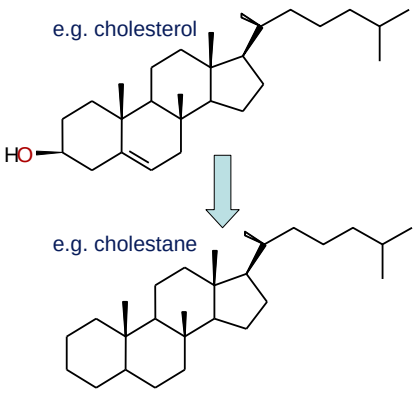
An Introduction to Molecular Markers

- **Key Reading**
- Killops S. and Killops V. (2005) An introduction to Organic Geochemistry, 2nd Edition. Blackwell Scientific. 393 pp.
- **Suggested Reading**
- Volkman J.K., Barrett S.M., Blackburn S.I., Mansour M.P., Sikes E.L. and Gelin F. (1998) Microalgal biomarkers: A review of recent research developments. Org. Geochem. 29, 1163-1179.
- Sinninghe Damste et al., 2004. The Rise of the Rhizosolenid Diatoms. Science. 304, 584-587.
- Coolen et al., 2004. Combined DNA and lipid analyses of sediments reveal changes in Holocene haptophyte and diatom populations in an Antarctic lake. EPSL, 223,225-239.
- Volkman J.K. 2005. Sterols and other triterpenoids: source specificity and evolution of biosynthetic pathways. Org. Geochem. 36, 139-159.

Life, Molecules and the Geological Record

- Life leaves molecular residues (*Chemical Fossils*) as well as visible shapes/objects (*Morphological Fossils*) in the sedimentary record.
- These molecular residues, when characterised as specific molecules (*Biomarkers*) by their structures and isotopic content, may give precise indications of their biosynthetic origins in particular organisms, as well as the environmental conditions that the organisms experienced.



Definition of a biomarker (or "molecular marker" or "geochemical fossil"):

"A molecule whose carbon skeleton can unambiguously be linked to that of a known biological precursor compound"

More generally:

"Organic compounds found in sediments which have properties that can be directly related to a known biological precursor"

Biological marker molecules

- Living organisms biosynthesize a very small subset of the billions of molecules that can be assembled in theory from C, H, O, N, S, P etc.
- These molecules can be regarded as biomarkers. Their presence in an environment reflects their synthesis by the parent organisms.
- Some biomolecules are produced only by a certain species or classes of organism, and hence indicate the presence or prior existence of those organisms (e.g., diatoms).
- Other biomolecules are produced by many species of organism and are indicative of the general level of biological activity (e.g., eukaryotes vs prokaryotes).
- Molecular signatures can comprise the only means to decipher past ecosystems and biological inputs for organisms composed only of soft parts (i.e., leave no morphological or isotopic imprint).

NB. This is an organic geochemist's definition of a biomarker

Molecular Characteristics of biomarkers

- Biomarkers are usually characterized by a high degree of order in their molecular structures, resulting from the specificity of the biosynthetic processes:-
  - Small molecule building blocks
  - Precise sequence of assembly
  - Chirality of carbon centers and stereochemistry of the units
  - Distribution of isotopes in the molecule
- Intra-molecular characteristics documented by structural identification and molecular isotope measurements.
- Inter-molecular variations assessed through compound distributions (e.g. abundance ratios).

Structural uniqueness

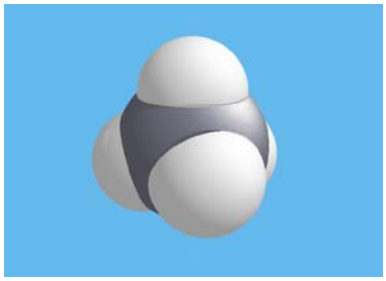
- molecular structure (carbon skeleton)
- stereochemistry

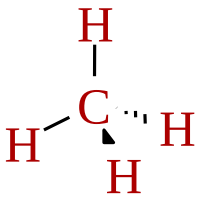
Example: Only three C_{31} hydrocarbons have been identified in plants (normal- iso- and anteiso-) although there are $>10^9$ possible isomers.

Distributional uniqueness

- isotopic composition (^{13}C, D/H)
- abundance

Methane, CH_4 - The Smallest Biomarker?



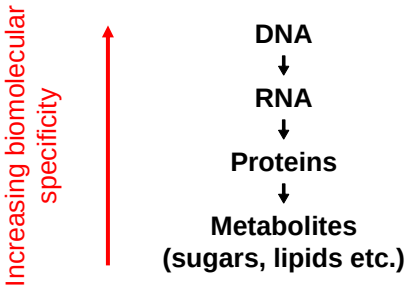


Isotopes:

Carbon: ^{12}C, ^{13}C, ^{14}C

Hydrogen: ^{1}H, ^{2}H, ^{3}H

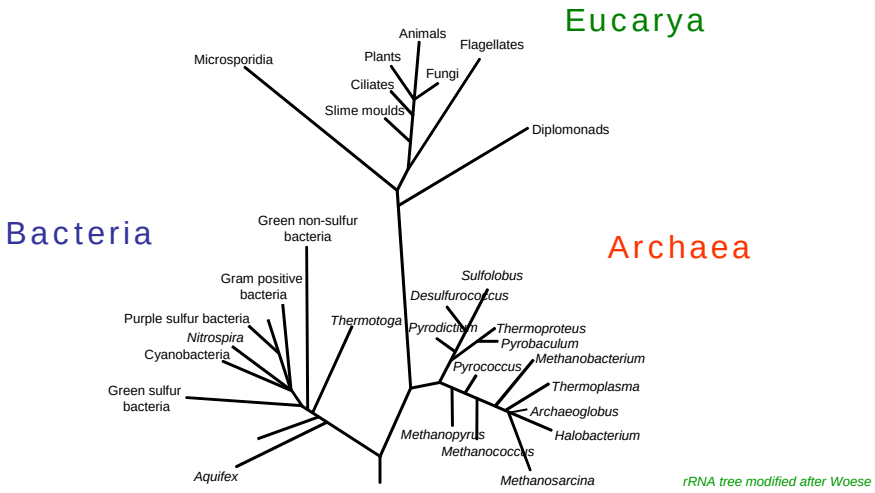
Universal biomolecular machinery



Key criteria:

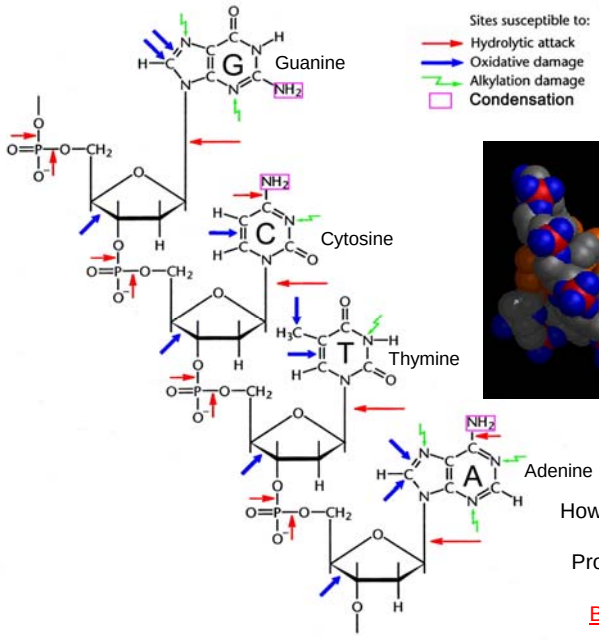
- Information content
- Robustness of molecule
- Ease of detection and analysis (both structural and isotopic)

DNA and the phylogenetic tree

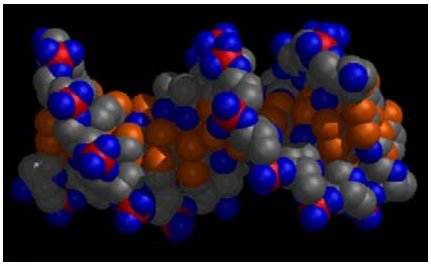


rRNA tree modified after Woese

DNA - The Biggest Biomarker!!



Sites susceptible to:
- Hydrolytic attack
- Oxidative damage
- Alkylation damage
- Condensation



How good is DNA as a biomarker?

Problem: It is not preserved well

BUT see Coolen et al., 2004

Lipids

- Lipids present in the water column and in sediments can originate from all three domains of life (i.e., eukaryotes, bacteria, archaea).
- Certain lipids are synthesized by only one domain.
  - Steroids are almost exclusively synthesized by eukaryotes
  - Hopanoids are exclusively synthesized by bacteria
  - Acyclic and cyclic isoprenoid ether lipids are restricted to the archaea.

