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Invited review

A review of molecular organic proxies for examining modern and ancient lacustrine environments

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ABSTRACT

Molecular organic geochemical proxies are increasingly being utilized to reconstruct past environmental conditions as new tools continue to be discovered and developed. To date, organic geochemical proxies have been developed mainly for use in marine systems and are widely used in paleoceanography. In contrast, organic proxies have been less commonly used on lacustrine sedimentary records. One reason for this is that the wide range in the physical and chemical properties of lakes complicates application of some organic geochemical proxies in lacustrine settings. Furthermore, in comparison to marine studies, presently only a small number of studies have conducted or are currently conducting fundamental research aimed at developing organic geochemical proxies for use in lacustrine settings. Despite this, an increasing number of (paleo)limnological studies are currently applying organic geochemical techniques to examine present and past environmental conditions. In this manuscript we review the use of a number of commonly utilized organic geochemical and isotopic proxies and discuss their potential for environmental reconstruction in Quaternary lacustrine deposits.

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1. Introduction

Organic matter preserved in geologic materials (e.g. sediment, rock, coal, petroleum) provides a direct indicator of environmental conditions at the time of deposition and thus is important to paleoenvironmental studies. Although only a small amount of the organic matter produced is eventually preserved in the geologic environment, the geochemical properties of this preserved organic matter can provide a wealth of information. The bulk geochemical properties of organic matter, including total organic carbon (TOC) content, carbon to nitrogen (C:N) ratios, and carbon ($\delta^{13}\text{C}$) and nitrogen ($\delta^{15}\text{N}$) isotopes, are routinely examined and applied to paleoenvironmental studies (e.g. Meyers, 2003). Examination of organic matter at the molecular level reveals contributions from the three domains of life: archaea, eukarya, and bacteria, and thus can often provide more specific information regarding past environmental conditions in comparison to bulk geochemical analyses, which reflect a mixture of allochthonous and autochthonous inputs.

Individual compounds, or compound classes, preserved in geological materials that can be traced to a particular source

organism, group of organisms, or to a particular process (e.g. photosynthesis) are called “biomarkers” (Peters et al., 2005). There are numerous reasons why biomarkers are of particular interest to paleolimnologists. Examination of organic matter at the molecular levels allows for the separation of terrestrial, aquatic and sedimentary components and thus environmental conditions in both the water column and the surrounding watershed can be examined simultaneously (Fig. 1). Biomarkers also allow for the examination of certain groups of algae and microorganisms that lack hard silica or carbonate shells and typically are absent from the fossil record. Diatoms are widely used as a paleolimnological proxy because their silica tests are often abundant and well preserved in lacustrine sediments. While diatoms are important contributors to primary productivity in many lakes, other types of phytoplankton also can be important. Biomarkers such as botryococenes and heterocyst glycolipids, produced by green algae and cyanobacteria, respectively, can be preserved in sediments thus providing information on the past abundance of these organisms, which are generally absent from the fossil record. Biomarkers of diatoms (e.g. loliolide) can be useful in conjunction with biogenic silica or diatom assemblage data, particularly in settings where silica dissolution has occurred. In recent years, organic geochemical proxies and compound-specific isotopic analyses have become increasingly utilized for paleoenvironmental reconstructions and have been used to address a wide range of research questions ranging from reconstructing

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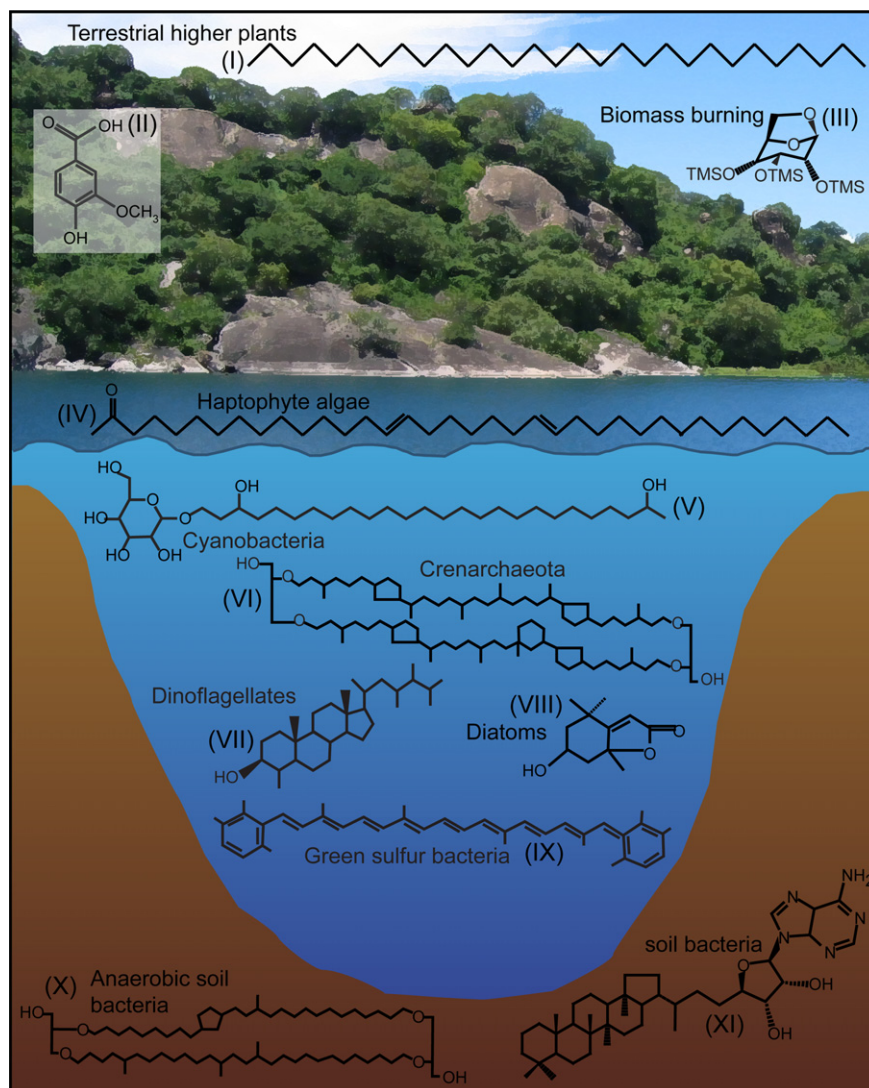


Fig. 1. The biomarker concept. Lacustrine sediments contain inputs from terrestrial and aquatic sources including higher land plants, microbes that live in the soils and sediments, and various types of algae and microorganisms that live in the water column. Using biomarkers, it is possible to examine contributions to sedimentary organic matter from the three domains of life, bacteria, archaea and eukarya, thus providing accurate paleoenvironmental information. There are also biomarkers of processes, such as photosynthesis or biomass burning. Structures shown include the (I) C_{29} *n*-alkane (a biomarker for terrestrial higher plants), (II) vanillic acid (a lignin phenol and biomarker for terrestrial higher plants) (Hedges and Mann, 1979), (III) levoglucosan (a biomarker for biomass burning) (Elias et al., 2001), (IV) the $C_{37:2}$ methyl alkenone (a biomarker for haptophyte algae), (V) 1-(*O*-hexose)-3,25-hexacosanediol (a heterocyst glycolipid and biomarker for N_2 -fixing cyanobacteria) (Bauersachs et al., 2009a), (VI) crenarchaeol (an isoprenoid glycerol dialkyl glycerol tetraether (GDGT) and biomarker for *Thaumarchaeota*) (Schouten et al., 2002), (VII) dinosterol (a biomarker for dinoflagellates) (Withers, 1983; Piretti et al., 1997), (VIII) loliolide (the anoxic degradation product of the pigment fucoxanthin, the major carotenoid present in diatoms) (Klok et al., 1984; Repeta, 1989), (IX) isorenieratene (a pigment produced by green sulphur bacteria that is used as an indicator of stratified conditions in ancient lake sediments) (Sinninghe Damsté et al., 2001a), (X) branched GDGT IIb (thought to be produced by anaerobic soil bacteria and a biomarker for soil organic matter) (Weijers et al., 2007b), and (XI) adenosylhopnane (an intact bacteriohopanepolyol (BHP) that has been proposed as a biomarker of soil bacteria) (Cooke et al., 2008a).

past terrestrial and aquatic ecosystems and environments (e.g. reconstructing lake surface temperatures, salinity, hydrological variability, vegetation type, soil organic matter input to aquatic environments, soil temperature and soil pH), for investigating biogeochemical cycling and carbon storage, and for examining the preservation and fate of organic matter in the geologic environment (Meyers, 1997; Killops and Killops, 2005; Peters et al., 2005; Eglinton and Eglinton, 2008). Biomarkers provide a wide range of opportunities for paleolimnological studies and the specific source information gained can complement or offer advantages over traditional paleolimnological proxies.

Lakes are sensitive recorders of climatic change and can provide excellent archives of past environmental conditions. In comparison to marine settings, many lakes are characterized by relatively high

sedimentation rates, offering possibilities for high-resolution paleoenvironmental reconstructions. Some lakes are characterized by a seasonally or permanently anoxic water column, thus increasing the preservation potential of organic matter. Lakes also provide opportunities for examining local or regional paleoenvironmental conditions and provide information on continental paleoclimate. The paleoclimatic importance of lacustrine sedimentary archives has long been recognized and in recent years there have been increased efforts to obtain long lacustrine paleoclimate records by drill coring. These records have provided both continuous and high-resolution climate archives from a variety of locations around the globe and geochemical studies of these lake sediments has provided, and is currently providing, a wealth of new paleoenvironmental information (e.g. Brigham-Grette et al., 2007;

Fritz et al., 2007; Hodell et al., 2008; Litt et al., 2009; Fawcett et al., 2011; Woltering et al., 2011).

The great majority of organic geochemical proxies that are of interest for reconstructing Quaternary paleoclimate were originally developed in and applied to marine systems, yet some can be directly applied to lacustrine environments. For example, primary producers such as diatoms, dinoflagellates and cyanobacteria are commonly found in both marine and lacustrine systems and there are diagnostic biomarkers for each of these groups (e.g. Volkman et al., 1998), allowing for past changes in primary productivity or shifts in algal community structure to be examined. However, for other types molecular proxies (e.g. paleotemperature proxies) application to lacustrine environments is not as straightforward due to additional complicating factors that are of importance in lacustrine systems: although the chemical properties of seawater vary somewhat between the different ocean basins, the range of variability in marine systems is quite small in comparison to the world's lakes. The physical (size, depth, basin morphology, catchment area) and chemical (temperature, salinity, pH, oxygen and nutrient concentrations) properties of lakes vary greatly. Here, we examine the application of commonly utilized organic geochemical and isotopic proxies to lacustrine sediments. Due to the extreme range in the chemical and physical properties of lakes it should be noted that the proxies and biomarkers discussed in this manuscript cannot simply be applied to all locations, and the applicability of many of the organic geochemical proxies to individual lakes needs to be examined on a site-to-site basis.

The application of organic geochemical proxies to lacustrine sediments has been previously reviewed by Meyers (1997, 2003), who also examined bulk geochemical techniques, which are not discussed here. However, since these reviews, several new molecular proxies have been developed, which can largely be attributed to the application and development of relatively new analytical methods such as high performance liquid chromatography mass spectrometry (HPLC-MS) and compound-specific carbon and hydrogen isotope analysis. Therefore, the main goal of this review is to provide an overview of recently developed molecular and isotopic proxies that can be used to reconstruct paleoenvironmental conditions from Quaternary lacustrine deposits. We also will highlight the differences, when necessary, between application of molecular proxies in marine and lacustrine settings and discuss additional factors that need to be considered for application of molecular proxies to lacustrine environments. Finally, we highlight some recently developed molecular proxies that have not yet been widely applied to lacustrine settings but that are likely to be useful for Quaternary studies.

1.1. A brief overview of analytical techniques for examining molecular fossils

Organic matter can be broadly separated into two classes: bitumen and kerogen. Bitumen is the solvent-extractable component of the organic matter while kerogen is the non-solvent-extractable component. This review focuses on biomarkers derived from bitumen since this is where the great majority of Quaternary paleoenvironmental studies have focused their research efforts, although we note that biomarkers are also present in kerogen that can provide paleoenvironmental information (see review of Vandenbroucke and Largeau, 2007).

The process of isolating biomarkers from a bulk sediment, culture, or water column filter sample typically begins by extraction with organic solvents followed by additional procedures to fractionate and purify compound classes of interest. Individual compounds (biomarkers) can be analyzed by a variety of techniques. Gas chromatography (GC) and gas chromatography-mass

spectrometry (GC/MS) are the most commonly applied techniques for quantifying and identifying individual compounds. GC-isotope ratio monitoring mass spectrometry (GC-IRMS) is a relatively newer technique (Hayes et al., 1990) and is used to measure the isotopic composition of individual compounds. Compound-specific carbon ($\delta^{13}\text{C}$), nitrogen ($\delta^{15}\text{N}$) and deuterium (δD) isotopes can be measured with a typical precision of 0.1–0.3‰, 0.3–0.7‰, and 2–5‰, respectively (see review of Sessions, 2006a). We also note that recently a technique for measuring compound-specific sulphur isotopes ($\delta^{34}\text{S}$) has been developed using GC together with multi-collector inductively coupled plasma mass spectrometry (MC-ICPMS), which has a precision better than 0.5‰ for samples containing as little as 6 pmol S (Amrani et al., 2009).

Another relatively new technique for quantifying and identifying biomarker lipids is high performance liquid chromatography (HPLC) with positive ion atmospheric pressure chemical ionization mass spectrometry (APCI-MS) (e.g. Hopmans et al., 2000). HPLC techniques have the advantage that they allow for larger (>1000 Da) and more polar compounds to be analyzed in comparison to traditional GC or GC/MS techniques. Additionally, HPLC coupled to mass spectrometry with electrospray ionization (HPLC/ESI-MSⁿ) allows for the detection of even more polar, charged compounds (Zink et al., 2003; Sturt et al., 2004) while HPLC with photodiode array UV–vis detection and APCI-MS can be used to examine sedimentary pigments (Keely et al., 1988; Eckardt et al., 1991; He et al., 1998; Ocampo and Repeta, 1999; Ocampo et al., 1999; Ains et al., 2001). Increased selectivity can be gained by using MS–MS (MS²) techniques (note that MS² techniques are also used to increase selectivity in GC/MS). HPLC techniques are currently being utilized by an increasing number of studies and in the last decade have detected previously unknown biomarkers and provided a number of novel proxies (e.g. Hopmans et al., 2006; Talbot and Farrimond, 2007; Bauersachs et al., 2009b).

1.2. Framework of review

In the following parts of this manuscript the discussion is divided into sections by individual compound classes. In the section headings, we have indicated the main environmental parameter (e.g. temperature) that can be examined based on individual compound classes but note that for certain compound classes multiple environmental parameters may be reconstructed. We also indicate the methods that are commonly used to examine individual compound classes although for some compound classes analysis by multiple techniques (e.g. GC/MS and HPLC/MS) is possible. Thus, the reader should refer to the cited references for detailed analytical information. Throughout this review we use Quaternary case studies to highlight recent research developments or new insights that have come from applying molecular or isotopic techniques to lacustrine sediments.

2. Paleotemperature, paleosalinity and haptophyte input from long-chain alkenones

One of the oldest and most widely-applied organic geochemical proxies, the U_{37}^k Index, is based on long-chain alkenones (LCAs) and provides a method for reconstructing past sea surface temperatures, and in some cases, lake surface temperatures. LCAs are found in many marine and an increasing number of lacustrine sediments and comprise a series of C₃₇–C₃₉ di-, tri-, and tetra-unsaturated methyl (Me) and ethyl (Et) ketones (Fig. 2), which are analyzed by GC and GC/MS techniques. The original U_{37}^k Index (Brassell et al., 1986) reflected the proportions of the di-, tri-, and tetra-unsaturated ketones (Table 1); however, subsequent research showed that there was no empirical benefit to including the tetra-

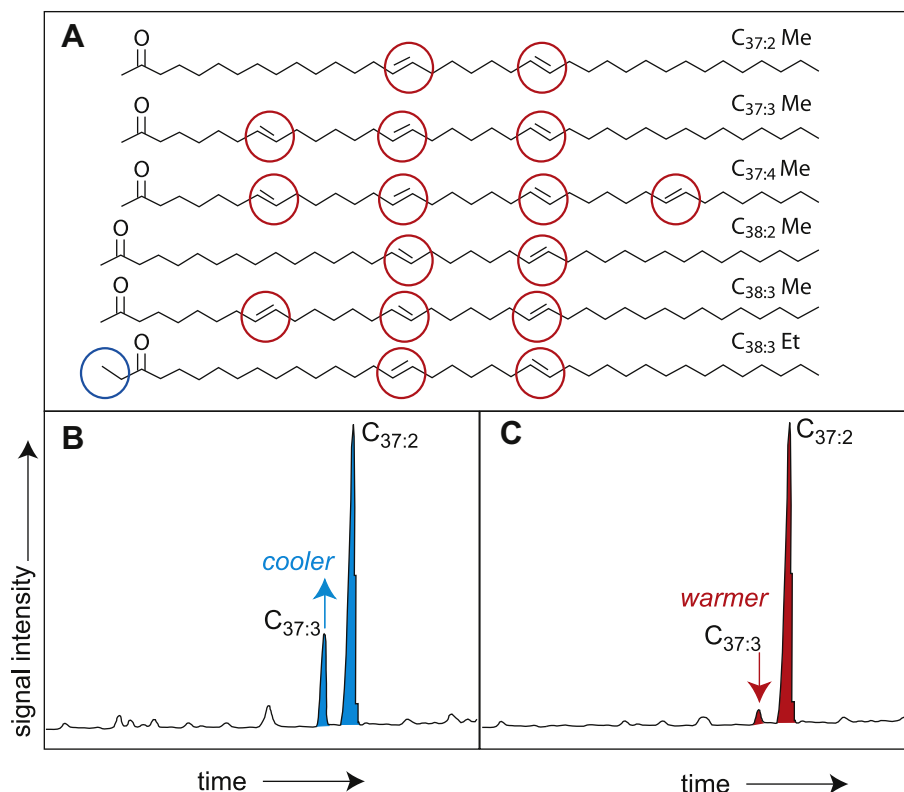


Fig. 2. A) Examples of structures of methyl (Me) and ethyl (Et; circled in blue) long-chain alkenones. These structures are somewhat similar but differ in the number of double bonds, which are indicated by the red circles. B) and C) Illustration of the U_{37}^k SST proxy. As temperature increases, the relative proportion of the $C_{37:3}$ alkenone decreases. Panel B illustrates the GC chromatogram of a sample with a relatively cold signal and C illustrates a sample with a relatively warm signal. Note that the uppermost limit of the U_{37}^k SST proxy is at $\sim 28^\circ\text{C}$ (e.g. $U_{37}^k = 1$, the corresponding temperature varies slightly depending on the calibration used), which is the point where no $C_{37:3}$ alkenone remains, or where its abundance is too low to quantify.

unsaturated ketone in the paleotemperature equation and the U_{37}^k Index (Table 1) was suggested (Prahl and Wakeham, 1987) and has been widely adopted. U_{37}^k ratios vary positively with growth temperature: as temperature increases, the relative proportion of the tri-unsaturated ketone decreases (Fig. 2). In marine environments, LCAs are mainly produced by two widely distributed species of haptophyte algae, *Emiliania huxleyi* (de Leeuw et al., 1979) and *Gephyrocapsa oceanica* (Conte et al., 1994; Volkman et al., 1995) and

have also been reported in the coastal species *Isochrysis galbana* and *Chrysolita lamellosa* (Marlowe et al., 1984). These organisms are all haptophyte algae of the class Prymnesiophyceae, which require sunlight for photosynthesis and generally reside in the upper photic zone (Herbert, 2003 and references therein). LCAs also have been reported from brackish and freshwater environments. Lacustrine LCAs were first reported from the English Lakes district in 1985 (Cranwell, 1985) but more recent studies have revealed

Table 1
Lacustrine U_{37}^k , U_{37}^k , U_{3738}^k and U_{38}^k calibrations and formulas.

Calibration equation	n	r^2	Calibrated to	Geographical location	Reference
$U_{37}^k = 0.02T - 0.121$	9	0.89	Summer lake temp	Germany	Zink et al. (2001)
$U_{37}^k = 0.0211T - 0.725$	9	0.68	Summer lake temp	Germany	Zink et al. (2001)
$U_{37}^k = 0.0328T + 0.126$	38	0.83	Mean annual air temp	China	Chu et al. (2005)
$U_{37}^k = 0.037T + 0.108$	14	0.9	Mean annual air temp	China: fresh-brackish lakes	Chu et al. (2005)
$U_{37}^k = 0.025T + 0.153$	24	0.67	Mean annual air temp	China: saline lakes	Chu et al. (2005)
$U_{37}^k = 0.0011T^2 - 0.0157T + 0.1057$	14	0.99	Growth temp ($10\text{--}22^\circ\text{C}$)	China: <i>C. lamellosa</i> culture	Sun et al. (2007)
$U_{37}^k = 0.0257T - 0.2608$	9	0.97	Growth temp ($14\text{--}22^\circ\text{C}$)	China: <i>C. lamellosa</i> culture (linear fit)	Sun et al. (2007)
$U_{3738}^k = 0.0464T - 0.867$	13	0.8	Mean autumn temp	Spain	Pearson E.J. et al. (2008)
$U_{38}^k = -0.753 + 0.0412T$	12	0.71	Mean autumn temp	Spain	Pearson E.J. et al. (2008)
$T = 39.9U_{37}^k + 36.418$	21	0.75	<i>in-situ</i> water temp	Lake George, North Dakota (USA)	Toney et al. (2010)
$T = 40.8U_{37}^k + 31.8$	34	0.96	<i>in-situ</i> water temp*	Braya Sø (Greenland)	D'Andrea et al. (2011)
$U_{37}^k = 0.033T + 0.044$	370	0.98	Annual mean SST	Global marine	Müller et al. (1998)

$$U_{37}^k = (C_{37:2} - C_{37:4}) / (C_{37:2} + C_{37:3} + C_{37:4}).$$

$$U_{37}^k = (C_{37:2}) / (C_{37:2} + C_{37:3}).$$

$$U_{38}^k = (C_{38:2} - C_{38:4}) / (C_{38:2} + C_{38:3} + C_{38:4}).$$

$$U_{38}^k = (C_{38:2}) / (C_{38:2} + C_{38:3}).$$

$$U_{3738}^k = (C_{38:2} - C_{38:4} + C_{37:2} - C_{37:4}) / (C_{38:2} + C_{38:3} + C_{38:4} + C_{37:2} + C_{37:3} + C_{37:4}).$$

*The calibration of D'Andrea et al. (2011) combines water filters from Braya Sø with the core top U_{37}^k calibration of Zink et al. (2001).

their presence in the water columns or sediments of lakes from around the world including Antarctica (Volkman et al., 1988; Coolen et al., 2004b; Jaraula et al., 2010), Europe (Cranwell, 1985; Thiel et al., 1997; Innes et al., 1998; Zink et al., 2001; Pearson E.J. et al., 2007, 2008), Asia (Li et al., 1996; Wang and Zheng, 1998; Sheng et al., 1999; Zink et al., 2001; Chu et al., 2005; Sun et al., 2007; Liu et al., 2008), North America (Zink et al., 2001; Toney et al., 2010), Africa (Kristen et al., 2010), South America (Theissen et al., 2005) and Greenland (D'Andrea and Huang, 2005; D'Andrea et al., 2011).

2.1. Lacustrine alkenone producers

Understanding the ecology of a particular organism, or group of organisms, is essential for interpretation of its associated molecular signature. In contrast to marine environments where the dominant alkenone producers are well known (*E. huxleyi* and *G. oceanica*), the organism(s) responsible for producing lacustrine alkenones are currently poorly understood although genetic studies provide some insights. Coolen et al. (2004b) examined preserved 18S rDNA in Holocene anoxic sediments from Ace Lake (Antarctica) and found six novel phylotypes related to known alkenone-biosynthesizing haptophytes, with the species *I. galbana* UIO 102 as their closest relative (98.3–99.5% sequence similarity). D'Andrea et al. (2006) also examined 18S rDNA in water filtrate, surface and Late-Holocene sediment samples from lakes in West Greenland. Their results suggest alkenones in these lakes are synthesized by a distinct phylotype, differing from both marine haptophytes and those from Ace Lake. Sun et al. (2007) isolated *C. lamellosa* Anand from a cultured mixture of phytoplankton (grown using filtered natural lake water with a Na⁺ concentration of 8.78 g/L) collected from saline Lake Xiarinur (Inner Mongolia, China). DNA analysis of water column samples of Lake Fryxell (Antarctica) indicated 18S rDNA sequences closely related to *Isochrysis* spp., *C. lamellosa* and the haptophyte sequences previously reported from the west Greenland lakes and Ace Lake (Jaraula et al., 2010). Theroux et al. (2010) examined 18S rDNA from 15 alkenone-containing lake surface sediments from diverse geographical locations. They found that all haptophyte sequences grouped within the order Isochrysidales and none of the sequences branched directly with the marine species *E. huxleyi* or *G. oceanica*. An especially important finding of this study is that individual lakes contained DNA from multiple haptophyte species (Theroux et al., 2010). Clearly, the main alkenone producers vary from lake to lake, which in combination with the observation that multiple haptophyte species may exist in individual lakes, likely explains some of the variability noted in lacustrine LCA distributions and relationships observed with temperature or salinity (discussed in Sections 2.3 and 2.4).

2.2. Lacustrine LCA homologue distributions

Early work suggested that a characteristic feature of lacustrine LCA distributions was relatively high amounts of the C_{37:4} Me ketone (Fig. 2) (Thiel et al., 1997). In contrast, marine environments are generally characterized by abundant di- and tri-unsaturated compounds (C_{37:2} and C_{37:3}) whereas the C_{37:4} homologues are typically present in minor abundances or are absent, except for more polar environments. Indeed, the presence of the C_{37:4} Me ketone has been noted in nearly all lacustrine sediments from which compositional data is reported and in many cases, C_{37:4} is the dominant homologue (Cranwell, 1985; Volkman et al., 1988; Li et al., 1996; Thiel et al., 1997; Wang and Zheng, 1998; Sun et al., 2004; Coolen et al., 2004b; Chu et al., 2005; D'Andrea and Huang, 2005; Liu et al., 2006b, 2008; Toney et al., in press). However, the C_{37:4} homologue was not detected in many Spanish lakes (Pearson

E.J. et al., 2008) or in lakes of the Nebraska Sand Hills (Toney et al., 2010) containing LCAs and is a minor component in Lake Tswaing, South Africa (Kristen et al., 2010). Thus, elevated abundances of the C_{37:4} homologue appear to be a characteristic feature of many, but not all, lacustrine alkenone distributions. It has also been suggested that dominance of C₃₈ homologues, and in particular of the C_{38:3} homologue, possibly can be used as an indicator of alkaline, evaporitic or saline settings based on observations from several sites (Pearson E.J. et al., 2008; Kristen et al., 2010).

2.3. Relationships between lacustrine LCAs and salinity

It has been noted that the occurrence of lacustrine LCAs appears to be related to salinity as LCAs have been reported from many lakes in evaporative settings (e.g. Li et al., 1996; Thiel et al., 1997; Wang and Zheng, 1998; Sun et al., 2004, 2007; Chu et al., 2005; D'Andrea and Huang, 2005; Theissen et al., 2005; Liu et al., 2008; Pearson E.J. et al., 2008). Additionally, high concentrations of LCAs have been reported from both sulphate (Sun et al., 2004; Pearson E.J. et al., 2008; Toney et al., 2010, in press) and carbonate-dominated lakes (Chu et al., 2005; D'Andrea and Huang, 2005; Toney et al., 2010). In west Greenland lakes, unusually high LCA concentrations are found in carbonate-dominated oligosaline lakes but LCAs are absent in nearby freshwater lakes (D'Andrea and Huang, 2005). Similarly, 12 of 13 LCA-containing lakes of the Northern Great Plains (5 lakes, all sulphate-dominated) and Nebraska Sand Hills (8 lakes, 6 carbonate-dominated, one sulfate-dominated and one lacking anion chemistry data) are characterized by elevated salinity (Toney et al., 2010). LCAs are also present in prairie lakes of interior Canada with salinities >1.5 g/L (Toney et al., in press). Pearson E.J. et al. (2008) note that LCA occurrence in Spanish lakes seems to be restricted to brackish-hypersaline sites and that LCAs are more abundant in sulphate-dominated lakes. However, LCAs have also been reported from some freshwater lakes (Zink et al., 2001; Toney et al., 2010) indicating that elevated salinity itself is not a strict requirement for LCA occurrence. *Vice versa*, LCAs also are not detected in all lakes with elevated salinity (Pearson E.J. et al., 2008; Toney et al., in press).

