

**Do all biogeochemical cycles work at  
elevated temperatures that exist at  
deep-sea hydrothermal vents ?**

Anne Godfroy & Daniel Prieur

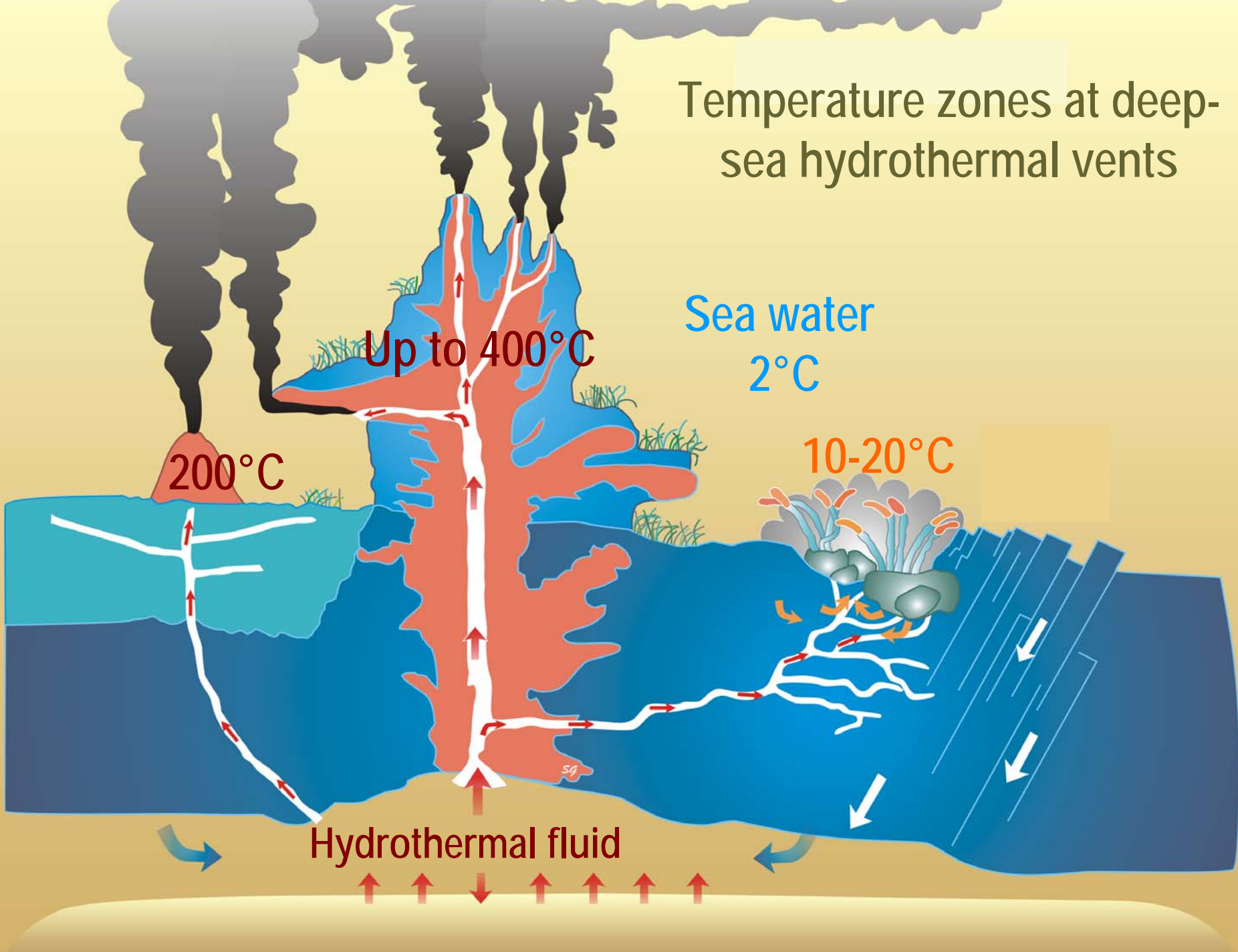
Laboratoire de Microbiologie des  
Environnements extrêmes

Ifremer, CNRS, Université de Bretagne Occidentale, Brest,  
France

# Outline

- Background
  - Microbial physiology
  - Microbial metabolism
  - Methods in microbial ecology
- Biogeochemical cycles at high temperature
  - Carbon
  - Nitrogen
  - Sulfur
  - Iron
- Novel approaches
- Conclusions

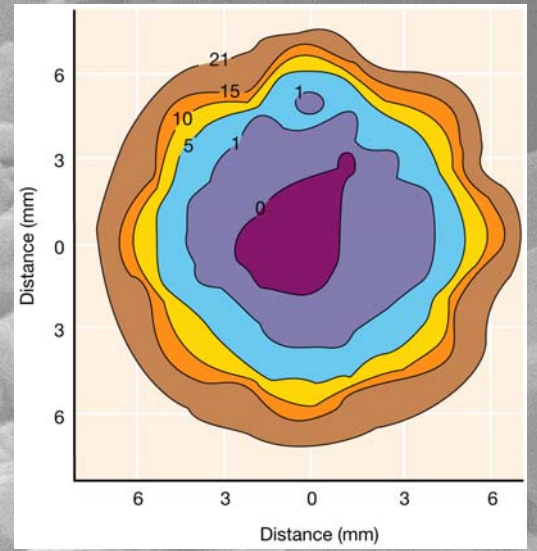
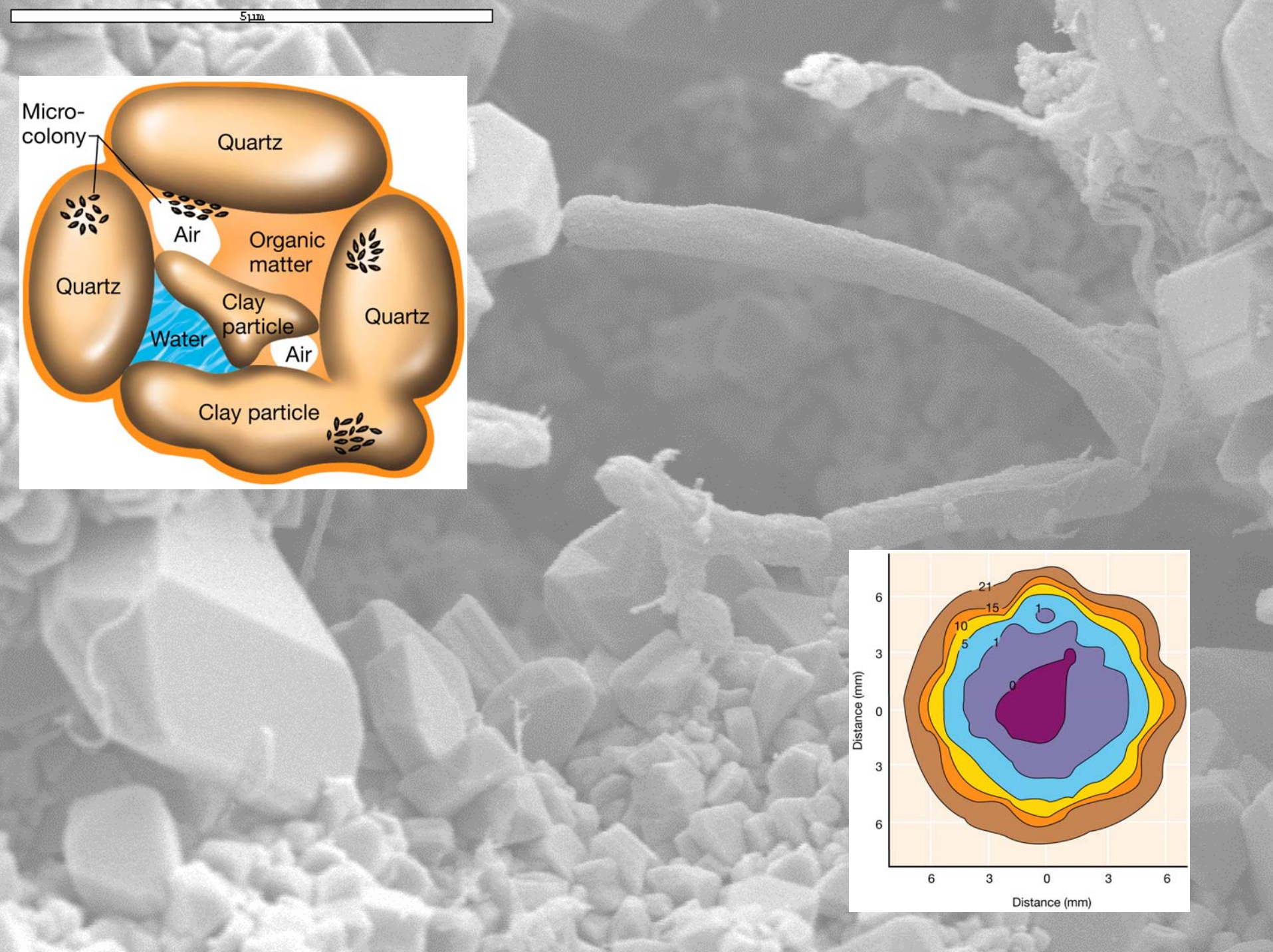
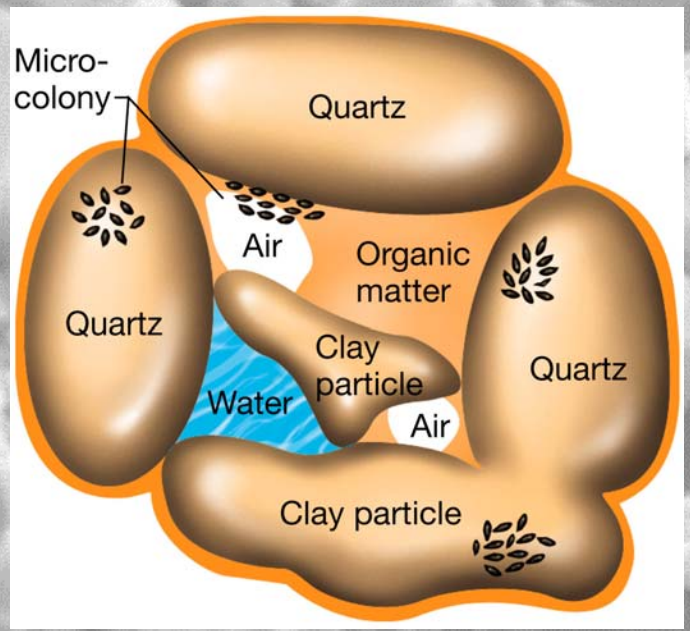
# Temperature zones at deep-sea hydrothermal vents



