 1, 2, and 5). Both a PCAPCA and correlation matrix were performed using the of the MI, Ca/Na, %GDGT-0/cren, %GDGT-2/cren, and [2]/[3][2]/[3] compared to compare to to the REEs rare earth elements (REEs) La, Ce, Nd, Sm, Eu, Tb, Yb, and Lu (Fig. 5Fig. 3). Increased values of REEs are characteristic of sodic systems influenced by hydrothermal springs, namely Mono Lake in California and this system (Johannesson and Lyons, 1994; Owen et al., 2019).~~

A PCA (Fig. 5Fig. 4a) and non-parametric Spearman correlation matrix (Fig. 5Fig. 4b) were performed 410 to quantify the relationship between REEs, MI, Ca/Na and [2]/[3][2]/[3]. The PCA showed that ~~MI~~Ca/Na and [2]/[3] loaded positively on PC1 and PC2 and, each of the methane indices loaded positively on PC1 and negatively on PC2. ~~while The REEs La, Yb, Tb, Eu, and SMLa loaded positively on PC2 and negatively on PC1 while-and the REEs Ce, Nd, and La~~ Ce, Nd, Sm, Eu, Yb, and Lu loaded negatively on PCs 1 and 2. This indicates a negative relationship between ~~the negatively loaded REEs and a high~~ 415 ~~MI~~Ca/Na. and [2]/[3]. Similarly, the correlation matrix of REEs and methane indices, MI, and [2]/[3] showed a negative relationship between each index and REE, except for the relationship of [2]/[3][2]/[3] and Nd ($r=0.02$) and %GDGT-2/crem ($r=-0.04$), which showed no observable linear trend ($r=-0.02$). The REEs and methane indices did not load on the same PC axis showing that there was also not a nonlinear trend associated with the REEs and methane indices.

420 **4 Discussion****4.1 Lake Magadi ~~microbial-archaeal~~ community shifts**

The abrupt changes in isoprenoid GDGT-based indices in the sediment record of Lake Magadi indicate 425 shifts in the archaeal communities present (Fig. 3Fig. 2). Shifts between two distinct communities were inferred using a combination of the Methane Index (MI) and ratios of GDGT-0 and GDGT-2 normalized to crenarchaeol (Eqs. 2 and 3; ~~GDGT structures in Fig. 2~~). We denote these shifts as either “MI-on

periods”, characterized by $MI > 0.5$ during intervals 2, 4, and 6, and “MI-off periods”, characterized by $MI < 0.5$ during intervals 1, 3, and 5. Oscillations between these two environmental states are discussed in detail in the following sections.

4.1.1 MI-on periods

430 In Lake Magadi, during the MI-on periods (Fig. 3Fig. 2; Intervals 2, 4, and 6), the MI is persistently greater than 0.83 and displays more ^{13}C -depleted $\delta^{13}\text{COM}$ values compared to MI-off periods, indicating periods of enhanced methane cycling. AOM is a likely mode of methane cycling in Interval 6 as well as parts of Interval 2 because SRB and AOM archaea live in a consortium together at the sulfate methane transition zone, or SMTZ (Boetius et al., 2000; Hinrichs and Boetius, 2002; Werne et al., 2004).~~– and~~
435 ~~biomarkers of SRB (FAMEs) were identified in those intervals.~~ Thus, in intervals of the Magadi core where a SMTZ is suspected, such as in parts of Interval 2 and most of Intervals 4 and 6, there should be an increase in indices related to methanotrophy such as high MI and %-GDGT-2/~~+~~cren (Weijers et al., 2011). Additionally, whereas methanogens and methanotrophs appear to be present in a consortium based on both the methane indices as well as bulk $\delta^{13}\text{COM}$, the majority of the contributions are coming from
440 methanogens, as seen in the ternary plot in Fig. 6Fig. 5. This may seem counter-intuitive as the MI has been typically used to describe samples exhibiting a high predilection towards methanotrophy, but a high MI value does not necessarily exclude methanogenesis and conversely neither does a low MI, rather the low MI value suggests a predominance of Thermoproteota over Euryarchaeota (Zhang et al., 2011). High
445 %-GDGT-0/~~+~~cren and %-GDGT-2/~~+~~cren index values in Intervals 2, 4, and 6 (Fig. 3Fig. 2) show that methanogenesis is co-occurring with AOM. The [2]/~~+~~[3] index is also useful in understanding the proportion of methanotrophs in sediments, even in intervals with high MI values like those discussed herein (Table 4Table S1; Fig. 3Fig. 2). Values of the GDGT [2]/~~+~~[3] ratio track nearly identically to the MI values (Fig. 3Fig. 2), indicating that the MI is influenced by GDGT-2, which is characteristic of methanotrophs (Pancost et al., 2001; Schouten et al., 2003; Zhang et al., 2011).

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Typically, methanogenesis in sulfate-rich systems is suppressed in favor of sulfate reduction caused by competition for both H_2 and organic substrates (Fazi et al., 2021; Sorokin et al., 2015). However, reports

of methanogenesis co-occurring with SRB have been noted when methanogens are using non-competitive substrates such as methanol, or when sulfate levels are low (Oremland et al., 1982; Giani et al., 1984
455 Hoehler et al., 2001; Bebout et al., 2004; Arp et al., 2008, 2012; Jahnke et al., 2008; Smith et al., 2008; Robertson et al., 2009). Furthermore, pyrite ~~nodules~~ ~~cubes~~ are common and scattered throughout ~~occur in~~
most of the intervals where high index values are observed, indicating that there was a substrate for SRB, though it may have been in low concentration (Table 1~~Table S1~~). SRB can also be traced with FAMEs, specifically odd, short-chained FAMEs such as C_{15:0} and C_{17:0} in addition to their iso- and ante-iso forms
460 (Boon et al., 1977; Ueki and Suto, 1979; Parkes and Taylor, 1983). The high relative abundance of C_{17:0} (Tables 1 and 2) in the same intervals characterized by high index values suggests that methanotrophy is occurring in these sediment intervals. Thus, the combined evidence of higher FAMEs and pyrite ~~nodules~~ in intervals with high GDGT-based indices (e.g. MI, %-GDGT-0/cren, and %-GDGT-2/cren [2]/[3]) indicates the presence of a SMTZ that supports AOM with the co-occurrence of methanogenesis.

