

Archaea - Exploration of Thermophilic Adaptations

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- ▶ Historically, living organisms were categorized into five kingdoms
 - ▶ Plantae, Animalia, Fungi, Protista and Monera (Eubacteria)
- ▶ 1977 - Carl Woese identifies third form of life
 - ▶ The Archaea (prokaryotic)
 - ▶ Primary evidence lies in rRNA sequence patterns
 - ▶ Resulted in elimination of Monera Kingdom

- ▶ Archaea are generally categorized into three functional groups
 - ▶ Methanogens
 - ▶ Unique form of carbon fixation (reduction) utilizing CO or CO₂ to produce methane
 - ▶ Halophiles
 - ▶ Thermophiles
- ▶ May be extremely misleading

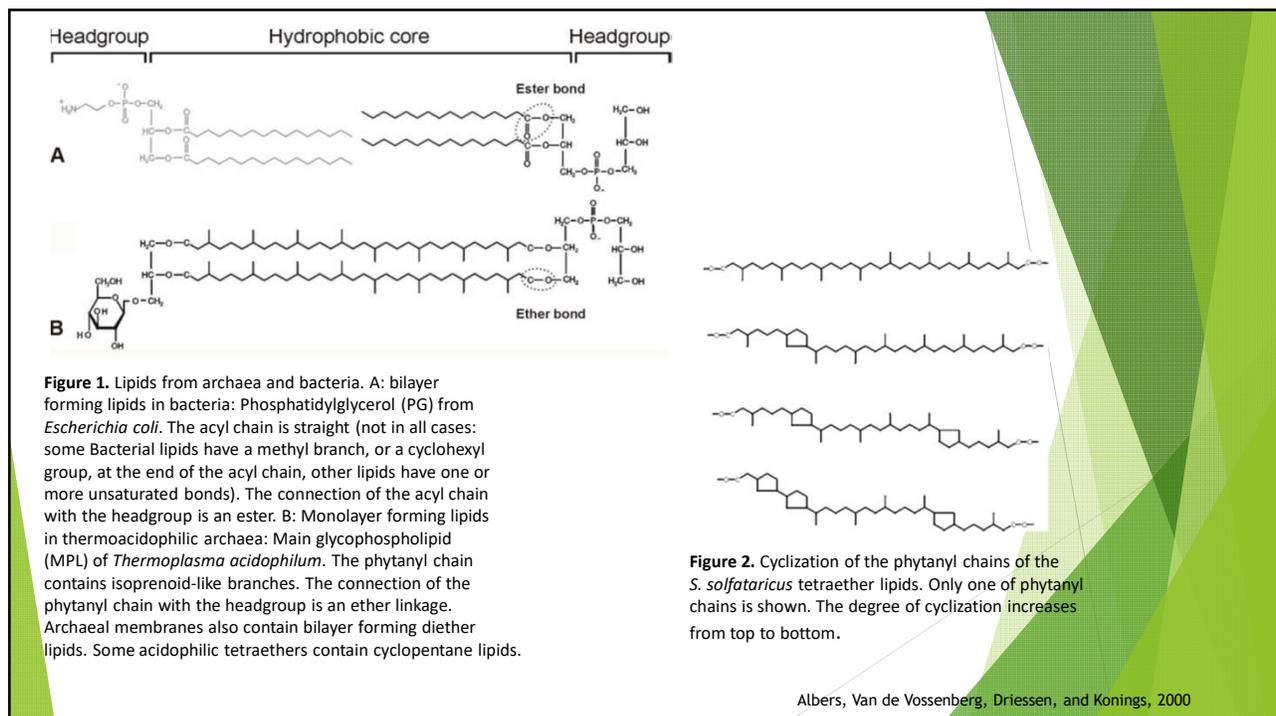
- ▶ They are fairly ubiquitous
 - ▶ Mesophilic inhabitants of soil and water
 - ▶ Form biofilms with bacterial species
 - ▶ Symbionts in the digestive tract of animals
 - ▶ Comprise 4% of human skin microbiota
 - ▶ None have been identified as pathogens
 - ▶ Also include psychrophiles, barophiles and acidophiles

