

Lipid Markers for Marine Organic Matter

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Abstract An astonishing variety of different lipids have been found in marine sediments and the water column attesting to the diversity of biosynthetic pathways employed by aquatic organisms. Many of these compounds have distinctive structures allowing them to be used as biomarkers for particular sources of organic matter in marine ecosystems. Microalgae synthesize many unusual compounds, such as long-chain alkenones, alkenoates and alkenes, long-chain alkyl diols, highly branched isoprenoid alkenes as well as distinctive sterols and unsaturated fatty acids, thus enabling inputs of microalgal organic matter to be easily recognized. The input of terrestrial organic matter to marine environments can be recognised from lipids of higher plant origin, such as long-chain alcohols, alkanes and fatty acids, and C₂₉ sterols, although marine sources for some of these compounds are now recognised. Bacteria synthesize a diverse range of compounds, such as branched fatty acids, hopanoids and isoprenoids, many of which are particularly stable, for instance those that contain an ether bond. Qualitative assignments of organic matter sources are thus reasonably straightforward, although even now lipids can be found for which no source is known. However, quantitative assessments are more difficult since lipid contents vary greatly between different organisms and lipids display a wide range of reactivities. The combination of lipid biomarker data with information from stable isotopes can provide good estimates of organic matter sources, provided that the isotope signatures of the contributing sources (end-members) are known. This chapter provides a review of biomarkers commonly found in sediments together with notes on their identification and source specificity.

Keywords Biomarkers · Lipids · GC-MS · Phytoplankton · Higher plants

1

Introduction

The lipid constituents of seawater and marine sediments have been the subject of numerous investigations over several decades. Some of the first work described the use of branched alkanes to identify algal contributions to sediments [1]. The lipids in phytoplankton and zooplankton communities were studied by Jeffries [2], who showed how specific fatty acids might be useful in food-web studies. Later work examined the occurrence of lipids in the surface microlayer of the oceans [3]. From these early beginnings a large body of research has developed – variously called environmental organic geochemistry or marine organic chemistry or marine lipid chemistry – that has demonstrated the value of studying individual compounds (biomarkers) as well as the bulk organic matter in order to understand carbon sources and cycling.

Organic geochemists have discovered many new and unusual lipid components during their studies of Recent and ancient sediments. Information ob-

tained from the distribution of components within a variety of lipid classes in such samples has led to searches for the same compounds in possible source organisms, in particular microalgae and bacteria. Indeed, the identification of many compounds in sediments occurred some years before a biological source was recognized [4]. This chapter provides an overview of the types of lipids found in marine ecosystems, and information on how to identify those lipids that are commonly found, together with some notes on their significance. Reference is made to review articles and in particular some early papers to illustrate how the field has developed.

2

Lipid Extraction

2.1

Extractable Lipids

Most studies of lipids in sediments and marine organisms use the Bligh and Dyer method of extraction based on mixtures of chloroform, methanol and water [5], or some modification of it such as the additional use of sonication to liberate more tightly bound lipids. Also used is the earlier Folch extraction method [6]. Recently, these two methods have been compared. In animal tissues containing < 2% lipids both methods give similar yields, but at higher lipid contents the Bligh and Dyer method appears to be less effective than the Folch method [7]. Some geochemical studies just use a hydrocarbon solvent such as hexane, but it should be noted that this is only suitable for extraction of hydrocarbons and even here the solubility of long-chain alkanes in hexane is quite low and this could lead to a biased distribution. Chlorinated solvents such as methylene chloride (DCM) and chloroform provide good extraction of neutral lipids, but are not suitable for the extraction of complex lipids such as phospholipids where recoveries can be as low as 60–65% [8].

The extracts obtained by solvent extraction need to be further fractionated in order to provide fractions with simpler distributions that are more amenable to analysis by gas chromatography-mass spectrometry (GC-MS) or high performance liquid chromatography. Saponification is often used to separate fatty acids from “neutral” components such as hydrocarbons, sterols, alcohols and the like. This process breaks down polar lipids such as triacylglycerols and phospholipids into their constituent fatty acids, plus glycerol, phosphate and sugar groups. Intact polar lipids can be analyzed by HPLC (see later) or by Iatroscan TLC-FID [9, 10].

Fatty acids are converted into methyl esters for GC-MS analysis: diazomethane is often used although the need to prepare the reagent each time is seen as a disadvantage. Dry methanol containing HCl gives good results, but the use of BF_3 in methanol can lead to artifacts [11]. Extracts are often further

fractionated into compound classes either by column chromatography, thin-layer chromatography or in a few cases by HPLC. The book by Christie [12] provides a good overview of methodologies available for lipid analysis.

2.2

Free and Bound Lipids

A few researchers have examined both the lipids that are readily extracted into organic solvents (usually designated as free or unbound lipids) and those that are bound more tightly to the sediment matrix (bound lipids). Several approaches have been employed, but most treat the extracted sediments with base and acid (Fig. 1) to liberate esterified (OH^- -labile) compounds and amide-linked (H^+ -labile) compounds [13–15], while a few studies have further examined the remaining residue by pyrolysis or chemolysis with tetramethylammonium hydroxide (TMAH) to liberate ether- or polymer-bound lipids [15]. It is commonly observed that the distributions of lipids in each fraction show some significant differences, and that the bound lipids usu-

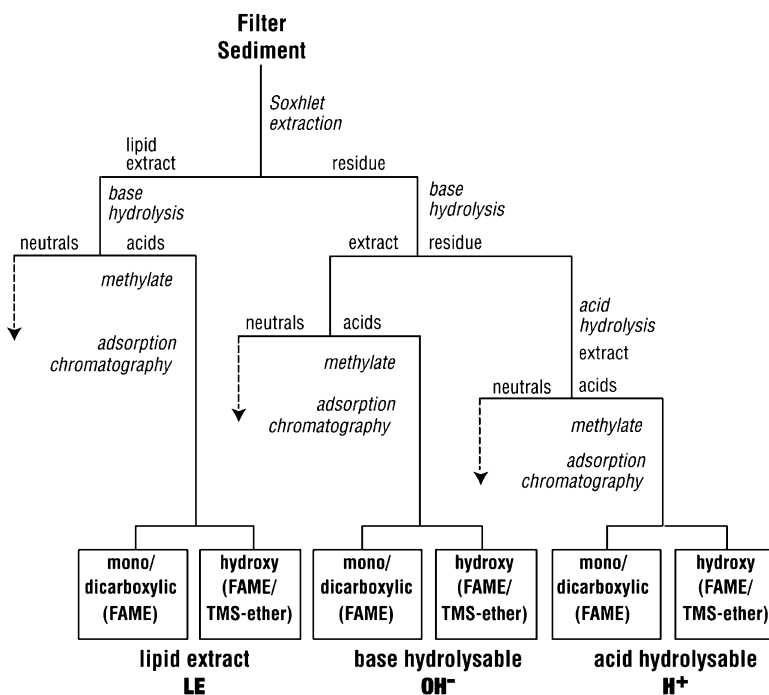


Fig. 1 Lipid analysis scheme from Wakeham [14] which provides fractions contain OH^- labile (i.e. ester-linked) and H^+ -labile (i.e. amide-linked) compounds. Many variants on this scheme exist (e.g. sonication is used instead of soxhlet extraction) and so direct comparisons of data between studies can be difficult

ally contain a higher proportion of labile lipids presumably indicating that adsorption to sediments protects the lipids to some extent from microbial attack. Such studies are much more time consuming than those that involve analyses of only the extractable lipids, but they do reveal much more about the chemical forms in which lipids exist in the sample.

3

Identification of Biomarkers in Sediments and Seawater

3.1

Hydrocarbons

Hydrocarbons are those compounds that contain only carbon and hydrogen. This simple description conceals the great variety of chemical structures that can be found in marine organisms and the multitude of structures formed by degradation of functionalized lipids. For further information, the reader is referred to recent books [16–20].

3.1.1

Straight-Chain Alkanes

Alkanes isolated from marine environments typically fall into two categories. Those with odd-chains such as n -C₁₅, n -C₁₇ and n -C₁₉ are indicative of algal inputs [1] and are often accompanied in higher abundance by the corresponding alkenes. Long-chain (n -C₂₀–C₃₅₊) alkanes that display a strong predominance of odd-chain lengths indicates a contribution from terrestrial plants. Alkanes from petroleum show little or no predominance of either odd- or even-chain lengths [21]. Erosion of ancient sediments can also be a source of similar n -alkane distributions in some aquatic sedimentary environments [22]. Hydrocarbons from eroded sediments often display distinctive sterane and hopane distributions [23].

n -Alkanes give characteristic mass spectra showing a monotonic series of C_{*n*}H_{2*n*+1} ions which decrease in abundance with increasing m/z value. Molecular ions are usually obvious, as is the M⁺-15 ion. These distributions are usually visualized using m/z 57, 71, or 85 mass fragmentograms, but these cannot be used directly for quantification since the proportion of these ions in the total ion current decreases with chain-length.

3.1.2

Branched Acyclic Alkanes (Including Isoprenoids)

Simple branched alkanes such as 7- and 8-methylheptadecane are found in many species of cyanobacteria [24], and in algal mats and lagoonal sedi-

ments. Series of longer mid-chain monomethyl alkanes have been identified in Precambrian oils and kerogens [25], testifying to the early evolution of these compounds. Some cyanobacteria contain more complex distributions of mono-, di- and trisubstituted methyl alkanes [26–28], but these are only infrequently encountered in sediments. Long-chain *iso*- and *anteiso*-branched alkanes found in some higher plants [29], are rarely seen in sediments. An exception is the report of abundant C_{20} – C_{30} *anteiso*-alkanes in some Antarctic rocks, but these are derived from cryptoendolithic microbial communities composed of microalgae, cyanobacteria, black and colorless fungi and heterotrophic bacteria [30].

The C_{19} isoprenoid alkane pristane is common in marine samples, reflecting its abundance in some zooplankton species [31]. The C_{20} isoprenoid phytane can also be found, either reflecting petroleum inputs or a contribution from Archaeobacteria [32]. Two isomeric C_{25} isoprenoid alkanes are commonly encountered: 2,6,10,14,18- and 2,6,10,15,19-pentamethylcosane in sediments (Fig. 2). The former is known from halophilic bacteria [33], while the latter is usually considered to be a biomarker for methanogenic bacteria [34]. However, the finding of exceptionally light $\delta^{13}C$ values for this C_{25} isoprenoid and the C_{20} isoprenoid crocetane (2,5,10,14-tetramethylhexadecane) in reducing sediments of the Hydrate Ridge of the Cascadia continental margin implies a methanotrophic source, at least in this environment [35], while other work suggests the possibility of algal sources [36].

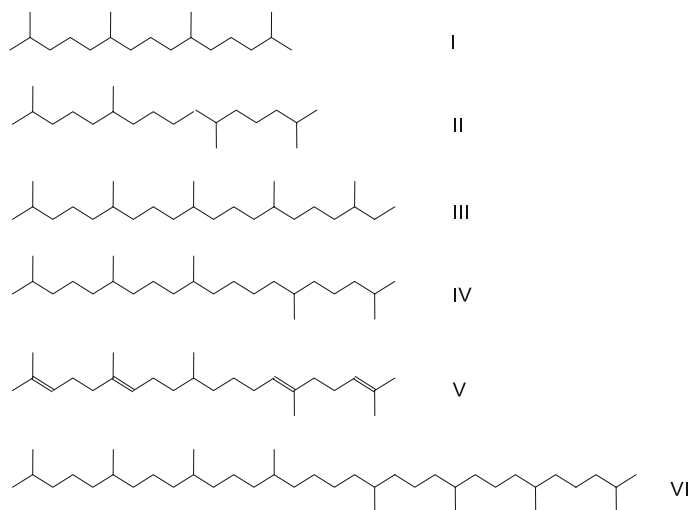


Fig. 2 Structures of some of the isoprenoid hydrocarbons found in sediments and seawater. I: pristane; II: crocetane; III: 2,6,10,14,18-pentamethylcosane; IV: 2,6,10,15,19-pentamethylcosane (PME: mainly methanogenic bacteria); V: 2,6,10,15,19-pentamethyl-icos-2,6,14,18-tetraene (methanogens); VI: lycopane

Polyunsaturated alkenes based on the 2,6,10,15,19-pentamethylcosane skeleton have been identified in methanogenic archaea [37]. For a review of the lipids in archaeobacteria, see the comprehensive compilation of De Rosa and Gambacorta [38].

