

# MICROBIOLOGY OF EXTREME ENVIRONMENTS

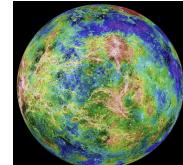
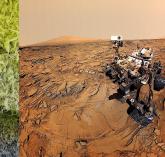
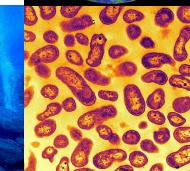
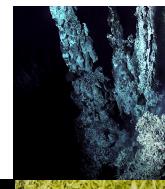
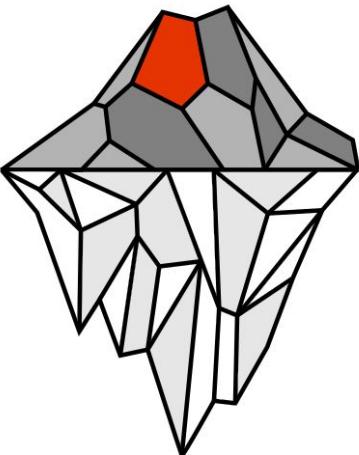
**Pressure as a parameter for bacterial growth, Xerophiles and radiation resistant extremophiles**

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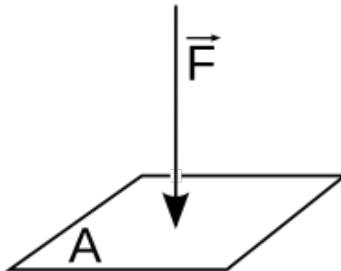




Class attendance form for contact tracing

# Pressure

Pressure is the force applied perpendicular to the surface of an object per unit area over which that force is distributed