Several studies have reported relationships between salinity and the percentage of the C_{37:4} alkenone (%C_{37:4}) as a proportion of the total C₃₇ alkenones. Initially a correlation ($r^2 = 0.78$) between C_{37:4} production and salinity was reported from the North Atlantic and Nordic Seas (Rosell-Melé, 1998; Rosell-Melé et al., 2002; Sicre et al., 2002). However, applying the marine calibrations (Rosell-Melé et al., 2002; Sicre et al., 2002) to West Greenland lakes produced salinity values an order of magnitude larger than the measured salinity values (D'Andrea and Huang, 2005) and subsequent marine research suggested that the %C_{37:4} salinity relationship was complicated by co-varying environmental factors (Bendle et al., 2005). Nevertheless, a study of Lake Qinghai (China) and its surrounding lakes, where large changes in surface salinity occur, suggests that %C_{37:4} can qualitatively indicate regional salinity changes (Liu et al., 2008, 2011). Here, it was found that %C_{37:4} increases with decreasing surface salinity (Liu et al., 2008, 2011) while U₃₇^K values do not change significantly across the large salinity range. The authors hypothesize that at a local or regional scale the LCA-biosynthesizing species may be the same allowing for the use of C_{37:4} as a salinity indicator. However, this remains to be tested. The study of Theroux et al. (2010), which examined 15 lakes from diverse geographical regions (many of these lakes were previously examined in the studies referenced above), also does not support a linear relationship between %C_{37:4} and salinity across multiple lakes and species. The data of Toney et al. (2010) indicate no clear relationship between salinity and C_{37:4} presence or absence but instead suggest that the sulfate:carbonate ratio of the

lake water may be related to the presence/absence of $C_{37:4}$. Lakes of the Northern Great Plains and Nebraska Sand Hills lacking the $C_{37:4}$ homologue were found to have significantly lower sulfate:carbonate ratios (0.01–2.59) compared to lakes containing abundant $C_{37:4}$ (15–117) (Toney et al., 2010). Although high sulfate:carbonate ratios >3.5 also are noted in Canadian prairie lakes where the $C_{37:4}$ homologue is present (Toney et al., in press), this relationship needs to be examined at other locations. Thus, at this point, it seems the relative abundance of $C_{37:4}$ may, at best, be a qualitative indicator for salinity provided that a regional calibration is made.

2.4. Relationships between lacustrine LCAs and temperature

Relationships between water temperature and the U_{37}^k or U_{37}^k Index have been reported for many of the LCA-containing lakes and several studies have provided calibrations (Table 1; Fig. 3) providing a method for reconstructing past lake surface temperature. Marine calibrations cannot be applied to lacustrine settings since predominance of $C_{37:4}$ over $C_{37:2}$ in many lakes results in unrealistically low temperature estimates (Zink et al., 2001). Zink et al. (2001) examined surface sediments of lakes in Germany and found the best correlation between the U_{37}^k Index and the summer average lake surface temperature ($r^2 = 0.90$; Table 1). Chu et al. (2005) also examined surface sediments of lakes in China and found a good correlation between U_{37}^k and mean annual air temperature (MAAT) ($r^2 = 0.83$), which generally correlates well with annual mean lake surface temperature (Fig. 3). Chu et al. (2005) also separated the China lakes into two groups, fresh-brackish and saline (salinity >3.0 g/L), and note the best correlation between U_{37}^k in fresh-brackish lakes and mean annual air temperature ($r^2 = 90$). Interestingly, the correlation for saline lakes is considerably lower ($r^2 = 0.67$). Pearson E.J. et al. (2008) were unable to calculate the U_{37}^k and U_{37}^k Indices for surface sediments in of many lakes in Spain due to low concentrations of the C_{37} homologues. Instead they substitute C_{38} compounds, which are abundant, into the U_{37}^k and U_{37}^k Indices (see Table 1 for equations) and find a significant relationship between U_{38}^k and mean autumn air temperature ($r^2 = 0.71$). They also calculated a combination of the C_{37} and C_{38} indices (U_{3738}^k ; Table 1) and find the strongest

relationship between U_{3738}^k and mean autumn air temperature ($r^2 = 0.8$). Sun et al. (2007) examined U_{37}^k and U_{37}^k values from a cultivated strain of *C. lamellosa* isolated from a lake in China. They found that U_{37}^k values were strongly correlated with growth temperature varying from 10 to 22 °C. A second order polynomial regression of the U_{37}^k index vs. water temperature produced the best correlation ($r^2 = 0.99$; Table 1) but a linear regression also results in a similar correlation ($r^2 = 0.97$). Interestingly, the slopes and intercepts of the calibration equations for the lacustrine *C. lamellosa* culture are considerably different from the calibration of a marine *C. lamellosa* species (Fig. 3). Toney et al. (2010) provide an *in-situ* calibration for U_{37}^k for Lake George, ND covering a temperature range from 2 to 22 °C based on filtered water samples. They found a significant relationship between the U_{37}^k Index and water temperature ($r^2 = 0.75$; Table 1) but in contrast to other studies, they found no relationship between U_{37}^k and water temperature ($r^2 = 0.14$). Toney et al. (2010) also note that the proportion of the $C_{37:4}$ alkenone, which is dominant in Lake George, is also linearly correlated to temperature ($r^2 = 0.73$). D'Andrea et al. (2011) created a U_{37}^k calibration for lake Braya SØ (Greenland) by combining data from filtered water samples with the core top U_{37}^k calibration of Zink et al. (2001) ($r^2 = 0.96$; Table 1). Thus, at this point there is no uniform calibration of U_{37}^k or U_{37}^k against lake surface temperature, which is likely related to the presence of different haptophyte producers at different sites. The presence of multiple haptophyte species within a lake is also a concern for lacustrine LCA calibrations; however, it has been suggested that if the haptophyte species are closely related, a single U_{37}^k or U_{37}^k calibration can be used for paleoclimate reconstructions if an *in-situ* or culture calibration is available (Theroux et al., 2010).

In addition to variability resulting from different haptophyte producers, a number of other factors are known to influence the relationship between U_{37}^k or U_{37}^k and temperature in marine environments (see Herbert, 2003 for a review), which likely also affect lacustrine environments. For example, seasonality may influence temperature reconstructions based on LCAs due to a seasonal production maximum for LCA-producing organisms. Sun et al. (2007) note the presence of *C. lamellosa* in phytoplankton samples collected from Lake Xiarinur (China) in spring and autumn whereas water column filtering of Lake George, ND, reveals alkenone production after ice out in the early spring (Toney et al., 2010). Similarly, alkenone fluxes to sediment traps in a Greenland lake indicates the haptophyte bloom occurring between June and mid-July (D'Andrea et al., 2011). Indeed, some of the available lacustrine LCA calibrations also report the best correlation with a seasonal temperature rather than with MAAT (Zink et al., 2001; Pearson E.J. et al., 2008). Thus, seasonal influences on temperature reconstructions from lacustrine alkenones are likely and should be evaluated for individual sites.

Non-thermal physiological growth factors including nutrient and light limitation are known to lead to significant variations in U_{37}^k values independent of temperature, and have been observed in culture experiments and in marine sediments (e.g. Prahl et al., 2006; Placencia et al., 2010). Batch culture experiments with *E. huxleyi* show that U_{37}^k values decrease under nutrient stress conditions and increase under prolonged dark stress conditions (Prahl et al., 2006). By changing nitrate and phosphate concentrations from 83.1 and 11.5 μM to 0.0 and 0.5 μM , respectively, Prahl et al. (2006) observed that U_{37}^k values decreased from 0.579 to 0.493, equivalent to a temperature change of 3 °C using the calibration of Müller et al. (1998). In the dark stress experiment at 15 °C a sample grown under a 12:12 light:dark cycle was compared to samples grown under 24 h of darkness, and it was found that U_{37}^k values increased by up to 0.042, equivalent to a temperature change of 1.3 °C (Prahl et al., 2006). Placencia et al. (2010) found

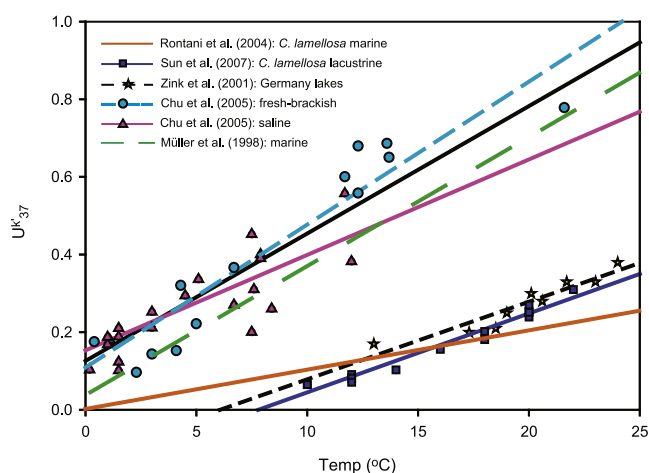


Fig. 3. Comparison of lacustrine alkenone calibration equations for the U_{37}^k Index. The marine calibrations of *E. huxleyi* (Prahl and Wakeham, 1987) and *C. lamellosa* (Conte et al., 1994) are shown for comparison. Lacustrine U_{37}^k calibrations include a calibration developed for surface sediments of lakes in China (solid black line; Chu et al., 2005), the same set of lakes divided into fresh-brackish and saline lakes (Chu et al., 2005), the calibration for Lake Braya SØ, Greenland (Theroux et al., 2010 and references therein), the calibration for lakes in Germany (Zink et al., 2001) and a calibration (linear and polynomial fit) for a cultivated strain of *C. lamellosa* isolated from a lake in China (Sun et al., 2007).

anomalously cold U_{37}^k values along the Peru-Chile margin where permanent upwelling and complete nutrient utilization by phytoplankton results in low surface nitrate ($<2 \mu\text{M}$) and chlorophyll *a* ($<1 \text{ mg m}^{-3}$) concentrations. Thus, it seems likely that nutrient concentrations and upwelling can also influence lacustrine alkenone distributions while light limitation may be especially important to ice-covered lakes. These processes probably account for some of the scatter seen in lacustrine alkenone calibrations.

Finally, several marine and laboratory studies suggest diagenesis, and in particular, oxic degradation may influence U_{37}^k records, with preferential degradation of the $C_{37:3}$ ketone occurring, resulting in slightly warmer temperatures. For a review of these issues we refer to reader to Herbert (2003) as, to our knowledge, these issues have not yet been specifically examined for lacustrine environments.

2.5. Paleoclimate reconstructions based on lacustrine LCAs

There is particular interest in utilizing lacustrine LCAs for paleotemperature reconstructions, and indeed, several studies have utilized this approach. For example, Liu et al. (2006b) utilized LCAs to examine both temperature and salinity changes in Late-Holocene sediments of Lake Qinghai (China). U_{37}^k values revealed fluctuating periods of warmer and cooler temperatures that could be related to known climate events including warming during the 20th century, the Little Ice Age, the Medieval Warm Period, the Dark Ages Cold Period, and the Roman Warm Period (Fig. 4a). Fluctuations in $\%C_{37:4}$ provided evidence of salinity changes, which

were found to be coupled to temperature changes, with fresher lake waters noted during warm periods. The authors related this pattern to the Asian monsoon exerting a large influence on the climate of the Lake Qinghai region (Liu et al., 2006b) (Fig. 4a).

D'Andrea et al. (2011) created U_{37}^k records for two lakes (Braya Sø and Lake E) in West Greenland after developing an in-situ temperature calibration from Braya Sø (Fig. 4b,c). They found the U_{37}^k records of both lakes to be in close agreement, displaying temperature changes of up to 5.5°C over the past 5600 years and an overall pattern similar to that of the GISP2 ice core. Interestingly, major and abrupt temperature changes, occurring within decades, were found to coincide with archaeological records indicating the settlement and abandonment of Saqqaq, Dorset and Norse cultures (Fig. 4b). D'Andrea et al. (2011) hypothesize that these significant temperature variations may be associated with the North Atlantic Oscillation.

2.6. Outlook on the application of lacustrine LCAs to Quaternary studies

Despite the fact that LCAs have been reported from globally distributed lakes, to date, relatively few studies have utilized lacustrine LCAs for paleoclimate reconstructions and to our knowledge, no studies have yet used alkenone concentrations and an indicator of past haptophyte productivity although this is commonly done in marine settings. There are several reasons for the small number of paleoclimate reconstructions based on lacustrine LCAs. First, alkenone unsaturation ratios correspond

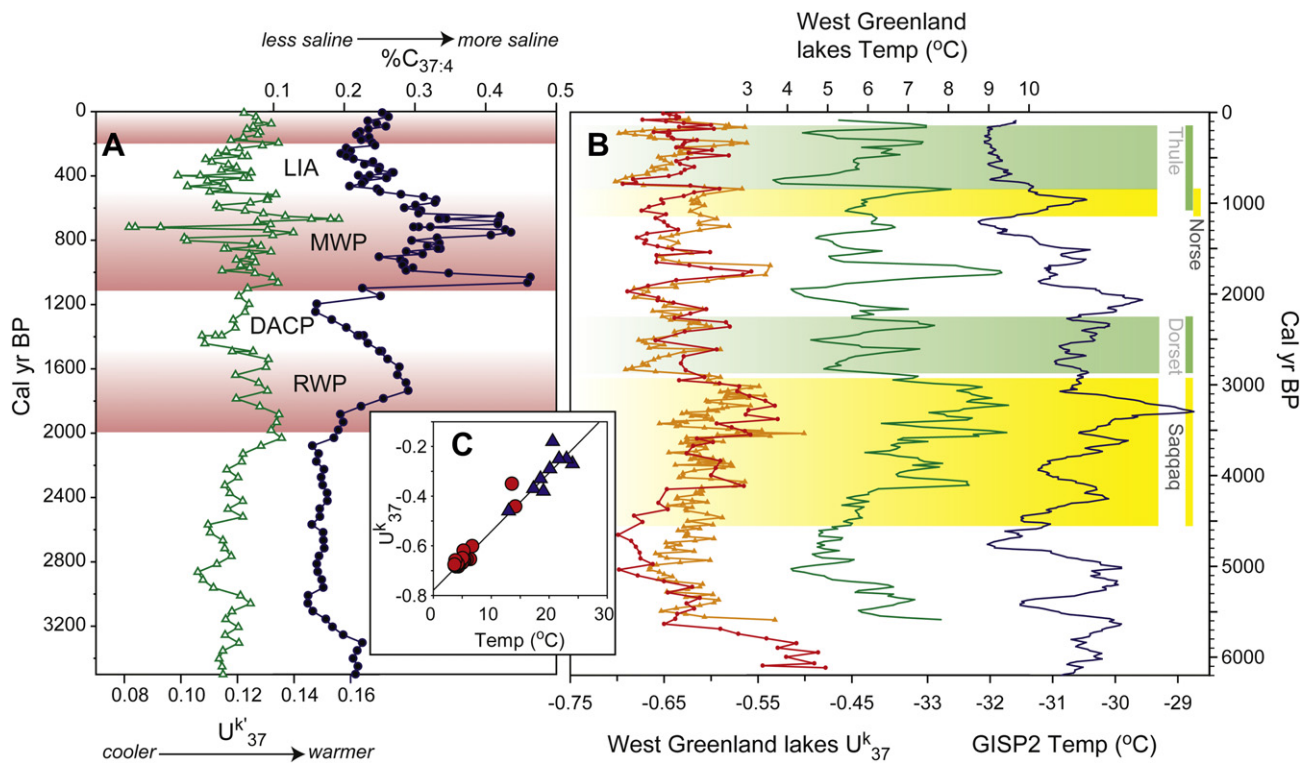


Fig. 4. Examples of lacustrine alkenone records. A) U_{37}^k and $\%C_{37:4}$ records of Lake Qinghai, China (data from Liu et al., 2006b). These records reveal a climate response to known climate events including 20th century warming, the Little Ice Age (LIA), the Medieval Warm Period (MWP), the Dark Ages Cold Period (DACP), and the Roman Warm Period (RWP). Warmer temperatures are found in association with fresher lake waters and is likely related to the influence of the Asian monsoon on the climate of Lake Qinghai. B) U_{37}^k temperature records of lakes in West Greenland (data from D'Andrea et al., 2011). The plot on the left shows U_{37}^k values for Braya Sø (red circles) and Lake E (orange triangles) in the Kangerlussaq region of West Greenland. The middle plot (green line) shows the stacked U_{37}^k records for these lakes, converted to temperature using a calibration developed by the authors (shown in C). The plot on the right (dark blue) shows a temperature reconstruction for the GISP2 Ice Core (Alley, 2004). The presence of the Saqqaq, Dorset Norse and Thule cultures is indicated by the yellow and green shading. C) U_{37}^k temperature calibration for Greenland lake Braya Sø ($r^2 = 0.96$), which was created by combining data from filtered water samples (red circles) with the core top U_{37}^k calibration of Zink et al. (2001) (blue triangles). Data from D'Andrea et al. (2011).

with temperature in only a small subset of LCA-containing lakes. Second, the occurrence of LCAs is episodic in some lakes and thus cannot be used to generate continuous climate records (Zink et al., 2001; Theissen et al., 2005). Furthermore, there is no generally applicable correlation of LST with alkenone distribution and one has to rely on regional calibrations. Nevertheless, several studies have generated paleoclimate records based on LCAs and for sites lacking calibrations, either U_{37}^k or U_{37}^k values have been used to examine relative temperature changes (Thiel et al., 1997; e.g. Wang and Zheng, 1998; Theissen et al., 2005; Liu et al., 2006b). Studies such as that of D'Andrea et al. (2011) demonstrate that high quality temperature records can be produced from lacustrine LCA distributions, providing important insights into past continental temperature variability. However, it is important to recognize that before these temperature reconstructions could be made, extensive background research was conducted on the Greenland lakes including filtering water column samples over multiple years to develop a temperature calibration (D'Andrea et al., 2011) and genetic studies to demine the main alkenone producing organism (D'Andrea et al., 2006). Such studies are essential for examining relationships between lacustrine LCA distributions and temperature (or salinity) and similar studies need to be conducted in a wider variety of locations. Additionally, identifying the main alkenone producers in other lakes where LCAs have been reported, and isolating and bringing them into culture, likely would facilitate utilizing these compounds as paleoenvironmental proxies.

3. Reconstructing temperature and soil organic matter input from glycerol dialkyl glycerol tetraethers (GDGTs)

Isoprenoid glycerol dialkyl glycerol tetraethers (GDGTs) are membrane lipids diagnostic for certain groups within the Archaea. These GDGTs consist of an isoprenoid carbon skeleton ether bonded to a glycerol-moiety containing 0–3 cyclopentane rings (GDGTs I–III) (Fig. 5) and are of interest since they provide proxies for reconstructing marine and lake surface temperatures, and for examining inputs of soil organic matter to marine or lacustrine environments. A large number of phylogenetic groups within the domain Archaea synthesize isoprenoid GDGTs (Schouten et al., 2007b and references therein). Archaeal 16S rRNA sequences are also ubiquitous in normal marine waters and are comprised mainly of Thaumarchaeota (formerly Group I Crenarchaeota; see Brochier-Armanet et al., 2008; Spang et al., 2010) and Group 2 Euryarchaeota (DeLong, 1992; Fuhrman et al., 1992; Karner et al., 2001; Lipp et al., 2008). The GDGT “crenarchaeol” (GDGT IV), which contains four cyclopentane moieties and a cyclohexane moiety (Fig. 5), is suggested to be specific to the Thaumarchaeota (Pitcher et al., 2011 and references therein). 16S rRNA gene sequences related to the marine Thaumarchaeota have also been reported in lakes (e.g. Keough et al., 2003; Sinninghe Damsté et al., 2009) in agreement with the finding of crenarchaeol and GDGTs 0–III in the water columns and sediments of lakes (e.g. Powers et al., 2004, 2010; Tierney et al., 2008; Blaga et al., 2009; Sinninghe Damsté et al., 2009).

Another group of GDGTs, the branched GDGTs, contain methyl-substituted C_{28} *n*-alkyl side chains that contain 4, 5 or 6 methyl substitutes (Fig. 5) (Sinninghe Damsté et al., 2000) and are of interest since these compounds provide proxies for reconstructing mean annual soil temperature and soil pH (Weijers et al., 2007b). Branched GDGTs were first identified in peats (Sinninghe Damsté et al., 2000) and were later found to be ubiquitous and dominant in soils (Weijers et al., 2006b). In contrast to the isoprenoid GDGTs that are produced by Archaea, it is thought that branched GDGTs are biosynthesized by anaerobic soil bacteria (Weijers et al., 2006a; Sinninghe Damsté et al., 2011) although branched GDGTs have not

yet been identified as the main membrane lipids in cultures and the exact source organism(s) remains unknown.

GDGTs provide good biomarkers as they are found in sediment back until at least the late Jurassic (Carillo-Hernandez et al., 2003). The various proxies based on isoprenoid and branched GDGTs (discussed in Sections 3.1–3.3) are all relatively new, beginning with the study of Schouten et al. (2002), and therefore much remains to be learned. However, these proxies have proven to be important indicators of past climate and thus have been widely applied to both marine and lacustrine sediments. Here, we examine application of these proxies to lacustrine environments.

3.1. The BIT Index as a soil organic matter tracer

The Branched and Isoprenoid Tetraether (BIT) index provides a proxy for aquatic vs. soil organic matter input and is based on relative abundances of soil derived (branched) GDGTs vs. crenarchaeol (a biomarker for aquatic Thaumarchaeota) (Fig. 5; Table 2) (Hopmans et al., 2004). BIT values range from 0 to 1, with a value of 0 indicating a purely marine (or aquatic) source in which only crenarchaeol is present, while a value of 1 represents pure soil OM source, consisting of only the branched GDGTs (Hopmans et al., 2004). However, since small to moderate amounts of crenarchaeol also may be present in soils and peat, soil OM is generally characterized by BIT values of >0.9 (Weijers et al., 2006b). In marine settings, high BIT values are noted near river mouths and decrease offshore while low BIT values are observed in open marine settings (Hopmans et al., 2004; Kim et al., 2010a). Thus, the BIT index can be useful for examining the delivery of soil organic matter to marine or lacustrine systems. Branched GDGTs are transported from terrestrial to aquatic environments by riverine input or surface runoff and are present in low abundance in open marine settings due to small amounts of *in-situ* production (Peterse et al., 2009). To date, branched GDGTs have not been detected in dust samples (Hopmans et al., 2004). It is important to note that in contrast to other terrestrial proxies (e.g. $\delta^{13}C_{org}$, C/N ratios), the BIT Index does not carry a large imprint of vegetation (Huguet et al., 2007; Walsh et al., 2008; Belicka and Harvey, 2009; Smith et al., 2010) as branched GDGTs are not produced by plants. Therefore, examining the BIT index in combination with other terrestrial proxies or biomarkers provides opportunities for examining the transport of soil organic matter and vegetation to marine or lacustrine environments.

Many lakes, especially smaller lakes, are generally characterized by a large watershed relative to the surface area of the lake. Thus, it is not surprising that many lakes are characterized by high BIT Index values. In a global survey of GDGTs in lakes (Powers et al., 2010) and a survey of GDGTs in European lakes (Blaga et al., 2009), it was found that $>75\%$ had high BIT values (>0.5). However, in lakes, high BIT values may also result from *in-situ* production of branched GDGTs in the water column. Several studies of lakes in East Africa (Sinninghe Damsté et al., 2009; Tierney et al., 2010b), Indonesia (Tierney and Russell, 2009); New Zealand (Zink et al., 2010) and Europe (Blaga et al., 2009; Bechtel et al., 2010) have concluded that *in-situ* production of branched GDGTs likely occurs in lakes, suggesting that they are biosynthesized by a source in addition to anaerobic soil bacteria. Branched GDGTs have been found in the water columns of both oxic and anoxic lakes (Sinninghe Damsté et al., 2009; Bechtel et al., 2010) and in oligotrophic and eutrophic lakes (Bechtel et al., 2010). As it appears that *in-situ* production of branched GDGTs is widespread in lakes, the BIT Index may not solely reflect soil organic matter input to all lakes, and in many lakes it is currently not clear what the signal represents.

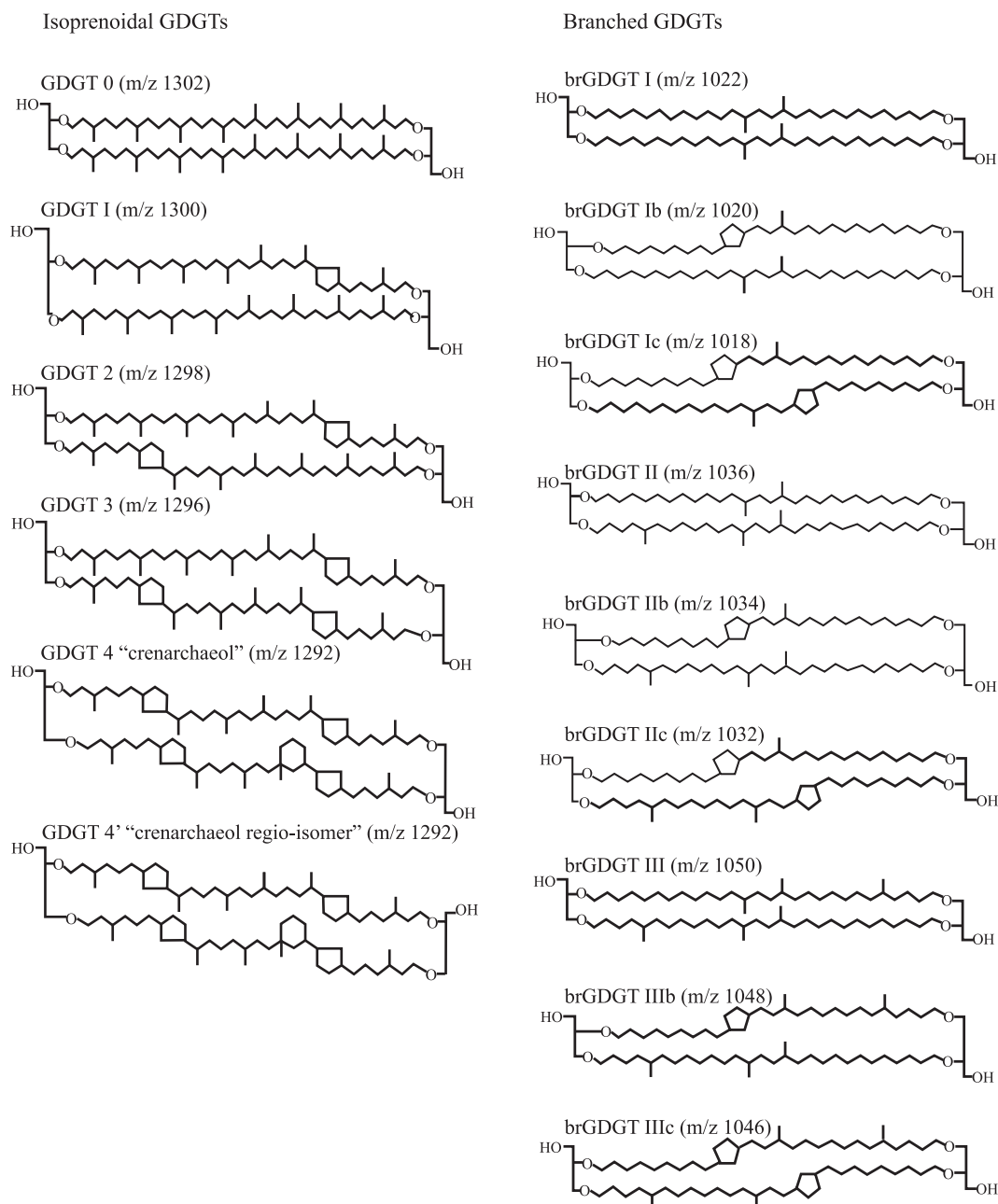


Fig. 5. Structures of the isoprenoid and branched glycerol dialkyl glycerol tetraethers (GDGTs) used for calculating TEX_{86} , the BIT Index, and MBT/CBT.

