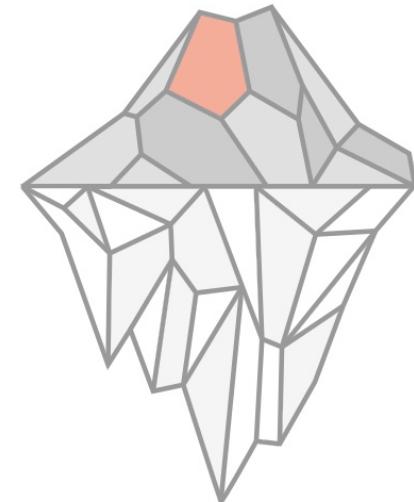


MICROBIOLOGY OF EXTREME ENVIRONMENTS



APPLICATIONS OF EXTREMOPHILES

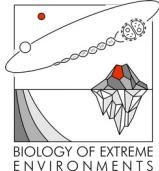
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BIOREMEDIATION



Bioremediation

Bioremediation is the use of biology to degrade or immobilize chemical compounds. It can be theoretically applied to the removal of any undesired compounds (organic or inorganic). Bioremediation has been used at a number of sites worldwide, including Europe, with varying degrees of success.

1972 - First commercial application: Sun Oil pipeline spill in Ambler, Pennsylvania

mid-1980s

Emphasis on bioengineering organisms for bioremediation. This technology did not live up to its initial promise

1990s

Emphasis switched to greater reliance on natural microorganisms and techniques to enhance their performance



Bioremediation: applications

Types of impact where bioremediation can be applied:

Petroleum hydrocarbons

Chlorinated solvents

Polynuclear aromatic hydrocarbons

Pesticides

TNT and a variety of other organic chemicals

Inorganic nitrogen (nitrate, ammonia)

Excessive nutrients and OM

Heavy metals (Pu, Np, Cr, As)

Radioactive metals (U, Th)







POP's

(Persistent Organic Pollutants)
from Industry, Agriculture, Vessels, etc.

Flame retardants, fertilizers, industrial chemicals & other pollutants such as:

- PCBs
- PCDDs (Polychlorinated dibenzo-p-dioxins)
- PCDFs (Polychlorinated dibenzofurans)
- OCs (Organochlorine pesticides)
- PAHs (Polycyclic aromatic hydrocarbons)
- PBDEs (Polybrominated diphenyl ethers)
- APs (Alkylphenol)
- APnEOs (Alkylphenol ethoxylates)
- CPs (Chlorinated paraffins)
- PCNs (Polychlorinated naphthalenes)
- FOCs (Fluorinated organic compounds)

- Bitumen, Condensate, petroleum products
- Pesticides: from forestry, agriculture, industry
- Metals
- PPCPs (Pharmaceuticals & personal care products)
- Biological contaminants

Bioaccumulation from prey

TOXINS



Toxin effects of increased mortality of prey

TOXINS



Whales must eat 50% more fish to survive.

Environmental toxins result in poor fish reproduction (fewer fish hatch), lower lipids (fats) in each fish (each fish is less healthy), and greater fish mortality (more fish die). Whales need to eat up to 50% more fish to satisfy their nutritional requirements, and the fish they are eating are more toxic, resulting in a higher toxin load in whales.

Direct exposure to air and sea borne toxins via contact, breathing, and injection.

TOXINS

Exhaust fumes from vessels - including whale watching boats.

- CO_x (Carbon oxides)
- SO_x (Sulfur oxides)
- NO_x (Nitrogen oxides)
- PM (Particulate Matter)

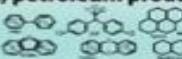
• HC (Hydrocarbons:
unburned fuel; aromatics,
alkanes, and alkenes; partially
oxidized phenols and carbonyls)

Whales do not have an olfactory system or facial sinuses. They cannot detect or avoid engine exhaust, oil slicks or other toxins. Whales cannot prevent water-soluble toxins from entering their lungs.

Oil and chemical slicks on water surface

Bitumen
Tar balls,
and other
chemical
emulsions

Direct exposure to and ingestion of
chemicals, petroleum products



Adverse effects of toxins in marine mammals include: increased inflammation, toxicity; reduced reproductive, immune, and behavioural health, and increased mortality.



Bioremediation: ups

1. Bioremediation can be “natural process” and is therefore perceived by the public as an acceptable waste treatment process for contaminated material. Microbes able to degrade the contaminant increase in numbers when the contaminant is present.
2. Theoretically, bioremediation is useful for the complete destruction of a wide variety of contaminants. The complete destruction of target pollutants is theoretically possible.
3. Bioremediation can often be carried out on site, often without causing a major disruption of normal activities. This also eliminates the need to transport quantities of waste off site and the potential threats to human health and the environment that can arise during transportation.
4. Bioremediation can prove less expensive than other technologies that are used for cleanup of hazardous waste.



Bioremediation: downs

1. Bioremediation is generally very costly, labor intensive, and time consuming considering the extent of the impact considered.
2. Bioremediation is limited to those compounds that are biodegradable. Not all compounds are susceptible to rapid and complete degradation.
3. There are some concerns that the byproducts of biodegradation may be more persistent or toxic than the parent compound.
4. Biological processes are often highly specific.
5. It is difficult to extrapolate from bench and pilot-scale studies to full-scale field operations.
6. Bioremediation often takes longer than other treatment options, such as excavation and removal of soil or incineration.
7. Regulatory uncertainty remains regarding acceptable performance criteria for bioremediation.



Bioremediation: *in situ* and *ex situ*

In situ bioremediation means there is no need to excavate or remove soils or water in order to accomplish remediation. *In situ* biodegradation can involve the addition of microorganisms, oxygen and nutrients. In its intrinsic version is the most widely applied type of bioremediation, although has several drawbacks.

Ex situ bioremediation requires excavation or removal of contaminated soil, sediments or waters (permanent or temporary) to facilitate microbial degradation. This technique is more expensive but it is able to overcome some of the disadvantage of *in situ* approaches.



Bioremediation: intrinsic vs engineered

Intrinsic bioremediation

This approach deals with stimulation of indigenous or naturally occurring microbial populations by feeding them nutrients and oxygen to increase their metabolic activity (Biostimulation). Chemotaxis is important to the study of *in situ* bioremediation because microbial organisms with chemotactic abilities can move into an area containing contaminants.

Engineered bioremediation

Involves the introduction of external (exogenous) microorganisms to enhance degradation (Bioaugmentation).

Basic Bioremediation Approaches

Natural attenuation – naturally occurring degradation of the pollutants by naturally occurring microbial populations **not supported** by human intervention

Biostimulation – Naturally occurring degradation of the pollutants by naturally occurring microbial population **supported by the additions** of nutrient, stimulants and surfactants

Bioaugmentation – Introduction of allochthonous engineered populations of microorganisms to enhance the degradation of the pollutants

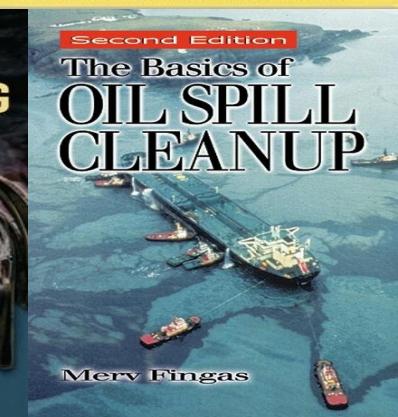
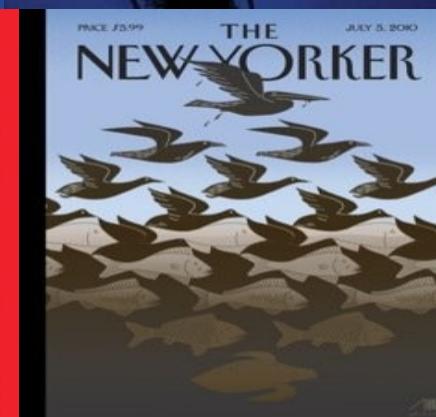
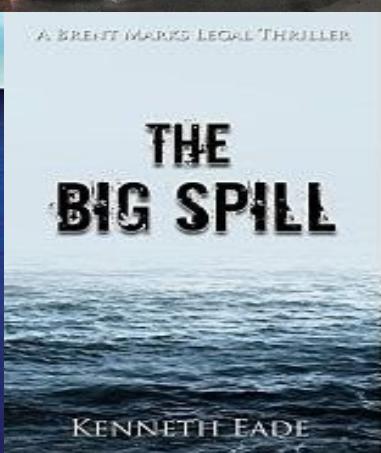
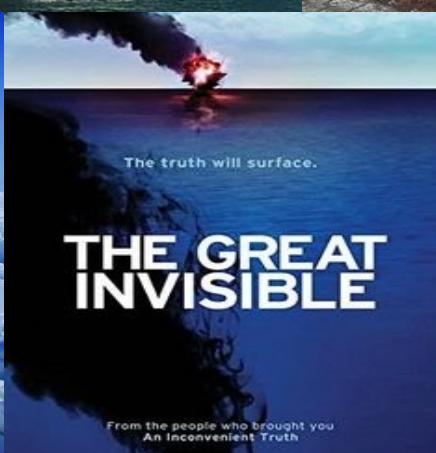
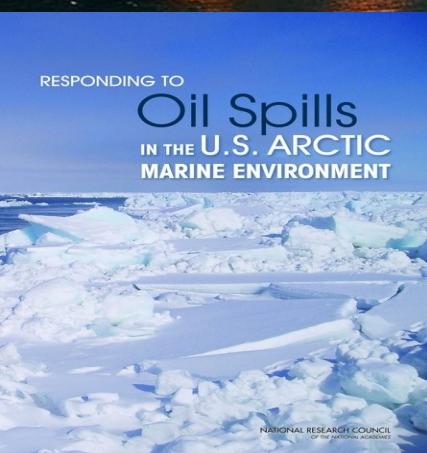


Microbial groups involved in degradation

Aerobic: Examples of aerobic bacteria recognized for their degradative abilities are *Pseudomonas*, *Alcaligenes*, *Sphingomonas*, *Rhodococcus*, and *Mycobacterium*. These microbes have often been reported to degrade pesticides and hydrocarbons, both alkanes and polyaromatic compounds. Many of these bacteria use the contaminant as the sole source of carbon and energy.

Anaerobic. Anaerobic bacteria are not as frequently used as aerobic bacteria. There is an increasing interest in anaerobic bacteria used for bioremediation of polychlorinated biphenyls (PCBs) in river sediments, dechlorination of the solvent trichloroethylene (TCE) and chloroform.

Methylotrophs. Aerobic bacteria that grow utilizing methane for carbon and energy. The initial enzyme in the pathway for aerobic degradation, methane monooxygenase, has a broad substrate range and is active against a wide range of compounds, including the chlorinated aliphatic trichloroethylene and 1, 2 dichloroethane.