Another isoprenoid that is ubiquitous in sediments is lycopane [39]. It seems to be particularly abundant in sediments that were deposited under anoxic conditions, suggesting that the lycopane/ C_{31} *n*-alkane ratio could be used as a proxy for oxic conditions [39]. The origin of lycopane is still unknown, but it has been suggested that a marine photoautotroph may be the source [39, 40].

Branched alkanes produce similar spectra to those of *n*-alkanes, but the ion due to α -cleavage is more intense so the position of branching is usually easy to identify [30]. More highly branched alkanes can also be identified from their mass spectra [26–28] (and references therein), but possible problems with co-elution must be taken into account. For example, Han et al. [24] showed that the “7,9-dimethylhexadecane” previously identified in some cyanobacteria was in fact a mixture of 7- and 8-methylheptadecane. In such cases, retention indices can be very helpful to distinguish between possible isomers [26–28, 41]. Isoprenoid alkanes are generally identified using a m/z 183 or 113 mass fragmentogram, since these ions are enhanced in abundance compared with *n*-alkanes. The C_{40} isoprenoid lycopane co-elutes with the *n*- C_{35} alkane on most non-polar capillary columns [39], but can be recognized by enhanced ions due to cleavage next to the methyl groups to give characteristic ions at m/z 477, 406/7, 336/7, 308/9 and 183.

3.1.3

n-Alkenes

A variety of unsaturated hydrocarbons (alkenes) are found in marine samples. The most common is the hexa-unsaturated alkene *n*- $C_{21:6}$, which is produced by many species of microalgae by decarboxylation of the 22:6(*n*-3) fatty acid [42]. It is particularly abundant in diatoms, which explains why these species rarely have significant amounts of the 22:6 PUFA. Conversely it is a relatively minor component in dinoflagellates which have abundant 22:6 fatty acid. It is not found in cyanobacteria or chlorophytes that lack longer-chain PUFA. In diatoms and dinoflagellates this alkene can be accompanied by the 21:5 and 21:4 alkenes [43]. The presence of such highly labile alkenes in a sediment usually points to the presence of intact (perhaps living) algal cells.

Microalgae and some prokaryotes are likely sources of long-chain alkenes in marine environments [44]. These include, *inter alia*, diunsaturated C_{31} , tri- and tetraunsaturated C_{33} alkenes (with *cis*-double bonds), in some haptophytes (e.g. [45–47]) and di- and tri-unsaturated C_{37} and C_{38} alkenes (with *trans*-double bonds) which occur in the haptophyte *Emiliania huxleyi* [45]. These alkenes are found in marine sediments [48] and fauna such as filter-

feeding mussels [49]. Saturated and polyunsaturated $C_{14} - C_{31}$ hydrocarbons with a strong odd-over-even carbon predominance have been isolated from two marine *Nannochloropsis* species [50]. A North Atlantic strain of *Rhizosolenia setigera* biosynthesizes a C_{25} HBI alkene (see below) as well as C_{25} and C_{27} *n*-polyenes with six or seven double bonds [51]. These alkenes may be the precursors for C_{25} and C_{27} 2-*n*-alkylthiophenes found in sediments from palaeo upwelling regions [51]. Prokaryotes are also potential sources, although their contributions are more obvious in specialized environments such as algal mats. For example, $C_{19} - C_{29}$ alkenes occur in the cyanobacterium *Anacystis montana* [52], and green photosynthetic bacteria of the genus *Chloroflexus* contain a $C_{31:3}$ alkene all-*cis* hentriaconta-9,15,22-triene [53].

Under electron impact conditions, alkenes give mass spectra with prominent molecular ions (provided that the number of double bonds is 4 or less) and major ions at C_nH_{2n-1} . It is usually not possible to assign the position or geometry of the double bond, but derivatization techniques such as formation of DMDS adducts [53] can help identify double bond positions, although mass spectra can be difficult to interpret where the number of double bonds is greater than 3.

3.1.4

Highly Branched Isoprenoid (HBI) Alkenes

An unusual class of highly branched polyunsaturated alkenes (now termed HBI alkenes) have been recognized in many studies of marine sediments; Gearing et al. [54] seems to be the first report. HBIs have an unusual coupling of C_5 isoprene units producing a "T" shaped molecule and typically have 2–4 double bonds (for C_{25} alkenes; Fig. 3) or 4–6 double bonds (for C_{30} alkenes). A C_{35} HBI with 7 double bonds has also been observed in sediments from the Arabian Sea [55]. The parent C_{20} alkane was first noted in the Rozel Point crude oil [56], but in modern marine environments it is much more common to find C_{25} and C_{30} chain-lengths, with saturated and monounsaturated C_{20} hydrocarbons much less frequent [57]. A diatom origin was suggested by the high abundance of C_{25} HBIs in sediments and seawater from the Peru upwelling [58] and the isolation of a diunsaturated C_{25} HBI from sea-ice communities [59]. This was confirmed when high contents of C_{25} highly branched isoprenoid alkenes were identified in laboratory cultures of the diatom *Haslea ostrearia* and C_{30} HBI alkenes were found in *Rhizosolenia setigera* [43]. Since then the C_{25} and C_{30} HBI alkenes have been found in several species of *Haslea*, *Navicula*, *Pleurosigma* and *Rhizosolenia* (e.g. [51, 60–63]).

Genetic differences between strains, environmental factors and growth stage appear to be important determinants of relative abundances of the various homologues identified [64]. Recently, HBI-related monocyclic compounds containing a single cyclohexane ring (Fig. 3) have also been identified

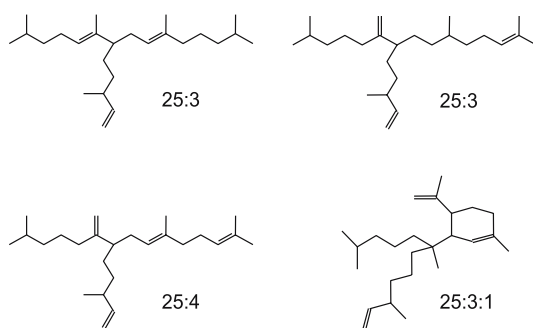


Fig. 3 Structures of a few of the many highly branched isoprenoid alkenes identified in sediments. Note that the 25:3:1 nomenclature here refers to 3 double bonds and 1 ring. The latter compound is difficult to hydrogenate and thus has been confused as a diene with 2 rings. Massé et al. [66] give full details of structures and retention times

in *Rhizosolenia setigera* [65]. Even though these are relatively minor constituents they can be readily recognized in the hydrocarbon fraction after hydrogenation. Massé et al. [66] have shown that these monocyclic alkenes also occur in a variety of sedimentary environments along with the corresponding HBI alkenes.

An extensive literature is now available to identify HBI alkenes in marine samples. Retention data are a good guide to which particular isomer is present [43, 60]. Mass spectra have been reported for the parent alkanes [60] and for most of the alkene isomers [51, 67, 68]. The paper by Massé et al. [66] provides a very good summary of RRT data and structures for HBI alkenes and their cyclohexyl analogs. NMR data for some of the hydrocarbons isolated in nearly pure form are also available through the work of Rowland and co-workers [61, 62, 69, 70].

3.1.5

Other Branched Alkenes

Another class of unusual isoprenoid alkenes are the botryococcenes formed by green microalgae of the genus *Botryococcus* [71–73]. Some strains of this alga also synthesize unusual lycopadienes [74]. *Botryococcus* is found in terrestrial freshwater aquatic ecosystems, and so will not be discussed further here. However, it should be noted that compounds thought to be derived from *Botryococcus* have been detected in ancient hypersaline environments [75], so a marine source for this compound type cannot be entirely discounted.

C₃₂ – C₃₆ polymethylene isoprenoid-like branched alkenes have been found in Black Sea sediments and attributed to an unknown photoautotroph source [76]. Their structures suggest that they were biosynthesized by methylation of an unsaturated *n*-C₃₁ precursor at specific positions. The mass

spectra of the corresponding alkanes show characteristic ion pairs due to cleavage α to the methyl group.

3.1.6

Polycyclic Alkenes and Petroleum Biomarkers

In seawater and Recent sediments one usually finds a mixture of steroidal alkenes (ster-2-enes, ster-4-enes and ster-5-enes) as degradation products of sterol and/or steroidal ketone precursors [77–79]. The presence of fully saturated steranes is indicative of thermally altered organic matter and/or petroleum. Petroleum hydrocarbons are ubiquitous in the marine environment: they originate from many sources such as shipping accidents, ships' discharge, urban and industrial effluents and natural oil seeps [21]. Sterane distributions are characterized from m/z 217 and 218 mass fragmentograms, hopanes from m/z 177, 191 and 205 mass fragmentograms and diasteranes from m/z 259 mass fragmentograms. Aromatic steroidal hydrocarbons with 1 or 3 aromatic rings are also useful compounds for fingerprinting petroleum samples.

3.2

Fatty Acids

3.2.1

Monocarboxylic Fatty Acids

Straight-chain fatty acids are often the most abundant lipids found in seawater particulate matter and in Recent marine sediments. Sources include bacteria, microalgae, higher plants and marine fauna; each of which has a distinctive fatty acid profile so that sources can usually be assigned. However, some fatty acids such as palmitic and stearic acids (16:0 and 18:0 respectively) are ubiquitous. Terrestrial plants are usually considered the main source of long-chain (C_{20} – C_{30}) fatty acids, showing a strong even over odd predominance, but some microalgae and/or bacteria are also suspected to be a minor source of these acids [80].

Bacteria are the major source of *iso*-, *anteiso*-, cyclopropyl and mid-chain branched fatty acids in marine ecosystems [81], but it is often overlooked that bacteria can also be a significant source of palmitoleic (16:1($n-7$)) and palmitic (16:0) acids in addition to the more commonly used biomarker *cis*-vaccenic acid (18:1($n-7$)) [82–84]. A typical example of a fatty acid distribution found in a modern marine sediment is shown in Fig. 4.