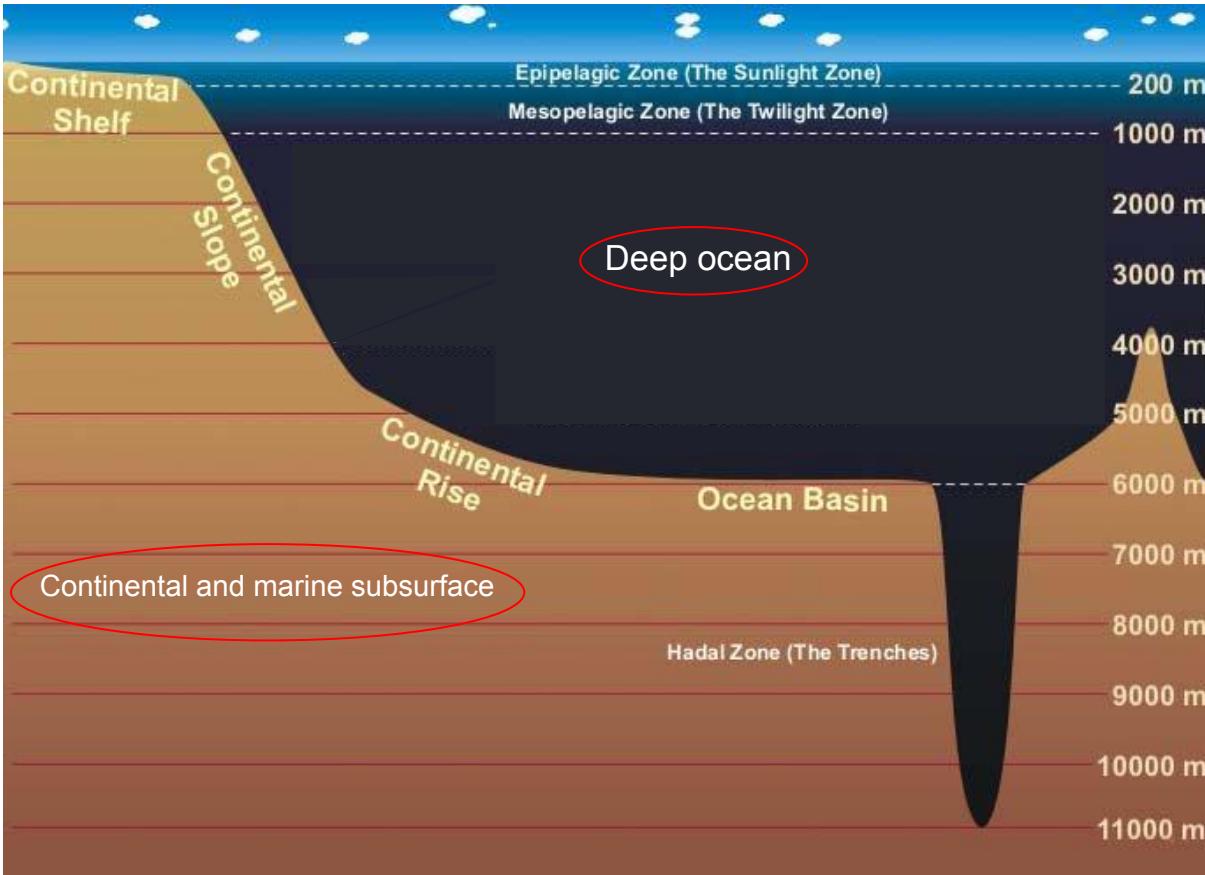
SI unit for pressure is **Pascal**

Atmospheric pressure (1 atm) = 1 bar = 101,325 Pa = **0.1 MPa**

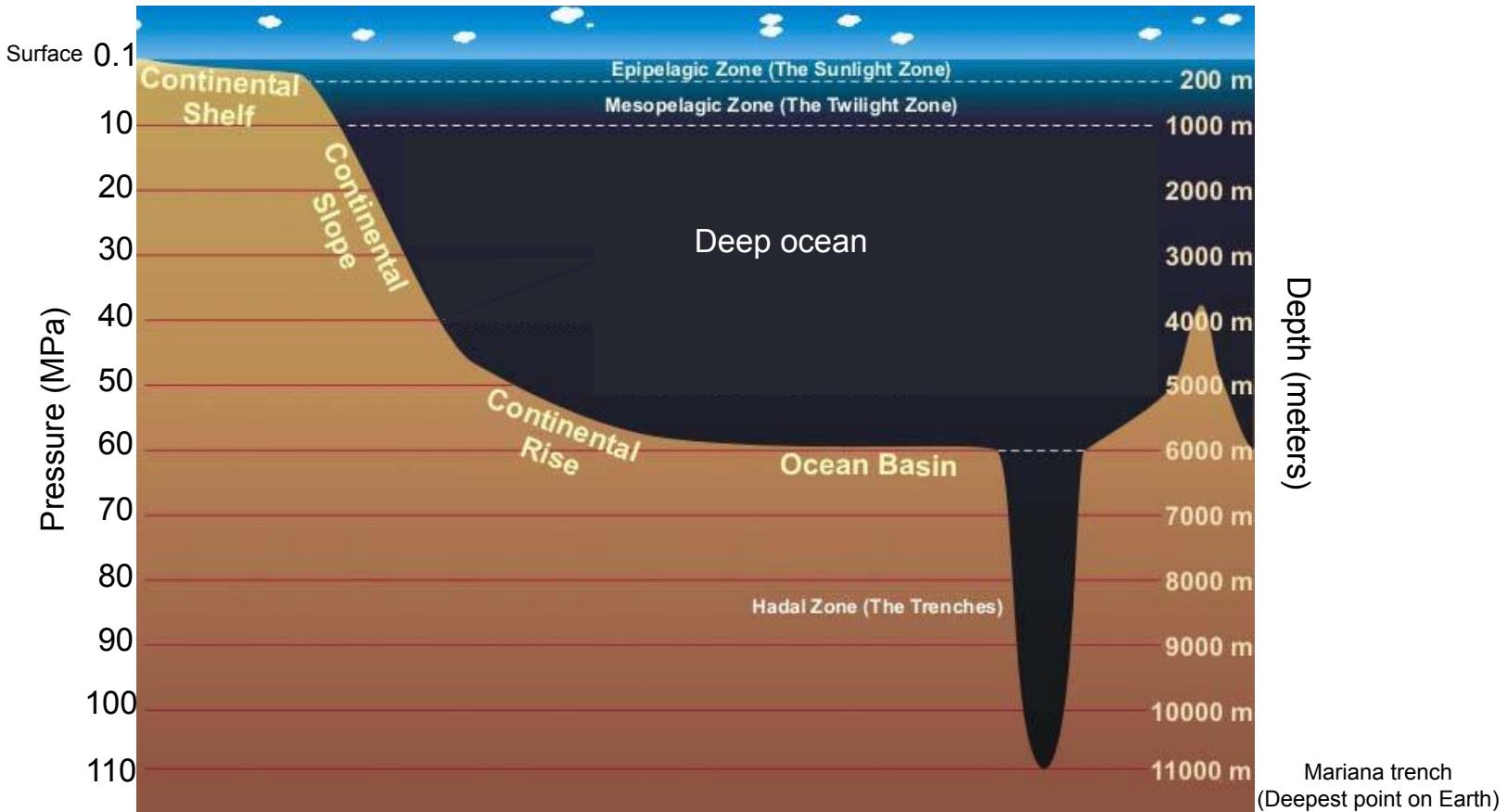
Hydrostatic pressure = 10 MPa/km

Lithostatic pressure = 25 MPa/km

# High pressure environments

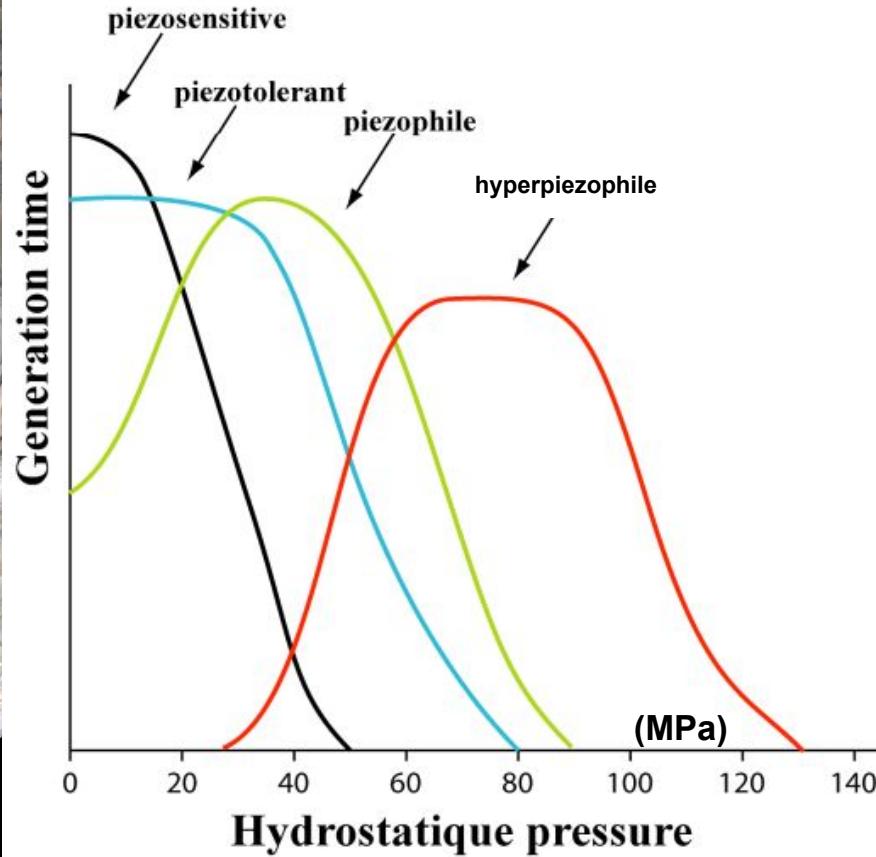


# Pressure range on Earth' surface



# Pressure effect on microbes growth

Piezo derive from Greek and it means to press

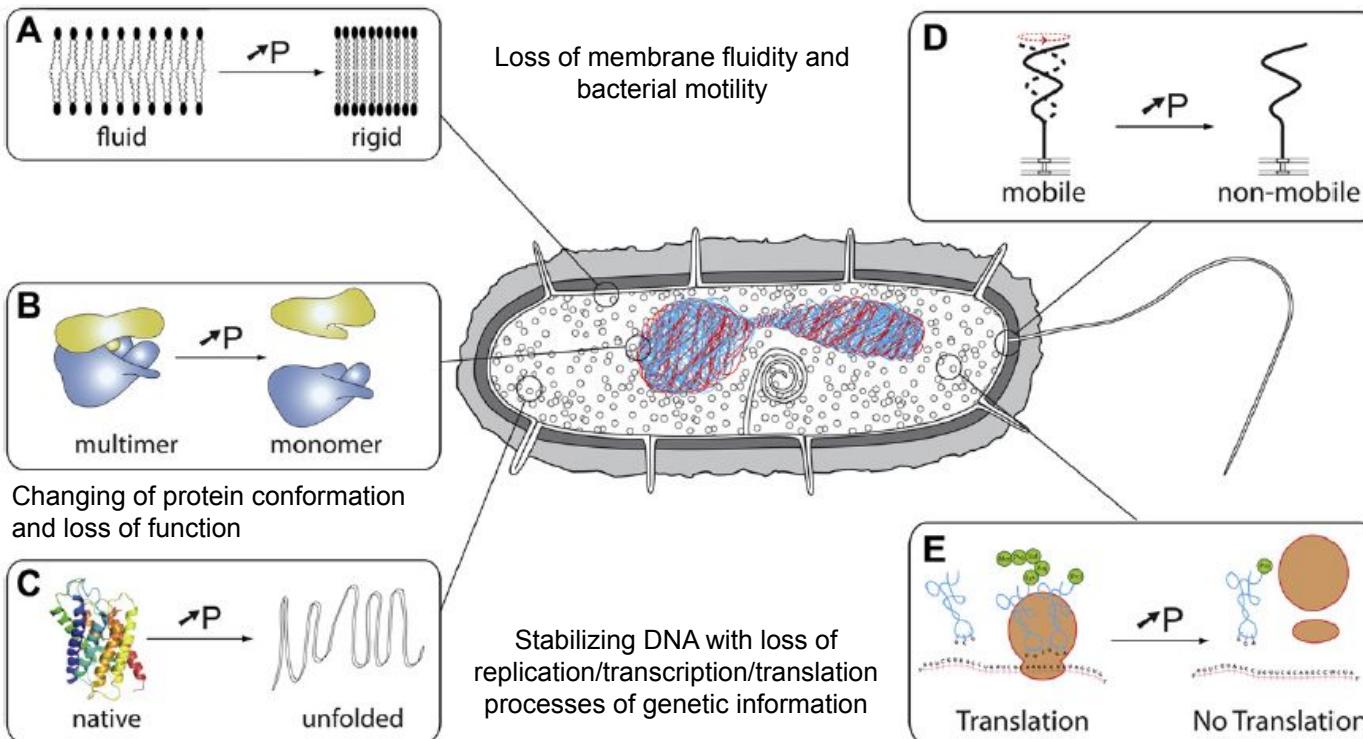


- Piezosensitive → 0.1 MPa
- Piezotolerant → 0.1 - 10 MPa
- Piezophile → 10 - 50 MPa
- Hyperpiezophile → >50 MPa

# Effect of pressure on microbes

Conformational changing of macromolecules with subsequent loss of function

Does not have effect on covalent bonds (e.g. Temperature)



# Effect of pressure on microbes

Table 1

Cellular processes/structures impaired by high hydrostatic pressure in *E. coli*.

Process	Pressure-abolishing process (MPa)	Reference
Motility	10	Meganath and Marquis (1973)
Substrate transport (isopropylthiogalactopyranoside)	26	Landau (1967)
Cell division	20–50	Zobell and Cobet (1962, 1963)
Growth	50	Yayanos and Pollard (1969)
DNA replication	50	Yayanos and Pollard (1969)
Translation	60	Yayanos and Pollard (1969)
Transcription	77	Yayanos and Pollard (1969)
Viability	200	Pagan and Mackey (2000)

Most of what we know about the effect of pressure on bacteria comes from experiment with *Escherichia coli*, a **mesophilic piezotolerant bacterium** → **DEFINITELY NOT AN EXTREMOPHYLE**

Piezophilic microbes are difficult to isolate and cultivate

# Strategies to cope with extreme pressures

Cell membrane packed with unsaturated fatty acids → Increase membrane fluidity at high pressure

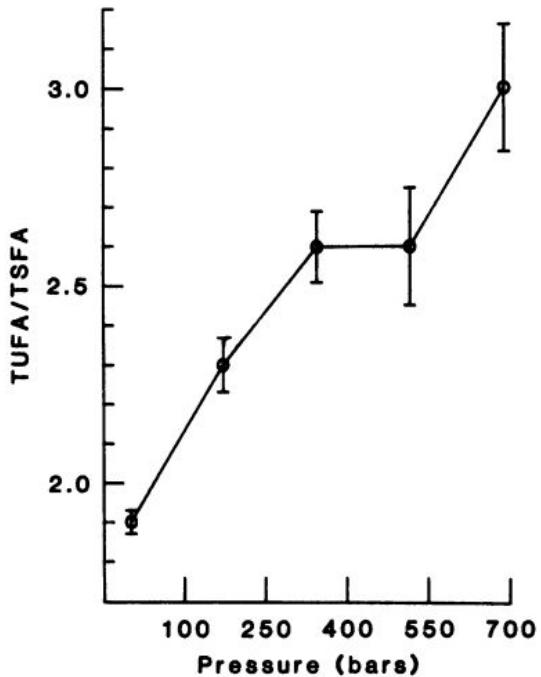
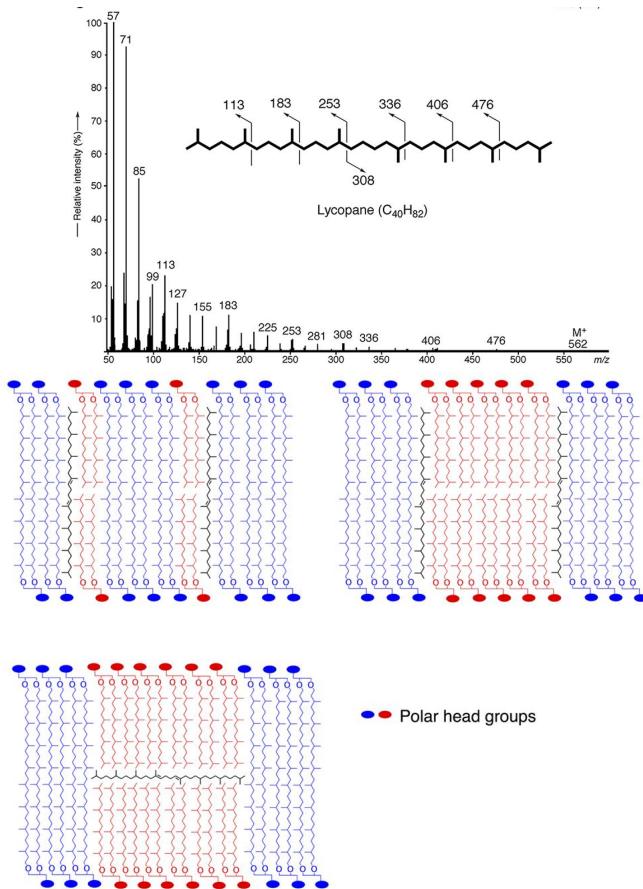


Fig. 1. Ratio of total unsaturated fatty acids (TUFA) to total saturated fatty acids (TSFA) as a function of pressure. Data points are means  $\pm$  standard errors for five separate experiments.



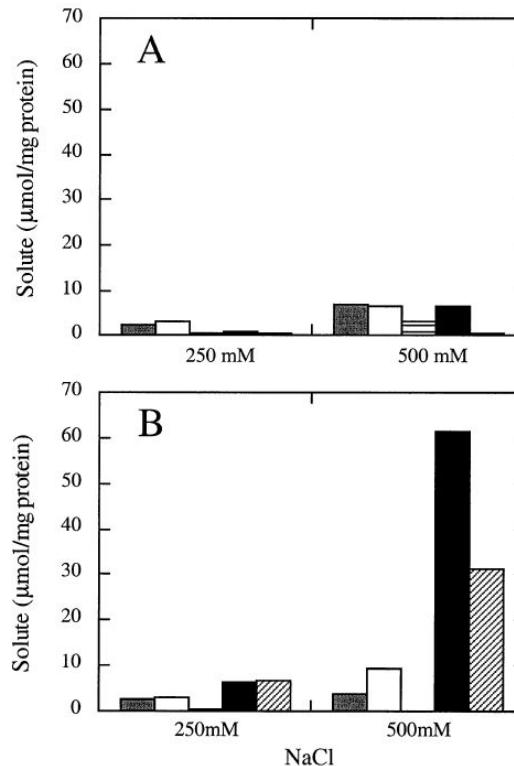
# Strategies to cope with extreme pressures

## Intracellular salt content and osmolyte regulation

At high pressure and high salt content, bacteria increase the intracellular concentration of osmolytes

$\beta$ -hydroxybutyrate ( $\beta$ -HB)

High concentration of osmolyte → integrity of the cell



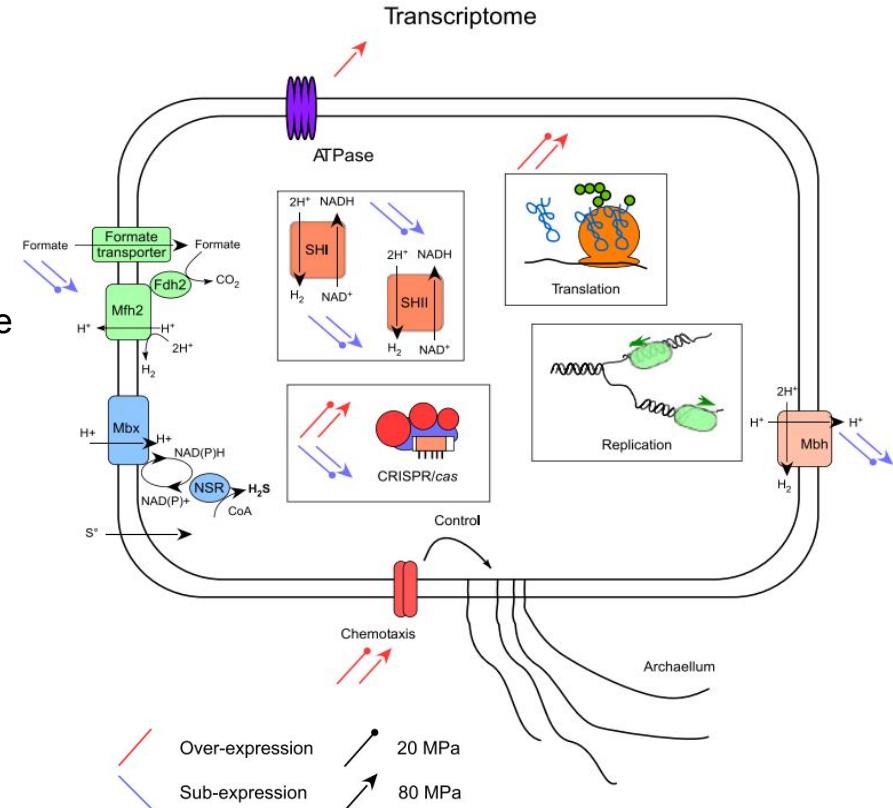
**Fig. 5.** Distribution of intracellular solutes in *P. profundum* cells grown at **A** 0.1 MPa or **B** 28 MPa to early stationary phase as a function as external NaCl: ■, betaine; □, glutamate; ▨, alanine; ■,  $\beta$ -HB; ▨,  $\beta$ -HB oligomer