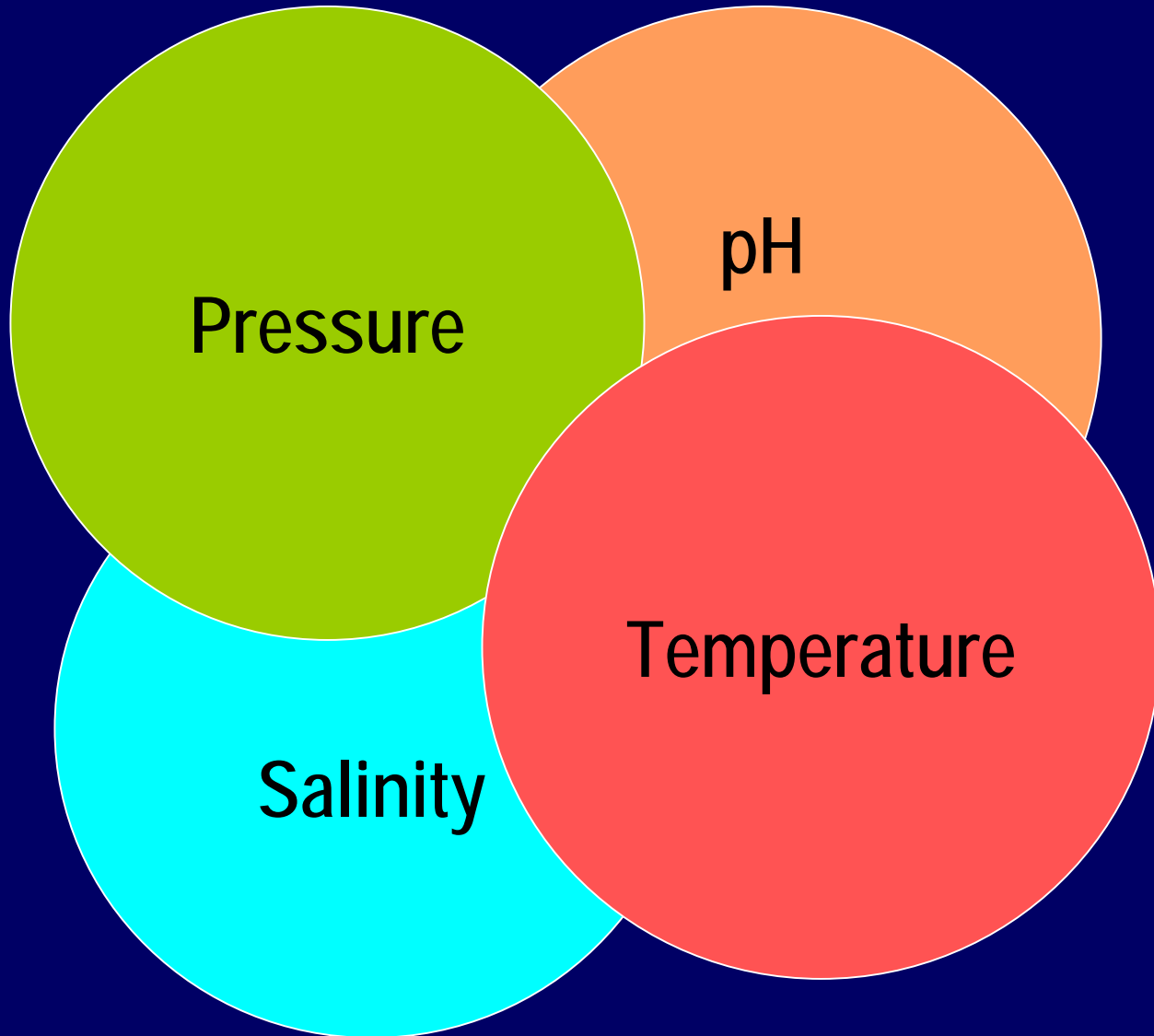
1  $\mu$ m



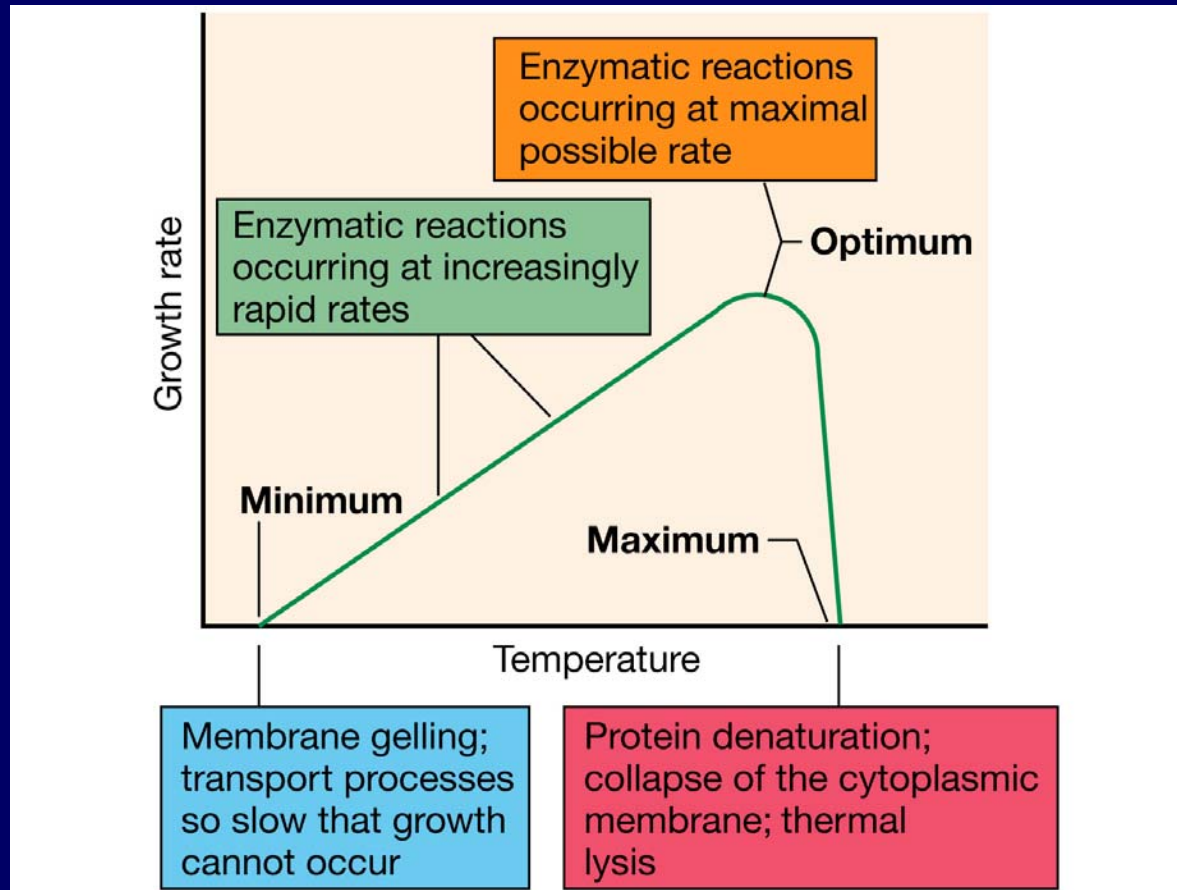
5µm

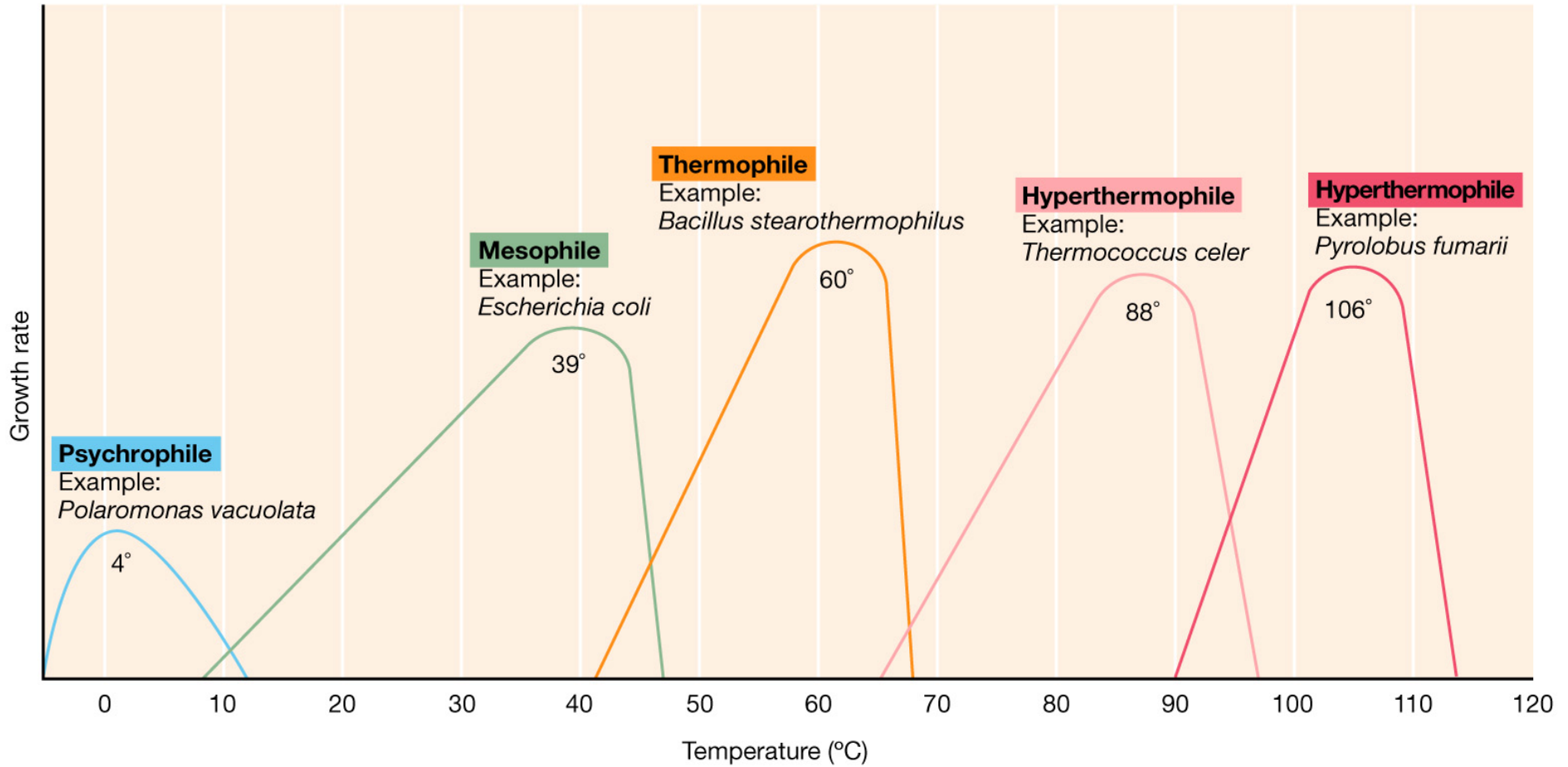


# Environmental parameters affecting microbial life



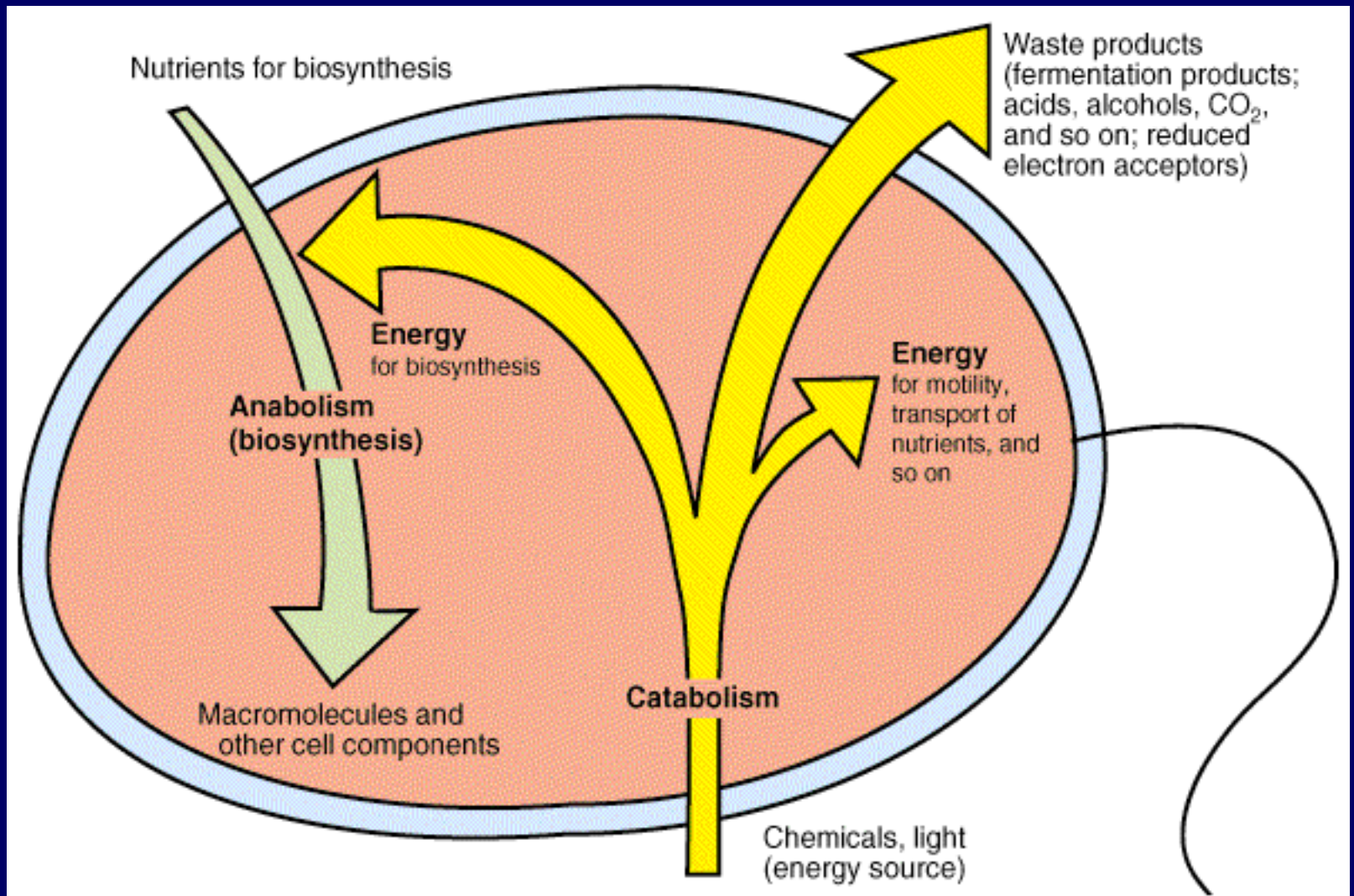
# Responses of microorganisms to temperature