A fascinating recent finding is the identification in planctomycete-like bacteria capable of anaerobic ammonium oxidation of highly unusual fatty acids and related compounds containing concatenated cyclobutane rings [85]. These lipids can contain up to 5 linearly fused cyclobutane rings joined by

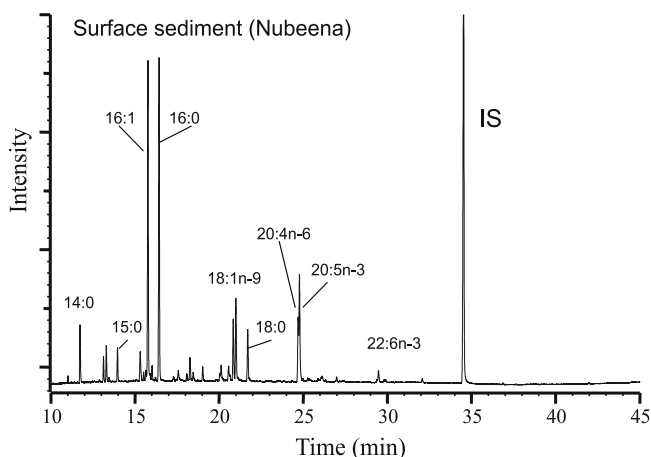


Fig. 4 Capillary gas chromatogram of total fatty acids (as methyl esters) in a surface sediment from coastal Tasmania (Nubeena), Australia. The predominance of 16:0, 16:1 and 20:5 is typical of sediments where diatoms are a major source of organic matter

cis-junctions, hence giving rise to the common name of ladderanes for these compounds [85]. These anammox bacteria contain a unique organelle called the anammoxasome in their cytoplasm [86], and it is postulated that the close packing of the ladderane lipids in the cell membrane reduces its permeability to toxic intermediates (hydrazine and hydroxylamine) produced in the anammox reaction [85]. Examples of this lipid type include fatty acid methyl esters, *sn*-2-alkyl glycerol monoethers, alcohols, *sn*-1,2-dialkyl glycerol diethers and *sn*-2-O-alkyl, *sn*-1-acyl glycerol [86]. Illustrative structures are shown in Fig. 5. Recent field work has established that coupled nitrite reduction and ammonia oxidation can be an important source of dinitrogen gas production in many aquatic systems including the Black Sea [87].

C₁₈ – C₂₂ polyunsaturated fatty acids (PUFA) are found in almost all marine organisms, often in amounts comparable to the saturated fatty acids. Their primary source is from the phytoplankton. Marine animals obtain most of their PUFA from the food-web and further modify the distributions by chain-elongation and desaturation of the ingested fatty acids. Many studies have used fatty acid distributions to identify food items [88]; one interesting example used fatty acids in benthic foraminifera to show that these animals selectively fed on settled phytodetritus to secure their needs for PUFA [89].

The PUFA composition of microalgae has been well studied [80] and a comprehensive review of fatty acids in marine organisms has recently been published [90]. Diatoms and eustigmatophytes are rich in 20:5(*n*-3) (EPA) and produce small amounts of arachidonic acid, 20:4(*n*-6) (AA) with negligible amounts of 22:6(*n*-3) (DHA). Diatoms synthesize unusual C₁₆ PUFA such as 16:4(*n*-1) and 16:3(*n*-4). In contrast, dinoflagellates have high con-

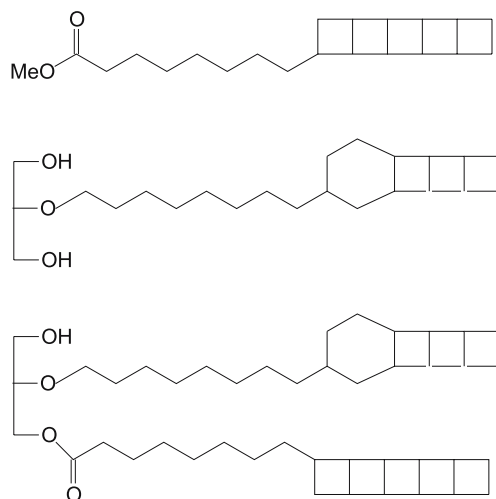


Fig. 5 Structures of ladderane lipids found in planctomycete-like anammox bacterium *Candidatus B. anammoxidans*. Modified from [86]

centrations of DHA and moderate to high proportions of EPA and precursor C_{18} PUFA [18:5($n-3$) and 18:4($n-3$)]. Haptophytes also contain EPA and DHA with EPA the dominant PUFA. Cryptomonads are a rich source of the C_{18} PUFA 18:3($n-3$) (ALA α -linolenic acid) and 18:4($n-3$) (stearidonic acid), as well as EPA and DHA. Chlorophytes (green microalgae) typically contain low concentrations of C_{20} and C_{22} PUFA, but have abundant 18:3($n-3$) and 18:2($n-6$), and are also able to make 16:4($n-3$). Two very-long-chain highly unsaturated (C_{28}) VLC-HUFA: octacosaeptaenoic acid [28:7($n-6$) (4,7,10,13,16,19,22)] and octacosaoctaenoic acid [28:8($n-3$) (4,7,10,13,16,19,22,25)] were discovered in seven marine dinoflagellates [91]. More recent work has shown that the glycolipid fraction of the dinoflagellate *Karenia brevis* contains 18:5($n-3$), while the phospholipid fraction contained small amounts of both 28:8($n-3$) and 28:7($n-6$) [92, 93].

Saturated fatty acid methyl esters (FAME) give characteristic mass spectra showing a base peak at m/z 74 and important ions at m/z 87, 143, M^+ , M^+-29 , M^+-31 and M^+-43 . *Iso*- and *anteiso*-branched acids also show weak ions due to α -cleavage at the branching point. Thus, *anteiso*-fatty acids show an enhanced M^+-29 ion [30]. Monounsaturated FAME show a reduced abundance for the m/z 74 and 87 ions and an enhanced m/z 55 ion. The molecular ion is readily discerned as is the M^+-32 ion. The geometry of the double bond has minimal effect on the mass spectrum, but E (*trans*) and Z (*cis*) isomers can be identified from their relative retention times (E-isomers elute just after Z-isomers on non-polar columns) [94]. Much information is available from the human nutrition and clinical literature [94, 95] due to the health concerns about *trans*-fatty acids in margarines. Double bond positions can be

confirmed by mass spectrometry of dimethyl disulphide (DMDS) adducts as described by [95–99]. Mass spectra of the adducts show ions due to cleavage between the two SCH₃ groups. E and Z isomers can be distinguished, since the erythro isomer (originally E) elutes after the threo isomer (originally Z).

Polyunsaturated fatty acids may be more difficult to identify since they can exist as different positional isomers. It is often helpful to analyse these on two capillary columns differing in polarity. Extensive sets of relative retention times (RRTs) are available [98]. A useful rule-of-thumb is that on non-polar columns, PUFA elute as pairs if they differ by one double bond and the position of the last double bond differs by three carbon atoms. Thus 20:4(*n*-6) elutes just before 20:5(*n*-3), 16:2(*n*-4) elutes with 16:1(*n*-7), 18:2(*n*-6) elutes just before 18:3(*n*-3) etc. Mass spectra are less useful for positive identification, since the intensity of the molecular ion decreases with increasing degree of unsaturation and may not be observed in FAME having 4 or more double bonds (although this is somewhat instrument dependent). The base peak is usually *m/z* 79.

Other useful techniques are argentation TLC, column chromatography or HPLC [99, 100]. A typical TLC procedure involves double development in hexane-Et₂O-HOAc (94 : 4 : 2) on silica gel (7 g) plates loaded with 3% AgNO₃ (w/w). Bands can be visualized under 366 nm UV light after spraying with 2',7'-dichlorofluorescein, and extracted into hexane-CHCl₃ (4:1, ×3). The FAME extract needs to be washed with saturated NaCl and 2 M NH₃ to remove Ag⁺ and 2',7'-dichlorofluorescein, respectively, before further analysis.

A range of derivatization methods are available for PUFA analysis, although their utilization by marine chemists and organic geochemists is still infrequent. A detailed account of the methods available is given by Christie [101]. Double bonds can be located by EI GC-MS of picolinyl esters or 4,4-dimethyloxazoline (DMOX) derivatives of the PUFA. Both techniques can be useful and should be considered complementary [101], although the picolinyl esters have long retention times and thus DMOX derivatives are often preferred. A general rule has been formulated to interpret the mass spectra of DMOX derivatives. It states that: if an interval of 12 Dalton, instead of the regular 14 Dalton, is observed between the most intense peaks of clusters of fragments containing *n* and *n* - 1 carbon atoms in the mass spectrum, then the double bond occurs between carbons *n* and *n* + 1 in the molecule (reviewed in [102]). Examples of DMOX mass spectra are given by Christie [101] and Spitzer [102].

A recent example of the application of these techniques is provided by Méjanelle et al. [103]. These authors used a combination of DMOX and DMDS derivatizations to identify the FAME in marine flagellates (smaller than 10 μm) that colonize marine particles in deep Atlantic waters. Among the compounds identified was a novel non-methylene-interrupted fatty acid 20 : 3Δ^{7,13,17}, which occurred in quite high amounts.

Nichols and Davies [104] report a highly sensitive method based on analysis of 2-oxo-phenylethyl esters by high-performance liquid chromatography-

mass spectrometry (LC-MS) combined with ultra violet (UV) detection. The technique was applied to identify PUFA in *Shewanella* bacteria. It is now recognized that some bacteria, including deep-sea and coastal species, can synthesize PUFA such as 20:5($n-3$) and thus can be a source of PUFA in particular sedimentary environments (e.g. [105]).

3.2.2

α,ω -Dicarboxylic Fatty Acids

C₁₆ and C₁₈ α,ω -dicarboxylic fatty acids are major constituents of the higher plant polyester cutin and C₁₆ – C₂₂ even-chain dicarboxylic fatty acids are often abundant in the plant biopolymer suberin [106–108]. Seagrasses also contain α,ω -dicarboxylic fatty acids, so most sediments containing higher-plant organic matter can be expected to contain these long-chain dicarboxylic acids (e.g. [109]). It is usually necessary to hydrolyze the sediments to release these esterified fatty acids [14].

Shorter-chain dicarboxylic fatty acids such as azelaic acid (C₉) occur as natural constituents of some plants and as degradation products from oxidative scission of the double bonds in unsaturated fatty acids. Azelaic and other short-chain dicarboxylic acids have been found in marine sediments and in aerosols [110]. The presence of long-chain α,ω -dicarboxylic fatty acids in Black Sea sediments in which shorter-chain α,ω -dicarboxylic fatty acids were absent suggested to Wakeham [14] that they were derived from plant matter rather than oxidation of hydroxyl fatty acids in the sediments.

3.2.3

Monohydroxy Monocarboxylic Fatty Acids

Hydroxylated fatty acids have been found in many sediments (see [44] for a review), but most reports are from lacustrine settings. Aliphatic α - and β -monohydroxy fatty acids (i.e. 2- and 3-hydroxy monocarboxylic acids) occur in a wide range of organisms [111] and are typically produced as intermediates in the α - and β -oxidation of monocarboxylic fatty acids. β -Oxidation occurs more widely than α -oxidation, although the latter is known in plants, animals and bacteria. In yeasts, α -hydroxy fatty acids are intermediates in fatty acid biosynthesis [112]. Long-chain hydroxy fatty acids are rarely reported in microalgae, although C₂₂ to C₂₆ saturated and monounsaturated α -hydroxy fatty acids have also been found as major lipid components of the cell wall of three marine chlorophytes [50] and a series of saturated α - and β -hydroxy acids ranging from C₂₄ to C₃₀ with C₂₈ predominating was detected in some freshwater eustigmatophytes [113].

Shorter-chain (C₁₂ – C₁₈) β -hydroxy fatty acids are commonly found in Gram-negative bacteria as amide-bound constituents of the lipid A component of the cell wall polysaccharide [114, 115]. The C₁₄ β -hydroxy fatty

acid usually predominates, but C₁₅ and C₁₇ *iso*- and *anteiso*-branched β -hydroxy fatty acids predominate in the common marine bacterium *Desulfovibrio desulfuricans* [116]. These biomarkers can be used to characterize microbial communities [116], and they provide a useful complement to phospholipid fatty acid analysis [117].

C₃₀ – C₃₄ mid-chain hydroxy fatty acids were identified in several marine eustigmatophytes of the genus *Nannochloropsis* [118]. The predominant positional isomer contained an hydroxy group at the ω 18 position, suggesting that the series was produced by chain-shortening or elongation of a single major precursor.