Table 2

Formulas for GDGT-based proxies including the BIT Index (Hopmans et al., 2004), TEX_{86} (Schouten et al., 2002), $\text{TEX}_{86}^{\text{H}}$ and $\text{TEX}_{86}^{\text{L}}$ (Kim et al., 2010b) and MBT and CBT (Weijers et al., 2007b).

$$\text{BIT} = \frac{[\text{GDGT I} + \text{GDGT II} + \text{GDGT III}]}{[\text{crenarchaeol} + \text{GDGT I} + \text{GDGT II} + \text{GDGT III}]}$$

$$\text{TEX}_{86} = \frac{[\text{GDGT2} + \text{GDGT3} + \text{GDGT4}']}{[\text{GDGT1} + \text{GDGT2} + \text{GDGT3} + \text{GDGT4}']}$$

$$\text{TEX}_{86}^{\text{L}} = \log \frac{[\text{GDGT2}]}{[\text{GDGT1} + \text{GDGT2} + \text{GDGT3}]}$$

$$\text{TEX}_{86}^{\text{H}} = \log \frac{[\text{GDGT2} + \text{GDGT3} + \text{GDGT4}']}{[\text{GDGT1} + \text{GDGT2} + \text{GDGT3} + \text{GDGT4}']}$$

$$\text{MBT} = \frac{[\text{GDGT I} + \text{GDGT Ib} + \text{GDGT Ic}]}{[\sum \text{all branched GDGTs}]}$$

$$\text{CBT} = \frac{[\text{GDGT Ib} - \text{GDGT IIb}]}{[\text{GDGT I} + \text{GDGT II}]}$$

A study from Lake Challa (Kenya) in which water column, sediment trap and surface sediment samples were analyzed for GDGTs offers some insights into processes influencing the distribution of GDGTs in lakes. In Lake Challa, the major settling flux of branched GDGTs coincides with the period of maximum precipitation and it was also observed that surface sediments located near the periphery of the lake had higher BIT values than surface sediments located in the middle of the lake (Sinninghe Damsté et al., 2009). Both of these observations support the hypothesis that delivery of branched GDGTs to this lake occurs through erosion of soils in the lake catchment (Sinninghe Damsté et al., 2009). However, it was also found that concentrations of branched GDGTs were 1–2 orders of magnitude higher in the deep waters of Lake Challa compared to the surface waters. The authors conclude that sediment resuspension is unlikely and thus either *in-situ* production occurs in the anoxic lower water column or branched GDGTs

are delivered to the deep waters of the lake via groundwater inputs (Sinninghe Damsté et al., 2009).

Verschuren et al. (2009) used the BIT Index record as a proxy of rainfall-induced surface runoff at Lake Challa. They argue that because the lake is a steep-sided crater basin, even major fluctuations in lake level do not significantly change the catchment area and thus variations in the BIT Index mainly reflect changes in surface runoff. Variations in the BIT record, complemented by evidence from seismic-reflection stratigraphy, were found to be in agreement with lake-level changes noted at other lakes in the region. Thus, the authors suggest that on longer time scales BIT Index values are proportional to rainfall over the Lake Challa drainage basin. The Lake Challa records indicate that monsoonal rainfall in East Africa varied at half-percentage intervals, which were in phase with orbitally controlled insolation forcing (Verschuren et al., 2009).

When interpreting BIT Index records, it is important to note that variability in crenarchaeol concentrations may also drive changes in the BIT Index rather than the influx of branched GDGTs. This has been noted at some marine (Castañeda et al., 2010; Smith et al., 2010) and lacustrine (Tierney et al., 2010b) sites where BIT values were found to mainly reflect crenarchaeol production rather than the branched GDGT flux. Therefore, in order to utilize the BIT index to examine soil organic matter input to marine or lacustrine environments it is essential to determine absolute concentrations of GDGTs (Huguet et al., 2006b). Furthermore, one should be aware that *in-situ* contributions of branched GDGTs may complicate interpretation of past BIT variations, especially when branched GDGT concentrations are low.

3.2. MBT and CBT as proxies for air temperature and soil pH

In addition to the BIT Index, two more recently developed proxies also based on branched GDGTs (Fig. 5) are the Cyclisation of Branched Tetraethers (CBT) and the Methylation of Branched Tetraethers (MBT) (Table 2) (Weijers et al., 2007b). CBT provides a proxy for soil pH and is based on the observation that the relative number of cyclopentane moieties of branched GDGTs is exponentially negatively correlated with soil pH (Weijers et al., 2007b). MBT is primarily positively correlated with mean annual soil temperature, which in many cases is similar to mean annual air temperature (MAAT), but, to a lesser extent, is also negatively correlated with soil pH (Weijers et al., 2007b). To reconstruct MAAT, soil pH is first obtained from the CBT equation and then this value is used to correct the MBT equation (Tables 2 and 3) for the influence of soil pH (Weijers et al., 2007b).

The MBT/CBT calibrations are based on a dataset of globally distributed soils consisting of 134 samples from 90 different locations (Weijers et al., 2007b). The uncertainties with reconstructing soil pH from CBT are $\sim \pm 1$ pH unit and ~ 5 °C for calculating absolute MAAT from MBT/CBT. However, it has been noted that local or regional calibrations may be necessary for reconstructing absolute MAATs (Weijers et al., 2007b; Sinninghe Damsté et al., 2008). It is also worth noting that a number of factors are currently poorly constrained with the MBT/CBT proxy. The

relatively large uncertainties in the calibration equations suggest that other factors besides temperature and soil pH are affecting the distribution of branched GDGTs. Indeed, Weijers et al. (2007b) noted a weak correlation ($r^2 = 0.47$) between MBT and precipitation.

Initially, application of the MBT/CBT proxies in lake sediments to reconstruct continental MAAT appeared particularly promising in small lakes with high BIT values where TEX_{86} could not be applied. However, recent research demonstrated that *in-situ* production of branched GDGTs from within the water column of lakes also affects MBT/CBT, as it does the BIT Index (Sinninghe Damsté et al., 2009; Tierney and Russell, 2009; Bechtel et al., 2010; Blaga et al., 2010). Tierney and Russell (2009) investigated branched GDGT distributions in Lake Towuti, Indonesia, and in soils from its watershed. They found that CBT-inferred pH values appeared to reflect the pH of the lake water rather than the surrounding soils, predicted temperatures from MBT/CBT were significantly colder than MAAT, and that concentrations of branched GDGTs were an order of magnitude higher in the lake sediments than the surrounding soils. Based on these observations, they suggested that branched GDGTs are produced *in-situ* in lakes, similar to the results of Peterse et al. (2009) who noted *in-situ* production of branched GDGTs in a Svalbard fjord. Blaga et al. (2009) investigated branched GDGT distributions in the surface sediments of 82 globally distributed lakes of variable water depth and size and conclude that branched GDGTs in lakes are derived from mixed (soil and *in-situ*) sources. Tyler et al. (2010) examined MBT/CBT in Lake Lochnagar (Scotland) and found that CBT-inferred pH was in good agreement with diatom-inferred pH but that MBT/CBT significantly underestimated MAAT by ~ 5 °C. Bechtel et al. (2010) examined GDGTs in eutrophic Lake Lugano and oligotrophic Lake Brienz, Switzerland, in cores covering the interval from ~ 1900 AD to the present. They found that branched GDGTs at Lake Brienz appear to be mainly soil derived and that reconstructed MAAT was comparable to nearby weather station data. In contrast, at Lake Lugano, where the branched GDGTs appear to be at least partially *in-situ* produced, MBT/CBT significantly underestimates MAAT (by up to ~ 10 °C) but instead resembles the temperature of the deep lake quite well (Bechtel et al., 2010).

Despite the fact that *in-situ* production of branched GDGTs likely occurs in variable amounts in different lakes, several lacustrine studies suggest that it is still possible to reconstruct MAAT using the MBT/CBT proxies. Zink et al. (2010) examined branched GDGT distributions in eleven New Zealand lakes ranging from 101 to 2000 m above sea level. Interestingly, they found that in these lakes, where *in-situ* production of branched GDGTs appears significant, the CBT Index was mainly controlled by temperature, rather than pH, and that both MBT and CBT indices indicated relationships with MAAT. They used their data to produce new calibration equations for MBT ($n = 10$, $r^2 = 0.74$) and CBT ($n = 9$, $r^2 = 0.72$; Table 3) and found that reconstructed MAATs show a clear covariance with altitude, in general agreement with weather station data and chironomid-based temperature reconstructions. Zink et al. (2010) also applied their MBT calibration to sediments of Alpine Lake to examine temperature changes during the past

Table 3
MBT and CBT calibrations for New Zealand and East African lakes compared to the global soils calibration dataset of Weijers et al. (2007).

Calibration equation	<i>n</i>	r^2	Geographical location	Reference
$\text{MAAT} = 55.01 \times \text{MBT} - 6.055$	10	0.74	New Zealand	Zink et al. (2010)
$\text{MAAT} = -6.567 \times \text{CBT} + 12.228$	9	0.72	New Zealand	Zink et al. (2010)
$\text{MAAT} = 50.47 - 74.18 \times (f\text{GDGT III}) - 31.60 \times (f\text{GDGT II}) - 34.69 \times (f\text{GDGT I})^a$	38	0.94	East Africa	Tierney et al. (2010b)
$\text{CBT} = 3.33 - 0.38 \times \text{pH}$	134	0.70	Global soils	Weijers et al. (2007b)
$\text{MBT} = 0.122 + 0.187 \times \text{CBT} + 0.020 \times \text{MAAT}$	134	0.77	Global soils	Weijers et al. (2007b)

^a Note that "*f*" denotes the fractional abundances of the branched GDGTs relative to the total branched GDGTs (includes GDGTs I, Ia, Ib, II, IIa, IIb, III, IIIa, IIIb).

29,000 years and found their results were similar to a chironomid-based temperature reconstruction from a nearby lake. Tierney et al. (2010b) examined branched GDGTs in 46 East African lakes. They noted that applying the global soils calibration to the East African lake sediments produced cooler than expected MAATs and thus present a different type of calibration, i.e. by using weighing factors for different GDGTs, for these lakes ($n = 38$, $r^2 = 0.94$; Table 3). Tierney et al. (2010b) find that temperature and pH are the main factors that can be related to variations in the distribution of branched GDGTs and suggest that despite being from mixed sources, branched GDGTs can be used to examine MAAT in the East African lakes.

Although uncertainties exist regarding the origin of branched GDGTs in lakes, at some sites there is strong evidence that the MBT/CBT paleothermometer indeed reflects MAAT. Fawcett et al. (2011) conducted a multi-proxy study of a lacustrine sediment core from Valles Caldera (New Mexico) and used MBT/CBT to examine MAAT and soil pH during Marine Isotope Stages (MIS) 10–14 (~368–552 Kyr). They found that the glacial stages were characterized by millennial-scale temperature variations of up to 7 °C. Stadials were characterized by cooler temperatures and increased percentages of boreal *Picea* + *Abies* pollen while interstadials were marked by warmer temperatures and increased percentages of *Quercus* and *Juniperus* pollen (Fig. 6). In addition, CBT reconstructions of soil pH indicate that arid conditions occurred during warm intervals (MIS 11 and 13) while calcium counts from core scanning XRF indicate closed-basin conditions at these times (Fig. 6). The Valles Caldera records display remarkable

agreement between branched GDGT, pollen and core scanning data, and show that extended “dust-bowl-like” megadroughts, lasting for centuries to millennia, occurred in the southwestern United States during MIS 11 and 13 (Fawcett et al., 2011).

3.2.1. Outlook for lacustrine MBT/CBT reconstructions

Overall, many uncertainties currently exist regarding sources of branched GDGTs to lakes and thus the application of the MBT/CBT paleothermometer to lakes. Studies such as Zink et al. (2010) and Tierney et al. (2010b) are promising as they suggest that branched GDGTs can be used to reconstruct MAAT, using local calibrations. A key factor in determining whether MBT/CBT is applicable for reconstructing MAAT in lakes may be the relative proportion of soil- and lacustrine-derived GDGTs. Although branched GDGTs in lakes reflect mixed sources (Blaga et al., 2010; Tierney et al., 2010b) lakes where either soil- or lacustrine-derived GDGTs dominate may be particularly suitable for reconstructing MAAT or water column temperatures, respectively. However, lake water temperatures are often strongly correlated with MAAT (Livingstone and Lotter, 1998) and thus lakes with mixed sources of branched GDGTs may also be suitable for reconstructing MAAT, depending on the geographical location of lake or the water depth of maximum GDGT production. Tropical lakes may be better suited for MBT/CBT temperature reconstructions because surface water temperatures are generally similar to MAAT and also smaller differences exist between surface and bottom water temperatures. In subtropical or temperate lakes, where the temperature difference between surface and bottom waters is greater, larger differences between MBT/CBT-inferred

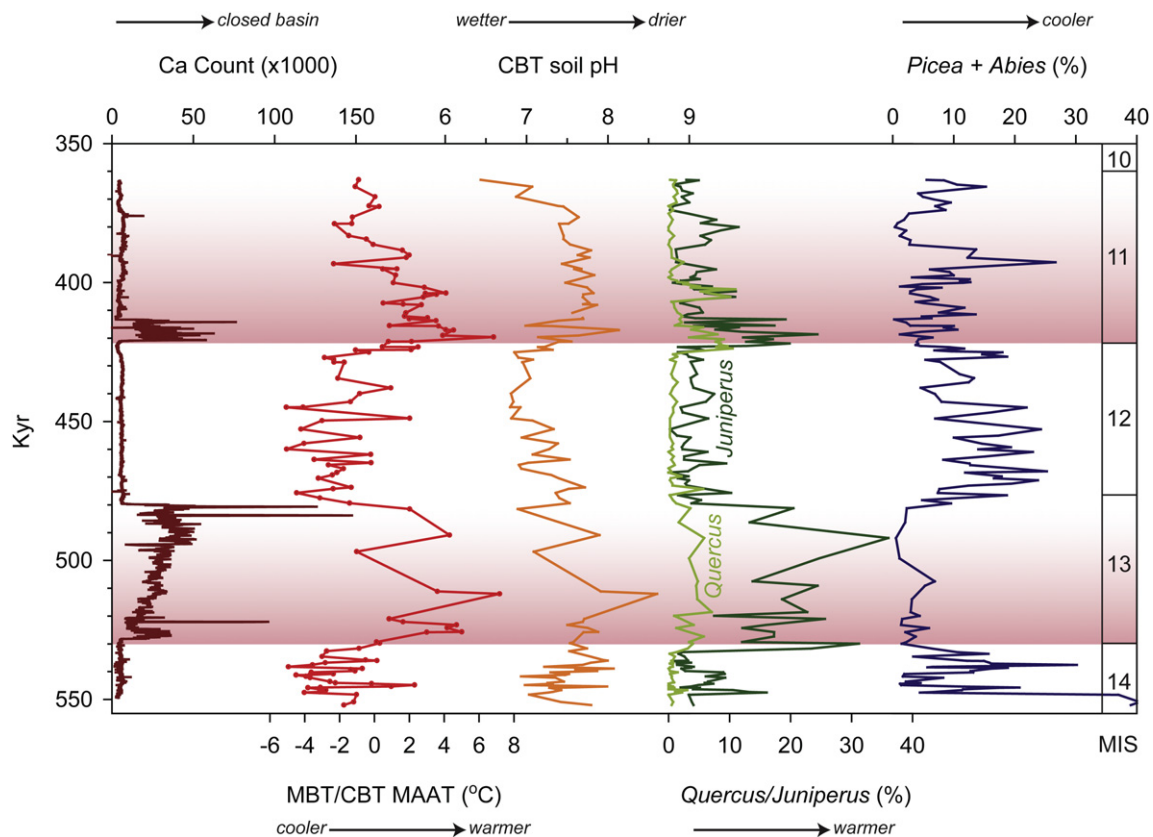


Fig. 6. Multi-proxy data from Valles Caldera, New Mexico (Fawcett et al., 2011). Calcium counts from core scanning XRF are plotted in dark red. MBT/CBT reconstructions of mean annual air temperature (MAAT) are plotted in red, CBT reconstructions of soil pH are plotted in orange. Percent *Quercus* and *Juniperus* pollen are plotted in light and dark green, respectively, while boreal pollen (*Picea* + *Abies*) is plotted in dark blue. Marine Isotope Stages (MIS) 10–14 are indicated on the right side of the graph and interstadials are highlighted in red. High calcium concentrations indicate periods of closed-basin conditions during MIS 11 and 13 and coincide with warmer MAAT, elevated soil pH (indicating arid conditions) and increased percentages of *Quercus* and *Juniperus* (warm) pollen. In contrast, boreal pollen displays increased percentages during the stadials (MIS 12 and 14).

temperature and MAAT may exist, depending on the location of branched GDGT productivity within the water column. It is also likely that seasonal variations in GDGT production could exist for some lakes and may influence MAAT reconstructions. Given that these proxies are only a few years old (Weijers et al., 2007b), a number of outstanding questions exist that remain to be answered by future studies. Clearly, the applicability of the MBT/CBT proxies needs to be assessed for individual sites but may be useful for temperature reconstructions in some lakes.

We note that when evaluating MBT/CBT records, it is important to consider that the age of the branched GDGTs may differ in comparison to marine (or lacustrine) derived components. Branched GDGTs are formed on land and thus may arrive in the marine (or lacustrine) realm with a significant time lag, although Weijers et al. (2007a) did not note a substantial time lag for Late Quaternary marine sediments collected from the Congo River fan. A related issue is that the sources of soil OM to marine (or lacustrine) sediments may have changed over time. This has been noted by an MBT/CBT study of the Amazon River fan, in which warmer MBT/CBT MAATs were noted during the LGM compared to the present and attributed to a change in source region from the Amazon flood plain to the Andes mountains (Bendle et al., 2010). Therefore, the transport pathways of branched GDGTs to marine or lacustrine settings should be evaluated for individual sites.

3.3. The TEX_{86} paleothermometer

The TEX_{86} (TetraEther index with 86 carbon atoms) (Schouten et al., 2002) temperature proxy is based on the relative abundance of isoprenoidal GDGTs, which are mainly biosynthesized by Thaumarchaeota (Fig. 5). The basic premise behind the TEX_{86} is that the number of cyclopentane moieties increases with increasing growth temperature (Fig. 7). The two most abundant isoprenoidal GDGTs, GDGT-0 and crenarchaeol (Fig. 5), are excluded from the

TEX_{86} to avoid their overpowering influence on the index since their abundances are generally much higher compared to the other GDGTs (Schouten et al., 2002). In addition, GDGT-0 is known to be produced by other types of Archaea, including methanogenic and methanotrophic Euryarchaeota, such as those living in anoxic sedimentary environments (Schouten et al., 2007b).

3.3.1. TEX_{86} temperature calibrations

The most recent marine calibration for TEX_{86} is based on a dataset of 426 core top samples and utilizes a log calibration (Kim et al., 2010b). Since it appears that the crenarchaeol regio-isomer is insensitive to temperature changes in the polar oceans, two calibration equations were developed for TEX_{86} for application in low ($<15\text{ }^{\circ}\text{C}$; TEX_{86}^L) and high ($>15\text{ }^{\circ}\text{C}$; TEX_{86}^H) temperature marine settings (Table 4; Kim et al., 2010b). TEX_{86} has also been shown to work in some lakes (e.g. Powers et al., 2004, 2010; Tierney et al., 2008; Blaga et al., 2009). In contrast to the relatively large dataset for the marine calibration, the present linear calibration of TEX_{86} in lakes is based on a much smaller dataset (Table 4). Powers et al. (2004) first investigated application of TEX_{86} in four lakes and found that the samples fit on the initial marine calibration of Schouten et al. (2002) and reflected mean annual lake surface temperature. More recently, Powers et al. (2010) generated a TEX_{86} calibration for lakes based on 12 samples (Fig. 8), which derive mostly from medium to large lakes ($r^2 = 0.86$; Table 4). Tierney et al. (2010a) added five additional samples, all with temperatures $>23\text{ }^{\circ}\text{C}$, to the calibration of Powers et al. (2010) and removed samples with temperatures $<10\text{ }^{\circ}\text{C}$ to produce a lacustrine TEX_{86} calibration covering a temperature range from 11 to $30\text{ }^{\circ}\text{C}$ ($n = 13$, $r^2 = 0.92$; Table 4). Combining all lakes used in the calibrations of Powers et al. (2010) and Tierney et al. (2010a) also yields a temperature calibration with a good correlation ($n = 19$, $r^2 = 0.89$; Fig. 8). The root mean square error for the different TEX_{86} calibrations ranges from 1.7 to $4.0\text{ }^{\circ}\text{C}$ (Table 4) while TEX_{86} measurements

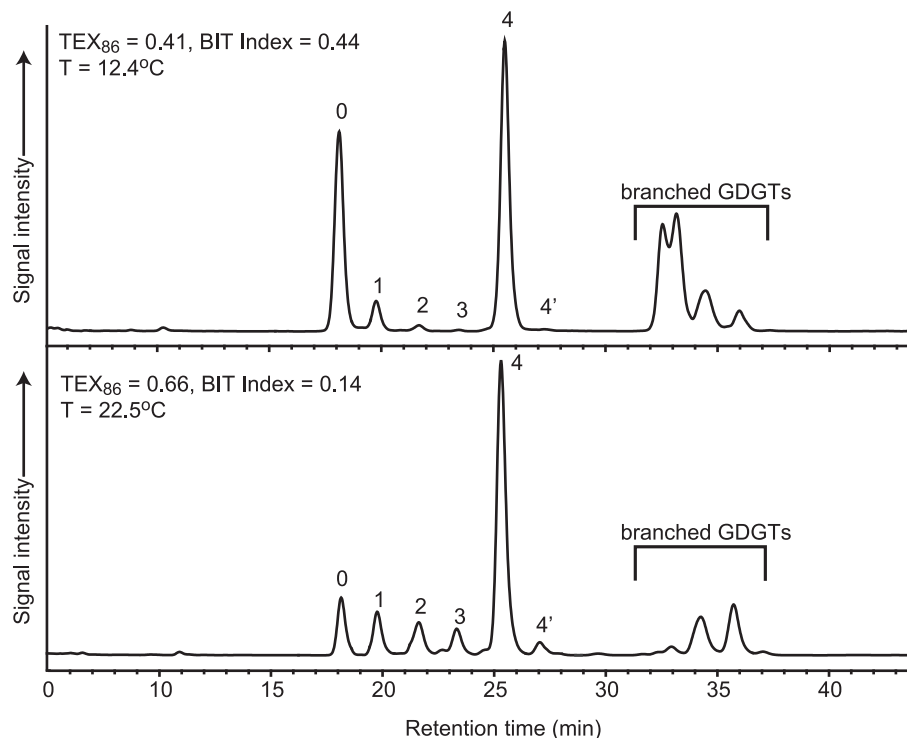


Fig. 7. HPLC chromatogram of the GDGT distribution of a cold sample from Lake Superior with a BIT Index value of 0.44 (shown in the upper panel) and a warm sample from Lake Turkana with a low BIT Index value of 0.14 (shown in the bottom panel). Applying the Powers et al. (2010) lacustrine TEX_{86} calibration, the Lake Superior sample yields a temperature of $8.7\text{ }^{\circ}\text{C}$ and the Lake Turkana a temperature of $22.5\text{ }^{\circ}\text{C}$.

Table 4

Marine and lacustrine TEX₈₆ calibrations. The formulas for calculating TEX₈₆, TEX₈₆^H and TEX₈₆^L values are provided in Table 2. See Fig. 8 for details on the combined calibration.

Calibration equation	n	r ²	Range (°C)	Error (°C)	Geographical location	Reference
TEX ₈₆ = 0.15T + 28	44	0.92	0–30	±2.0	Global marine surface sediments	Schouten et al. (2002)
T = -10.78 + 56.2 × TEX ₈₆	223	0.94	8–30	±1.7	Global marine surface sediments	Kim et al. (2008)
T = 67.5 × TEX ₈₆ ^L + 46.9	396	0.86	-3–30	±4.0	Global marine surface sediments	Kim et al. (2010b)
T = 68.4 × TEX ₈₆ ^H + 38.6	225	0.87	5–30	±2.5	Global marine surface sediments	Kim et al. (2010b)
T = 55.231 × TEX ₈₆ - 13.955	12	0.86	5–30	±3.62	Global lake surface sediments; BIT < 0.5	Powers et al. (2010)
T = TEX ₈₆ × 38.874 - 3.4992	13	0.92	11–30	±2.1	Global lake surface sediments with T > 10 °C; BIT < 0.6	Tierney et al. (2010a)
T = 54.889 × TEX ₈₆ - 13.363	19	0.89	5–30	±3.1	Global lake surface sediments	Combined calibration

can be determined with a reproducibility of ±0.004, equivalent to ±0.3 °C (Schouten et al., 2007a).

3.3.2. Factors potentially influencing TEX₈₆ temperature reconstructions

Although the current lakes calibrations for TEX₈₆ are based on a relatively small number of samples, surface sediments of a considerably larger number of lakes have been analyzed for TEX₈₆. For example, Powers et al. (2010) examined 46 lakes while Blaga et al. (2009) examined 47 lakes for GDGTs. However, several factors complicate application of the TEX₈₆ in many lakes. Both Powers et al. (2010) and Blaga et al. (2009) note that not all lakes contain the full suite of GDGTs needed to calculate a TEX₈₆ value. Second, some of the isoprenoid GDGTs used to calculate TEX₈₆ are also found in small amounts in soils, which are produced not only by soil Thaumarchaeota but also likely by other Archaea (Weijers et al., 2006b), and can be transported to oceans or lakes. The presence of soil-derived isoprenoid GDGTs may lead to either a warm or a cold bias in TEX₈₆ temperatures, and the direction of the bias is not predictable. In sediments with high BIT Index values, it is likely that crenarchaeol and the other isoprenoid GDGTs are partially derived from soils (Weijers et al., 2006b). Thus, it has been suggested for marine environments that TEX₈₆ only be applied where the BIT Index is ~ <0.3 (Weijers et al., 2006b). However, due

to potential *in-situ* production of branched GDGTs (Section 3.1), the BIT index may not accurately reflect soil organic matter input to lakes. It is also worth noting that in some cases, samples with relatively high BIT Index values can produce reliable TEX₈₆ estimates since TEX₈₆ temperatures have been found to agree closely with other temperature proxies such as the U₃₇^k Index (Castañeda et al., 2010). The degree of bias is dependent on the 'TEX₈₆' value of GDGTs brought in from soils to the lakes: if this value is similar to that of the TEX₈₆ produced in the lake then high input of soil organic matter (and thus high BIT values) will not have a large impact. The most appropriate testing may be to check for coinciding changes of BIT with TEX₈₆; in these cases care should be taken in interpreting TEX₈₆ temperature estimates.