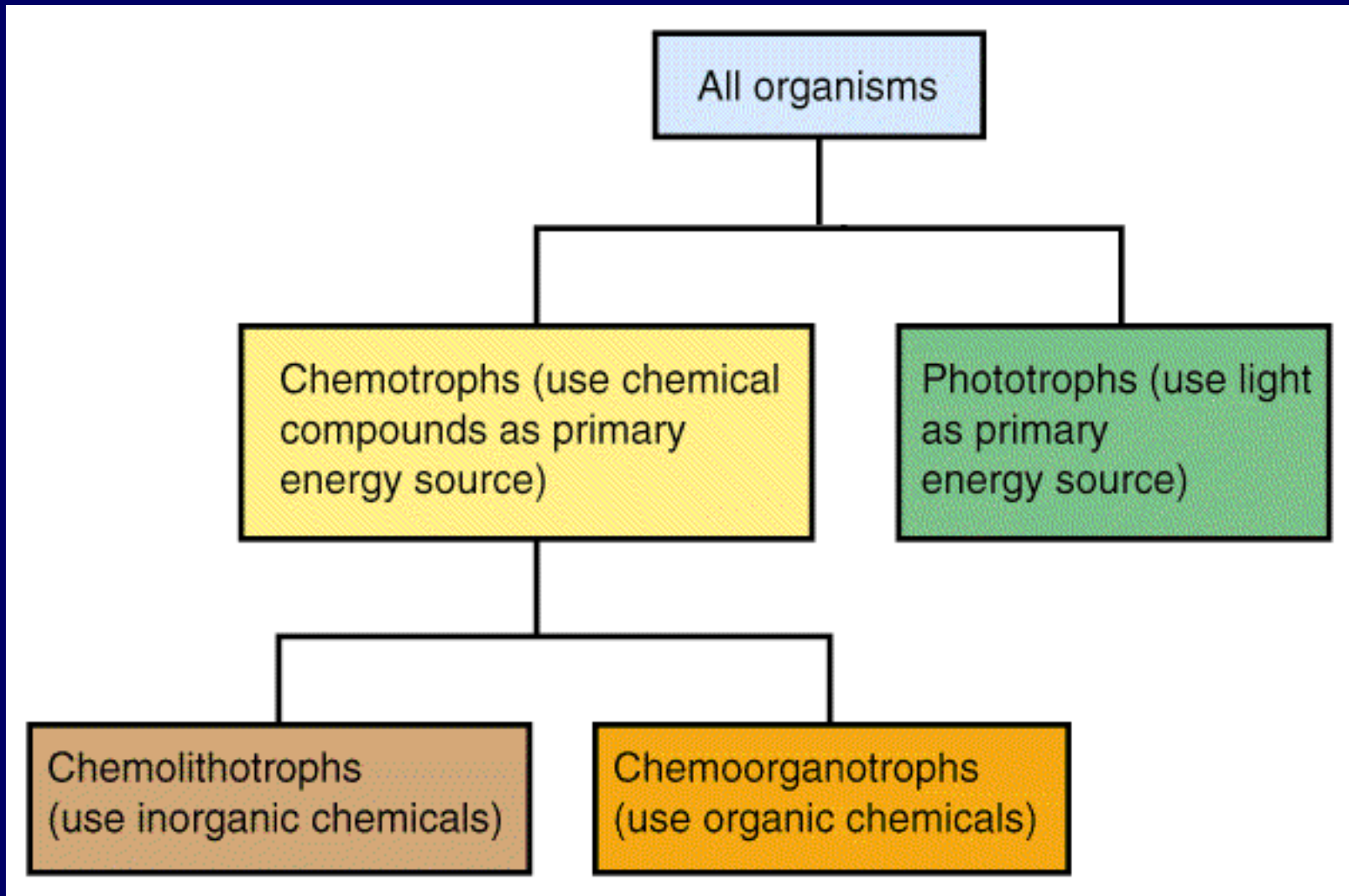




# Microbial cell metabolism

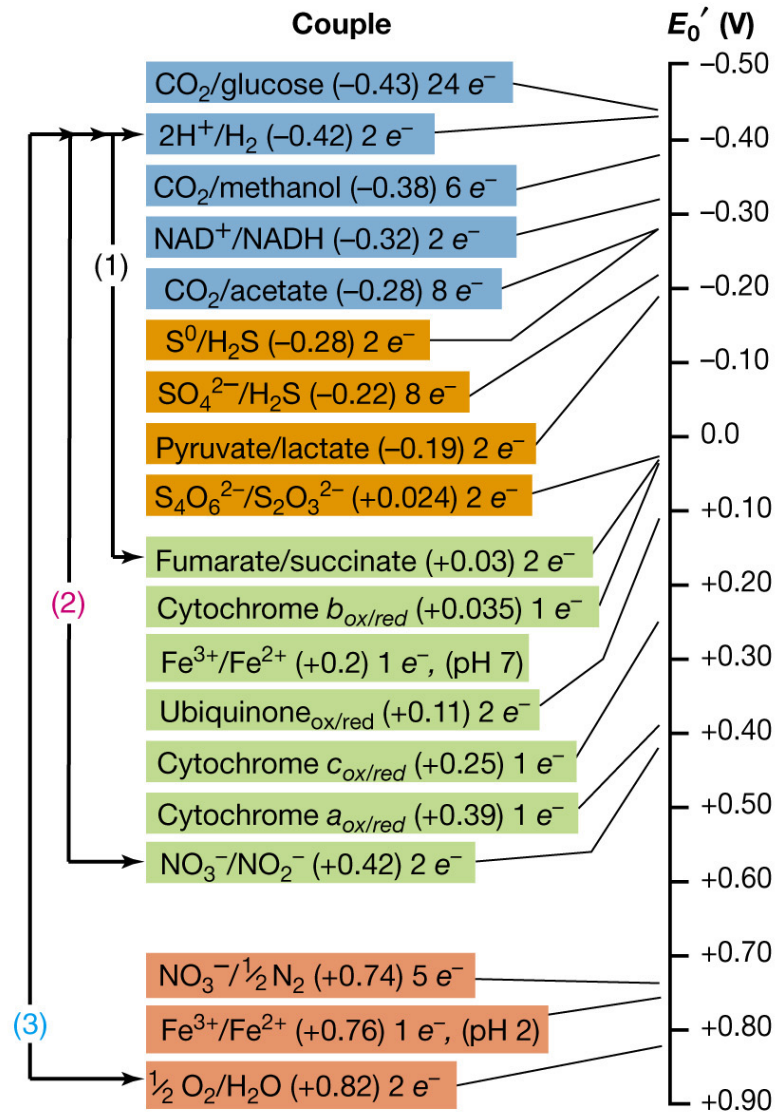
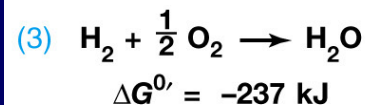
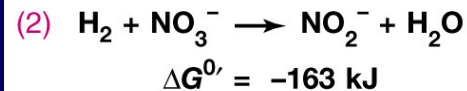
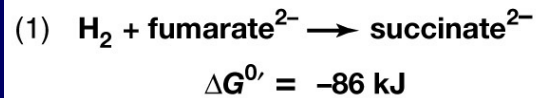


# Energy sources

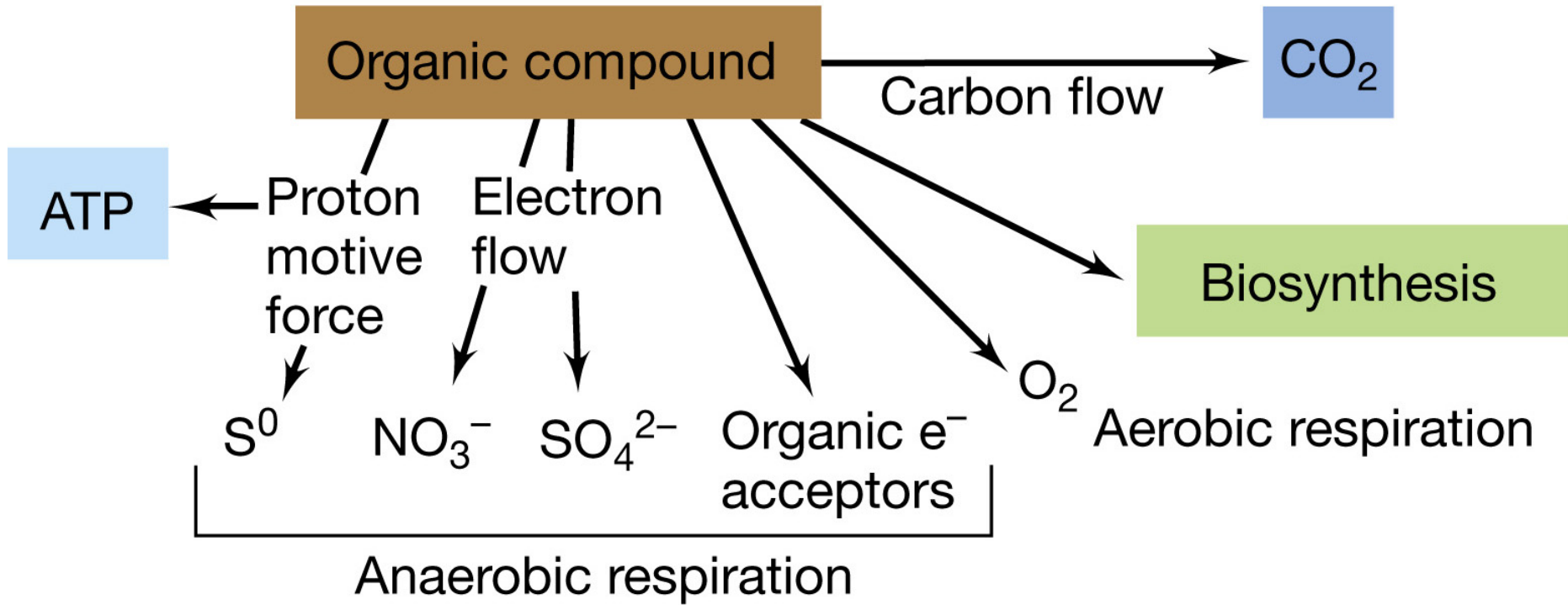


# Redox couples: electron tower

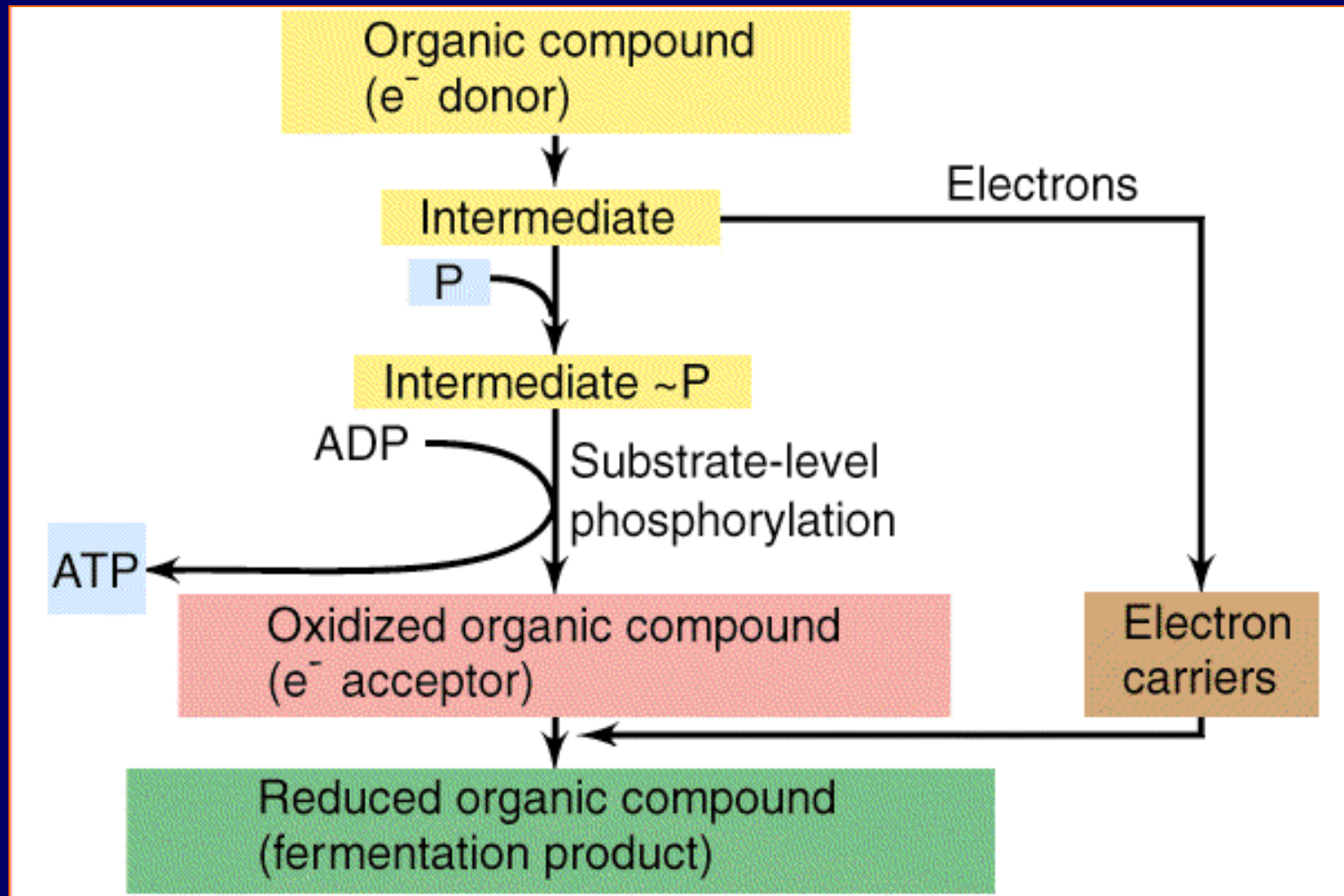
Examples of reactions  
with H<sub>2</sub> as e<sup>-</sup> donor



