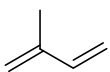


# Topics

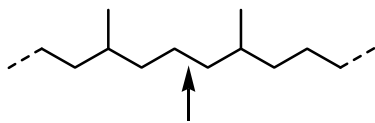
## Complex lipids of extant archaea

- Structural peculiarities of archaeal lipids
- LCMS methods
- Molecular Signatures of HT Methanogens
- Low temperature Chrenarchaea
- *Ignicoccus* sp., *Nanoarchaeum equitans*

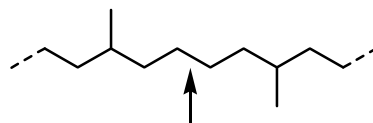
# Common Acyclic Isoprenoids



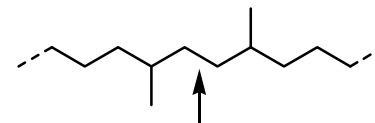
**isoprene**



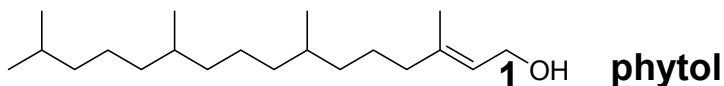
head-to-tail



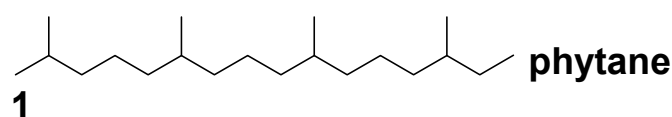
tail-to-tail



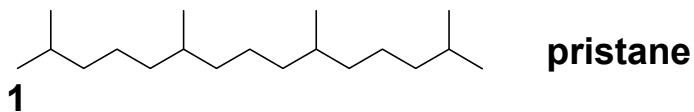
head-to-head



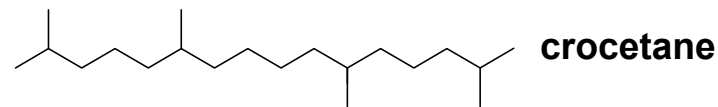
**phytol**



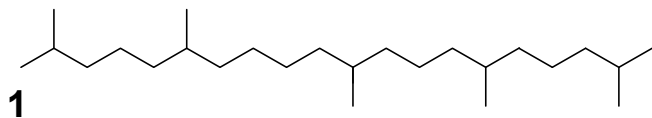
**phytane**



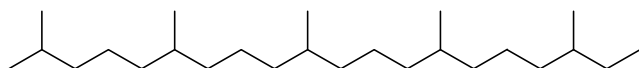
**pristane**



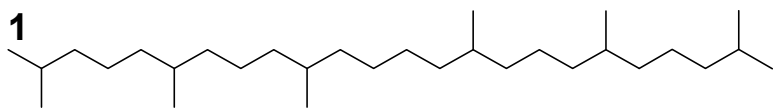
**crocetane**



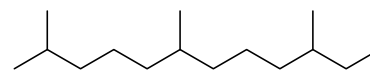
**2,6,10,15,19-pentamethylcosane (PMI)**



**2,6,10,14,18-pentamethylcosane**

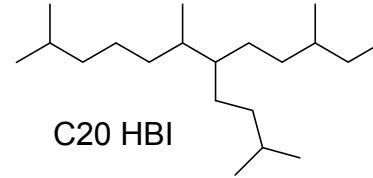
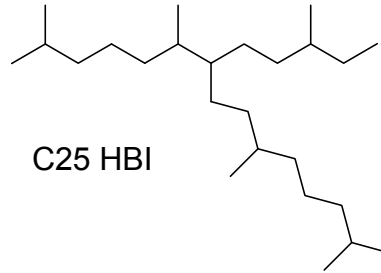
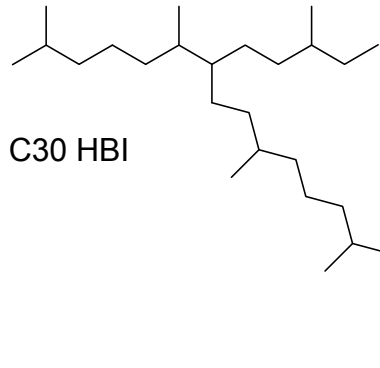


**squalane**



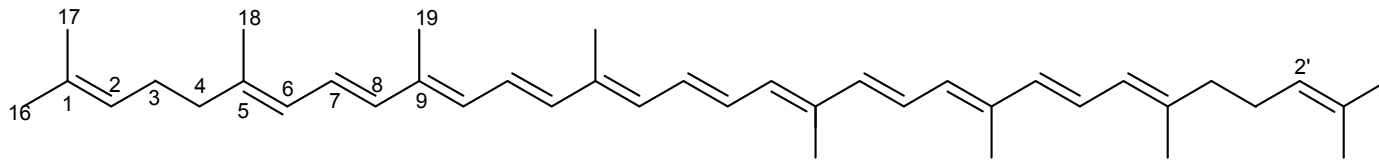
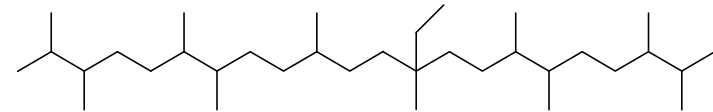
**farnesane**

# Less Common Acyclic Isoprenoids



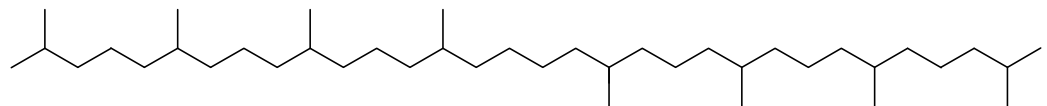
Diatom sources

*Botryococcus braunii*

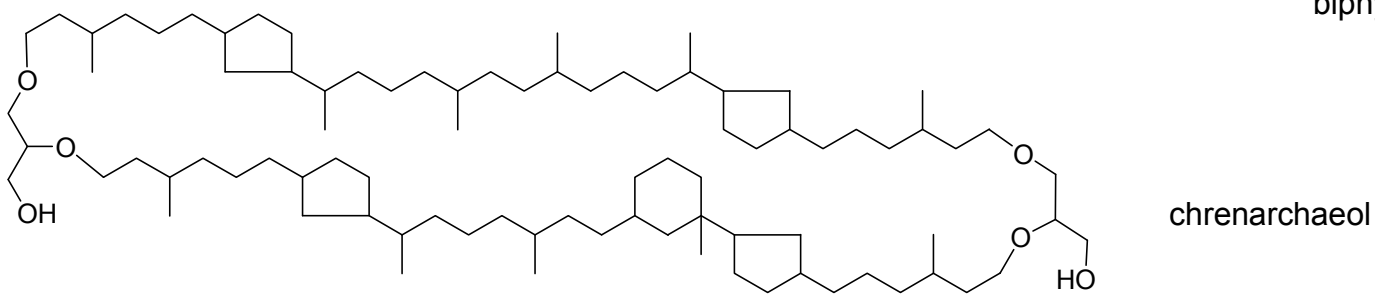
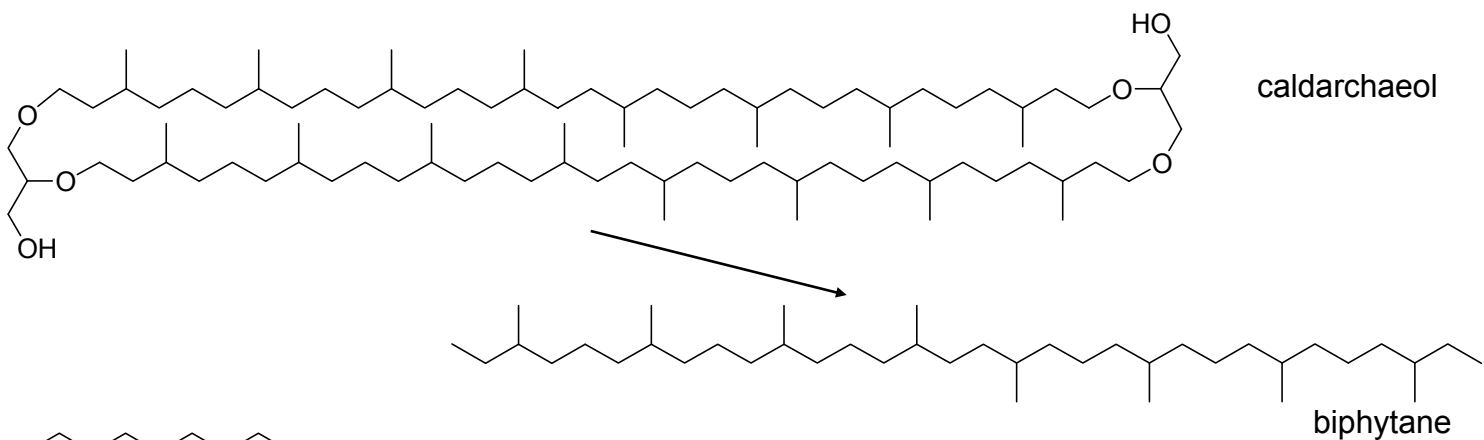
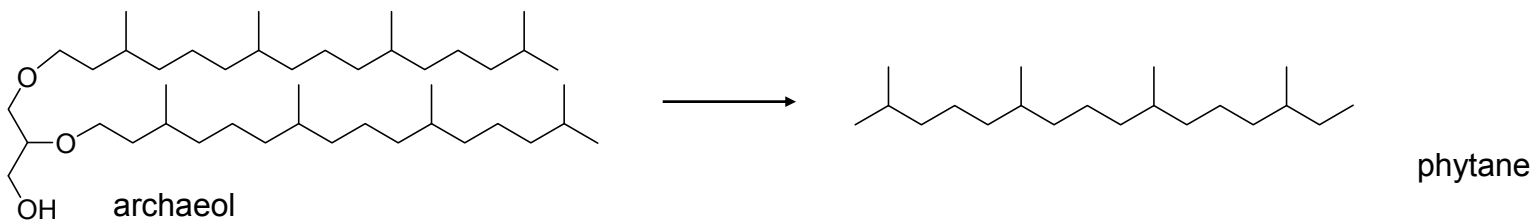