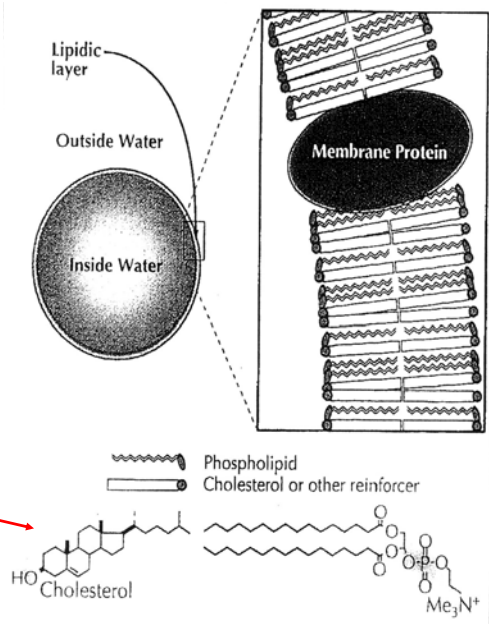
Occurrence:

- Ubiquitous
- 10-20% of TOC in most organisms
- Extensively studied classes of compounds
- - analytically accessible
- - diagenetically and chemically [relatively] stable
- - structurally extremely diverse (high potential as "biomarkers")

Function:

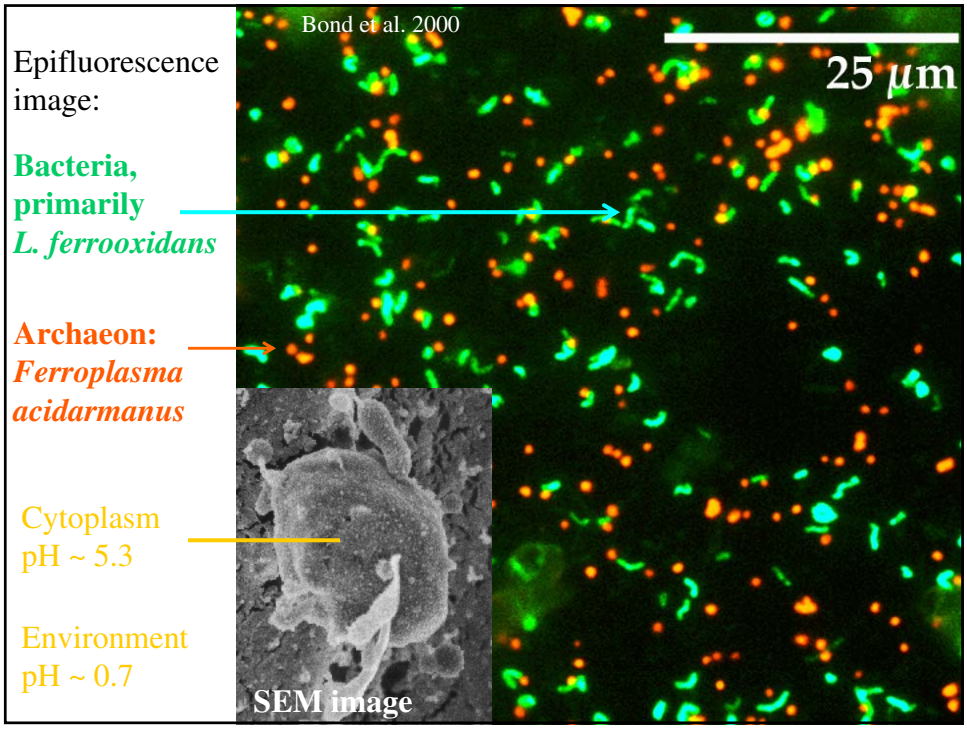
- - Long-terms energy storage
- - membrane fluidity regulators
- - membrane rigidity/barrier to proton exchange
- - pigments
- - hormones
- - vitamins

All-important cellular membrane: Source of lipids in the environment



Steroids in Eukaryotes
Hopanoids in Bacteria
Acyclic isoprenoid ethers in Archaea

Ourisson and Nakatani, 1994



Lipid Structures:

Several major compound classes:

- Fatty acids
- Fatty alcohols
- Hydrocarbons
- Terpenoids

Fall into two main groups:

- Polyketide lipids
- Polyisoprene lipids

Occur as “free” compounds or chemically bound (ester or ether linkages) to other biochemical components (e.g., glycerol).

Lipid distributions in plankton

Composition of lipid fraction of diatoms (Clarke and Mazur, 1941)

	%
Uncombined (free) fatty acids	59-82
Combined (bound) fatty acids	1-17
Non-saponifiable (tightly bound) lipids	12-29
Fatty Alcohols	3-7
Hydrocarbons	3-14

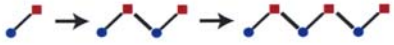
Composition of lipid fraction (%) of copepods (Lee et al., 1970)

	C. helgolandicus	*G. princeps*
Hydrocarbons	Tr	Tr
Wax esters	37-30	73
Triglycerides	5	9
Polar lipids	14-17	
(free acids, sterols,	50-60	17
phospholipids)		
Total lipid (% dry wt.)	12-15	29

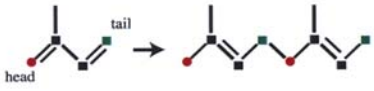
Lipid biosynthesis

Occurs via two main pathways:

1. Polyketide Biosynthesis: The polymerization of acetate; products typically have even carbon numbers.



2. Isoprenoid synthesis: The polymerization of isoprene; products typically have 10, 15, 20 ….. carbon atoms.



Group I. Polyketide lipids

Compounds whose structure is based on repeat units of acetate.

Products usually have an even number of carbons.

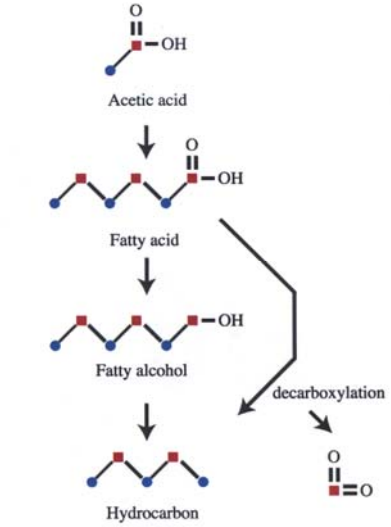
Common compound classes:

- Fatty acids
- Fatty alcohols
- Hydrocarbons (*n*-alkanes)

Nomenclature

- *Normal* alkanes and alcohols are denoted: *n*-C_x where x = number of carbon atoms
- e.g. *n*-C_{15} = normal pentadecane
- Fatty acids are often unsaturated $C_{x:y}$ where y = number of double bonds
- double bonds are usually *cis*
  Δ denotes the position of unsaturation from the COOH end of the molecule
  - e.g. Oleic acid is $^{9}\Delta$ $C_{18:1}$

  ω denotes unsaturation from the methyl end
- Carbon Preference Index (CPI), Odd/Even Predominance (OEP) and Average Chain Length (ACL) used to describe distributions.



Polyketide lipids - Fatty Acids

- Fatty acids are abundant in most organisms (often the most abundant lipid type).
- Sources include bacteria, microalgae, higher plants and marine fauna (e.g., zooplankton).
- Each source has a distinctive profile although some fatty acids are ubiquitous (e.g., $C_{16:0}$, $C_{18:0}$), and profile may change with growth status/stress of the organism.
- Bacteria are a major source of branched fatty acids (iso-, anteiso-, mid-chain branched) and can also be a major source of $C_{16:1n-7}$ and *cis*-vaccenic acid ($C_{18:1n-7}$).
- Microalgae and bacteria are the major sources of fatty acids found in most sedimentary environments.
- Different microalgal inputs can potentially be distinguished based on fatty acid distributions, especially based on # and positions of double bonds.
- Some microalgae contain high concentrations of specific long-chain essential fatty acids (e.g., $C_{20:5n-3}$; $C_{22:6n-3}$).

Biosynthesis of Fatty Acids

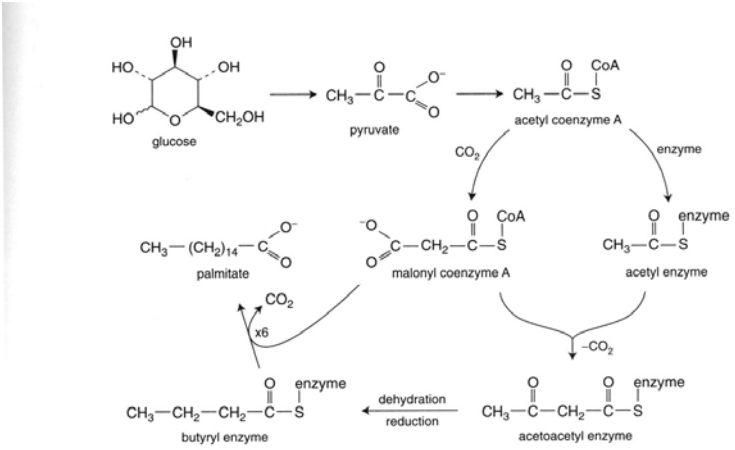


Fig. 2.13 Biosynthesis of saturated fatty acids in plants and animals. Palmitate is formed by successive additions of malonyl coenzyme A to the enzyme-bound chain, with CO_2 being lost at each addition. This results in chain elongation by a $(CH_2)_2$ unit at each step. Details of the formation of butyryl (C_4) from acetyl (C_2) are shown, while the subsequent six further additions, terminating in palmitate, proceed similarly.