# Chemoorganotrophic metabolisms



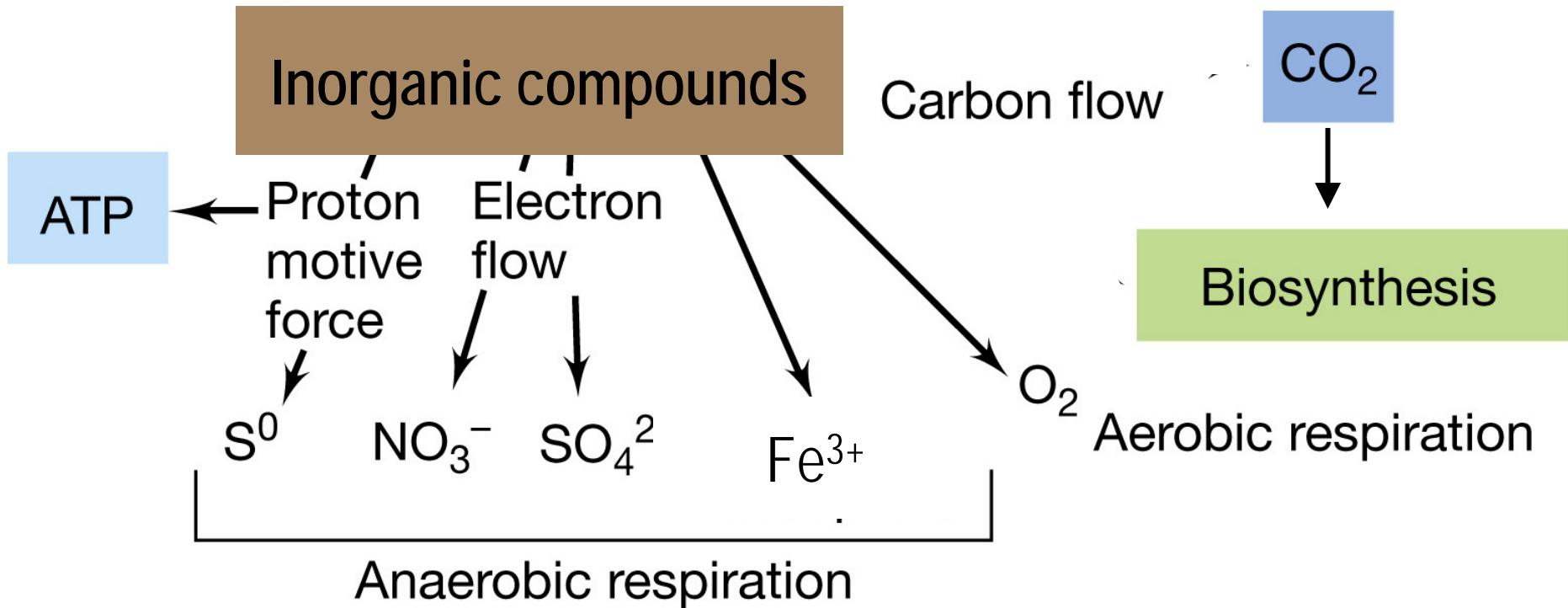
# Fermentation



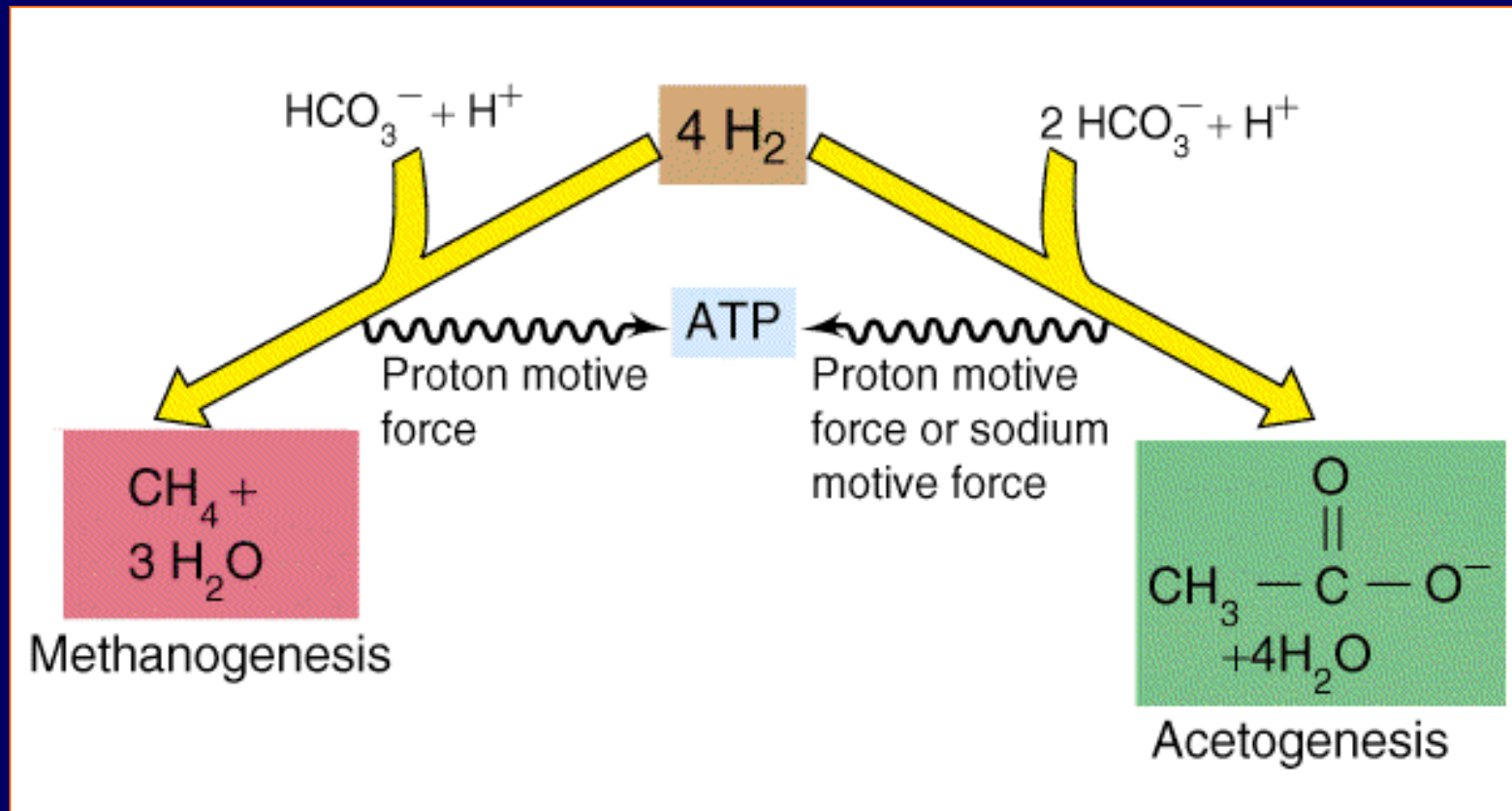
# Organic carbon sources/electron donors

- Hexoses, pentoses
- Polysaccharides
- Proteins
- Amino acids
- Organic acids
- Lipids
- Hydrocarbons

# Chemolithotrophic metabolisms



# CO<sub>2</sub>: electron acceptor

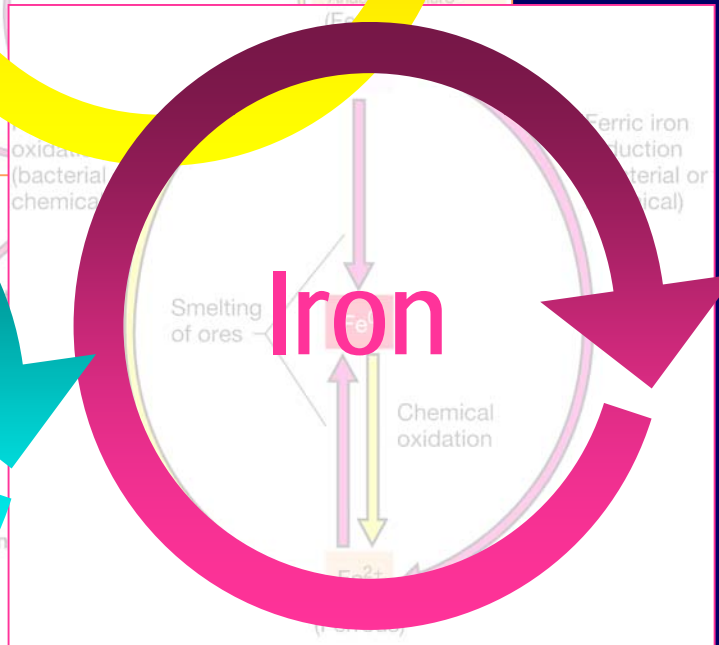
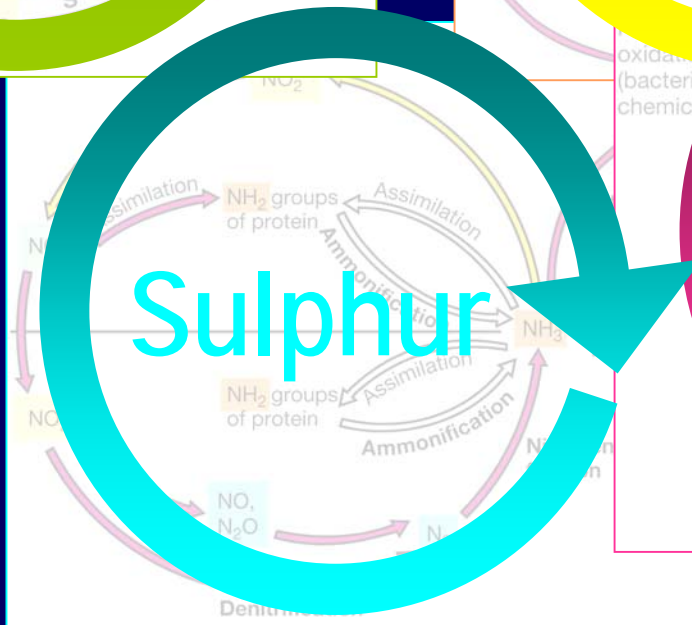
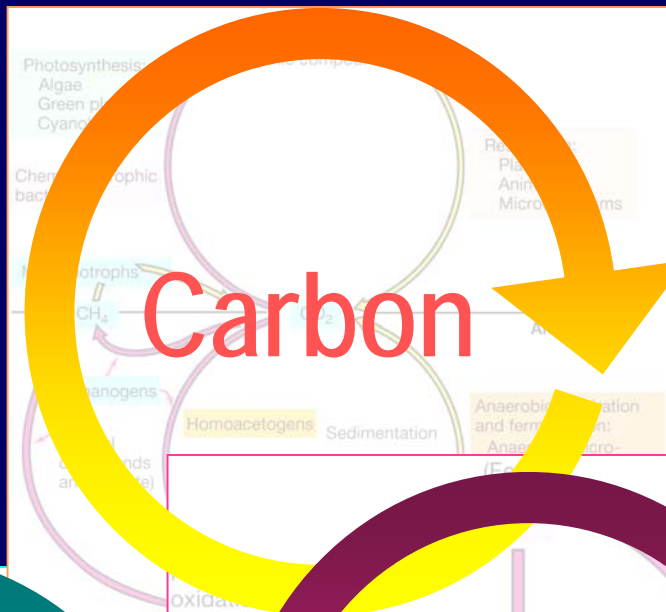
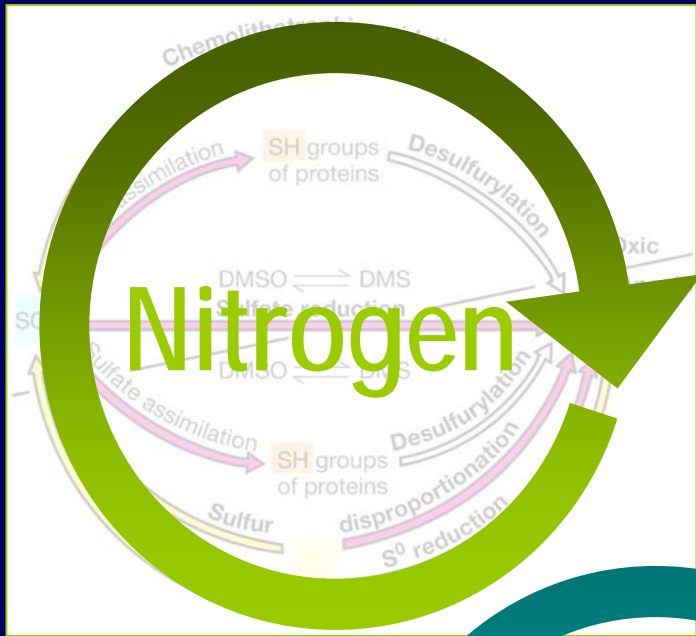