A third factor that can influence TEX₈₆ values is additional contribution of isoprenoid GDGTs from groups other than Thaumarchaeota, such as methanogenic Euryarchaeota (Blaga et al., 2009). GDGT-0, a common Archaeal membrane lipid, is found in thermophilic Crenarchaeota and Euryarchaeota, methanogenic Euryarchaeota (Koga et al., 1993) and Euryarchaeota mediating the anaerobic oxidation of methane (Pancost et al., 2001; Blumenberg et al., 2004) in addition to Thaumarchaeota (Sinninghe Damsté et al., 2002a). In lakes, the most likely sources of GDGT-0 are from methanogenic Euryarchaeota and Thaumarchaeota (Blaga et al., 2009). It has been proposed that the ratio of GDGT-0 to crenarchaeol can be used to assess contributions from methanogenic Euryarchaeota because GDGT-0 and crenarchaeol are produced both by Thaumarchaeota and methanogenic Euryarchaeota whereas methanogenic Euryarchaeota produce GDGT-0 but lack crenarchaeol (Blaga et al., 2009). If the GDGT-0/crenarchaeol ratio is >2, a substantial methanogenic origin for GDGT-0 is likely (Blaga et al., 2009). Anoxic conditions are required for methanogenesis and thus the process can occur in sediments as well as in anoxic water columns. Indeed, many lake sediment and water column samples have been observed with GDGT-0/crenarchaeol values >2, which is perhaps not surprising given that methanogenesis is the dominant anaerobic mineralization process in lake sediments. Interestingly, Blaga et al. (2009) note that some samples from aerobic part of the water column were also characterized by GDGT-0/crenarchaeol ratios >2, suggesting an additional, possibly allochthonous, source of GDGT-0 in some settings.

A fourth factor that can influence TEX₈₆ values is the seasonal timing of the Thaumarchaeota bloom. In marine systems it has been noted that marine Thaumarchaeota are generally abundant in surface waters at times of the year when the majority of phytoplankton are not blooming (e.g. Murray et al., 1999; Wuchter et al., 2006a) and thus, TEX₈₆ temperatures may reflect this period in the seasonal cycle. Because most of the Thaumarchaeota seem to be nitrifiers (Francis et al., 2005; Könneke et al., 2005; Wuchter et al., 2006a) their growth rates are likely related to ammonium availability, derived from the decay of organic matter from primary producers. Indeed, it has been noted at several marine sites that TEX₈₆ reflects either summer or winter SST (Herfort et al., 2006; Menzel et al., 2006; Wuchter et al., 2006a; Castañeda et al., 2010;

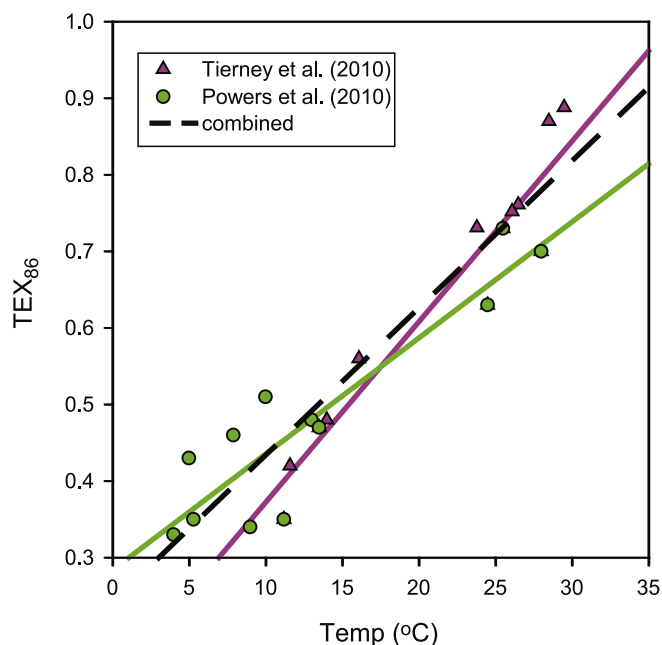


Fig. 8. Lacustrine TEX₈₆ calibrations. Data points used for the calibrations of Powers et al. (2010) and Tierney et al. (2010a) are shown by the green circles and pink triangles, respectively, while the solid lines of the same colors represent the linear fit. The black line represents the linear fit of all data points used in both the calibrations of Powers et al. (2010) and Tierney et al., (2010a). See Table 4 for equations and correlation coefficients of these calibrations.

Leider et al., 2010; Richey et al., 2011), which is likely the cause of some of the scatter in the TEX₈₆ core top calibration (Kim et al., 2010b). Similar seasonal influences have also been noted in lakes. At Lake Lucerne (Switzerland) the highest concentrations of isoprenoid GDGTs in suspended particulate material from surface waters (2 m) is observed in January–February (Blaga, 2010). Maximum fluxes of isoprenoid GDGTs occurs during the winter at Lake Challa (Sinninghe Damsté et al., 2009). Similarly, at Lake Superior fluxes of isoprenoid GDGTs towards the sediment mainly occurs in winter and late spring/early summer and particle transport initiates during overturning conditions (Woltering et al., 2010). Powers et al. (2010) also note a statistically significant difference between TEX₈₆ and winter surface temperature ($r^2 = 0.84$) in comparison to summer surface temperature ($r^2 = 0.71$). While presently limited to a few studies, it appears seasonal variability in Thaumarchaeota production likely exerts an influence on lacustrine TEX₈₆ values.

Sub-surface production by Thaumarchaeota is a fifth factor that can influence TEX₈₆ values. Thaumarchaeota are likely chemotrophs and therefore are not required to reside within the photic zone (Karner et al., 2001; Sinninghe Damsté et al., 2002a). In fact, Thaumarchaeota are distributed throughout the marine (Murray et al., 1999; Karner et al., 2001; Herndl et al., 2005) and lacustrine (Woltering et al., 2010) water columns yet studies of TEX₈₆ in sinking particle fractions in marine systems indicate that GDGTs mostly derive from the upper water column and that Thaumarchaeota living in deeper waters do not contribute substantially to the GDGT flux and thus the TEX₈₆ signal (Wuchter et al., 2005, 2006b). In some locations, Thaumarchaeota appear to reside in the upper water column but at greater depths than the sea surface. Indeed, TEX₈₆ has been reported to reflect sub-surface temperatures in several locations including the Benguela upwelling system (>40 m depth) (Lee et al., 2008), the Santa Barbara Basin (100–150 m depth) (Huguet et al., 2007), offshore NW Africa (~30 m depth) (Lopes dos Santos et al., 2010) and possibly in the Pigmy Basin, Gulf of Mexico (Richey et al., 2011). Sub-surface production has also been noted in lakes. Woltering et al. (2010) examined Archaeal 16S rRNA genes and GDGTs in sediment trap and filtered samples from the water column of Lake Superior over three annual cycles. They found that both GDGTs and Archaeal 16S rRNA genes are concentrated around the hypolimnion when Lake Superior is thermally stratified but are distributed throughout the water column during overturning periods. Woltering et al. (2010) also note that TEX₈₆ derived temperatures of the settling particles appear to represent the water temperature of the hypolimnion. Similarly, in Lake Lucerne (Switzerland) maximum production of isoprenoid GDGTs occurs within the aphotic zone, at ~50 m depth (Blaga, 2010). Interestingly, in a Lake Lucerne sediment core, an offset to higher temperatures is noted prior to ~1960 AD and is hypothesized to be related to the recent eutrophication of the lake, which likely resulted in migration of the Thaumarchaeota to a deeper ecological niche (Blaga, 2010).

Another factor which may need particular consideration is pH, which can be highly variable in lakes. In marine environments, the potential effects of pH on GDGT distributions are likely minor due to relatively small pH variations, even on geologic time scales. However, lakes display a wide range of pH values. For example, the volcanic crater lake Kawah Ijen (Indonesia) is extremely acidic with a pH < 0.3 (Lohr et al., 2005) while alkaline Lake Magadi (Kenya), which is recharged mainly by saline hot springs, has a pH of 10 (Jones et al., 1977). Even within individual lakes, some are known to have experienced large pH variations in the past. Pearson A. et al. (2008) found that GDGT distributions in hot springs displayed a good correlation with pH. Although hot springs contain a wide variety of (thermophilic) archaea with different GDGT distributions,

it is possible that pH may influence TEX₈₆ values. Culture experiments with thermophilic Euryarchaeota also show a decrease in cyclopentane moieties with decreasing pH (Shimada et al., 2008). Therefore, pH variations may be a potential concern for application of TEX₈₆ to lacustrine environments and there is a need for future studies to further investigate the potential effects of pH on GDGT distributions.

The impact of diagenesis on TEX₈₆ is not well-constrained although the few studies that have investigated the effects of oxic degradation suggest no substantial effect. Schouten et al. (2004) examined Arabian Sea sediments that were deposited in contrasting redox conditions but did not find a significant effect on TEX₈₆ values. Similarly, Kim et al. (2009) conducted a water column degradation experiment over a one year and did not find a significant effect of oxic degradation on TEX₈₆. Huguet et al. (2006a) also found no significant effect of herbivory on TEX₈₆ values. Thus, at this point it appears that diagenesis does not have a significant effect on TEX₈₆; however, differences in BIT Index (Section 3.1) values have been noted. Huguet et al. (2008) examined BIT Index values across the oxidation fronts of deep-sea turbidites (~127 Ka to 14 Ma) deposited at the Madeira Abyssal Plain. They note increasing BIT values on long-term exposure to oxygen and attribute this pattern to the enhanced preservation of soil organic matter due to sorptive protection, rather than to differences in degradation kinetics for the branched and isoprenoid GDGTs. We note that the impact of diagenesis on the MBT/CBT proxies (Section 3.2) have not yet been investigated and additional studies are needed to further examine the impacts of diagenesis on both branched and isoprenoid GDGTs and their associated proxies.

Finally, different populations of Thaumarchaeota may reside in different lakes, requiring local calibrations for TEX₈₆. In the high salinity and temperature environment of the Red Sea, Trommer et al. (2009) found that TEX₈₆ in surface sediments from the northern basin has a very different relationship between TEX₈₆ and temperature when compared to the global marine calibration. They speculated that this is due to a different population of Thaumarchaeota residing in this region. Indeed, Ionescu et al. (2009) found that Thaumarchaeota in the Gulf of Aqaba, northern Red Sea, were genetically distinct from those of the open ocean. It is possible that regional calibrations are necessary for increased accuracy of temperature predictions using TEX₈₆ in lakes, similar to the U₃₇^k index.

3.3.3. Lacustrine TEX₈₆ temperature reconstructions and outlook

Despite the many uncertainties with applying TEX₈₆ in lakes, the proxy has been applied to several lakes and importantly, has provided a method to examine continental paleotemperature history. TEX₈₆ has proven particularly successful in the large lakes of the East African Rift Valley, where isoprenoid GDGTs are particularly abundant, and to date, LST records have been generated from Lake Malawi (Powers et al., 2005, 2011; Woltering et al., 2011), Lake Tanganyika (Tierney et al., 2008, 2010a) and Lake Turkana and Lake Victoria (Berke et al., 2009, 2010). The temperature records of these lakes exhibit several common features indicating a regional, rather than a local, response to climate variability. These include a Last Glacial Maximum (LGM) present-day temperature difference of 3.5–4 °C, early warming following the LGM (Powers et al., 2005; Tierney et al., 2008; Woltering et al., 2011), a mid-Holocene warming centred at ~5 kyr (Powers et al., 2005; Tierney et al., 2008; Berke et al., 2009) and a recent warming on the order of ~2.5 °C occurring within the past 100 years (Tierney et al., 2010a; Powers et al., 2011) (Fig. 9). Thus, TEX₈₆ provides a useful tool for reconstructing past LST and for examining continental paleoclimate; however, its applicability to lakes needs to be assessed using the constraints outlined above.

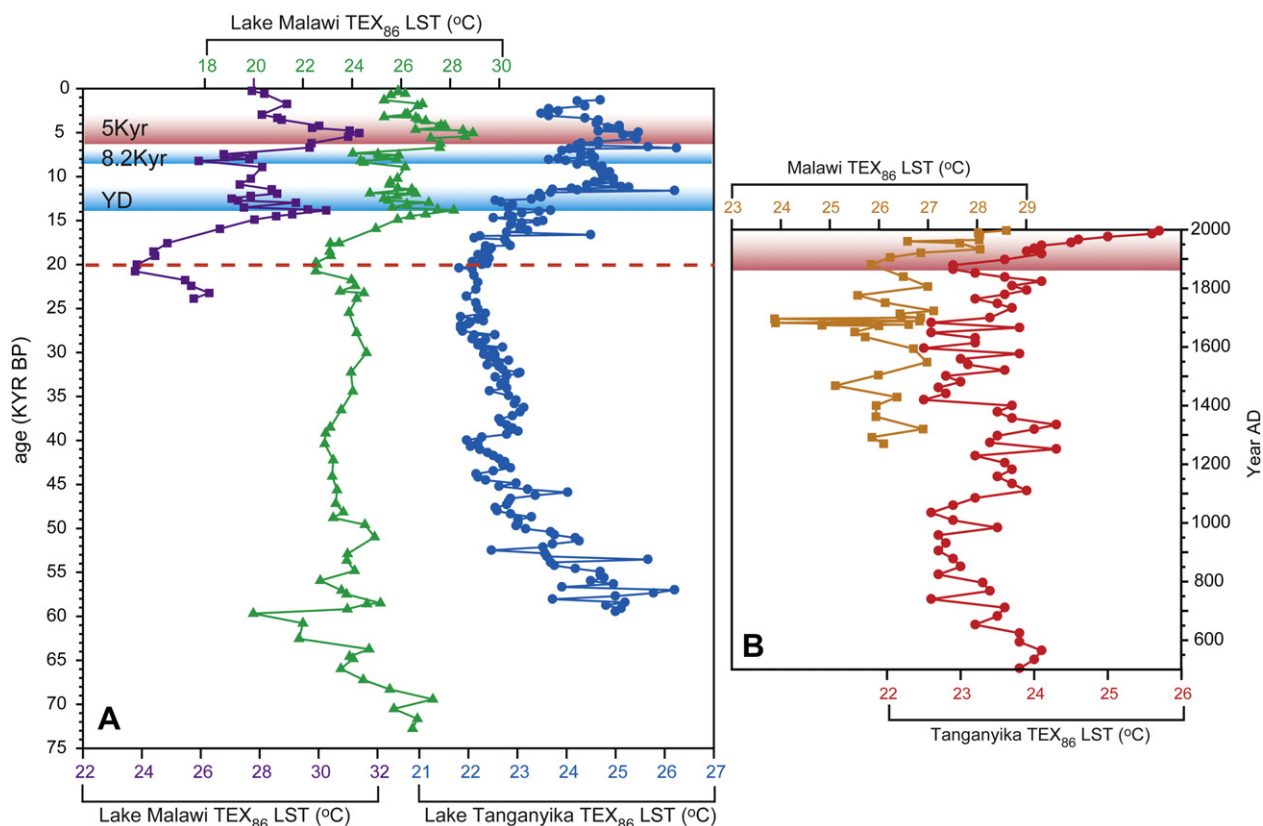


Fig. 9. TEX₈₆ lake surface temperature (LST) reconstructions from East Africa. In panel A, TEX₈₆ data from Lake Malawi core M98-1P (Powers et al., 2005) is shown with the TEX₈₆ data from Lake Malawi core MAL05-2A (Wolterring et al., 2011) and the Lake Tanganyika TEX₈₆ record (Tierney et al., 2008). A continuous record from Lake Tanganyika was generated from cores NP04-KH04-3A-1K and NP04-KH04-4A-1K (Tierney et al., 2008). All TEX₈₆ records indicate early warming initiating at ~20,000 years (indicated by the dashed red line) followed by a period of generally increasing temperatures until ~14,000 years. Variable and cooler conditions are noted until ~12,000 years, spanning the interval of the Younger Dryas (YD) (blue shading). A minor cooling is noted in all records around 8000 years, and may be related to the 8.2 Kyr cold event. An interesting feature of these records is a mid-Holocene warm period, centred at approximately 5000 years (red shading). Panel B shows two recent TEX₈₆ records: the record of cores M98-11MC and M98-2PG from Lake Malawi (Powers et al., 2011) and Lake Tanganyika cores KH1 and MC1 from Lake Tanganyika (Tierney et al., 2010a). Both of these records indicate a substantial >2.5 °C warming since ~1900 AD (red shading).

4. Reconstructing C₃ vs. C₄ vegetation and primary productivity using compound-specific isotopes ($\delta^{13}\text{C}$)

The *n*-alkanes, *n*-alkanols and *n*-alkanoic acids are straight-chain hydrocarbons produced by many organisms and the dominant chain lengths, carbon number distributions and isotopic compositions vary depending on the source organism. In many organisms the *n*-alkanes exhibit strong odd over even carbon number predominance whereas the *n*-alkanols and *n*-alkanoic acids exhibit strong even over odd carbon number predominance. The long-chain (C₂₇–C₃₅) *n*-alkanes (and the C₂₆–C₃₆ *n*-alkanols and *n*-alkanoic acids) are a main component of the epicuticular waxes of higher plants (Eglinton and Hamilton, 1967). In contrast, aquatic algae are dominated by shorter-chain homologues (C₁₇–C₂₁ *n*-alkanes) (Giger et al., 1980; Cranwell et al., 1987) while the mid-chain homologues (C₂₃–C₂₅ *n*-alkanes) are a dominant component of submerged aquatic macrophytes (Ficken et al., 2000). The *n*-alkanes (*n*-alkanols and *n*-alkanoic acids) are generally well preserved in sediments and plant leaf waxes are transported to marine or lacustrine sediments via water or wind erosion. These compounds are identified and quantified by GC and GC/MS, while their isotopic composition is determined by GC-IRMS. The carbon and deuterium isotopic composition of both plant leaf waxes and algal lipids can provide important paleo-environmental information, which is discussed in detail below and in Section 5.

Ratios based on *n*-alkanes of differing chain lengths have been used to examine relative inputs of aquatic macrophytes or terrestrial organic matter into lake sediments. For example, the *P*_{aq} ratio is the abundance of mid-chain *n*-alkanes over the total sum of the abundances of the mid plus long-chain *n*-alkanes and is used to discriminate sources of submersed and emersed vegetation (Ficken et al., 2000). The ratio of the long chain (C₂₇ + C₂₉ + C₃₁) to short-chain (C₁₅ + C₁₇ + C₁₉) *n*-alkanes can be used to assess terrestrial vs. aquatic sources of organic matter (Meyers, 1997). In addition to being able to differentiate between source organisms, the distribution patterns of *n*-alkanes can sometimes be used to distinguish between different vegetation types (e.g. Schwark et al., 2002; Hanisch et al., 2003). The C₃₁ *n*-alkane tends to be dominant in grasses while the C₂₇ and C₂₉ *n*-alkanes are dominant in deciduous trees (Cranwell, 1973). Additionally, the carbon preference index (CPI) (Bray and Evans, 1961) of *n*-alkanes is used to examine the odd over even carbon number predominance, which can be used to distinguish terrestrial plant from bacterial or petroleum sources, while the average chain length (ACL) of *n*-alkanes has been observed to increase with increasing aridity or temperature at some, but not all, locations (e.g. Peltzer and Gagosian, 1989; Rommerskirchen et al., 2003).

Although longer-chain lengths are generally produced by higher plants and the shorter-chain lengths by aquatic algae, some notable exceptions have been observed. For example, it was found that *Betula* (birch) leaves contain significant amounts of the C₂₃ *n*-

alkane (Sachse et al., 2006) while long-chain C_{27} and C_{29} n -alkanes have been reported from emersed aquatic macrophytes (Aichner et al., 2010b). Mid-chain n -alkanes have also been detected in *Sphagnum* species (Baas et al., 2000). Certain types of algae such as *Botryococcus braunii* may also be a source of long-chain n -alkanes in sediments (Lichtfouse et al., 1994) while some algae also produce unsaturated alkanes (e.g. C_{27} and C_{29} n -alkenes in *Rhizozolenoid* diatoms (Sinninghe Damsté et al., 1999a)). Thus, care should be taken when examining n -alkane distributions and when assigning sources to particular homologues.

4.1. Reconstructing C_3 vs. C_4 vegetation using plant leaf wax $\delta^{13}C$

The carbon isotopic composition ($\delta^{13}C$) of the long-chain n -alkanes (or other compounds derived from higher plants, such as lignin phenols) can be used to distinguish between vegetation utilizing different photosynthetic pathways. Plants utilizing the C_3 (Calvin–Benson) photosynthetic pathway tend to have average n -alkane carbon isotopic compositions of -34.7‰ for the C_{29} n -alkane (-35.2‰ for the C_{31} n -alkane), while plants utilizing the C_4 (Hatch–Slack) photosynthetic pathway are isotopically enriched with average n -alkane carbon isotopic compositions of around -21.4‰ for the C_{29} n -alkane (-21.7‰ for the C_{31} n -alkane) (Fig. 8) (Castañeda et al., 2009a and references therein). A third photosynthetic carbon-fixation pathway, the Crassulacean Acid Metabolism (CAM) pathway, has isotopic values intermediate between those of C_3 and C_4 plants (Deines, 1980). Thus, the $\delta^{13}C$ of plant leaf waxes preserved in sediments can be used to examine past vegetation type, which in turn can be related to climatic conditions such as temperature, aridity and atmospheric carbon dioxide (pCO_2) concentrations (Cerling et al., 1993; Collatz et al., 1998; Kuypers et al., 1999; Pagani et al., 1999; Huang et al., 2001). These endmember values for C_3 and C_4 plants can be used to create a binary mixing model for examining the percentage of C_4 plants present in past vegetation assemblages (e.g. Schefuss et al., 2003; Castañeda et al., 2007; Collins et al., 2011).

A recent study of bulk leaf carbon isotopes of 334 species of globally distributed C_3 trees and shrubs reports wide range of values from -34.9‰ to -21‰ (Diefendorf et al., 2010). Although n -alkane $\delta^{13}C$ was not examined, this study suggests that within the C_3 plants a wide range in n -alkane $\delta^{13}C$ values may also occur, and if so, assigning an endmember value for C_3 plants for use in a binary mixing model becomes difficult. Thus, the wide range in leaf $\delta^{13}C$ values noted in the relatively large dataset of Diefendorf et al. (2010) suggests that the error associated with converting n -alkane $\delta^{13}C$ values to % C_4 vegetation may be larger than the currently estimated error of $\pm 20\%$ (Castañeda et al., 2009a), which reflects uncertainty in the endmember values reported from approximately 50 samples each for C_3 and C_4 plants. However, additional surveys of n -alkane $\delta^{13}C$ values in modern vegetation are needed in order to constrain this uncertainty. Furthermore, it may be the case that a smaller range in endmember $\delta^{13}C$ values exists for particular ecosystems or biomes but this would require regional surveys of local biomes. Nevertheless, these uncertainties do not affect interpretation of the overall trends where relatively enriched (depleted) n -alkane $\delta^{13}C$ values indicate increased (decreased) inputs from C_4 plants (Fig. 10).

A number of lacustrine studies have utilized compound-specific $\delta^{13}C$ to examine past vegetation change and these studies have provided important insights into past environmental conditions (StreetPerrott et al., 1997; Ficken et al., 1998; Brincat et al., 2000; Huang et al., 2001; Boom et al., 2002; Castañeda et al., 2007, 2009a; Feakins et al., 2007; Tierney et al., 2010c). For example, Huang et al. (2006) examined the $\delta^{13}C$ of long-chain n -alkanes and

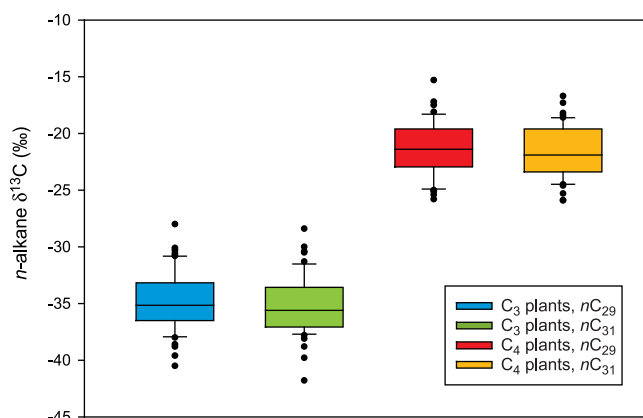


Fig. 10. Carbon isotopic composition ($\delta^{13}C$) of C_{29} and C_{31} n -alkanes in C_3 and C_4 plants. The box and whisker plots show the 25th (bottom of the box) and 75th (top of the box) percentile (the lower and upper quartiles) while the line in the middle of the boxes represents the median (the 50th percentile). The whiskers (error bars) above and below the box indicate the 90th and 10th percentiles, respectively. The black dots represent outliers. Each of these boxes represents approximately 50 samples of modern day vegetation and the data was compiled from values reported from the literature (see supplementary dataset of (Castañeda et al., 2009a)).

n -acids in a $\sim 62,000$ year sedimentary record from Lake Tulane (Florida) and compared the results to existing pollen data for the same lake. Instead of using a binary mixing model to reconstruct % C_4 vegetation, they created a three endmember mixing model because Lake Tulane receives inputs from pines and oaks. Although both pine and oak are C_3 plants, leaf waxes from pine (or other evergreens) are $4\text{--}5\text{‰}$ enriched compared to oak (or other angiosperm) leaf waxes (Huang et al., 2006 and references therein). Therefore, Huang et al. (2006) used the fraction of pine pollen as an input parameter and estimated the input of C_3 and C_4 plants while taking into consideration the differing $\delta^{13}C$ values of evergreen and deciduous trees. They found that the stable carbon isotope values of the long-chain n -alkanes and n -acids were in good agreement with the pollen record showing the highest % C_4 input during the LGM (Fig. 11), which they attributed to more arid conditions and lower pCO_2 levels. Despite low pCO_2 levels during the deglacial period, an increase in pine pollen was noted and accompanied by a shift to lower % C_4 values, which resulted from increased precipitation. Small changes of $\sim 5\text{‰}$ are noted during the Holocene and are attributed to changes in the relative abundance of pine and oak, rather than changes in C_3 and C_4 vegetation, supported by the pollen data (Huang et al., 2006). This study demonstrates the power of combining both plant leaf wax $\delta^{13}C$ and pollen data; however, it also should be noted that in some cases these proxies may not yield similar results since n -alkane inputs to lake sediments have been mainly attributed to local (within the watershed) sources while pollen may reflect more regional conditions due to long-distance atmospheric transport (e.g. Zheng et al., 2009). Conversely, long-distance atmospheric transport of n -alkanes can also occur and thus sources of n -alkanes and pollen need to be examined for individual sites. Some types of plant do not produce pollen and this may also lead to differences between n -alkane and pollen records.

4.2. Examining primary productivity using the $\delta^{13}C$ of short-chain n -alkanes (n -acids, n -alcohols)

The carbon isotopic composition of bulk organic matter ($\delta^{13}C_{OM}$) is widely used as a proxy for aquatic productivity in lakes (Meyers, 2003; Eglinton and Eglinton, 2008), with enriched $\delta^{13}C_{OM}$ values generally indicating increased productivity (Hollander and

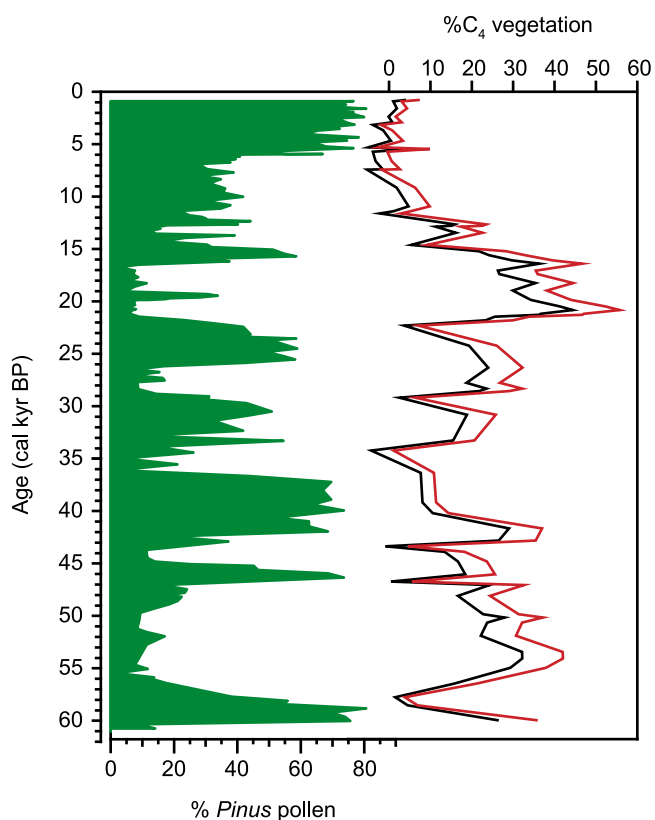


Fig. 11. Vegetation reconstruction from Lake Tulane, Florida. Percent *Pinus* pollen (C_3 plant) and the percent C_4 vegetation estimated from n -alkane $\delta^{13}C$ values. The percentage input from C_4 plants was calculated with two different isotopic end-member assumptions. The black line is the percentage C_4 input using endmember values of $\delta C_4 = -20\text{‰}$, and $\delta_p = -33\text{‰}$ and $\delta C_3 = -38\text{‰}$, where δC_4 , δ_p , and δC_3 are endmember $\delta^{13}C$ values for leaf waxes C_4 plants, pine, and C_3 plants (excluding pine, including oak and other C_3 trees and grasses), respectively. The red line represents endmember values of $\delta C_4 = -23\text{‰}$, $\delta_p = -33\text{‰}$, and $\delta C_3 = -39\text{‰}$. Data from Huang et al. (2006).