Polyketide lipids - Fatty Acids

- In most marine organisms, fatty acids occur predominantly as polar lipids, such as glyco- or phospholipids. [*Also sulfo-lipids in picoplankton: see Van Mooy et al., PNAS, 2006, 103, 8607–8612.]*
- Free fatty acids are rarely abundant in living organisms, but in sediments they can be the major form due to rapid chemical or enzymatic hydrolysis of polar lipids.
- In the water column and contemporary sediments, intact esterified lipids are usually associated with the indigenous organisms.
- Fatty acid distributions in sediments have been used successfully to characterize bacterial populations (phospholipid fatty acid, PLFA) analysis. Measurement of intact polar lipids now feasible – provides more specificity.
- A common feature of fatty acid distributions in sediments is the presence of C_{20}-C_{30} saturated straight-chain fatty acids that show a strong predominance of even chain lengths. In many cases these are probably derived from higher plant leaf waxes. However, algae and bacteria can produce these lipids, albeit in small amounts relative to C_{14}-C_{20} acids.

Esterified lipids

Diglyerides and Triglycerides ("fats")

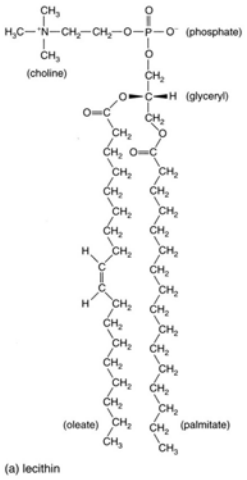
- Esters of glycerol + fatty acids
- formed by condensation (-H_2O) reactions
- fatty acids usually straight-chain (C_{14}-C_{18}) with various levels of unsaturation

Wax esters

- Esters comprising of a fatty acid + fatty alcohol

Phospholipids

- Fatty acid(s) + phosphoric acid + glycerol (+ basic nitrogen)



Phospholipid

Polyketide lipids - Long-chain alcohols

- Microalgae are not a major source of these lipids in most sediments.
- Long-chain diols occur in most marine sediments, and in some cases (e.g., Black Sea Unit I) they can be major lipids.
- A microalgal source was discovered when $C_{30\text{-}32}$ alcohols and diols were identified in marine eustigmatophytes from the genus nannochloropsis (although the distn differed significantly from that in sediments).
- It is suggested that these diols are the building blocks of novel aliphatic biopolymers produced by these microalgae.
- Most studies of mid-chain diols report presence of C_{30} and C_{32} saturated constituents having a predominance of 1,15 isomers.

C30-C32 alkyl diols and unsaturated alcohols in microalgae of the class Eustigmatophyceae

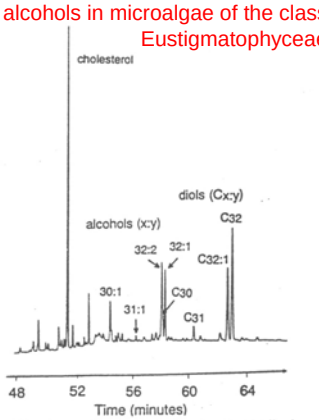


Fig. 1. Partial gas chromatogram showing the distribution of higher molecular weight components in *N. oculata* (strain CS-179) after acid hydrolysis of the extract and derivatization with BSTFA. Alcohols are designated *x*:*y* and diols $C_{x:y}$ where *x* is the number of carbon atoms and *y* is the number of double bonds. Note that the C_{30} diol elutes on the downslope of the peak due to the 32:2 alcohol. Quantitative data are shown in Table 2.

Polyketide lipids - Hydrocarbons

- Biogenic alkanes and alkenes are a common feature of the hydrocarbon distributions in sediments.
- Many microalgae contain the highly unsaturated alkene *n*-$C_{21:6}$ (*n*-heneicosa-3,6,9,12,15,18-hexaene) formed by decarboxylation of the $C_{22:6n-3}$ fatty acid. However, this compound is rarely found in sediments, probably because it is rapidly degraded.
- There are several reports of shorter-chain *n*-C_{15}, n-C_{17} and n-C_{19} alkanes (and monounsaturated alkenes) in algae. The n-C_{17} alkane is common in contemporary and ancient sediments, and if frequently interpreted as a biomarker of algal input.
- Several microalgae contain very long chain alkenes, including $C_{31:2}$, $C_{33:3}$ and $C_{33:4}$, and $C_{37:2}$ and $C_{37:3}$ (also C_{38} counterparts).

Generalized Lipid Distributions

	Acids	**Alcohols**	**Hydrocarbons**
Phytoplankton	even/odd CPI 16:0, 16:1 18:0, 18:1	even/odd CPI	odd/even nC17, nC18
Bacteria	iso + anteiso	?	CPI = 1 nC13-nC30 nC17-nC20
Zooplankton	same as phyto		same as phyto
Higher plants	even/odd CPI max C28-C30	C28, C30, C32	odd/even max C29-C31

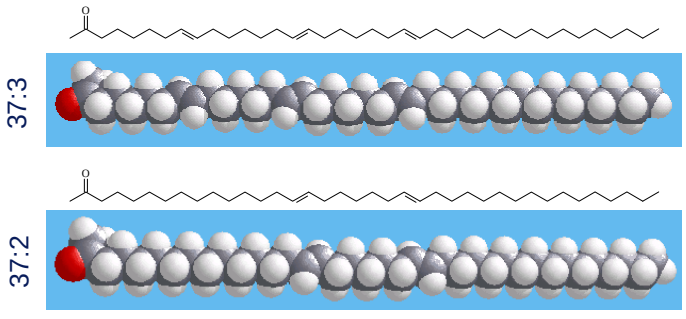
- Lipids in higher plants mainly associated with leaf cuticles ("waxes")
- Serve as physical protection against mechanical abrasion, microbial attack and water loss.

Polyketide lipids - Hydroxy fatty acids

- A wide range of hydroxyl fatty acids has been found in sediments, these compounds have received relatively little attention from organic geochemists.
- These compounds can be separated into different categories based on the # and position of the hydroxyl groups.
- Aliphatic α- and β-monohydroxyfatty acids occur in a wide range of organisms.
  - α-hydroxy fatty acids are intermediates in fatty acid biosynthesis in yeasts.
- Bacterially-derived β-hydroxy fatty acids are found in many recent sediments. The carbon # range is typically from C_{10}-C_{20}, typical of carbon # distribution for lipopolysaccharide cell walls of gram negative bacteria. Bacteria also contribute significant amounts of *iso* and *anteiso* branched C_{12}-C_{18} β-hydroxy fatty acids.
- Higher plant cutin and suberin can also be a significant source of esterified C_{16}-C_{22} α-, β-, and ω-monohydroxy fatty acids.
- Recent work suggests microalgae are also a potential source of monohydroxy fatty acids.
- C_{30}-C_{34} mid-chain hydroxyl fatty acids were identified in hydrolyzed extracts of marine eustigmatophytes of the genus Nannochloropsis (related to the long-chain diols).
- C_{22}-C_{26} saturated and monounsaturated α-hydroxyfatty acids have also been found as major lipid components of the cell wall of several marine chlorophytes.

Polyketide lipids - Long-chain ketones, esters

- Long-chain unsaturated ketones (alkenones) have been identified in several species of haptophytes, esp. the widely distributed coccolithophorids *Emiliania huxleyi* and *Gephyrocapsa oceanica*.
- These compounds will be the focus of a separate lecture.

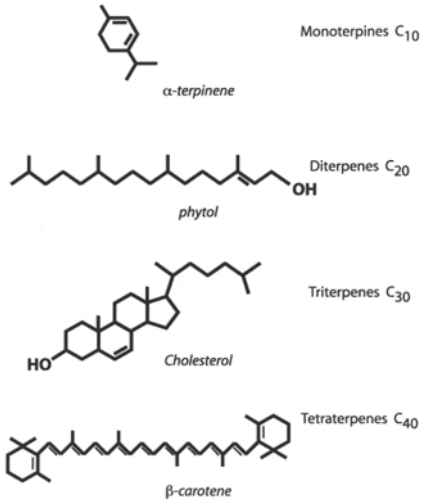


E. huxleyi

Group II. Polyisoprenoid lipids

Biosynthetically related to the polymerization of isoprene (C5)

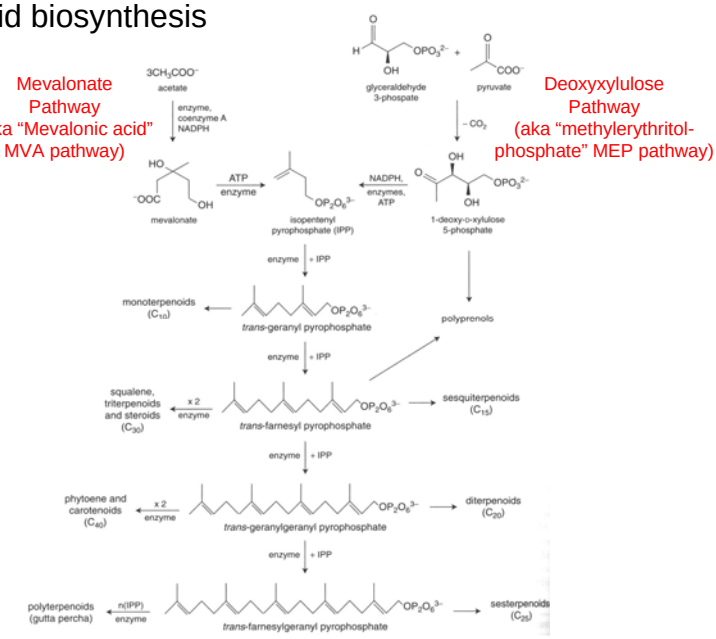
- dimer - monoterpene (C_{10})
- e.g. essential oils
- trimer - sesquiterpenes (C_{15})
- e.g. farnesol, abeitic acid
- tetramer - diterpenes (C_{20})
- phytol
- hexamer - triterpenes (C_{30})
- steroids, hopanoids
- octamer - tetraterpenes (C_{40})
- carotenoids, ether lipids
- polymer - polyterpenes
- e.g. natural resins (polycadinene)