# Strategies to cope with extreme pressures

Specific high-pressure gene expression

Under high pressure:

- ↓ Mfh2 hydrogenases → repression of H<sub>2</sub> metabolism
- ↑ Mbx sulfur-dependant hydrogenases → alternative energy supply
- ↑ chemotaxis machinery → due to change in membrane fluidity
- ↑ Translation genes → due to disruptive effect of pressure on ribosomes

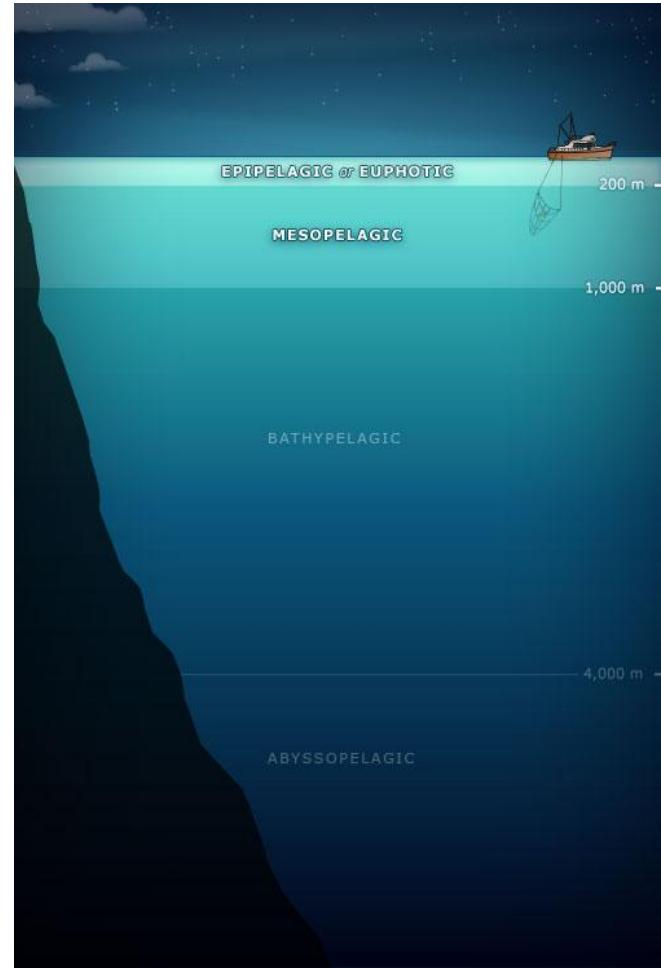




# **WHERE DO PIEZOPHILIC MICROORGANISMS LIVE?**

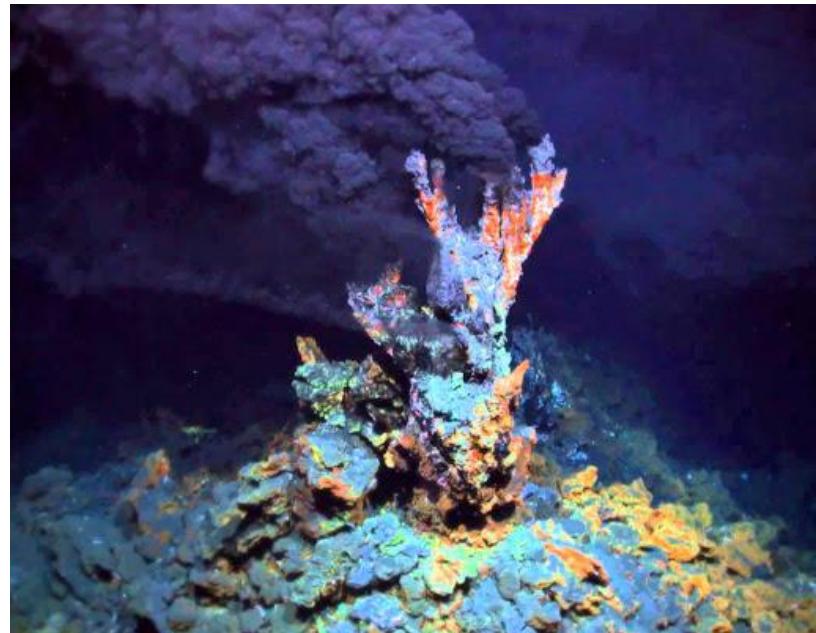
# Piezophiles in cold deep ocean

- Deep-sea covers 65% of Earth's surface and holds 88% of total biomass
- Only 5% percent of the oceans have been explored and the 0.01% has been sampled
- Average depth of the ocean → 3,800 meters
- Average pressure in the ocean → 38 MPa
- Average deep ocean temperature → 2-4 °C
- No sunlight → No photosynthesis → Chemolithoautotrophy
- Oligotrophic water = poor in nutrients
- Psychrophilic piezophilic Bacteria and Archaea



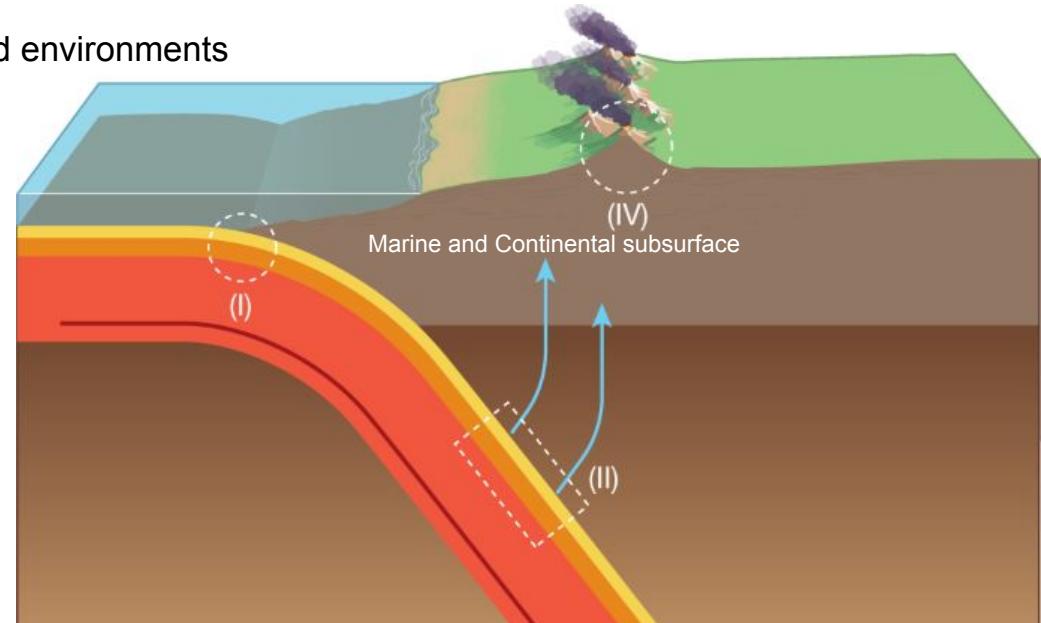
# Piezophiles in deep-sea hydrothermal vents

- Environmental fluctuations (e.g. Temperature, salinity, pH)
- Temperatures range between 460 and 2 °C in just a few cm
- Salinity ranges from 0.1 % to 8 %
- No sunlight → No photosynthesis → Chemolithoautotrophy
- Hyperthermophilic piezophilic Archaea and Bacteria



# Subsurface environments

- It is estimated that the subsurface holds about 15% of the total Earth's biomass
- Here, pressures >110 MPa have been recorded
- Highest pressure = 900 MPa at the top of a subducting plate, Mariana Forearc
- Resident microbial communities play important roles on mediating biogeochemical cycles
- However, it is one of the most unexplored environments





**Are we forgetting something?**



# Low-pressure environments

- Mount Everest → highest peak on Earth with the lowest pressure on the planet (0.0033 MPa)
- Low pressure seems to not have detrimental effect on microbes
- Space → lowest pressure. It ranges between  $10^{-13}$  and  $10^{-10}$  MPa (space vacuum)
- Long exposure to space vacuum cause dehydration of DNA





**WHO ARE THEY?**

# Piezophilic isolates

To date, only about 56 facultative and obligate piezophile, both Archaea and Bacteria, are known

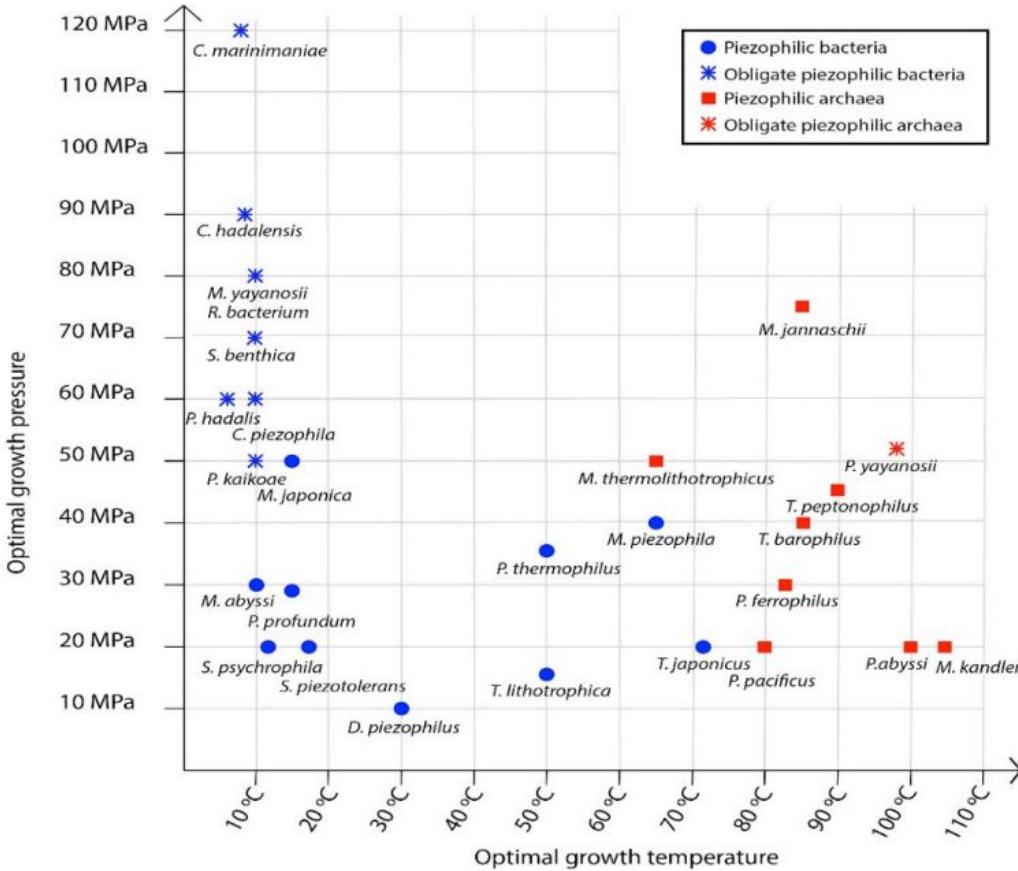
## Archaea

- Euryarchaeota
- Crenarchaeota
- Asgardarchaeota

## Bacteria

- Gammaproteobacteria (*Colwellia*, *Moritella*, *Photobacterium* and *Shewanella*)
- Deltaproteobacteria (*Desulfovibrio*)
- Epsilonbacteria (*Marinitoga*)
- Bacilli (*Carnobacterium*)