Probable algal hydrocarbon  
? from lycopodiene

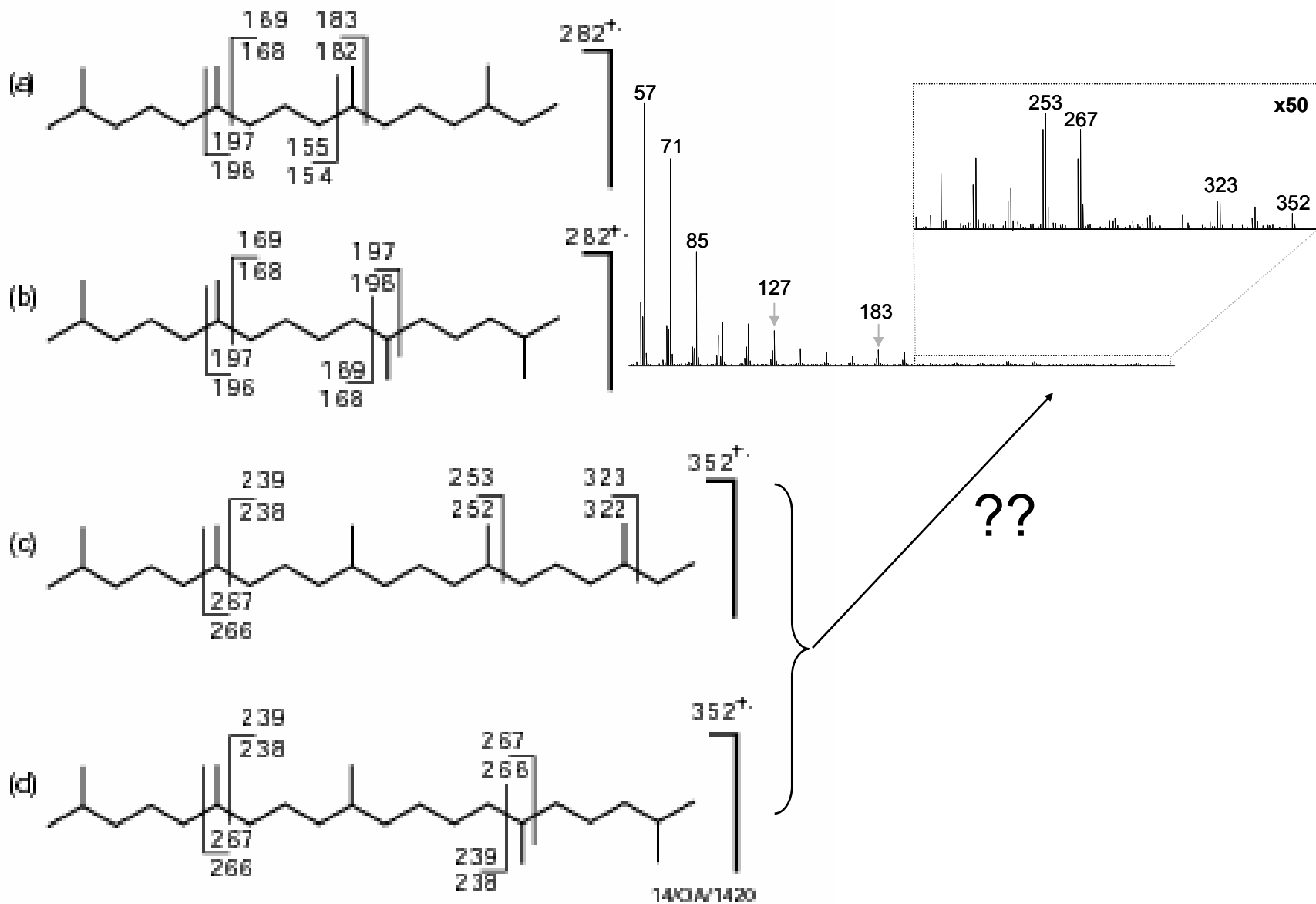
lycopane



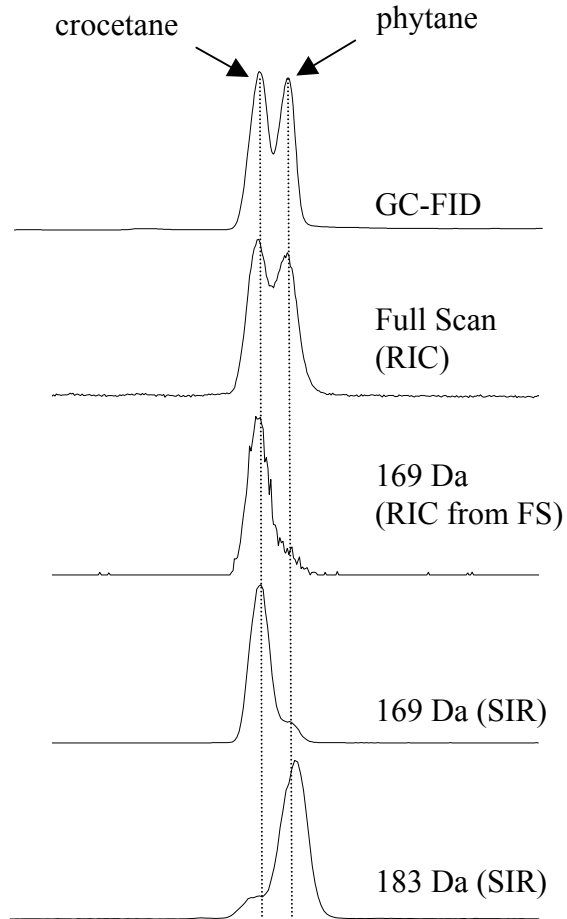
# Polar Lipid Precursors of Acyclic Isoprenoids



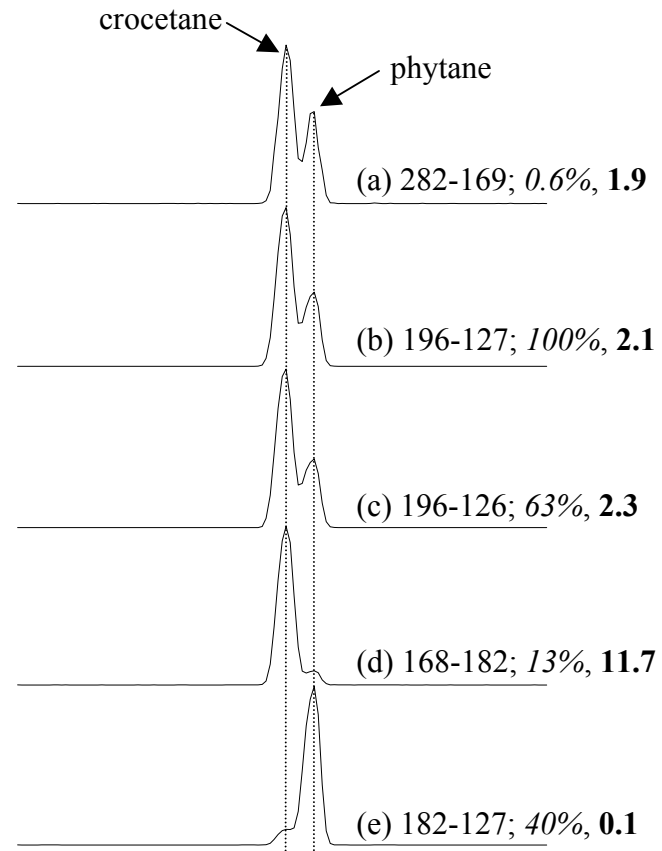
# Favored Mass Spectrometric Fragmentations



# Crocetane – Phytane Distinction



GC and GC-MS (SIR)