The cost of cleaning

Different costs need to be take into account while considering bioremediation approaches vs natural attenuation

Economic cost of bioremediation effort

Economic cost of the damage caused by the no-intervention alternative

Economic cost of the post bioremediation dismantling

Ecological cost of bioremediation effort

Ecological cost of the post bioremediation dismantling

Economic and ecological cost of long term effects of the bioremediation effort

Table 2 Estimated costs of remediation strategies for hydrocarbon-polluted soils and sediments

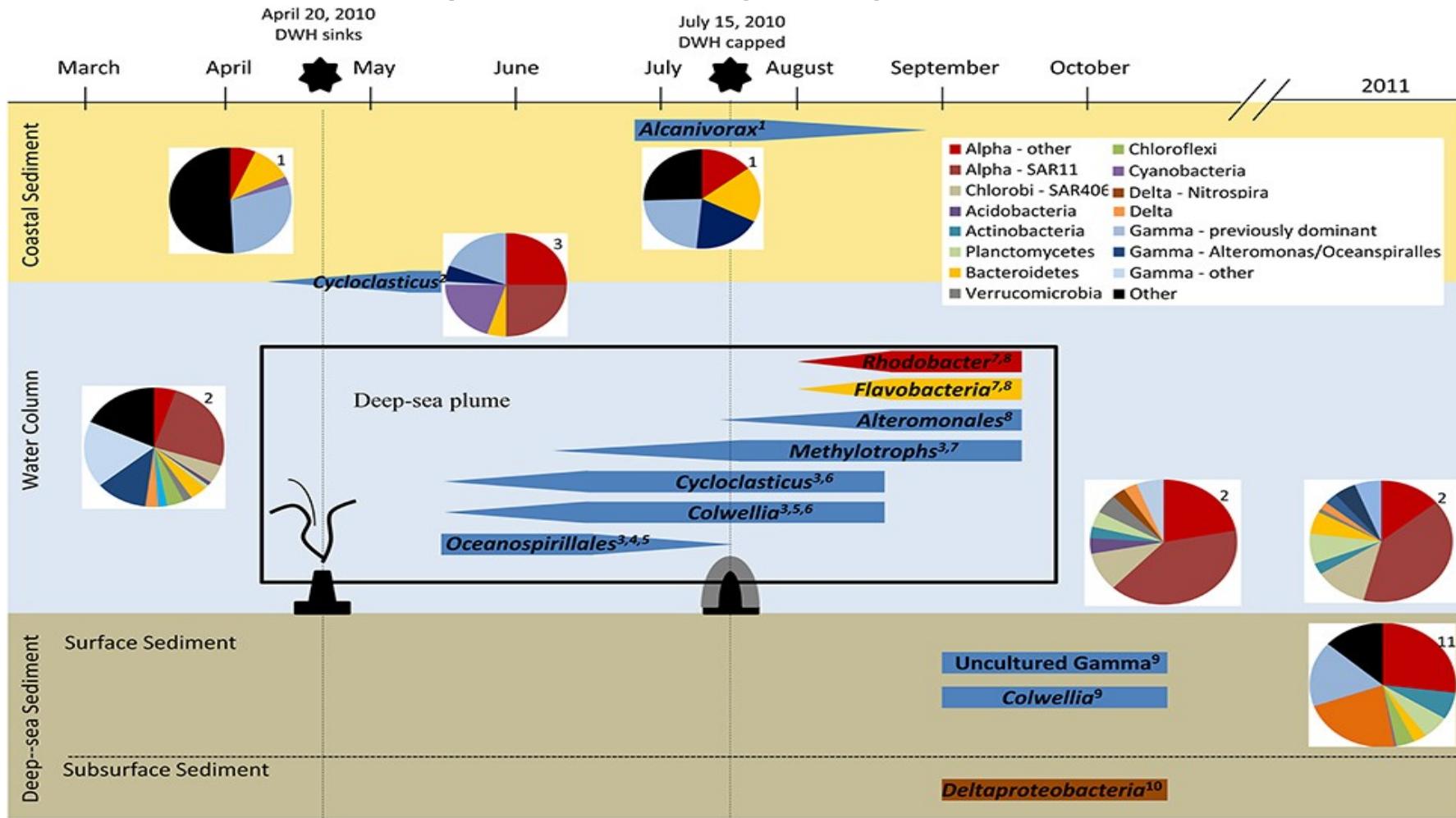
Remediation strategy	Treating site	Cost (US \$/m ³) ^a
Biological		
Biostimulation	In situ	30–100
Bioaugmentation	In situ	30–100
Bioventing	In situ	79–970
Biopiles	Ex situ	130–260
Composting	Ex situ	630–757
Landfarming	Ex situ	30–70
Physicochemical		
Vapor extraction	In situ	405–1,485
Thermal desorption	Ex situ	44–252

^a Data from the US Federal Remediation Technologies Roundtable (2014).



BP DWH Spill – 61 Billion USD

Naturally occurring degraders





Obligate Hydrocarbonoclastic Bacteria

The ecology of a special group of **ubiquitous marine hydrocarbon-degrading bacteria** offers a model of microbial dynamics after oils spills.

Obligate hydrocarbonoclastic bacteria (OHCB) are an unusual ecological group that plays a key role in biodegradation of petroleum hydrocarbons in marine environments. OHCB are able to grow on a reduced spectrum of hydrocarbons.

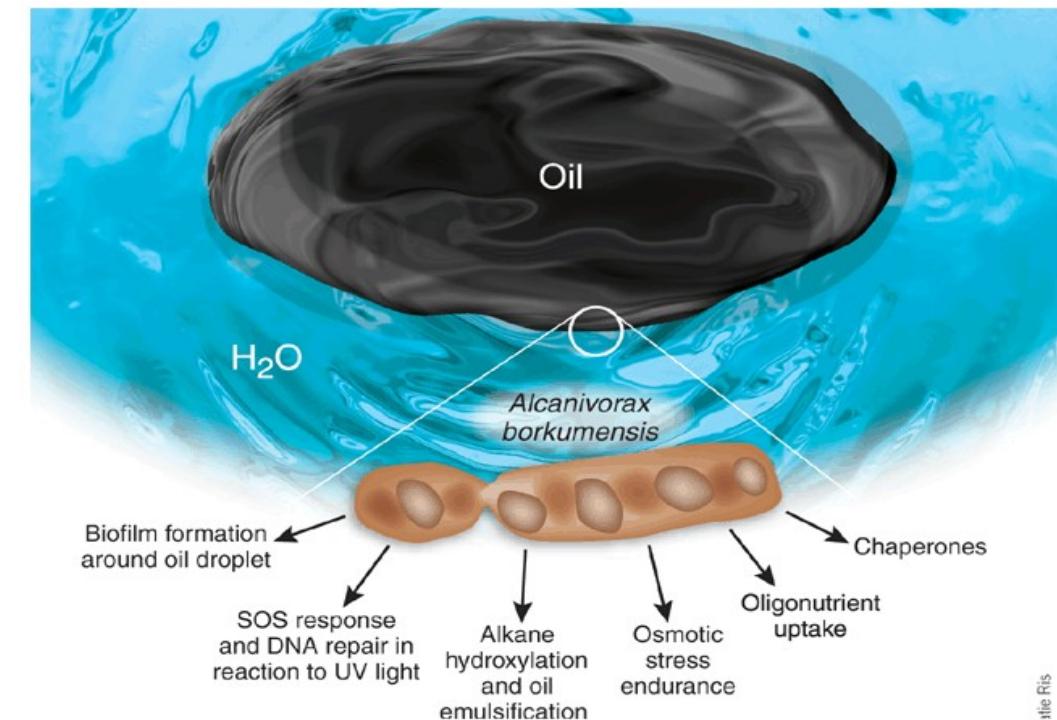
In non-polluted marine environments, OHCB are found in **extremely low numbers**, but after an oil pollution event, members of this group undergo a **bloom**. OHCB growth is favored in hydrocarbon-polluted seawater because they can breakdown substrates that are useless for most bacteria.

The model OHCB is ***Alcanivorax borkumensis*** SK2, able to degrade n-alkanes of chain-length up to C32, long-chain isoprenoids, phytane, pristane, and alkyl-aromatic hydrocarbons and to synthesize **biosurfactants** and **exopolysaccharides** probably involved in **biofilm formation**.

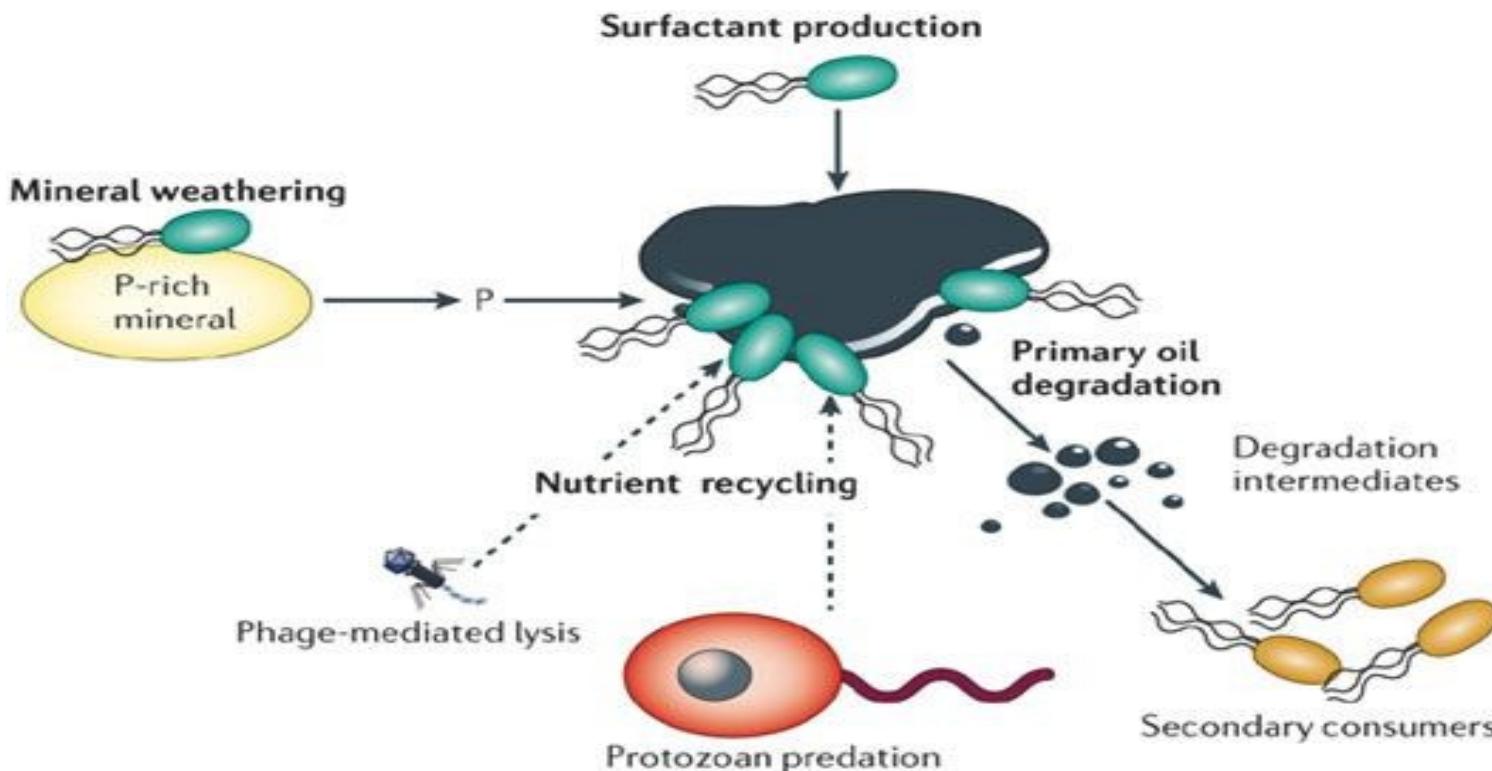
Alcanivorax borkumensis

A. *borkumensis* genome revealed **multiple systems for hydrocarbon degradation pathways**, i.e., two alkane hydroxylase systems, AlkB1 and AlkB2, and three P450 cytochromes.

The **absence** of genes belonging to **glucose breakdown** pathways highlights its amazing metabolic specialization.