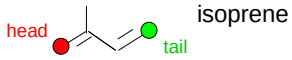
Isoprenoid lipid biosynthesis

Mevalonate Pathway (aka "Mevalonic acid" MVA pathway)

Deoxyxylulose Pathway (aka "methylerythritol-phosphate" MEP pathway)



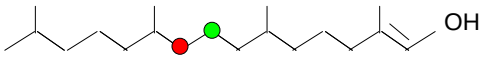
Isoprenoid lipids - Configurations



isoprene

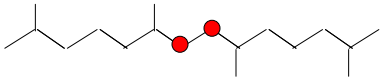
Regular isoprenoids

- head-to-tail
  - e.g. phytol

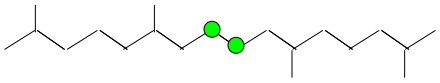


Irregular isoprenoids

- head-to-head



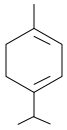
- tail-to-tail



Isoprenoid lipids

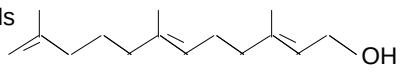
Monoterpenes (C_{10})

- Abundant in both higher plants and algae
- not extensively used as biomarkers (very volatile, so not well preserved in sediments)

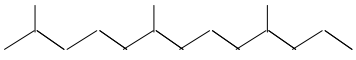

α-terpinene

Sesquiterpenes (C_{15})

- farnesol
- esterified to bacterial chlorophylls

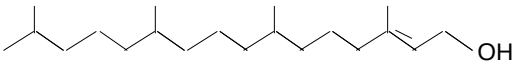


- ancient analogue - "farnesane"



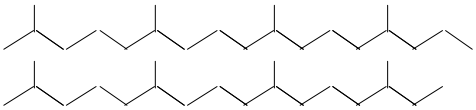
Isoprenoid lipids

Diterpenes (C20)
Acyclic diterpenoids
- phytol: *trans*-3,7(R),11(R),15-tetramethylhexadec-2-ene-1-ol



sources:
- esterified to chlorophyll-*a,b*
- phytanylethers (archaebacteria)

ancient analogues
- Phytane (C_{20})
- Pristane (C_{19})

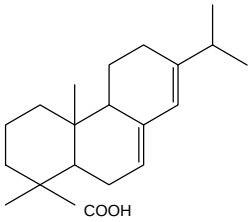


origin of pristane
- chlorophyll ?
- tocopherol (vitamin e)?

- Lots of literature on pristane/n-C_{17}, phytane/n-C_{18}, and pristane/phytane ratios.
- Used to assess depositional environment and degradation of petroleum

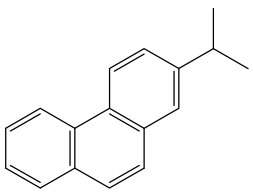
Isoprenoid lipids

Cyclic diterpenoids
- abietic acid



- from gymnosperm (conifer) resin/gum
- not widely found in marine algae
- therefore, excellent "biomarker" for higher plant input

<u>ancient analogue (also combustion product):</u>
- retene



Highly branched isoprenoid alkenes

- Highly branched unsaturated C_{20}, C_{25}, C_{30} alkenes (containing between 3 and 6 double bonds) are observed in most marine sediments. These compounds appear to derive exclusively from certain types of diatoms, although the function and bioactivity of these compounds is yet to be established.

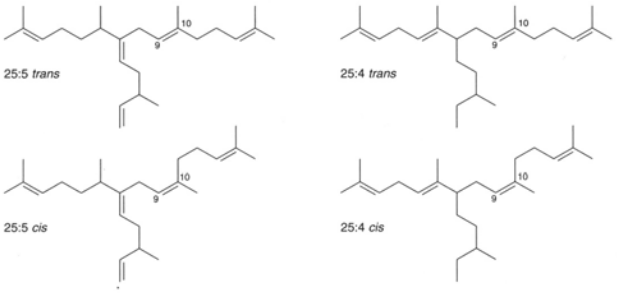


Fig. 2.18 Examples of highly branched sesterterpenes isolated from planktonic diatom cultures (*trans* and *cis* refer to configuration at double bond between C-9 and C-10; after Belt et al. 2001).

Isoprenoid lipids

Cyclic Triterpenoids (C30)

- Squalene ($C_{30}H_{50}$) - main biosynthetic precursor to cyclic triterpenes.
- An irregular isoprenoid (tail-to-tail)

- *Pentacyclic triterpenoids*
- oleanane type
- ursane type
- lupane type
- hopane type
- gammacerane type
- arborane type
- primary sources: bacteria, higher plants

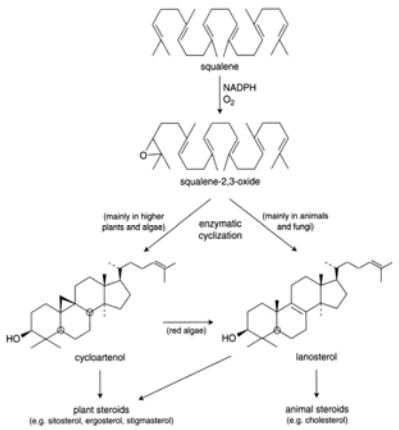


Fig. 2.21 Steroid biosynthesis from squalene (after Harwood & Russell 1984).

Pentacyclic triterpenoids

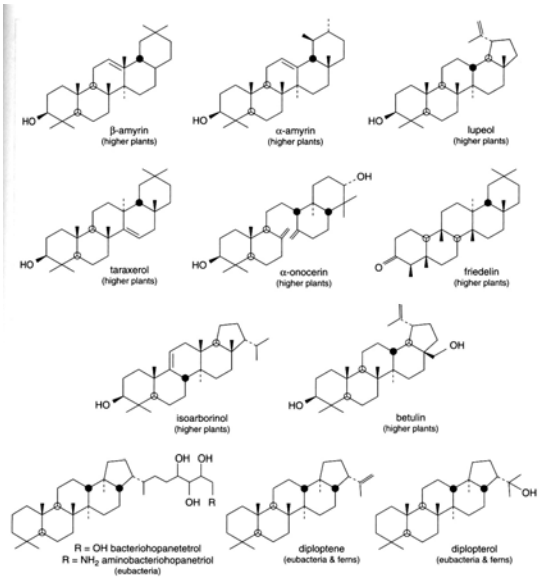


Fig. 2.19 Some geochemically important polycyclic triterpenoids and their major sources.

Carbon numbering of steroids and hopanoids

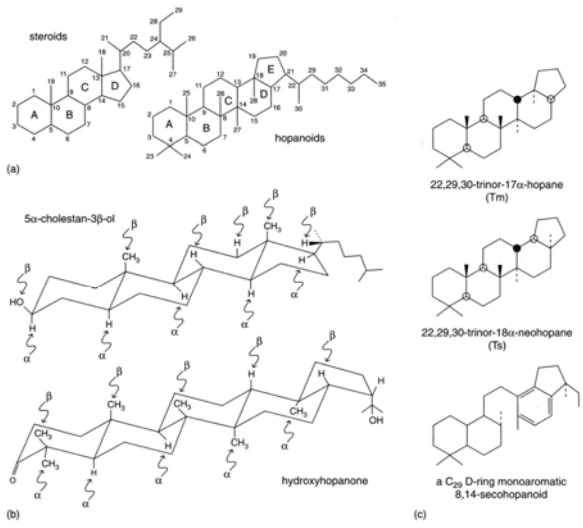


Fig. 2.20 (a) R ing numbering conventions for steroids and hopanoids; (b) examples of 'all-chair' conformations (with *trans* ring junctions) for steroids and hopanoids; (c) examples of application of hopanoidal nomenclature system. (Ts may also be called 17-methyl-18,22,29,30-tetranor-17α,18α-hopane.)