# Pressure Vs Temperature



# Pressure Vs Temperature

Polyextremophile

**Table 1. Classification scheme of piezophiles based on optimal growth temperature and pressure<sup>a</sup>**

$P_{kmax} \setminus T_{kmax}$	< 15 °C	15–45 °C	45–80 °C	>80 °C
Piezotolerant (< 10 MPa)	Psychro-piezotolerant	Meso-piezotolerant	Thermo-piezotolerant	Hyperthermo-piezotolerant
Piezophilic (10–50 MPa)	Psychro-piezophile	Meso-piezophile	Thermo-piezophile	Hyperthermo-piezophile
Hyperpiezophilic (> 50 MPa)	Psychro-hyperpiezophile	Meso-hyperpiezophile	Thermo-hyperpiezophile	Hyperthermo-hyperpiezophile

# ***Colwellia* sp., first obligate piezophile psychrophilic bacterium**

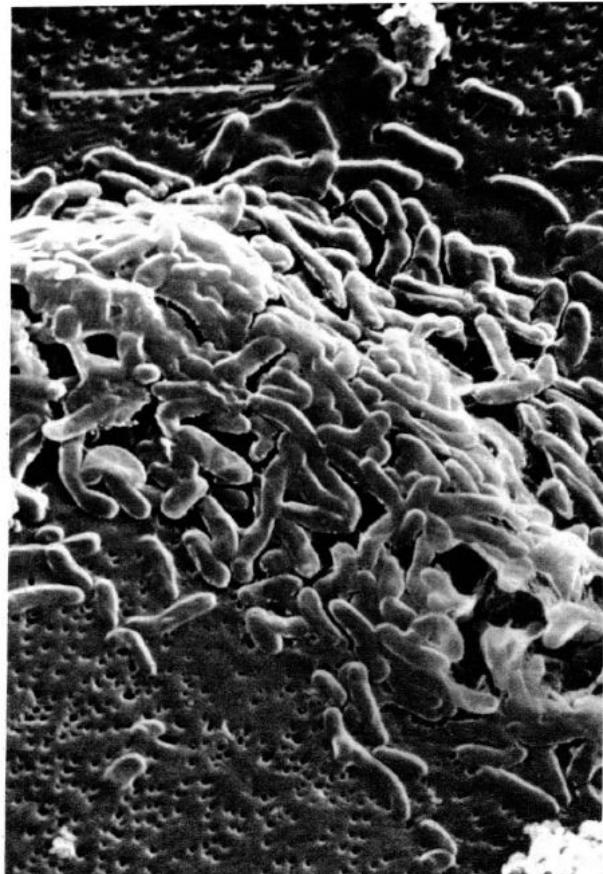
Gram negative bacterium

Strain MT-41

Isolated from a decaying amphipod found at the bottom of the Mariana trench  
(11,000 meters depth)

Optimal growth pressure = 70 MPa

Optimal growth temperature = 2 °C



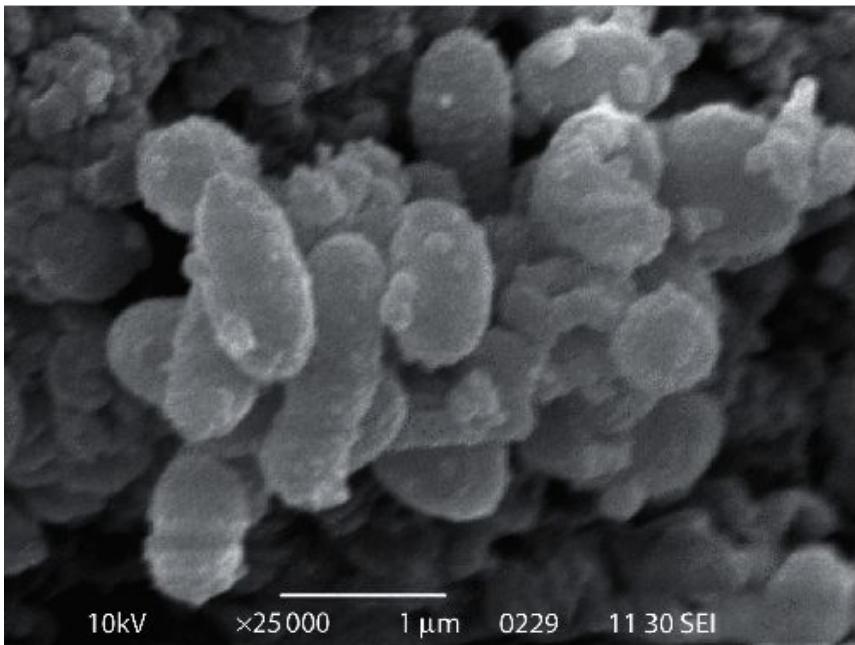
# *Colwellia marinimaniae*, obligate piezophile bacterium with a record

Gram negative bacterium

Rods shape

Optimal growth pressure = 120 MPa

Highest growth pressure = **140 MPa**



# *Pyrococcus yayanosii* – first obligate piezophilic hyperthermophilic archaeon

Archaeon

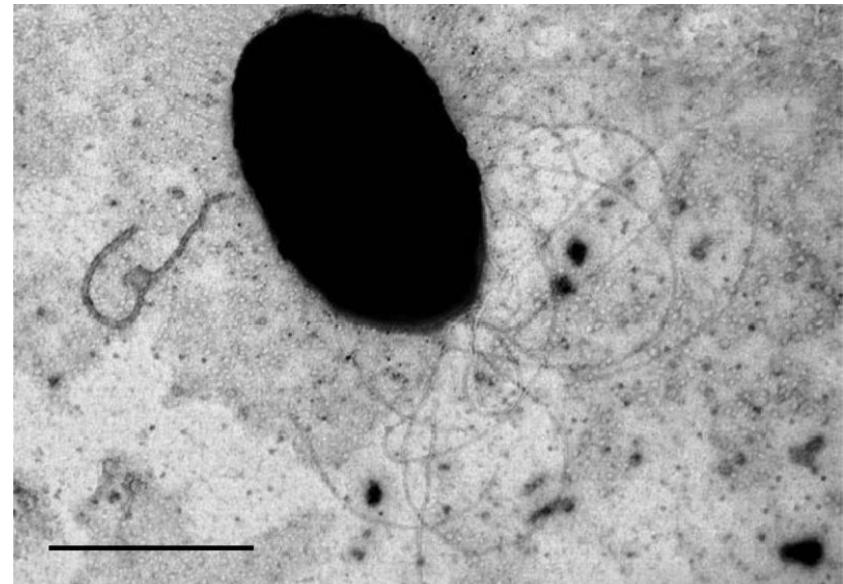
Strain CH1

Isolated from Ashadze site (Mid-atlantic ridge), the deepest hydrothermal vent field explored so far (4 km depth)

Optimal growth pressure = 52 MPa

Optimal growth temperature = 98 °C

Highest pressure growth = 120 MPa

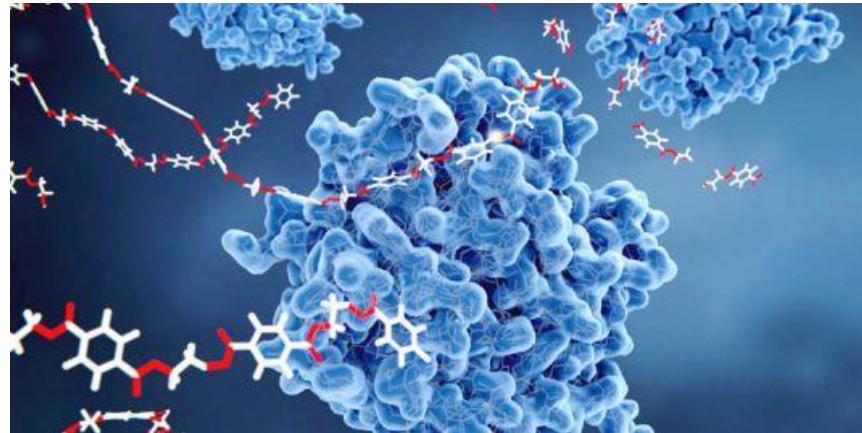


# Biotech applications and advancements

Food industry



Extremozymes

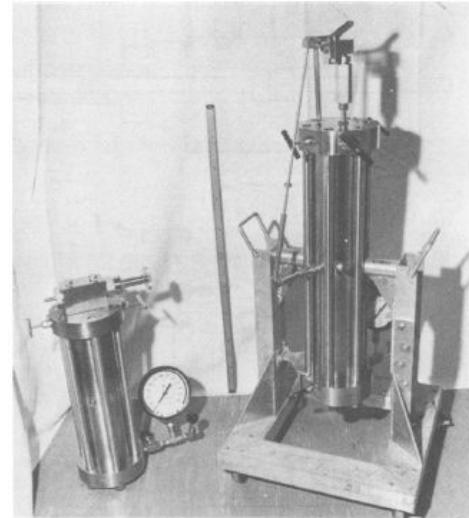


# Biotech applications and advancements

Deep-sea exploration



Pressure-retaining samplers



Jannasch et al. 1976



# Suggested readings

- Lauro and Bartlett (2008) - Prokaryotic lifestyles in deep sea habitats
- Oger and Jebbar (2010) - The many ways to cope with pressure
- Jebbar et al. 2005 - Microbial diversity and adaptation to high hydrostatic pressure in deep-sea hydrothermal vents prokaryotes
- Cario et al 2019 - Exploring the Deep Marine Biosphere: Challenges, Innovations, and Opportunities

# Xerophiles

Microorganisms that have adapted to survive in environments in which water is scarce like hyper-arid deserts and desiccated foods.