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However, when looking at Fig. 6~~Figure 4~~, it appears that GDGT-0 is the dominant GDGT compared to GDGT-2 and cren, indicating that this interval is likely methanogen-dominant rather than ANME dominant. Interval 2 (Figs. 3 and 5) of the Magadi core appears to be more influenced by methanogenesis than AOM, resulting from a more prevalent %-%GDGT-0/cren signal accompanied by a high %-GDGT-
470 2/cren signal, high [2]/[3] ratios, and a more ¹³C-depleted bulk δ¹³COM signal (average = -35.1 ‰; median = -28.6 ‰). Values of bulk δ¹³COM are similarly ¹³C-depleted in AOM-dominant Euryarchaeotal systems ranging from active mud volcanoes (~-27 ‰; ANME-1), a Danish freshwater lake (average ~-29.7 ‰; ANME-2), and the Sea of Galilee (~-30 ‰; ANME-2) in Israel (Lee et al., 2018; Norði et al., 2013; Sivan et al., 2011). However, when looking at Fig. 6, it appears that GDGT-0 is the dominant
475 GDGT compared to GDGT-2 and cren, indicating that this interval is likely methanogen-dominant rather than ANME dominant. At points where the bulk δ¹³COM values are at their lowest (e.g., -89 ‰), they are accompanied by a lower %-GDGT-2/cren at ca. 95 % and an elevated %GDGT-0/cren at > 99.5 %. This is in line with the literature as Summons et al. (1998) reported values between -53.4 and -48.7 ‰ in the total lipid extract of methylotrophic methanogens using non-competitive substrates in anoxic
480 hypersaline environments. Furthermore, as these waters are typically sulfate-limited, it is likely

that acetoclastic and/or hydrogenotrophic methanogenesis is acetoclastic and/or hydrogenotrophic methanogenesis is likely dominant when evidence for SRB is lacking (i.e., pyrite, $\text{C}_{17:0}$ FAMEs). Zhuang et al. (2016) performed compound-specific isotope analysis on several archaeol compounds from the Orca Basin and found archaeol and hydroxyarchaeol using H_2 or CO_2 (diagnostic of 485 methanogens and methanotrophs) were relatively depleted (ca. -80 to -60 ‰) compared to the bulk $^{13}\text{C}_{\text{OM}}$ (ca. -22 ‰). Zhuang et al. (2016) concluded that acetoclastic and/or hydrogenotrophic methanogenesis was unlikely due to high SO_4^{2-} concentrations in the Orca Basin, which may be the case in Lake Magadi. In Interval 2, there is no evidence of visible pyrite cubes and we did not have a priority at the time of 490 sampling to check a thin section of each sample for smaller pyrite aggregates and a limited number of samples with $\text{C}_{17:0}$ FAMEs, which. This indicates that there may be other Euryarchaeotal communities with other Euryarchaeotal communities may have different forms of AOM occurring in the sediments. These other forms of AOM include nitrate/nitrite reduction and iron coupled to AOM (in't Zandt et al., 2018). This is further bolstered by the evidence outlined by Kim and Zhang (2023) that not only 495 quantitatively linked AOM to high MI values, but also to non-Group I ANME Euryarchaea because other non-GDGT producing ANME (e.g. ANME-2 and ANME-3) were shown to co-exist with Group I ANME. In the intervals that are missing pyrite (i.e., most of Interval 2; see Table 4 Table S1; Ferland, 2017), the pyrite nodules may have either been too small to see with the naked eye, or the excess H_2S could have 500 been incorporated into the kerogen by reacting with labile organic matter. As for the dearth of FAMEs observed, save for the samples at 58.80 and 58.74 m, there is limited evidence for sulfate reduction. From 59.40 to 58.80 m, values of the bulk $\delta^{13}\text{C}_{\text{OM}}$ dip as low as -89.4 ‰ (Table 4 Table S1; Fig. 3 Fig. 2), which aligns well with methanogenic archaeal biomass (Norði et al., 2013). However, as discussed above there is likely acetoclastic and/or hydrogenotrophic methanogenesis co-occurring in these high index intervals and is likely the dominant process where sulfate-dependent AOM is absent, and the sulfate-dependent AOM is likely replaced by a coupling to either nitrate/nitrite or iron reduction.

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Samples in Interval 4 (Table 4 Table S1) of the Magadi core have high index values, but no evidence for sulfate-dependent methanotrophy except for high MI values. These intervals are thus interpreted as being methanogenic, rather than methanotrophic. It is interesting to note that the abundance

of pyrite in the four samples with low MI values (Table 1 Table S1; 104.10 to 103.16 m), indicating sulfate reduction not linked to AOM. This is not observed in any other location level of the core and a hypothesized series of reactions is described below, which may be linked to an abundance of SRB, anaerobic ammonium oxidizing (anammox) bacteria, and Thermoproteota (AOA aerobic ammonia oxidizing archaea, AOA) in the overlying water column. Due to periodic influxes of freshwater in Magadi, in addition to a permanent meromixis present in virtually almost all samples post 380 ka, the water column would have been oxic in the upper portion and anoxic below the chemocline. Freshwater pulses would have also brought nutrients to the lake such as ammonia (NH_4^+) and sulfate (SO_4^{2-}). The oxic portion of the water column would have supported microaerophilic AOA that oxidize NH_4^+ to nitrite (NO_2^-), which is then transported to the anoxic part of the water column (Straka et al., 2019). Here, anammox bacteria are using excess NH_4^+ and the NO_2^- from the AOA and converting these to N_2 . Excess SO_4^{2-} is simultaneously being used by SRB, creating HS^- that is reacting with iron species in the sediments and being buried as pyrite. Ladderane lipids characteristic of marine anammox bacteria (Jetten et al., 2009) were not studied in Magadi sediments. However, there is both 16S rRNA and lipid evidence for the production of ladderanes in hot springs in the western United States suggesting that anammox bacteria can persist in hot spring environments (Jaeschke et al., 2009). Additionally, Kambura et al. (2016) found evidence for *Planctomyces* in both microbial mat and water samples surrounding the hot springs of Lake Magadi, lending credence to the hypothesis of AOA persisting in Lake Magadi. Without other lines of evidence, however, these are simply hypothesized hypothetical reactions for explaining excess pyrite in the sediments without accompanying MI values, yet. Nonetheless, this explanation has some merit because of the high relative abundance of both crenarchaeol and cren'.