McKenzie, 1991; e.g. Eglinton and Eglinton, 2008). However, bulk organic matter reflects contributions from terrestrial, aquatic and bacterial sources and a variety of processes can influence $\delta^{13}C_{COM}$ values. For example, enriched $\delta^{13}C_{COM}$ values can be produced by bacterial processing of terrestrially derived C_4 organic matter (Hedges et al., 1997). When dissolved CO_2 is limited in lakes, many organisms utilize dissolved bicarbonate (HCO_3^-) as the carbonate source, leading to enriched isotopic compositions that can be heavier than those of land plants (Meyers, 2003). In anoxic systems, anaerobic respiration of organic matter produces ^{13}C -depleted methane and respired CO_2 (Woltemate et al., 1984; Whiticar et al., 1986), which can be incorporated into surface waters during upwelling events and produce phytoplankton that are ^{13}C -depleted during a time of enhanced productivity (Hollander and Smith, 2001). Other environmental factors including pH, temperature, salinity, growth rate, nitrogen limitation, eutrophication and physical diffusion barriers (such as ice cover) can also influence phytoplankton $\delta^{13}C$ values (e.g. Beardall et al., 1982; Takahashi et al., 1990; Schelske and Hodell, 1991, 1995; Fogel and Cifuentes, 1993; Hinga et al., 1994; Laws et al., 1995). In lakes that receive significant terrestrial inputs the $\delta^{13}C_{COM}$ record may mainly reflect vegetation changes in the watershed rather than changes in primary productivity. Examining the carbon isotopic composition of both terrestrial and aquatic biomarkers may help shed light on these processes.

Many lacustrine studies have examined the $\delta^{13}C$ of aquatic biomarkers, such as short-chain n -alkanes, to examine past changes in primary productivity or shifts in the dominant carbon source (e.g. Ostrom et al., 1998; Huang et al., 1999a; Filley et al., 2001; Castañeda et al., 2009b; Aichner et al., 2010a). In some cases, aquatic biomarker $\delta^{13}C$ records have been found to differ considerably from bulk $\delta^{13}C_{COM}$ records, which is not surprising given the various factors and processes that can influence bulk isotopic composition, as outlined above. For example, in Mud Lake (Florida) distinct $\delta^{13}C$ depth relationships are observed for the short-, mid- and long-chain n -alkanes, which differ from that of the bulk $\delta^{13}C_{COM}$ (Filley et al., 2001). In Lake Malawi it was found that $\delta^{13}C_{COM}$ mainly reflects changes in C_3 vs. C_4 vegetation in its watershed rather than changes in aquatic productivity (Castañeda et al., 2009b). However, compounds such as the short or mid-chain n -alkanes, while more source specific than bulk organic matter, likely still represent inputs from multiple source organisms and thus shifts in their carbon isotopic signatures can be quite difficult to interpret.

To date, only a few studies have examined the $\delta^{13}C$ of other, more source specific, algal lipids (e.g. sterols, long-chain n -alkyl diols or botryococcenes). Huang et al. (1999b) examined the $\delta^{13}C$ of multiple compounds (short-, mid- and long-chain n -alkanes, n -alcohols, n -acids and botryococcenes produced by the green alga *B. braunii*) from Sacred Lake, Kenya. They conclude that during the LGM, the relative contribution of biogenic CO_2 to carbon pool of Sacred Lake decreased, the ^{13}C content of the lake increased due to a shift to C_4 vegetation dominance in the catchment, and lower ambient atmospheric carbon dioxide concentrations were prevalent. The combination of these factors likely led to severe inorganic carbon limitation, thus producing algal biomass highly-enriched in ^{13}C . Huang et al. (1999b) also conclude that *B. braunii* must have used biocarbonate as its main carbon source. Although the authors were able to gain important insights into past carbon cycling at Sacred Lake, their study also illustrates the multitude of processes influencing $\delta^{13}C$ signatures, which can interfere with using $\delta^{13}C$ as a primary productivity proxy.

Aichner et al. (2010a) examined $\delta^{13}C$ values of total organic carbon ($\delta^{13}C_{TOC}$), total inorganic carbon ($\delta^{13}C_{TIC}$) and of a variety of biomarkers in a sediment core from Lake Koucha (Tibetan Plateau), a shallow lake densely populated by aquatic macrophytes that preserves carbonates. They found that the ^{13}C values of mid-chain n -alkanes derived from aquatic macrophytes tracked carbon limitation in the lake. They suggest that due to considerable variation in the isotopic composition of the carbon source, rather than examining shifts in the $\delta^{13}C$ of the C_{23} n -alkane alone, the offset between $\delta^{13}C$ values of the C_{23} n -alkane and TIC may provide a better indication of carbon limitation, and thus could provide an indicator of lake productivity (Aichner et al., 2010a). However, they caution that the main contributors to the sedimentary organic carbon pool must be known in order to interpret offsets between biomarker $\delta^{13}C$ and TIC (Aichner et al., 2010a).

Overall, deciphering lipid biomarker $\delta^{13}C$ records is very complex in many settings due to the numerous processes influencing $\delta^{13}C$ values (Huang et al., 1999b; Castañeda et al., 2009b; Kristen et al., 2010; Aichner et al., 2010a). For this reason, some studies of compound-specific $\delta^{13}C$ have led to inconclusive results (Russell et al., 2009; e.g. Castañeda et al., 2009b). Studies that have combined examining biomarker concentrations in conjunction with $\delta^{13}C$ records (of both aquatic and terrestrial sources) appear to be the most successful for assessing shifts in the dominant carbon source and primary productivity. However, in many cases, as outlined above, compound-specific $\delta^{13}C$ records cannot be used to specifically examine past changes in primary productivity.

5. Examining hydrological variability using compound-specific isotopes (δD)

5.1. Reconstructing hydrological variability using plant leaf wax δD

The deuterium isotopic composition (δD) of long-chain plant leaf waxes (e.g. *n*-alkanes or *n*-alkanoic acids) is being increasingly utilized as a tool for examining changes in aridity and precipitation δD values. The isotopic composition of meteoric water, the hydrogen source for terrestrial plants (Gat, 1996), is geographically influenced, reflecting environmental parameters including temperature, the source and amount of precipitation, elevation, and distance from the ocean, and exhibits an overall trend of decreasing δD values in precipitation with increasing latitude (Craig and Gordon, 1965; Bowen and Revenaugh, 2003). Indeed, at continental and global scales, studies have shown that precipitation δD values are an important factor controlling the δD of plant leaf waxes (δD_{LW} ; Fig. 12), reflecting the “latitude effect” (Bi et al., 2005; Sachse et al., 2006; Smith and Freeman, 2006; Hou et al., 2008; Liu and Yang, 2008; Polissar and Freeman, 2010). Shifts in δD_{LW} from a single site have been qualitatively interpreted as reflecting evapotranspiration (e.g. Schefuß et al., 2005) or a shift in precipitation δD (e.g. Liu and Yang, 2008) (e.g. a negative δD shift would indicate wetter conditions in both cases).

While the δD_{LW} is dependent on the δD of the source water, a number of other factors may influence δD_{LW} including an isotopic offset caused by fractionation during *n*-alkane biosynthesis, evapotranspiration from soil and leaf water, relative humidity, plant life form (e.g. tree, shrub, grass), and physiological differences such as differing photosynthetic pathways or water-use efficiencies (Craig and Gordon, 1965; Sessions et al., 1999; Smith and Freeman, 2006; Hou et al., 2008; Liu and Yang, 2008; McInerney et al., 2011). Furthermore, widely variable δD_{LW} values (up to 70‰) have been reported from the same site (Bi et al., 2005; Liu et al., 2006a; Hou et al., 2007c) and also in a single plant utilizing a water source with constant δD value where δD_{LW} values were found to vary throughout the different seasons (Sessions, 2006b). These factors complicate use of δD_{LW} as a paleohydrological proxy and must be considered before paleoclimatic interpretations can be made. Below we discuss these factors in more detail.

5.1.1. Influence of vegetation type on leaf wax δD

Quantitative reconstructions of the δD of meteoric water rely on the existence of a constant biosynthetic fractionation (“net fractionation” or “apparent fractionation”) between meteoric water and δD_{LW} values. However, differences in the net fractionation between meteoric water and δD_{LW} are known to exist between plant species (Chikaraishi and Naraoka, 2003; Smith and Freeman, 2006; Hou et al., 2007c; Feakins and Sessions, 2010a) and the available evidence suggests that a constant fractionation factor is unlikely (Chikaraishi and Naraoka, 2003; Liu and Yang, 2008). Net fractionations ranging from -73‰ to -242‰ for grasses (average -156‰) and from -57‰ to -220‰ for woody species (average -120‰) are reported based on a compilation of data from globally distributed sites (Liu and Yang, 2008 and references therein). Additionally, somewhat smaller net fractionations (-90‰) between meteoric water and δD_{LW} have been noted in semi-arid to arid environments (Feakins and Sessions, 2010a). Despite these complications, it is thought that within watersheds these differences are integrated and thus might allow for a constant offset to be used for paleoenvironmental reconstructions (Sachse et al., 2006; Hou et al., 2008; Feakins and Sessions, 2010a).

Differences in δD_{LW} have been noted between plants utilizing the C_3 vs. C_4 photosynthetic pathways in several studies, although some of the results are contradictory. While most studies have shown that *n*-alkanes from C_4 grasses are slightly enriched in δD in comparison to *n*-alkanes from C_3 grasses (in the range of $<35\text{‰}$) (Smith and Freeman, 2006; Liu et al., 2006a; Liu and Yang, 2008; McInerney et al., 2011), in *n*-alkanes from Japan and Thailand the opposite was observed with C_4 plants having slightly lower δD_{LW} values compared to C_3 plants (Chikaraishi and Naraoka, 2003) whereas no significant differences in δD_{LW} were observed between C_3 and C_4 plants collected from South China (Bi et al., 2005). Liu et al. (2006a) and Liu and Yang (2008) suggest that plant ecological life forms (e.g. trees, shrub, grass) exert a greater influence on δD_{LW} rather than differences in photosynthetic pathway. These authors argue that the various ecological life forms utilize different source waters, which leads to variations in δD_{LW} on the order of $60\text{--}70\text{‰}$, and find that grasses are more deuterium depleted compared to woody plants (trees and shrubs). Since trees and shrubs typically have longer and deeper roots they utilize water from deeper soil horizons (with potentially more enriched δD values) in comparison to grasses with short roots that use surface

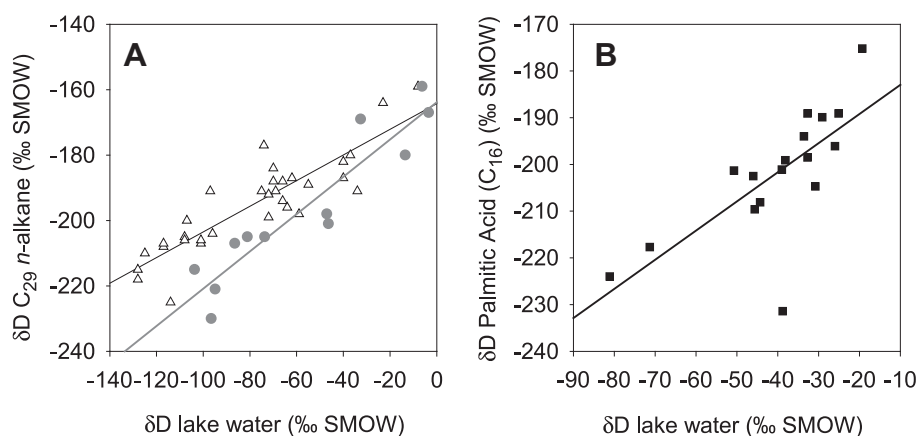


Fig. 12. Examples of relationships between δD of lipids and δD of lake water. A) The δD of the C_{29} *n*-alkane vs. δD of lake water from sites in South and North America (data from Polissar and Freeman, 2010) is indicated by the open triangles ($r^2 = 0.79$, $n = 31$). The gray circles indicate the δD of the C_{29} *n*-alkane vs. δD of lake water ($r^2 = 0.87$, $n = 12$) from a north–south transect of lakes in Europe (data from Sachse et al., 2004). Despite the fact that these lakes span a wide range of climatic conditions, elevations, and ecosystem types and structures, a significant relationship is noted between δD of lake water and *n*-alkane δD (combining the data for the American and European lakes yields a correlation with an $r^2 = 0.72$). B) The δD of palmitic acid (C_{16} *n*-acid) vs. the δD of lake water ($r^2 = 0.53$, $n = 17$) from an east–west precipitation gradient in South Dakota and Minnesota (USA) (data from Shuman et al., 2006).

soil water (potentially more depleted δD values). Succulent plants using the crassulacean acid metabolism (CAM) photosynthetic pathway, which have the ability to switch between using the CAM and C_3 pathways (Osmond et al., 1989), were found to display a wide variability ($\sim 90\%$) in δD_{lw} ranging from -193% to -107% even though environmental growth conditions were controlled (Feakins and Sessions, 2010b). Feakins and Sessions (2010b) suggest that coupled leaf wax $\delta^{13}C$ and δD data can be used to gain insights into the importance of CAM ecology in the past, in environments where it is possible to independently constrain the isotopic composition of meteoric waters. They note that with C_3 plants, leaf wax $\delta^{13}C$ and δD values are usually inversely correlated (Bi et al., 2005; Hou et al., 2007a) whereas they are positively correlated in CAM plants (Feakins and Sessions, 2010b).

5.1.2. Influence of transpiration on leaf wax δD

Relative humidity, via evaporative enrichment of soil and/or leaf water (transpiration), is another factor that can influence δD_{lw} values but its importance is currently debated. Hou et al. (2008) grew trees and grasses in humidity-controlled growth chambers and found that changing relative humidity from 80% to 40% produced only a $\sim 7\%$ increase in δD_{lw} relative to the source water. The authors concluded that most of the D-enrichment should occur through soil water evaporation. McInerney et al. (2011) conducted both growth chamber and field experiments of grasses, which produced contrasting results. In the growth chambers they did not find an effect of transpirational D-enrichment on δD_{lw} values; however the field samples provided clear evidence of D-enrichment, correlating with variations in relative humidity at the sites. Their observations, supported by a Craig–Gordon isotopic fractionation model, suggest that evaporation from soils and/or stems affects δD_{lw} but transpiration from leaves does not. McInerney et al. (2011) suggest that these observations can be explained by enrichment of the grass source water by evaporation from soils and/or stems for the field samples. In contrast, Feakins and Sessions (2010a) examined δD values of precipitation, groundwater, plant xylem water to examine their impact on δD_{lw} in an arid ecosystem. They found that leaf transpiration is the process responsible for most of the D-enrichment rather than soil water evaporation and observe that many species take up groundwater or precipitation without significant fractionation. Polissar and Freeman (2010) examined δD_{lw} in 28 watersheds spanning a range of precipitation δD , vegetation and climate types. They find that net fractionations between δD_{lw} and precipitation varies with ecosystem type and structure within the watershed and observe that δD_{lw} values from open grasslands are more sensitive to changes in aridity whereas those from closed, forested ecosystems are less sensitive due to reduced soil evaporation (Polissar and Freeman, 2010).

5.1.3. δD of leaf waxes: applications in paleoclimate reconstruction

Despite the apparent complications in interpreting the δD of plant waxes, they have been applied as a qualitative indicator for continental hydrology (e.g. Schefuß et al., 2005; Weijers et al., 2007a; Tierney et al., 2008, 2011). Tierney et al. (2008) examined δD_{lw} (the C_{28} *n*-alkanoic acid) and TEX_{86} (see Section 3.3) in a sediment core from Lake Tanganyika (East Africa), spanning the past $\sim 60,000$ years. They found that both temperature and hydrology in the Tanganyika basin were highly variable: δD_{lw} values spanning a range of $\sim 50\%$ and temperature changes of $\sim 4^\circ C$ were noted, indicating that Lake Tanganyika experienced abrupt changes in both temperature and hydrology (Fig. 13). Interestingly, temperature and precipitation were found to co-vary on orbital time scales whereas millennial-scale arid intervals noted in the δD_{lw} record were not accompanied by cooler lake surface temperatures. Their results suggest that precipitation at Lake

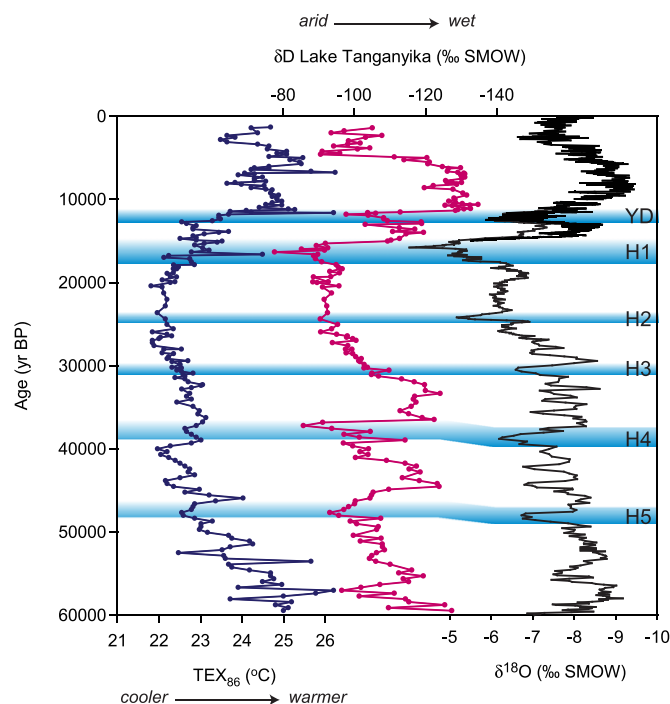


Fig. 13. Temperature and hydrologic history of Lake Tanganyika (East Africa) from TEX_{86} and leaf wax (C_{28} *n*-acid) δD analyses (Tierney et al., 2008). Oxygen isotope data from Hulu and Dongge Caves is shown for comparison (Wang et al., 2001; Dykoski et al., 2005). The blue shading indicates the arid intervals of the Younger Dryas (YD) and Heinrich Events H1–H5. Note that the TEX_{86} data of Tierney et al. (2008) was recalibrated using the newer lacustrine calibration of Tierney et al. (2010a).

Tanganyika is mainly influenced by Indian Ocean sea surface temperatures and the winter Indian monsoon rather than by past migrations of the position of the Intertropical Convergence Zone (Tierney et al., 2008).

5.1.4. Outlook: leaf wax δD as a hydrological proxy

Overall, it is clear that many outstanding questions exist with regard to the biological processes controlling δD_{lw} . Yet despite these complexities, δD_{lw} is emerging as a powerful proxy for examining continental hydrological variability (e.g. Schefuß et al., 2005; Weijers et al., 2007a; Tierney et al., 2008, 2011), at least qualitatively. Although more studies are certainly needed to better understand controls on δD_{lw} , it is apparent that evaluating the influence of vegetation change on δD in conjunction with δD_{lw} data is essential, a point that has been stressed by many authors (e.g. Liu et al., 2006a; Hou et al., 2007c; Liu and Yang, 2008; Polissar and Freeman, 2010; Feakins and Sessions, 2010b; Pu and Weiguo, 2011). Sites where vegetation change can be constrained by leaf wax $\delta^{13}C$ and/or pollen data or where large vegetation/ecosystem changes have not occurred in the past may be better suited for applying δD_{lw} in paleohydrological reconstructions.

5.2. Reconstructing hydrologic variability using the δD of algal lipids

Similar to the long-chain plant leaf waxes, δD measurements of algal lipids is also used as a tool for examining past hydrological fluctuations. Algal lipids are particularly well-suited for δD studies because all hydrogen in algae is derived from water and algae do not transpire (Zhang and Sachs, 2007), a process that can overprint the δD signal in land plants (see Section 5.1.2). The sensitivity of algal lipids δD as a recorder of source water δD values has been

demonstrated by both culture and surface sediment studies. Studies of algal cultures have examined a variety of lipids including alkenones, *n*-alkanes, *n*-alkanoic acids, and bytrococcones, and have demonstrated that the δD of algal lipids can closely track ($R^2 > 0.99$) water δD values (Englebrecht and Sachs, 2005; Zhang and Sachs, 2007). Studies of δD values of core top sediments also reveal a good correlation with the δD of the lake source water (Fig. 12; Sauer et al., 2001; Huang et al., 2004; Sachse et al., 2004). For these reasons, lipid biomarker δD is emerging as a potentially powerful paleoclimate proxy for lakes that do not preserve carbonates and where traditional paleolimnological proxies, such as oxygen isotope ratios of lacustrine carbonates, cannot be applied. In comparison to the oceans, lakes are much more sensitive to changes in precipitation/evaporation due to their smaller size and thus the δD of algal lipids can be a sensitive recorder of environmental conditions. While the δD of kerogen from bulk organic matter (e.g. Krishnamurthy et al., 1995; Tiljander et al., 2006; Lovan and Krishnamurthy, in press) or cellulose from aquatic plants (e.g. Edwards and Fritz, 1988; Edwards and Mcandrews, 1989; Wolfe et al., 2007) has been utilized for lacustrine paleoclimate studies, both can have significant allochthonous components possessing a different isotopic signature relative to the aquatic organic matter (Sauer et al., 2001). Therefore, δD measurements of bulk organic matter or cellulose should not be applied to sediments that receive large or varying inputs of allochthonous material. It also should be noted that in lakes where snowmelt is a major contributor to the lake water, the δD signal can be complicated by changing meltwater dynamics (e.g. Enders et al., 2008).

5.2.1. Influence of biosynthetic fractionations and other environmental factors on δD of algal lipids

Although culture and surface sediment studies have demonstrated that δD of various algal lipids exhibit good relationships with source water δD over large δD gradients, a key assumption to utilizing δD as a paleoclimatic proxy, like for plant leaf waxes, is a constant net apparent fractionation for each biomarker lipid. Thus, if the same lipid is analyzed for δD throughout a sediment core the biosynthetic fractionations will be similar and will not affect overall trends in δD . However, culture studies suggest this assumption is unlikely to be correct (Zhang and Sachs, 2007; Zhang et al., 2009). Zhang and Sachs (2007) examined hydrogen isotopic fractionation in five species of cultured freshwater green algae, including three strains of *B. braunii*, and demonstrated that algal lipid δD values reflect source water δD values but within a single species the hydrogen isotopic fractionation was found to vary systematically between lipids and lipid homologues. Importantly, they also note large differences (~ 90 – 100%) in the hydrogen isotopic fractionation of a single lipid between families of green algae and also relatively small (10 – 15%) differences between different species of green algae. They caution that if a non-source specific lipid is used to reconstruct water δD values, a shift in the dominant algal group could affect the lipid δD signature and be incorrectly interpreted as a shift in the δD of lake water (Zhang and Sachs, 2007). Given this, it is surprising that several studies have reported strong relationships between the δD of short-chain *n*-alkanes or *n*-alkanoic acids in surface sediments and source water δD (Huang et al., 2004; Sachse et al., 2004, 2006; Shuman et al., 2006; Hou et al., 2007b; Henderson et al., 2010) since these compounds are produced by multiple sources. However, Gao et al. (2011) examined D/H ratios in a wide range of plants in and around Blood Pond (MA) and created a mathematical model to quantitatively assess aquatic inputs of the mid-chain *n*-alkyl lipids to the lake sediments. They found that 97% of the total mid-chain *n*-alkyl lipids in Blood Pond derived from aquatic macrophytes. Thus, it appears that for some lakes a dominant source of the *n*-alkyl lipids

exists, facilitating the use of their δD as a paleoenvironmental proxy.

Other environmental factors such as temperature, salinity, growth rate, nutrient availability and light levels can also influence D/H fractionation. Zhang et al. (2009) investigated the effects of temperature and cultured two species of freshwater green algae at 15 and 25 °C and found increased D/H fractionation (2 – 4% per °C) for all lipids analyzed at the higher temperature. They also investigated the effects of nitrogen limitation and found that fatty acids from both nitrogen replete and nitrogen-limiting cultures had similar δD values whereas sterols in the nitrogen-limiting culture were enriched in δD by 37% compared to the nitrogen replete culture. These results suggest that growth rate likely does not influence D/H fractionation in acetogenic lipids and thus *n*-alkanes, fatty acids, alkenones, alkenadienes and other linear lipids may be good targets for δD analysis (Zhang et al., 2009).

A study of surface sediments from 28 saline and hypersaline lakes on Christmas Island (central Pacific Ocean), where cyanobacteria are the major source of the organic matter, indicates that salinity also can induce a large effect on D/H fractionation. Sachse and Sachs (2008) examined δD values of a variety of lipids and found that as salinity increased, the lipids became increasingly deuterium enriched. They note that lipid δD values span a range of 100% whereas the δD of the source waters exhibited relatively little variability, spanning a range of 12% . Thus, the net D/H fractionation between the lipids and the source water decreases with increasing salinity, by a factor of approximately 0.8% – 1.1% for every 1-unit increase in salinity (Sachse and Sachs, 2008). Schouten et al. (2006) reported a somewhat larger D/H fractionation of 4 – 5% for every 1 unit of salinity for long-chain alkenones produced by *E. huxleyi* and *G. oceanica*. These studies suggest that δD might be used to develop a paleosalinity proxy if it is possible to make additional constraints on the source water δD . Given the observed effects of temperature and salinity on lipid D/H fractionation, it is probably best to target lakes that have not undergone significant temperature or salinity variations in the past for δD reconstructions.

5.2.2. δD as a proxy for evapotranspiration or lake water balance

It has been suggested that the difference between long- and short-chain *n*-alkane δD may serve as a proxy for evapotranspiration or water balance in some lakes. Sachse et al. (2004) examined a north–south transect of small, groundwater fed lakes in Europe and noted that long-chain *n*-alkanes recorded the δD value of meteoric water but were enriched by $\sim 30\%$ compared to the C_{17} *n*-alkane, reflecting evapotranspiration processes in plant leaves. They suggested that the difference between δD values of aquatically and terrestrially derived *n*-alkanes could serve as an evapotranspiration proxy in small, closed lake systems that are fed by meteoric water (Sachse et al., 2004, 2006). However, an important assumption to this proxy is that the water source remains constant. Subsequent research by Mügler et al. (2008) examined both lacustrine and higher plant *n*-alkanes of sediments in semi-arid to arid lakes in Tibet (Lakes Nam Co and Co Jiana) and of a humid lake in Germany (Lake Holzmaar). At humid Lake Holzmaar they found that terrestrial *n*-alkanes (C_{29}) were isotopically enriched in δD by $\sim 30\%$ compared to aquatic *n*-alkanes (C_{23}) in agreement with the results of Sachse et al. (2004). In contrast, in the semi-arid to arid Tibetan lake sediments terrestrial *n*-alkanes were isotopically depleted in comparison to aquatic *n*-alkanes, with an enrichment of $\sim 60\%$ observed for the aquatic *n*-alkanes. Mügler et al. (2008) note that the lake water, precipitation and inflow water are in isotopic equilibrium at humid Lake Holzmaar whereas lake water at the Tibetan sites is enriched by 30 – 50% compared to the inflow water, reflecting evaporation. The authors suggest that a positive C_{29} – C_{23} δD difference can potentially indicate humid environments

whereas semi-arid and arid conditions are reflected by a negative C_{29} – C_{23} δD difference. They further suggest that the C_{29} – C_{23} δD difference can be used to estimate evaporation to inflow (E/I) ratios and to reconstruct the lakes past water balance.