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- Archaea



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- Archaea
- Bacteria



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- Archaea
- Bacteria
- Fungi



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# Xerophiles



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~~Xerophiles~~

Xerotolerant





# Xeric stress limits

A xeric stress can be caused either by:



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- **Desiccation**, lack of water



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# Xeric stress limits

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- **Desiccation**, lack of water
- **Hypertonicity**, excessive solute concentrations in the surrounding environment that removes water from cells

**How do we measure the amount of  
water available in an environment?**

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# Xeric stress limits



Water activity

# Xeric stress limits

**Water activity**

The vapor pressure of water in a system divided by that of pure water at the same temperature



# Xeric stress limits

**Water activity**

The vapor pressure of water in a system divided by that of pure water at the same temperature

It's a measure of how much of that water is free and unbound, and thus available for microorganisms to use for growth

# Xeric stress limits

Species	Lowest water activity ( $a_w$ ) for cell division	Environmental source	Refs
<b>Bacteria and archaea</b>			
<i>Haloarchaea GN-2</i>	0.635	Solar salterns, Mexico	3
<i>Haloarchaea GN-5</i>	0.635	Solar salterns, Mexico	3
<i>Halorhabdus utahensis</i> DSM 12940	0.647	Salt Lake, USA	3
<i>Halobacterium</i> strain 004.1	0.658	Brine pool, UK	3
<i>Halorhodospira halophila</i> DSM 244	0.66	Salt lake, USA	3
<i>Salinibacter ruber</i> DSM13855	0.725	Solar salterns, Spain	3
<i>Salisaeta longa</i> DSM 21114	0.747	Dead Sea, Israel	3
<b>Fungi</b>			
<i>Aspergillus penicillioides</i>	0.585	Raisins, Australia	7,106
<i>Xeromyces bisporus</i>	0.637	Antique wood, Thailand	3,6,7,107
<i>Aspergillus amstelodami</i> FRR2792	0.656	Dates, Australia	7
<i>Xerochrysum xerophilum</i> FRR 0530 (formerly known as <i>Chrysosporium xerophilum</i> )	0.686	High-moisture prunes, Australia	7
<i>Aspergillus chevalieri</i> PIL 119	0.71	Soiled prunes, Australia	7

Lebre, P. H., De Maayer, P., & Cowan, D. A. (2017). *Nature Reviews Microbiology*, 15(5), 285-296.



# Where?



B

Hyper-arid deserts





# Where?



B

Hyper-arid deserts



# Where?



Salt-cured food



B



Hyper-arid deserts



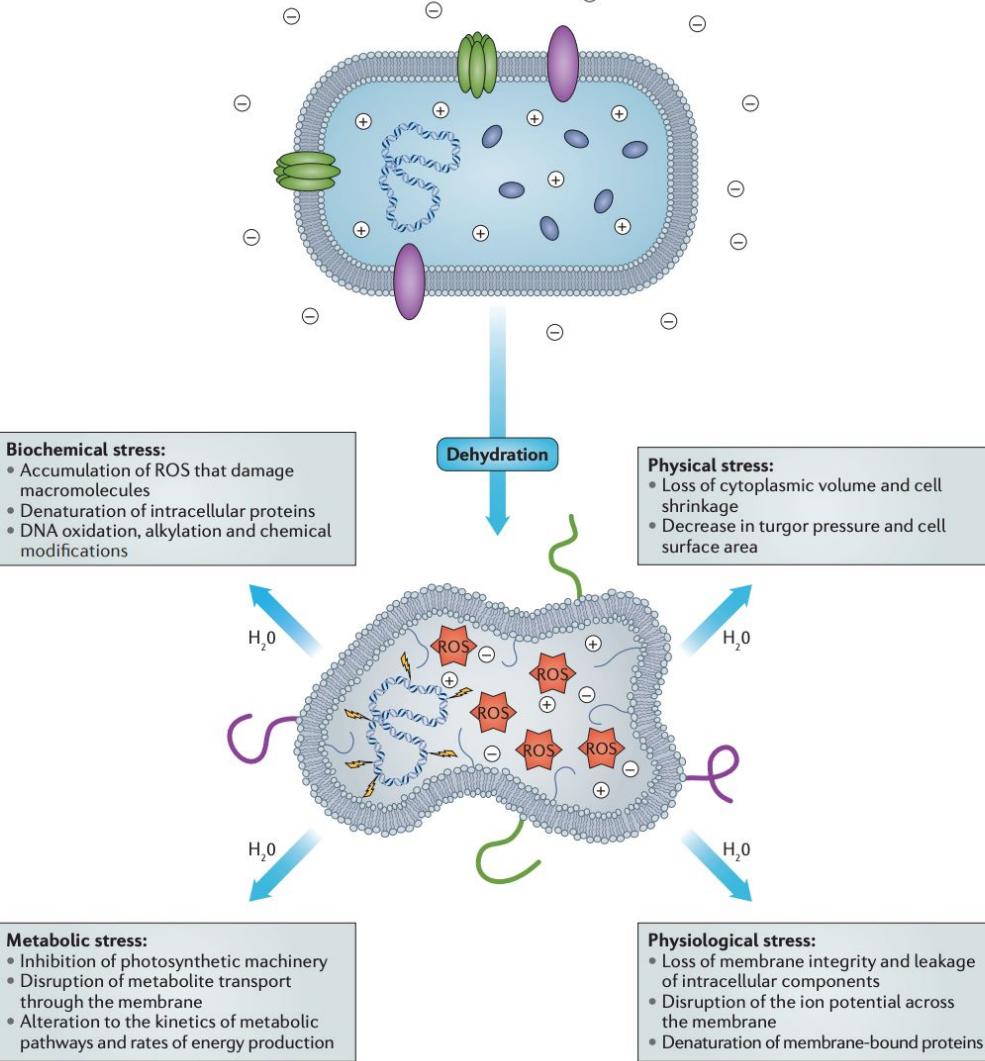
Hyper-arid deserts



Hypersaline aquatic environments



**What happens when the water activity  
is too low?**



Lebre, P. H., De Maayer, P., & Cowan, D. A. (2017). *Nature Reviews Microbiology*, 15(5), 285-296.



# How do they survive?

To counteract the negative morphological, physiological and biochemical consequences of desiccation, xerotolerant microorganisms have developed a broad range of adaptive strategies.



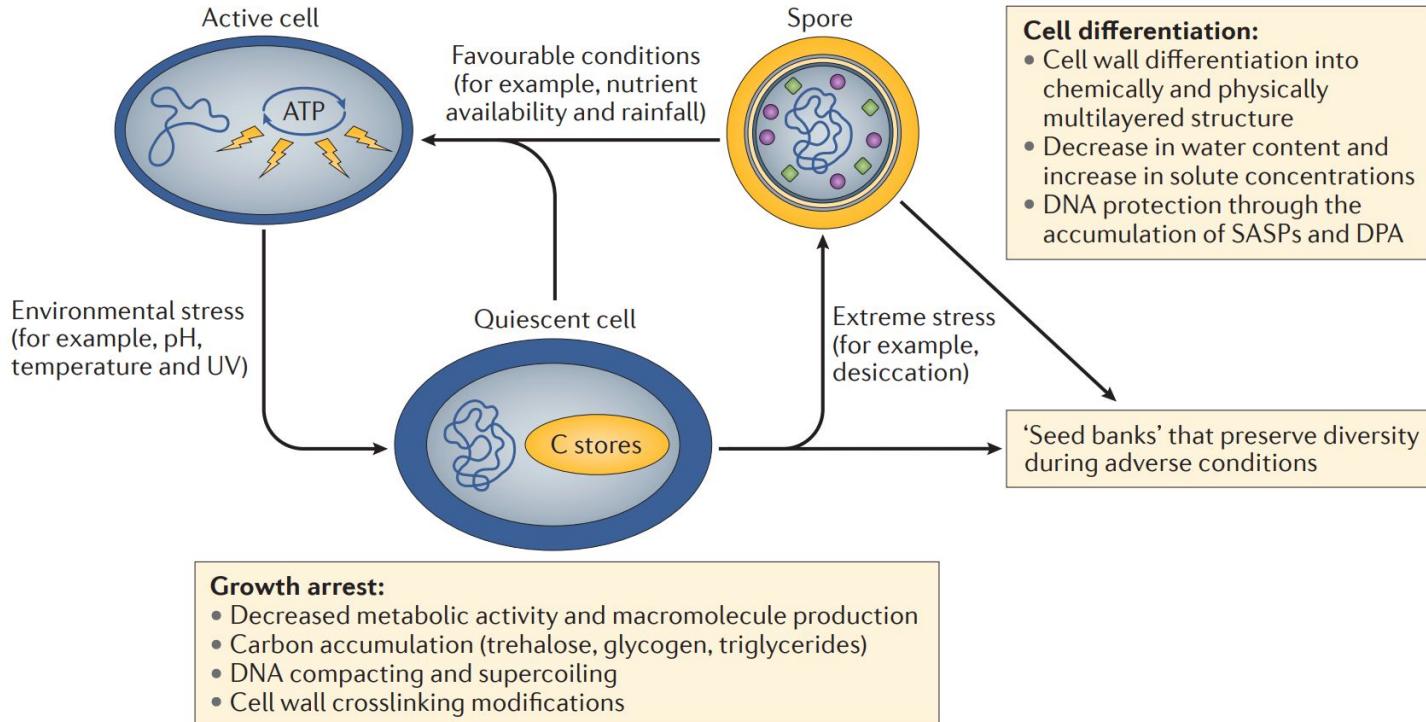
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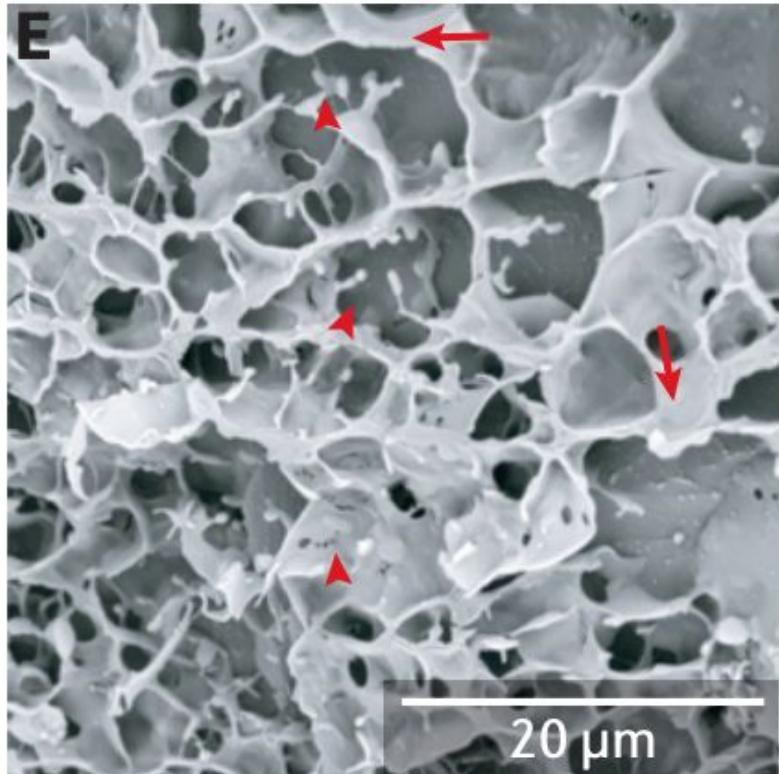
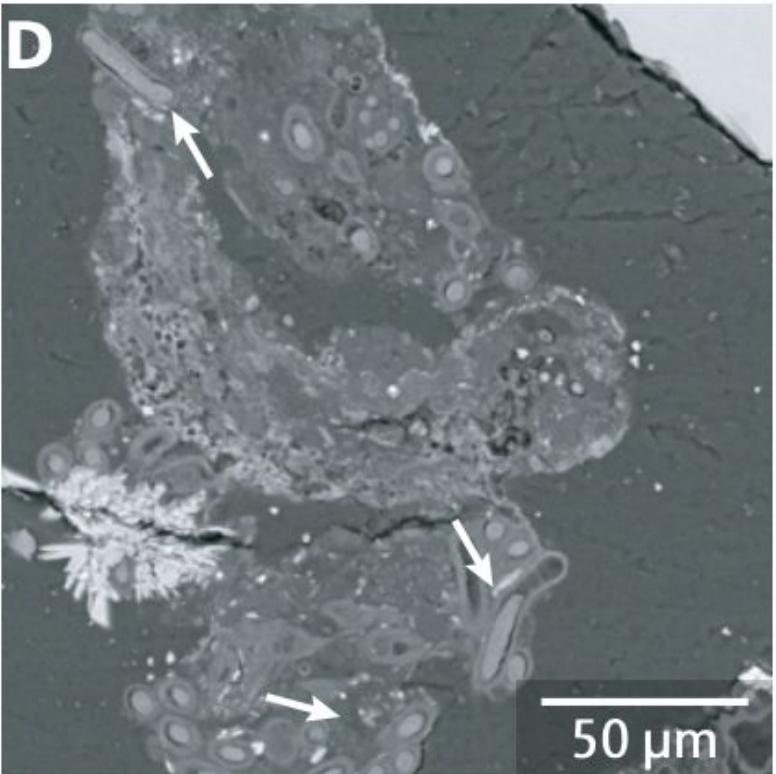
They have evolved a series of **behavioural, physiological and molecular mechanisms**