Community Response



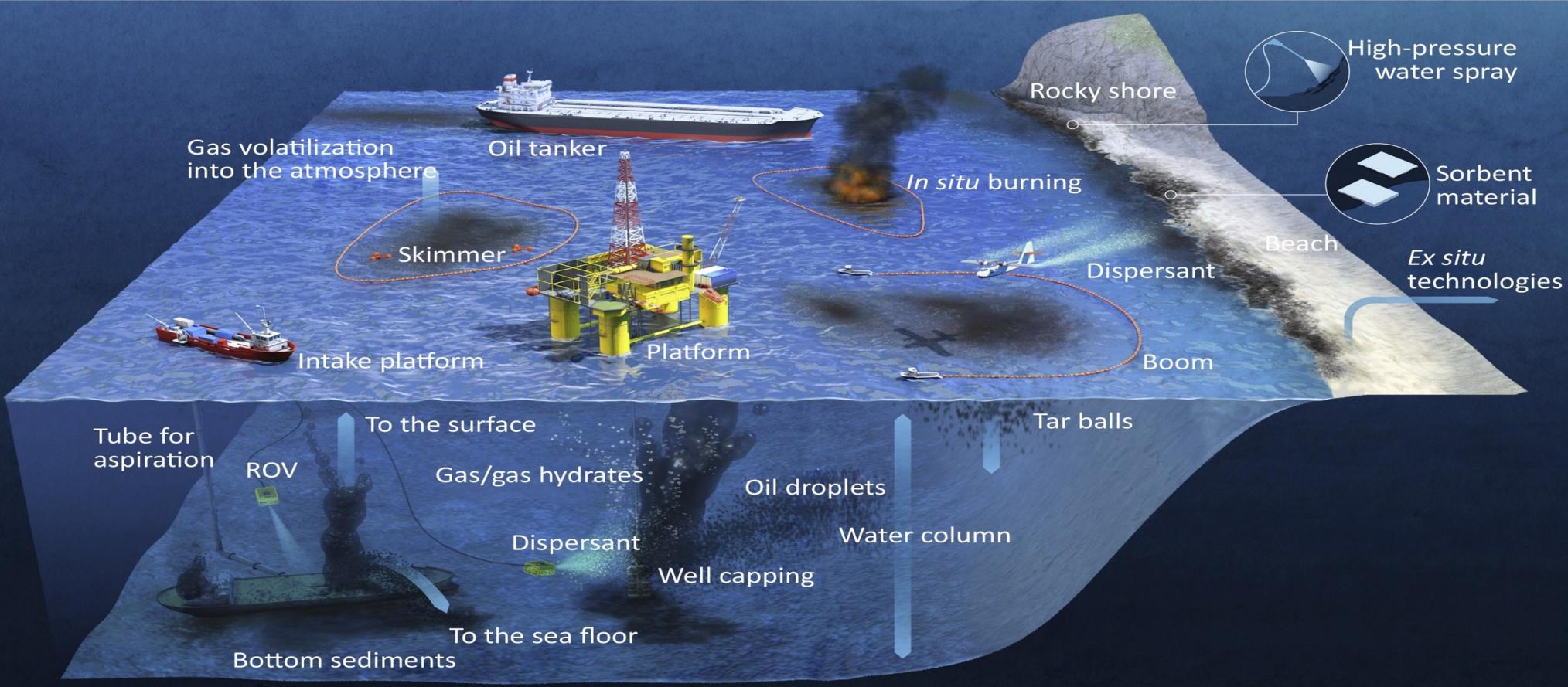
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Nature Reviews | Microbiology

Head et al., 2006 NatRevMicrobiol

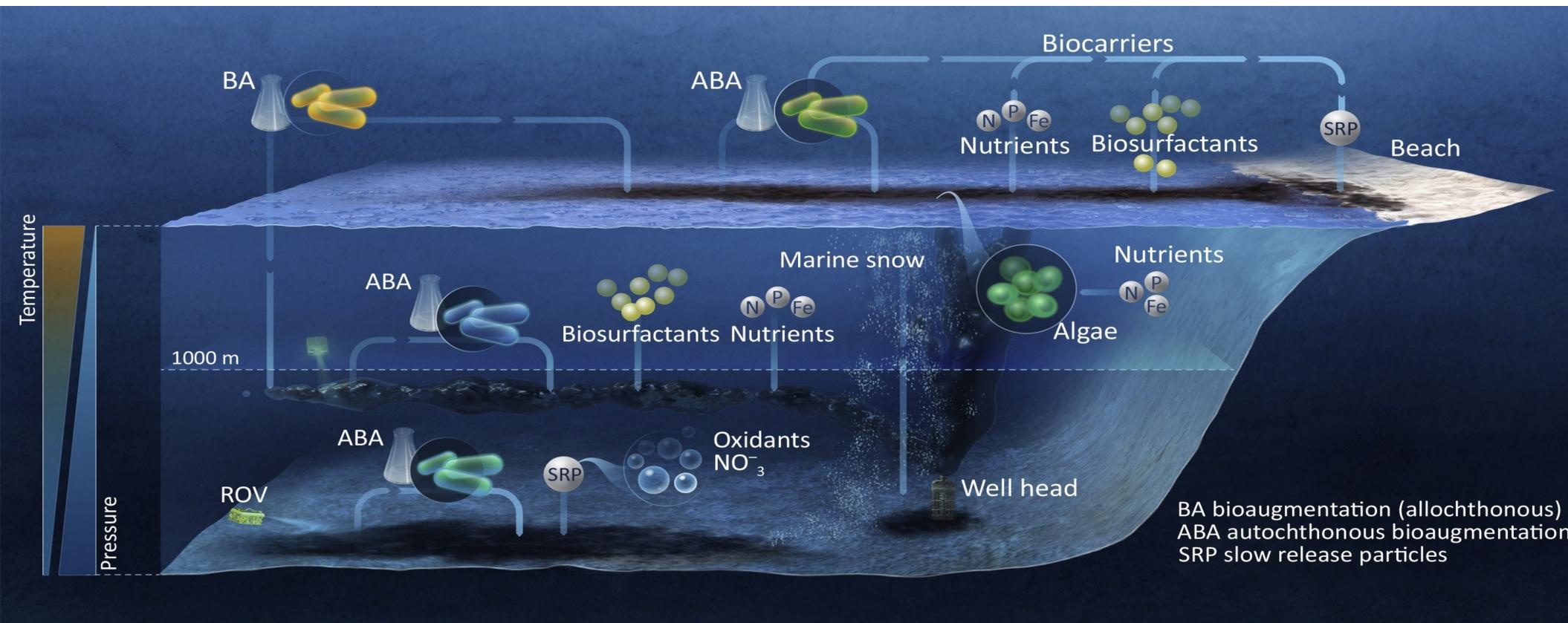
Enriching/Isolating oil degrading microbes



Oil spill response

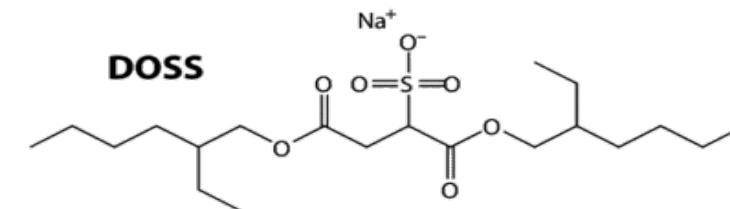
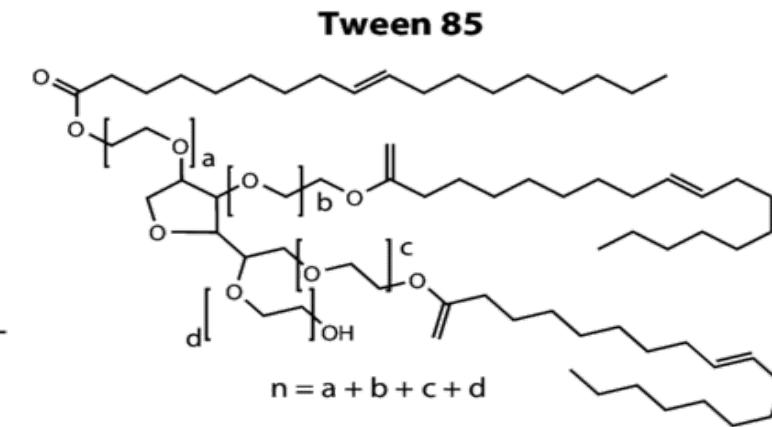
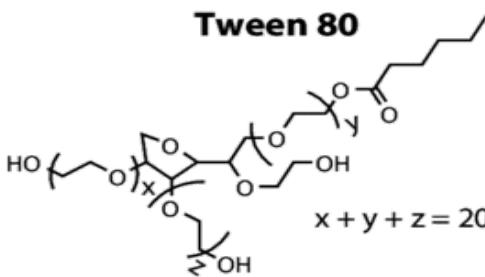
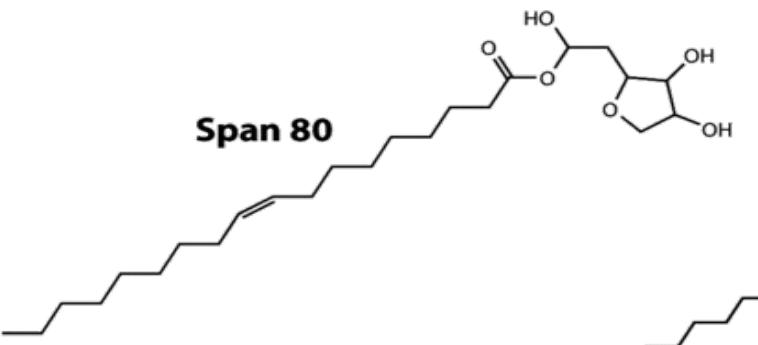


Hydrocarbon decontamination during spills



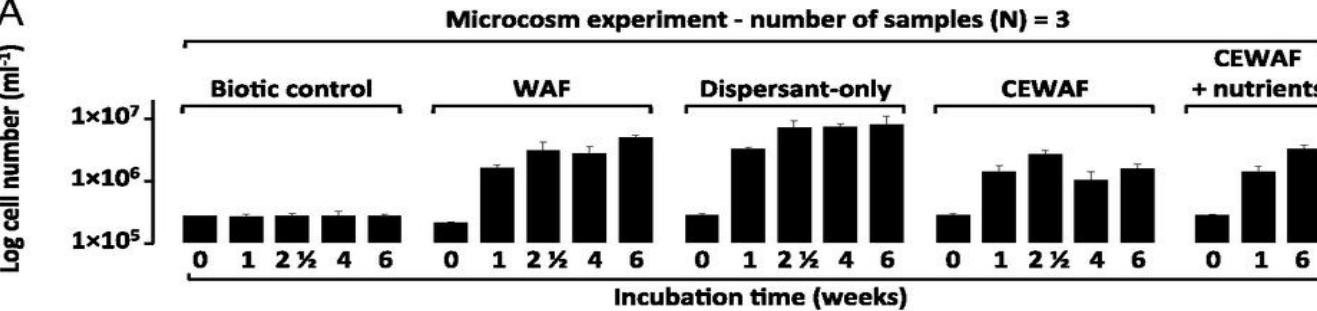


Unexpected effects of dispersant



Unexpected effects of dispersant

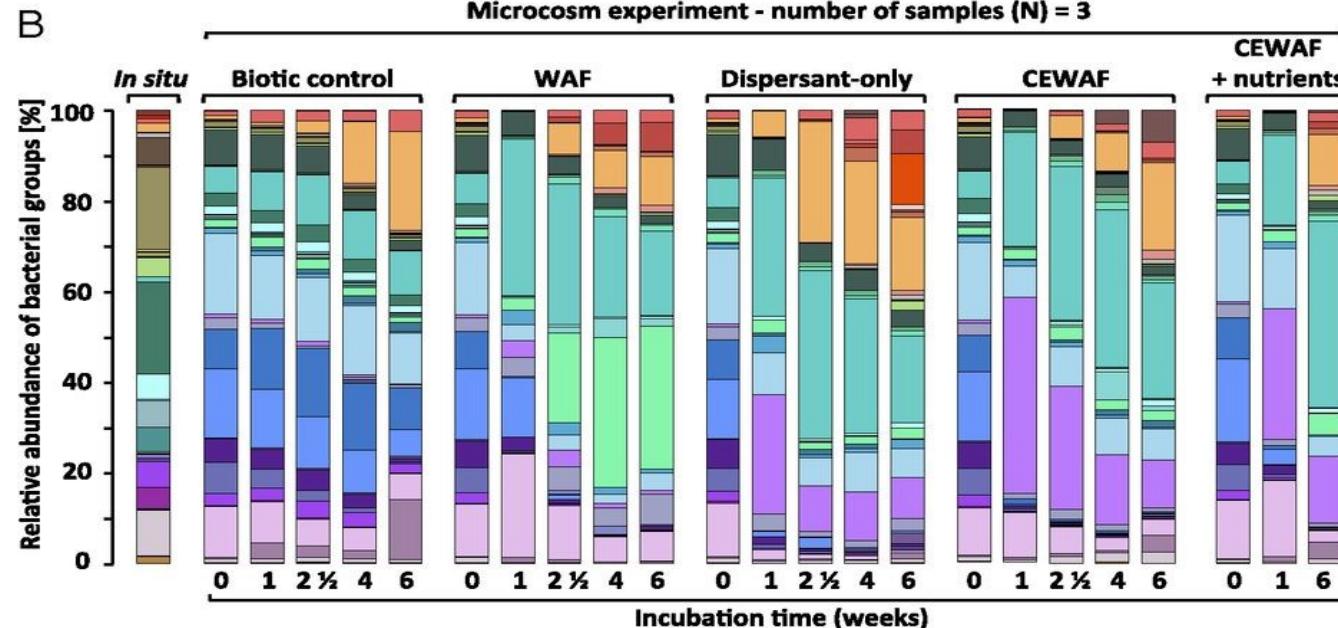
A



Taxonomic identification

Flavobacteria	Acidobacteria
	Actinobacteria
α	Crocinitomix
	Owenweeksia
δ	Bizonia
	Flavobacteriaceae
ε	Psychroserpens
	Kordia
β	Ulvibacter
	Balneola
Bacteria	Other Bacteriodetes
	Chloroflexi / SAR202
γ	Deferrabacteres / SAR406
	Phyllobacteriaceae
Proteobacteria	Rhodobacteraceae
	SAR11
α	Other Alphaproteobacteria
	Bdellovibrionaceae
δ	Marinobacter
	SAR92
ε	Teredinibacter
	Other Alteromonadaceae
β	Uncult. Alteromonadaceae
	Colwellia
γ	Alkanivorax
	Amphritea
Proteobacteria	Balneatrix
	Oceaniserpentilla
α	Oceanospirillaceae
	Oleiphilus
δ	Oceanospirillales
	Cycloclasticus
ε	Methylophaga
	Other Gammaproteobacteria
β	Verrucomicrobia

B





BIO MASS CONVERSION

Production from Biomass

Biomass are defined as **organic products from biological material** such as plants, animals, microorganisms and municipal waste. They can be directly the bodies of the organisms or byproducts of other human-mediated processes.