Aichner et al. (2010b) further tested using the δD difference between terrestrial and aquatic *n*-alkanes as a proxy to reconstruct past lake water balance. They examined surface sediment samples from a number of sites on the Tibetan Plateau and a sediment core from Koucha Lake, in the NE Tibetan Plateau. In contrast to the results of Mügler et al. (2008), it was found that most of the surface sediment samples did not display a significant offset between δD values of aquatic and terrestrially derived *n*-alkanes. A similar result was found in the downcore record, where only minor offsets between δD values of aquatic and terrestrially derived *n*-alkanes were observed. They note that in arid environments evapotranspiration alters the hydrogen isotope signal of leaf and soil water, in addition to evaporation enriching lake water δD . Thus this paleo-aridity proxy may not be applicable to all settings since for it to work δD -enrichment of soil and leaf water must be outweighed by lake water evaporation (Aichner et al., 2010b).

5.2.3. Outlook: δD of algal lipids as hydrologic proxy

From the above discussion it is clear that more studies are needed to investigate the effects of biosynthetic fractionations and other environmental parameters including salinity, temperature, growth rate, nutrient availability and light limitation on lipid D/H fractionation in a wider variety of algal species and over a wider range of environmental conditions. Nevertheless, several studies have demonstrated that the δD of algal lipids can be used to examine past hydrologic variability from lake sediment, at least qualitatively, and have provided important paleoenvironmental information (Shuman et al., 2006; Hou et al., 2007b; Mügler et al., 2008; Henderson et al., 2010; Aichner et al., 2010b).

Considering the observation that large difference in hydrogen isotopic fractionation exist between single lipids produced by different algae (Zhang and Sachs, 2007), and the fact that many lipids have multiple source organisms, the choice of a lipid to analyze for δD reconstructions is an important one. Lipids that have been identified as being the most promising for δD reconstructions include C_{34} botryococcene, which produced solely by the B race of *B. braunii* (Zhang and Sachs, 2007), and lacustrine alkenones, which derive from haptophyte algae and provide a lake water signal (D'Andrea and Huang, 2005). Unfortunately, in many lakes these lipids are either not present or their abundances are too low to allow for δD analysis (Sachse et al., 2004; Castañeda et al., 2009b). Huang et al. (2004) suggested that the δD of palmitic acid, a C_{16} fatty acid that is abundant in many lake sediments, also captures the δD signals of lake water. From a north–south transect of lake surface sediments from eastern North America, Huang et al. (2002) demonstrated a strong correlation between lake water δD and the δD of palmitic acid. They also found that downcore δD values from Crooked Pond (MA) exhibited the same trends as pollen-inferred temperature data. Similarly, Hou et al. (2007c) examined δD of behenic acid (C_{22} *n*-acid) in a sediment core from Blood Pond (MA) and showed that the δD record displays a remarkable similarity with Greenland ice core records and also a pollen-inferred temperature record from Blood Pond as well as a carbonate $\delta^{18}O$ record from a nearby lake (Fig. 14). However, the results of Zhang and Sachs (2007) strongly argue against using palmitic acid, or any compound that derives from multiple sources, for δD reconstructions.

Clearly, there is a need to identify a suitable aquatic freshwater biomarker for δD reconstructions that is both widespread and abundant in lakes. We suggest that the long-chain (C_{28} – C_{32}) 1,15 *n*-alkyl diols (discussed in section 8.4), biomarkers for

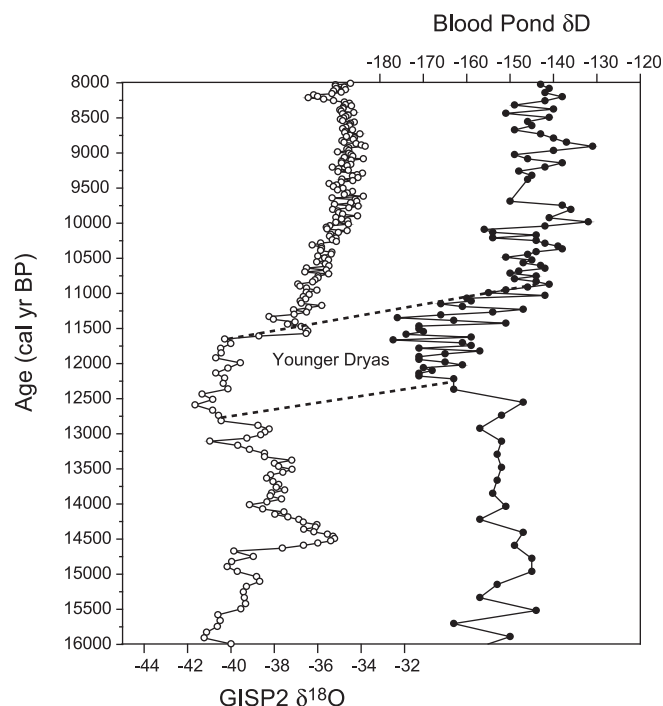


Fig. 14. Deuterium isotope (δD) reconstruction from Blood Pond, MA. Data from Hou et al. (2007b). The δD of behenic acid (C_{22} *n*-acid) is shown next to the oxygen isotope ($\delta^{18}O$) record from Greenland Ice Sheet Project 2 (GISP2) for comparison. The dashed lines indicate excursions noted during the interval of the Younger Dryas cold period in both records.

eustigmatophyte algae (Volkman et al., 1992; Versteegh et al., 1997), may offer possibilities for lacustrine δD reconstructions these compounds appear to be both very abundant and widespread in lake sediments (Xu et al., 2007; Xu and Jaffé, 2008; Castañeda et al., 2009c; Shimokawara et al., 2010). Another potential candidate would be the δD of ether-bound isoprenoid from archaeal membranes using chemical degradation and GC/pyrolysis/IRMS (Kaneko et al., 2011). This approach, although time consuming, allows for the determination of δD of biphytane moieties of GDGTs (Kaneko et al., 2011). Since crenarchaeol is very abundant in some lakes, it may provide a suitable biomarker for δD reconstructions in lakes that receive little terrestrial input (i.e. where BIT index values are low).

6. Investigating aquatic microbial community composition and UV radiation using sedimentary pigments

Pigments such as chlorophylls, carotenoids and their derivatives, preserved in sediments have long been utilized for paleolimnological studies (e.g. Fox, 1944; Vallentyne, 1954, 1956, 1957). Algae, microbes and higher plants synthesize a variety of pigments for use in photosynthesis and while some are broadly-distributed (e.g. chlorophyll *a*, β -carotene, pheophytin *a*), others provide a high degree of taxonomic specificity. Sedimentary pigments have been used to address a wide range of research questions including changes in algal and bacterial community composition, changes in trophic status, food–web interactions, lake acidification, and anthropogenic impacts on aquatic ecosystems (Table 5) (Sanger, 1988; Leavitt, 1993; Millie et al., 1993; Leavitt and Hodgson, 2001). In lakes, the main pigment sources are planktonic and benthic algal and microbial communities, aquatic macrophytes, and higher plant materials from the surrounding catchment. Chlorophyll *a* is the most common pigment and has been used to estimate

Table 5
Common sedimentary pigments and some of the environmental parameters they can be used to reconstruct. Table modified from Leavitt and Hodgson (2001). The main sources of each pigment are indicated: planktonic (P), littoral (L), terrestrial (T) or sedimentary (S; post-depositional derivatives). Derivatives of chlorophylls and carotenoids can provide important environmental information regarding water column and sedimentary processes that transform pigments including grazing, anoxia, stratification, and light intensity. Other proxies, such as the ratio of absorbance at 430 and 410 nm provides information on the extent of chlorophyll to pheopigments, and has been used as proxy for lake acidification (Guilizzoni et al., 1992).

Pigment	Source	Affinity	Potential for proxy reconstruction
Carotenoids			
β, β-carotene	P, L, T	Higher plants, algae, some phototrophic bacteria	Total algal abundance
β, α-carotene	P, L	Cryptophytes, chrysophytes, dinoflagellates, some chlorophytes	Total algal abundance
Alloxanthin	P	Cryptophytes	Cryptophyte productivity
Fucoxanthin	P, L	Diatoms, chrysophytes, some dinoflagellates,	Siliceous algae and dinoflagellate abundance
Diatoxanthin	P, L, S	Diatoms, dinoflagellates, chrysophytes	Algal abundance
Diadinoxanthin	P, L, S	Diatoms, dinoflagellates, chrysophytes, cryptophytes	Algal abundance
Dinoxanthin	P	Dinoflagellates	Dinoflagellate abundance
Peridinin	P	Dinoflagellates	Very labile compound, rarely recovered from sediments
Echinenone	P, L	Cyanobacteria	Cyanobacteria abundance
Zeaxanthin	P, L	Cyanobacteria	Cyanobacteria abundance
Canthaxanthin	P, L	Colonial cyanobacteria, herbivore tissues	Cyanobacteria abundance
Myxoxanthophyll	P, L	Colonial cyanobacteria	Cyanobacteria abundance
Oscillaxanthin	P, L	Cyanobacteria (Oscillatoriaceae)	Cyanobacteria abundance
Aphanizophyll	P, L	N ₂ -fixing cyanobacteria (Nostocales)	Cyanobacteria abundance
Lutein	P, L, T	Green algae, euglenophytes, higher plants	Algal abundance
Neoxanthin	L	Green algae, euglenophytes, higher plants	Algal abundance; very labile compound
Violaxanthin	L	Green algae, euglenophytes, higher plants	Algal abundance; very labile compound
Okenone	P	Purple sulphur bacteria	Photic zone anoxia, water column stratification
β-isorenieratene	P	Chlorobiaceae (green sulphur bacteria)	Photic zone anoxia, water column stratification
Isorenieratene	P	Chlorobiaceae (brown varieties)	Photic zone anoxia, water column stratification
Chlorobactene	P	Chlorobiaceae (green sulphur bacteria)	Photic zone anoxia, water column stratification
Astaxanthin	P, L	Invertebrates, N-limited Chlorophyta	Indicator of herbivory (grazing); labile compound
Scytonemin	P, L	Colonial cyanobacteria	Past UV radiation environments
Total carotenoids	P, L, T, S	All of the above	Total phosphorus concentrations in lake water
Mycosporine-like amino acids	P	Diatoms, cyanobacteria, haptophytes, dinoflagellates	Past UV radiation environments
Chlorophylls			
Chlorophyll <i>a</i>	P, L	Photosynthetic algae, higher plants	Total algal abundance
Chlorophyll <i>b</i>	P, L	Green algae, euglenophytes, higher plants	Chlorophyte abundance
Chlorophyll <i>c</i>	P, L	Dinoflagellates, diatoms, chrysophytes	Total algal abundance; very labile compound
Chlorophyll degradation products			
Chlorophyll <i>a'</i>	P, L, T, S	chl <i>a</i> derivative, forms as an artefact of extraction	Total algal abundance
Chlorophyllide <i>a</i>	P, L, T, S	chl <i>a</i> derivative	Algal abundance; product of senescence
Chlorophyllone <i>a</i>	P	chl <i>a</i> derivative	Indicator of zooplankton herbivory (grazing)
Chlorophyll <i>a</i> allomers	P, L, T, S	chl <i>a</i> derivative, produced by autoxidation reaction	Grazing, photooxidation
Pheophytin <i>a</i>	P, L, T, S	chl <i>a</i> derivative (general)	Total algal abundance
Pheophytin <i>b</i>	P, L, T, S	chl <i>b</i> derivative (general)	Chlorophyte abundance
Pheophorbide <i>a</i>	P, L, S	chl <i>a</i> derivative (grazing, senescent diatoms)	Indicator of zooplankton herbivory (grazing)
Pyro-pheo(pigments)	L, S	Derivatives of <i>a</i> and <i>b</i> -phorbins	
Deoxyphyloerythroetioporphyryn (DPEP)	S	Defunctionalized product of chl <i>a</i>	
Purpurin-18 phytol ester	P	chl <i>a</i> derivative	Oxidative transformations associated with senescence or particle sinking
Sterol chlorin esters (SCEs)	S	Sterol group originates from zooplankton or phytoplankton	Indicator of zooplankton herbivory (grazing); formed by biological reaction at the time of deposition
Bacteriochlorophylls			
Bacteriochlorophyll <i>a</i>	P	Purple photosynthetic bacteria	Photic zone anoxia, water column stratification
Bacteriochlorophyll <i>b</i>	P	Purple photosynthetic bacteria	Photic zone anoxia, water column stratification
Bacteriochlorophyll <i>c</i>	P	Green sulphur bacteria	Anoxygenic photosynthesis, photic zone anoxia
Bacteriochlorophyll <i>c</i> ₂	P	Green non-sulphur bacteria (Chloroflexi)	Oxic conditions
Bacteriochlorophyll <i>d</i>	P	Green sulphur bacteria	Anoxygenic photosynthesis, photic zone anoxia
Bacteriochlorophyll <i>e</i>	P	Green sulphur bacteria	Anoxygenic photosynthesis, photic zone anoxia
Bacteriochlorophyll <i>g</i>	T	Heliobacteria	Anoxic conditions
Bacteriochlorophyll degradation products			
Me Et maleimides	P	Phytoplanktonic chlorophyll	
Me n-Pr maleimides	P	Bacteriochlorophylls <i>c</i> , <i>d</i> , and <i>e</i>	Anoxygenic photosynthesis, photic zone anoxia
Me i-Bu maleimides	P	Bacteriochlorophylls <i>c</i> , <i>d</i> , and <i>e</i>	Anoxygenic photosynthesis, photic zone anoxia

phytoplankton biomass in the water column (e.g. Steele, 1962; Cullen, 1982). Aquatic organisms produce carotenoid compounds for photosynthesis and a number of different carotenoids (e.g. alloxanthin, lutein, echinenone, fucoxanthin, peridinin) can be used

to examine changes in specific algal classes (Leavitt and Hodgson, 2001 and references therein). Certain benthic algae produce unique pigments when exposed to ultraviolet (UV) radiation and these pigments can be used to examine past changes in UV

radiation in lakes (Leavitt et al., 1997, 1999). Specific pigments and bacteriochlorophylls derived from photosynthetic green and purple sulphur bacteria provide reliable biomarkers for water column anoxia since they require both light and sulphide. For example, purple sulphur bacteria produce bacteriochlorophyll *a* or bacteriochlorophyll *b* as well as several characteristic carotenoids including okenone, which is highly specific to this group (Sinninghe Damsté and Schouten, 2006 and references therein). Green sulphur bacteria produce bacteriochlorophyll *c*, *d* or *e* as well as the aromatic carotenoids chlorobactene and isorenieratene (Koopmans et al., 1996; Sinninghe Damsté and Schouten, 2006 and references therein). The presence of these compounds in ancient lake sediments has been used as an indicator of stratified conditions (e.g. Ariztegui et al., 2001; Hanisch et al., 2003; Itoh et al., 2003; Mallorquí et al., 2005).

Although it is well-known that most pigments are degraded within the water column and surface sediments (e.g. Furlong and Carpenter, 1988) where they undergo a series of transformation reactions, in many cases the diagenetic and catagenetic products retain enough specificity for them to be used as environmental indicators (Summons and Powell, 1987; Koopmans et al., 1996; Sinninghe Damsté et al., 2001a). Sedimentary pigments have thus been used to examine past phytoplankton dynamics and water column anoxia by countless studies, which are too numerous to review here. We note that sedimentary pigments have been previously reviewed (Sanger, 1988; Leavitt, 1993; Millie et al., 1993; Leavitt and Hodgson, 2001; Sinninghe Damsté and Schouten, 2006) and that they continue to be utilized for many paleolimnological investigations (e.g. Squier et al., 2002; Waters et al., 2005; Soma et al., 2007; Romero-Viana et al., 2009; Guilizzoni et al., in press; Hobbs et al., 2010; Aichner et al., 2010b). Here, we briefly highlight a couple of recent research developments.

6.1. Scytonemin and mycosporine-like amino acids as indicators of past UV receipt

Certain types of photoautotrophs produce ultraviolet (UV)-absorbing pigments to protect against cellular damage from UV radiation, and these compounds have the potential to serve as proxies for examining past solar irradiance (Leavitt et al., 1997; Sinha et al., 1998; Quesada et al., 1999; Squier et al., 2004; Hodgson et al., 2005). Scytonemin (Fig. 15) is a sheath pigment produced by certain types of cyanobacteria and variations in its abundance have been used to examine past UV variations because cyanobacteria increase extra-cellular concentrations of photoprotective compounds when exposed to elevated UV radiation (Hodgson et al., 2005). Although absolute intensities of UV radiation cannot be reconstructed from scytonemin concentrations, relative changes can be examined. For example, Hodgson et al. (2005) examined the ratio of total scytonemin to total sedimentary carotenoids in Lake Reid, East Antarctica, and concluded that during the last glacial benthic cyanobacteria were exposed to significantly higher levels of UV radiation (more than three times higher) than during the Holocene. Similarly, Verleyen et al. (2005) examined scytonemin in a sedimentary record spanning the past ~2000 years from a shallow lake, Pup Lagoon, in the Larsemann Hills of East Antarctica. The authors used the ratio of total scytonemin concentrations to the total cyanobacterial carotenoid concentration as a proxy for light intensity in the UV radiation wavelengths. This ratio excludes influences from changes in the abundances of cyanobacteria and other co-dominant algal groups (Verleyen et al., 2005). They found that several reconstructed periods of high UV radiation coincide with solar minima noted in historical records. Antarctic lakes are particularly well-suited for analysis of photoprotective compounds since dissolved organic matter (DOM), which absorbs UV radiation, is generally absent

(Hodgson et al., 2005). In other Arctic and high-latitude lakes, fluctuations in DOM inputs from vegetation and soils have been related to significant variations in UV radiation receipt (Leavitt et al., 2003).

Another group of UV-screening compounds, the mycosporine-like amino acids (MAAs), are known from over 380 species of marine organisms (including bacteria, algae, corals, and fish) and also in freshwater organisms including cyanobacteria, microalgae and zooplankton (Helbling et al., 2002; Tartarotti et al., 2004 and references therein). MAAs reported from marine phytoplankton occur predominately in members of Dinophyceae (dinoflagellates), Bacillariophyceae (diatoms), and Haptophyceae (Sinha et al., 1998). Interestingly, MAAs provide protection against UV radiation to both organisms that produce these compounds, as well as to their primary and secondary consumers (Sinha et al., 1998 and references therein; Helbling et al., 2002). Over 20 different types of MAAs have been described but typically a glycine subunit is found at the third carbon atom of the ring system (Sinha et al., 1998). It should be noted that while higher concentrations of MAAs are noted in organisms exposed to intense solar radiation (e.g. Tartarotti et al., 2004) it is thought that MAAs may also play other roles in cellular metabolism (Sinha et al., 1998 and references therein), which may influence MAA concentrations.

MAAs have been discovered embedded in the frustules of marine diatoms from the Southern Ocean in both plankton and sediment samples, which appears to protect them from chemical degradation (Ingalls et al., 2010). These MAAs are stable when associated with the diatom frustules and after being released by HF digestion they can be quantified by HPLC-MS techniques. Ingalls et al. (2010) suggest that it might be possible to develop an MAA-based indicator of past solar irradiance, assuming that there is a reliable relationship between frustule-bound MAA concentrations and irradiance. As diatoms are a major contributor to primary productivity in many lakes, frustule-bound MAAs may be useful for paleolimnological studies of past solar irradiance. Ingalls et al. (2010) also suggest that MAAs are well-suited for paleoenvironmental reconstructions based on compound-specific nitrogen ($\delta^{15}\text{N}$) and carbon ($\delta^{13}\text{C}$ and $\Delta^{14}\text{C}$) isotopes.

6.2. Sedimentary pigments as a proxy for past phosphorous concentration in lakes

Recently, it has been proposed that sedimentary pigments can be used to infer past phosphorous concentration in lakes. Guilizzoni et al. (2011) spectrophotometrically measured total carotenoids (TC) in sediment samples from 28 Italian lakes and found a significant positive relationship ($r = 0.78$; $p < 0.001$) between TC concentrations (relative to organic matter content) and total phosphorous (TP) concentrations of lake water measured at overturn. They developed a transfer function from this dataset and then used TC concentrations in sediment cores to estimate past TP concentrations. At Lake Maggiore, where two other diatom-based reconstructions of TP were available in addition to measured TP values during recent times, the pigment and diatom-based TP estimates were in good agreement (Fig. 16) and it was found that the TC method provided the best approximation of the measured TP values (Guilizzoni et al., 2011). The lakes they examined are characterized by a variety of climate conditions, morphologies, and nutrient conditions but despite this, it found that the pigment model correctly estimated TP in all lakes except for those rich in submerged aquatic macrophytes and epiphytes, which were outliers in the TP vs. TC relationship. Thus, the approach of using sedimentary pigments to examine past TP concentrations appears promising but requires additional testing in a wider variety of geographical locations.

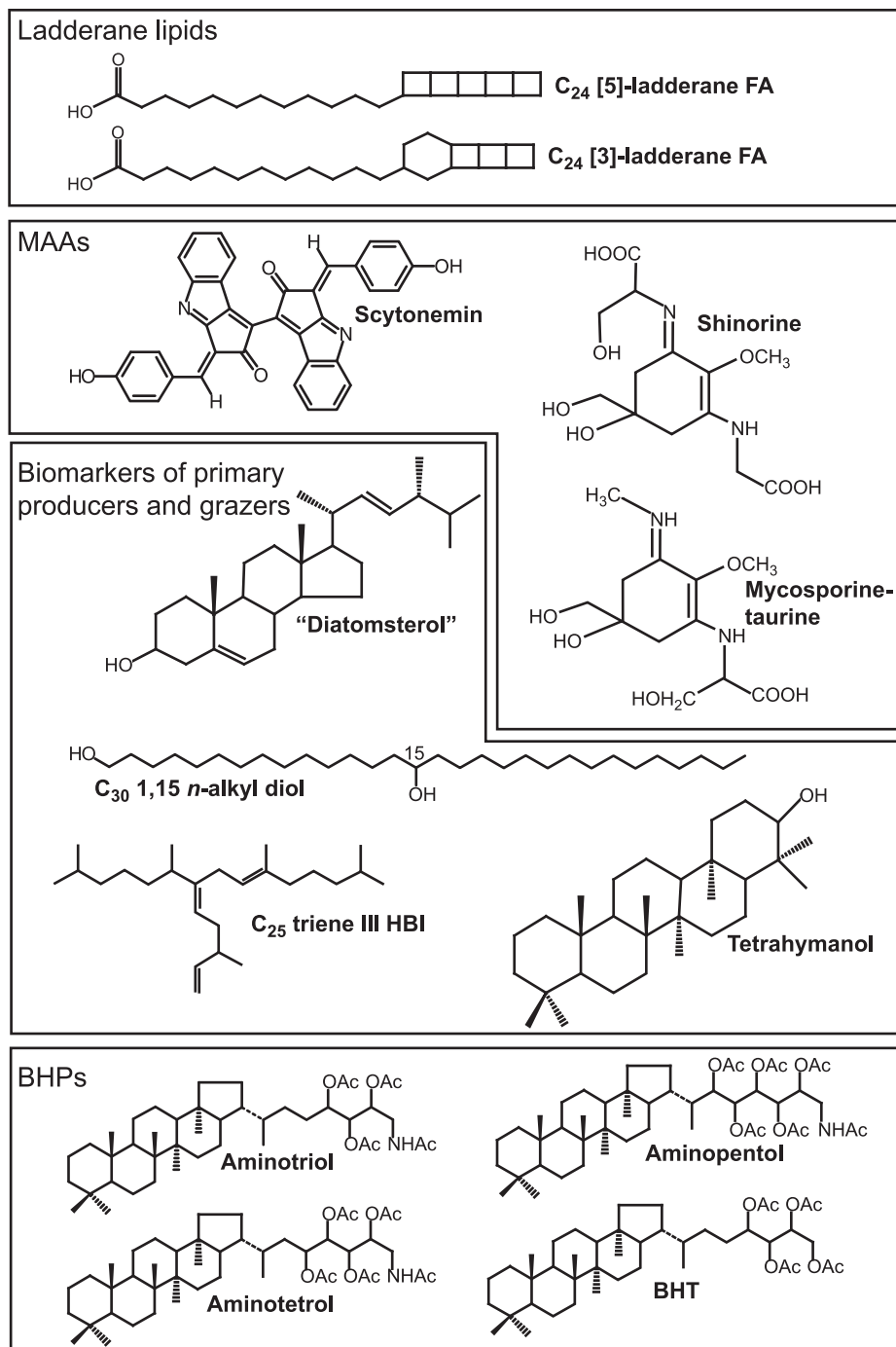


Fig. 15. Structures of selected biomarkers discussed in the text. Structures shown include the C₂₄ [5]-ladderane fatty acid (FA) and the C₂₄ [3]-ladderane FA; the compound scytonemin; examples of the mycosporine-like amino acids (MAAs) including scytonemin, shinorine and mycosporine-aurine; the C₃₀ 1,15 *n*-alkyl diol, BHPs including aminotriol, aminopentol, aminotetrol and bacteriohopane-32,33,34,35-tetrol (BHT); a highly branched isoprenoid (HBI) alkane, C₂₅ triene III; and, tetrahymanol.

7. Exploring soil, cyanobacteria and methanotrophic bacteria inputs and processes using bacteriohopanepolyols

Intact bacteriohopanepolyols (BHPs) are pentacyclic triterpenoids (Fig. 15) that comprise a diverse group of membrane lipids produced by many (but not all) bacteria including α -, β -, and γ -proteobacteria, cyanobacteria and planctomycetes (Talbot and Farrimond, 2007; e.g. Cooke et al., 2009; Welander et al., 2010). BHPs are also the major precursors of sedimentary geohopanooids (e.g. hopanes, hopenes, hopanoic acids, hopanols) (e.g. Ourisson

and Albrecht, 1992). A wide variety of BHP structures are known, which vary in the number, position and nature of the functional groups on the side chain (Talbot and Farrimond, 2007). The most common BHP in soils and sediments is bacteriohopane-32,33,34,35-tetrol (BHT), which is known to be produced by a diverse suite of organisms (Talbot and Farrimond, 2007) and has been reported in sediments up to 50 Ma (van Dongen et al., 2006). While some BHPs are produced by numerous organisms, others are more specific and can be related to groups of organisms or environments and thus provide biomarkers for examining bacterial

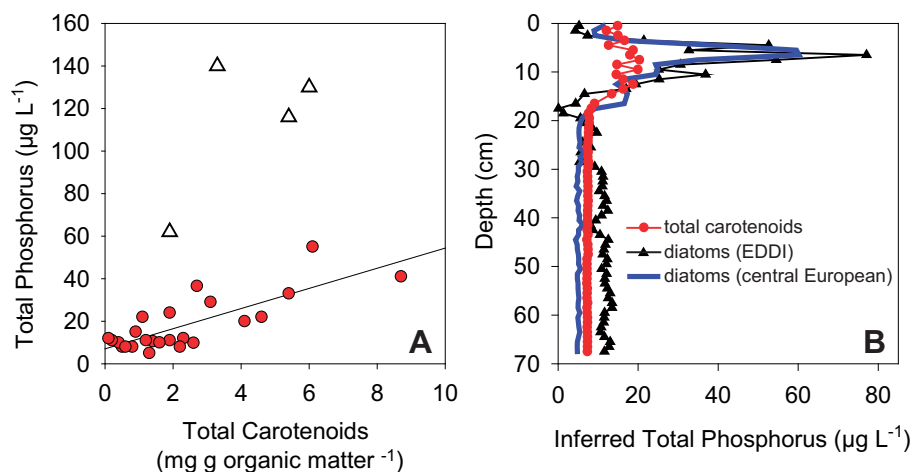


Fig. 16. Total carotenoids as a proxy for total phosphorus concentration in lake water. A) Correlation between total phosphorus of water samples with total sedimentary carotenoid concentrations ($r^2 = 0.78$, $n = 24$) in lakes from Italy (data from Guilizzoni et al., 2011). Four shallow lakes were outliers (indicated by the open triangles) and were not included in the correlation. B) Inferred total phosphorus reconstructions from a Lake Maggiore sediment core (data from Guilizzoni et al., 2011). The total phosphorus reconstruction based on total carotenoids is indicated by the red circles while two models based on diatoms remains for the European Diatom Database (EDDI) (Battarbee et al., 2000) and a regional central-European dataset (Wunsam and Schmidt, 1995) are shown by the black triangles and the solid blue line, respectively.

populations and processes in modern and ancient environments (e.g. Talbot et al., 2003c; Talbot and Farrimond, 2007; Cooke et al., 2009; Handley et al., 2010). Although sedimentary geohopanoids, analyzed by GC–MS techniques, are widely recognized (e.g. Ourisson and Albrecht, 1992) the investigation of intact BHP structures by APCI LC–MSⁿ is relatively new (Talbot et al., 2003a,b). Thus, the diversity of BHP structures and their potential as microbial markers is just beginning to be elucidated. Here, we highlight a few intact BHPs that seem potentially useful to (paleo)limnological studies but note that the reader should refer to Talbot and Farrimond (2007) for a detailed overview of the diversity of BHP structures and related source organisms.