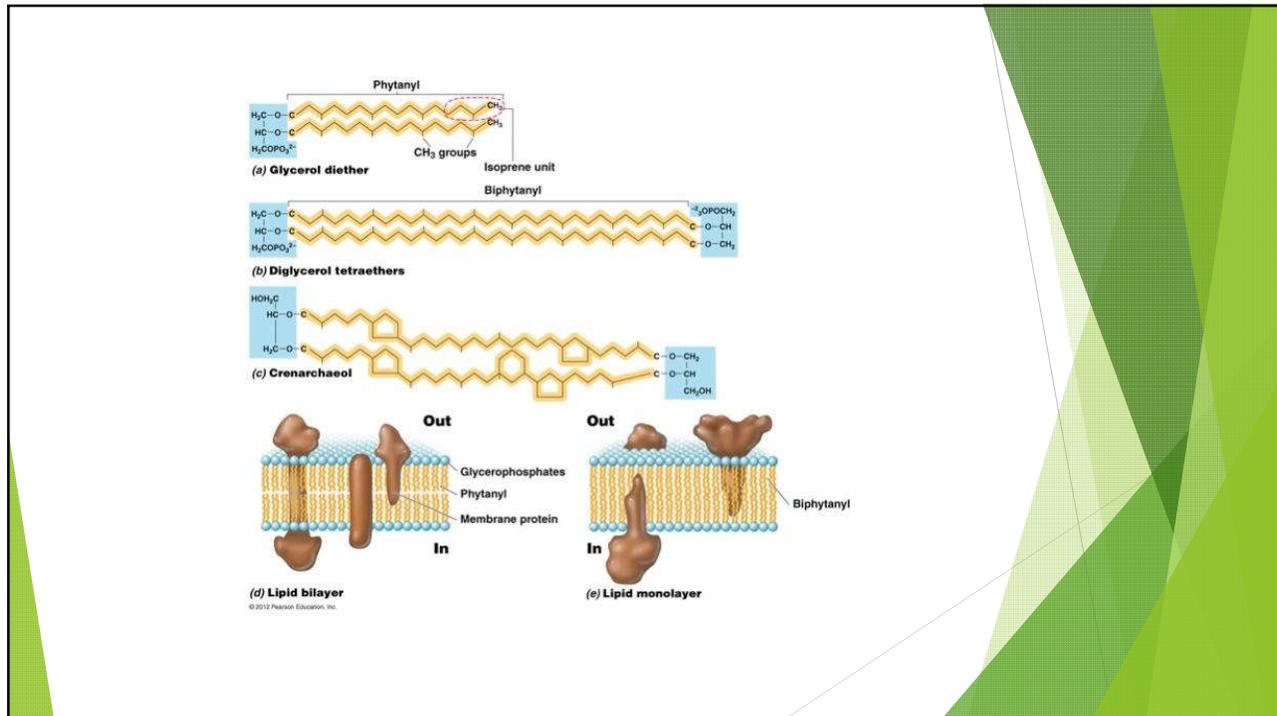
- ▶ Hyperthermophiles are commonly associated with hot springs, geysers and volcanic vents (including hydrothermal vents)
 - ▶ Unique features of these environments:
 - ▶ Reduced available minerals due to precipitation
 - ▶ Low oxygen availability (most are anaerobes)
 - ▶ Steep temperature gradients
 - ▶ Extreme acidity
 - ▶ Extreme pressure

- ▶ Hyperthermophiles are adapted to grow optimally in temperatures ranging from 80°C-106°C
 - ▶ Highest recorded critical temperature for an Archaeal species is 121°C
- ▶ The most extreme hyperthermophiles are also barophiles and/or acidophiles associated with hydrothermal vents
 - ▶ Steam (400°C) meets sea water and rapidly cools
 - ▶ Results in precipitation of minerals (Black Smokers)
 - ▶ High pressure allows water to remain liquid at extremely high temperatures