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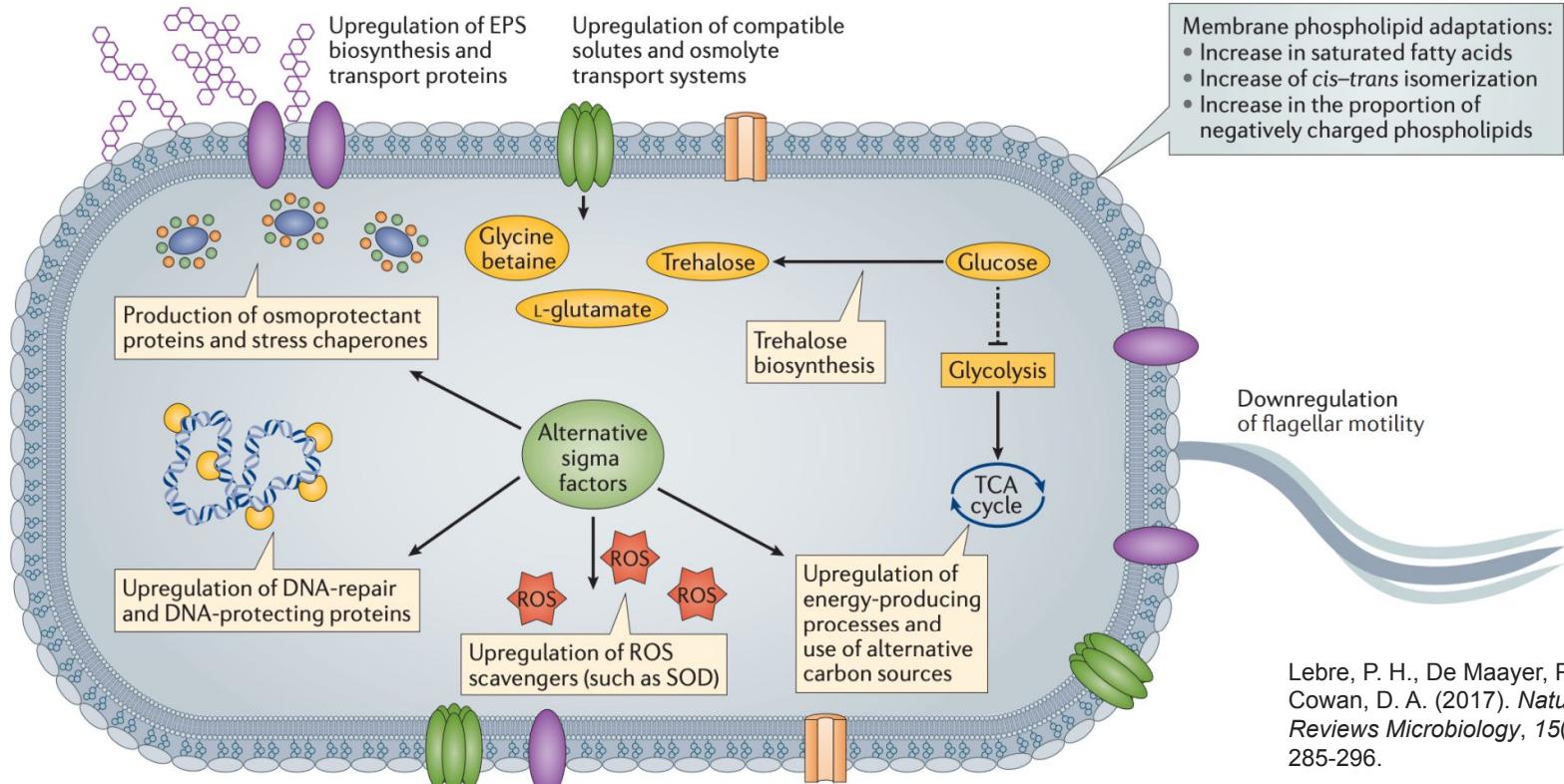
# Behavioural adaptations: Dormancy and sporulation



# Behavioural adaptations: EPS and biofilm formation



# Physiological adaptations: Cell membrane adaptations, accumulation and biosynthesis of compatible solutes



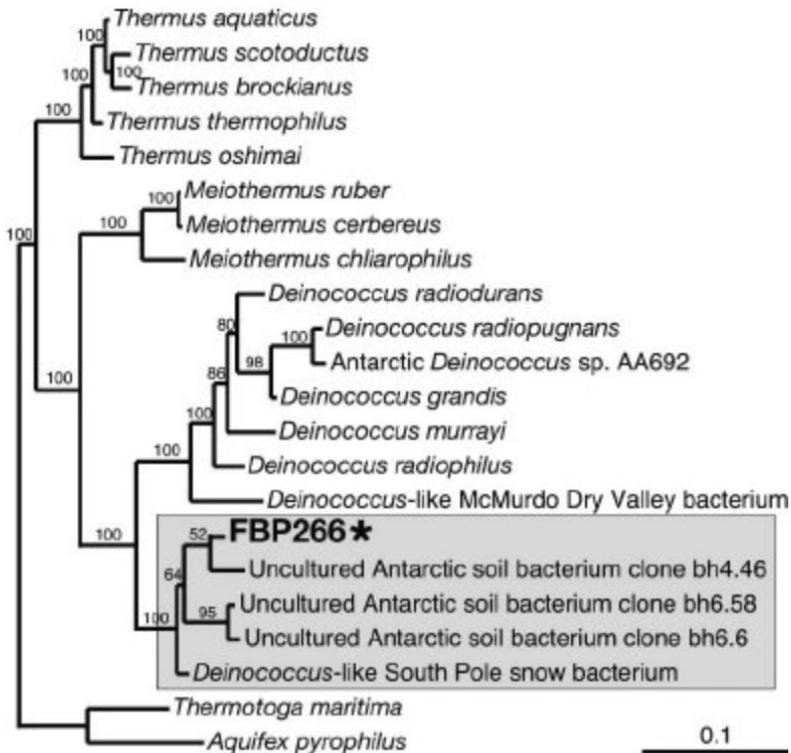


# Metabolic and molecular adaptations

- They can differentially upregulate and downregulate various metabolic pathways, which can lead to a shift from anabolic to catabolic metabolism
- They can express a range of proteins to counteract the effects of low water activity es. Late embryogenesis abundant (LEA)