7.1. Adenosylhopane, a tracer of soil OM to aquatic environments

Adenosylhopane (Fig. 1), 2-methyladenosylhopane and structurally related compounds have been proposed as being synthesized solely by soil bacteria (Cooke et al., 2008a). Thus far, adenosylhopane has been identified in two proteobacteria, the nitrogen-fixing bacterium *Bradyrhizobium japonicum* and the ammonia oxidizing bacterium *Nitrosomonas europaea* while 2-methyladenosylhopane has been identified in *B. japonicum* (Talbot and Farrimond, 2007; Cooke et al., 2009 and references therein). Adenosylhopane, 2-methyladenosylhopane and structurally related compounds have been found to be abundant in soils from a widely distributed geographical locations (Talbot and Farrimond, 2007, and references therein) and thus can serve as tracers of soil organic matter input to aquatic environments (Talbot and Farrimond, 2007; Cooke et al., 2008b, 2009; Handley et al., 2010; Kim et al., 2011), similar to the BIT Index (see Section 3.1). However, recently adenosylhopane has been identified as the first intermediate in hopanoid side chain synthesis in *Methylobacterium* (Bradley et al., 2010). The authors suggest that adenosylhopane may also occur as an intermediate in all hopanoid-producing bacteria, which, if proven correct, may limit its use as marker specifically for soil bacteria.

Adenosylhopane, 2-methyladenosylhopane and structurally related compounds also have been reported from a few lacustrine environments including Lake Windermere (UK), Lake Nkunga (Kenya) and Loch Ness (UK) (Talbot and Farrimond, 2007). High terrestrial inputs occur at both Lake Nkunga and Loch Ness (Ficken et al., 1998; Talbot and Farrimond, 2007) and thus these compounds

may be present in other lakes with similarly high terrestrial inputs. However, Talbot and Farrimond (2007) re-analyzed 24 lacustrine sediments with multistage MS that they had previously analyzed using single stage MS (Talbot et al., 2003c) but did not find adenosylhopane or 2-methyladenosylhopane at other lacustrine sites. Therefore, these BHPs can potentially serve as a tracer for soil organic matter to lacustrine environments but it remains to be seen whether these compounds are common in lacustrine sediments.

To date, the diagenetic alteration of intact BHPs has not been systematically investigated. Although many of the intact BHPs appear to be well preserved on long time scales (e.g. the presence of BHT in 50 Myr old sediments) (van Dongen et al., 2006), it has been noted that other intact BHPs are likely influenced by diagenesis and some also may be diagenetically produced. For example, in their 1.2 Ma record from the Congo deep-sea fan, Handley et al. (2010) found that the ratio of BHT to (BHT plus adenosylhopane) increased gradually with depth suggesting a potential diagenetic source. In a 70 Kyr sedimentary record from the Benguela Upwelling System, a decoupling between BHPs and geohopanoids is noted (Blumenberg et al., 2010). Here, it appears that the geohopanoids likely reflect laterally-transported fossil organic matter whereas the source of BHPs is from the water column thereby demonstrating that sedimentation processes can complicate using geohopanoids and BHPs for paleoenvironmental reconstructions (Blumenberg et al., 2010).

7.2. BHP markers for cyanobacteria

Cyanobacteria are important primary producers in many lacustrine systems and some types produce BHPs. Methylation at C-2 is a well-known structural feature of cyanobacterial BHPs (e.g. Summons et al., 1999; Talbot et al., 2008) and species producing C-2 methylated structures occur in all five divisions of cyanobacteria (Chroococcales, Pluerocapsales, Oscillatoriales, Nostocales and Stigonematales) (Talbot et al., 2008). However, C-2 methylated structures are not produced by all cyanobacteria. A study of 58 strains of cyanobacteria found that while most (49 strains) produce BHPs including BHT and aminotriol (discussed in Section 7.3), the C-2 methylated BHPs are not as widespread and are noted in only 21 strains (Talbot et al., 2008 and references therein). C-2 methylated structures also can be produced by other organisms including facultative methylotrophs and some nitrogen-fixing bacteria and

a purple non-sulfur bacteria grown anaerobically (Summons et al., 1999 and references therein; Rashby et al., 2007; Talbot et al., 2008; Welander et al., 2010). The hydrocarbon skeletons of BHPs are resistant to biodegradation and are extremely refractory, allowing for incorporation into kerogen or sulphur-linked macromolecules (Summons et al., 1999). Indeed, 2-methyl-BHP derivatives, the 2 α -methylhopanes have been found to be abundant in organic-rich sediments up to 2500 Myr old, where they were used as biomarkers for cyanobacterial oxygenic photosynthesis (Summons et al., 1999). However, more recently Welander et al. (2010) identified the protein responsible for hopanoid methylation at C-2 (*S*-adenosylmethionine (SAM) methylase encoded by *hpnP*) in cyanobacteria but also in acidobacteria and a subclade of the rhizobiales within α -proteobacteria. Thus, the findings of Welander et al. (2010), that *hpnP* is found in many modern bacteria that do not utilize oxygenic photosynthesis, argues against using 2-methylhopanoids as biomarkers for this process.

In addition to methylation at C-2, a number of other side chain structures have been reported for cyanobacteria (Talbot et al., 2008) and some of these structures seem to be unique to cyanobacteria. For example, specific cyanobacterial biomarkers may be provided by a 2-methyl cyclitol ether, which is currently only reported from the cyanobacterium *Anacystis montana*, and bacteriohopanepentol (BHpentol), which has only been reported to occur in *Nostoc muscorum* and *Nostoc* sp. (Talbot et al., 2008 and references therein). Other side-chain structures that are only known from cyanobacteria include an irregular pentol, an alternative tetrol glycoside, and *O*- β -3,5-anhydrogalacturonopyranosyl tetrol (Talbot et al., 2008). Additionally, novel structures have been identified in *O. amphigranulata*, *Calothrix* sp. and in an enrichment culture of *Gloeocapsa* (see Talbot et al., 2008 for structures). Some of these cyanobacterial BHPs have been reported from lacustrine sediments. For example, 2-methyl cyclitol ether was found in Loch Ness (UK) and in La Piscina de Yuriria (Mexico) while BHpentol was found in Lake Druzhby (Antarctica) and also in Loch Ness (Talbot and Farrimond, 2007).

7.3. BHP markers for methanotrophs

Methanotrophs are subdivided into two types depending on their carbon assimilation pathway; Type I methanotrophs are members of the γ -proteobacteria and utilize the ribulose monophosphate pathway whereas Type II methanotrophs are members of the α -proteobacteria and utilize the serine pathway (Hanson and Hanson, 1996; Talbot et al., 2003c). In aquatic settings methane consumption occurs by either anaerobic oxidation of methane (AOM) or aerobic methanotrophy (e.g. Zhu et al., 2010 and references therein). To date relatively few studies have conducted organic geochemical investigations of aerobic methanotrophy in lacustrine environments; however, this can be investigated by examining BHPs. In particular, 35-aminobacteriohopane-30,31,32,33,34-pentol (aminopentol), has been identified as a BHP specific for Type I methane oxidizing bacteria (Talbot and Farrimond, 2007 and references therein). Others BHPs, while somewhat less specific, can also be related to methane oxidizing bacteria. 35-aminobacteriohopane-31,32,33,34-tetrol (aminotetrol) is present in varying amounts in all methanotrophs (Talbot and Farrimond, 2007) but also in trace amounts in some species of sulfate reducing bacteria (Blumenberg et al., 2006). 35-Aminobacteriohopane-32,33,34-triol (aminotriol) is produced by all Type II methanotrophs, some Type I methanotrophs, as well as a diverse range of other microorganisms (Talbot and Farrimond, 2007 and references therein). Zhu et al. (2010) used these BHPs to trace aerobic methane oxidation across a large methane concentration gradient in surface sediments from the lower Yangtze River and the East China Sea.

The occurrence of aminopentol, aminotetrol, and aminotriol BHPs has also been noted in a number of lacustrine sediments. Talbot et al. (2003c) examined BHP distributions in a number of lakes including temperate, Antarctic, high-altitude, and saline lakes. The aminotriol BHP, presumably derived from methanotrophs, was found to be one of the major components in many of these sediments. The relative proportion of hexafunctionalized BHPs (such as aminopentol) has been observed to be high in small, stratified and highly productive lakes but low in marine settings (Farrimond et al., 2000; Watson and Farrimond, 2000; Talbot et al., 2003c). While sulphate reduction dominates over methanogenesis in marine sediments, methanogenesis is the dominant process in highly productive lakes and Type I methanotrophs, producers of hexafunctionalized BHPs, are known to occur in the water columns of these types of lakes (Talbot et al., 2003c).

7.4. Application of BHPs in an Antarctic lacustrine record

An interesting study of Ace Lake (Antarctica) examined both BHPs and 16S ribosomal RNA genes (16S rDNA) to help reveal sources for sedimentary BHPs and examine if changing environmental conditions are reflected in BHP distributions (Coolen et al., 2008). Ace Lake has a unique history; it was initially a meltwater filled freshwater lake but due to sea level rise following deglaciation, it was then connected to the ocean and became a stratified fjord with sulfidic bottom waters, and later, due to isostatic rebound, the lake was again re-isolated (Coolen et al., 2008 and references therein). Presently Ace Lake is saline with anoxic bottom waters that are characterized by high methane and sulphide concentrations (Coolen et al., 2008). Coolen et al. (2008) found that when Ace Lake was a fresh meltwater lake, the main BHPs present were aminotriol, aminotetrol and aminopentol BHPs (Fig. 17). Although aminotriol is produced by a wide range of organisms, compound-specific carbon isotope analysis of these lipids revealed a highly depleted ^{13}C content, confirming a methanotrophic origin for these BHPs (Jahnke et al., 1999; Coolen et al., 2008). In these sediments, phylotypes related to Type I methanotrophs of the genus *Methylomonas* and to Type II methanotrophs of the genus *Methylocystis* were detected (Coolen et al., 2008). When Ace Lake became a stratified fjord, a number of major changes are noted. Significantly lower levels of total BHPs are observed and 16s rDNA of methanotrophs comprises a smaller portion of the total bacterial 16S rDNA pool (Coolen et al., 2008). These observations are consistent with sulphate reduction likely dominating over methanogenesis at this time (Coolen et al., 2008). Subsequently, when Ace Lake returned to being an isolated lake, there is again evidence for an active methane cycle. Aminotetrol and aminotriol BHPs were found to be present in these sediments while aminopentol was absent (Fig. 17). While this distribution was previously considered a good indicator for Type II methanotrophs, the 16S rDNA data indicated that phylotypes related to Type II methanotrophs were not present (Coolen et al., 2008). Instead, there was evidence that a phylotype related to methanotrophic gill symbionts of deep-sea cold-seep mussels was the likely source of aminotetrol and aminotriol in these sediments. These phylotypes also produce ^{13}C -depleted 4-methylated sterols, compounds also found in methanotrophs isolated from Ace Lake (Schouten et al., 2001a).

7.5. Outlook: BHPs for lacustrine paleoenvironmental reconstructions

The use of BHPs such as aminopentol and 2-methyl cyclitol ether as biomarkers of methanotrophs and cyanobacteria, respectively, is of particular interest to lacustrine studies as these organisms are dominant in many lakes and their biomarkers may provide

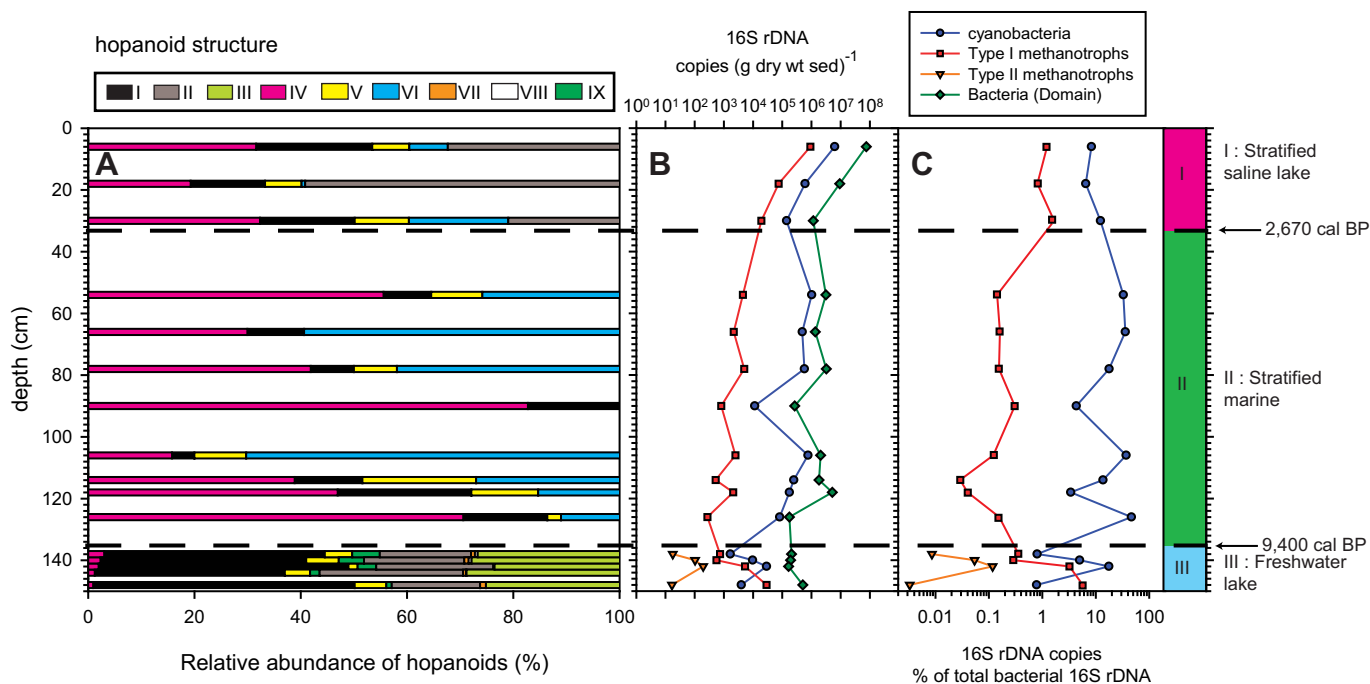


Fig. 17. BHP and 16S rDNA data from Ace Lake, Antarctica. Data from Coolen et al. (2008). A) The relative abundance of intact BHPs in Ace Lake sediments. BHPs structures indicate sources from: I (aminotriol), II (aminotetrol), III (Aminopentol), IV (BHT), V (bacteriohopanetetrol cyclito ether), VI (bacteriohopanetetrol glycoside), VII (bacteriohopanepentol cyclito ether), VIII (bacteriohopanepentol glycoside), IX (2-methylBHT) (see Coolen et al., 2008 for more details). B) Quantitative and C) relative distribution of sedimentary 16S rDNA of cyanobacteria, Type I methanotrophs, Type II methanotrophs, and bacteria (general). Shifts in the relative distribution of BHPs and 16S rDNA coincide with the depositional history of Ace Lake reflecting changes from a freshwater lake to stratified marine system, to a stratified saline lake.

opportunities for investigating past methane or nitrogen cycling. However, at present only a few studies have examined BHPs in lakes and thus more investigations are required to determine how widespread and abundant BHPs are in lacustrine sediments. In addition, further studies are certainly needed to better assess diagenetic and lateral transport effects on BHPs as well as the occurrence of BHPs (e.g. adenosylhopane) in other hopanoid producing bacteria (Bradley et al., 2010).

8. Lipid biomarker indicators of primary producers and grazers

Microalgae are generally the major sources of lipids in lacustrine environments and there are biomarkers for a number of different types microalgae. Above we already discussed the examples of alkenones as biomarkers for haptophyte algae, carotenoids for different types of photoautotrophs and BHPs for cyanobacteria. Here, we highlight a few examples of other biomarkers that can be used to examine primary productivity of selected algal classes in many lacustrine environments. We note that most of these microalgal biomarkers have been previously reviewed by Volkman et al. (1998) and Volkman (2003), and the reader should refer to these publications for detailed information on the wide variety of lipids present in microalgae. For a detailed review on the distribution of sterols in diatoms we refer the reader to the comprehensive study of Rampen et al. (2010). The algal lipids discussed in Sections 8.1–8.4 are generally analyzed by GC and GC/MS.

8.1. Dinosterol, a biomarker for dinoflagellates

Sterols are compounds that occur in all eukaryotes and are membrane rigidifiers, and the specificity of some of these compounds for different phytoplankton groups is well known (Volkman, 1986, 2003; Volkman et al., 1998). The compound

dinosterol (4 α ,23,24-trimethyl-5 α -cholest-22-en-3 β -ol; Fig. 1) is found in many dinoflagellate (Pyrrophyta) species (Withers, 1983; Piretti et al., 1997) and is commonly used as a biomarker for these organisms (Boon et al., 1979; Robinson et al., 1984; Volkman, 2003). While many sterols are produced by both aquatic algae and terrestrial plants, dinosterol is not synthesized by higher plants and is thus recognized as a robust biomarker for dinoflagellates (Volkman et al., 1999). However, it should be noted that dinosterol is not found in all dinoflagellate species (Kokke et al., 1981) and in fact a study by Leblond and Chapman (2002) only found it in about half of 43 dinoflagellate species examined. Dinosterol can also be produced by a few diatoms and haptophyte algae (Volkman et al., 1993, 1997). Nevertheless, dinoflagellates are an important primary producer in many lakes and since dinosterol is present in many lake sediments, it may be useful for examining past changes in dinoflagellate productivity in some systems (e.g. Castañeda et al., 2011).

8.2. Diatom biomarkers

A number of different sterols are often used as biomarkers for diatoms. For example, the compound 24-methylcholesta-5,22E-dien-3 β -ol is known as “diatomsterol” (Fig. 15); however, this sterol and other common diatom sterols are produced by a large number of algal classes (Volkman, 2003) thus restricting their use as biomarkers for diatoms. In fact, a recent study that analyzed sterols in 106 diatom cultures revealed that all of the major sterols found in diatoms are also common sterols of other algal groups (Rampen et al., 2010). The most common sterols found in diatoms are 24-methylcholesta-5,24(28)-dien-3 β -ol and the Δ^5 sterols, cholest-5-en-3 β -ol (cholesterol), 24-methylcholest-5-en-3 β -ol, and 24-ethylcholest-5-en-3 β -ol (Rampen et al., 2010). Notably, “diatomsterol” was found to be absent in some diatoms and relatively high concentrations of this compound were only observed in pennate

diatoms (Rampen et al., 2010). Furthermore, some of the major sterols found in diatoms are also produced by higher plants (Nishimura and Koyama, 1977) thus restricting their utility as an aquatic biomarker in many systems. However, in lakes that receive little terrestrial input, a number of sterols could potentially serve as diatom biomarkers. In lakes where diatoms are one of the dominant primary producers, sterol distributions could provide information regarding the class of diatoms present (Rampen et al., 2010).

The compounds loliolide (Fig. 1) and isololiolide are the anoxic degradation products of the pigment fucoxanthin, the major carotenoid present in diatoms (Klok et al., 1984; Repeta, 1989), and can provide a diatom biomarker in some environments. When fucoxanthin undergoes anoxic degradation, loliolide (isololiolide) is produced on a mole-to-mole basis (Repeta, 1989) and this process can occur both in the water column and in the sediments. Although dinoflagellates and haptophyte algae can also contain fucoxanthin (Klok et al., 1984; Jeffrey and Vesk, 1997), loliolide and isololiolide can provide a reliable biomarker for diatoms in lakes where dinoflagellates are not a major contributor to primary productivity and where haptophyte algae are not present (e.g. Castañeda et al., 2011).

Highly branched isoprenoid alkanes (HBIs) (Fig. 15) are highly specific biomarkers for diatoms and are produced by four genera: *Haslea* (Volkman et al., 1994; Belt et al., 1996; Wraige et al., 1997, 1998; Allard et al., 2001; Sinninghe Damsté et al., 2004), *Rhizosolenia* (Volkman et al., 1994; Sinninghe Damsté et al., 1999b; Rowland et al., 2001; Massé et al., 2004; Sinninghe Damsté et al., 2004), *Pleurosigma* (Belt et al., 2000; Belt et al., 2001a; Grossi et al., 2004), and *Navicula* (Belt et al., 2001b; Sinninghe Damsté et al., 2004). While the majority of known HBI producers are marine (both benthic and planktonic species) diatoms, the freshwater diatom *Navicula sclesvicensis* produces a C₂₅ HBI triene (Belt et al., 2001b). HBIs have been noted in sediments from several lakes including saline lake Small Meromictic Basin in Ellis Fjord, Antarctica (Sinninghe Damsté et al., 2007), and freshwater Mud Lake, Florida (Filley et al., 2001), Lake Koucha (Aichner et al., 2010a) and Lake Lugano (Bechtel and Schubert, 2009). In a sediment core from Lake Lugano (Switzerland) spanning the interval from 1900 AD to the present, increased concentrations of HBIs, in agreement with high 24-methylcholesta-5,22-dien-3b-ol (brassicasterol) concentrations, provided evidence for periods of increased input from diatoms (Bechtel and Schubert, 2009). In a 16 cal ka record from Lake Koucha (Tibetan Plateau), the appearance of an unsaturated C₂₅ HBI at 4.7 cal ka BP is noted along with increased concentrations of other algal biomarkers since ~6.1 cal ka BP (Aichner et al., 2010a). Lake Koucha transitioned from a saline to a freshwater system at ~7.2 cal ka BP and after this time a number of phytoplankton biomarkers are observed whereas previously the lake had been dominated by aquatic macrophytes (Aichner et al., 2010a).

Overall, molecular biomarkers, and in particular the highly specific HBIs, can be useful for examining past diatom productivity, especially in lakes where silica dissolution has occurred. However, diatoms are often well preserved in many lacustrine sediments and thus highly specific species level information can be gained from examining diatom assemblages, which in many cases cannot be gained from lipid analysis. Nevertheless, for studies targeting past changes in diatom productivity it could be useful to examine diatom biomarkers in conjunction with biogenic silica or diatom assemblage data.

8.3. Tetrahymanol, a biomarker for bacterivorous ciliates

The compound tetrahymanol (gammaceran-3β-ol; Fig. 15) is abundant in many lake sediments (e.g. Hanisch et al., 2003;

Castañeda et al., 2011) and is produced by bacterivorous ciliates, such as the freshwater ciliate *Tetrahymena* (Mallory et al., 1963; Harvey and McManus, 1991). More specifically, it is recognized that ciliates biosynthesize tetrahymanol only when their diet does not contain sterols (Harvey and McManus, 1991) and in modern stratified environments, ciliates feed on organisms found at or below the chemocline such as purple sulphur bacteria and sulphide-oxidizing bacteria (Sinninghe Damsté et al., 1995 and references therein). Tetrahymanol can also be produced by the anaerobic purple bacterium *Rhodospseudomonas palustris* (Kleemann et al., 1990), anaerobic rumen fungus (Kemp and Lander, 1984), and in small amounts by a fern (Zander et al., 1969). Sources from rumen fungus and ferns are unlikely in many lake sediments but anaerobic purple bacteria could be a potential source of tetrahymanol in some environments. However, in aquatic sediments tetrahymanol typically derives from ciliates (Harvey and McManus, 1991) and previous studies have attributed its occurrence to the presence of a stratified water column (e.g. Hanisch et al., 2003; Xu and Jaffé, 2008) since bacterivorous ciliates are typically found at the oxic–anoxic boundary where large bacterial populations are present (Sinninghe Damsté et al., 1995; Thiel et al., 1997). In a study of Lake Valencia (Venezuela), it was found that an abrupt increase in tetrahymanol concentration occurred at ~7260 cal BP whereas prior to this time its abundance was below the detection limit (Xu and Jaffé, 2008). The authors attributed the increase in tetrahymanol concentrations to the establishment of an oxic–anoxic boundary in the water column of Lake Valencia, supported by independent evidence for lake-level changes between the Late Pleistocene and early Holocene (Xu and Jaffé, 2008 and references therein). It also should be noted that the compound gammacerane, a C₃₀ triterpane which has been found to be abundant in many lacustrine deposits, is produced from tetrahymanol by diagenesis and early catagenesis (ten Haven et al., 1989).

8.4. Long-chain diols as a biomarker for Eustigmatophyte algae

Long-chain (C₂₈–C₃₂) 1,15 *n*-alkyl diols (Fig. 11) are produced by Eustigmatophyte (yellow-green) algae (Volkman et al., 1992; Versteegh et al., 1997), a class that relatively little is known about (Ott and Oldham-Ott, 2003). Long-chain *n*-alkyl diols also have been noted in many lacustrine sediments, and often they are present in high abundances (Xu et al., 2007; Xu and Jaffé, 2008; Castañeda et al., 2009c; Shimokawara et al., 2010). Interestingly, long-chain *n*-alkyl diols have been reported from many lakes although eustigmatophyte algae have not yet been positively identified from any these lakes, to the best of our knowledge. Most species of eustigmatophytes are very small (2–4 mm) and can be easily confused with coccoid forms of Chlorophyceae or Xanthophyceae and thus this group may have been overlooked in algal surveys (Ott and Oldham-Ott, 2003). In addition, the class Eustigmatophyceae includes members that inhabit terrestrial soils (Ott and Oldham-Ott, 2003) and so terrestrial sources to lakes may be possible but previous studies point to aquatic sources for the long-chain *n*-alkyl diols (Xu et al., 2007; Xu and Jaffé, 2008; Castañeda et al., 2009c; Shimokawara et al., 2010). Furthermore, a study of the phytoplankton in Lake Tanganyika (East Africa) based on polymerase chain reaction-amplified 18S ribosomal DNA, provides evidence for the presence of eustigmatophytes possessing a sequence similar to marine and freshwater members of *Nannochloropsis* (De Wever, 2006). Alternatively, it has been suggested that cyanobacteria may be a source of the long-chain 1, 15 *n*-alkyl diols in lakes (Xu et al., 2007). At Lake Valencia (Venezuela), one of the most productive lakes in the world, primary productivity is dominated by cyanobacterial species, which comprise >90% of the biomass (Xu et al., 2007). The exceptionally high abundance of diols

in this lake, supported by similar $\delta^{13}\text{C}$ values of the diols and other cyanobacterial biomarkers, suggests that cyanobacteria may be a source of these compounds in Lake Valencia (Xu et al., 2007). Given widespread occurrence of the long-chain *n*-alkyl diols in lakes, and the fact that they are often present in high abundances, there is a need to identify the producer(s) of these lipids in freshwater environments.