C₁₆ and C₁₈ ω -hydroxy fatty acids are common constituents in plant cutin while C₁₆ – C₂₂ ω -hydroxy fatty acids are found in suberin [119]. Long-chain ω -hydroxy fatty acids and α -hydroxy fatty acids also occur in the seagrass *Zostera* [109]. (ω – 1)-Hydroxy long-chain fatty acids have been identified in some marine and lacustrine environments, but their origin is still debated [116, 120]. Methane-utilizing bacteria are a possible source of C₂₆, C₂₈ and C₃₀ (ω – 1)-hydroxy fatty acids [121], and a C₂₆ (ω – 1) hydroxy acid has also been found in the two cyanobacteria, *Anabaena cylindrica* [122] and *Aphanizomenon flos-aquae* [123].

Hydroxy fatty acids (as methyl esters) can be separated by silica gel TLC into different categories according to the number and position of the hydroxyl groups. These fractions are then derivatized with BSTFA (N,O-*bis*-(trimethylsilyl)-trifluoroacetamide) to convert the free OH group(s) into the TMSi-ether. α - and β -hydroxy acids coelute even on the high resolution columns [106, 113], but they can be identified from their characteristic mass spectra. Mass spectra of β -hydroxy acids (as methyl esters, TMSi-ethers) show major ions at m/z 175 [(CH₃)₃SiO = CHCH₂CO₂CH₃] and M⁺-15 (base peak) with minor ions at m/z 103, 89, 133, 159 and M⁺-31 [106]. The mass spectra of α -hydroxy fatty acids (as methyl esters, TMSi-ethers) show major ions at m/z 73 (base peak), M⁺-15 and M⁺-59 with minor ions at m/z 89, 103, 129, 159 and M⁺-43 [106]. α - and β -hydroxy fatty acids in soils, sediments and biofilms have also been identified as the methyl ester with the OH group underivatized [117].

Quantification of α - and β -hydroxy fatty acids can be problematic since they co-elute. Volkman et al. [113] estimated the contribution of α -hydroxy acids to each peak in the hydroxy fatty acids of some freshwater eustigmatophytes from the proportion of the M⁺-59 ion in the background-subtracted mass spectrum using the fact that this was about 11.5% of the total ions in the mass spectra of peaks containing only α -hydroxy acids. This value was determined from the mass spectrum of the C₂₆ saturated α -hydroxy fatty acid. Note that this value may vary under different GC-MS conditions and there will be a small change in the abundance of this ion with increasing chain length.

3.2.4

Polyhydroxy Monocarboxylic Fatty Acids

C₁₆ – C₂₂ polyhydroxy fatty acids are common constituents of higher plants where they occur inter-esterified in the cutin and suberin biopolymers. C₁₆ and C₁₈ polyhydroxy fatty acids are common constituents in plant cutin while C₁₆ – C₂₂ fatty acids are found in suberin. In cutin, the most abundant ω -hydroxy fatty acid is usually 16-hydroxyhexadecanoic acid, which is formed from palmitic acid by ω -oxidation [124]. This in turn is hydroxylated at the 10- or 9-position to form 10,16- and 9,16-dihydroxy hexadecanoic acids. Oxidation of the ω -hydroxy fatty acid forms the corresponding α,ω -dicarboxylic acid. The C₁₈ fatty acids are usually less abundant than the C₁₆ and here the main precursor is oleic acid [18:1(*n*-9)] giving rise to 18-hydroxyoctadec-9-enoic acid which is further epoxidized to 18-hydroxy-9,10-epoxy octadecanoic acid which in turn is converted to 9,10,18-trihydroxyoctadecanoic acid [107, 125]. These hydroxy fatty acids, C₁₆ – C₂₂ dicarboxylic acids and ω -hydroxy fatty acids can be excellent markers for terrestrial organic matter in sediments [106], especially when combined with compound-specific isotope data [126]. Note, however, that two dihydroxy fatty acids (identified as 15,16-dihydroxy-dotriacontanoic acid and 16,17-dihydroxytritriacontanoic acid) have been found in a marine eustigmatophyte [118], so this general type of biosynthesis is not unique to plants.

Polyhydroxy fatty acids are usually identified as their methyl ester, TMSi-ether. Strong ions occur due to α -cleavage to the hydroxy group. Thus 10,16-dihydroxyhexadecanoic acid (FAME, TMSi-ether) gives a weak molecular ion at *m/z* 446 and major ions at *m/z* M-15, M-31, M-47, *m/z* 373 and 275. The presence of any co-eluting 9,16-dihydroxy isomer is shown by ions at *m/z* 259 and 289.

3.3

Intact Esterified Lipid Classes

The fatty acids in phytoplankton occur in a variety of more complex polar lipids. Neutral lipids such as triacylglycerols (TAG; also referred to as triglycerides in the older literature), and glycolipids such as monogalactosyldiacylglycerols (MGDG), and digalactosyldiacylglycerols (DGDG) are often predominant with lesser amounts of phospholipids (PL) and so-called acetone mobile polar lipids [127]. Wax esters are very uncommon in phytoplankton, although abundant in many zooplankton [128]. Information about the polar lipids in seawater and natural phytoplankton is surprisingly sparse so we are heavily reliant on data published for microalgal species cultured in the laboratory. Often the conditions used are very different from those occurring in the field, and it is well documented that the proportions of lipid classes is very dependent on environmental conditions (e.g. [129] and references therein).

3.3.1

Triacylglycerols

Lipids of microalgae are often rich in polar lipids during logarithmic phase growth, but many species accumulate triacylglycerol during stationary phase when nitrogen is limiting [130]. The fatty acid amount and composition is dependent on both the growth conditions and the physiological state (e.g. [131]). In contrast, there are very few reports of TAG in bacteria or other prokaryotes so TAG in marine samples is almost always due to eukaryotic organisms. Triacylglycerols are also used by many marine animals as an energy reserve [132], but the proportion of PUFA is often quite low compared with either the triacylglycerols in microalgae or the phospholipids of zooplankton [133, 134]. Wax esters (WE; see later) are another lipid store, especially used by some species of zooplankton. The composition of fatty acids in these TAG or WE can provide useful clues about prey species of the animal in question (e.g. [135]).

Triacylglycerols are readily isolated by column chromatography on silica [135]. Their fatty acid composition can be determined after saponification or transesterification. If information on the composition of intact triacylglycerols is required then high temperature GC and GC-MS methods are available [134–137], although loss of the more unsaturated triacylglycerols through adsorption on the GC column can be a problem. An alternative technique is probe MS analysis of the entire TAG fraction isolated by TLC or HPLC. This was used by Boon et al. [138] to demonstrate the presence of intact TAG in a diatomaceous ooze from off-shore Namibia.

3.3.2

Phospholipids

Phospholipids are key components of biological membranes. They contain glycerol (glycerophosphatides) with a least one O-acyl, O-alkyl or O-alkyl-1'-enyl attached to the glycerol plus a polar head group (nitrogenous base, glycerol or inositol). Common examples containing one nitrogenous base include PC (phosphatidylcholine), PE (phosphatidylethanolamine) and PS (phosphatidyl serine), or inositol (PI: phosphatidyl inositol) or two glycerol molecules (PG: phosphatidylglycerol; and DPG: diphosphatidylglycerol or cardiolipin). Related compounds include the sphingosyl phosphatides.

The research group of Professor D.C. White pioneered the use of total phospholipid fatty acids (PLFA) as a measure of microbial biomass in sediments and other environments (e.g. [139–141]). Dobbs and Findlay [141, 142] report a set of values for converting lipid-P concentrations to carbon biomass for bacteria, cyanobacteria, yeasts and some microalgae. Viable microbes have an intact membrane which contains phospholipids and their constituent fatty acids. When these cells die, enzymes hydrolyze the phosphate group within minutes to hours leaving a diacylglycerol (diglyceride: DG) [143]. The

resulting DG has the same signature fatty acids as the phospholipids (until it degrades), so a comparison of the ratio of PLFA to DG provides an indication of the proportions of viable and non-viable microbes.

A development of this approach is to analyze the intact phospholipids directly as a marker for intact viable cells in marine sediments. Rütters et al. [144, 145] have shown that intact phospholipids isolated from a sediment extract can be identified using liquid chromatography-electrospray ionization-mass spectrometry (HPLC-ESI-MS). The combined analysis of phospholipid types and their fatty acid substituents allows various groups of microorganisms living in the sediment to be differentiated. Note that phospholipids and their degradation products (free fatty acids, mono- and diacylglycerols) are rapidly recycled in sediments and so their abundances show a sharp decline with sediment depth [145, 146].

3.3.3

Glycolipids

Glycolipids are important constituents of cell membranes. Common glycolipids include monogalactosyldiacylglycerol MGDG, digalactosyldiacylglycerol DGDG, and sulfoquinovosyldiacylglycerol SQDG (e.g. [128, 147, 148]), but many other more complex glycolipids (i.e. polar molecules containing sugars) exist. New galactolipids based on a sugar galactopyranoside and 16:1 and 20:5 fatty acids have been identified in the diatom *Nitzschia* sp. [149]. Glycolipids can be purified by chromatography on DEAE-cellulose and silica-gel columns [150], since most glycolipids are less polar than most phospholipids [151]. Very few data have been published on glycolipids in marine waters or sediments, reflecting their low concentration due rapid

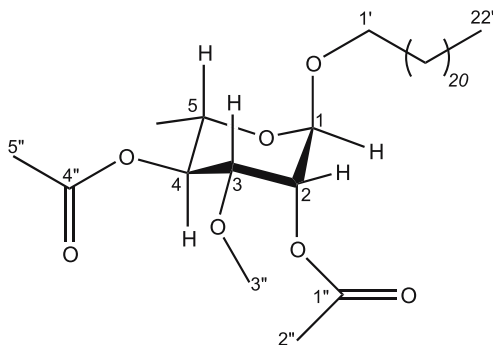


Fig. 6 Structure of the unusual glycolipid found in sediments from Ace Lake Antarctica [55]. The presence of C_{22} and C_{24} carbon chains ether-linked at position 1 of the sugar shows that some common biomarkers (in this case C_{22} and C_{24} alcohols) may be derived from quite unexpected molecules. The source organism of this compound is still unknown

hydrolysis following their liberation from the microbial cell (e.g. [151]). An interesting exception was the isolation of intact novel glycolipids from anoxic sediments in an Antarctic lake [152]. These unusual compounds were identified as docosanyl 3-O-methyl- α -rhamnopyranoside and docosanyl 3-O-methylxylopyranoside (Fig. 6). These compounds were a major and unexpected source of C₂₂ and C₂₄ alcohols in the sediment.

Analysis of the polar lipids in the Dead Sea biomass during a mass bloom of halophilic archaea in 1992 showed one major glycolipid to be present in the extracts, corresponding with the sulfated diglycosyl diether lipid (S-DGD-1) characteristic of the genus *Haloferax* [153]. Other glycolipids indicative of *Halobacterium sodomense* or *Haloarcula marismortui*, or other *Halobacterium* species were not found. These data indicated that the major archaea in the lake was not one of those that had been cultured.

3.3.4

Betaine Lipids

DGCC (1,2-diacylglyceryl-3-(O-carboxyhydroxymethylcholine)) is a betaine lipid which is a common constituent of the Haptophyceae [154]. It also occurs in dinoflagellates and in at least one diatom [154]. DGTA (1,2-diacylglyceryl-O-2'-(hydroxymethyl)-(N,N,N-trimethyl)- β -alanine) was detected in five out of 16 species of Haptophyceae, whereas DGTS (1,2-diacylglyceryl-O-4'-(N,N,N-trimethyl)homoserine) was not detected. There are no reports of betaine lipids in seawater, particulate matter or sediments, but it is doubtful that analysts would have been looking for these compounds.