Distinctly Unique Features of Archaea

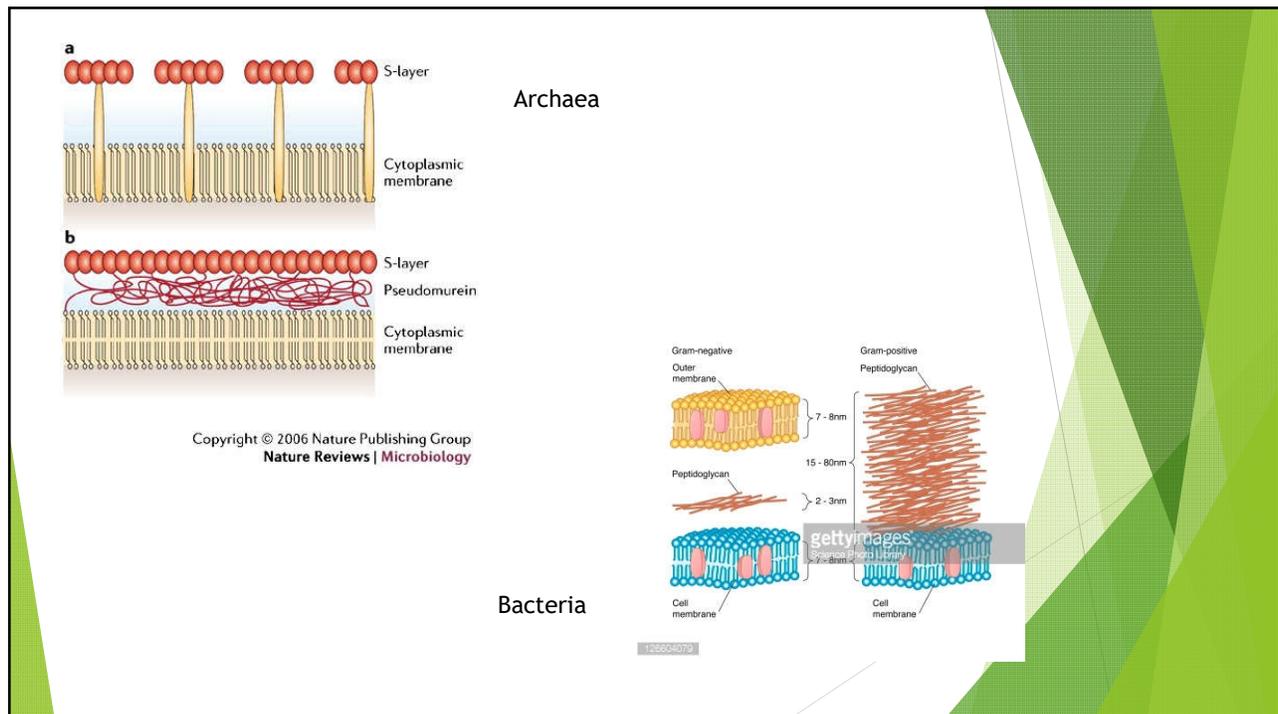
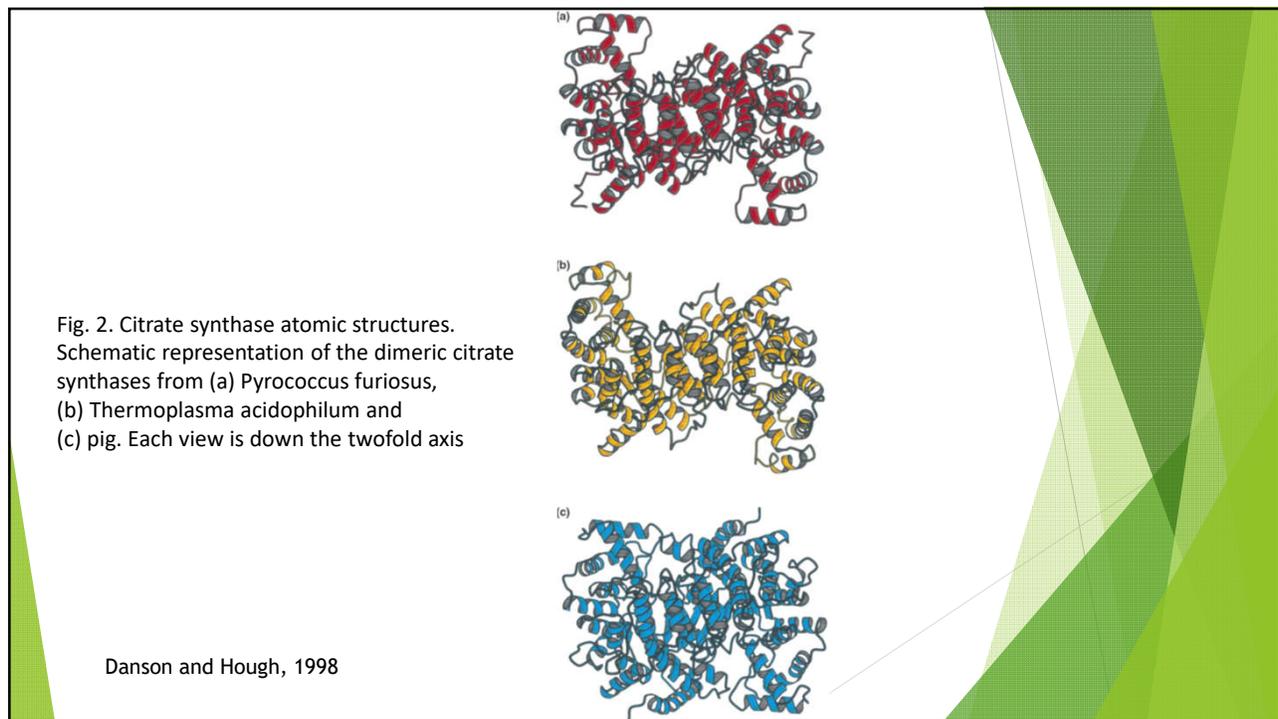
- ▶ Isoprenoid membranes
 - ▶ Glycerol isomer with ether linkage
 - ▶ Branched isoprenoid hydrocarbon chains
 - ▶ May cyclize to form ring structures or cross-link to form rigid tetraether monolayers
 - ▶ Thermal stability and reduced H⁺ permeability
 - ▶ H⁺ permeability is vital to bioenergetics
 - ▶ Reduced membrane permeability reduces energy stress