# Thermus/Deinococcus phylum

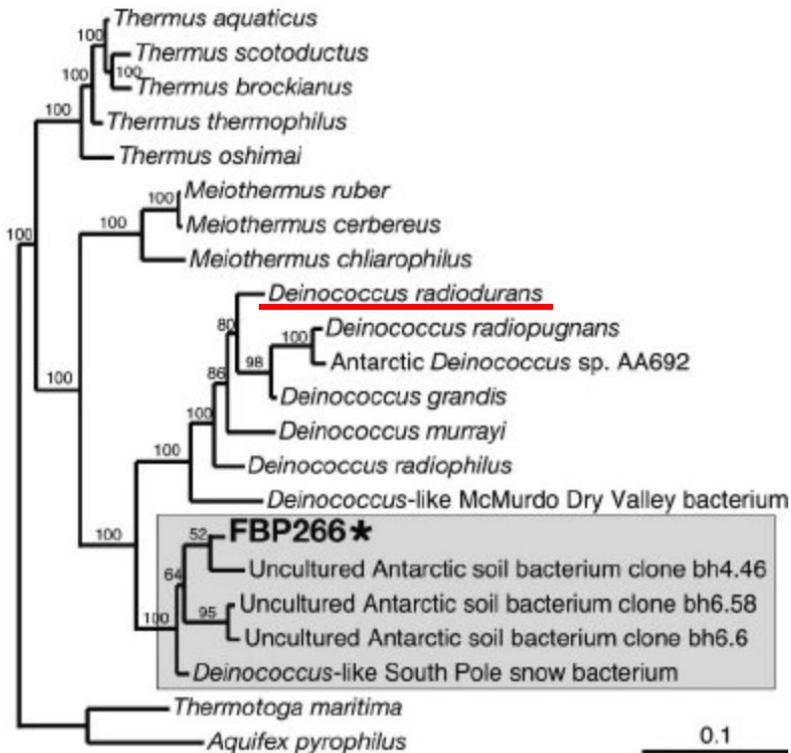
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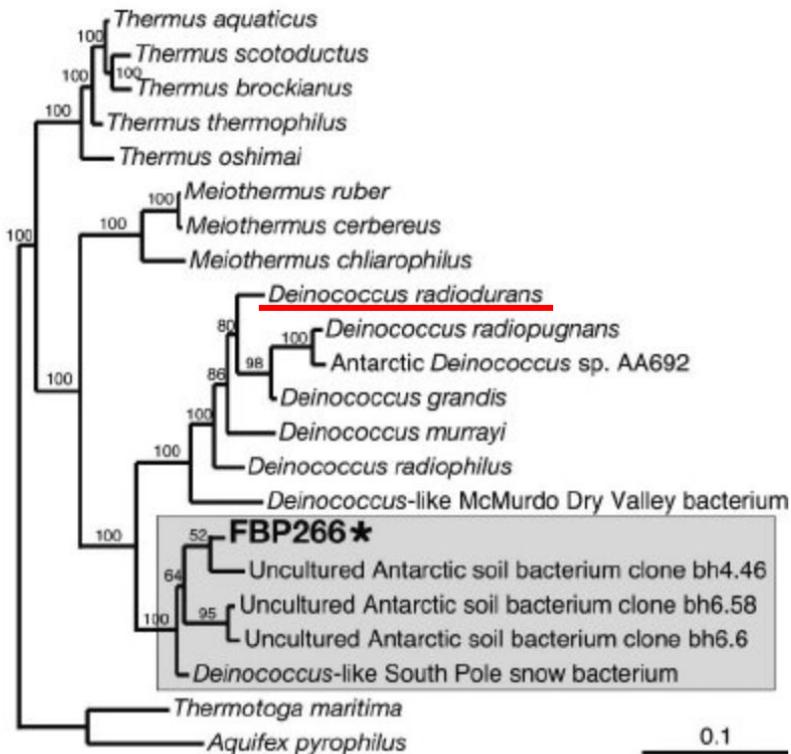
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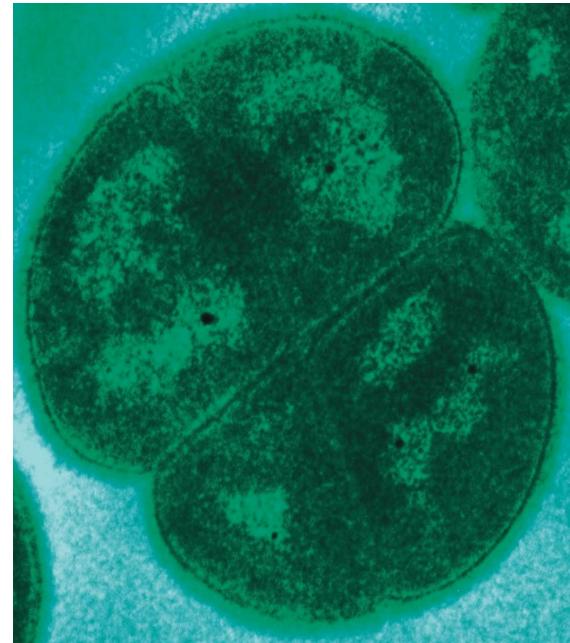
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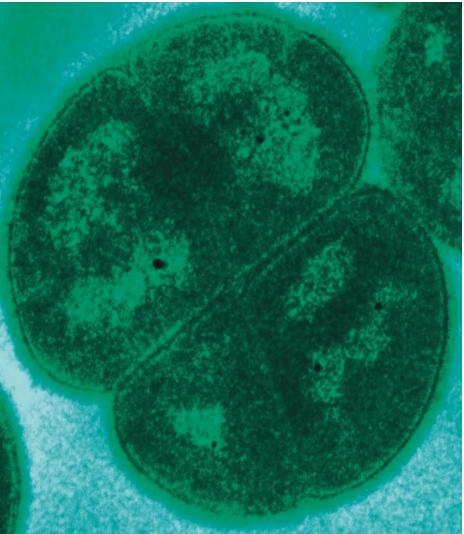


## Deinococcus radiodurans



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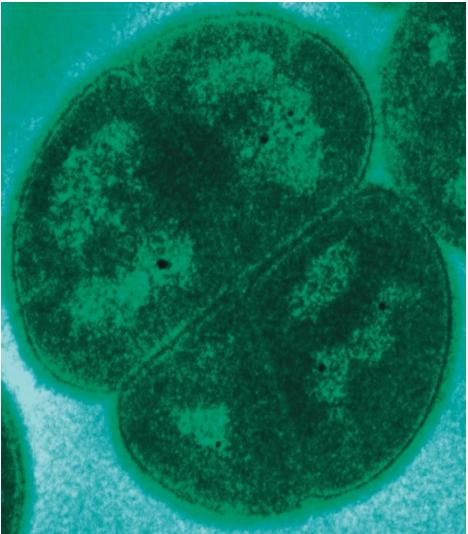
# *Deinococcus radiodurans*



It is a **polyextremophile**:

- one of the most **radiation-resistant** organisms known
- It can survive cold, dehydration and vacuum

# *Deinococcus radiodurans*



It is a **polyextremophile**:

- one of the most **radiation-resistant** organisms studied
- It can survive cold, dehydration and vacuum
- Humans cannot survive a whole-body exposure of 10 Gy
- 200 Gy is lethal for most bacteria
- *Deinococcus radiodurans* can survive acute exposures of 15,000 Gy and chronic exposures of 60 Gy/h



# Radiation-resistance

UV radiation, X-rays and gamma rays are different types of ionizing radiation that can impact microbial cells.

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UV radiation, X-rays and gamma rays are different types of ionizing radiation that can impact microbial cells.

## Radiation-resistance limits

- *Thermococcus gammatolerans* EJ3 and *Deinococcus hohokamensis* have been shown to resist up to 30 kGy of  $\gamma$ -radiation

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- *Psychrobacter pacificensis* L0S3S-03b can resist up to 100–1000 J/m<sup>2</sup> of UV254

# Radiation-resistance evolution

There are no naturally occurring environments known that result in exposures exceeding 400 mGy per years

**How it has evolved?**

Radiations

Other stresses  
like desiccation

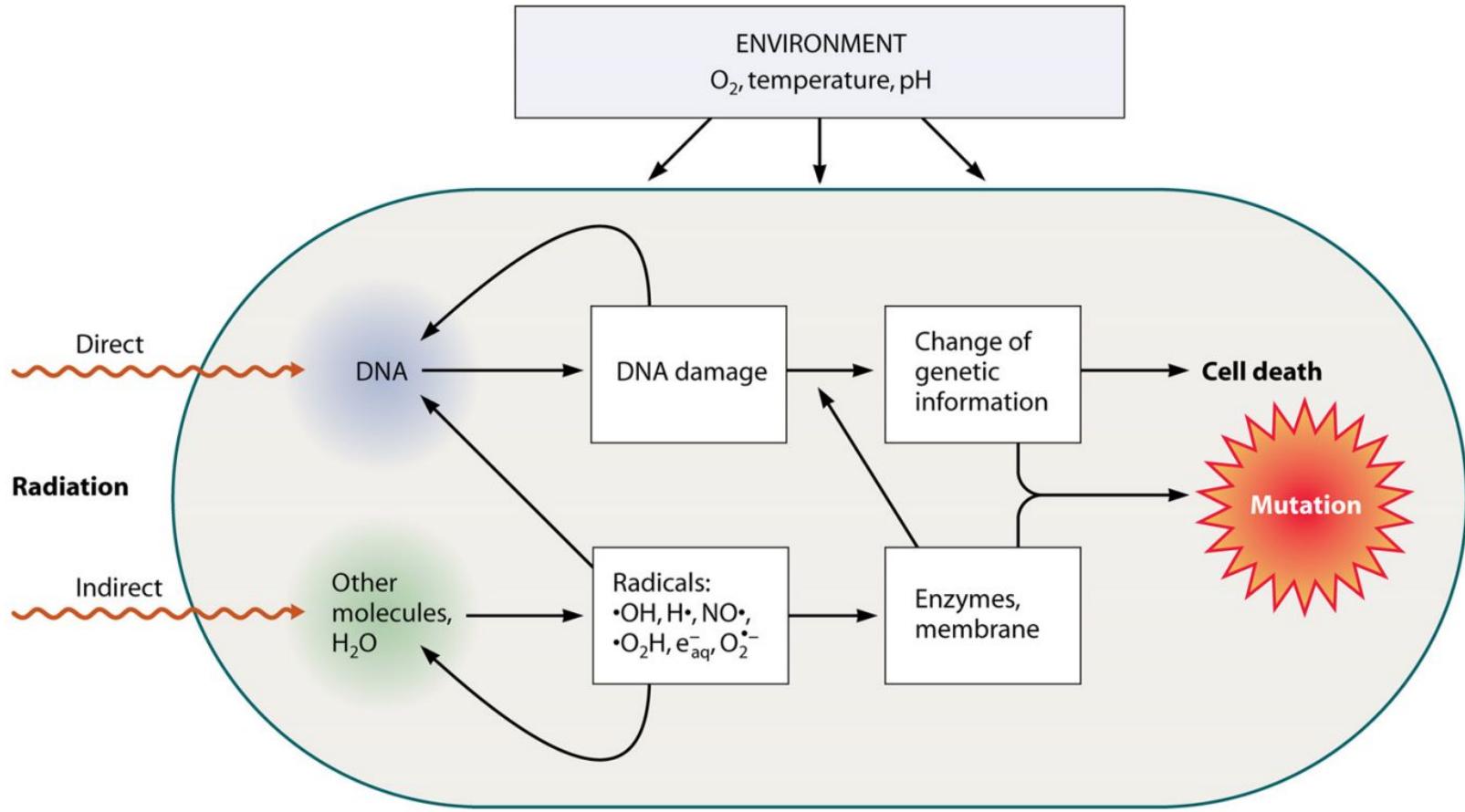
They cause similar  
damages



# The effects of ionizing radiations

The effects of ionizing radiation on living cells include:  
the formation of large numbers of **ROS**, **DNA SSBs**, **DNA DSBs** and  
extensive **base modifications**; combinations of all of these induce  
the so-called **complex (clustered) DNA damage**

A dose of 6 kGy is expected to induce approximately **200 DSBs**,  
**over 3000 SSBs** and many more sites of DNA base lesions per *D. radiodurans*' genome. In addition, no significant differences have been found in genomic damage accumulation between radioresistant and radiosensitive bacteria.



Horneck, G., Klaus, D. M., & Mancinelli, R. L. (2010). *Microbiology and Molecular Biology Reviews*, 74(1), 121-156.



# Survival strategies

They need two main response systems:



# Survival strategies

They need two main response systems:

- DNA and protein protection mechanisms



# Survival strategies

They need two main response systems:

- DNA and protein protection mechanisms
- Efficient DNA repair mechanisms to restore genome functionality



# Survival strategies

- **Small non-coding RNAs** (sRNAs) regulate DNA repair gene expression
- Genome redundancy
- Novel two-steps DNA repair processes

Published: 27 January 2009

# A new perspective on radiation resistance based on *Deinococcus radiodurans*

Michael J. Daly

*Nature Reviews Microbiology* 7, 237–245 (2009) | [Cite this article](#)

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## Abstract

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In classical models of radiation toxicity, DNA is the molecule that is most affected by ionizing radiation (IR). However, recent data show that the amount of protein damage caused during irradiation of bacteria is better related to survival than to DNA damage. In this Opinion article, a new model is presented in which proteins are the most important target in the hierarchy of macromolecules affected by IR. A first line of defence against IR in extremely radiation-resistant bacteria might be the accumulation of manganese complexes, which can prevent the production of iron-dependent reactive oxygen species. This would allow an irradiated cell to protect sufficient enzymatic activity needed to repair DNA and survive.