3.3.5

Wax (Alkyl) Esters

Commonly used chromatographic lipid separation schemes typically provide a fraction containing long-chain wax esters, steryl ethers and steryl esters (e.g. [135, 137]). Although these quite different lipid classes have quite different origins and geochemical significance they are often analyzed together.

Wax esters (or more correctly alkyl esters) are particularly common in the marine environment as they are used as an energy store by many species of zooplankton [133, 155]. Wax esters of zooplankton origin have been identified in sediments deposited under the upwelling cells of Walvis Bay [137], the Peru margin [155], and the Arabian Sea [55]. The fatty acid composition of these wax esters generally resembles that of the phytoplankton diet and contains PUFA such as 20:5(*n*-3) and 22:6(*n*-3) [133]. The alcohols consist mainly of saturated and monounsaturated C₁₆ – C₂₂ moieties. Indeed, elevated levels of 20:1 and 22:1 alcohols in sediments is usually indicative of hydrolysis of marine wax esters. Note that the distributions of wax esters found in lacustrine and terrestrial sediments usually show a much higher proportion of saturated

fatty acids and alcohols extending to much longer chain-lengths (e.g. [156]). Some bacteria also synthesize wax esters [157], but it seems that bacteria are rarely a significant source of these lipids in aquatic environments.

Wax esters can be analyzed on low bleed non-polar capillary columns able to operate up to 350 °C. They give readily interpretable mass spectra [137, 159]. A wax ester of general formula $R_F\text{COOR}_A$ gives a strong molecular ion and cleavage ions that include the fatty acid moiety $[R_F\text{COO} + 2\text{H}]^+$ and the alcohol moiety $[R_A + \text{H}]^+$. Thus within each set of wax esters having the same overall carbon number, the various combinations of alcohols and fatty acids present can be readily identified and quantified from the abundance of the cleavage ions.

3.4

Alicyclic Alcohols

$C_{22} - C_{32}$ even-chain *n*-alkanols are found in most marine sediments and are usually attributed to an input from plant waxes, especially when they are accompanied by *n*-alkanes showing a strong predominance of odd chain-lengths. Marine sources of *n*-alkanols include hydrolysis of the wax esters of zooplankton which gives rise to saturated and unsaturated alcohols from C_{14} to C_{22} . These distributions often show a predominance of 16:0, 20:1, and 22:1 alcohols [128]. A specific, but unidentified, marine source of the 22:0 alcohol has also been suggested (Fig. 6; [44, 152]). An interesting finding is the presence of straight-chain monounsaturated C_{32} and C_{30} alcohols and a diunsaturated C_{32} alcohol in acid hydrolysates of the total extract from marine eustigmatophytes [160]. The relative proportions of the $C_{30} - C_{32}$ alcohols were very similar to those of the corresponding alkyl diols, having the same carbon number but one less double bond. Free alcohols were present only in trace amounts, suggesting that the alcohols are bound to extractable polar lipids by bonds that are resistant to base hydrolysis (perhaps by amide bonds).

Alcohols are usually examined as their TMSi-ethers although some researchers prefer to use acetate derivatives because of their greater stability. The TMSi-ethers give a strong M-15 ion, weak M^+ and typically show a base peak at m/z 75 and a major ion at m/z 103.

3.5

Long-Chain Alkyl Diols

$C_{30} - C_{32}$ alkyl 1,15-diols were first discovered in Black Sea sediments [161], which was soon followed by numerous reports on their occurrence in other sediments (e.g. [162-164]). After initial suggestions that the alkyl diols were derived from cyanobacteria, it was shown that saturated and monounsaturated $C_{30} - C_{34}$ alkyl diols occur in marine eustigmatophytes from the

genus *Nannochloropsis* [160] together with alcohols of the same chain-length (Fig. 7). These compounds are probably building blocks for biopolymers called algaenans found in these microalgae [50]. Algaenans such as this are quite resistant to bacterial degradation and thus they can survive into the sediment record ultimately to become an important source of hydrocarbons in some crude oils [165, 166]. The biosynthesis of these alkyl diols remain unknown although some clues are provided by the occurrence of long-chain alcohols and hydroxy fatty acids in the same species (e.g. [167]).

The distribution of alkyl diols in eustigmatophytes is not an exact match with those found in some sediments, indicating that other organisms might synthesize these compounds [160]. An additional source has now been found with the identification of C_{28} , $C_{28:1}$, C_{30} , and $C_{30:1}$ alkyl 1,14-diols, together with C_{27} and C_{29} 12-hydroxy methyl alkanoates, as major neutral lipids in rhizolenid diatoms belonging to the widespread diatom genus *Proboscia* [168]. These components were abundant in sediment traps and sediments, especially in areas with elevated primary production such as upwelling regions of the Arabian Sea.

Alkyl diols are often accompanied by the corresponding long-chain keto-ols where a carbonyl group replaces the mid-chain hydroxy group (Fig. 7). These have no known biological source, but they occur ubiquitously in marine sediments. It seems highly likely that they are formed by oxidation of long-chain diols, both in the water column and in the sediment [169].

Alkyl diols are readily identified from their mass spectra as TMSi-ethers which show intense ions due to cleavage α to the mid-chain hydroxyl group. Illustrative mass spectra are provided by de Leeuw et al. [161] and Volkman et al. [160]. The mass spectrum of the TMSi-ether derivative of the C_{32} saturated 1,15-diol shows a molecular ion at m/z 626, and major ions at m/z 341 and 387 due to cleavage either side of the C-15 hydroxy group. The mass spectrum of the of the $C_{32:1}$ 1,15-diol (as the TMSi-ether) shows cleavage ions at m/z 339 and 387, and an $M^+ - 90$ ion at m/z 534 due to loss of the OTMSi group. Alkyl

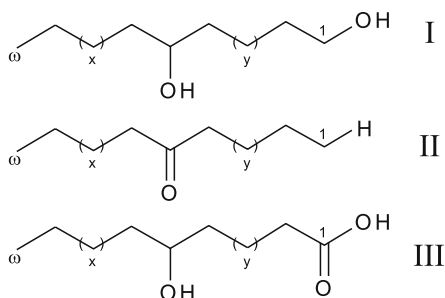


Fig. 7 Structures of alkyl keto-ols (I), alkyl diols (II) and corresponding hydroxyl fatty acids (III). In eustigmatophytes the chain-lengths are usually C_{28} , C_{30} or C_{32} and the mid-chain hydroxyl group is on C-15. In diatoms, the hydroxyl group is at C-14

keto-ols are easily identifiable from their mass spectra which show a base peak at m/z 130 and major ions at m/z 143 and $M^+ - 15$ [161].

3.6 Isoprenoid Ether Lipids

The lipids of organisms from the archaea contain ether-linked lipids [38] rather than the more common ester-linked lipids of eukaryotic organisms. A common constituent is archaeol (Fig. 8) in which C_{20} isoprenoid chains are ether-linked to the glycerol backbone. Related compounds with a hydroxyl group on the isoprenoid chain (*sn*2- or *sn*3-hydroxyarchaeols; Fig. 8) are typical constituents of the methanogenic order Methanosarcinales [170, 171]).

Glyceryl dialkyl glyceryl tetraethers (GDGTs) are specific to the archaea. A variety of chemical forms are known having different numbers of cyclopentyl rings usually designated by a numeral after GDGT. Thus GDGT-0 is the base skeleton having no cyclopentyl rings (Fig. 8). Wakeham et al. [173] showed that bicyclic and tricyclic GDGTs and their constituent ^{13}C -depleted

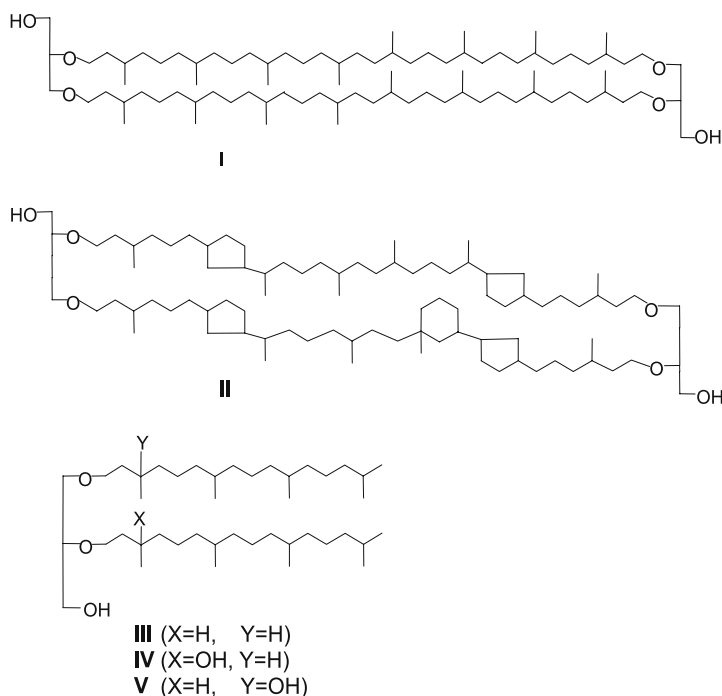


Fig. 8 Structures of some isoprenoid ether lipids of prokaryotic origin. I: GDGT-0 (from methanogenic archaea); II: Crenarchaeol; III: archaeol (X = H, Y = H) (from methanogenic archaeobacteria); IV: *sn*-2-hydroxyarchaeol (X = OH, Y = H); V: *sn*-3-hydroxyarchaeol (X = H, Y = OH) (from methanogens)

monocyclic and bicyclic biphytanes (^{13}C down to -67%) in the Black Sea were indicative of archaea involved in anaerobic oxidation of methane (AOM).

An exciting discovery was an unusual glycerol dibiphytanyl glycerol tetraether (GDGT) termed crenarchaeol in both oxic and anoxic marine waters and sediments (e.g. [174]). Its structure (Fig. 8) was identified using high field two-dimensional NMR techniques which demonstrated the presence of an unusual cyclohexane ring and four cyclopentane rings formed by internal cyclization of the biphytanyl chains [175]. Its structure is thus similar to that of GDGTs biosynthesized by (hyper)thermophilic crenarchaeota apart from the cyclohexane ring. This compound is now recognized as derived from the membrane lipids of cosmopolitan pelagic crenarchaeota. In the Black Sea, GDGT-0 and crenarchaeol dominated the distributions of ether lipids in the oxic surface and shallow anoxic waters reflecting their origin from planktonic crenarchaeota.

Compound-specific isotope analyses of the carbon skeletons suggest that planktonic archaea utilize an isotopically heavy carbon source such as algal carbohydrates and proteins or dissolved bicarbonate. Due to their high preservation potential, these lipids provide a fossil record of planktonic archaea and suggest that they have thrived in marine environments for more than 50 million years [174]. Sinninghe Damsté et al. [176] suggest that planktonic Crenarchaeota are probably facultative anaerobes and that the world's oceans contain ca. 10^{28} cells of planktonic Crenarchaeota.

Intact tetraether lipids in archaeal cell material and sediments can be analyzed by high performance liquid chromatography in combination with atmospheric pressure chemical ionization mass spectrometry [177, 178]. Non-isoprenoid dialkyl diglycerol tetraethers containing 13,16-di- or 5,13,16-trimethyloctacosanyl moieties have been identified in peats and coastal marine and lake sediments using similar techniques of HPLC-MS and high-field NMR spectroscopy [178, 179]. The ratio of these branched isoprenoids to themselves plus crenarchaeol (BIT index) has been used to estimate the relative amounts of terrestrial and marine organic inputs to marine sediments [178].

3.7

Aliphatic Ketones

A surprising diversity of aliphatic compounds containing carbonyl groups has been found in sediments and seawater. Some of these are natural products, while others are formed by diagenetic reactions.