GC-MS-MS

# Crocetane – Phytane Distinction

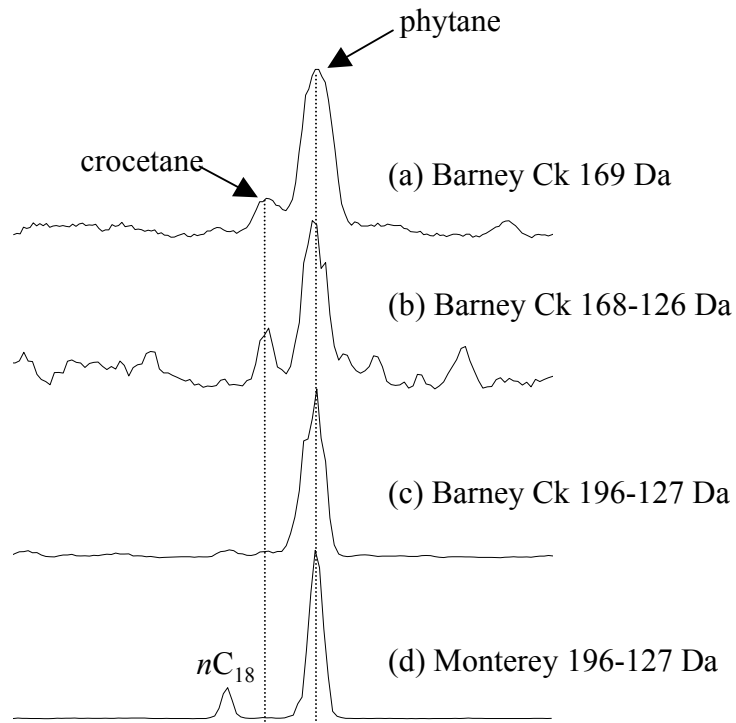
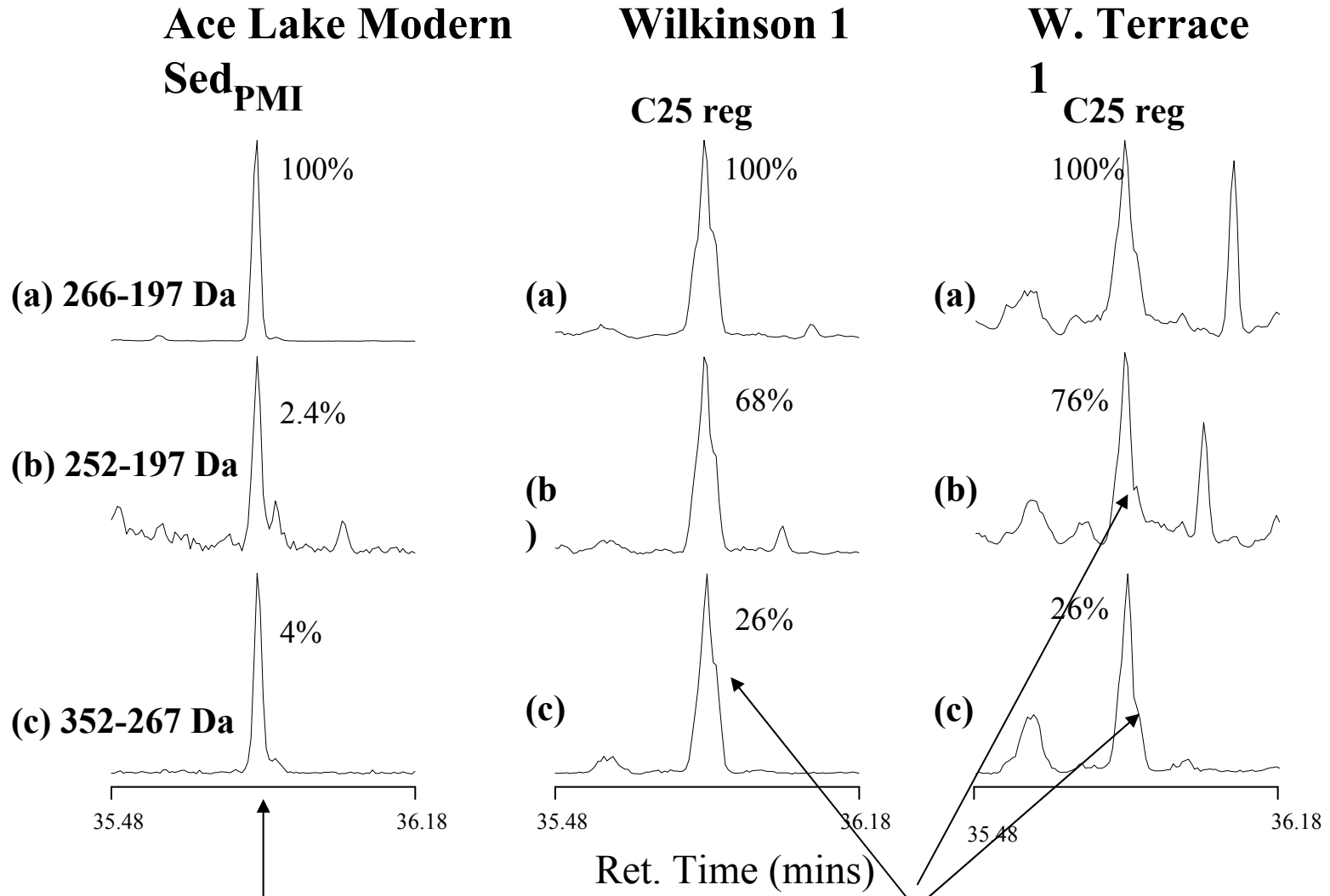


Figure 4

# Regular C<sub>25</sub> vs PMI Distinction

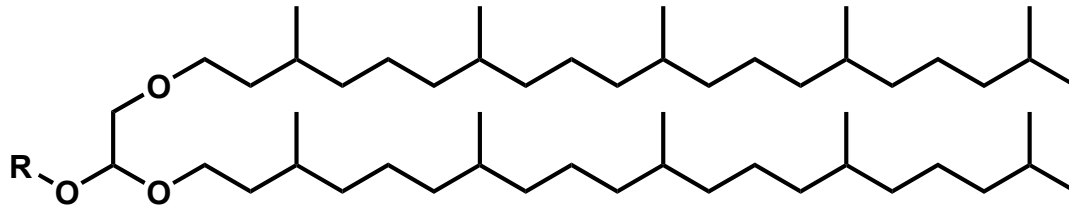


One thin peak+ one compound

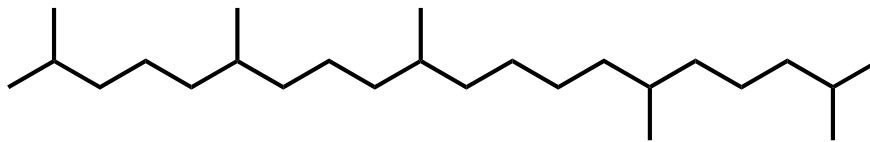
All fat peaks = more than one compound



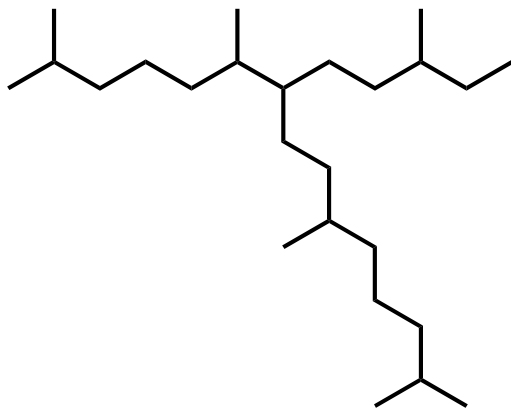
# Regular C<sub>25</sub> vs PMI Distinction



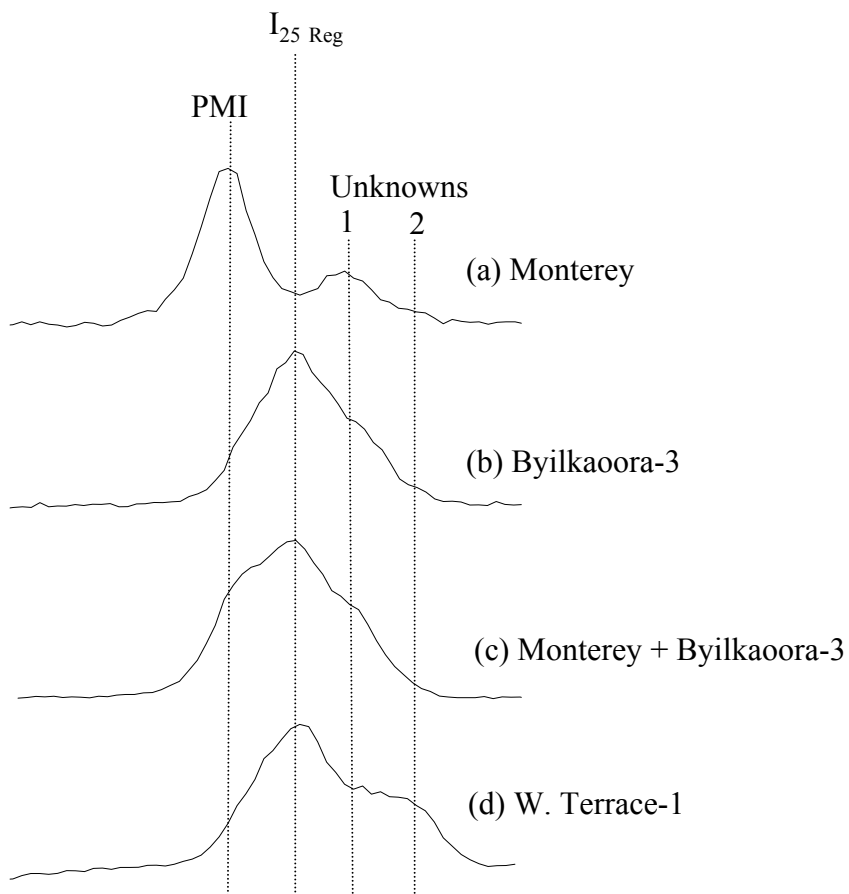
2,6,10,14,14-pentamethylcosane  
Carbon chains of *Halobacterium* core lipid