In nearly all of Interval 6 (Table 1 Table S1), there is evidence for a higher proportion of methanotrophic archaea from 128.74 to 119.64 m (increased %-GDGT-2/cren and [2]/[3]) and methanogenesis in the intervals from 130.21 to 129.77 m (Table 1 Table S1; higher %-GDGT-0/cren compared to %-GDGT-2/cren). Samples from 123.43 to 119.64 (Table 1 Table S1) are of note because the [2]/[3] values are lower than the MI values whereas every other MI and [2]/[3]/[2]/[3] values aligned nearly 1:1. This is likely due to GDGT-2 not being the dominant control of the MI and while both %-GDGT-0/cren and %

GDGT-2/cren are equally high, there may be other factors in the water column exporting GDGT-2 to the sediments, possibly from deep-dwelling Group I.1b Thermoproteota, although this is unlikely due to limitations of depth (Taylor et al., 2013). The 656 m paleoshoreline reported by Casanova (1986, 1987) and Casanova and Hillaire-Marcel (1987) would imply a maximum water depth of ~ 50 m during the Late Pleistocene (African Humid Period: AHP) based on present topography. However, sedimentary evidence for such a high paleoshoreline is not seen throughout the Magadi Basin. ~~E~~arlier water depths are also unclear because accommodation space was changing as the axial rift developed with faulting and subsidence (Owen et al., 2024). This is not deep enough (> 1 km) to support ~~the~~ Group I.1b Thermoproteota per the constraints outlined in Taylor et al. (2013).

4.1.2 MI-off periods

In intervals characterized by low MI, %-GDGT-0/cren and %-GDGT-2/cren values (Fig. 3Fig. 2; MI-off intervals outlined in green checkered patterns~~are odd numbers highlighted in blue~~), the $\delta^{13}\text{C}_{\text{OM}}$ values are ^{13}C -enriched relative to those intervals characterized by higher index values (MI-on: ~~3~~(Fig. 3Fig. 2)).

Since the methane cycling indices (%-GDGT-0/cren and %-GDGT-2/cren) are both predominantly influenced by the availability of crenarchaeol, MI-off periods are marked by ~~an~~ increased production in crenarchaeol. Typically, crenarchaeol is produced in open ocean systems, freshwater lakes, and soils by the mesotrophic aerobic ammonium oxidizing phylum Nitrosphaerota. However, they can also be found in other environments, such as hot spring mats in Thermoproteota (Pearson et al., 2004, 2008; Schouten et al., 2013). As Thermoproteota require oxygen to oxidize ammonium to nitrate, the increased presence of crenarchaeol in the MI-off intervals, therefore, suggests periods ~~of time during which~~when conditions were more oxic, at least in the upper water column.

The MI-off periods in Lake Magadi are the core intervals where there is an increase in the relative abundance of crenarchaeol in the sediments, driving the MI below the 0.5 threshold that defines methanotrophy (Zhang et al., 2011). The low MI values are accompanied by equally low values in the other indices and relatively ^{13}C -enriched $\delta^{13}\text{C}_{\text{OM}}$ values (Fig. 3). The increase in crenarchaeol, as well as the low ~~[2]/[3]~~[2]/[3] index values, suggest that more Thermoproteota ~~are~~were present in Lake Magadi

in thosese periods. As mentioned in Section 4.1.1, three groups of AOA are of interest for interpreting 565 what which archaeal groups are found in low index intervals of Lake Magadi. Averages of [2]/[3][2]/[3] from the global dataset in Rattanasriampaipong et al. (2022) are as follows: hot spring mats (avg. = 1.00), shallow AOA cultures (avg. = 1.16), and shallow core tops (avg. = 2.64). Placing these on a continuum, we can approximate the environment from [2]/[3][2]/[3] averages in Magadi, though it should be noted that the shallow AOA and shallow core-top values in Rattanasriampaipong et al. (2022) are based on 570 marine core-tops, while the hot spring mats are based on terrestrial hot springs like those observed around Lake Magadi (i.e., pH > 6.5).

Interval 1 captures a transition from a more arid East Africa to a wetter period at the onset of the African Humid Period (AHP). During wetter periods, more allochthonous material is carried transported to into 575 the lake, which includes vegetal remainstion that impacts the overall bulk $\delta^{13}\text{COM}$ values. This allochthonous vegetation enriches the overall bulk $\delta^{13}\text{COM}$ values more significantly compared to than other intervals in the Magadi core. Average values of bulk $\delta^{13}\text{COM}$ are -17.7 ‰ in Interval 1, which correspond to the $\delta^{13}\text{COM}$ values of aquatic sedges mixed with a terrestrial signal of grassy woodland (Sikes, 1994; Reiffarth et al., 2016). The average ACL₍₂₅₋₃₃₎, CPI_{alk}, and CPI_{FA} values in Interval 1 are

580 29.4, 2.0, 3.2, respectively, suggesting a C4 vegetation origin of the long chain n-alkanes as indicated by the value of 29.4 and moderate overall degradation overall as implied by the low CPI_{alk} and CPI_{FA} values (Table 2). This degradation is likely why no n-alkanes shorter than C₂₂ were identified in the core.
Furthermore, pollenPollen records in Lake Magadi indicate that a mixture of C4 grassy woodlands and C4 aquatic sedges were predominant in the landscape that surrounded Lake Magadi at this time (Muiruri 585 et al., 2021). Supporting the pollen record, the $\delta^{13}\text{COM}$ values are likely reflecting $\delta^{13}\text{C}$ values similar to those observed by Garcin et al. (2014) in equatorial regions of Cameroon. The bulk $\delta^{13}\text{COM}$ is likely recording a mixture of C4 grasses and C4 sedges similar to $\delta^{13}\text{C}$ values of C₂₇ to C₃₃ n-alkanes obtained from C4 grasses and sedges in Cameroon which ranged from -18.2 to -17.6 ‰ (Garcin et al., 2014). This all suggests that the bulk $\delta^{13}\text{COM}$ signal is dominated by terrestrial biomass, unlike other sections of the 590 core, and there does not appear to be a significant influence from the benthic microbial community (i.e., methane cyclers or SRB).