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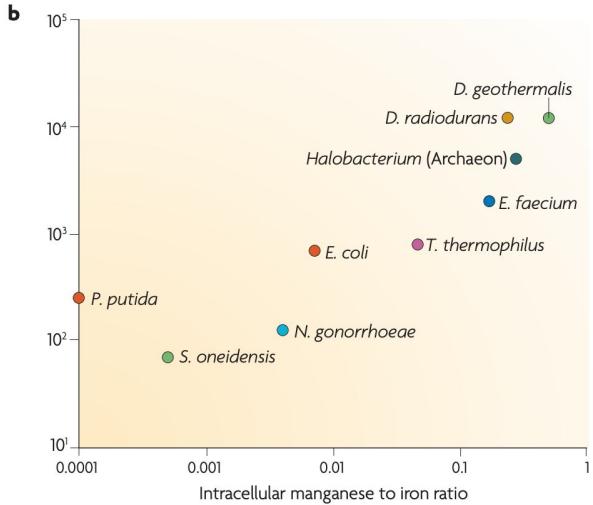
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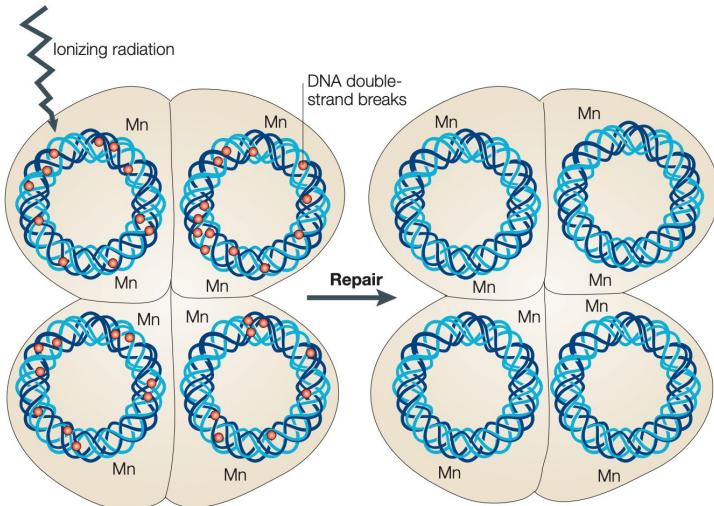
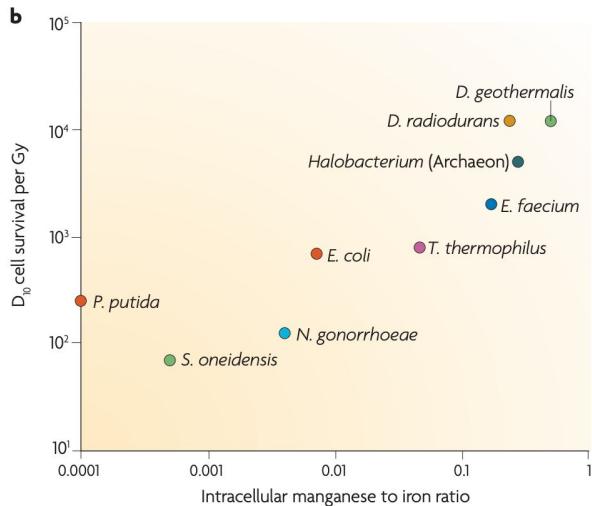
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## Abstract

In classical models of radiation toxicity, DNA is the molecule that is most affected by ionizing radiation (IR). However, recent data show that the amount of protein damage caused during irradiation of bacteria is better related to survival than to DNA damage. In this Opinion article, a new model is presented in which proteins are the most important target in the hierarchy of macromolecules affected by IR. A first line of defence against extremely radiation-resistant bacteria might be the accumulation of manganese complexes which can prevent the production of iron-dependent reactive oxygen species. This would allow an irradiated cell to protect sufficient enzymatic activity needed to repair DNA and survive.



# Where?

**Table 1** Radiation-resistant extremophiles isolated from different types of environments

Organism	Environment	Radiation	Reference
<i>Cellulosimicrobium cellulans</i> UVP1;	Elevated land	UVR-type C	Gabani et al. (2012a)
<i>Bacillus pumilus</i> UVP4			
<i>B. pumilus</i> SAFR-032	International Space Station	UVR	Vaishampayan et al. (2012)
<i>Bacillus subtilis</i>	Earth's magnetosphere	UVR	Nicholson et al. (2011)
<i>B. subtilis</i> HA101	Simulated Martian environment	UVR	Kerney and Schuerger (2011)
<i>Hymenobacter xinjiangensis</i>	Desert	UVR and gamma	Zhang et al. (2007a)
<i>Rubrobacter radiotolerance</i>	Unknown	Gamma	Terato et al. (2011)
<i>Sphingomonas</i> sp. RB2256	Unknown	UVR-type B	Joux et al. (1999)
<i>Chroococcidiopsis</i> sp.	Desert and hypersaline	X-ray	Billi et al. (2000)
<i>Halobacterium salinarum</i>	Unknown	X-ray	Robinson et al. (2011)
<i>Deinococcus radiodurans</i> R1	Canned meat	X-ray, UVR, and gamma	Shukla et al. (2007)
<i>Bacillus megaterium</i>	Lake	UVR	Zenoff et al. (2006)
<i>Staphylococcus saprophyticus</i>	Lake	UVR	Zenoff et al. (2006)
<i>Acinetobacter</i> sp. Ver3, Ver5, and Ver7	Lake	UVR-type B	Di Capua et al. (2011)
<i>Streptomyces radiopugnans</i>	Radiation-pollution	Gamma	Mao et al. (2007)

Table 2 | **Species of ionizing-radiation-resistant bacteria**

<b>Species</b>	<b>Representative D<sub>10</sub> value*</b>	<b>Phylum</b>	<b>Refs</b>
<i>Methylobacterium radiotolerans</i>	1,000 Gray	α-Proteobacteria	80,81
<i>Kocuria rosea</i>	2,000 Gray	Actinobacteria	4,8
<i>Acinetobacter radioresistens</i>	2,000 Gray	γ-Proteobacteria	82
<i>Kineococcus radiotolerans</i>	2,000 Gray	Actinobacteria	83
<i>Hymenobacter actinosclerus</i>	3,500 Gray	Flexibacter–Cytophaga–Bacteroides	84
<i>Chroococcidiopsis</i> spp.	4,000 Gray	Cyanobacteria	11
<i>Rubrobacter xylanophilus</i>	5,500 Gray	Actinobacteria	85
<i>Deinococcus radiodurans</i> R1	10,000 Gray	Deinococcus–Thermus	86
<i>Rubrobacter radiotolerans</i>	11,000 Gray	Actinobacteria	85

Cox, M. M., & Battista, J. R. (2005). *Deinococcus radiodurans—the consummate survivor*. *Nature Reviews Microbiology*, 3(11), 882-892.

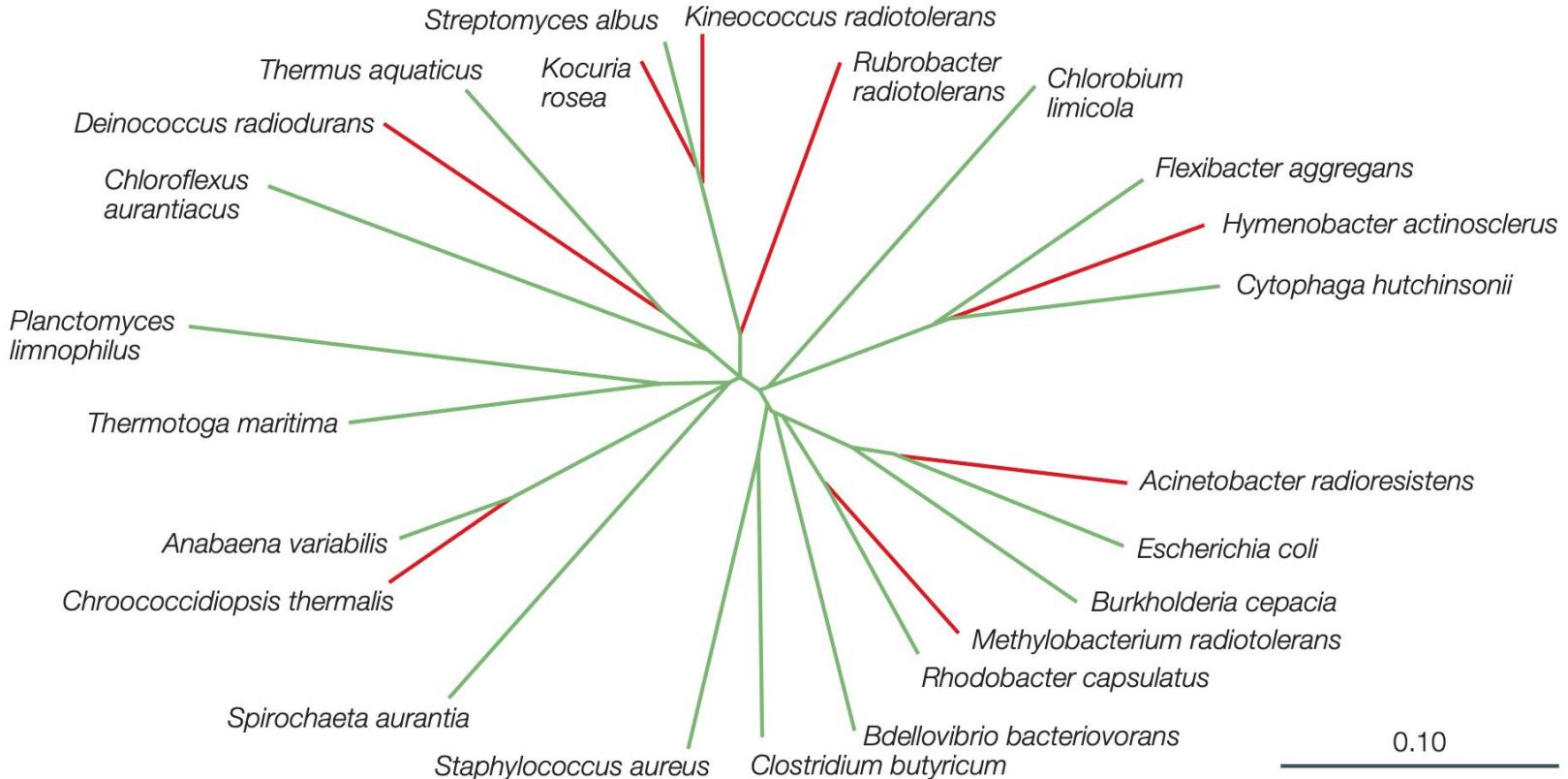


Figure 2 | **A 16S-rRNA-gene-sequence-based phylogeny of the main lineages of the domain Bacteria.** The branches in red are those in which ionizing-radiation-resistant taxa have been described. The scale bar represents 10 inferred nucleotide substitutions per 100 nucleotides.



# Two hypotheses



Radioresistance is a vestige of DNA-repair mechanisms that were present in ancestral species retained in those organisms that continue to require this phenotype



This phenotype has arisen in unrelated species through horizontal gene transfer, or possibly convergent evolution



# Applications

- Bioremediation of nuclear waste
- *D. radiodurans* has been proven effective in bioremediation of heavy metals from acidic and neutral water
- Several UVR-protective compounds have been isolated from UVR-resistant extremophiles



## Reads

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- Gabani, P., & Singh, O. V. (2013). Radiation-resistant extremophiles and their potential in biotechnology and therapeutics. *Applied microbiology and biotechnology*, 97(3), 993-1004.
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