2,6,10,15,19-pentamethylcosane (PMI)  
Found as a free hydrocarbon in some methanogens



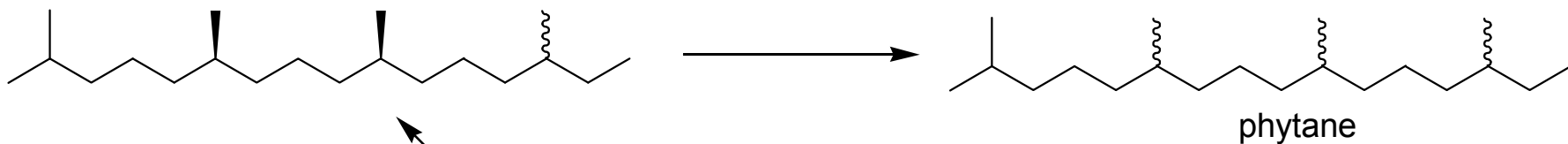
A 'highly branched isoprenoid' (HBI)  
from a diatom



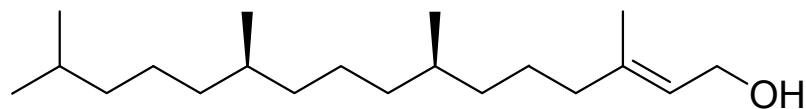
## Distinguishing C25 Isoprenoids

note peak shapes

Partial 183 Da (SIR) chromatograms of (a) Monterey Formation showing elution position of PMI; (b) Byilkaora-3 showing elution position of I<sub>25</sub> reg; (c) Monterey + Byilkaora-3 mixture showing relative elution order of PMI and I<sub>25</sub> reg isomers (NB. only partially resolved); (d) West Terrace-1 which has a peak at the same position as the I<sub>25</sub> reg isomer and no peak at the earlier retention time of PMI. Unknown peaks 1 (Monterey) and 2 (West Terrace-1) elute after I<sub>25</sub> reg. Chromatogram time range = 36 sec.

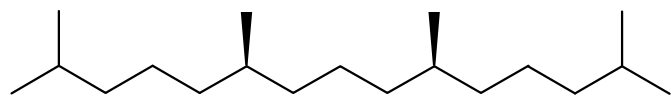


reduction/dehydration/reduction



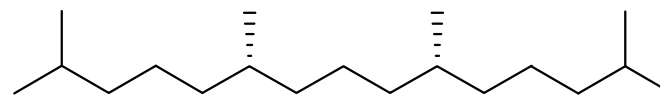
E-3, 7R, 11R, 15-tetramethylhexadec-2-enol = phytol

oxidation/decarboxylation/reduction

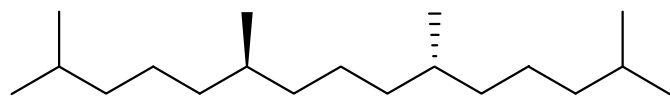


6(R), 10(S) - pristane

=

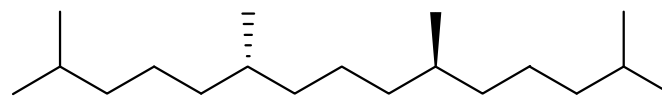


6(S), 10(R) - pristane

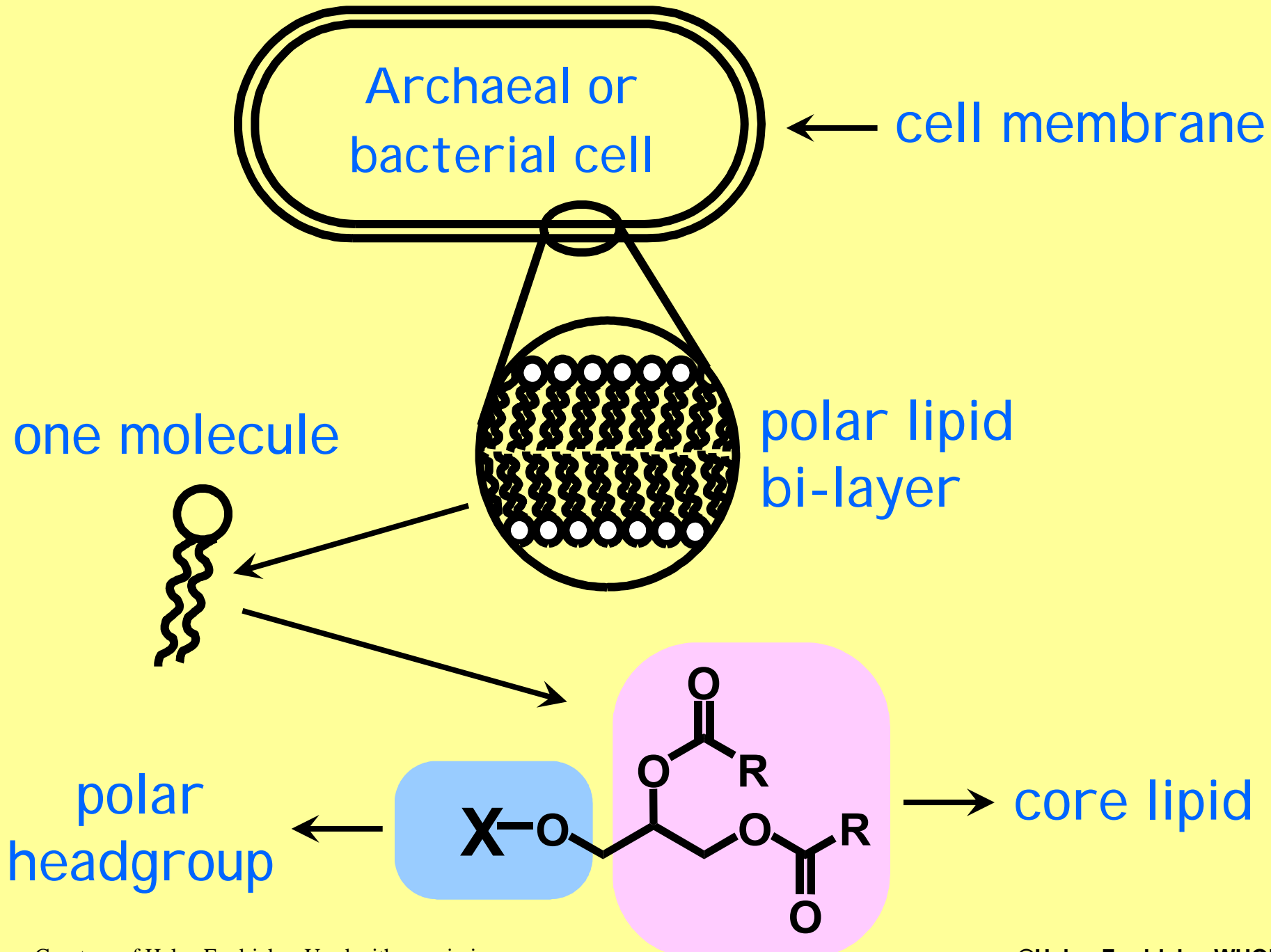


6(R), 10(R) - pristane

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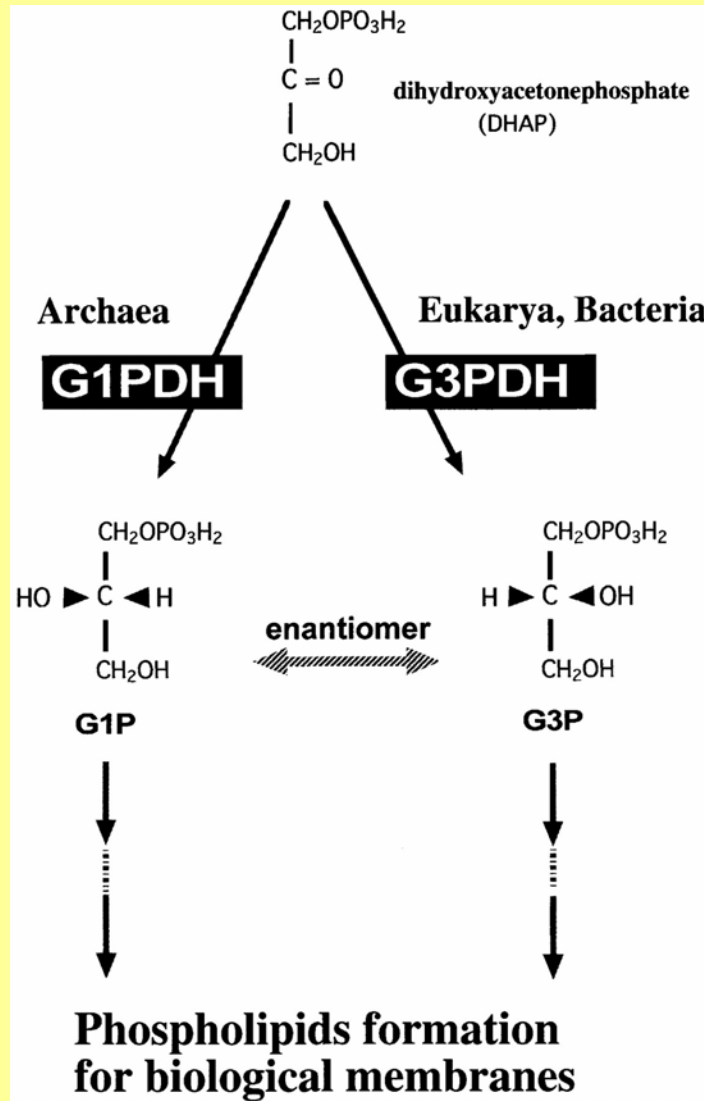