Biomass can be classified in many different types, depending from the provenience, the composition, the state (liquid, solid) and the lability.



Examples of Biomass



Plant oils or sugary
/starchy parts of crops



Trees, wood chippings,
or straw



Waste cooking oil

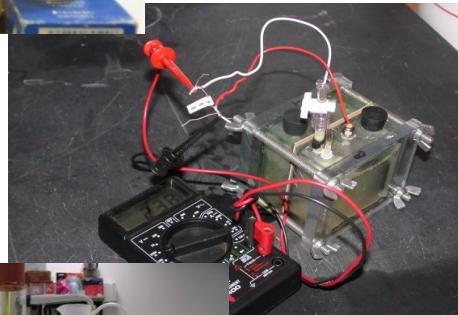


Algae oil

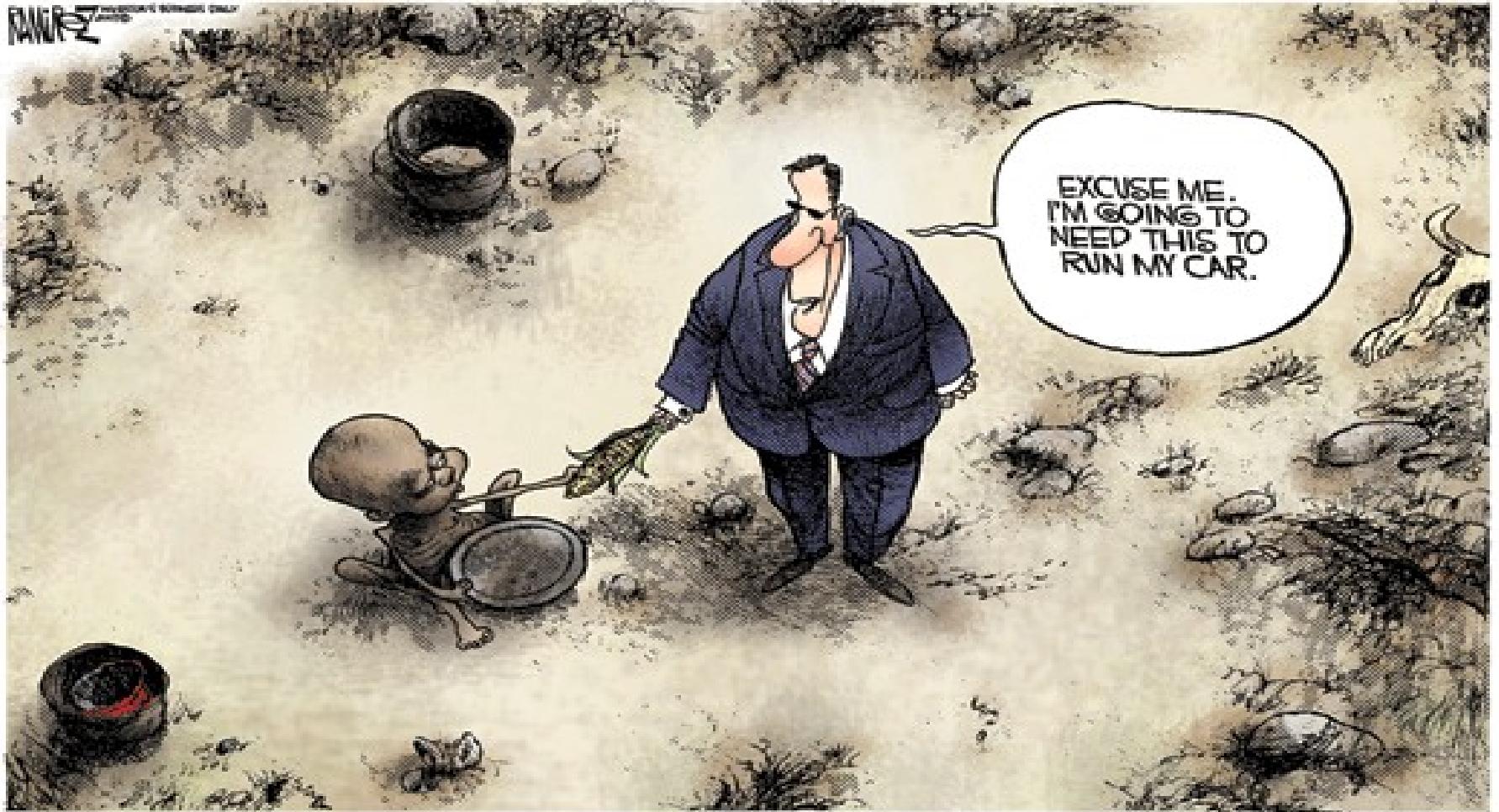


Animal waste

Methods of Biomass to Energy Conversion



- **Thermal conversion**
 - Combustion
 - Gasification
 - Pyrolysis
- **Chemical conversion**
 - Transesterification (biodiesel)
- **Biological conversion**
 - Fermentation (methanol, ethanol, butanol)
 - Anaerobic digestion (methane)
 - Anaerobic respiration (bio-battery)



Biogas

Anaerobic digestion is a commercially proven technology and is widely used for recycling and treating wet organic waste and waste waters. It is a type of fermentation that converts **organic material** into **biogas**, which mainly consists of **methane** (approximately 60%) and carbon dioxide (approximately 40%) and is comparable to landfill gas.

Virtually **any biomass except lignin** (a major component of wood) can be converted to biogas—including animal and human wastes, sewage sludge, crop residues, industrial processing byproducts, and landfill material.

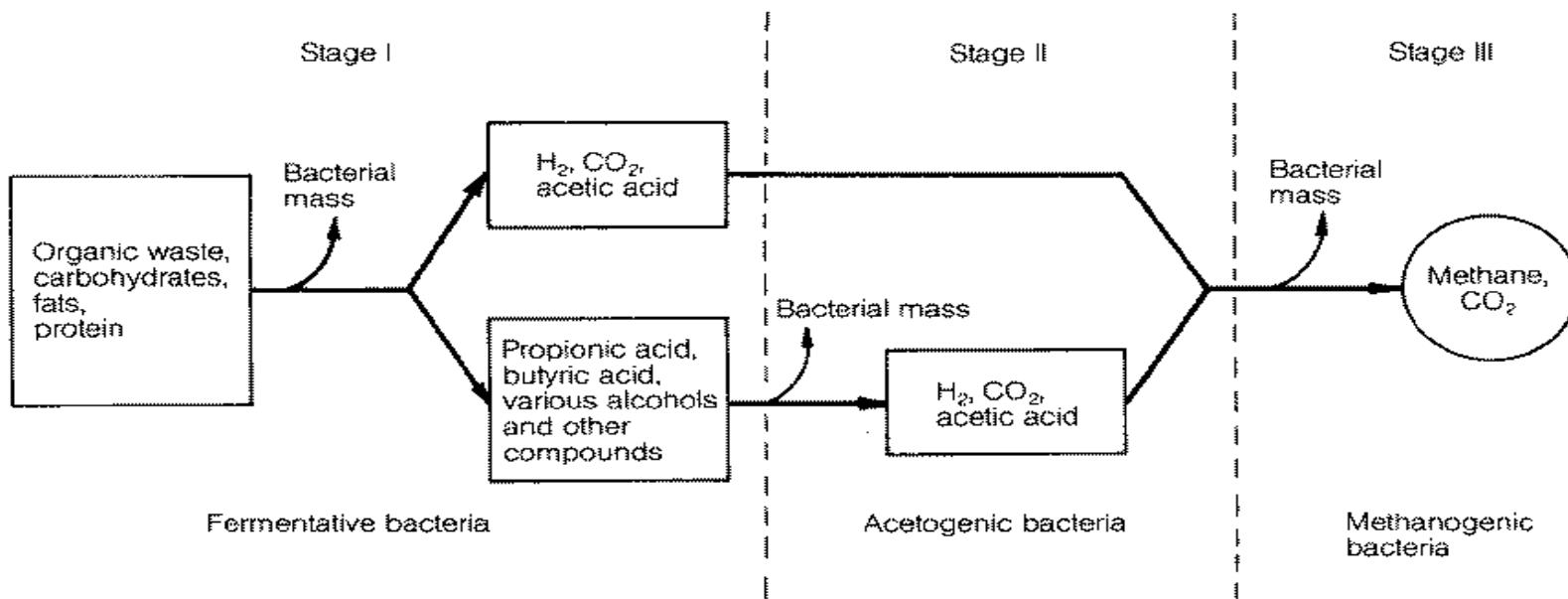
The conversion of animal wastes and manure to methane/biogas can yield significant health and environmental benefits.

Biogas Production



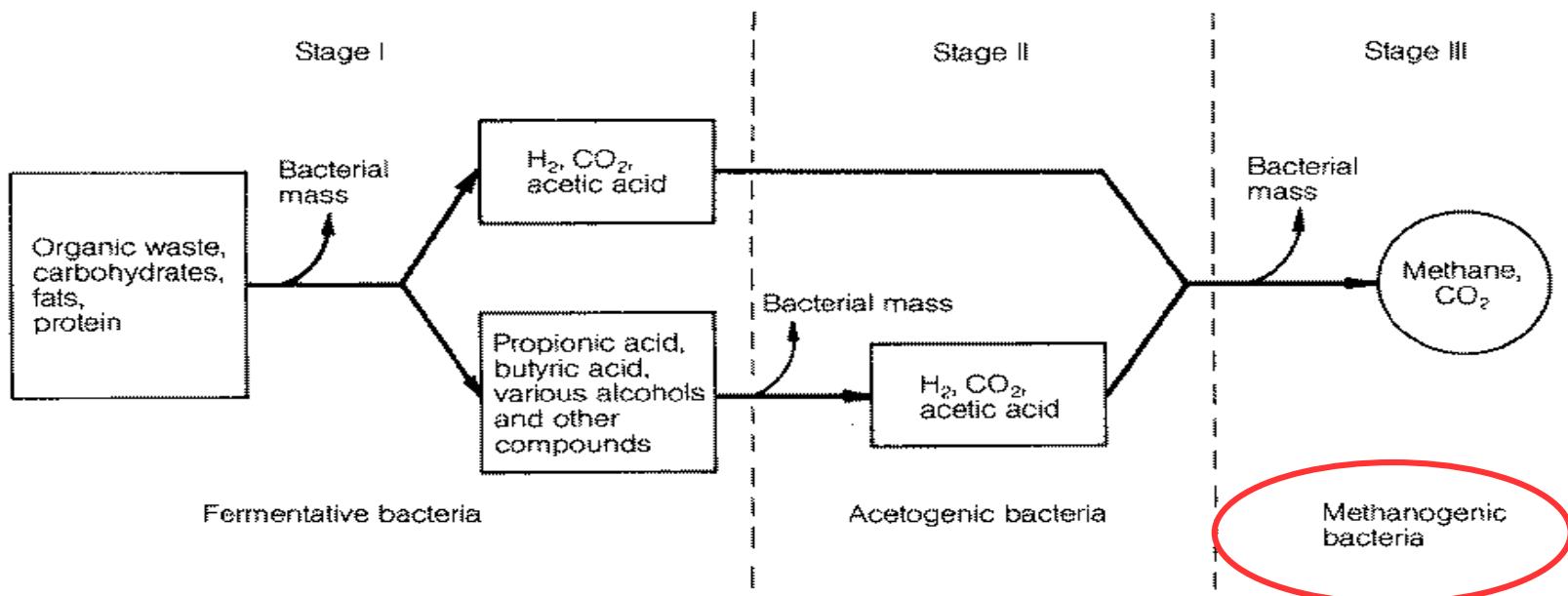
Biomass is fermented in a multistep process to produce secondary wastes (together with bio-fertilizers and other compounds) and methane (together with other compounds that need to be removed)