# Other electron acceptors

- Chlorate ( $\text{ClO}_3^-$ ) => Chlorine
- $\text{Mn}^{4+}$  =>  $\text{Mn}^{2+}$
- $\text{Fe}^{3+}$  =>  $\text{Fe}^{2+}$
- Selenate => Selenite
- Arsenate => Arsenite
- DMSO => DMS
- Fumarate => Succinate

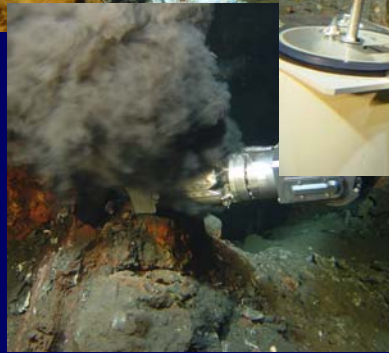
# Biogeochemical cycles



# Microbial ecology of hydrothermal vent chimneys



Sample preservation



Chimney sampling

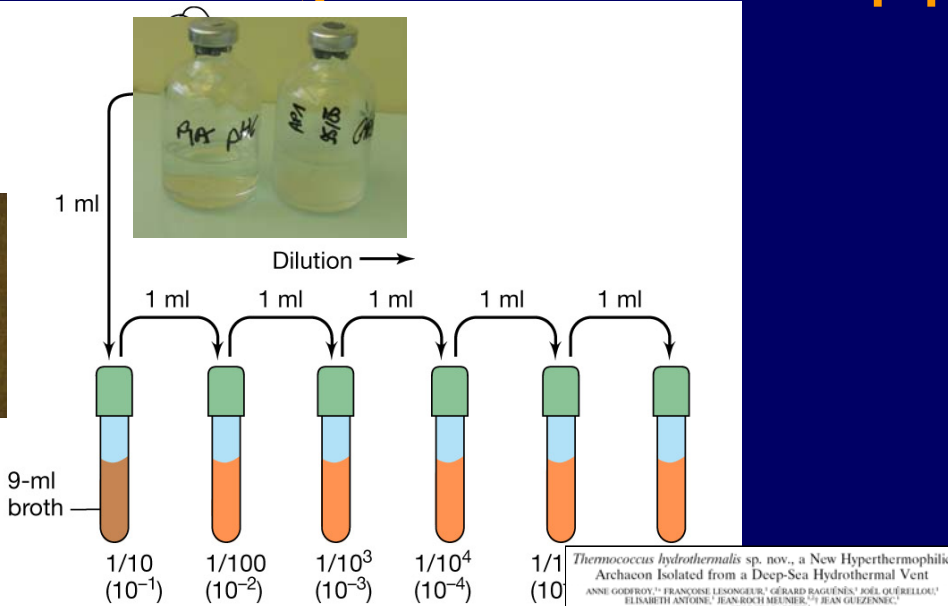


Activity

Molecular diversity studies

Cultures

# Microbial diversity in hydrothermal vent chimneys : cultural approaches



*Thermococcus hydrothermalis* sp. nov., a New Hyperthermophilic Archaeon Isolated from a Deep-Sea Hydrothermal Vent  
 ANNE GODFREY<sup>1</sup>, FRANCISSE LEJONNEUR<sup>1</sup>, GÉRARD BAGUENES<sup>1</sup>, JOËL OUBRELOU<sup>1</sup>, ELISABETH ANTOINE<sup>1</sup>, JEAN ROCH MEUNIER<sup>1,2</sup>, JEAN GUZENNEC<sup>1</sup>, and GEORGES BARBIER<sup>1</sup>



**Description of *Thermococcus hydrothermalis* sp. nov.** *Thermococcus hydrothermalis* (hydrothermalis, N.L. adj. hydrothermalis, pertaining to a hydrothermal vent). Cells are cocci (diameter, 0.8 to 2 μm) that are motile by means of polar flagella. Cell division occurs by constriction. Obligately anaerobic. Grows optimally in the presence of 30 to 40 g of Sea Salt per liter and at a pH around 6. Growth occurs at 55 to 100°C, and the optimum temperature is around 85°C. Obligately chemoorganotrophic. Grows preferentially on proteolysis products, a mixture of amino acids, and maltose. Sulfur is not necessary for growth but greatly enhances growth. The results of 16S rRNA sequence comparisons place *Thermococcus hydrothermalis* in the *Thermococcales*. Type strain AL662 (= CNCMI1319 [Collection Nationale de Cultures de Microorganismes, Institut Pasteur, Paris, France]) was isolated from an active chimney wall fragment recovered from a hydrothermal site on the East Pacific Rise at a latitude of 21°N.

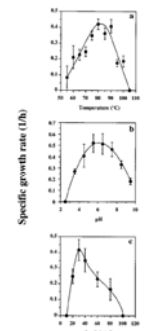


FIG. 1. Temperature, pH, and Sea salt concentration optima for growth of strain AL662<sup>T</sup> on 2216-S medium. (a) Specific growth rate as a function of temperature (in the presence of 30 g of Sea Salt per liter at pH 7.5). (b) Specific growth rate as a function of pH (in the presence of 30 g of Sea Salt per liter at 85°C). (c) Specific growth rate as a function of salinity (at 85°C and pH 7). Growth rates were calculated by performing a linear regression analysis along the logarithmic parts of the growth curves. If enough data were available, inferential about the precision of the growth rate is given (regression coefficient ± standard errors).

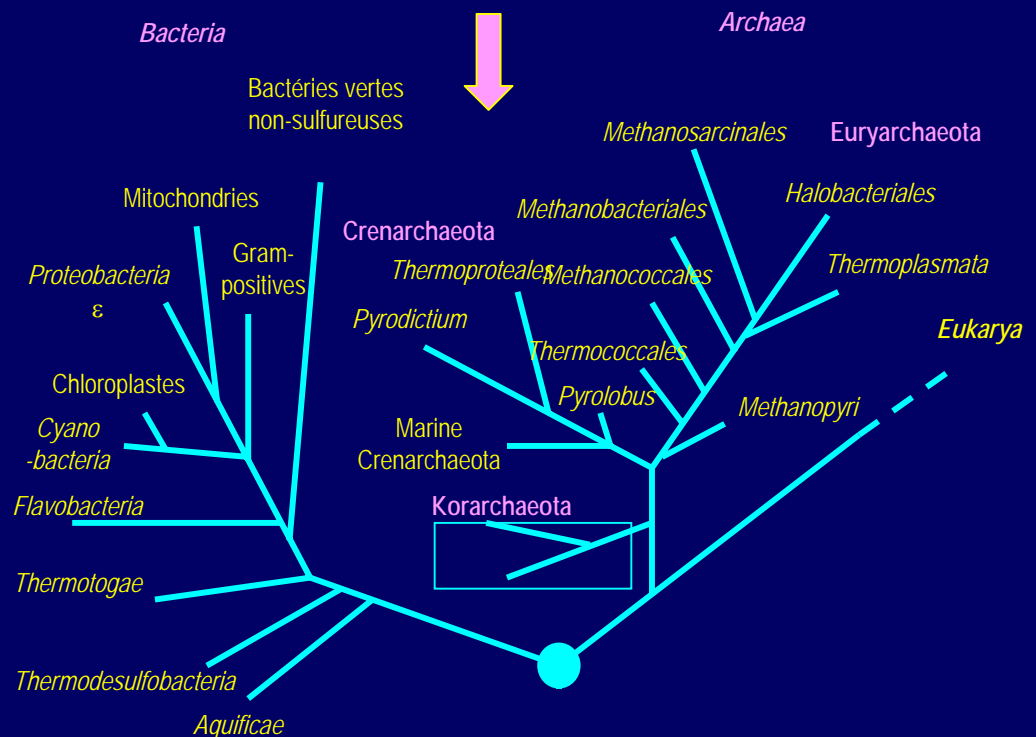
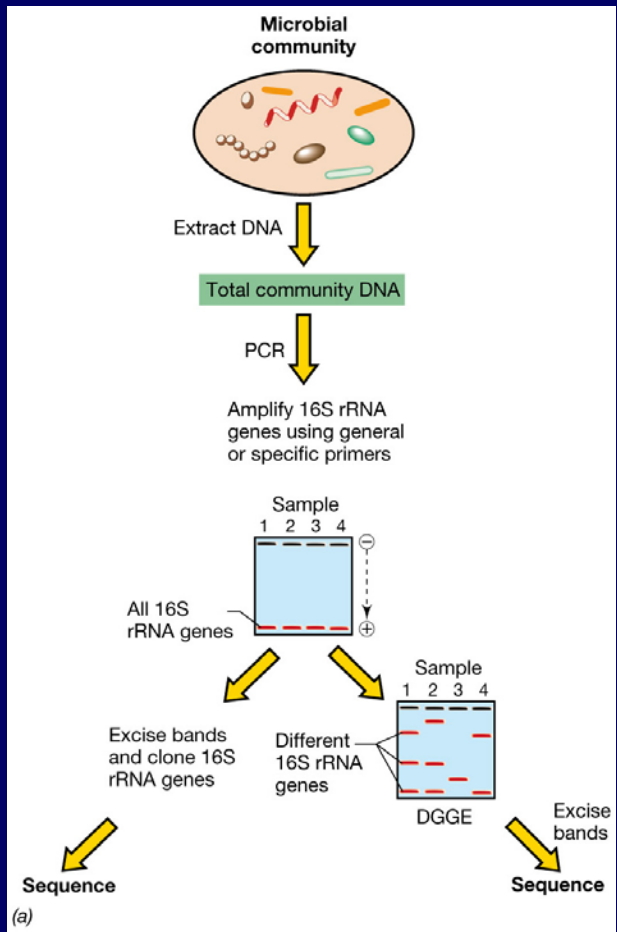
TABLE 1. Growth requirements of strain AL662<sup>T</sup> (Growth after\*)

Compound(s)	Growth after*		
	8 h	15 h	24 h
Carbon sources			
Yeast extract-peptone (2216-S medium)	+++	+++	ND
Meat extract	++	+++	ND
Yeast extract	+	++	+++
Malt extract	++	+++	+++
Brain heart infusion (BHS medium)	+++	+++	+++
Casamino Acids	++	+++	ND
Cysteine	+	++	++
Glucose	ND	—	—
Maltose	—	—	—
Sucrose	+++	ND	+++
Cellulose	+	ND	+++
Starch	—	—	—
Ethanol	—	—	—
Mannitol	—	—	—
Acetate	—	—	—
Pyruvate	—	—	+
20 Amino acids (20AAS medium)	++	+++	+
None (under H <sub>2</sub> -CO <sub>2</sub> )	—	—	—
BHS-S medium (under H <sub>2</sub> -CO <sub>2</sub> )	+++	+++	ND
Electron acceptors			
Sulfur	+++	ND	+++
None (under N <sub>2</sub> )	++	ND	++
None (under N <sub>2</sub> -CO <sub>2</sub> 80:20)	++	ND	++
None (under N <sub>2</sub> -CO <sub>2</sub> 80:20)	—	ND	—
Cysteine	++	+++	ND
Polyallylde	++	ND	ND

\* +, ++, +++: 10<sup>6</sup>, 10<sup>7</sup>, 10<sup>8</sup> cells per ml of culture (final concentration); —, ND: 10<sup>6</sup> to 10<sup>8</sup> cells per ml of culture (final concentration); —, 2: 10<sup>6</sup> to 10<sup>8</sup> cells per ml of culture (final concentration); —, no growth (< 10<sup>6</sup> cells per ml of culture); ND, not determined.