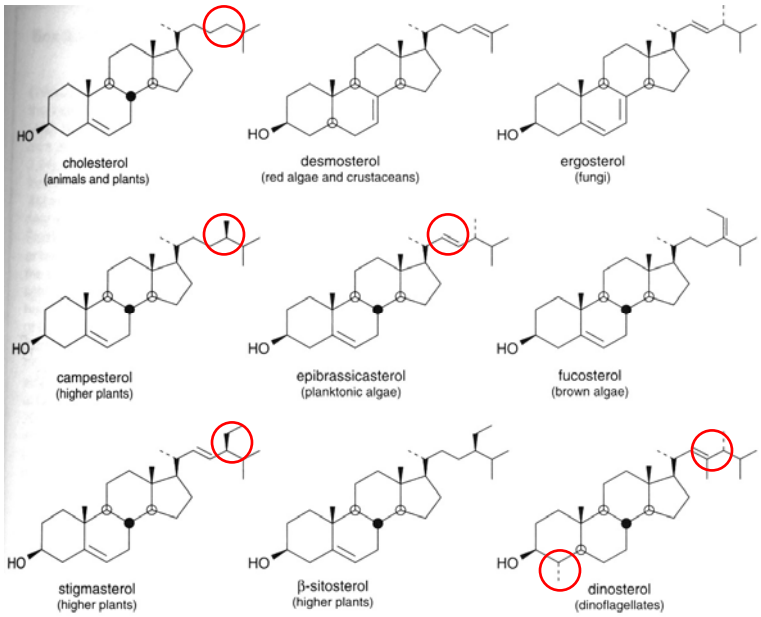
Isoprenoid lipids - Tetracyclic triterpenoids (Sterols)

Numbering and nomenclature:
- -, - below (dashed arrow) and above (bold arrow) the ring
- *R/S* stereochemistry at a ring juncture and in side chain
- sterol/stenol - unsat'd alcohol
- stanol - sat'd alcohol
- sterene - unsat'd alkene
- sterane - sat'd alkane

Occurrence:
- very widely distributed in plants and animals
- As a rule, bacteria do not make sterols

Structure:
- Mainly C27, C28, C29 (also C26,C30)
- Basic steroid skeleton is modified through oxidation and alkylation
- Hundreds of natural products based on this skeleton have been identified.
- cholesterol C27 (universally distributed)
- β-sitosterol C29 (higher plants)
- brassicasterol (diatoms)
- dinosterol C30 (dinoflagellates)
- fucosterol (brown algae)

Steroids



Sterols (continued)

- A great diversity of sterols are found in microalgae. Distributions range from the predominance of a single sterol, such as cholesterol in marine eustigmatophytes and 24-methylcholesta-5,22-dien-3β-ol in some diatoms and haptophytes to mixtures of 10 or more 4-methyl and desmethylsterols.
- Some sterols are widely distributed but others are chemotaxanomic markers.
- The diatoms display considerable diversity in sterol composition, and given the importance of diatoms as a source of organic matter in marine systems it is not surprising that sediments display complex and varying sterol distributions.
- Sterols derived from dinoflagellates are often major constituents of the sterol distributions.
- The sterol composition of dinoflagellates is dominated by 4α-methyl sterols, including dinosterol (4α,23,24-trimethyl-5α-cholest-22E-en-3β-ol) – often used as an indicator of dinoflagellate inputs to sediments.
- Sterols with a fully saturated ring system (5α(H)-stanols) often occur in dinoflagellates but are not common in other marine microalgae. Hence dinos are the major direct source of stanols to marine sediments, supplementing those formed by bacterial reduction of stenols.

Other Steroids

- *Related compounds:*
- Steroidal ketones – primarily intermediates in the microbially or chemically mediated degradation in sediments of stenols to sterenes. A direct biological source is also possible.
- Steroidal diols – one species of the genus Pavlova (Haptophyta) contains novel 3,4-dihydroxy-4α-methyl sterols. Source specificity not yet known.

Tetraterpenoids (C40)

- **A. Ether lipids**
- Occurrence:
- Archaea
- methanogens
- thermoacidophiles
- extreme halophiles
- eurythermal archaeota
- Function:
- Membrane rigidifiers
- Structure:
- unusual linkage type (mainly head-to-head)
- Ancient analogues:
- head-to-head acyclic isoprenoid alkanes.

- Focus of a separate lecture

Tetraterpenoids - B. Carotenoid pigments

- Occurrence:
- Universally distributed in all photosynthetic organisms
- Function:
- Accessory pigments, antioxidants
- Structure:
- Many different structures (>100 identified to date)
- Bacillophyceae
- fucoxanthin
- diadinoxanthin
- diatoxanthin
- β-carotene
- Dinophyceae
- peridinin
- Ancient analogue:
- β-carotane

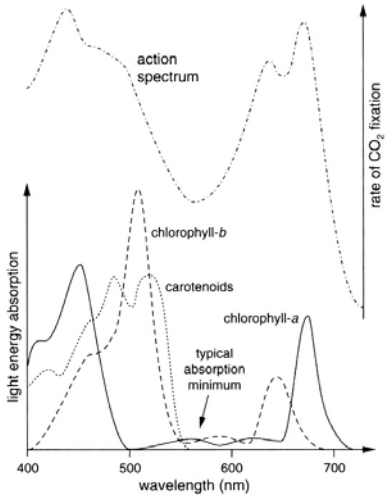
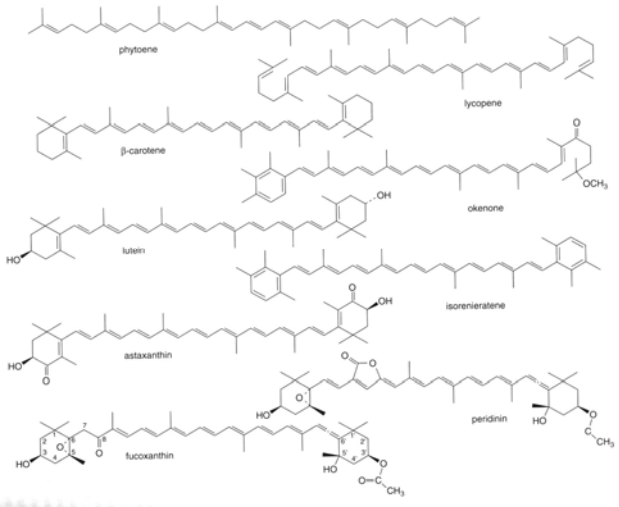


Fig. 2.24 Light absorption characteristics of some photosynthetic pigments and their relationship with utilization of light energy during photosynthesis (action spectrum).

Carotenoids



Because of their high degree of unsaturation, carotenoids are usually not well preserved in the sedimentary record

Chlorophylls

Occurrence:

- Universally distributed in all photosynthetic organisms

Function:

- Used for photosynthesis

Structure:

- All are tetrapyrroles
- Chl-a,b,c1,2,3; a1+a2; b1+b2, d,e, - oxygenic photosynthetic organisms
- bchl-a,b,c,d,e - bacteriochlorophylls

Abundance:

- Ratio of carbon/chl = 60 for phytoplankton

Ancient analogue:

- Porphyrins were the first molecules to be recognized in ancient sediments and petroleum as of biological origin - structurally related to chlorophylls (Treibs, circa 1934). This was seen as the beginning of organic geochemistry.

chlorophyll	structure	R^1	R^2	R^3	R^4
chlorophyll-*a*	I	$-CH_2=CH_2$	$-CH_3$	$-CH_2-CH_3$	phytyl
chlorophyll-*b*	I	$-CH_2=CH_2$	$-C(=O)H$	$-CH_2-CH_3$	phytyl
chlorophyll-a_2	I	$-CH_2=CH_2$	$-CH_3$	$-CH_2=CH_2$	phytyl
chlorophyll-b_2	I	$-CH_2=CH_2$	$-C(=O)H$	$-CH_2=CH_2$	phytyl
chlorophyll-c_1	II	$-CH_2=CH_2$	$-CH_3$	$-CH_2-CH_3$	H
chlorophyll-c_2	II	$-CH_2=CH_2$	$-CH_3$	$-CH=CH_2$	H
chlorophyll-*d*	I	$-C(=O)H$	$-CH_3$	$-CH_2-CH_3$	phytyl
bacteriochlorophyll-*a*	III	$-C(=O)CH_3$	$-CH_3$	$-CH_2-CH_3$	phytyl, farnesyl or geranylgeranyl
bacteriochlorophyll-*b*	III	$-C(=O)CH_3$	$-CH_3$	$=CH-CH_3$	
bacteriochlorophyll-*e*	IV	$-C(H)(OH)CH_3$	$-C(=O)H$	$-CH_2-CH_2-CH_2-CH_2-CH_2-CH(CH_3)-CH_3$	farnesyl