3.7.1

n-Alkan-2-ones

Distributions of long-chain ($\text{C}_{19} - \text{C}_{35}$) ketones having a carbonyl at the 2-position (i.e. methyl ketones) have been found in soils [180], peats [181],

lacustrine sediments [182] and some coastal marine sediments (e.g. [58]). These compounds can be derived from oxidation of *n*-alkanes *via* the intermediate alkan-2-ols [182], but the mismatch of chain-lengths between alkanes and ketones in coastal sediments suggests that they are not produced *in-situ*, but rather they are transported with terrestrial organic matter to the marine environment [58]. Hernandez et al. [183] reported distributions of alkan-2-ones maximizing at C₂₅ in a subtropical estuary for which they proposed a seagrass source. It is not yet clear whether this could be a source of alkan-2-ones in other coastal environments. *n*-Alkan-2-ones are readily identified by GC-MS. Their mass spectra are similar to those of *n*-alkanes, but show a base peak at *m/z* 58 which shifts to *m/z* 59 with increasing chain-length [58].

3.7.2

Mid-Chain Ketones

Long-chain ketones having a carbonyl group near the centre of the chain (mid-chain ketones) are infrequently found in marine samples. Boon and de Leeuw [137] reported a series of these compounds, dominated by C₃₈ and C₄₀ ketones with the carbonyl at C-19, in Walvis Bay diatomaceous ooze. The origin of these compounds remains unknown, but the predominance of even carbon atoms rules out higher plants which typically contain odd-chain mid-chain ketones such as those that have been found in lacustrine settings [158].

Mid-chain ketones occur in the same chromatographic fraction as wax esters and have similar GC retention times and thus may be overlooked. A simple procedure is to re-examine the "wax ester" fraction after saponification or transesterification to see whether long-chain components are still present. Mid-chain ketones R₁COR₂ give prominent ions from cleavage α to the carbonyl group together with H and CH₃ rearrangement ions [R₁CO]⁺, [R₂CO]⁺, [R₁CO + H]⁺, [R₂CO + H]⁺, [R₁COCH₃]⁺, [R₂COCH₃]⁺, [R₁COCH₃ + H]⁺, [R₂COCH₃ + H]⁺ [137].

3.7.3

Alkenones

Very long-chain (C₃₅ – C₄₀) unsaturated methyl and ethyl ketones termed alkenones, having a carbonyl at the 2 or 3 position respectively, are found in haptophytes including the coccolithophores *Emiliana huxleyi* and *Gephyrocapsa oceanica* [45, 48, 184]. Alkenones are straight-chain alkyl lipids with *trans* double bonds. The most abundant are C₃₇ and C₃₈ chain-lengths with 2 or 3 double bonds with the carbonyl group at C-2 or C-3. Recent work has demonstrated the presence of several new alkenones in these microalgae, including monounsaturated homologs, as well as the corresponding long-chain

alkenols [185]. Alkenones are ubiquitous in marine sediments (e.g. [162, 186–189]), and the ratio of components has been found to vary systematically with the seawater temperature in which the microalgae grow [186, 189]. This has prompted many paleoceanographic studies that have used the ratio of concentrations of tri- to di-unsaturated C_{37} ketones in sediments (some over 100 million years old) to estimate the paleotemperature when the sediments were deposited (e.g. [187, 189, 190] and references therein). Environmental factors other than temperature such as salinity may also influence alkenone distributions (e.g. [191]), but these effects are not well understood or quantified [188].

3.8

Steroidal Compounds

Sterols and compounds derived from them by diagenetic reactions are ubiquitous in sediments. Their structures contain a number of unique features such as positions of double bonds, alkylation in the ring system and side-chain, and stereochemistry (Fig. 9) which makes them ideal for assigning sources of organic matter and for studying its short-term fate (e.g. [44, 192–194]). A diversity of steroidal compounds are found in marine waters and sediments, reflecting the variety of sterol distributions found in microalgae, plants and animals.

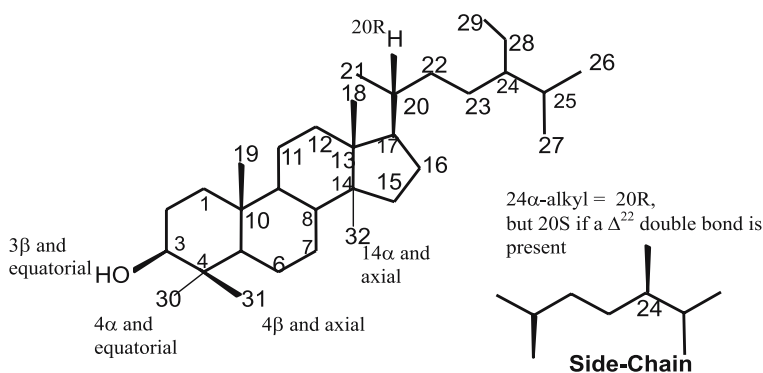


Fig. 9 Generalized structure of a 4,4-dimethyl, 24-ethyl sterol showing numbering system and stereochemistry at important centre; adapted from [194]

3.8.1

Sterols (Stenols, Stanols)

Microalgae are the primary source of sterols in the sea. Some species show a predominance of a single sterol, such as cholesterol in marine eustigmatophytes and 24-methylcholesta-5,22E-dien-3β-ol in some diatoms and haptophytes, to complex mixtures of 4-desmethyl and 4-methylsterols found in

some species of dinoflagellates [192]. Some sterols are widely distributed, but others are useful chemotaxonomic markers. For example, the C₂₈ sterol 24-methylcholesta-5,22E-dien-3 β -ol (*epi*-brassicasterol or brassicasterol depending on the stereochemistry of the methyl group at C-24) is sometimes incorrectly thought of as a unique marker for diatoms, but it only occurs in some diatoms and it is abundant in haptophytes and cryptophytes [192, 195]. A more specific marker is 24-methylcholesta-5,24(28)-dien-3 β -ol, which is abundant in some centric diatoms such as *Thalassiosira* and *Skeletonema* [196]. Some diatoms have a high content of the cholesta-5,22E-dien-3 β -ol, while others have abundant 24-ethylcholesta-5,22E-dien-3 β -ol, or cholesterol or cholesta-5,24-dien-3 β -ol [196]. Given the diversity of sterols now known to be present in diatoms and the importance of diatoms as a source of organic matter in marine systems, it is not surprising that the sterol distributions found in many marine sediments show a variety of structures.

Other algal groups display characteristic sterol distributions. For example, 24-*n*-propylidenecholesterol seems to be a marker for some chrysophytes [197] and eustigmatophytes are a source of cholesterol [160, 167]. Green microalgae can have a wide variety of sterols, many of which have double bonds at Δ^7 [198].

Dinoflagellates are the major source of 4-methyl sterols in marine systems and the C₃₀ sterol 4 α ,23,24-trimethyl-5 α -cholest-22E-en-3 β -ol (dinosterol) is often used as a biomarker for dinoflagellates (although it does occur as a minor constituent of a few diatoms [199]). However species from the genus *Pavlova* (Prymesiohyceae) also contain 4-methyl sterols and 5 α (H)-stanols as well as 3,4-dihydroxy-4 α -methylsterols termed pavlovols, but these rarely occur in marine sediments [200, 201]. An important consideration is that not all dinoflagellates have high contents of 4-methyl sterols [202] and some species lack dinosterol (Fig. 10). For example, *Amphidinium* spp. synthesize amphisterol (4 α ,24-dimethyl-5 α -cholesta-8(14),24(28)-dien-3 β -ol) as their major sterol [203] whereas several *Gymnodinium* species have 4 α ,24-dimethylcholestanol as a major sterol and contain little dinosterol [204]. An illustration of the diversity in proportions of 4-desmethyl sterols and 4-methyl sterols, even in closely related species, is shown in Fig. 10.

An uncommon example where a 4-methylsterol may not be derived from dinoflagellates comes from work by Santos et al. [205], who found high contents of 4-methylcholestanol in abyssal sediments from the Porcupine Abyssal Plain. This was also the most abundant sterol in the hindgut of holothurians (sea cucumbers), *Oneirophanta mutabilis*, suggesting that this sterol might be a marker for holothurian fecal matter in the sediments. Indeed, the significant role of benthic animals in reprocessing deposited organic matter is often overlooked in organic geochemical studies of surface sediments.

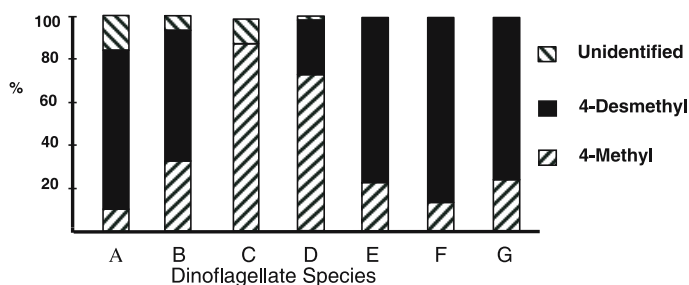


Fig. 10 Although many dinoflagellates contain a high proportion of 4-methylsterols, often dominated by dinosterol, this example shows the great range in the proportion of 4-methyl and 4-desmethylsterols in these microalgae. The dinoflagellates analysed were: A: *Gymnodinium sanguineum*; B: *Symbiodinium microadriaticum*; C: *Gymnodinium* sp.; D: *Scripsiella* sp.; E: *Fragilidium* sp.; F: *Prorocentrum mexicanum*; G: *Prorocentrum micans*. Selected data were taken from [113] and [202]

Many microalgae synthesize C_{29} sterols and a number of examples are now known where the major sterol is 24-ethylcholesterol or 24-ethylcholesta-5,22E-dien- 3β -ol [196], both of which are more commonly associated with higher plants [192]. In many near-shore sediments it is likely that much of the 24-ethylcholesterol present is derived from plants. Figure 11 shows an example where the content of the C_{29} sterol in sediments from the Huon estuary in southern Tasmania [206] is plotted against the proportion of terrestrial organic matter determined independently from stable isotope data. The good correlation shows that little of the 24-ethylcholesterol present in this case is derived from microalgae. However, in off-shore sediments where much of the organic matter is from phytoplankton, then algal sources can also be important contributors [58].

Sterols with a fully saturated ring system (5α (H)-stanols) occur in all marine sediments where they are thought to be formed by bacterial reduction of stenols. A typical distribution of stenols and stanols in a modern coastal sediment is shown in Fig. 12. Direct inputs from dinoflagellates are also possible [207], as well as minor inputs from diatoms (e.g. [196, 208]) and some haptophytes [200, 201]. The presence of 5β (H)-stanols in sediments is often taken as evidence for the presence of fecal-derived organic matter [209], since the 5β (H)-stanol coprostanol constitutes approximately 60% of the total sterols found in human feces. More detailed analyses of the carbon number distributions of 5β (H)-stanols indicate that human-derived fecal matter can be distinguished from other sources [210]. Note, however, that 5β (H)-stanols can also be formed in sediments under highly reducing conditions [192] and they can be derived from marine mammals [211]. Stanols with 3α (OH) stereochemistry can also be found; epicoprostanol (5β -cholestan- 3α -ol) is particularly abundant in Antarctic sediments and in the feces of baleen whales [210].