# Microbial diversity in hydrothermal vent chimneys : Molecular approaches

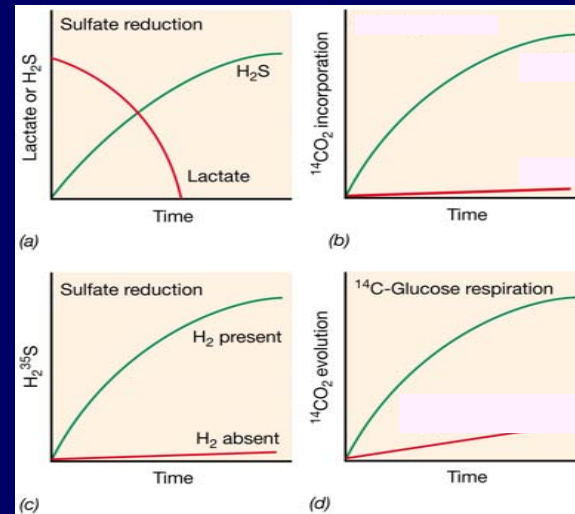
16S rRNA genes  
or  
functional genes



# Microbial diversity in hydrothermal vent chimneys : metabolic activities



Samples



Incubation with labeled substrates  
(stable or radioactive isotopes or fluorescent molecules)



Hydrolytic activity  
Sulphate reduction  
etc...

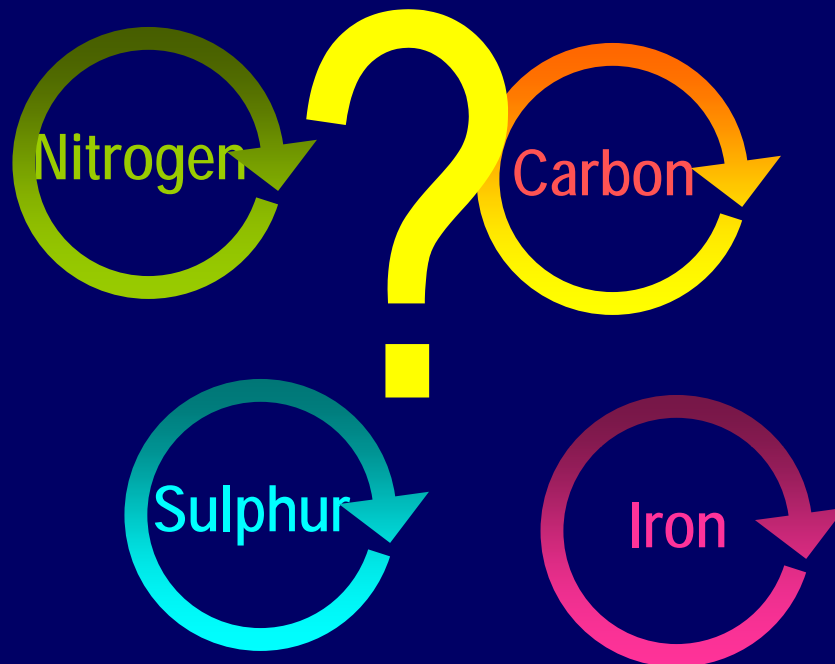
# Conclusions (1)

For a given strain, cultural approaches give informations about carbon sources, electron donors and acceptors, and suitable environmental conditions for this strain.

Molecular approaches give informations about phylogeny (sometimes linked to metabolism) and/or functions (functional genes).  
Environmental conditions for a given clone are uncertain.

Measurements of metabolic activities confirm this activity exists for the conditions of the assay.

# What do we know about biogeochemical cycles at high temperature in deep sea-hydrothermal vent chimneys



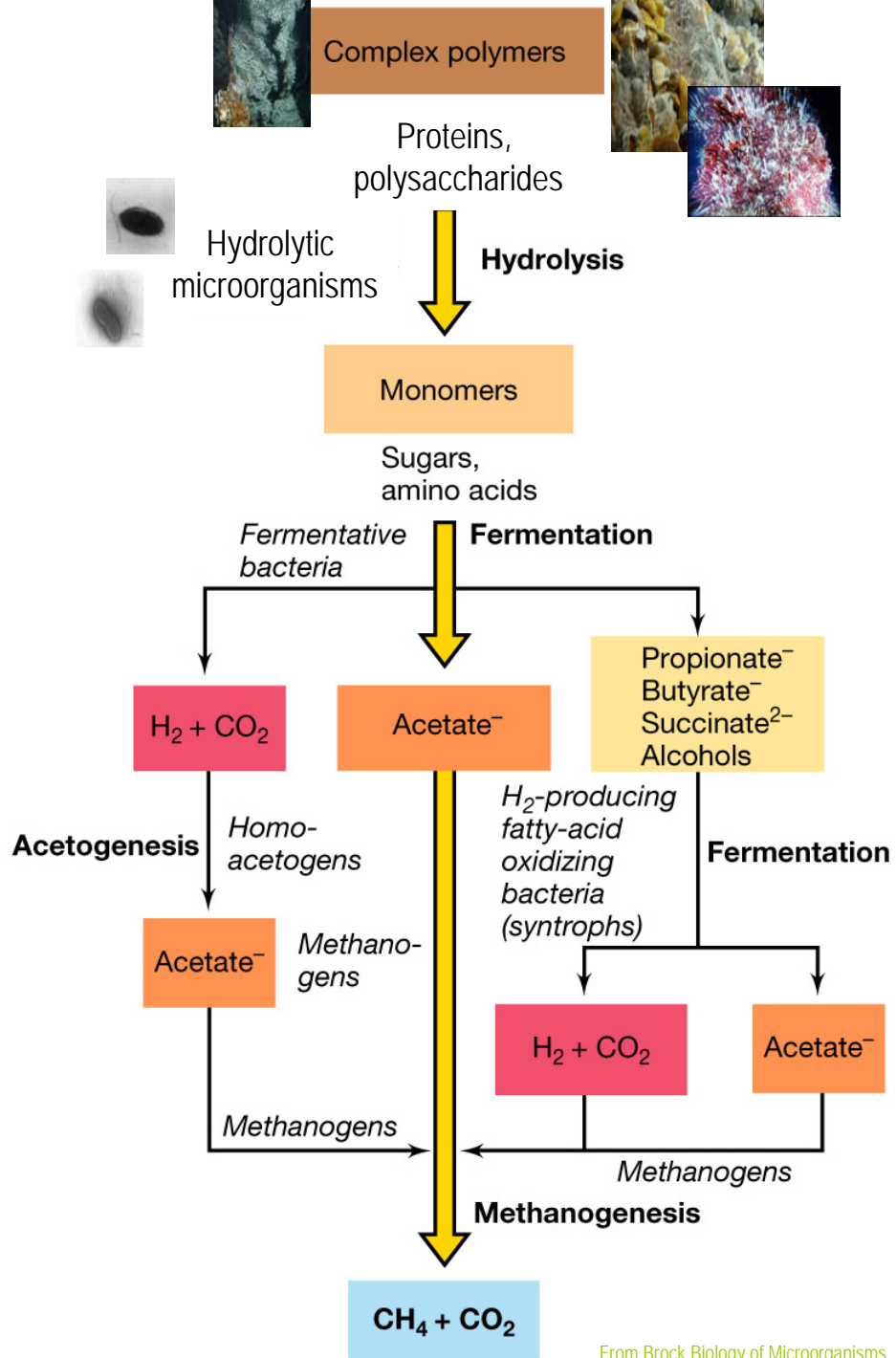


# Please, note that

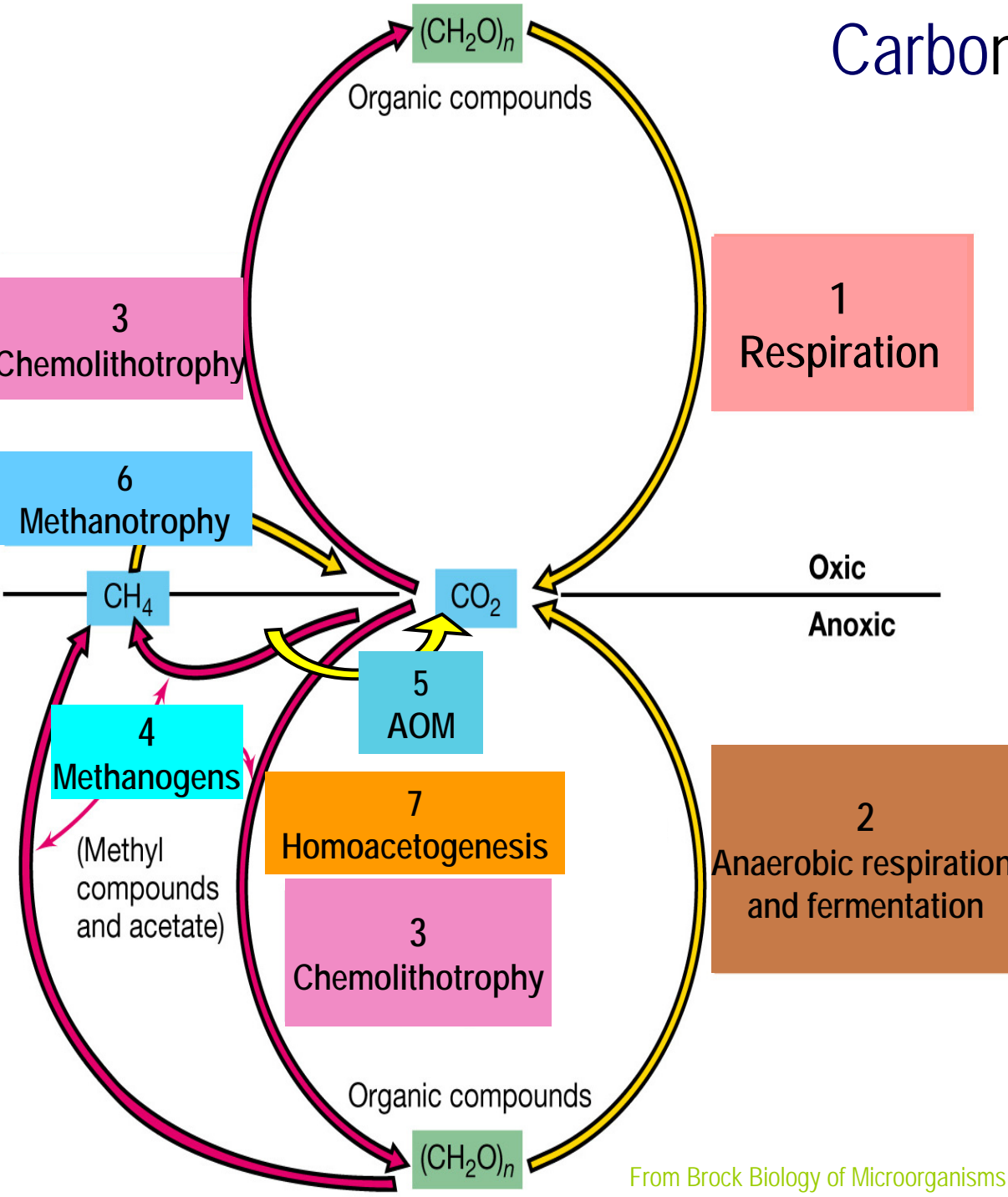
- Data used for this presentation were collected from published papers and pooled.
- Locations of vent sites were not taken into account.
- We apologize for possibly missing data.
- Please let us know...

# Carbon cycle

Organic matter degradation/  
Organic matter synthesis



# Carbon cycle

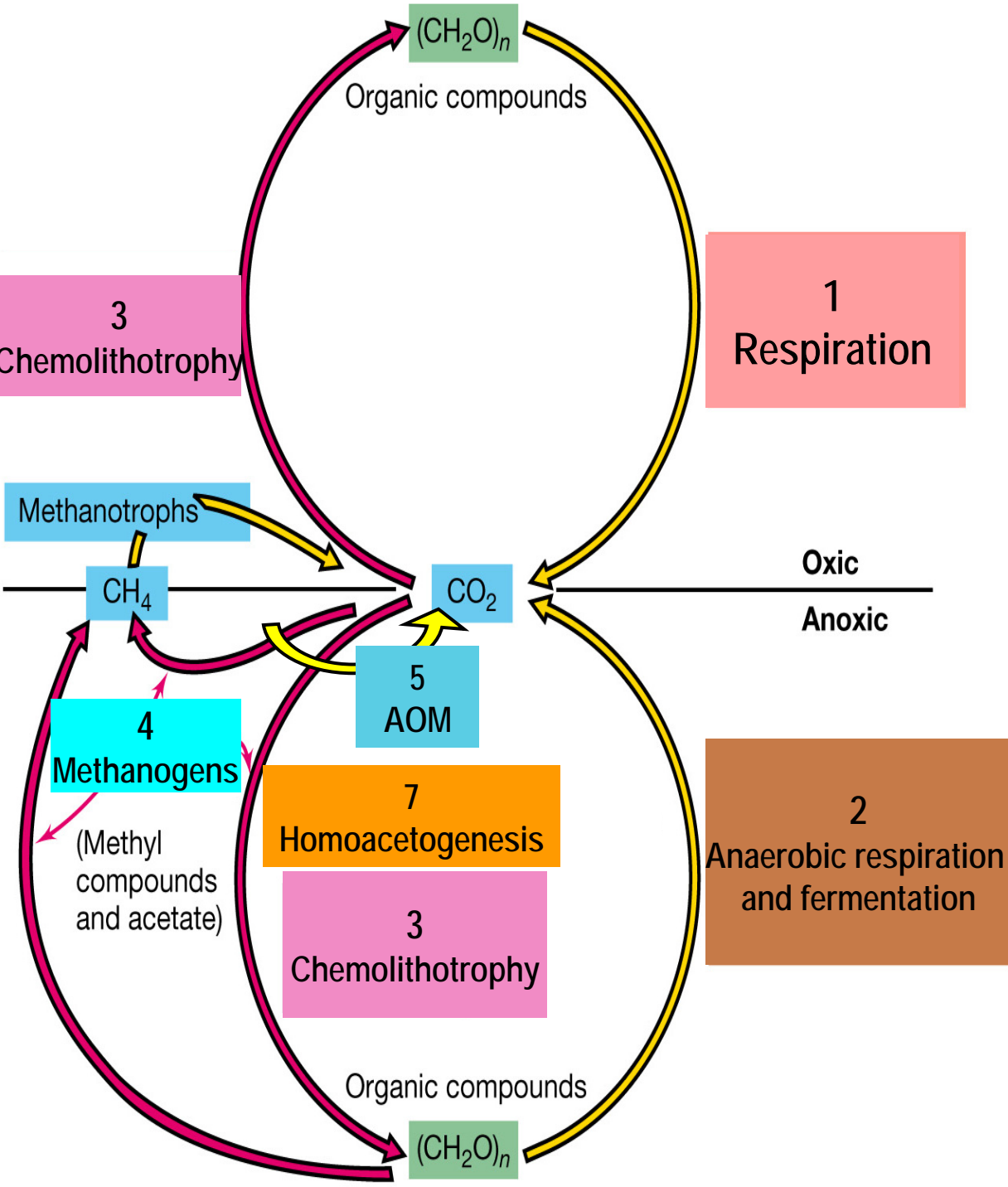


- 1**
- Aerobes and Microaerophiles
  - Oceanithermus*
  - Vulcanithermus*
  - Aeropyrum camini*
  - Thermus sp.*
  - T° op max 85°C