6(S), 10(S) - pristane



# Glycerol Stereochemistry ?

*sn*-glycerol-1-phosphate dehydrogenase (G1PDH)



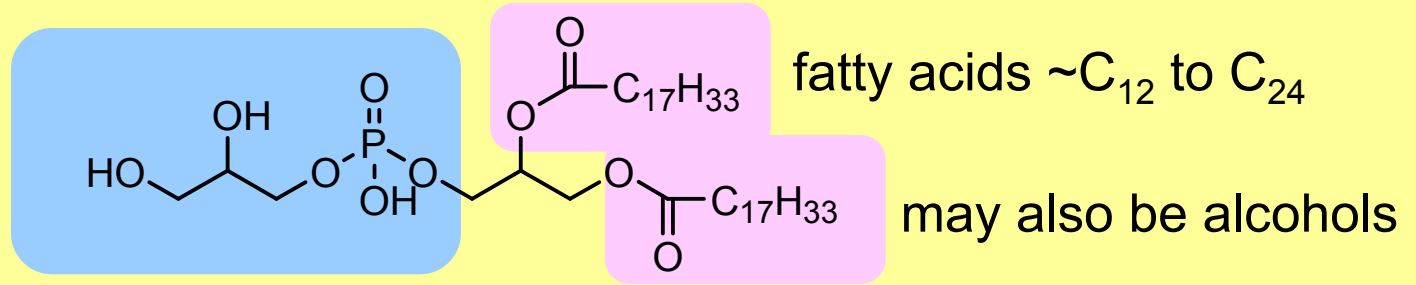
*sn*-glycerol-3-phosphate dehydrogenase (G3PDH)

# I PL variety analyzed by HPLC-ESI-MS<sup>n</sup>

## bacterial

10+ different headgroups identified...

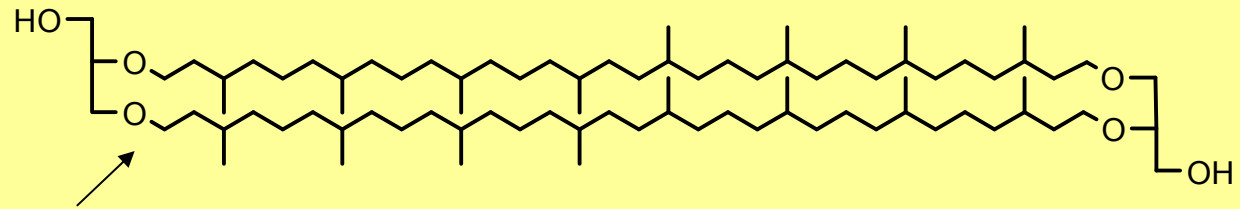
phospholipids & glycolipids, some highly specific



## archaeal

similar variety of headgroups but highly specific core lipids

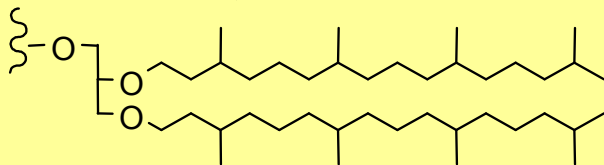
2 basic types with some variations



archaeol

caldarchaeol (GDGT: glycerol dialkyl glycerol tetraether)

Data on I PL distributions in biogeochemically relevant prokaryotes are very sparse



# *Ignicoccus islandicus* & *Ignicoccus pacificus*

**Sources of samples.** At the Kolbeinsey Ridge, north of Iceland, eight samples of submarine sandy sediments and venting water (original temperatures around 90 °C) were taken by the research submersible 'Geo' at depths between 103 and 106 m (Fricke *et al.*, 1989; Burggraf *et al.*, 1990).

Furthermore, black smoker samples were obtained during dive 3072 of the submersible 'Alvin' at the East Pacific Rise at 9° N, 104° W at a depth of 2500 m.

Arch Microbiol (2004) 182: 404-413  
DOI 10.1007/s00 203-004-0725-x

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ORIGINAL PAPER

**Ulrike Jahn, Roger Summons, Helen Sturt  
Emmanuelle Grosjean, Harald Huber**

**Composition of the lipids of *Nanoarchaeum equitans* and their  
origin from its host *Ignicoccus* sp. strain KIN4/I**

Received: 22 January 2004 / Revised: 30 June 2004 / Accepted: 5 August 2004 /  
Published online: 14 September 2004 © Springer-Verlag 2004





## **Genome Structure**

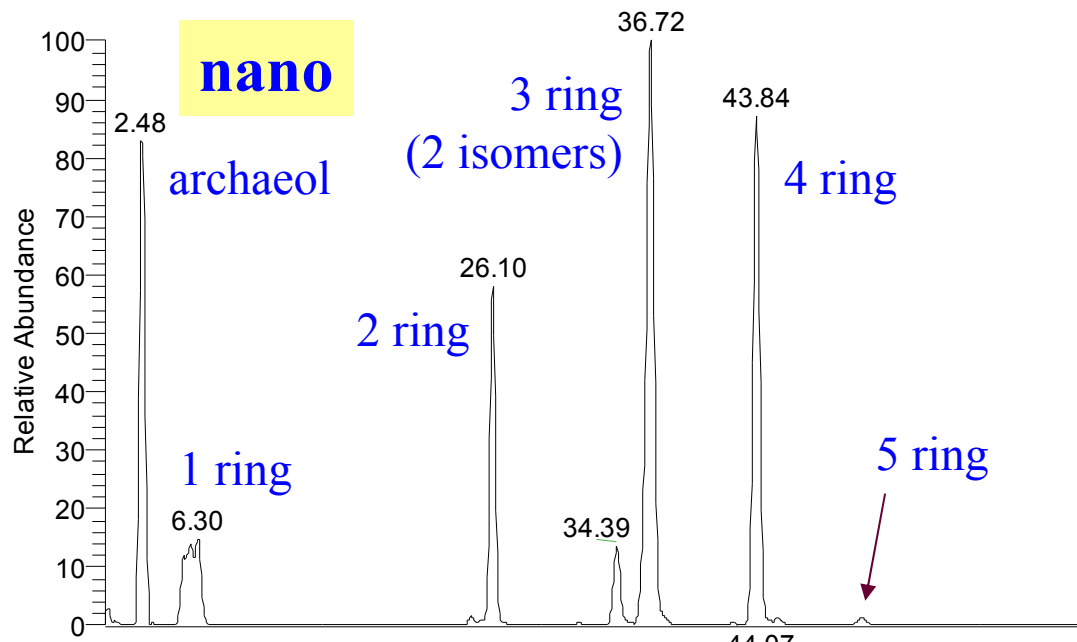
*Nanoarchaeum* has a genome with only 490 kb, which represents the smallest archaeal genome to date. Comparing ss rRNA sequences, it was noted that sequence identities were more like archaeon than bacterial species. There was no difference, however, in the sequence identity to the Crenarchaeota, Euryarchaeota, and 'Korarchaeota', indicating it represents a new archaeal phylum.

## **Cell Structure and Metabolism**

These coccus cells are only 400 nm in diameter and are covered by an S-layer. They require cell-cell contact with an actively growing *Ignicoccus* cell in order to grow.

Other nanoarchaeotal 16S rRNA genes have been obtained from the East Pacific Rise (pH 6.5), the Obsidian Pool in Yellowstone National Park (80°C, pH 6.0), and Caldera Uzon in Kamchatka, Russia (85°C, pH 5.5).