Values of the $\frac{[2]}{[3]} / \frac{[2]}{[3]} / \frac{[2]}{[3]}$ index average 2.1 in Interval 1 with some values as high as 3.74 and 4.63 at 33.28 and 33.03 m, respectively (Table 4 Table S1). The higher values are closer to what is captured from 595 deep oceanic suspended particulate matter (SPM) and deep ocean core-tops below the pycnocline, though caution should be used when comparing lacustrine and oceanic sediments (Rattanasriampaipong et al., 2022). The increase in %-GDGT-0/4-cren (50.6 and 54.3 %; Table 4 Table S1) and the slightly increased MI values (0.37 and 0.41; Table 4 Table S1) suggest imply that these samples were deposited in a deeper lacustrine environment. Evidence for a deeper paleolake at ~40 m-feet (~17–18 m) above the modern 600 lakeshore (Baker, 1958) is also observed in the High Magadi Beds Formation (ca. 17.7 to 10.8 ka) indicating that there was fresh water flowing into the lake during the period of deposition in Interval 1, likely creating a fresher water cap on the meromictic Lake Magadi (Barker et al., 1991; Behr, 2002; Owen et al., 2019). However, excluding the high $\frac{[2]}{[3]} / \frac{[2]}{[3]} / \frac{[2]}{[3]}$ index values in Interval 1, the average is 1.6, which is closer to the hot spring mats and shallow AOA cultures (Rattanasriampaipong et al., 2022). 605 Likely, the higher $\frac{[2]}{[3]} / \frac{[2]}{[3]} / \frac{[2]}{[3]}$ index values mean represent periods of increased methanogenesis occurring in the sediments, with AOA input from the upper water column likely induced by proportionally increases in the amount of hydrothermal activity inflow to the lake (see Section 4.2.1). Fig. 4 shows that the Ca/Na is anti-correlated with REEs in both the PCA and correlation matrix. Since the proportion of Ca/Na decreases when REEs increase, we can say that statistically, when it is drier (and thus 610 proportionately more hydrothermally influenced) the Ca/Na decreases, REE values increase, and the methane indices are suppressed. In the periods of lower $\frac{[2]}{[3]} / \frac{[2]}{[3]} / \frac{[2]}{[3]}$ values, the community is interpreted as being dominated by AOA and thermophilic AOA cultures (i.e., Thermoproteota; Rattanasriampaipong et al., 2022) and is further supported by high % cren and % cren². Kumar et al. (2019) described similarly low $\frac{[2]}{[3]} / \frac{[2]}{[3]} / \frac{[2]}{[3]}$ values in the water column of Lake Malawi that are akin to 615 values observed in Lake Magadi in both Intervals 1 and 3. They found that values of a lower normalized $[2] / [2+3]$, ranging from 0.55 to 0.59, in Lake Malawi were associated with the shallower Thermoproteota (Thaumarchaeota) Group I.1b. This is compared to greaterhigher values of $[2] / [2+3]$ in the deeper dwelling Thermoproteota Group I.1a, which means that most samples in Interval 1 are likely sourced from Group I.1b (Kumar et al., 2019). The interpretations of Kumar et al. (2019) concluded that Group

620 I.1b Thermoproteota were contributing to the lower [2] / [2+3] values, while the more deeply dwelling
Group I.1a Thermoproteota were more prevalent in aphotic portions of the water column (Kumar et al.,
2019). The normalized [2] / [2+3] used by Kumar et al. (2019), with values ranging from 0.55 to 0.65,
approximates values of $\frac{[2]}{[2]+[3]}$ in the 1.30 to 1.65 range as described in this paper. More recently,
Baxter et al. (2021) found that Thermoproteota I.1b are more prevalent in the upper oxygenated portion
625 of the water column within the photic zone as evidenced by a higher relative abundance of crenarchaeol
and lower relative abundance of GDGT-2. Thus, our interpretations of thaumarcheotal AOA in Lake
Magadi sediments are consistent with data from Baxter et al. (2021) and Kumar et al. (2019). This
interpretation is consistent with Interval 1 being a period of proportionately more freshwater and HCO_3^- -
rich hydrothermal input and a deeper lake overall, which would explain the accompanying increase in
630 crenarchaeol.

Interval 3 $\frac{[2]}{[2]+[3]}$ averages are lower overall (Table 1/Table S1; avg. = 5.4), with only one outlying
high value (ca. 77.32 m at a value of 36.7). Excluding this high index value, the $\frac{[2]}{[2]+[3]}$ average
drops to 1.5, which is closer to what is observed in shallow AOA cultures and hot spring mats. With most
635 samples being closer to unity (i.e., $\frac{[2]}{[2]+[3]} = 1.0$), it is likely that hot springs had a greater influence
on the community composition in these intervals. Samples that are closer to unity (70.78, 70.86, and 71.08
to 75.93 m) also have a relatively ^{13}C -enriched $\delta^{13}\text{C}_{\text{OM}}$ values (avg. = -21.8‰) compared to samples with
a higher $\frac{[2]}{[2]+[3]}$ (averaging 1.5 excluding the outlying value of 36.7). This average is closer to
shallow Group I.1a Thermoproteota as described previously. Average isotope values in Interval 3 are
640 between oceanic hydrothermal vents (avg. = -19.0 ‰) and terrestrial alkaline hot spring systems such as
the Bison Pool hot spring in Yellowstone National Park (avg. = -24.9 ‰) (Shah et al., 2008; Schubotz et
al., 2013). Since elevated amounts of GDGT-2 (i.e., relative abundance > 45%) are associated with
Euryarchaeota, and values in Intervals 1, 3, and 5 are much lower than 45% (Table 1/Table S1), these
645 intervals are likely dominated by Thermoproteotal AOA (Pancost et al., 2001; Turich et al., 2007; Taylor
et al., 2013). Archaeal community composition in Intervals 1, 3, and 5 is independent of these external
factors and is related to hydrothermal flows. CPI_{alk} , CPI_{FA} , and $\text{ACL}_{(25-33)}$ average 7.0, 5.0, and 30.9,
which indicates a higher terrestrial input that tracks with the aridity during this period. This further

supports the hot springs driving the lake archaeal community composition as there was less overall precipitation and the Thermoproteotal communities were more abundant during Intervals 1, 3, and 5.

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Lastly, in-Interval 5, which only has 4 samples, has similarly low values of [2]/[3]/[2]/[3] (average = 1.4) like Intervals 1 and 3, which is likely indicative of Thermoproteotal AOA cultures. The CPI_{alk} and CPI_{FA} averages were 4.6 and 5.0, ~~which indicates more of a~~ ~~indicates more~~ terrestrial input. So, while these values are lower than Interval 3, and closer to the values in Interval 1, these still indicate a higher terrestrial input during this time-frame.