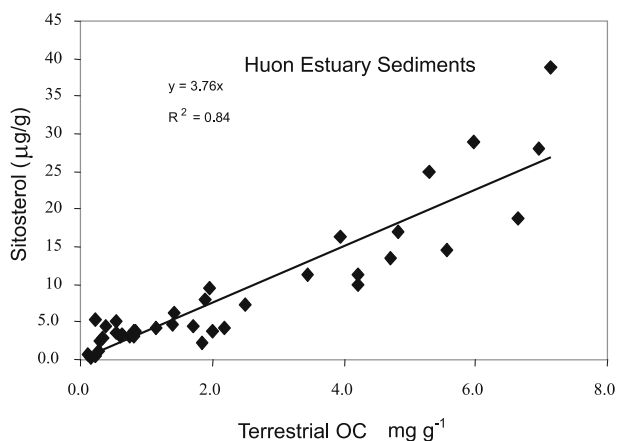


Fig. 11 Plot of sitosterol (24-ethylcholesterol) contents in sediments from the Huon estuary, Tasmania against an independent measure of terrestrial organic carbon derived from $\delta^{13}\text{C}$ values. In this instance, sitosterol provides a quantitative index of plant organic matter in the sediments, whereas in most marine sediments a contribution from microalgae may also be important. Data from Butler et al. [206]

Sterols are readily identified from their mass spectra and are usually studied as their acetate or trimethylsilyl-ether derivative, although free sterols can also produce useful mass spectra [212]. TMSi-ethers are generally preferred because they produce more diagnostic mass spectra, but are less stable than acetates [213, 214]. Acetate derivatives, however, may show a weak molecular ion, or even no molecular ion in which case the $M^+ - 60$ ion is abundant [214].

In analyses of TMSi-ethers, a few simple rules can help identify particular series. For example sterols with Δ^5 -unsaturation show a base peak at m/z 129 and fragmentation ions at $M^+ - 90$ and $M^+ - 129$. Fully saturated sterols elute slightly later (Fig. 12) and show mass spectra with m/z 215 as the base peak (m/z 229 if a 4-methyl group is present). $5\beta(\text{H})$ -stanols elute much earlier than $5\alpha(\text{H})$ -stanols. Sterols with Δ^7 , Δ^8 or $\Delta^9(11)$ lack the ion at m/z 129, and show major ions at m/z 213, 229 and 255. These mass spectra are very similar [214], but retention indices can be used to identify which isomer is present [215]. The presence of a stenol/stanol pair of peaks is good indication for a Δ^5 double bond in the first eluting compound, and conversely the absence of a second peak indicates that the ring double bond must be at a position other than Δ^5 . Sterols with $\Delta^{5,24(28)}$ unsaturation produce a major ion at m/z 386 (c.f. m/z 388 if the double bond at Δ^5 is absent). Sterols with $\Delta^{5,22}$ unsaturation show a base peak at m/z 255 (c.f. m/z 257 if the Δ^5 double bond is absent). Comprehensive compilations of retention time data on various GC phases are available for many of the commonly encountered structures [215, 216].

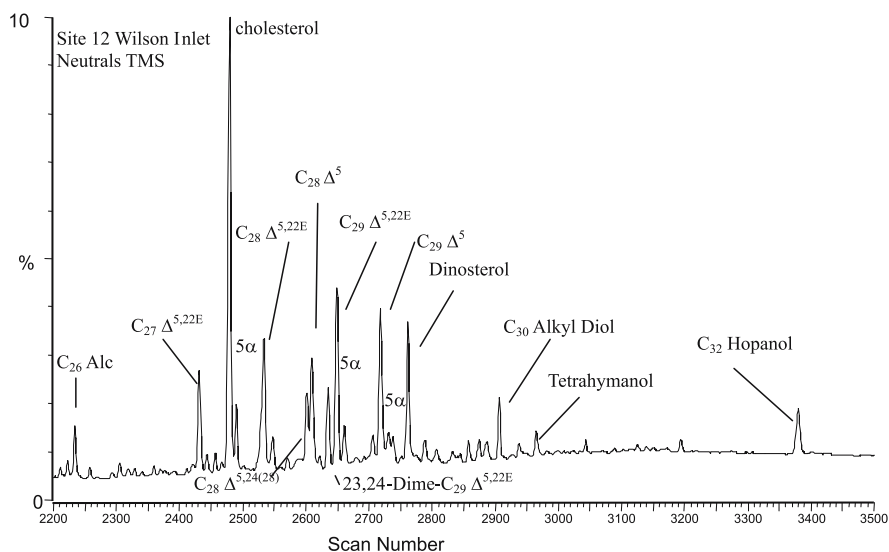


Fig. 12 Section of the chromatogram from GC-MS analysis of total neutrals in a surface sediment from Wilson Inlet, Western Australia. The sterols show a diversity of structures indicative of mixed inputs from algae, fauna, and plants. Note that each Δ^5 sterol is accompanied by smaller amounts of the corresponding 5α (H)-stanol which elutes immediately afterwards. Also shown are C_{30} alkyl diols (microalgae), tetrahymanol (protozoans) and the C_{32} triterpenoid alcohol (bacteria and/or cyanobacteria). The 4-methyl sterols are dominated by dinosterol from dinoflagellates

3.8.2

Steroid Ketones

Steroidal ketones are often found in marine environments (e.g. [217]). These distributions reflect early stage oxidation of the sterols present [217], plus direct inputs from a small subset of microalgae, such as the dinoflagellates. For example, the dinoflagellate *Scrippsiella* contains at least 21 steroidal ketones including dinosterone ($4\alpha,23,24$ -trimethyl- 5α -cholest-22E-en-3-one), dinostanone and $4\alpha,23,24$ -trimethyl- 5α -cholest-8(14)-en-3-one [202, 218]. Mass spectra for these same steroid ketones isolated from a toxic dinoflagellate *Pfiesteria piscida* are provided by Leblond and Chapman [219].

Δ^5 -unsaturated sterols can be converted to Δ^4 -3-one stenones in sediments by microbial processes similar to those operating in the rumen [220]. These compounds are readily recognized by a base peak at m/z 124. Reduction of these unsaturated steroidal ketones can then give rise to both 5α (H)- and 5β (H)-stanols.

Steroid ketones with the carbonyl group at C-3 from oxidation of the sterol 3-hydroxy group are most common. Saturated ketones (stanones) have a base peak at m/z 231 while the 4-methyl equivalents (4-methylstanones) show

a base peak at m/z 245 (see examples in [217] and [219]). The mass spectrum of Δ^{22} -unsaturated steroid ketones show strong ions due to cleavage about the side-chain. The most common example of the latter is dinosterone which has major ions at m/z 69(b), 285, 287, 314, 383 and 436 [217, 219].

Benfenati et al. [221] reported that stanones such as coprostanone (5 β -cholestan-3-one) and cholestanone can be readily converted to the silylated enol ether with the silylating reagent MSTFA (*N*-methyl-*N*-trimethylsilyltrifluoroacetamide). The more commonly used reagent BSTFA is less effective and often leads to a mixture of silylated and non-silylated forms. These enol derivatives give a strong molecular ion, M-15 and M-29 ions and a major ion at m/z 143. This side-reaction needs to be considered when analyzing a total neutral extract containing sterols and steroidal ketones after silylation. Coprostanone is found in animal feces. On non-polar GC columns it elutes just before the *n*-C₂₈ alcohol and is readily distinguished from the 5 α (H)-isomer by the presence of an M-70 ion in its mass spectrum [210].

3.8.3

Steryl Chlorin Esters (SCEs)

Steryl esters of the chlorophyll a transformation product pyropheophorbide a (SCEs; also known as phorbin steryl esters – PSEs) occur in most Recent sediments and to a much lesser degree in the particulate matter in seawater [222, 223]. It is has demonstrated that they are formed from zooplankton grazing on microalgae. For example, the copepod *Calanus helgolandicus* fed the marine diatom *Thalassiosira weissflogii* produced fecal pellets containing SCEs in which all of the animal and algal sterol components were found, although in different proportions [223]. When similar experiments were carried out with the dinoflagellates *Prorocentrum micans* and *Alexandrium tamarense*, there was a clear discrimination against the uptake of 4-methyl sterols into the SCEs [224]. SCEs degrade more slowly than free sterols thus providing a means for preserving phytoplankton-derived sterols in sediments [224].

Intact SCEs are analyzed by HPLC and identified by HPLC-MS using APCI. Chlorins can be identified by electronic (UV-visible) spectroscopy and the sterols by GC-MS [223, 224].

3.8.4

Steryl Esters and Steryl Ethers

There are relatively few reports of steryl esters in seawater and marine sediments, and in some cases their presence has been inferred by isolation of sterols from the “wax ester” fraction after saponification (e.g. [161]). Intact steryl esters have been found in sediment trap samples from the North At-

lantic [134]. The most abundant constituents were C₁₆ and C₁₈ fatty acids esterified to cholesterol, which suggests a primary source from zooplankton. Many microalgae also synthesize steryl esters (although amounts vary greatly relative to free sterols depending on the species analyzed), so this should also be considered as a possible source in surface waters and some sediments (e.g. [137]).

Steryl ethers found in marine sediments tend to be dominated by cholesteryl ethers with a C₈ – C₁₀ alkyl chain ether-linked at C-3 of the cholesterol molecule. Reports of steryl ethers are very limited [55, 137] and, indeed, there are few reports of their occurrence in the biota. Trace amounts have been found in bovine cardiac muscle [225], although quite a bit is known of their chemical properties from compounds synthesized to study sterol absorption in animals (e.g. [172]). The noticeable occurrence of these compounds in sediments underlying waters with high abundances of diatoms and the disparity between the sterol moiety of the ethers compared to the free sterols studied by Schouten et al. [55] led these authors to suggest that diatoms may be a direct source of sterol ethers, but this still remains unproven despite the numerous lipid studies of diatoms.

Steryl ethers isolated from marine sediments elute in the same general region of the chromatogram as wax esters [137]. The major ions are the molecular ion and two ions at *m/z* 368 and 370 due to loss of the cholesteryl group. A prominent ion at *m/z* 329 occurs due to cleavage of the A-ring. Boon and de Leeuw [137] provide a spectrum of 3-nonoxycholest-5-ene, while Schouten et al. [55] show a spectrum for the C₁₀ analog.

Steryl esters elute much later than wax esters on non-polar GC columns and thus high temperature columns able to operate to 370 °C or higher are required for direct analysis [134, 136]. GC-MS analysis is not straightforward due to this temperature limitation, but has nonetheless been used to identify steryl esters in sediment trap material [134]. Steryl esters rarely give molecular ions in EI or CI mode, but useful fragment ions can be obtained using CI with methane as the reagent gas [134].

3.9

Triterpenoid Alcohols

This is a very broad collection of compounds and so the discussion here will be brief. The term “triterpenoid” alcohols is usually restricted to those compounds with a 6-membered E-ring (e.g. compounds based on the oleanane or ursane skeletons), but the term can be taken to include those with a 5-membered ring such as hopanoids (Figs. 13 and 14 [194]). Triterpenoids have proven to be particularly useful to assign sources of organic matter in sediments and petroleum.