► Cell Wall or Envelope Structure

- Many archaea lack a cell wall (atypical of prokaryotes)
- If cell wall is present, distinctly different composition than bacteria
 - Resistant to compounds that degrade bacterial cell walls
- May have an envelope created by fusion of S-layer of glycoproteins to cell wall or membrane
 - Highly protective to environmental stresses but very flexible

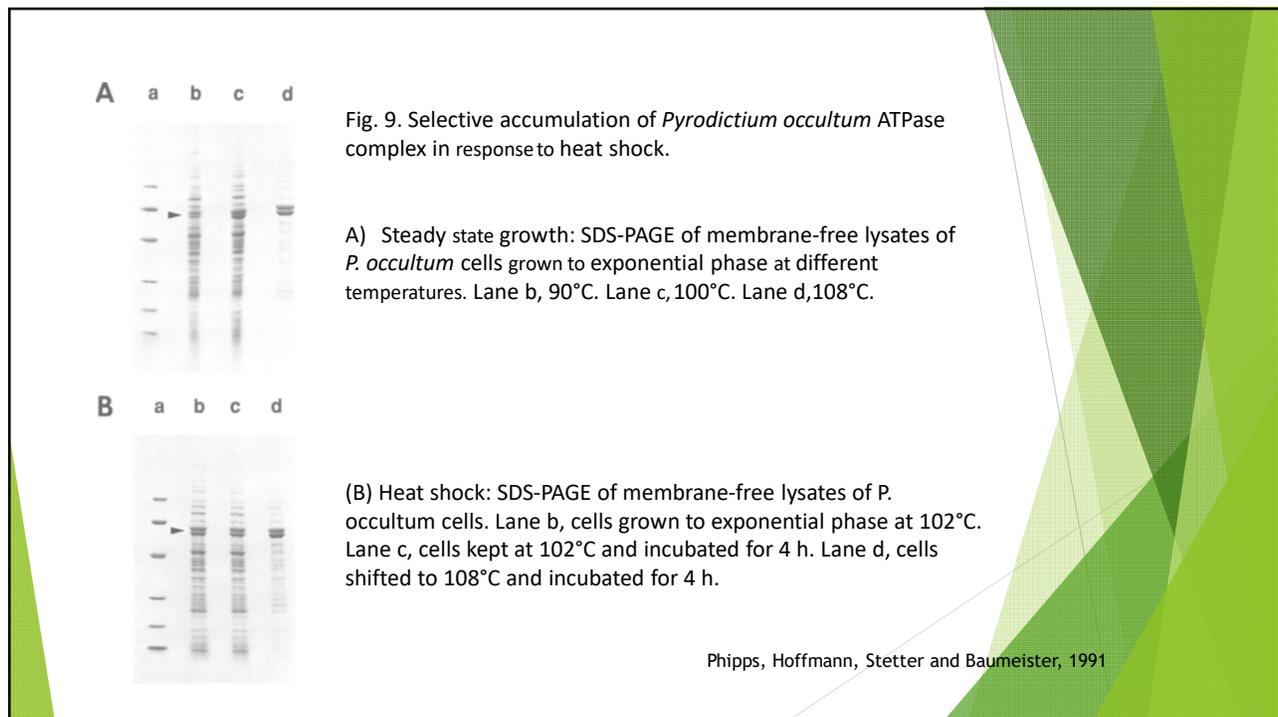


Sulfolobus (Crenarchaota)

- ▶ Requires both extreme heat (80-90°C) and extreme acidity (pH of 2-3)
- ▶ Tetraether mono-layer membrane with cyclic isoprenoid hydrocarbon chains
- ▶ Extensive S-layer protein envelope
- ▶ Autotrophic species oxidize H₂S
- ▶ Heterotrophic species use ferredoxin as an electron carrier
 - ▶ More stable at high temperatures than NAD

Pyrodictium (Crenarchaota)

- ▶ Anaerobes reducing precipitating sulfur to form H₂S
- ▶ Tetraether monolayer membrane
- ▶ S-layer protein envelope coated with zinc sulfides precipitating at the vent
- ▶ Produce a heat shock ATPase protein



Thermococcus and *Pyrococcus* (Euryarchaeota)

- ▶ Found around submarine volcanic vents
- ▶ Temperatures are above 90°C (up to 105°C)
- ▶ “Vent Polymerases” are replacing Taq Polymerase for PCR
 - ▶ Limited stability above 95°C

► Unique Genetic Composition

- Size and gene number similar to bacteria, under operon control
- Polymerase structure and action similar to eukarya
 - Have introns (interruptive sequences) and histones (DNA packaging proteins)
- 2/3 of genes in archaea have no homologue in bacteria or eukarya
- Reverse gyrase enzymes resulting in positive supercoiling of DNA
 - Only present in hyperthermophiles
 - Prevents unnecessary unwinding and separation of DNA strands at high temperatures

Table 1. Distribution of reverse gyrase in completely sequenced genomes from Archaea and hyperthermophilic Bacteria

	Optimal growth temperature (°C)	Presence of genes encoding reverse gyrase
<i>Pyrobaculum aerophilum</i>	100	Yes (2)
<i>Pyrococcus furiosus</i>	100	Yes
<i>Pyrococcus horikoshii</i>	98	Yes
<i>Methanopyrus kandleri</i>	98	Yes
<i>Pyrococcus abyssi</i>	98	Yes
<i>Aeropyrum pernix</i>	90-95	Yes (2)
<i>Aquifex aeolicus</i> (B)	85	Yes (2)
<i>Methanocaldococcus jannashii</i>	85	Yes
<i>Archaeoglobus fulgidus</i>	85	Yes
<i>Sulfolobus tokodaii</i>	85	Yes (2)
<i>Sulfolobus solfataricus</i>	85	Yes (2)
<i>Thermotoga maritima</i> (B)	80	Yes
<i>Methanothermobacter thermoautotrophicus</i>	60-65	No
<i>Thermoplasma volcanium</i>	55-60	No
<i>Thermoplasma acidophilum</i>	55-60	No
<i>Ferroplasma acidarmanus</i>	55-60	No
<i>Halobacterium</i> NRC1	50	No
All mesophiles (Archaea; Bacteria, Eukarya)	Under 50	No

Bacteria are denoted by (B).

Two reverse gyrase genes are present in species marked (2).

Forterre, 2002

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