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4.2 The influence of hot hydrothermal activityspring/runoff ratios on the microbial-archaeal methane cycles

In the Lake Magadi core, there are intervals in which sediments have a tuffaceous quality implying a more hydrothermal origin produced in situ (Fig. 3Fig. 2, data from Owen et al., 2019). Hydrothermal

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fluids in the basin are rich in carbonate and bicarbonate as well as Na⁺ ions, inferred to be a result as a

direct result of the weathering and alteration of trachytic (silica-rich) basement rock (Jones et al., 1977;

Allen et al., 1989), and mantle-derived CO₂ discharged mainly along faults (Lee et al., 2017; Muirhead

et al.). Ca²⁺ and Mg²⁺ are also very low with major impacts on sedimentation (Deocampo and Renaut,

2016Renaut and Owen, 2023). Renaut and Owen (2023) note that ~~the~~ hydrothermal waters become

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important contributors to lake recharge proportional influence of spring fluids becomes more significant

during arid phases when fluvial inflow is reduced~~declines~~. For example, Nasikie Engida, a small

hypersaline lake to the northwest of Lake Magadi, is partly maintained today almost entirely by

hydrothermal inputs, with trona, nahcolite and zeolites accumulating during dry periods when there is

little or no fluvial inflow (DeCort et al., 2019; Renaut et al., 2020; Renaut and Owen, 2023). Magadi The

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Core MAG14-2A Magadi palaeolakes lacks evidence (e.g., mud cracks, soils, calcrete) for complete

desiccation (e.g., mudcracks, soils, calcrete) and the lake appears to have been maintained surface

waters through multiple drought episodes during the last million years (Owen et al., 2019). In contrast,

separate lakes in the neighbouring Koora Basin (~10 km to the east), dried out many times, leading to soil

formation in that basin (Owen et al., 2024).

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Owen et al. (2019) also noted that during periods when highly saline, alkaline water dominated at Magadi ash was zeolitized, REE patterns developed significant anomalies, and pyrite developed in anoxic/euxinic bottom waters of a meromictic lake. They also noted the excellent preservation of diatom opaline silica in highly alkaline lakes after about 540 ka, which is suggests that very high levels of silica in lake 680 water to-preserved their frustules from dissolution. Hydrothermal springs at Magadi today contain high silica, but their waters require further evaporation to achieve concentrations that would preserve diatom silica under very high pH conditions. High silica concentrations brought about by strong evaporation in a lake maintained by spring inflows (meteoric or hydrothermal) may also partly help explain the abundance of chert through the Magadi sedimentary sequence.

685 Proportional increases in hot-spring water increased 'proportional' inputs of spring water during episodes periods of increased aridity would have favoured the development of highly saline, alkaline waters with significant impacts on archaeal communities.

There is abundant evidence for hydrothermal influences in the sediments in the Lake Magadi core (Owen 690 et al., 2019). Owen et al. (2019) in detail from trachyte 1,056 ka to modern alkaline pan. Here, the increases in hydrothermal activity are interpreted as a proportionally higher input of hydrothermal fluids due to concurrent aridity and there is overall less of an influence of other meteoric flows during these intervals. Hydrothermal activity, like in Nasikie Engida to the northwest, continually supplies its lake and during 695 periods of lower overall precipitation (DeCort et al., 2019; Renaut and Owen, 2023). This proportional increase in hydrothermal activity is likely why mud cracks and other evidence for a complete lakebed drying event was not recorded in the sampling core. Samples ranging from ~545 to 380 ka (Figs. 3 and 4; Interval 6) in Interval 6 marked represent a drier period when the lake area and volume had shrunk, partly due to tectonic influences (Owen et al., 2024), and lake floor anoxia was prevalent (Owen et al., 2019). Sediments in Interval 6 do not show any evidence of hot spring activity (i.e., there is a higher Ca/ 700 Na; Fig. 4Fig. 5), and the main zeolite in this interval is analcime, indicating more evaporitic saline as well as alkaline waters (Owen et al., 2019). In the core intervals between from ~380 to and 105 ka (Fig. 3Fig. 2; Interval 5 through mid-Interval 2), the system paleolake was largely frequently meromictic was flooded periodically with a freshwater- mixolimnion that supported freshwater planktonic diatoms while

the saline monimolimnion and lake bottom floor waters favoured alteration of ash to a variety of zeolites

705 (e.g., erionite, phillipsite, clinoptilolite, analcime) in different lake settings but maintained meromixis and a high proportion of zeolites forming in the lake waters with analcime as the dominant zeolite indicating a higher salinity (Owen et al., 2019). Starting at ca. 380 ka to modern day, hydrothermal fluids with abundant HCO_3^- and CO_3^{2-} ions and $\text{Na}^+ > \text{Ca}^{2+}$ have heavily influenced both deposition and chemical alteration of sediments (Owen et al., 2019). Evaporitic enrichment of alkaline hydrothermal fluids

710 circulating through trachytic basement rocks and the increase in their contribution to Lake Magadi are responsible for the alteration of the volcanic glasses. An abundance of analcime, combined with the $\text{Na}^+ > \text{Ca}^{2+}$, indicates a proportionally higher influx of hydrothermal waters in Intervals 5 through 2 (Fig. 4 Fig. 5; Owen et al., 2019). Similarly, from ~105 to 0 ka (Fig. 3 Fig. 2; mid-Interval 2 through Interval 1) there is evidence of a low Ca / Na, increased Br, and the abundant zeolite formation, namely analcime,

715 pointing to increased saline conditions. After about 80 ka, tectonic adjustments and increasing aridity led to desiccation in the Koora Basin suggesting that spring inflows were important in maintaining a hypersaline lake in the Magadi Basin, as they do today during dry seasons (hydrothermal flow in this interval as well (Fig. 4 Fig. 5; Owen et al., 2019). These hydrothermal flows are especially prevalent (low Ca/Na values) where at intervals where tuffaceous silts are the dominant lithology of the core, values of

720 Ca/Na have decreased, however, the values of Ca/Na ratios have steadily decreased over time, though not at a 1:1 rate, which aligns with methane index values and sudden increases of these indices. Fig. 4 shows that the Ca/Na and methane indices are statistically different from one another in both the correlation matrix and PCA plot. In Fig. 5b, Ca/Na is loaded positively on PC1 and PC2, while the methane indices ($\text{[2]}/[3]$, MI, %0/Cren, and %2/Cren) are loaded positively on PC1 and negatively on PC2. Furthermore,

725 The interval of tuffaceous silts occur at 96–102 m core depth, and is marked by increases in zeolitic alteration from saline water inflow and a low Ca/Na ratio (Fig. 4, Intervals 1, 3, and 5). While the tuffaceous silts at 30–36 and 74–76 m were changed to muds in Renaut and Owen (2023), REE data also still indicate appear to reflect our a increased proportions of hydrothermal activity origin of in the MI-

730 off intervals and are typically higher in and MI-on periods such as Intervals 1 and 3 (Fig. 5) as the REEs are anticorrelated in the correlation matrix (Fig. 5a) and loaded on different PC axes (Fig. 5b).