8.5. Examining past changes in productivity from biomarkers of primary producers: an example from Lake Malawi

Examining changes in the abundance of several different aquatic biomarkers provides a method for examining past changes in primary productivity. While biomarker abundances cannot be directly correlated to biomass due to complications arising from degradation and variations in cellular lipid concentrations, changes in the relative abundances of microalgal lipids can be used to examine past trends in aquatic ecosystem structure and productivity (e.g. Holtvoeth et al., 2010; Kristen et al., 2010). For example, in a Lake Malawi sediment core spanning the past 700 years, Castañeda et al. (2011) utilized the accumulation rates of loliolide, dinosterol, long-chain *n*-alkyl diols, and tetrahymanol as productivity tracers of diatoms, dinoflagellates, eustigmatophyte algae and bacterivorous ciliates, respectively (Fig. 18). In modern Lake Malawi, where dinoflagellates are a minor contributor to algal productivity and haptophyte algae are not present, it was found that accumulation rates of loliolide exhibited the same overall trends as biogenic silica, another proxy for diatom productivity. The biomarker records revealed several notable changes including an increase in dinoflagellate and bacterivorous ciliate biomarkers over the past few centuries, accompanied by a decrease in diatom lipids (Fig. 18). Independent evidence from Lake Malawi showed that a change in the dominant wind direction over the lake occurred at the end of the Little Ice Age (Brown and Johnson, 2005), with the

northern end of Lake Malawi experiencing stronger or more frequent northerly winds prior to this time. Diatom productivity in the northern basin of Lake Malawi is sensitive to changes in wind strength and direction, which control upwelling. The decreased diatom productivity noted in Lake Malawi after ~1780 AD coincides with a shift to less frequent or weaker northerly winds over the lake. A change in the dominant wind regime may also explain the increased abundances of dinoflagellate biomarkers after this time since studies have suggested that dinoflagellate blooms are favoured by a stable water column with minimal mixing (Pollinger and Zemel, 1981). In contrast to diatoms, which rely on turbulent mixing to remain in the photic zone, turbulent mixing can destroy or damage dinoflagellate cells (Pollinger and Zemel, 1981). Observations from quantitative algal surveys of Lake Malawi also support the increased presence of dinoflagellates with recent blooms observed (Hecky et al., 1999).

9. Fossil DNA indicators of primary producers and grazers

Ancient DNA preserved in sediments (fossil DNA) is a relatively recent tool that can be used to reconstruct past ecosystems and has been used to examine environmental changes in specific marine, lacustrine and terrestrial settings (Coolen et al., 2004b, 2008; D'Andrea et al., 2006; Willerslev et al., 2007; Boere et al., 2009). While the majority of lipid biomarkers cannot be used to provide taxonomic differentiation at the species level, fossil DNA is perhaps the ultimate biomarker as it can be used to provide species level information from past environments and thus there is much interest in developing these techniques for application in Quaternary science. However, DNA is fragile and degradation of ancient DNA to shorter fragments occurs relatively rapidly after deposition, within the first several thousand years (Coolen and Overmann, 1998). Nevertheless, the shorter DNA fragments (less than 500 base pair-long (Coolen and Gibson, 2009)) can be analyzed using

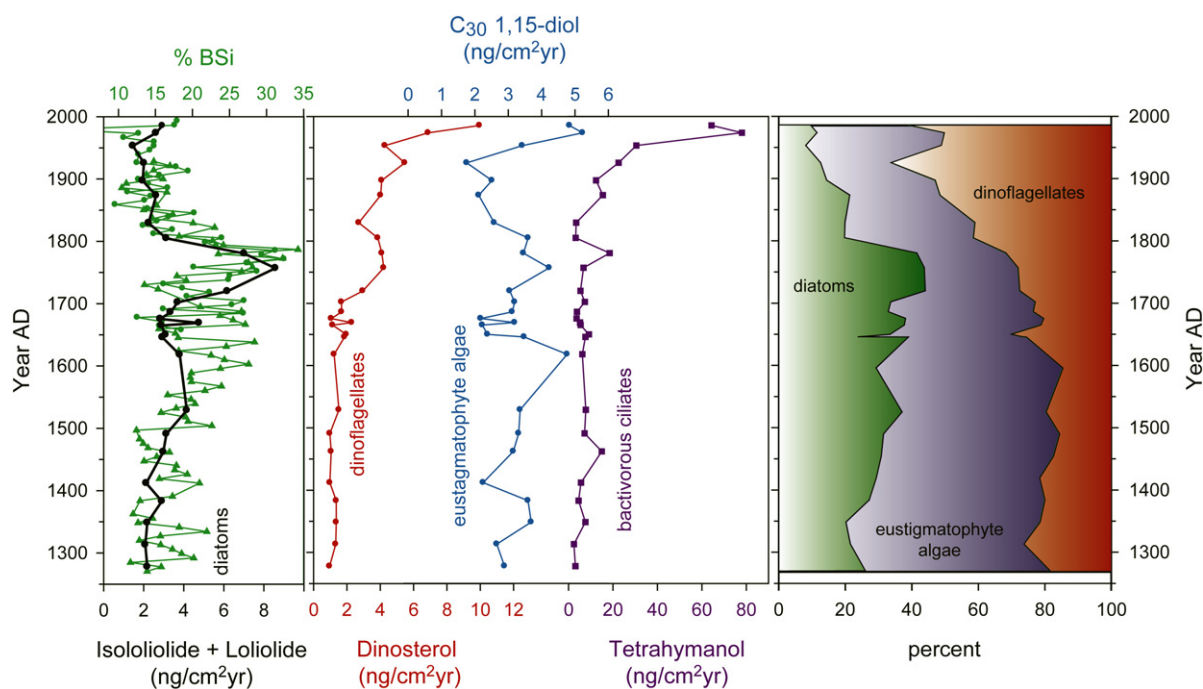


Fig. 18. Lake Malawi productivity biomarkers (data from Castañeda et al., 2011). Mass accumulation rates of isololiolide and loliolide, a diatom biomarker, are plotted with biogenic silica (data from Johnson et al., 2001), another diatom proxy. Mass accumulation rates of dinosterol (a biomarker of dinoflagellates), the C₃₀, 1,15 *n*-alkyl diol (a biomarker of eustigmatophyte algae) and tetrahymanol (a biomarker for bacterivorous ciliates) are also shown. The panel on the far right is a relative percentage plot of the main aquatic biomarkers present in Lake Malawi sediments. One compound representative of each group (diatoms, eustigmatophyte algae and dinoflagellates) was chosen and abundances were normalized to 100%. An increase in dinoflagellates accompanied by a decrease in diatoms is noted since ~1800 AD.

polymerase chain reaction (PCR) techniques to amplify DNA and produce billions of copies of a particular DNA sequence. Sedimentary conditions favourable to DNA preservation include cold, anoxic and light-free environments (Coolen et al., 2004b) and fossil DNA analysis has proven to be especially useful to Holocene studies of polar lakes (e.g. Coolen et al., 2004b, 2008). Another complicating factor with fossil DNA analysis is that metabolically active organisms in deep waters and sediments contribute to the DNA pool; however, species that express genes in deep waters and sediments at the time of sampling can be examined and excluded as paleo-environmental proxies (Coolen and Gibson, 2009). Additionally, fossil DNA analysis can be conducted along with lipid biomarker analysis to further ground-truth fossil DNA data and also provide insights into lipid biomarker distributions (Coolen and Gibson, 2009). To date, fossil DNA has commonly been used to examine the diversity of past planktonic communities including unicellular aquatic organisms such as copepods (Bissett et al., 2005; Gibson and Bayly, 2007).

An interesting study by Boere et al. (2009) utilized a multi-proxy approach to examine fossil DNA, lipid biomarkers and palynomorphs of dinoflagellates from Holocene sediments of Small Meromictic Basin, Ellis Fjord (Antarctica), which is in the advanced stages of a transition from a fjord system to a meromictic saline lake. Their fossil 18S rDNA-based survey revealed at least nine different dinoflagellate species have lived in Ellis Fjord and they found that fossil DNA was preserved for ~2700 years in organic carbon-rich and sulfidic sediments of Small Meromictic Basin. At ~1850 cal yr BP a community shift is noted and the autotrophic sea-ice dinoflagellate *Polarella glacialis* was found to be the main precursor of fossil dinoflagellate biomarkers in the younger sediments. In contrast, abundances of dinosterol, the only dinoflagellate-diagnostic lipid present in the sediments, suggested decreasing contributions of dinoflagellates to sediments in the youngest portion of the core. However, they note that *P. glacialis* was found not to produce dinosterol in culture studies and thus studies using only dinosterol as a dinoflagellate biomarker may lead to biased interpretations of past changes in dinoflagellate abundance. Boere et al. (2009) were not able to relate specific palynomorph morphotypes to the 18S rDNA sequences but they note that rare species that were present as cysts were not detected with the phylogenetic analysis. Thus, these three approaches provide complimentary information regarding past dinoflagellate succession in Small Meromictic Basin.

Overall, fossil DNA analysis is proving to be an extremely powerful approach for examining past species composition in polar lakes and also for examining biological sources of lipid biomarkers. However, many outstanding questions presently exist regarding the factors and conditions that control DNA preservation, which need to be addressed before fossil DNA techniques can be more broadly applied (Coolen and Gibson, 2009). Distinguishing DNA signals from living and fossil populations of microorganisms, as well as contamination, also continues to be a challenge and care must be taken when utilizing fossil DNA for paleoenvironmental studies.

10. New biomarkers and techniques potentially useful for lacustrine studies

New biomarkers and organic geochemical proxies are continually being discovered and developed. In this section we highlight a few recently discovered biomarkers and research developments that will likely be useful for lacustrine studies but that, to our knowledge, have not yet been widely applied to paleolimnological investigations. Several new biomarkers can be related to nitrogen or methane cycling and thus are of much interest for Quaternary studies. Nitrogen is often a limiting element for primary

productivity and therefore plays a major role in both ocean and lake biogeochemistry. Likewise, methane is an important greenhouse gas and biological production by methanogens (anaerobic members of the Archaea) is the source of approximately 70% of global methane (Schimel, 2004). In addition to these new biomarkers, we also discuss recent developments in using compound-specific radiocarbon analysis as a tool for improving lacustrine sediment chronologies.

10.1. Tracing past N_2 fixation

In addition to the BHPs discussed in Section 7.2, long-chain glycolipids (Fig. 1) have been recently reported as biomarkers of cyanobacteria, specifically for N_2 -fixing heterocyst cyanobacteria. Two intact glycolipids, docosanyl 3-*O*-methyl- α -rhamnopyranoside and docosanyl 3-*O*-methylxylopyranoside, were first reported in sediments from Ace Lake (Antarctica) and a cyanobacterial origin was suggested for these sedimentary glycolipids (Sinninghe Damsté et al., 2001b). Docosanyl 3-*O*-methylxylopyranoside was additionally reported from Lake Malawi (East Africa), where cyanobacteria are one of the dominant primary producers (Castañeda et al., 2009b). Both of these studies utilized GC and GC/MS techniques to analyze these glycolipids. More recently, HPLC/ESI-MS² techniques have been used to examine the glycolipid content of 34 axenic strains of cyanobacteria (Bauersachs et al., 2009a). It was found that species of the families Nostocaceae and Rivulariaceae, which are capable of biosynthesising heterocysts (specialized cells that contain the nitrogenase enzyme and that are the site of N_2 fixation), contain a suite of glycolipids that consist of a sugar moiety glycosidically bound to long-chain diols, triols, keto-ols and keto-diols (Bauersachs et al., 2009a). These compounds appear to be a unique structural component of the heterocyst cell envelope (Nichols and Wood, 1968; Gambacorta et al., 1998) and when found in sediments can be used as biomarkers of N_2 -fixing heterocystous cyanobacteria (Bauersachs et al., 2010). Indeed, long-chain heterocyst glycolipids have been found in both the water column and sediments of Lake Challa (Kenya) and in sediments from the Baltic Sea where heterocystous cyanobacteria form an important component of the present-day phytoplankton community (Bauersachs et al., 2010). Furthermore, long-chain heterocyst glycolipids are excellent biomarkers since they are well preserved in several lake surface sediments and have been reported from Oligocene sediments from Lake Enspel as well as in Eocene sediments from Lake Messel and the Green River Shale (Bauersachs et al., 2010). Since nitrogen-fixing cyanobacteria are present in many lakes, the analysis of long-chain heterocyst glycolipids provides an opportunity to examine past cyanobacterial production and nitrogen cycling.

10.2. Tracing past anaerobic oxidation of ammonium

The anaerobic oxidation of ammonium (anammox) to di-nitrogen gas with nitrate as an electron acceptor is a globally important microbial process in both marine and freshwater nitrogen cycling. The bacteria that perform the anammox process belong to a distinct phylogenetic group related to the *Planctomycetes* (Strous et al., 1999; Schmid et al., 2007). The anammox reaction occurs with a specialized intracellular compartment, the anammoxosome, which has a membrane consisting of a dense layer of ladderane lipids (Sinninghe Damsté et al., 2002b). These unusual lipids consist of three or five linearly linked cyclobutane rings (Fig. 15) that are either ester or ether bonded to a glycerol backbone (Sinninghe Damsté et al., 2002b) and can be analyzed by GC and GC/MS (Sinninghe Damsté et al., 2005) but analysis by HPLC/APCI-MS² is shown to be superior in sensitivity (Hopmans et al., 2006).

Ladderane lipids are unique to anammox bacteria and thus serve as excellent biomarkers for the detection of anammox in present and past environments (Sinninghe Damsté et al., 2002b; Jaeschke et al., 2007, 2009a,b; Rattray et al., 2008). Anammox bacteria have been detected in a wide variety of environments including marine and lacustrine environments, marshes, sea ice, and in permafrost and agricultural soils (Humbert et al., 2010 and references therein). In aquatic environments, they are active at redox transition zones such as oxygen-minimum zones (Kuypers et al., 2003; Jaeschke et al., 2009b).

Interestingly, it has been observed in culture studies that anammox bacteria alter their membrane lipid composition in response to temperature with increased amounts of shorter-chain ladderane fatty acids relative to the amount of longer-chain fatty acids occurring at lower temperatures (Rattray et al., 2010). An index, called NL₅, has been proposed to quantify this relative temperature change (Rattray et al., 2010). Rattray et al. (2010) note that most of the changes in ladderane lipid chain length occur between 12 °C and 20 °C with no significant changes observed at lower or higher temperatures. They suggest that the NL₅ index can be useful for distinguishing between fossil ladderane lipids that originate in the relatively warmer upper water column and those that are produced *in-situ* in relatively cold surface sediments, which is necessary for examining past anammox. Jaeschke et al. (2009b) utilized the NL₅ index to examine past anammox in a sediment core from the northern Arabian Sea spanning the last 140 kyr. They found that low concentrations of ladderane lipids, indicating low-anammox activity, coincided with periods when the oxygen-minimum zone was severely diminished while high concentrations of ladderane lipids coincided with periods characterized by an intense oxygen-minimum zone.

At present, little is known about the importance of anammox in past marine nitrogen cycles but essentially nothing is known about its importance in both present and past lacustrine nitrogen cycles. However, the anammox process has been reported from meromictic Lakes Tanganyika (East Africa) and Rassnitzer (Germany) (Schubert et al., 2006; Hamersley et al., 2009) and likely occurs in many lakes. To date, ladderane lipids have not yet been used to examine past anammox in lacustrine environments but thermally immature sediments from lakes with anoxic bottom waters are potentially excellent targets since the preservation potential of ladderane lipids is likely high in these environments. It should be noted that although not much is currently known about the diagenesis of ladderane lipids, a thermal degradation study revealed that structural alterations occur at relatively low temperatures of 120 °C and therefore, ladderane lipids are unlikely to occur in ancient sediments (Jaeschke et al., 2008). However, ladderane lipids will likely prove useful for examining past anammox at least on Quaternary time scales.

10.3. Compound-specific $\delta^{15}\text{N}$ of pigments as a tool to investigate past nitrogen cycling

The nitrogen isotopic composition ($\delta^{15}\text{N}$) of bulk sediments is a widely used paleolimnological proxy for examining sources of organic matter, reconstructing past productivity rates and for studying past nitrogen cycling (Meyers and Teranes, 2001; Meyers, 2003). However, interpretation of bulk $\delta^{15}\text{N}$ records can be complicated by a variety of factors including remineralization, input from terrestrial sources and shifts in phytoplankton or heterotroph assemblages. Thus, it is advantageous to examine the $\delta^{15}\text{N}$ of molecular fossils having a definitive biological origin. In aquatic environments pigments such as chlorophylls and bacteriochlorophylls are markers of photoautotrophic activity and these compounds also contain nitrogen, making them attractive targets

for $\delta^{15}\text{N}$ analysis. The first $\delta^{15}\text{N}$ measurements of sedimentary geoporphyryns, the tetrapyrrole skeletons of chlorophyll molecules, were conducted by purifying compounds (e.g. by using preparative HPLC and size exclusion chromatography) and analyzing them with elemental analyzer isotope ratio mass spectrometry (EA-IRMS) (Sachs et al., 1999). Using these methods, Sachs and Repeta (1999) found evidence for the widespread occurrence of nitrogen fixation in the Eastern Mediterranean Sea during Late Pleistocene sapropel events. However, this approach is time consuming and requires large sample sizes, limiting its application to paleoenvironmental studies. While compound-specific isotope analysis requires minimal amounts of material, compounds such as tetrapyrroles are not very volatile and are difficult to analyze by GC-IRMS.

Recently, new techniques have been developed that allow for more rapid compound-specific $\delta^{15}\text{N}$ measurements of smaller samples. Chikaraishi et al. (2008) developed a method for analysing the $\delta^{15}\text{N}$ of tetrapyrroles by applying a chemical treatment to degrade them into monopyrrole unites (maleimides) and then subsequently measuring their isotopic composition by GC combustion IRMS (GC/C/IRMS). They found that the isotopic composition of the maleimides reflected the isotopic composition of the individual tetrapyrroles and also that no substantial differences existed in the isotopic composition of maleimides derived from a single tetrapyrrole. Polissar et al. (2009) developed a method termed nano-EA-IRMS, which can measure the isotopic composition of nanomolar quantities of C and N (for example, from individual compounds isolated by HPLC techniques). Higgins et al. (2009) developed a method for rapid analysis of sedimentary porphyrins that involves first separating and purifying them from sediment samples using HPLC, oxidizing them to nitrate in a two-step process, followed by isotopic analysis of the nitrate. Their approach measures the $\delta^{15}\text{N}$ of a mixture of porphyrins, rather than the $\delta^{15}\text{N}$ of individually purified compounds, which reduces sample preparation time and also the amount of sediment needed. These methods appear promising for examining compound-specific nitrogen signatures, and thus, past nitrogen cycling.

10.4. $\delta^{13}\text{C}$ of GDGTs as tracer of methanotrophic archaea

The stable carbon isotopic composition ($\delta^{13}\text{C}$) of biphytanes derived from GDGTs can be used to indicate the presence of methanotrophic archaea. These organisms utilize ^{13}C -depleted methane and thus ^{13}C -depleted isoprenoid GDGTs have been used as indicators for methanotrophic archaea in a variety of environments (Pancost et al., 2001; Schouten et al., 2001b, 2003; Wakeham et al., 2003; Blumenberg et al., 2004; van Dongen et al., 2007). For example, Pancost et al. (2000) examined biphytanes with 0–2 cyclopentane moieties in Holocene peat deposits and found relatively ^{13}C -depleted values for biphytanes with 1 and 2 cyclopentane moieties (–34 to –34‰) whereas those with 0 cyclopentane moieties were relatively ^{13}C -enriched (–27 to –30‰). They suggest that the biphytanes with 1 and 2 cyclopentane moieties were derived from acetotrophic methanogens while those with 0 cyclopentane moieties were derived from CO₂-reducing autotrophic methanogens.

Schouten et al. (2001a) examined the $\delta^{13}\text{C}$ of a range of organic compounds present in Ace Lake, Antarctica including sedimentary biomarkers of methanogenic archaea and methanotrophic bacteria. These included the diether lipid archaeol, which occurs ubiquitously in archaea, *sn*2-hydroxyarchaeol, which occurs in methanogenic archaea within the orders Methanococoides and Methanosarcinales, and saturated and unsaturated 2,6,10,15,19-pentamethyleicosane (PME), which are also attributed to methanogenic archaea (Schouten et al., 2001a). Depleted ^{13}C values (–45‰ to –133‰) typically characterize lipids of methanotrophic

bacteria that oxidize methane aerobically. Schouten et al. (2001a) found relatively enriched isotopic compositions of archaeol (-17%) and polyunsaturated PME (-23% to -28%), and concluded that little anaerobic methane oxidation takes place in Ace Lake but rather the archaea are involved only in the methane generation process. They also note that the isotopic composition of 4 α -methyl-5 α -cholest-8(14)-en-3 β -ol and its diagenetic derivatives displayed depleted ^{13}C values (-57%), which is diagnostic for bacterial methanotrophy (aerobic consumption of methane). Coolen et al. (2004a) later provided additional support for bacterial methanotrophy in Ace Lake with a combined study of lipid biomarkers and 16S rDNA genes (see Section 7.4). These studies demonstrate that isotopic analysis of lipid biomarkers such as archaeol or GDGTs can provide detailed insights into past methane cycling but to date, these techniques have not been widely applied to lacustrine studies.

10.5. Advances in compound-specific radiocarbon analysis

A reliable sediment chronology is a fundamental requirement for paleoenvironmental investigations but creating an accurate age model can be difficult. Radiocarbon dating is a widely applied dating technique utilized in Quaternary studies; however, radiocarbon ages based on bulk organic matter can be unreliable since it consists of a complex mixture of organic carbon derived from multiple sources varying in radiocarbon age (e.g. Eglinton et al., 1997; Smittenberg et al., 2004). While AMS dating of macrofossils is a preferable approach for reconstructing sediment chronology, such macrofossils are not always present. An alternative approach is compound-specific radiocarbon analysis (CSRA), which utilizes accelerator mass spectrometry (AMS) techniques in combination with automated preparative capillary gas chromatography (Eglinton et al., 1996). CSRA is presently emerging as a powerful method for creating and improving marine and lacustrine chronologies.

Uchikawa et al. (2008) used CSRA to date long-chain *n*-alkanes in Holocene sediments from Ordy Pond (Hawaii) and compared these ages to those derived from bulk sediment and plant macrofossils. They found that ^{14}C ages of the *n*-alkanes and terrestrial plant macrofossils were in good agreement but note substantially older ^{14}C ages for the bulk sediment, which is mainly composed of algal material (Uchikawa et al., 2008 and references therein). They note that ^{14}C -depleted Pleistocene limestone likely contributes to the DIC pool in Ordy Pond and thus autochthonous material produced from the lake water should be avoided for ^{14}C dating.

More recently, Hou et al. (2010) developed a method for radiocarbon dating individual lignin phenols, which are produced by vascular plants and can be isolated using reversed phase HPLC. The authors argue that lignin phenols are a good choice for radiocarbon dating since they are abundant in lake sediments, they are generally transported to lake sediments by water and thus derive from within the lake catchment, and they are rapidly recycled in the soils making it is unlikely that ^{14}C -depleted lignin will reach lake sediments (Hou et al., 2010). Hou et al. (2010) constructed age models for several lakes based on ^{14}C dating of lignin phenols and found the ages of lignin phenols to be consistent with varve counts or ^{14}C ages of terrestrial plant fragments, supporting a short residence time of lignin phenols in the catchment. However, they note that in an arctic lake from southwest Greenland, which is surrounded by permafrost soils, the lignin phenol ^{14}C ages are significantly older than bulk sediment ages, which is likely due to remobilization or preserved lignin in soils (Hou et al., 2010). Thus, this method appears promising from developing more accurate chronologies although in some systems the long residence time of lignin phenols will hamper this approach. Although CSRA has not

yet been widely applied to dating lacustrine sediments, it is likely that age differences exist between terrestrial and aquatic components of organic matter in many systems and thus this approach will be useful for creating reliable age models, which are especially important to high-resolution paleoclimate studies.

In addition to improving sediment chronologies, CSRA is a useful tool for examining the sources, fixation, and transport and fate of organic carbon in the geologic environment. CSRA can also be used to elucidate microbial metabolic pathways or to distinguish anthropogenic and natural inputs of organic carbon. In paleoceanography, CSRA has been widely applied to examine lateral transport. While age offsets between foraminifera and organic compounds such as alkenones have been noted in the same marine sediment samples and related to differences in lateral transport (e.g. Ohkouchi et al., 2002), age offsets also have been noted between different types of organic compounds. For example, several marine studies have examined the ^{14}C ages of crenarchaeol and alkenones and have concluded that isoprenoid GDGT distributions are less affected by lateral transport than those of alkenones (Mollenhauer et al., 2007, 2008; Shah et al., 2008). Although lateral transport likely does not induce substantial age differences between different aquatic components of organic matter in smaller lakes, it could conceivably be an issue in larger lakes. CSRA thus provide an important tool for assessing time lags between soil or vegetation derived organic matter and aquatic organic matter in lake sediments.

11. Outlook for the application of organic molecular proxies in Quaternary lacustrine studies

Molecular organic proxies are becoming increasingly popular for both marine and lacustrine investigations as they provide detailed (paleo)environmental information. A number of recent developments are especially exciting for Quaternary lacustrine studies, such as organic geochemical proxies for reconstructing continental temperature (lacustrine alkenones, TEX₈₆, MBT/CBT), hydrology (δD) and CSRA as a tool for improving sediment chronologies. Many of these molecular organic proxies are relatively new and there are currently limitations and outstanding questions associated with each proxy that remain to be answered. Nevertheless, there have been major developments and advances over recent years and currently much research is focused on further development and testing of these proxies. Clearly, the application of organic geochemical proxies to Quaternary lacustrine studies is an exciting and developing area of research, and as more studies are done to further refine these methods for use in lacustrine settings, they undoubtedly will be applied to an increasing number of studies.

In this review we discussed biomarkers that can be traced to a specific organism, group of organisms or process. However, there is still a need to identify both more organism and process-specific biomarkers in lacustrine environments. This can be achieved by large-scale surveys of lacustrine environments for common biomarkers, or trends in biomarker ratios, which may be related to environmental parameters (e.g. core top studies of TEX₈₆) assuming that the physical and chemical properties of the system being studied are also well-constrained. It is even possible that there may be biomarkers or organic geochemical proxies that can be specifically developed for use in lacustrine systems, which may or may not be applicable to marine systems. There is certainly a need to identify the source organisms of lacustrine biomarker lipids that have been widely identified in environmental samples but where the producer remains unknown (e.g. the branched GDGTs). This may be achieved by combining lipid studies with genetic and proteomic studies. Conversely, in some cases the organism

producing a particular lipid has been identified yet there is a need to learn more about its ecology in order to improve paleoenvironmental interpretations (e.g. relatively little is known about the ecology of Thaumarchaeota, which produce isoprenoid GDGTs). Furthermore, there is currently a lack of detailed screening of biomarker lipids in important freshwater microbes. Such studies would shed light on which lipids could be used as biomarkers of specific organisms or processes, and may also identify new biomarkers. Finally, confidence in paleoenvironmental reconstructions based on organic geochemical proxies can be gained through multi-proxy and interdisciplinary investigations (e.g. by combining organic geochemical studies with inorganic proxies or molecular biological techniques), which are essential for both accurate paleoenvironmental reconstructions and for providing information on additional factors and processes that may influence proxy records.

A challenge facing paleolimnologists is that the information gained from certain proxies will differ from location to location and some proxies may not be applicable to all locations. This contrasts with paleoceanographic applications of organic proxies where, to a certain extent, conditions are more homogenous and proxies are more widely applicable. For example, lacustrine alkenone distributions may be related to temperature at some locations and to salinity at others. In some lakes the BIT Index may reflect soil organic matter input whereas in others it may mainly reflect *in-situ* production. At some sites TEX₈₆ may provide a robust tool for reconstructing past lake surface temperatures but at other sites the proxy may not work due to e.g. high soil organic matter input. Thus, in contrast to paleoceanographic proxies, paleolimnological proxies may not be generally widely applicable. Nevertheless, organic geochemical proxies can provide powerful tools for examining and reconstructing lacustrine environments but it is essential to be aware of limitations and to understand when and where they should, or should not, be used.

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