RT: 0.00 - 66.00 SM: 7B



NL: 9.84E7

Base Peak m/z=

670.0-671.0+832.0-833.0+

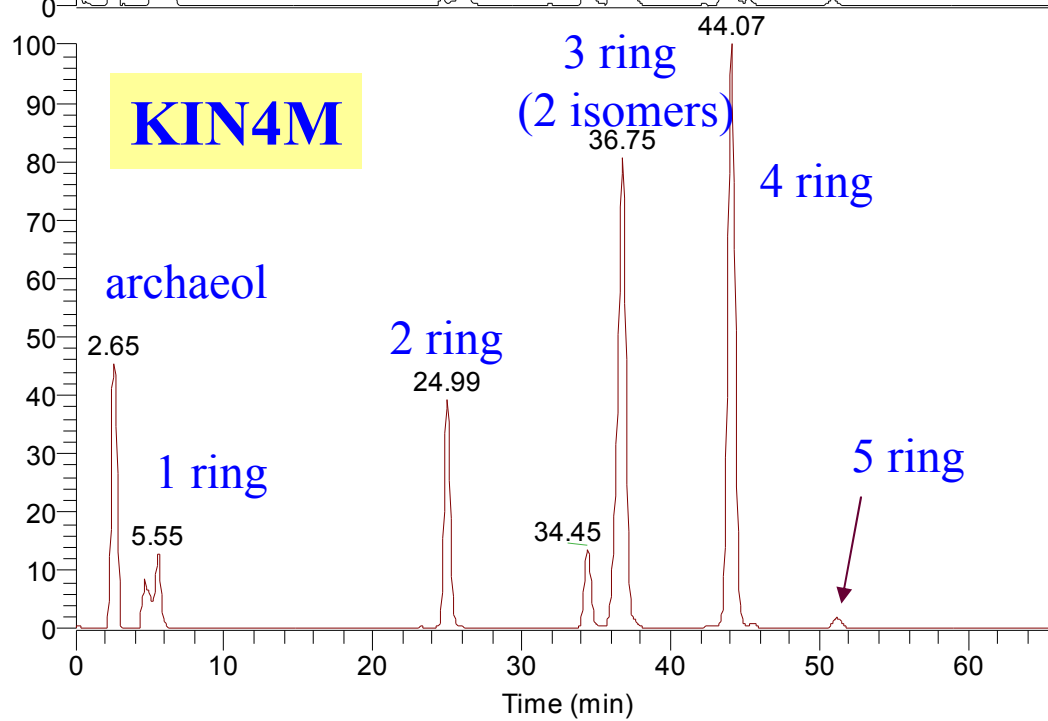
994.0-995.0+1156.0-1157.0+

1318.0-1319.0+1480.0-1481.0 F: + c

Full ms [ 500.00-2000.00] MS

Nano1pos 65 min

**this is an extracted ion chromatogram showing only archaeol and the glyco caldarchaeols**



NL: 1.77E8

Base Peak m/z=

670.0-671.0+832.0-833.0+

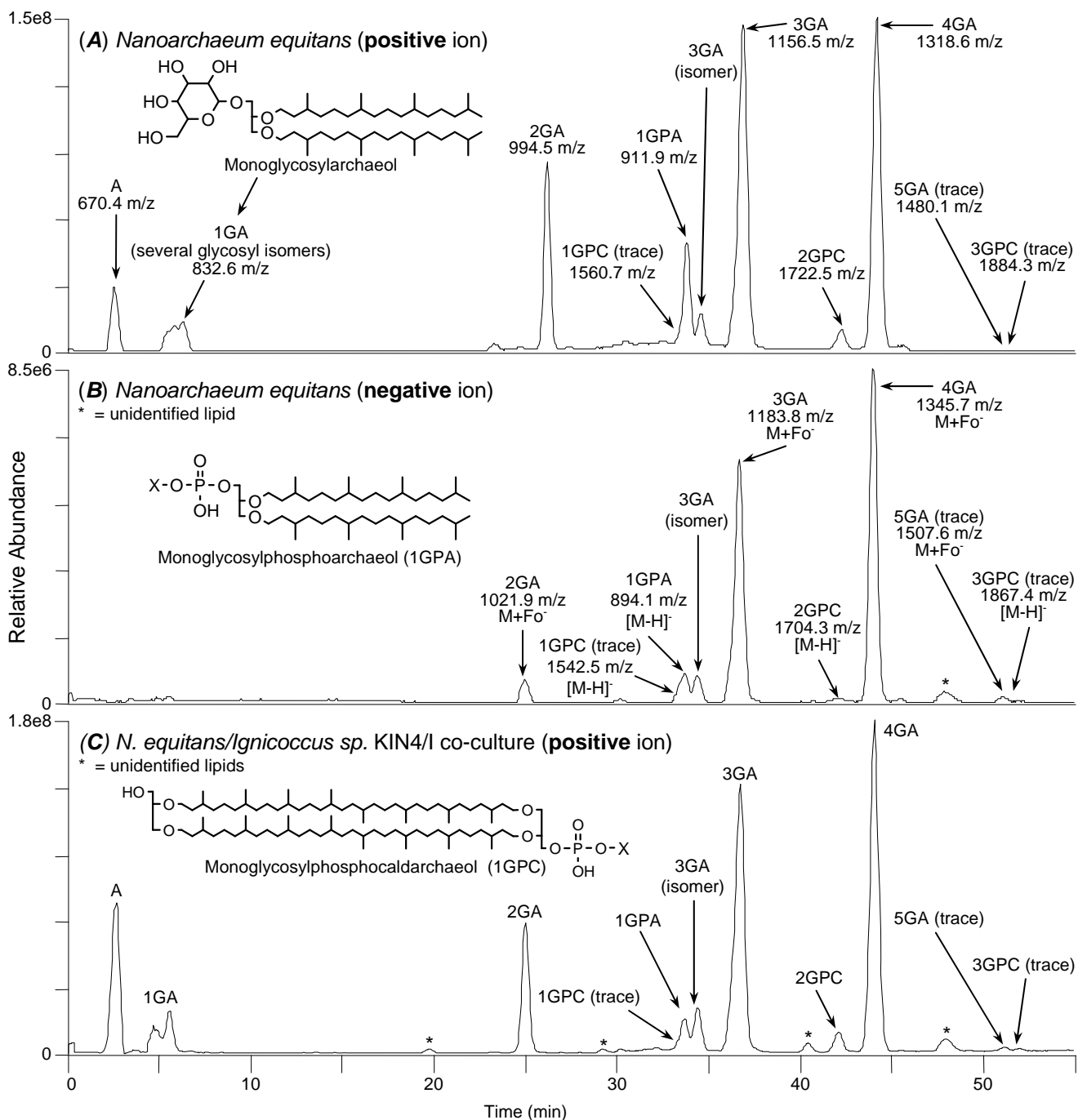
994.0-995.0+1156.0-1157.0+

1318.0-1319.0+1480.0-1481.0 F: + c

Full ms [ 500.00-2000.00] MS

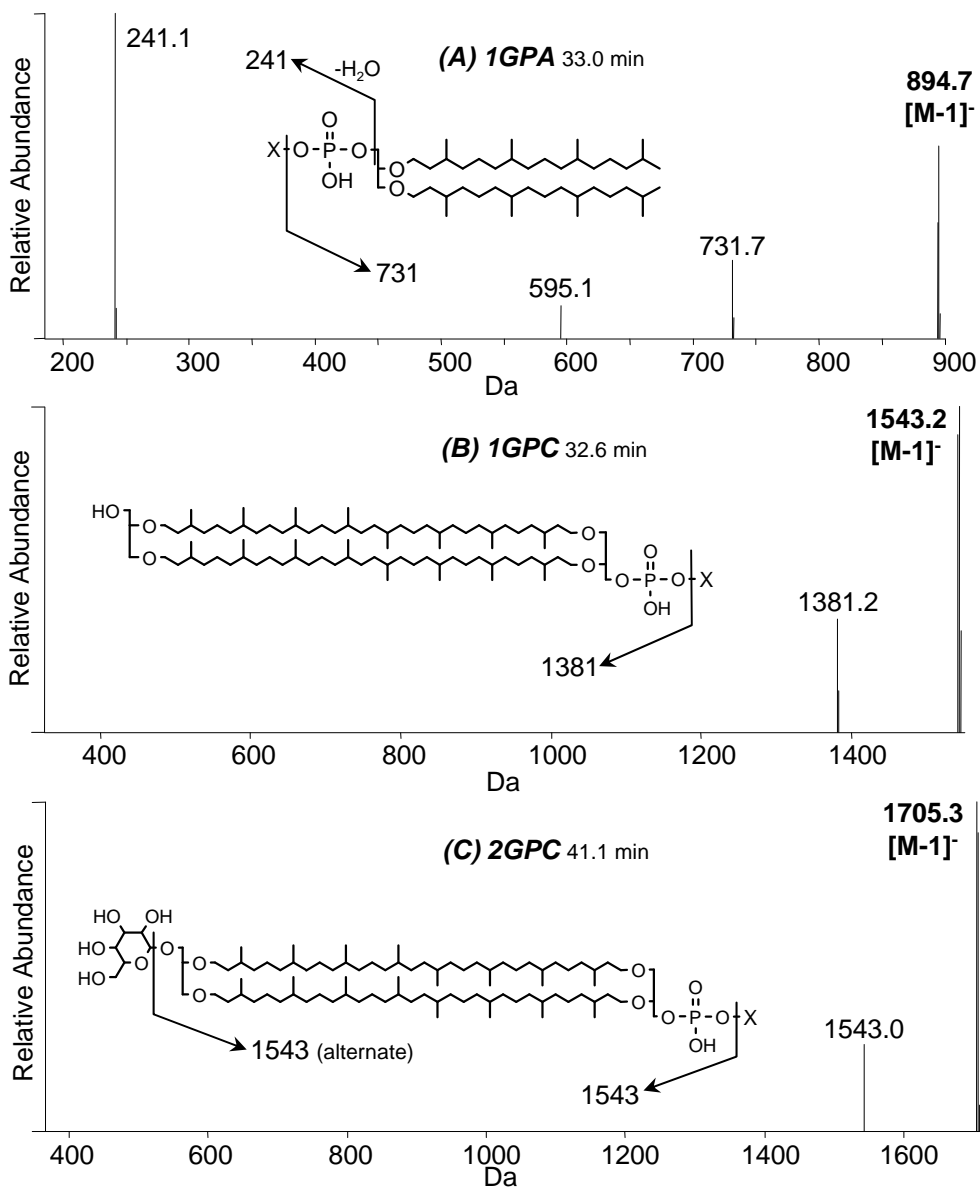
KIN4Mpos 65 min

**positive ion mode  
65 minute run  
time**



**Fig. 2** Base peak chromatograms of full-scan (650-2,000  $m/z$ ) mass spectra of

*N. equitans* (**a** positive ion; **b** negative ion) and a *N. equitans/ignicoccus* sp. strain KIN4/I co-culture (**c** positive ion). Identifications were made on the basis of MS<sup>N</sup> spectra in both positive and negative ion modes from LC-MS and from direct infusion of total lipid extracts. Peaks are labeled as follows: *I-5* number of glycosyl units, *G* glycolipid, *GP* phospholipid, *A* archaeol lipid core, *C* caldarchaeol lipid core and \* unidentified lipid. The dominant lipids are glycolipids with an archaeol core, with smaller amounts of phospholipids which consist of a phosphate group with glycosyl headgroup and either an archaeol or caldarchaeol lipid core.



**Fig. 4** Negative ion mode MS/MS spectra of three glycosylbearing phospholipid species observed in the total lipid extracts of both *N. equitans* (shown above) and *Ignicoccus* sp. strain KIN4/I. Unlike glycolipids, these phospholipids do not form formate adducts and are directly observed as [M-H]<sup>-</sup> ions. Spectrum a is identified as a glycosylphosphoarchaeol (at 33 min in Fig. 2), since the 241-Da ion

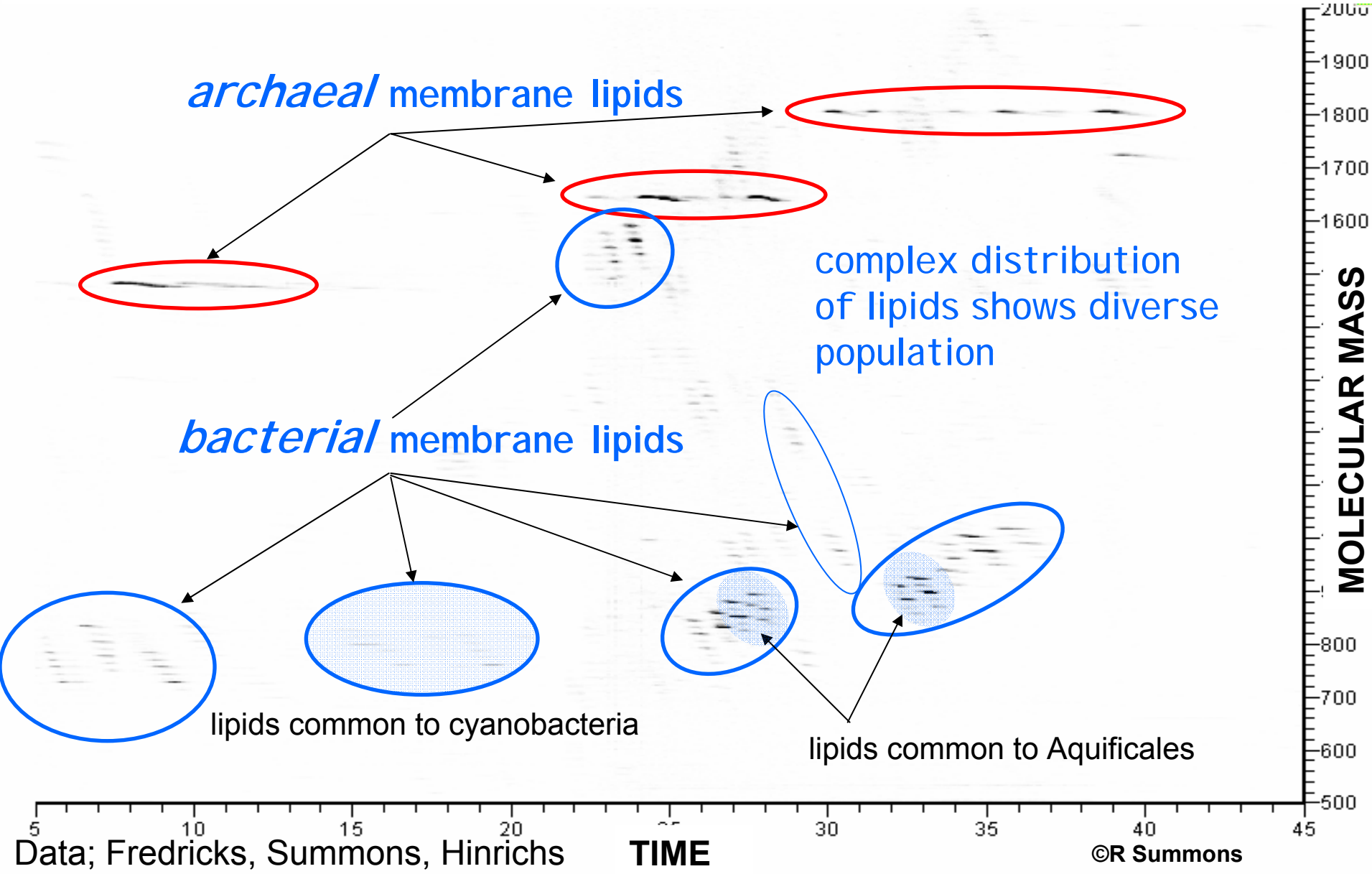
is diagnostic for phosphoinositol lipids and corresponds to the dehydrated glycoposphate headgroup. The 731-Da ion corresponds to an archaeol phosphate ion and is observed in phosphoarchaeol standards with different headgroups. The exact structure of the glycosyl unit attached to the phosphate is unknown since these  $MS^N$  analyses do not give detailed structural information on the nature of the glycosyl units. Spectra **b**, **c** are identified as phosphoglycosyl caldarchaeols with one phosphate and one (at 32.6 min in Fig. 2) or two (at 41.1 min in Fig. 2) glycosyl groups, respectively, attached either via the phosphate group or the glycerol moiety.