Consequently, we interpret the changes in salinity and alkalinity in the Magadi palaeolakes as reflecting the impact of climate on spring/runoff ratios into the Magadi Basin, which in turn have exerted significant impact on the archaeal communities through the last million years. Tuffaceous silt intervals align well overall with the low MI values; sampled material with low MI values is found wholly within the tuffaceous silts, indicating that these values are still likely related to increased hydrothermal inputs. The presence of chert in non-tuffaceous samples might also be a hydrothermal indicator, as the circulating hydrothermal fluids beneath Lake Magadi are silica rich (Eugster, 1969). Periods of proportionally increased hydrothermal activity were likely occurring throughout Lake Magadi's history as there is evidence for chert beds and nodules in the Quaternary sediments surrounding the lake—namely the Oloronga Beds, Green Beds, and High Magadi Beds (Behr, 2002; Renault and Owen, 2023). The chert-bearing Oloronga Beds were deposited from approximately 800 to 300 ka, while the Green Beds were actively precipitating chert from ~220 to 70 ka and more recently, the High Magadi Beds were being deposited between ~25 and 9 ka (Fairhead et al., 1972; Goetz and Hillaire-Marcel, 1992; Williamson et al., 1993; Behr and Röhricht, 2000; Owen et al., 2019; Reinhardt et al., 2019). Thus, samples in the low-MI intervals (ca. 32.61 to 35.67, 70.78 to 75.93, and 103.2 to 104.1 m) are likely reflecting these periods of proportionally increased hydrothermal activity/spring-runoff ratios at Magadi caused by increased due in part to more evaporation of the lakebed and decreased precipitation in the surrounding landscape.

750 5. Conclusions

Sediments in Lake Magadi track the environmentally driven changes in archaeal communities over the past ~456 ka. Using the MI to track the predominantly microbial-archaeal inputs at Lake Magadi, we have observed sudden and distinct shifts between mixed archaeal communities of Euryarchaeotal methanogens and methanotrophs transitioning to mesophilic AOA Thermoproteota communities and back again. This shift is driven, in part, by moisture balances in the East African Rift, with wetter conditions periodically leading causing freshwater floods into a saline lake to form a meromictic to lake stratification waterbody at Magadi, and with more archaea derived from the upper water column rather than the sediments, as evidenced by low MI, low [2]/[3][2]/[3], and relatively ^{13}C -enriched bulk $\delta^{13}\text{C}_{\text{OM}}$.

Methane indices were typically higher during periods of reduced hydrothermal activity, indicating more 760 Euryarchaeal communities, whereas Thermoproteota communities thrived during periods of higher hydrothermal activity. There This is also a clear relationship between low MI values, and enhanced hydrothermal activity whereby increased hydrothermal activity spring/runoff ratios, lake (as recorded by tuffaceous silts and REEs in the sediments) salinity, alkalinity and the development of maintains low overall index values, indicating that environmental conditions supported more mesophilic 765 Thermoproteota. In contrast, periods of lower hydrothermal activity are connected to high index values. Thus, the MI indicates a switch between high and low hot spring activity in Lake Magadi. The intervals of MI off periods coincide with the tuffaceous units of inferred high hot spring activity in the geologic record, while MI on periods occur during periods of lower hydrothermal activity. As this was Theis study represents one of the first studies to look at examine methane cycling in a soda lake over geologic time, 770 we have and gained presents provides valuable insights into how variable these systems can be. Soda lakes are important ecosystems for methane cyclers and should be studied more closely so that we can better improve understanding of global methane contributions both in the past, and better constrain sources in the future.