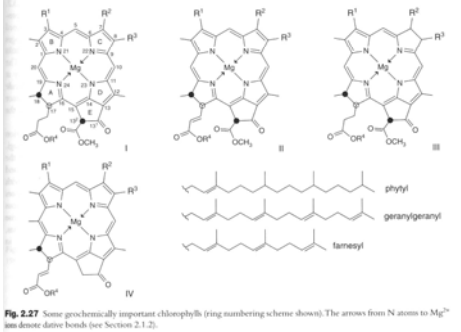
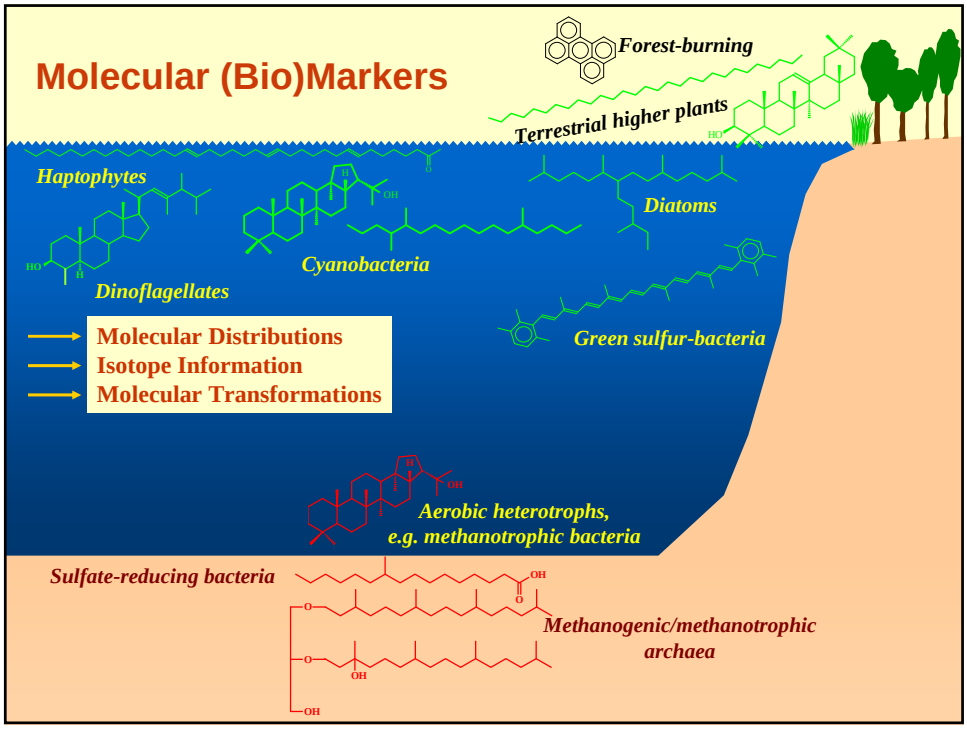


Fig. 2.27 Some geochemically important chlorophylls (ring numbering scheme shown). The arrows from N atoms to Mg^{2+} ions denote dative bonds (see Section 2.1.2).



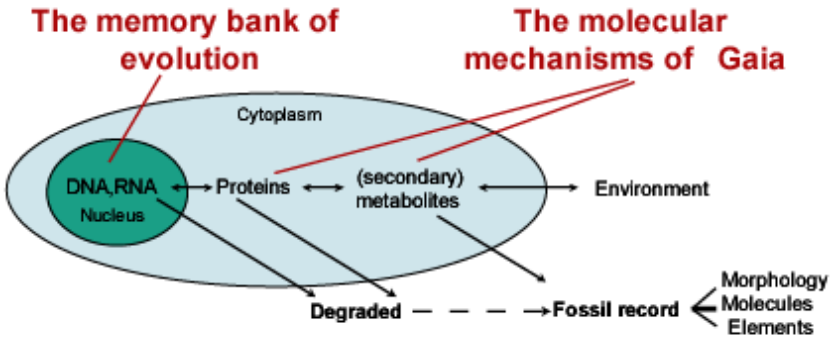
Biomarker Properties

We can measure and utilise their:

- Precise molecular structures, including stereochemistry
- Relative and absolute amounts
- Isotopic composition – C, H, N etc

Biomarker Research Areas

- Petroleum Exploration
- Environments & Ecology, including anthropogenic effects ("environmental forensics")
- Palaeoenvironment Reconstruction & Climate Change,
  - e.g. Palaeooceanography - Catastrophic events
- Carbon cycles studies
- Exploration of Biosphere – especially the Microbial World
  - e.g. Extreme environments, deep biosphere etc.
- Evolution of the Biosphere, Origin of Life & Archaean Studies
- Meteoritics, Exobiology & Planetary Studies
  - e.g. Moon, Mars
- Archaeology



DNA (Genotype) does not interact with environment directly

- Preservation potential of DNA/RNA is very low
- Instead: study living and recent organisms and make phylogenetic tree.

Secondary metabolites (phenotype) interact with environment

- Check contemporary DNA tree for key genes controlling biosynthetic pathways
- Links of secondary metabolites to the environment inform us about their function and can lead to selection of new biomarker proxies.
- Search fossil record for times of first appearance of the biomarkers

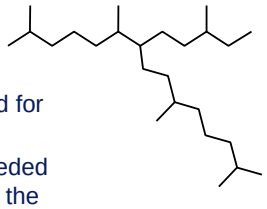
State of the Art: Integration of Lipid Biomarkers & Molecular Biology

DNA and biomarker chemotaxonomy

RNA and DNA sequence analysis of Groups of living organisms enables :

- Evolutionary trees to be constructed.
- Specific Biochemical Pathways to be investigated for their distributions in the Phylogenetic Group.
  - E.g. Identifying the gene for the enzymes needed to synthesise particular biomarkers, such as the highly branched isoprenoid (HBI) compounds.

C_{25} HBI



A prior limitation in assessments of the source specificity of lipid biomarkers is that it has been restricted to studies of cultured organisms

DNA and biomarker chemotaxonomy:

When did the Rhizosolenid Diatoms Evolve?

(Sinninghe Damste et al. 2004 Science,304, 584)

- Rhizosolenid diatoms are a very successful group of marine diatoms. They currently fix around half of the CO_2 flux in the oceans.
- They are the only Group of diatoms to make the HBI Biomarkers
  - 150 Diatom Species analysed for Molecular Phylogeny and HBI Production
  - The role of the HBIs in these Diatoms remains unknown!
- Their Evolutionary History certainly extends back millions of years. ***Therefore, cannot use Fossil DNA but can use Biomarker Distributions***.

DNA and biomarker chemotaxonomy:

When did the Rhizosolenid Diatoms Evolve?

- Genomic analyses of living diatoms gives the Phylogenetic Tree in which the deepest branching points for the Rhizosolenids can be seen.
- Analysis for HBI biomarkers of 81 well-dated petroleums and 700 ancient sediments, going back 0.7 Ma., reveals the first appearance of the HBI at 91.5 Ma in the Upper Turonian.
- So we can date the first appearance of the Rhizosolenids to 91.5 Ma, based on the first occurrence of the HBI biomarkers.

Recovery of Fossil DNA from aquatic sediments

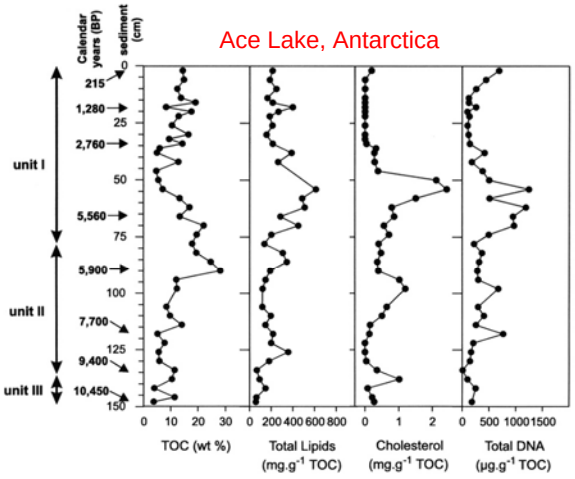


Fig. 3. Depth profiles of total organic carbon (TOC) contents (weight percent), total lipid contents (milligrams per gram TOC), cholesterol as a generic biomarker for overall algal productivity (milligrams per gram of TOC), and extractable DNA (micrograms per gram of TOC). Note that DNA was extractable from even the deepest sediment.

Coolen et al. 2004 EPSL 223, 225-239

Recovery of Fossil DNA from aquatic sediments

Ace Lake, Antarctica

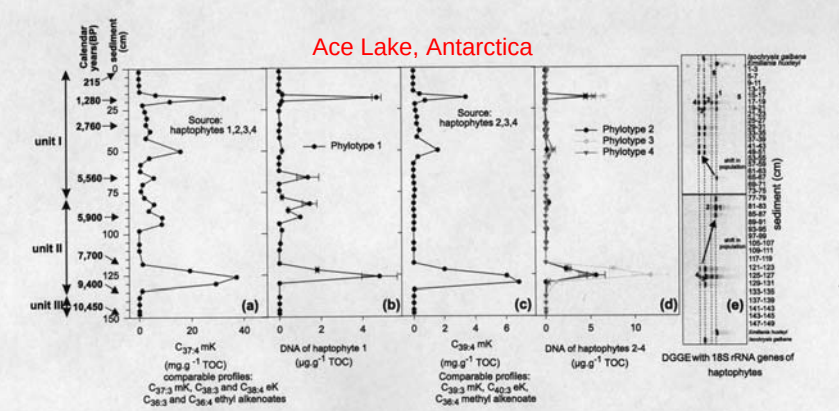


Fig. 4. Stratigraphy of biomarkers of haptophytes (18S rDNA, alkenones [mK, methyl ketone; eK, ethyl ketone], and alkenoates) recovered from the Holocene Ace Lake sediments. (a) Quantity of C_{37}:mK (mg g^{-1} TOC). Comparable profiles were found for $C_{37:3}$ mK, $C_{38:3}$ and $C_{38:4}$ eK as well as C_{36} ethyl alkanoates; (b) Quantity of 18S rDNA of haptophyte 1 (phylotype 1) (µg g^{-1} TOC); (c) quantity of $C_{39:4}$ mK (mg g^{-1} TOC). Comparable profiles were found for $C_{39:3}$ mK, $C_{40:3}$ eK, and a C_{36} methyl alkanoate; (d) quantity of 18S rDNA of haptophytes phylotypes 2–4 (µg g^{-1} sediment); (e) separation of PCR-amplified 18S rDNA of haptophytes from the Holocene sediment layers by DGGE resulted in six fragments with unique melting positions (haptophytes 1–6 numbered in the gel). Phylotypes 5 and 6 (e) were rare and therefore not used for quantitative stratigraphic analysis. The dashed vertical lines indicate identical melting behaviour of phylotypes 1–4 throughout the gel. For comparative analysis, PCR products of reference strains of the haptophytes *I. galbana* CCMP1323 and *E. huxleyi* str. L were separated by DGGE along with the sediment samples. The arrows in the DGGE indicate shifts in the haptophyte populations. Unit III: freshwater lacustrine period. Unit II: due to the rising sea level caused by Holocene deglaciation, marine waters with haptophytes and diatoms entered Ace Lake. Unit I: present-day meromictic saline lacustrine system with euxinic conditions as indicated by molecular remains of green sulfur bacteria (Coolen et al., unpublished results).