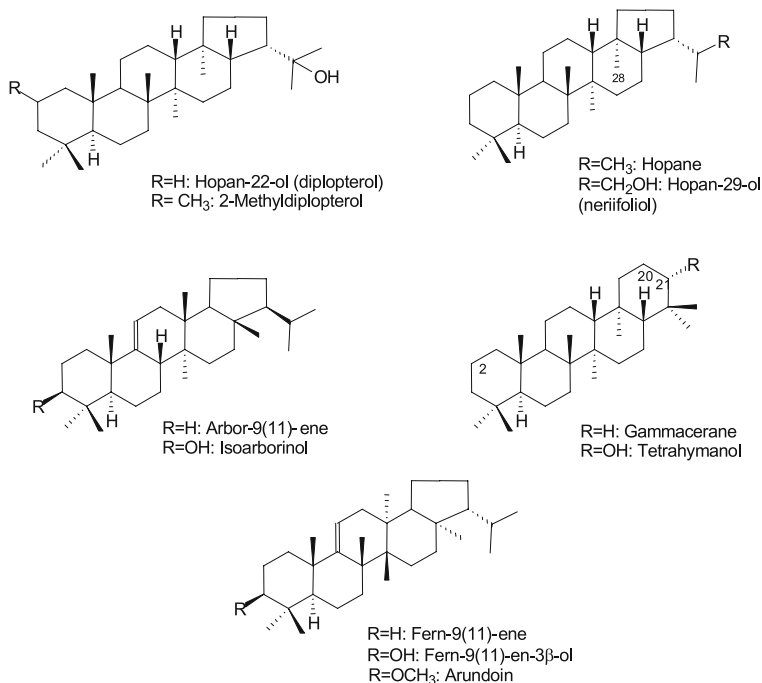


Fig. 13 Structures of some triterpenoids derived primarily from prokaryotes found in sediments

3.9.1

Hopanoids

The most common triterpenoid class in sediments is the hopanoids. Indeed, they may even be the most abundant natural product on earth [226]. Hopanoids are synthesized in bacteria from squalene which is cyclized by squalene-hopene cyclase (e.g. [227]). This reaction shares considerable similarities with that of the conversion of oxidosqualene to sterols (e.g. [194]). Hopanoids can occur in a variety of functionalized forms including alkenes, ketones, acids and alcohols (Fig. 13). Most are derived either directly from bacterial sources and have an extended alkyl chain at C-22 (e.g. [228]), or are diagenetic products. These diagenetic reactions occur very early in the diagenetic sequence in sediments (e.g. [229, 230]). For example, sediments of the highly productive lake (Priest Pot, UK) contain not only the parent bacteriohopanetetrol (BHT), but a series of hopanediols and triols such as trishomohopane-32,33-diol and bishomohopane-30,31,32-triol which represent intermediate stages in the conversion of biologically produced hopanoids to the diagenetic products, hopanes [231]. This process can be mimicked in the laboratory by oxidizing the biohopanoid in the presence of cupric chloride in pyridine [232].

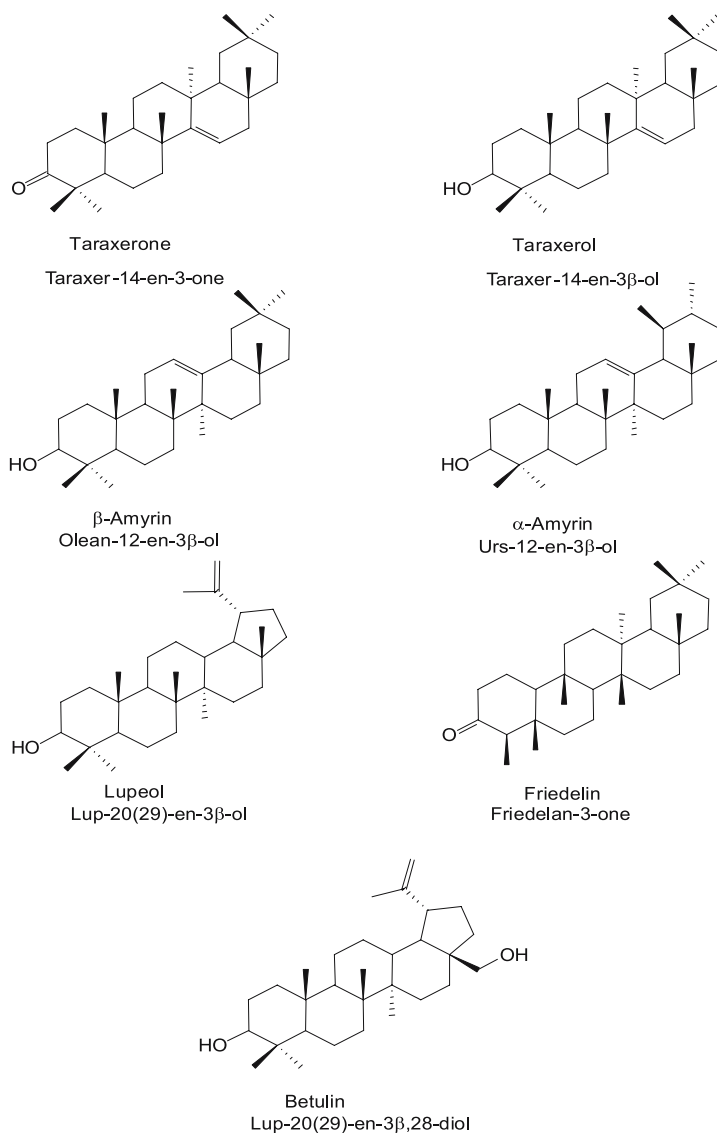


Fig. 14 Structures of some triterpenoids commonly found in higher plants

Unsaturated hydrocarbons (hopenes) such as hop-22(29)-ene, hop-17(21)-ene and 17 β (H)-hop-21-ene [233] and C₃₁ and C₃₂ $\beta\beta$ -isomers of diagenetically formed hopanes derived from polyhydroxyhopanoids of bacterial origin [234] are often present in immature sediments. Some unusual hopenes with the “moretane” stereochemistry such as 17 β (H)-moret-22(29)-ene have also been reported in sediments [235]. Fully saturated hopanes are rare in

Recent sediments and seawater and in most cases these are attributable to ancient organic matter or petroleum contamination [21], rather than biological sources. This is most clearly shown when the carbon number range extends from C₂₇ to C₃₅ and several 17,21 α - and β -isomers are present. A full discussion of these compounds would take a chapter in their own right, so the reader is referred to leading reviews [18–20].

Most hopanoids can be identified directly by straightforward GC-MS techniques (e.g. [216]), and are usually characterized with a m/z 191 mass fragmentogram (or m/z 205 for hopanoids with a methyl group in the A ring or m/z 177 for demethylated hopanes). The direct detection of most intact biohopanoids is not possible due to their highly functionalized and amphiphilic nature. Talbot et al. [236] have developed a new reversed-phase high-performance liquid chromatography method for the direct analysis of acetylated, intact bacteriohopanepolyols and applied it to solvent extracts of methanotrophic bacteria. This method was suitable for identifying the four biohopanoids: bacteriohopanetetrol and aminobacteriohopanetriol, -tetrol and -pentol.

3.9.2

“Higher Plant” Triterpenoids

A major source of triterpenoid alcohols in modern sediments is from the angiosperms, which are the most abundant plant group on the earth today. Angiosperms contain triterpenoids of the β -amyrin (oleananoid) type, which on diagenesis ultimately produces the C₃₀ triterpenoid hydrocarbon oleanane. The occurrence of oleanane in the geological record broadly correlates with the emergence of angiosperms in the Early Cretaceous and radiation in the Late Cretaceous and Tertiary (e.g. [237]).

Triterpenoids with a single hydroxyl group such as α - and β -amyrin, germanicol and taraxerol (Fig. 14), are widely used as terrestrial plant markers. For example, Killips and Frewin [238] used β -amyrin and taraxerol and their degradation products to trace organic matter from mangroves (*Rhizophora mangle*) in sediments from Florida Bay. Koch et al. [239] have had similar success in using such compounds as tracers. Volkman et al. [240] used a dihydroxylated triterpenoid betulin (Fig. 14) and other triterpenoid alcohols to show that sediments from the Wadden Sea contained organic matter from eroded peats. Note, however, that these compounds are not source-specific and can be found in a variety of plants [241], so care must be taken when using them to trace organic matter from specific plants.

3.9.3

Other Triterpenoids

Unfortunately, higher plants are not the only source of triterpenoid skeletons. For example, Smetanina et al. [242] identified the 3 β -methoxyolean-18-ene

(miliacin – the methoxy ether derivative of germanicol) in a marine fungus *Chaetomium olivaceum* (Fig. 13). α -Amyrin has been positively identified in the spores from 16 species of arbuscular mycorrhizal fungi (AMF) belonging to the order Glomales [243]. Triterpene methyl ethers have been reported in a few sediments (e.g. [244]), but their origins remain obscure. Methyl ethers of other common triterpene series (e.g. cylindrin – isoarborinol methyl ether; sawamilletin - taraxerol methyl ether; isosawamilletin – β -amyrin methoxy ether) are also known from a variety of plants, but most commonly from grasses [245, 246].

Fernene isomers of bacterial origin have also been observed in some Recent sediments (e.g. [233, 247]). The distribution of ferenes appears to follow progressive isomerization from the Δ^7 to the thermodynamically more stable Δ^8 and $\Delta^9(11)$ isomers (structures in Fig. 13) with depth, although the latter can also have natural sources [248]. These C_{30} triterpenes have a prominent m/z 243 ion in their mass spectra. Ferenes and ferenols are abundant in contemporary ferns (e.g. [249, 250]), but their most likely source in most sediments is from bacteria [233, 251]. Fern-7-ene and fern-9(11)-ene have been isolated from a purple, non-sulfur bacterium *Rhodomicrobium vanniellii* [248].

Another interesting class of triterpenoids is based on the isoarborinol skeleton (Fig. 13). Isoarborinol has been found in quite high concentration in sediments from the Permian and Triassic [252], but reports in contemporary environments are rare. It has been suggested that it is derived from a microbial source [253–255].

Gammacerane has been found in sediments from the Late Proterozoic which also contain some of the earliest examples of fossil protozoans [256]. It is presumed to be derived from tetrahymanol (gammaceran-21 α -ol) which is found in ciliates, ferns, fungi and bacteria [257, 258]. Tetrahymanol is both common and abundant in recent marine sediments (e.g. [259]).

3.10

Chlorophyll and Carotenoid Pigments

Microalgae contain a diverse array of carotenoid and chlorophyll pigments, including a number of newly discovered components. Some pigments are specific to one or a limited number of algal classes and thus can be used as biomarkers in the same way as lipids. Indeed, a valuable development has been the software program CHEMTAX [260], which uses specific pigment ratios to ascertain the abundance of microalgal groups in seawater samples. A comprehensive review of pigments is provided in the SCOR-UNESCO publication on phytoplankton pigments in oceanography [261]. This information is updated and summarized in the chapter by Wright and Jeffrey. The combination of pigment markers with lipid biomarkers such as fatty acids has proven to be very useful for elucidating sources of organic matter in the sea and for studying their rates of degradation [262].

3.11

Organic Sulfur Compounds

A great variety of organic sulfur compounds (OSCs) have been identified in marine sediments and in anoxic seawater. Most of these compounds result from incorporation of sulfur into functionalized biolipids, especially when the content of reactive iron in the sediments is low. Both intra- and intermolecular incorporation of sulfur has been documented. These OSCs thus contain biomarker information about the originally deposited organic matter. Further details can be found in recent papers [263–265].

4

Summary

A wide range of chemically distinct organic compounds can be found in marine sediments and seawater. The structures of many of these have been elucidated, although it seems that every year brings the discovery of some new class of compound. Some chemical structures can be related to specific groups of organisms, or even to a particular Class, Order or genus and thus they can be used as biomarkers for elucidating the sources of the organic matter. Many biomarkers degrade over time as a result of chemical and biological processes. The distributions of these degradation products can thus also provide a history of the breakdown of organic matter in the sediments or water column. Advances in analytical and chemical identification techniques now make it possible to identify very small amounts (ng to mg) in aquatic environment even when present in complex mixtures. Moreover, it is now possible to determine the ^{13}C and ^{14}C content in many compounds, thus providing additional information about their source and age.

Acknowledgements I wish to pay tribute to my colleagues for their assistance, advice and fruitful conversations over many years which has led to this overview of organic geochemical biomarkers. Funding from CSIRO Marine Research, the Fisheries Research and Development Corporation, Aquafin CRC, Antarctic Science Advisory Committee and the Australian Research Council is gratefully acknowledged.

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