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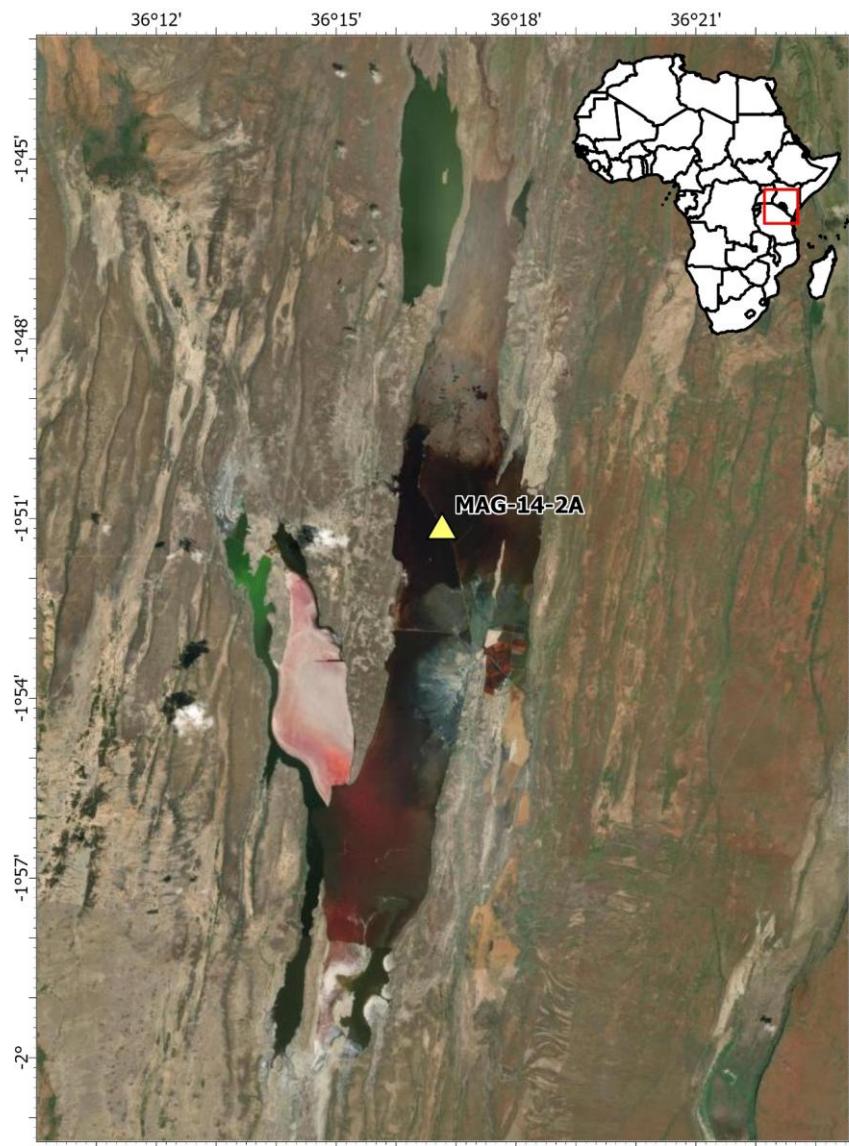
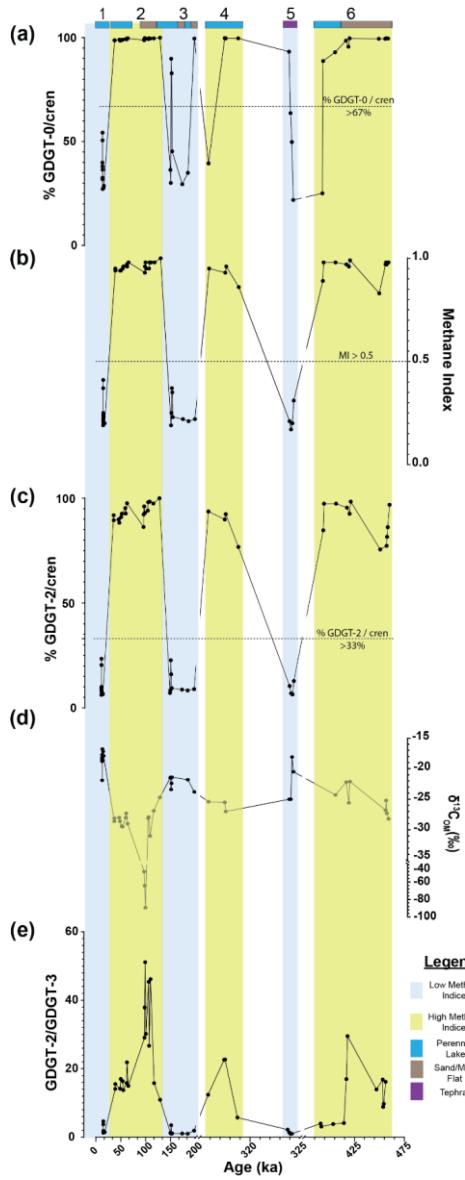
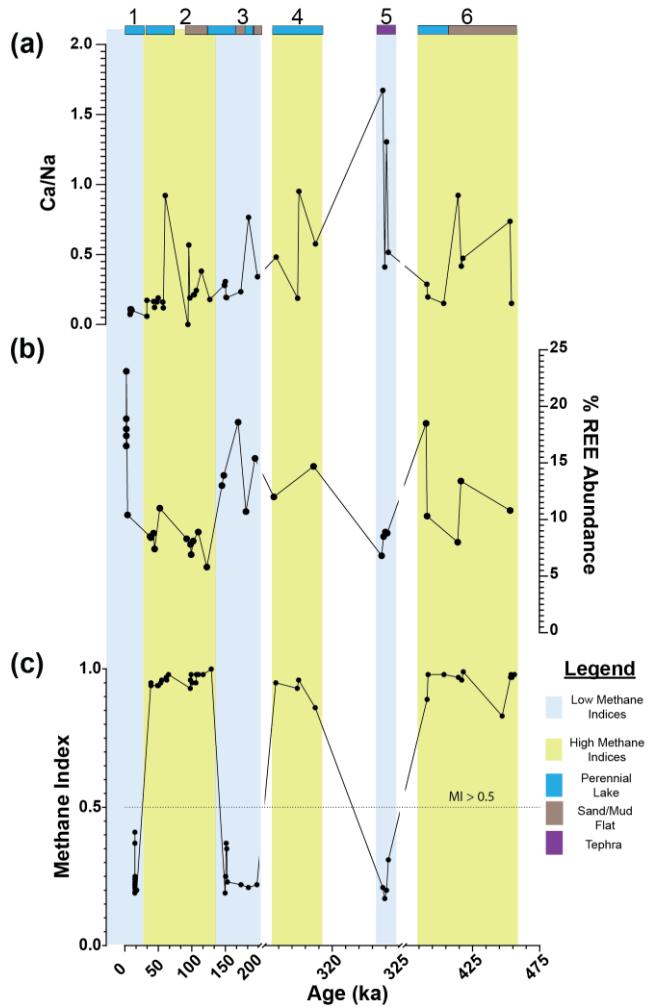


Figure 1. Map of the drilling location of MAG-14-2A (yellow triangle) in Lake Magadi for the Hominin Sites and Paleolakes Drilling Project (HSPDP).



1200 **Figure 2.** Downcore variations in Lake Magadi of the **a)** % 0 / Cren, **b)** MI, **c)** % 2 / Cren, **d)** bulk $\delta^{13}\text{C}_{\text{OM}}$,
and **e)** the GDGT-2 / GDGT-3 ([2] / [3]) values from ca. 14.9 to 456 ka. Sections 1, 3, and 5 are low MI
intervals outlined in blue, the high MI intervals in Sections 2, 4, and 6 are in yellow. Checkered patterns
indicate periods of tuffaceous silt deposit, which align with the low MI intervals. Bands at the top of the
graph indicate the inferred (via Renaut and Owen, 2023) lake levels and major inputs with dark blue
1205 indicating a perennial lake, brown indicating a sand or mud flat, and purple indicating tephra. Dotted lines
on each section denote the cut-off points for methane related indices MI (> 0.5), % GDGT-2 / cren (> 33
%), and % GDGT-0 / cren (> 67 %). See Section 2.3.2 for more details. Note the breaks in the X-axis
scale.



1210 **Figure 3.** Downcore plot for Lake Magadi of a) Ca/Na, b) % REE abundance, and c) MI. Values range from ca. 14.9 to 456 ka and Sections 1, 3, and 5 are outlined in blue reflecting a low MI interval, while high MI intervals are outlined in yellow. The checkered pattern is indicative of periods of higher inferred

hydrothermal flow. Bands at the top of the graph indicate the inferred (via Renaut and Owen, 2023) lake levels and major inputs with dark blue indicating a perennial lake, brown indicating a sand or mud flat, and purple indicating tephra. The dotted line on the MI plot (c) denotes the cutoff point > 0.5 for values significantly affected by methane cycling archaea. Note the breaks in the X-axis scale. REE values are from Owen et al. (2019).