Coolen et al. 2004 EPSL 223, 225-239

Recovery of Fossil DNA from aquatic sediments

Ace Lake, Antarctica

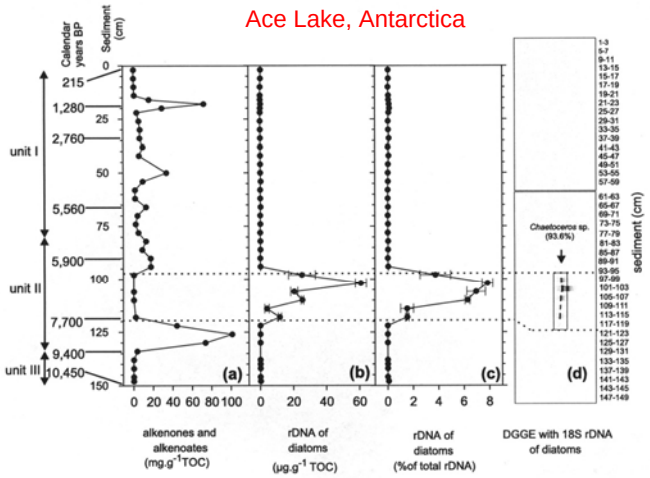


Fig. 6. Abundances of total alkenones and alkenoates (milligrams per gram of TOC) and DNA of diatoms expressed as micrograms per gram of TOC and of total community DNA. DGGE results are shown illustrating the occurrence of high contents of DNA from a diatom related to *Chaetoceros* sp. in the core section deposited after 7700 years BP. Note that this diatom did not become abundant until after the haptophyte populations waned.

Coolen et al. 2004 EPSL 223, 225-239

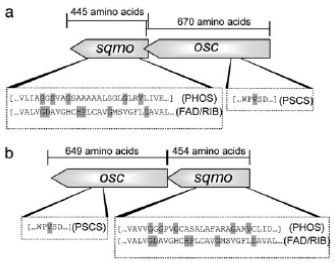


Fig. 1. The sterol biosynthesis genes found in *M. capsulatus* (*a*) and *G. obscuriglobus* (*b*), showing the direction of transcription (arrows) and the critical conserved amino acids (expanded view; critical residues are highlighted). Important functional regions (25) PSCS (protosteryl cation stabilizing), PHOS (pyrophosphate binding), and FAD/RIB (FAD-ribityl binding) are discussed in the text.

Sterol biosynthesis is viewed primarily as a eukaryotic process, and the frequency of its occurrence in bacteria has long been a subject of controversy. Two enzymes, squalene monooxygenase and oxidosqualene cyclase, are the minimum necessary for initial biosynthesis of sterols from squalene. In this work, 19 protein gene sequences for eukaryotic squalene monooxygenase and 12 protein gene sequences for eukaryotic oxidosqualene cyclase were compared with all available complete and partial prokaryotic genomes. The only unequivocal matches for a sterol biosynthetic pathway were in the proteobacterium, *Methylococcus capsulatus*, in which sterol biosynthesis is known, and in the planctomycete, *Gemmata obscuriglobus*. The latter species contains the most abbreviated sterol pathway yet identified in any organism. Analysis shows that the major sterols in *Gemmata* are lanosterol and its uncommon isomer, parkeol. There are no subsequent modifications of these products. In bacteria, the sterol biosynthesis genes occupy a contiguous coding region and possibly comprise a single operon. Phylogenetic trees constructed for both enzymes show that the sterol pathway in bacteria and eukaryotes has a common ancestry. It is likely that this contiguous reading frame was exchanged between bacteria and early eukaryotes via lateral gene transfer or endosymbiotic events. The primitive sterols produced by *Gemmata* suggest that this genus could retain the most ancient remnants of the sterol biosynthetic pathway.

Phylogenetic and biochemical evidence for sterol synthesis in the bacterium *Gemmata obscuriglobus*

Ann Pearson*†, Meytal Budin*, and Jochen J. Brocks‡

15352–15357 | PNAS | December 23, 2003 | vol. 100 | no. 26

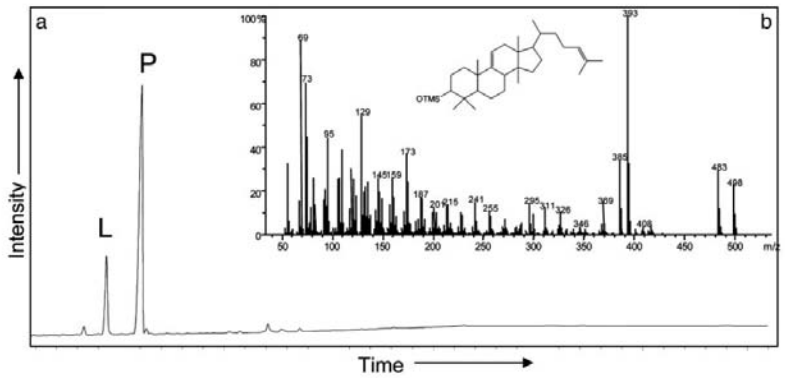


Fig. 2. (*a*) Chromatogram of the sterol region of acid-hydrolyzed whole cells, showing lanosterol (L) and parkeol (P). Minor peaks are nonsteroidal, with the possible exception of the peak immediately to the right of parkeol. (*b*) Structure and mass spectrum (70 eV; 1 eV = 1.602×10^{-19} J) of parkeol as its trimethylsilyl derivative. See Figs. 6 and 7 and Table 3, which are published as supporting information on the PNAS web site.

A polycyclic terpenoid that alleviates oxidative stress

T. Bosak*†, R. M. Losick‡, and A. Pearson§

PNAS | May 6, 2008 | vol. 105 | no. 18 | 6725–6729

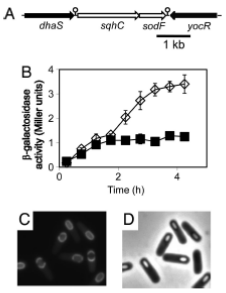


Fig. 1. Expression of *sqhC* and the intracellular localization of SqhC. (*A*) Organization of the *sqhC* operon. Hairpin symbols represent transcriptional terminators. (*B*) Accumulation of colorimetric marker-enzyme β-galactosidase expressed from P*sqhC-lacZ* in a wild-type strain (*amyE*::P_{sqhC}-*lacZ*, TB12; open diamond) and in a strain lacking the sporulation-dependent RNA polymerase subunit σ^F but harboring P*sqhC-lacZ* (*spoIIGA*::*tet*, *amyE*::P_{sqhC}-*lacZ*, TB19; filled square). Samples were collected at the indicated times after the beginning of sporulation (hour 0). Error bars correspond to 1 σ error on the mean value from triplicate samples. (*C*) Fluorescence micrograph of representative cells producing green fluorescent protein (GFP) fused in-frame to the C terminus of SqhC (*amyE*::$P_{Hyspank}$-*sqhC-gfp*, TB29) 4 h after the beginning of the sporulation. The expression of the fusion was driven by 1 mM IPTG 2.5 h after the induction of sporulation. (*D*) The same field of view as in *C* in transmitted light.

Polycyclic terpenoid lipids such as hopanes and steranes have been widely used to understand ancient biology, Earth history, and the oxygenation of the ocean–atmosphere system. Some of these lipids are believed to be produced only by aerobic organisms, whereas others actually require molecular oxygen for their biosynthesis. A persistent question remains: Did some polycyclic lipids initially evolve in response to certain environmental or metabolic stresses, including the presence of oxygen? Here, we identify tetracyclic isoprenoids in spores of the bacterium *Bacillus subtilis*. We call them sporulenes. They are produced by cyclization of regular polyprenes, a reaction that is more favorable chemically than the formation of terpenoids such as hopanoids and steroids from squalene. The simplicity of the reaction suggests that the *B. subtilis* cyclase may be analogous to evolutionarily ancient cyclases. We show that these molecules increase the resistance of spores to a reactive oxygen species, demonstrating a specific physiological role for a nonpigment bacterial lipid biomarker. Geostable derivatives of these compounds in sediments could thus be used as direct indicators of oxidative stress and aerobic environments.