Biogas Microbiology



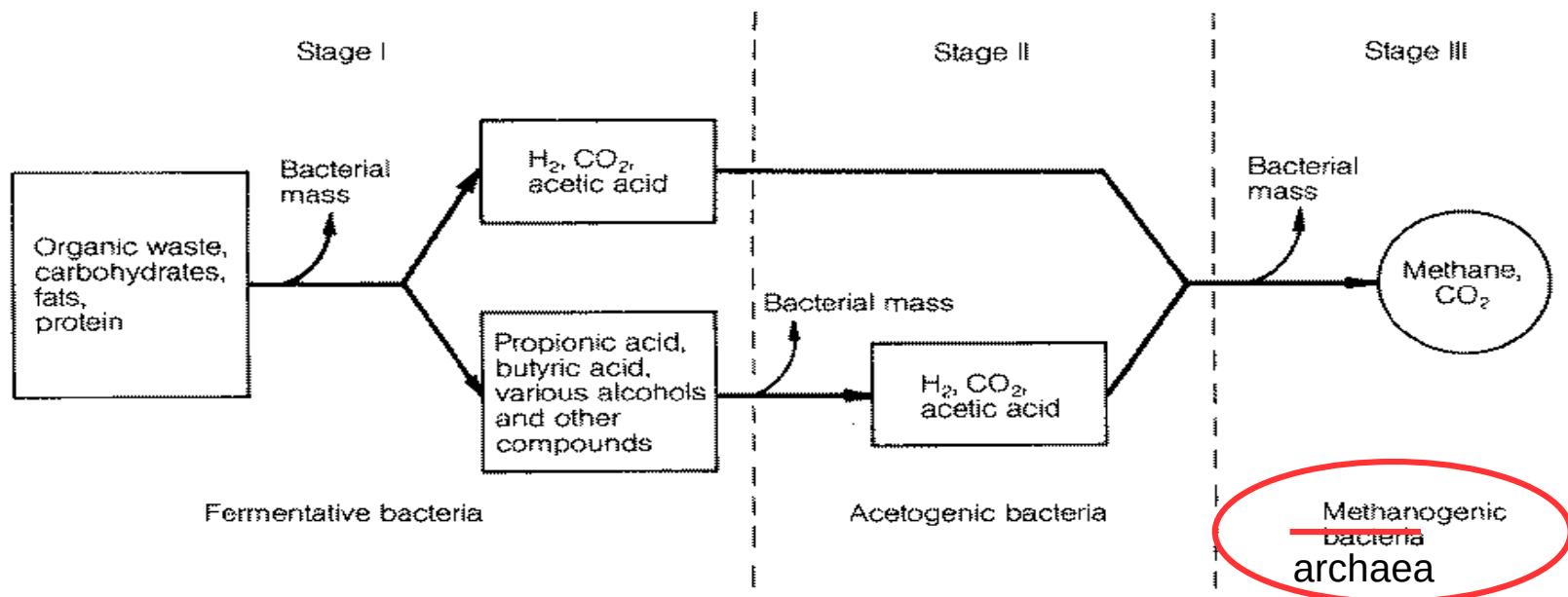
A complex consortia of microorganisms is able to convert waste into Biogas (mainly CH_4) for use in cogenerators and as energy source.

Biogas Microbiology



A complex consortia of microorganisms is able to convert waste into Biogas (mainly CH_4) for use in cogenerators and as energy source.

Biogas Microbiology



A complex consortia of microorganisms is able to convert waste into Biogas (mainly CH₄) for use in cogenerators and as energy source.



Biorefineries

An emerging concept to be aware of is **biorefineries**. A biorefinery involves the **co-production** of a spectrum of **bio-based products** (food, feed, materials, chemicals) **and energy** (fuels, power, heat) from biomass.

A biorefinery is a facility that integrates biomass conversion processes and equipment to produce fuels, power, and value-added chemicals from biomass. The biorefinery concept is **analogous to today's petroleum refinery**, which produces multiple fuels and products from petroleum.

A biorefinery could, for example, produce one or several low-volume, but high-value, chemical products and a low-value, but high-volume liquid transportation fuel such as biodiesel or bioethanol. At the same time, it can generate electricity and process heat, for its own use and perhaps enough for sale of electricity to the local utility.

Biobatteries

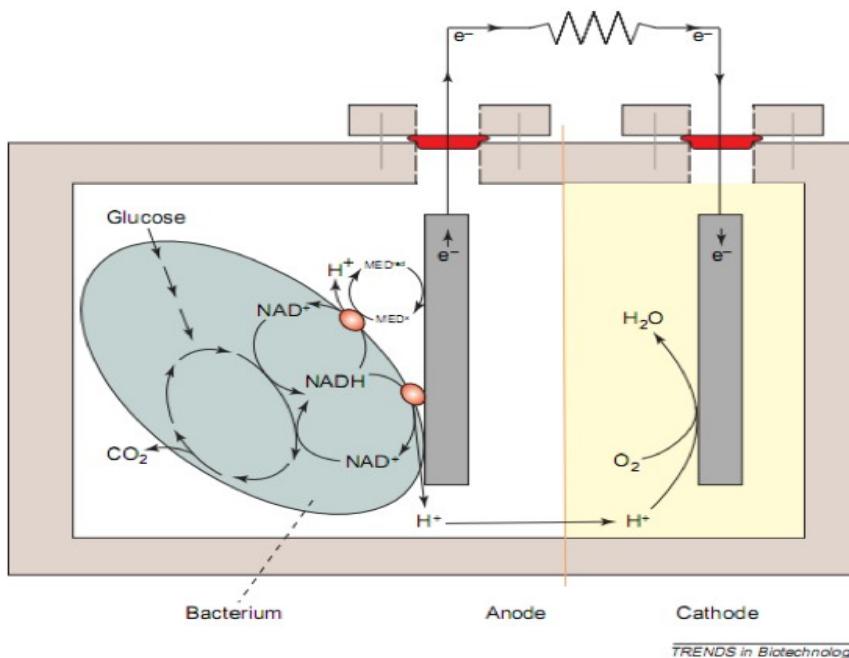
An emerging technology is the direct use of **microbial metabolism** to produce **electrical currents**, called Biobatteries.

This technology takes advantage of the ability of certain microbes to use a solid metal substrate as **electron acceptor of their respiratory chain**. This effectively couples the degradation of an organic substrate to the capacity of producing an electrical current.

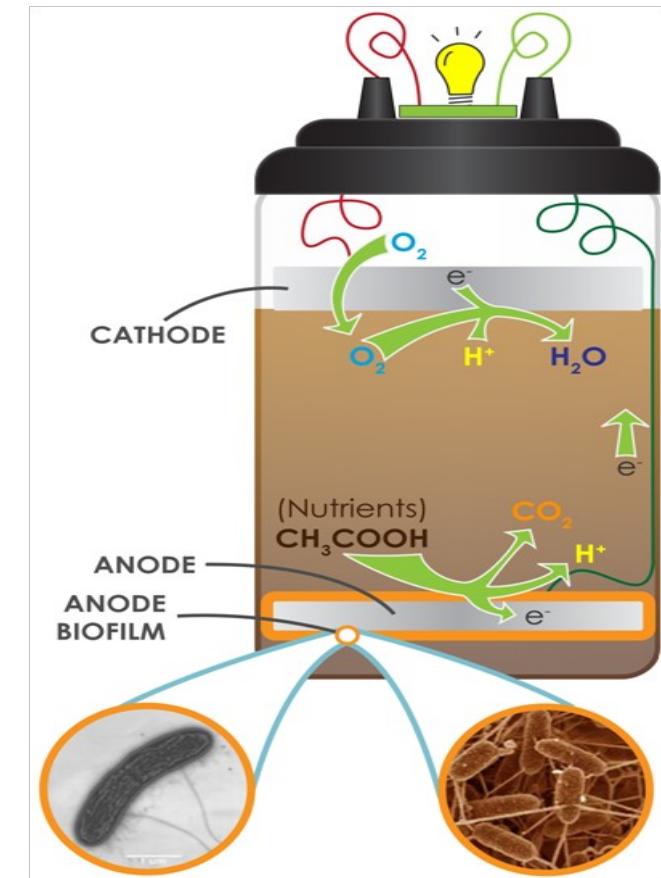
The technology is under development, however has the potential to be transformative, especially if coupled to waste biomass. Current biobatteries use sugars (glucose) as main source of energy.

Several prototypes and patents have already proven their ability to power small scale everyday electronic devices.

Biobatteries



Electrons flow from an anode through a resistor to a cathode where electron acceptors are reduced. Protons flow across a **proton exchange** membrane (PEM) to complete the circuit.





Back to the Future, 1985

Bioplastics: mainly PHA and PHB

Bioplastics are plastics derived from **renewable biomass** sources, such as vegetable fats and oils, corn starch, straw, woodchips, food waste, etc.

Many different types exist: **cellulose exters**, **celluloid**, **nitrocellulose**, **polylactic acid** (from corn mainly), **PA11** (from vegetable oils), **Polyethylene from bioethanol**, **Polyurethanes** and **epoxies** (from microalgae oils).

Current focus is on a **biobased AND biodegradable** type of biopolymer:

Polyhydroxyalkanoates (PHA) are a class of linear polyesters produced in nature by bacterial fermentation of sugar or lipids. They have characteristics similar to those of the petroplastic polypropylene.

Case Study: Bio-On Spa

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Case Study: Bio-On Spa

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il Resto del Carlino BOLOGNA

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Pubblicato il 14 gennaio 2020

Bio on, parla il fondatore Astorri. "Una rivoluzione, perciò ci hanno fermato"

Intervista esclusiva all'ex presidente (indagato) dopo il fallimento: "Il potenziale tecnologico resta enorme. Vittime di un attacco speculativo ben preparato"

di NICOLA BIANCHI E RICCARDO RIMONDI

Case Study: Bio-On Spa





CLIMATE ENGINEERING

SMOKING DOES NOT
CAUSE CANCER.

OKAY, IT DOES, BUT I'M
ADDICTED.

I'M ADDICTED, BUT I KNOW
I'M GOING TO HAVE TO QUIT.

I HAVEN'T QUIT YET, BUT
I'M CUTTING BACK.

I DIDN'T CUT BACK
VERY MUCH, SO NOW
I'M GOING TO QUIT.

I SHOULD HAVE QUIT
WHEN I SAID I WOULD,
BUT NOW IT'S TOO LATE.

UNIVERSAL PRESS SYNDICATE
2006 THE WASHINGTON POST

4-4-06

FOSIL FUELS DO NOT
CAUSE GLOBAL WARMING.

OKAY, THEY DO, BUT
I'M ADDICTED...