# Co-culture Isotopes

	<b>phytane</b>	<b>biphytane</b>
<b>nano</b>	-16.1185	-16.0045
<b>igni</b>	-16.023	-16.297
<b>igni 77C</b>	-15.219	-15.249
<b>igni 90C</b>	-42.087	-43.603
<b>igni 95C</b>	-18.623	-19.767



# Yellowstone National Park 'Ojo Caliente' hot spring sample of biomass from a 'streamer' community



APPLIED AND ENVIRONMENTAL MICROBIOLOGY,  
0099-2240/97/\$04.0010 Jan. 1997, p. 50–56 Vol. 63, No. 1  
Copyright © 1997, American Society for Microbiology

## **Vertical Distribution and Phylogenetic Characterization of Marine Planktonic *Archaea* in the Santa Barbara Channel**

RAMON MASSANA,<sup>1</sup> ALISON E. MURRAY,<sup>2</sup> CHRISTINA M. PRESTON,<sup>2</sup> AND EDWARD F. DELONG<sup>2\*</sup>

*Marine Science Institute*<sup>1</sup> and *Department of Ecology, Evolution and Marine Biology*,<sup>2</sup>*University of California, Santa Barbara, Santa Barbara, California 93106*

Received 1 August 1996/Accepted 11 October 1996

**Newly described phylogenetic lineages within the domain *Archaea* have recently been found to be significant components of marine picoplankton assemblages. To better understand the ecology of these microorganisms, we investigated the relative abundance, distribution, and phylogenetic composition of *Archaea* in the Santa Barbara Channel. Significant amounts of archaeal rRNA and rDNA (genes coding for rRNA) were detected in all samples analyzed.**

# A Few Cosmopolitan Phylotypes Dominate Planktonic Archaeal Assemblages in Widely Different Oceanic Provinces

RAMON MASSANA,<sup>1\*</sup> EDWARD F. DELONG,<sup>2</sup> AND CARLOS PEDRO´ S-ALIO´ 1

*Institut de Cie`ncies del Mar, CSIC, 08039 Barcelona, Catalunya, Spain,<sup>1</sup> and Monterey Bay*

*Aquarium Research Institute, Moss Landing, California 950392*

Received 4 October 1999/Accepted 2 February 2000

**We compared the phylogenetic compositions of marine planktonic archaeal populations in different marine provinces.....**