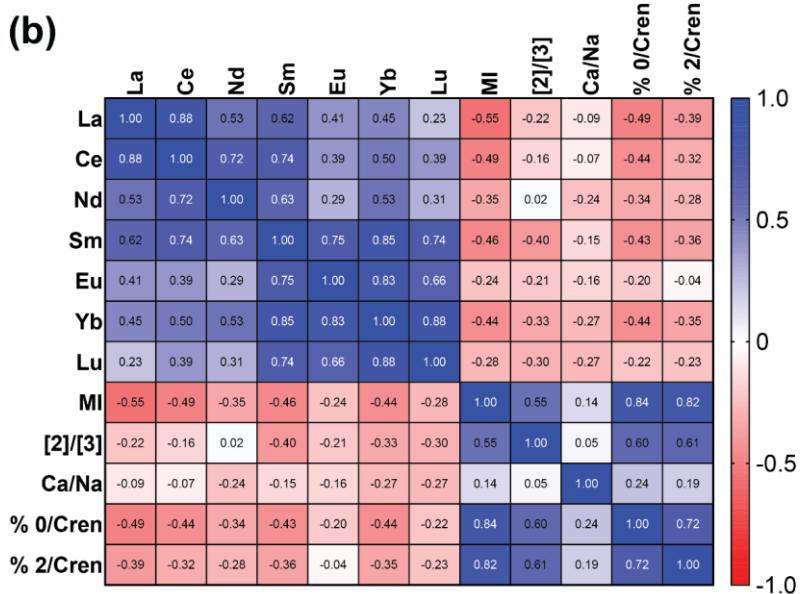
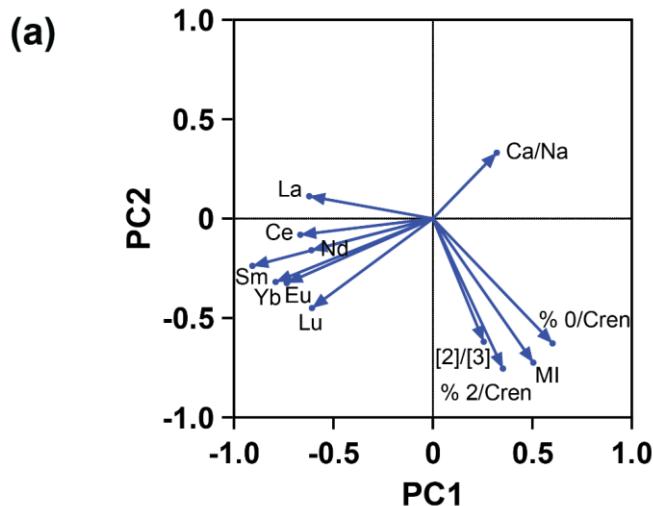


Figure 4. Both **a)** PCA and **b)** Spearman Correlation Matrix showing the relationship between methane related indices (MI and [2] / [3]) and REEs (La, Ce, Nd, Sm, Eu, Tb, Yb, and Lu) in the sampled intervals of the core. A negative relationship is seen between the methane indices and REEs as shown by opposing eigenvectors on the PCA (**a**) and negative r values on the correlation matrix (**b**). REE values are from Owen et al. (2019).

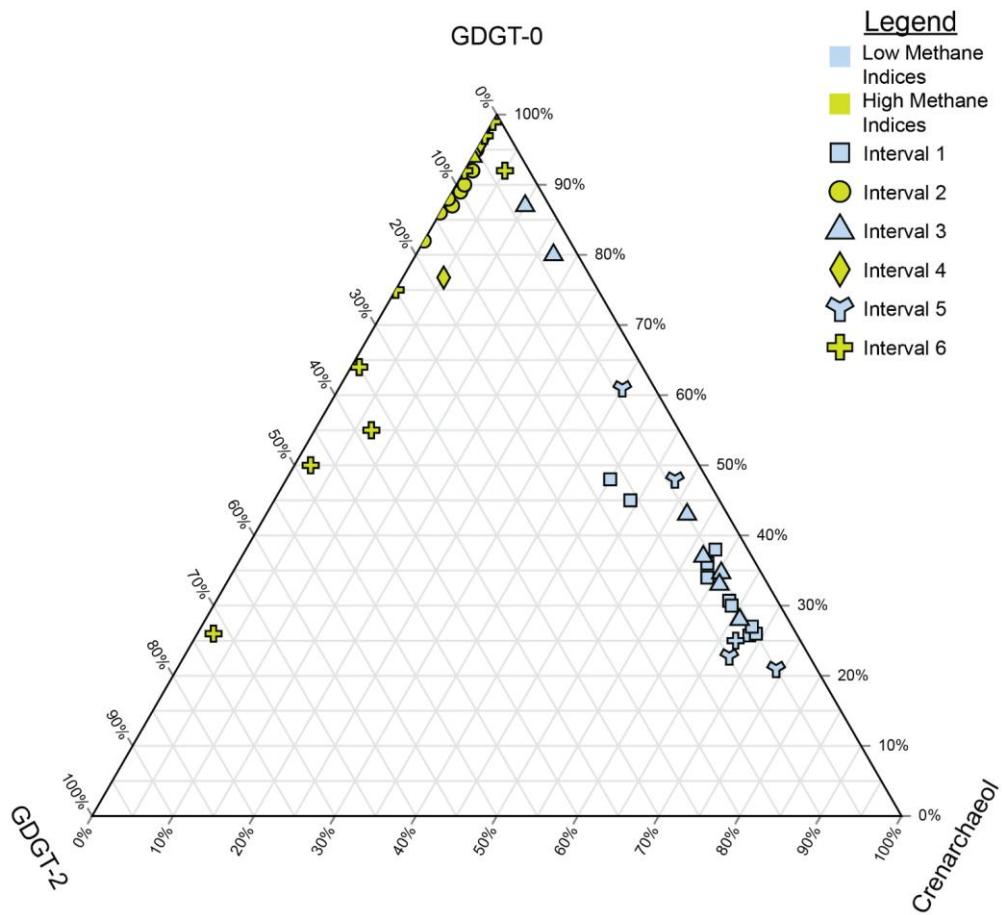


Figure 5. Ternary plot of crenarchaeol, GDGT-0, and GDGT-2, which are used to calculate the methane indices. Samples are split by both their interval (denoted by their shape) and whether they are from a high MI (yellow) or low MI (blue) interval. Higher proportions of GDGT-0 indicate methanogenic inputs, higher GDGT-2 indicate methanotrophy, and higher crenarchaeol indicates more mesophilic conditions influenced by hot springs.

1230