LIVE AND LEARN, OR VICE VERSA -

Climate Engineering: Geoengineering

Refers to actions **deliberately taken** in order to alter the natural functioning of processes that can **directly or indirectly** help with the **mitigation of the climate change crisis**, global warming and the increase of CO₂ in the atmosphere

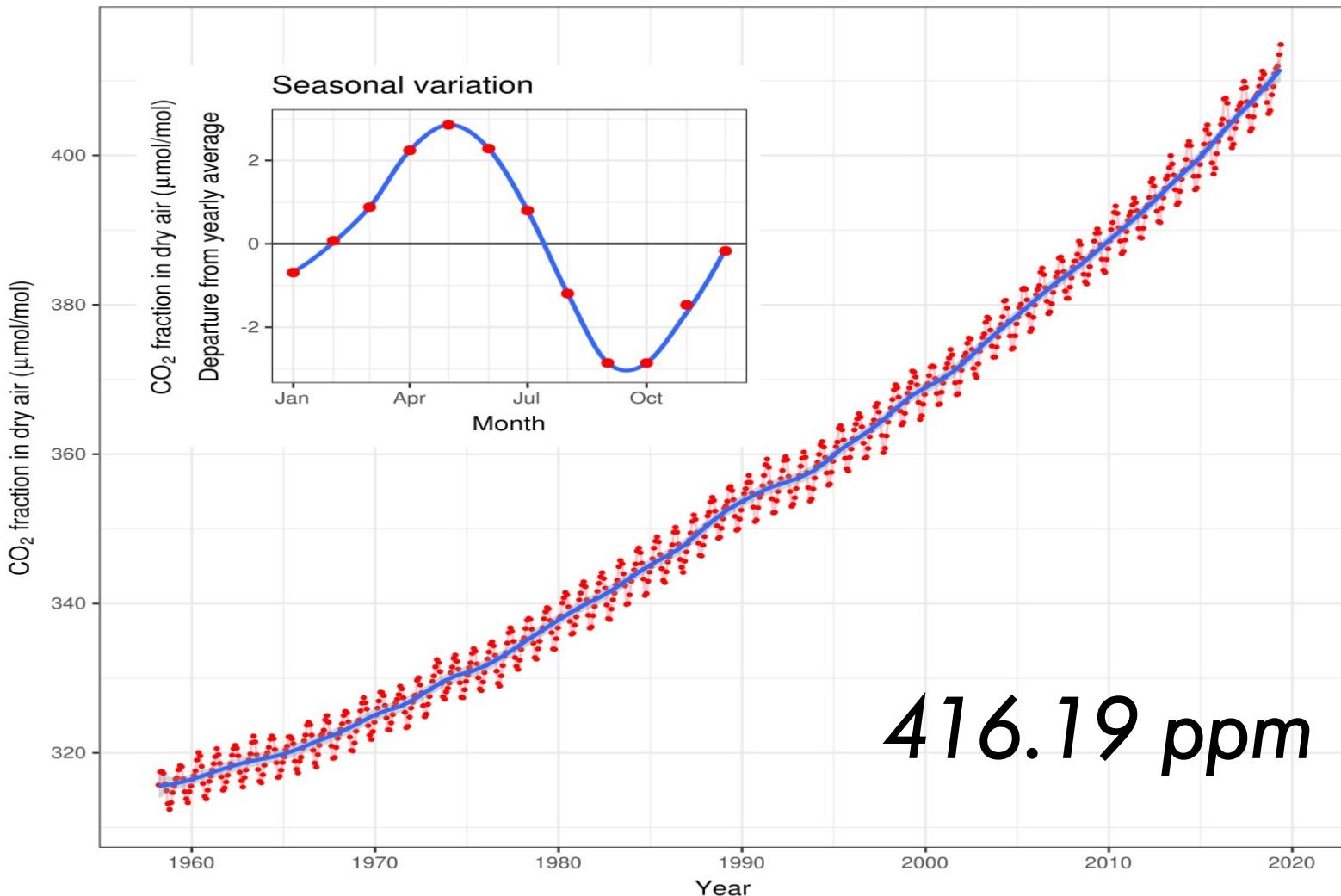
Geoengineering is the deliberate modification of Earth's environment on a large scale "**to suit human needs and promote habitability**"

Over the years it has focused on different aspect of the issue, tacking the problem with different technological approach

These can be broadly divided in **Solar Radiation Management** and **Carbon Sequestration**

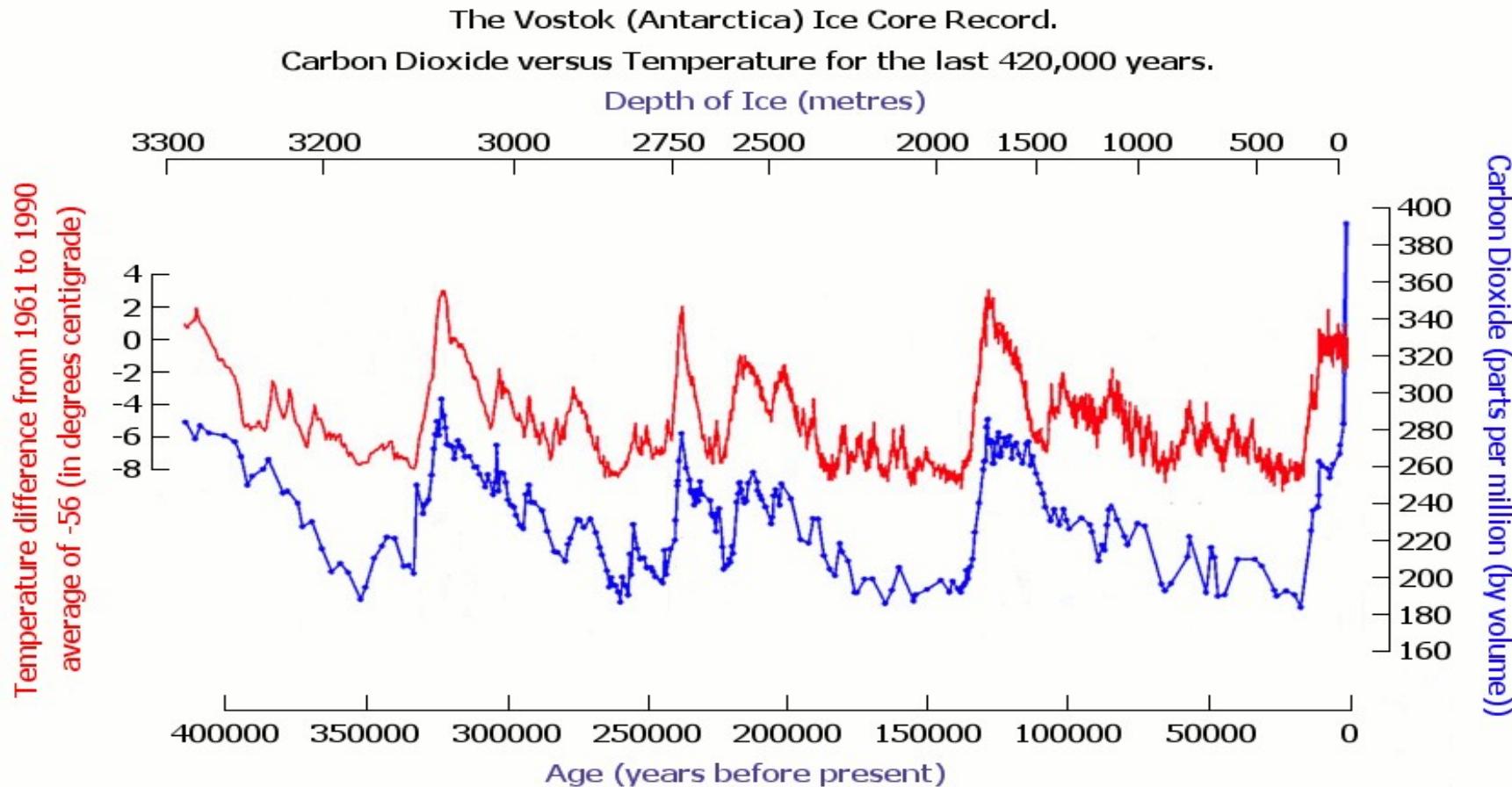
Monthly mean CO₂ concentration

Mauna Loa 1958 - 2019

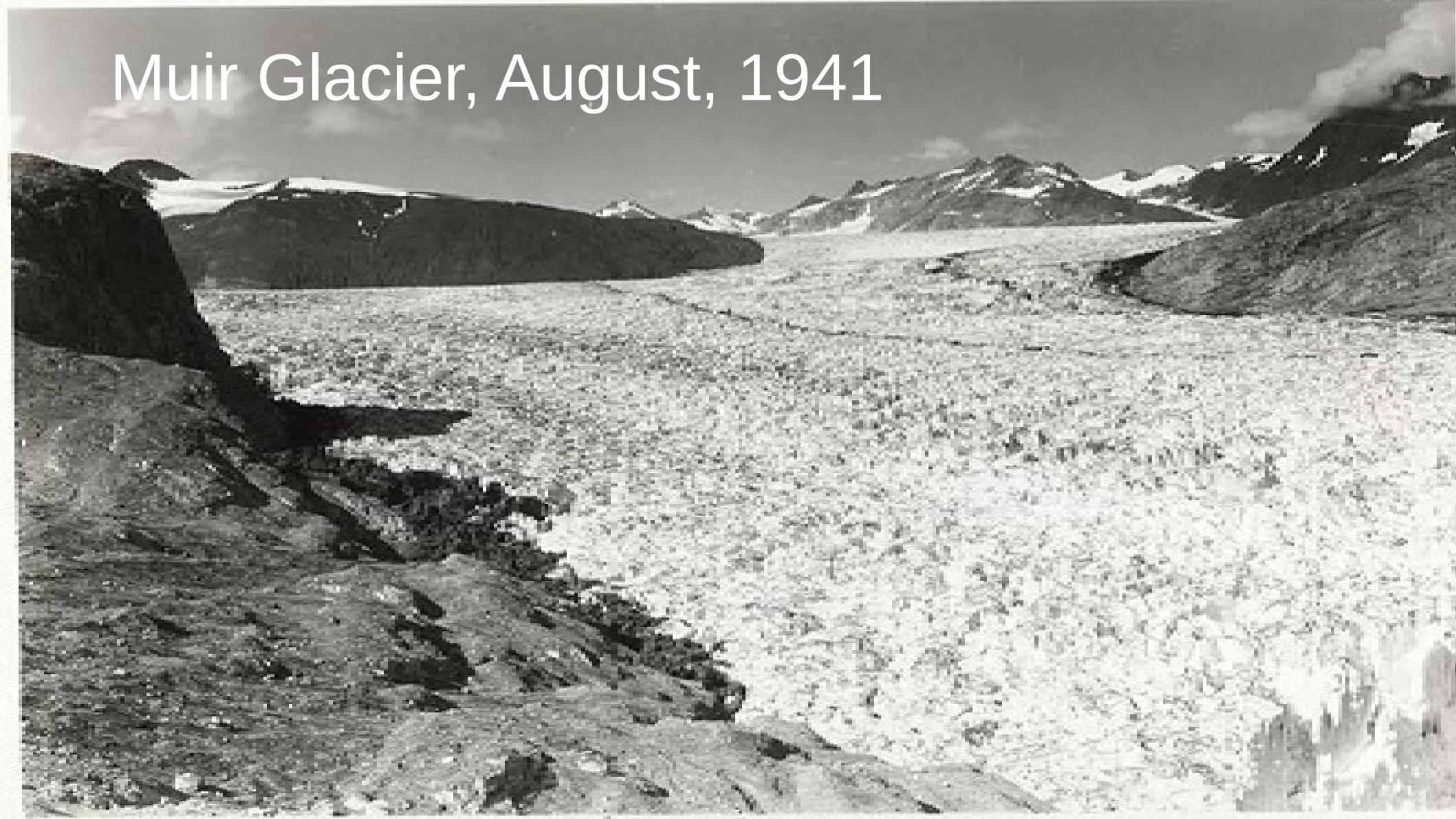


Data : R. F. Keeling, S. J. Walker, S. C. Piper and A. F. Bollenbacher
Scripps CO₂ Program (<http://scrippsc02.ucsd.edu>). Accessed 2019-07-20