- 2**
- Caminicella*
  - Vulcanibacillus*
  - Caloranaerobacter*
  - Thermosipho*
  - Marinotoga*
  - Tepidibacter*
  - Deferribacter desulfuri*
  - Sulfurospirillum*
  - Desulfurococcus sp.*
  - Staphylothermus*
  - Thermococcus*
  - Pyrococcus*
  - Palaeococcus*
  - Pyrodictium abyssi*
  - Aciduliprofundum*
  - T° op max 95°C

- 3**
- Archaeoglobus*
  - Persephonella*
  - Desulfurobacterium*
  - Balnearium*
  - Thermovibrio*
  - Deferribacter*
  - Caminibacter*
  - Nautilia*
  - Thermodesulfobacterium*
  - Thermodesulfatator*
  - Sulfurimonas*
  - Hydrogenimonas*
  - Lebetimonas*
  - Ignicoccus*
  - Pyrolobus*
  - T° op max 106°C



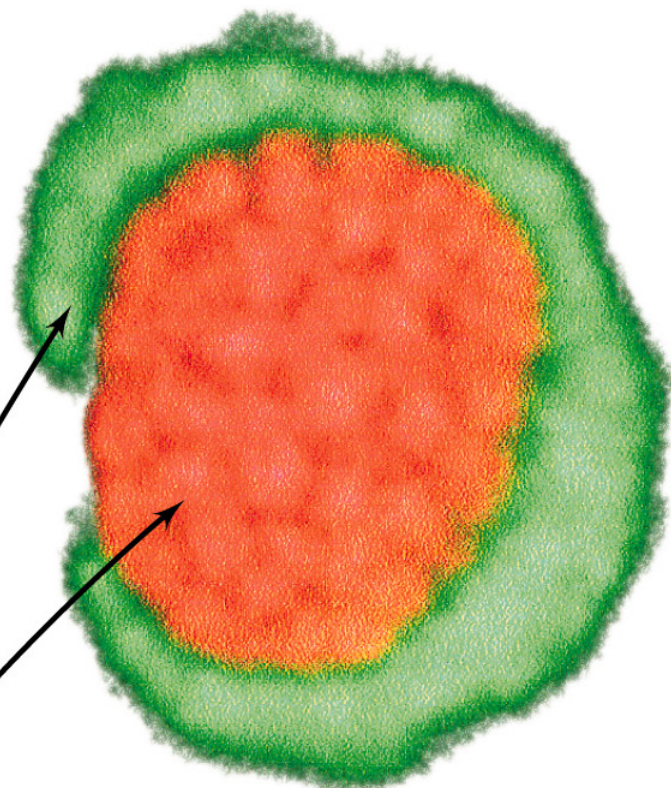


**4**  
 Methanoarchaea  
 ( $\text{H}_2$ )  
*Methanocaldococcus*  
*Methanotorrus*  
*Methanopyrus*  
 T° op max 98°C

**5**  
 AOM  
 Anaerobic oxidation of  
 methane  
 Molecular and activity  
 Evidence

**6**  
 Methanotrophy  
 $\text{CH}_4$  oxidation  
 NO

**7**  
 Homoacetogenesis  
 NO

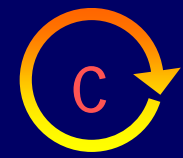


Antje Boetius and Armin Gieseke

(a)

Reaction	Organism	$\Delta G^{0'}$ (kJ)
$\text{CH}_4 + 2 \text{H}_2\text{O} \longrightarrow \text{CO}_2 + 4 \text{H}_2$	Methanogen	+131
$\text{SO}_4^{2-} + 4 \text{H}_2 + \text{H}^+ \longrightarrow \text{HS}^- + 4 \text{H}_2\text{O}$	Sulfate-reducer	-156
<hr/>		
Sum: $\text{SO}_4^{2-} + \text{CH}_4 \longrightarrow \text{HCO}_3^- + \text{HS}^- + \text{H}_2\text{O}$	Syntrophic reaction	-25

(b)



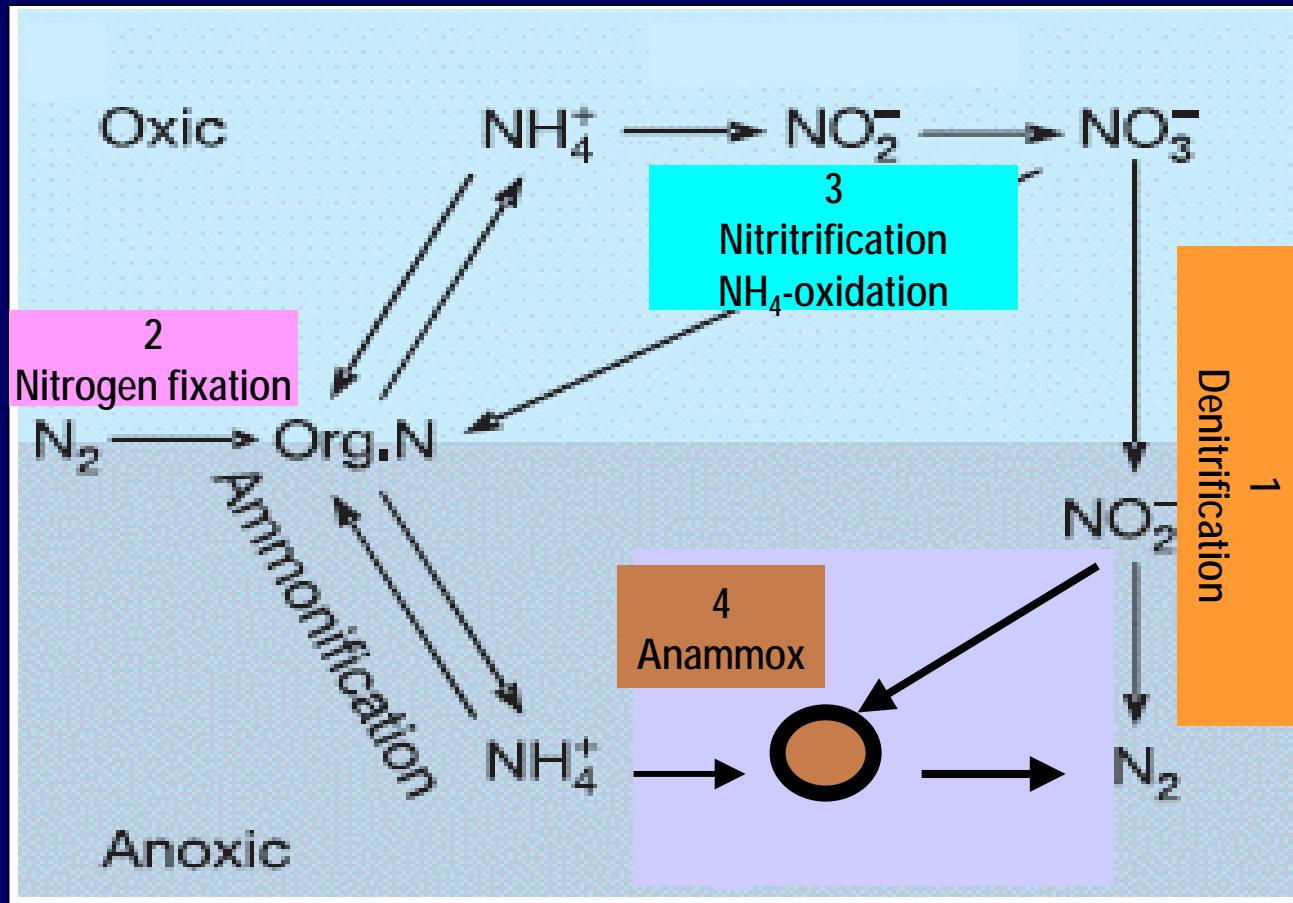
# Nitrogen Cycle

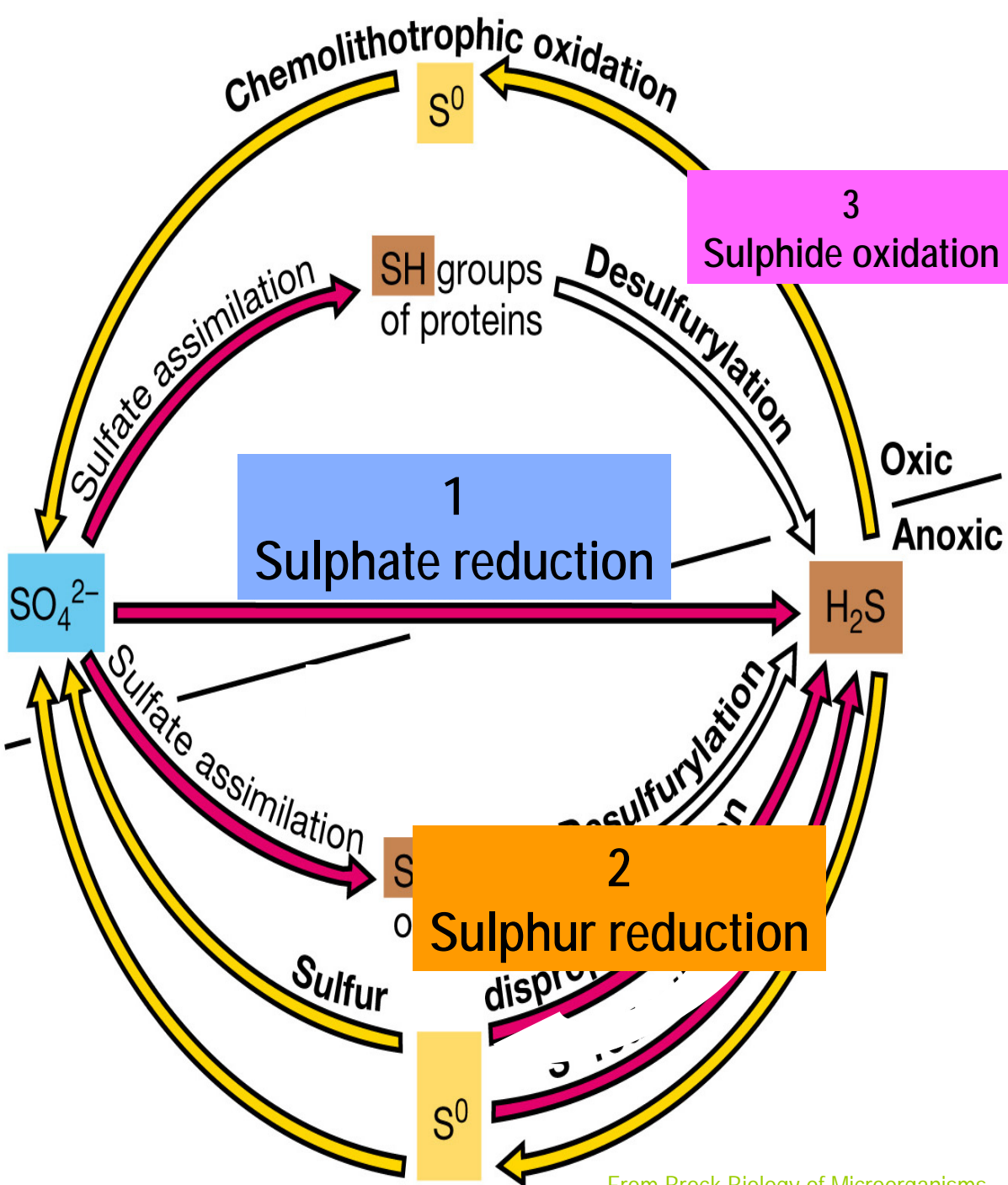
1  
Denitrification  
NO<sub>3</sub><sup>-</sup> reducers  
*Persephonella*  
*D. crinifex*  
*Deferribacter abyssi*  
*Caminibacter*  
*Sulfurimonas*  
*Pyrolobus*  
*Caldithrix*  
*Geothermobacter*  
T° op max 106°C