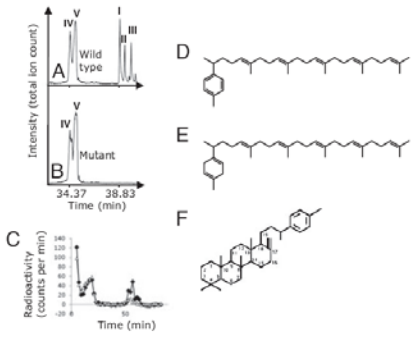


Fig. 2. SqhC produces polycyclic terpenoids. (*A*) Total ion chromatogram of the lipid extract of wild-type spores (PY79). Three conspicuous peaks (I, II, and III) were never present in the extracts of the strain lacking SqhC (Δ*sqhC* Δ*sodF*::*tet*, TB10). (*B*) Acyclic tetraprenyl curcumenes (IV and V) were abundant in the spores of both strains. (*C*) The radioactivity of eighty lipid fractions extracted from the cell lysates of the strain overexpressing SqhC (Δ*sqhC* Δ*sodF*::*tet amyE*::*PHyperspank-sqhC*, TB28; filled diamond) and the strain mutant for *sqhC* (TB10; open triangle) after incubation with ^{3}H-FPP. Fractions derived from TB28 that contained I–III were three times as radioactive as the corresponding fractions derived from TB10 (*). These fractions from TB10 still contained a nonnegligible amount of radioactivity that is probably due to the presence of unidentified polyprenoid lipids other than I–III. (*D*) Structure of acyclic C-35 polyprenoid tetraprenyl-ar-curcumene (IV) detected in the spores of *B. subtilis* both in the presence and the absence of the putative cyclase. (*E*) Structure of acyclic C-35 polyprenoid tetraprenyl-β-curcumene (IV) detected in the spores of *B. subtilis* in the presence and the absence of the putative cyclase. (*F*) Proposed structure of isomer (II) of tetracyclic C-35 polyprenoids. Isomers I–III are detected only in *B. subtilis* spores that contained the putative squalene hopene cyclase.

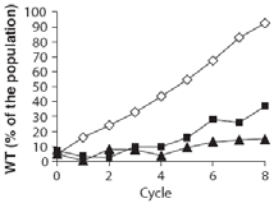
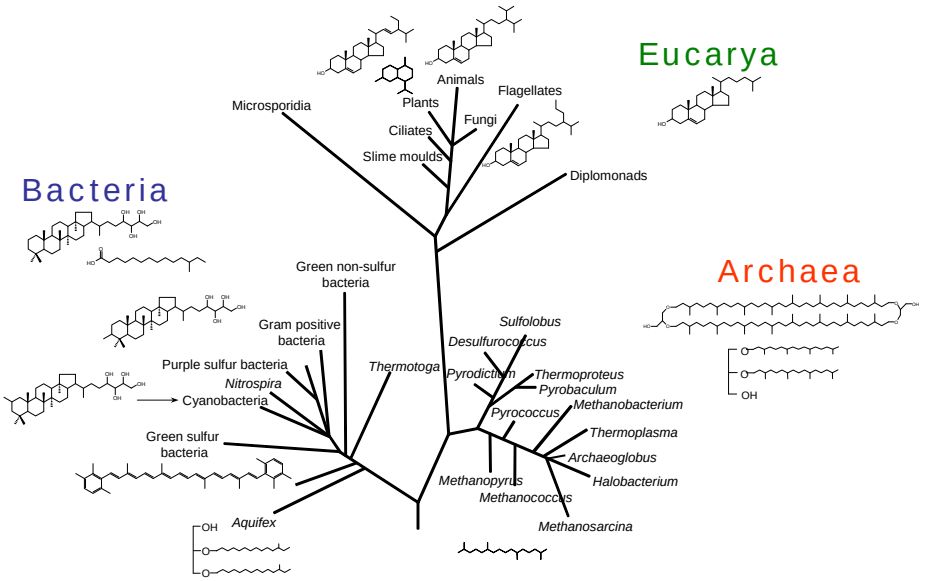


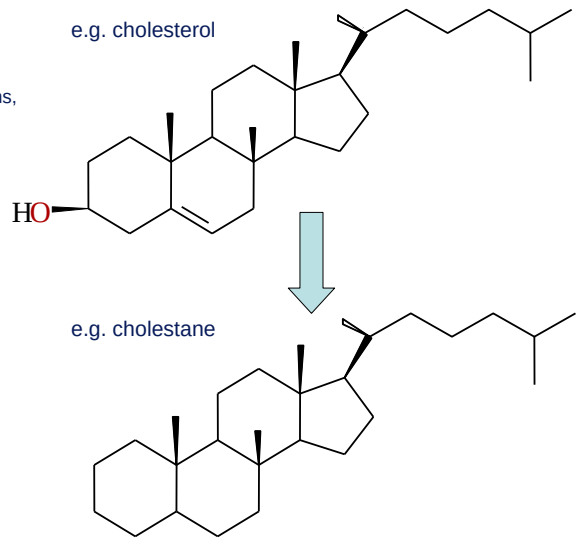
Fig. 3. Spores containing *sqhC* are more resistant to hydrogen peroxide. Wild-type colorimetrically marked cells (*amyE*::P*Spac-lacZ*) (TB38) and cells lacking *sqhC* (Δ*sqhC* Δ*sodF*::*tet*) (TB10) were mixed in 10:90 initial ratio in liquid sporulation medium. Purified spores were treated with heat and hydrogen peroxide. A small aliquot of spores was used to determine the ratio of wild-type and mutant cells in the population and the rest was used to inoculate the next cycle of the experiment for eight consecutive cycles. The treatment included the incubation of spores in the presence of 1% H_2O_2 (open squares) and the absence of H_2O_2 (filled triangles). The same experiment was repeated with a mixture of TB38 cells and cells mutant for *sqhC* and *sodF* that contained a functional copy of *sqhC* (Δ*sqhC* Δ*sodF*::*tet*, *thrC*::P*sqhC-sqhC*) (TB71, filled squares). The observed trend was not influenced by the presence of the gene encoding for the colorimetric marker *lacZ* (Fig. S5).

Summary:
The Biological Precursors…



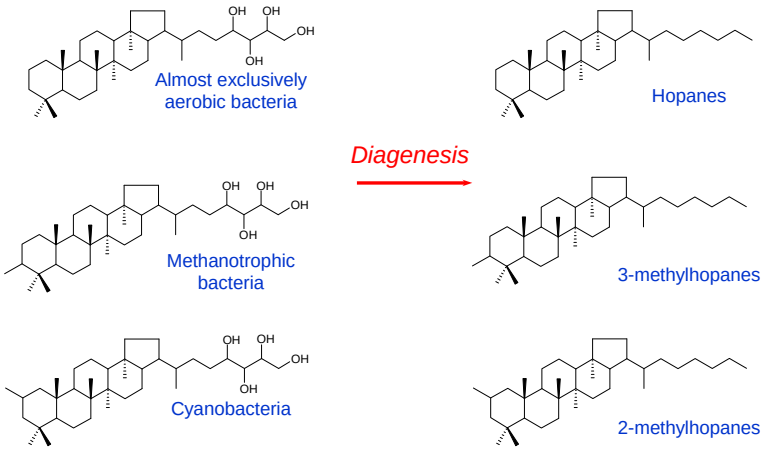
Biomarker Molecules in the Biosphere & Geosphere

Biosphere
Organisms synthesise molecules with highly ordered hydrocarbon skeletons, often carrying several functional groups containing O,N atoms etc.

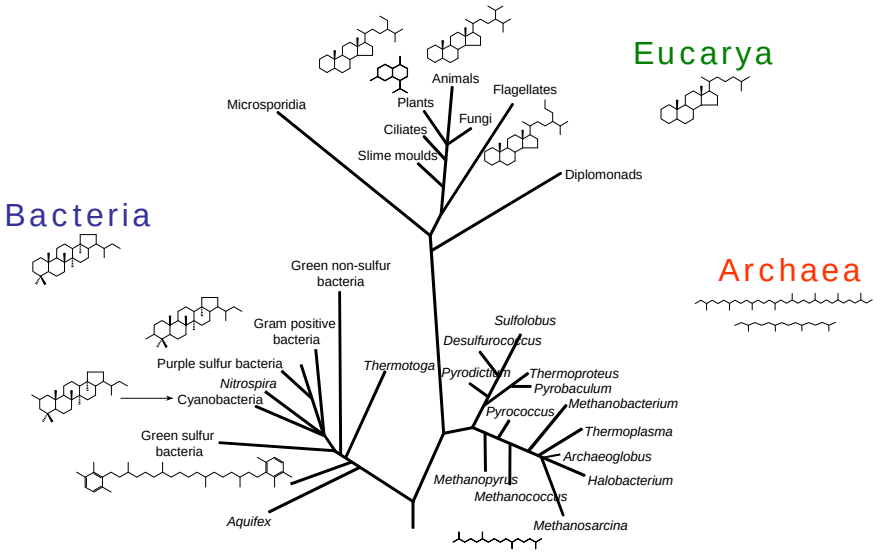


Geosphere
Sedimentary compounds often defunctionalised as parent hydrocarbons with same carbon skeleton.

Hopanoids - The 'Most abundant natural products on Earth?'



…and their Molecular Fossils



rRNA tree modified after Woese

The Microbial Record In The Geosphere