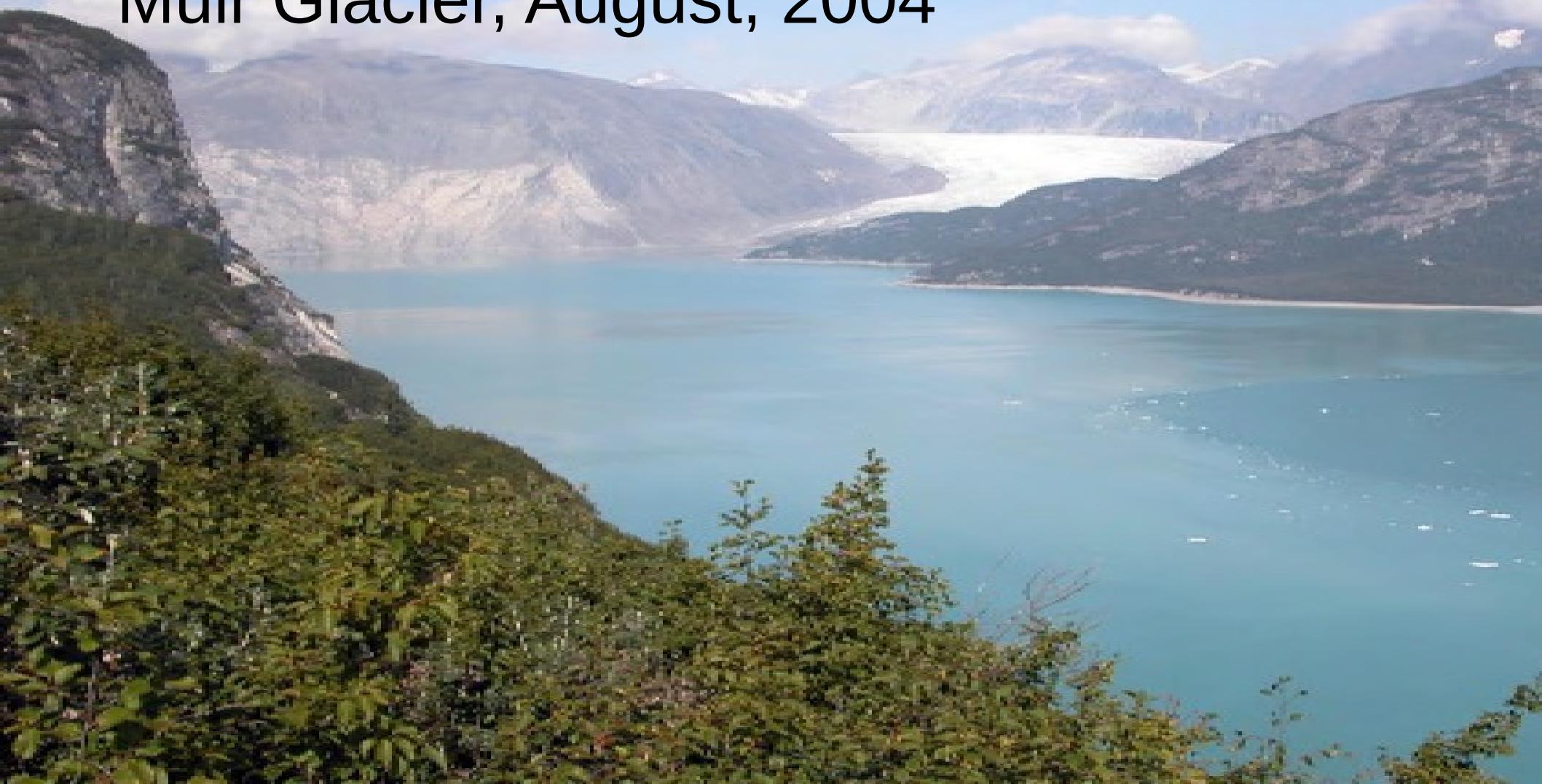
CO₂ – Temperature correlations

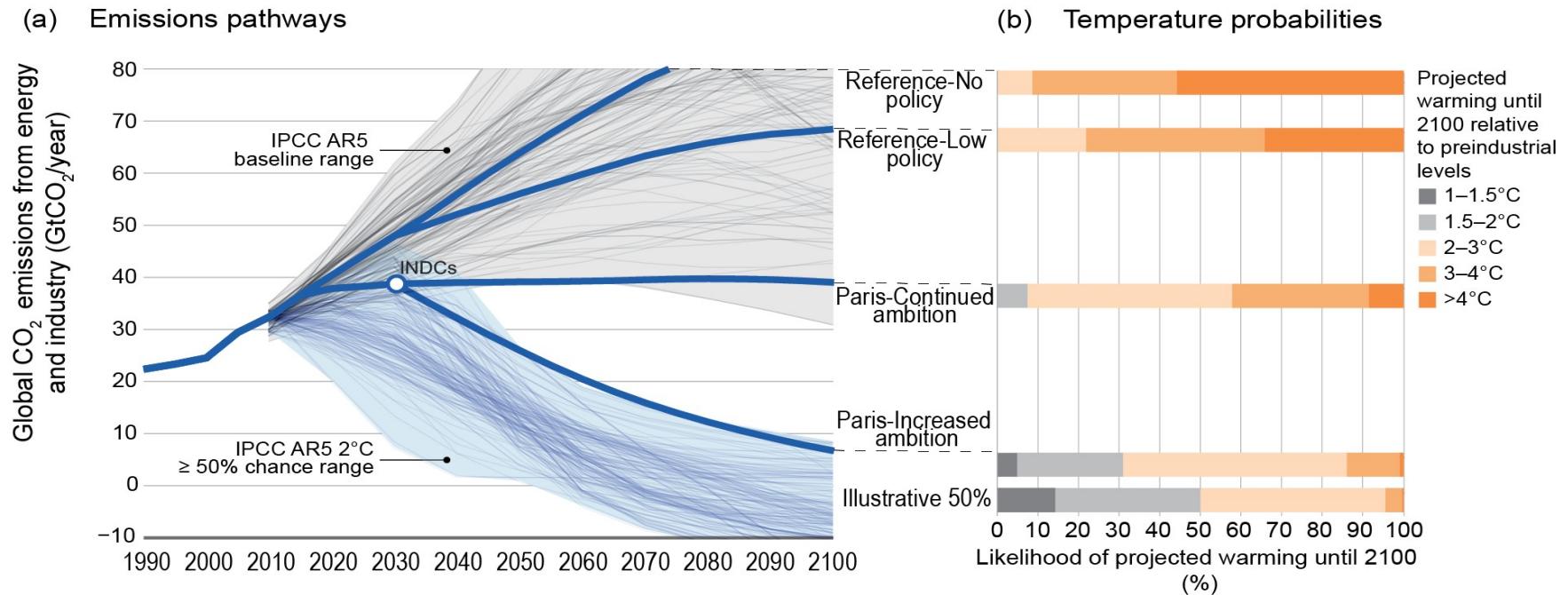


Muir Glacier, August, 1941



Muir Glacier, August, 2004



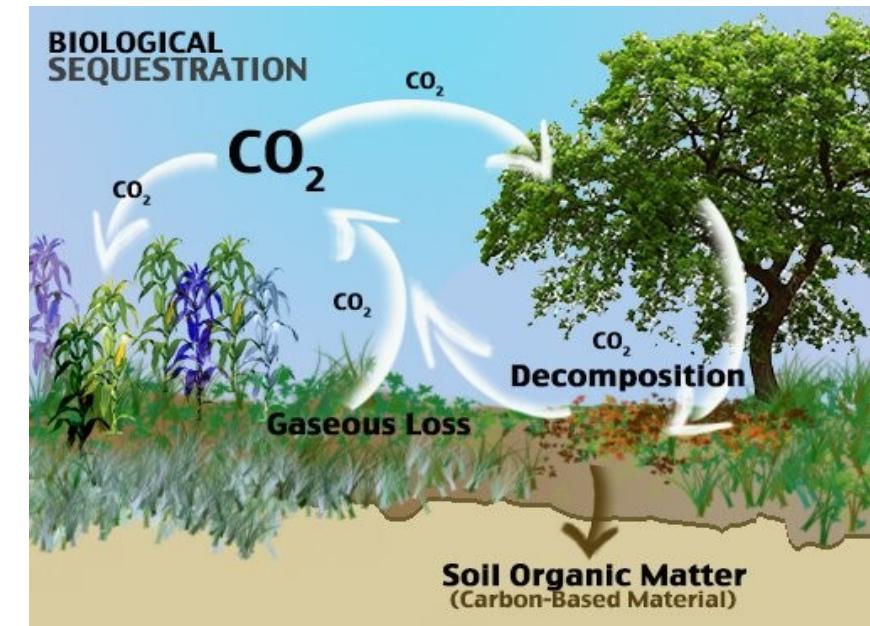


Biological Sequestration

Biological sequestration is the capture and storage of the atmospheric greenhouse gas carbon dioxide by biological processes

This may be by increased **primary productivity** (reforestation, biological carbon pump) or by **preventive measures** (avoid deforestation)

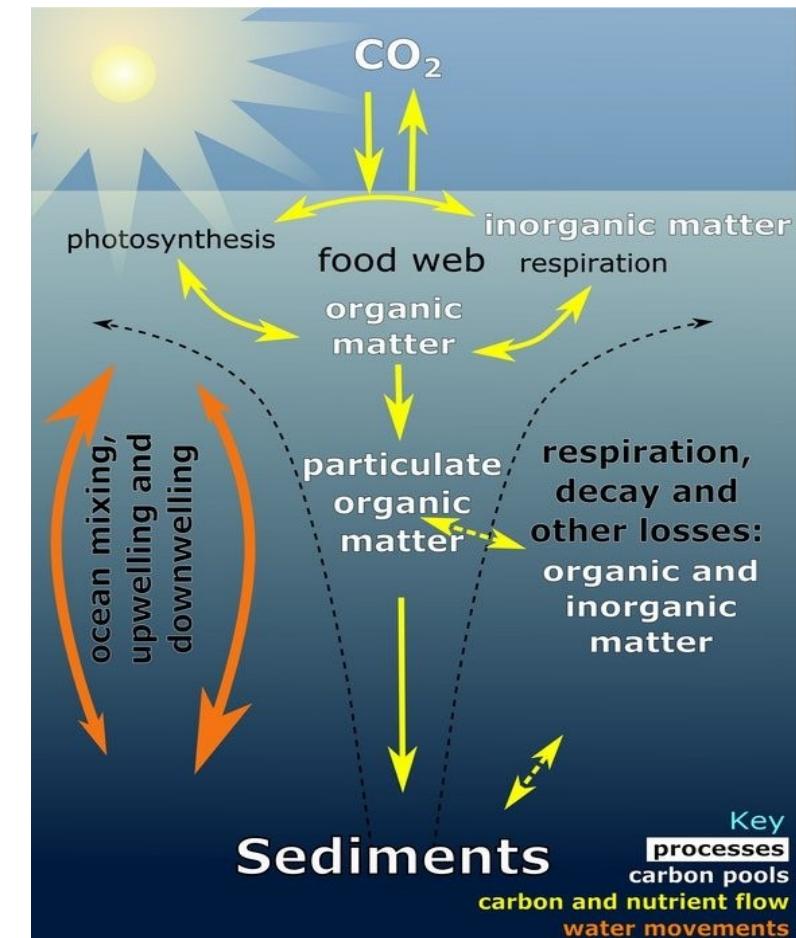
Storage can be achieved in different ways, such as through **burial** or **immobilization** in refractory fractions of organic matter



Biological Carbon Pump

The **biological pump**, in its simplest form, is the ocean's biologically driven sequestration of carbon from the atmosphere to the **ocean interior** and **seafloor sediments**

It is the part of the oceanic carbon cycle responsible for the sequestering organic matter formed mainly by phytoplankton during **photosynthesis** (soft-tissue pump), as well as the cycling of **calcium carbonate** (CaCO_3) formed into shells by certain organisms such as plankton and mollusks (carbonate pump)



Carbonation of Rocks

Carbon, in the form of CO₂ can be removed from the atmosphere by chemical processes, and stored in **stable carbonate mineral** forms. This process is known as 'carbon sequestration by mineral carbonation' or mineral sequestration.

The process involves reacting carbon dioxide with abundantly available metal oxides—either magnesium oxide (MgO) or calcium oxide (CaO)—to **form stable carbonates**.

These reactions are exothermic and **occur naturally** (e.g., the weathering of rock over geologic time periods).