## **letters to nature**

*Nature* **371**, 695 - 697 (2002); doi:10.1038/371695a0

### **High abundance of Archaea in Antarctic marine picoplankton**

Edward F. DeLong, Ke Ying Wu, Barbara B. Prézelin & Raffael V. M. Jovine

**ARCHAEA (archaebacteria) constitute one of the three major evolutionary lineages of life on Earth<sup>13</sup>. Previously these prokaryotes were thought to predominate in only a few unusual and disparate niches, characterized by hypersaline, extremely hot, or strictly anoxic conditions<sup>47</sup>. Recently, novel (uncultivated) phylotypes of Archaea have been detected in coastal<sup>8</sup> and subsurface<sup>9,10</sup> marine waters, but their abundance, distribution, physiology and ecology remain largely undescribed. Here we report exceptionally high archaeal abundance in frigid marine surface waters of Antarctica. Pelagic Archaea constituted up to 34% of the prokaryotic biomass in coastal Antarctic surface waters, and they were also abundant in a variety of other cold, pelagic marine environments. Because they can make up a significant fraction of picoplankton biomass in the vast habitats encompassed by cold and deep marine waters, these pelagic Archaea represent an unexpectedly abundant component of the Earth's biota.**

APPLIED AND ENVIRONMENTAL MICROBIOLOGY, 0099-2240/98/\$04.0010  
Mar. 1998, p. 1133–1138 Vol. 64, No. 3 Copyright © 1998, American Society for  
Microbiology

## **Dibiphytanyl Ether Lipids in Nonthermophilic Crenarchaeotes**

EDWARD F. DELONG,<sup>1\*</sup> LINDA L. KING,<sup>2</sup> RAMON MASSANA,<sup>1</sup> HENRY  
CITTONE,<sup>1</sup> ALISON MURRAY,<sup>1</sup> CHRISTA SCHLEPER,<sup>1</sup> AND STUART G.  
WAKEHAM<sup>2</sup>

*The Marine Science Institute, University of California, Santa Barbara, California  
93106,<sup>1</sup> and Skidaway Institute of Oceanography, Savannah, Georgia 31411<sup>2</sup>*

Received 30 September 1997/Accepted 23 December 1997

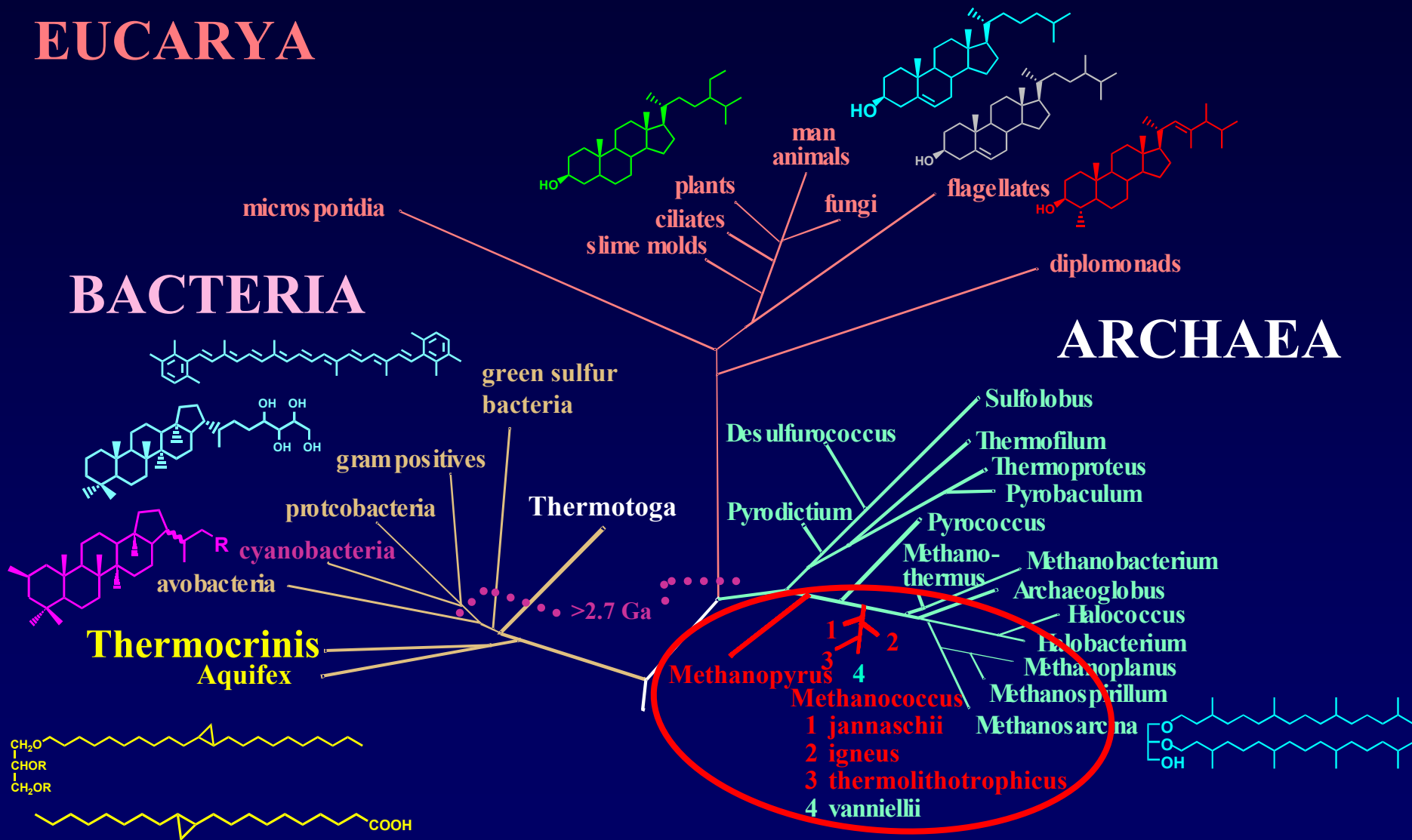
**The kingdom *Crenarchaeota* is now known to include archaea which inhabit a wide variety of low-temperature environments. We report here lipid analyses of nonthermophilic crenarchaeotes, which revealed the presence of cyclic and acyclic dibiphytanylglycerol tetraether lipids. Nonthermophilic crenarchaeotes appear to be a major biological source of tetraether lipids in marine planktonic environments.**

# Parallel Molecular Signatures

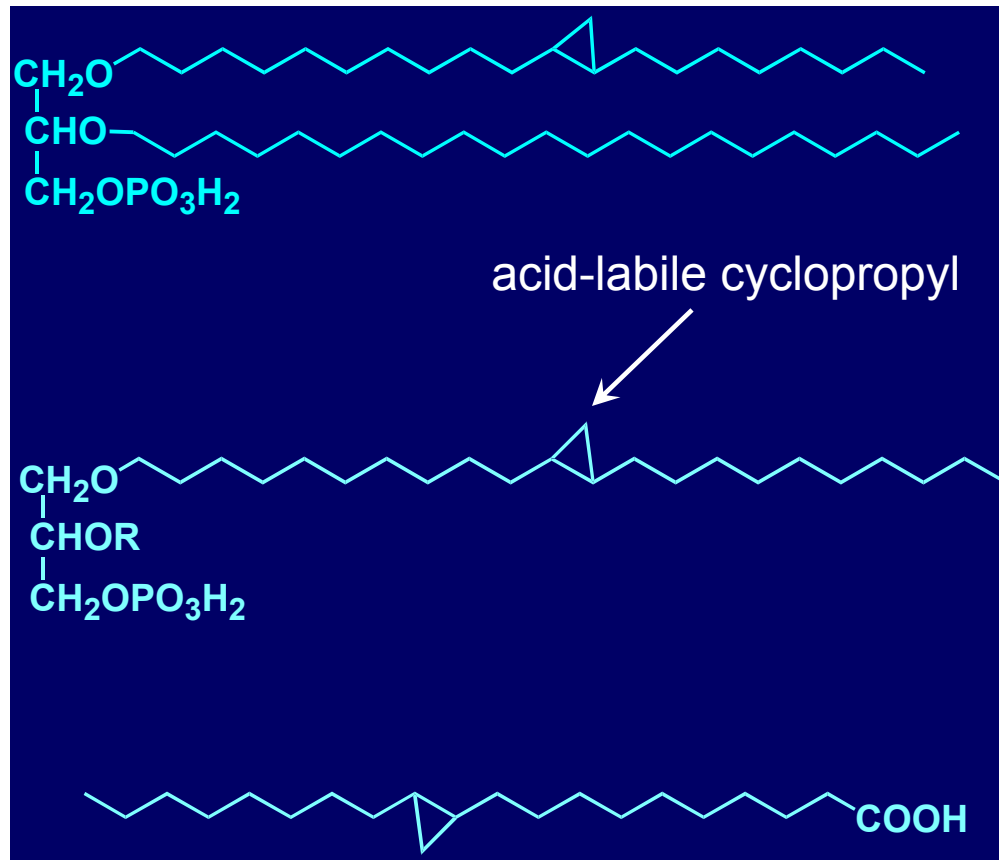
## EUCARYA

## BACTERIA

## ARCHAEA



# Acid-labile cyclopropyl lipids from Aquificales



**Alkaline methanolysis OK for FAME but Mild Acid Hydrolysis needed to liberate ether lipids**