3  
Aerobic  
NH<sub>4</sub>-oxidation

NO  
4  
Anammox

2  
Nitrogen fixation  
nifH genes detection  
and  
*M. Jannaschii*  
str FS406-22  
T° op max 90°C



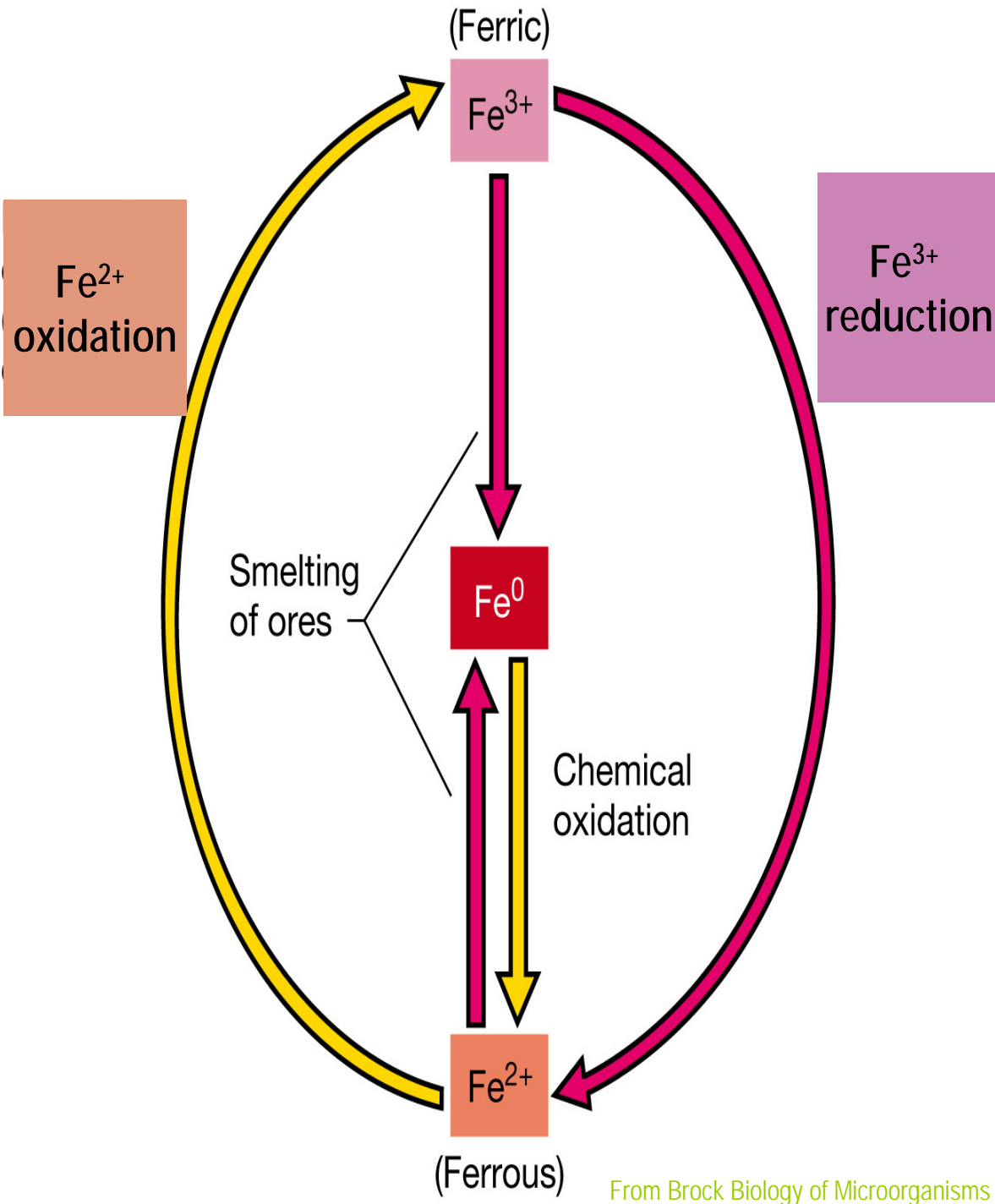


- 2 Sulphur reduction**
- Marinitoga*
  - Thermosipho*
  - Persephonella*
  - Desulfobacterium*
  - Tepidibacter*
  - Sufurospirillum*
  - Desulfurococcus*
  - Staphylothermus*
  - Pyrodictium abyssi*
  - Thermococcus*
  - Pyrococcus*
  - Palaeococcus*
  - Balnearium*
  - Thermovibrio*
  - Deferribacter*
  - Caminibacter*
  - Nautilia*
  - Sulfurimonas*
  - Hydrogenimonas*
  - Lebelimonas*
  - Ignicoccus*
  - T° max 97°C

- 3 Aerobic (microaerophilic) S° & thiosulphate oxidation**
- Persephonella*
  - T° op max 70°C

- 1 Sulphate reduction**
- Thermodesulfobacterium*
  - Thermodesulfatator*
  - Archaeoglobus*
  - T° op max 82°C





From Brock Biology of Microorganisms

# Iron cycle

$\text{Fe}^{2+}$   
Oxidation  
NO

$\text{Fe}^{3+}$   
reduction

*Geoglobus*  
*Geothermobacter*  
Strain 121  
T° max  
up to 121°C!!!





## Conclusions (2)

From available microbiology data,  
at elevated temperatures existing at deep-sea  
in hydrothermal vent chimneys,  
C, N, S and Fe cycles do not work completely.  
Particularly,  
methane, ammonium, and ferrous iron are not oxidized

# WANTED



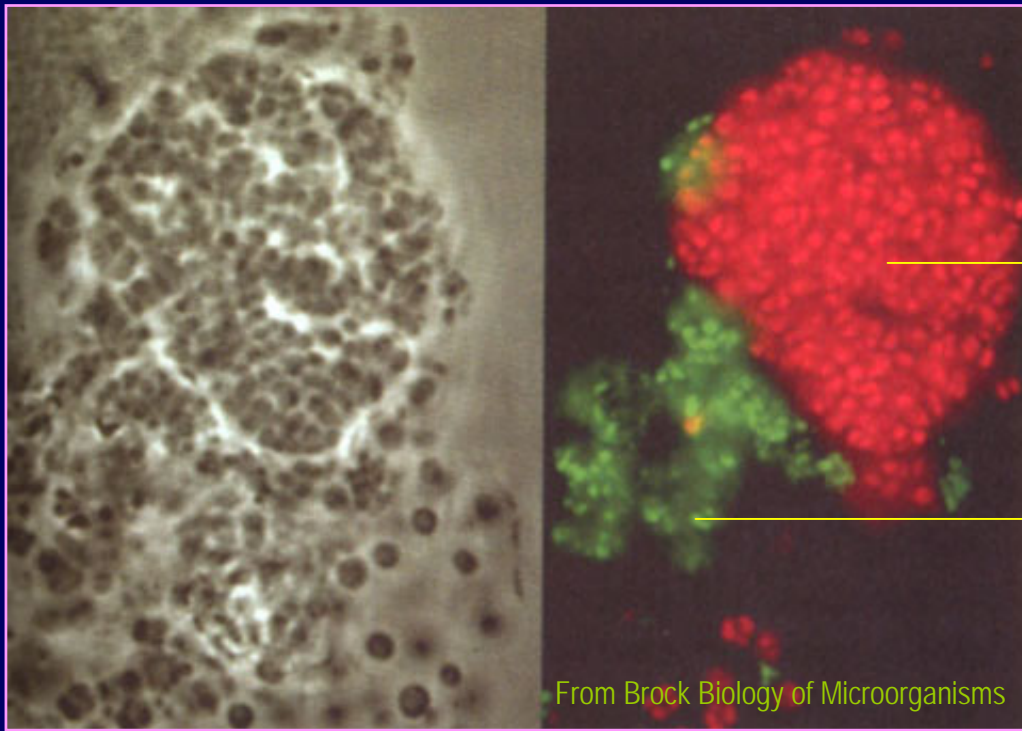
- Methanotrophs (aerobic oxidation of methane)
- Homoacetogens
- Ammonium-oxidizers (both aerobic and anaerobic)
- Sulphide-oxidizers
- $\text{Fe}^{2+}$  -oxidizers

# How to catch them ! (if they exist...)



Polyphasic approaches consisting of:

- *In situ* and "on board" activity measurements
- Phylogenetic and Functional gene analysis
- Innovative cultural approaches
  - Gradient culture (Winogradsky columns)
  - Microbial community cultivation in bioreactors
  - In situ* enrichment culture
  - High throughput cultivation techniques



FISH (Fluorescence  
*in situ* Hybridization)

Ammonium-oxidizers

Nitrification

Nitrite-oxidizers

From Brock Biology of Microorganisms

Developping co-culture techniques

# Microbial community cultivation in bioreactor

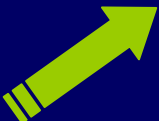
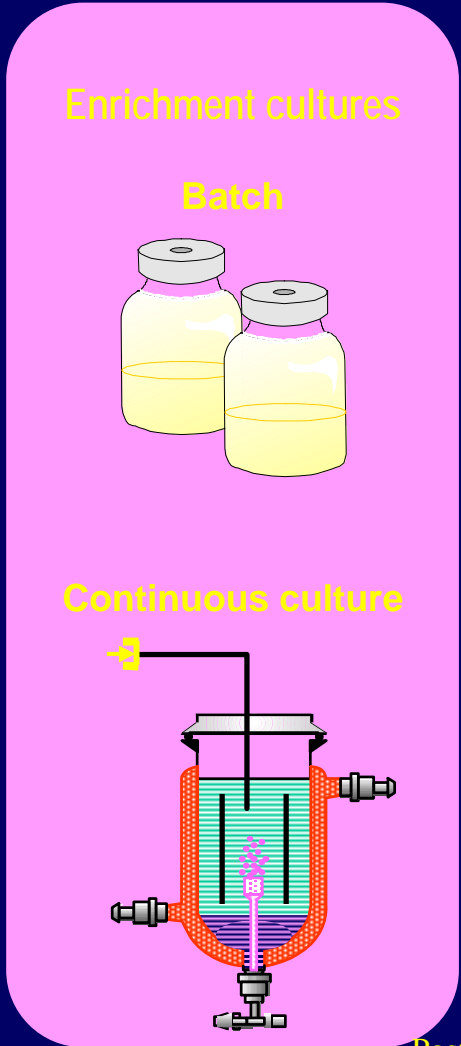


Hydrothermal chimney sample

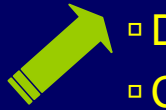
Inoculation

**Culture conditions**  
**Temperature**  
**pH**  
**Gas sparging**  
(electron donors and acceptors)  
**Dilution rate**  
(substrates concentration)

**Medium composition**  
Carbon sources  
(nature and concentration)  
Electron donors  
and acceptors



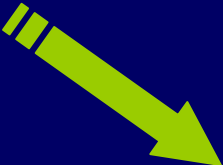
Microscopic observation  
Phase contrast  
epifluorescence



Molecular analysis :  
16SrRNA genes diversity  
□ DGGE/SSCP  
□ Cloning/sequencing  
□ Phylogenetic analysis  
□ *In situ* hybridization  
□ Quantitative PCR



HPLC analysis  
of medium  
GC analysis of  
gas exhaust



Sub-cultures and  
strains isolation

Postec et al., *Extremophiles* in press (2007).  
Postec et al., *Current Microbiology* 50, 138 (2005).

- Efficiency of such system to recover a largest diversity of cultivated thermophilic and hyperthermophilic microorganisms from a deep-sea chimney, compared to traditional cultures in vial
  - New species
  - Cultivate the uncultivable
- Efficiency to cultivate microbial communities
  - Population dynamics studies
  - Interactions between microbial populations
  - Influence of various parameters on cultivated microbial communities

# Conclusions (final)

- At elevated temperatures (at deep-sea vents) C,N,S Fe cycles do not work completely
- Novel approaches (including those suggested here) should contribute to fill the gaps
- In case of failure, it could be concluded that the high temperature ecosystems must rely on lower temperature ecosystems for recycling some compounds
- But, let's work first in a multidisciplinary approach to get more data!!