Some rocks are **more prone to carbonation**, and therefore are primary target for carbon storage studies



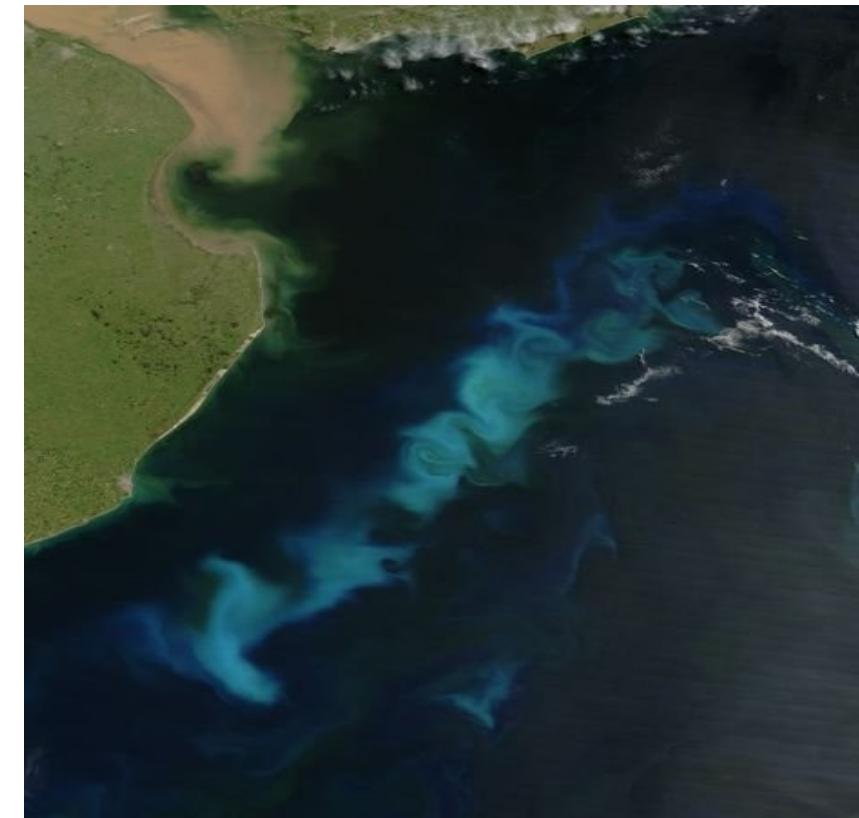
Carbon Sequestration: Ocean

Primary Productivity in the Ocean

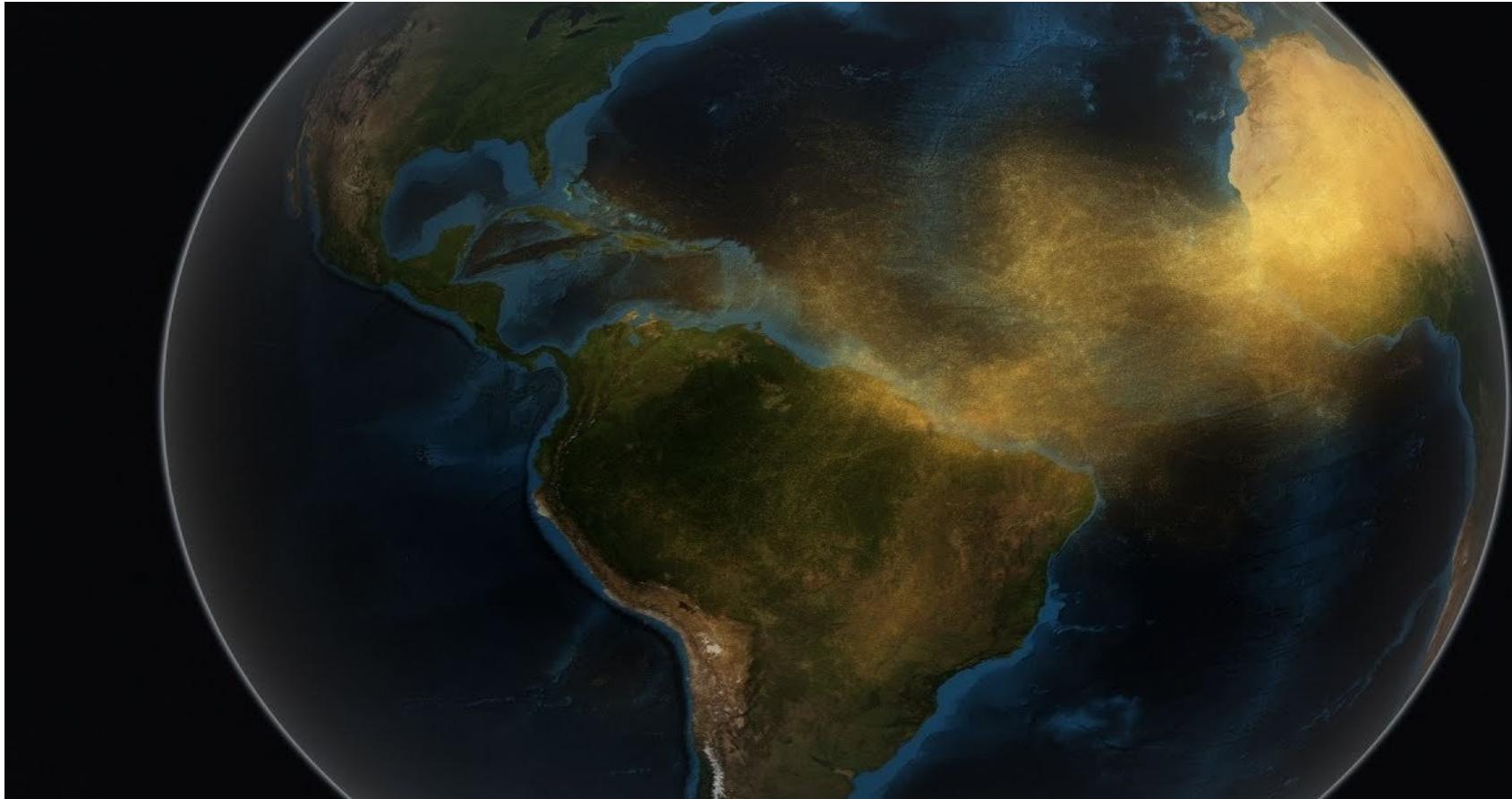
Primary productivity is limited in the ocean by the penetration of the **solar radiation**, **nutrient** availability and **iron** concentrations

Besides photosynthesis, plankton that **generate calcium or silica carbonate** skeletons, such as diatoms, coccolithophores and foraminifera, account for most direct carbon sequestration

When these organisms die 20-30% of their carbonate skeletons **sink to the seafloor** and get buried in sediments



Natural Iron Fertilization





“Give me a few oil tankers full of iron, and I’ll give you an ice age.”

– John Martin, WHOI Scientist

Iron Fertilization

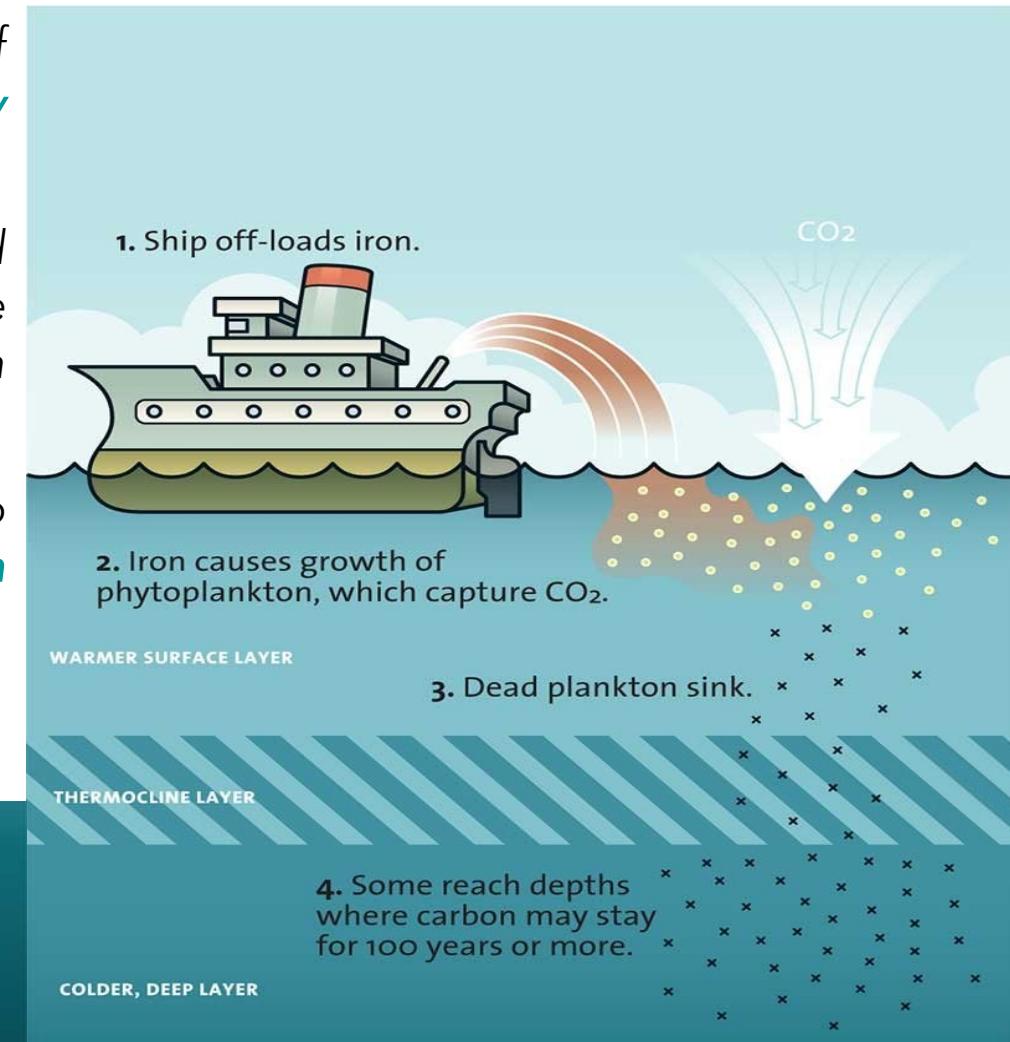
Iron fertilization refers to the release of iron in the ocean to **stimulate primary productivity**

Increased primary productivity should lead to the **increase of exports** to the seafloor with resulting increased carbon sequestration

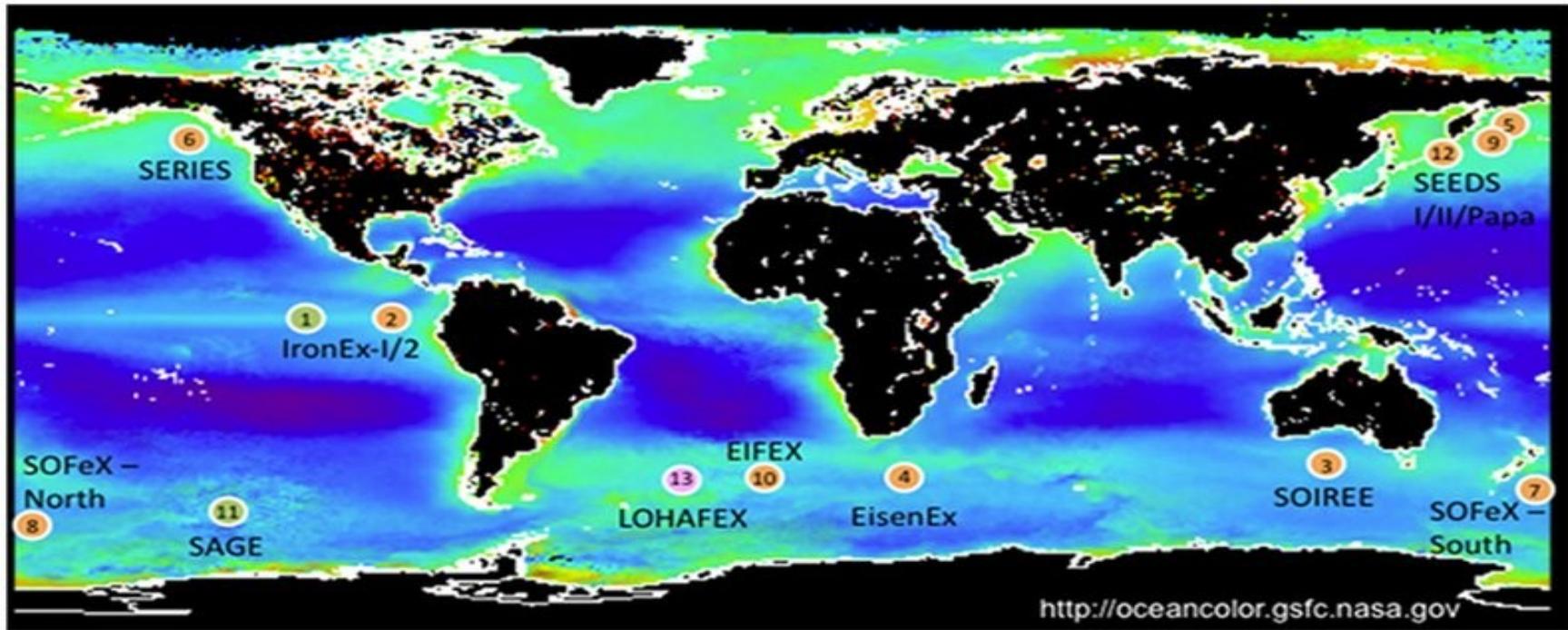
Increased sedimentation rates due to higher exports should ensure **long term burial** of the sequestered carbon

The Fantasy:
Plankton populations rebound to historic levels, reviving fisheries and sequestering vast amounts of carbon.

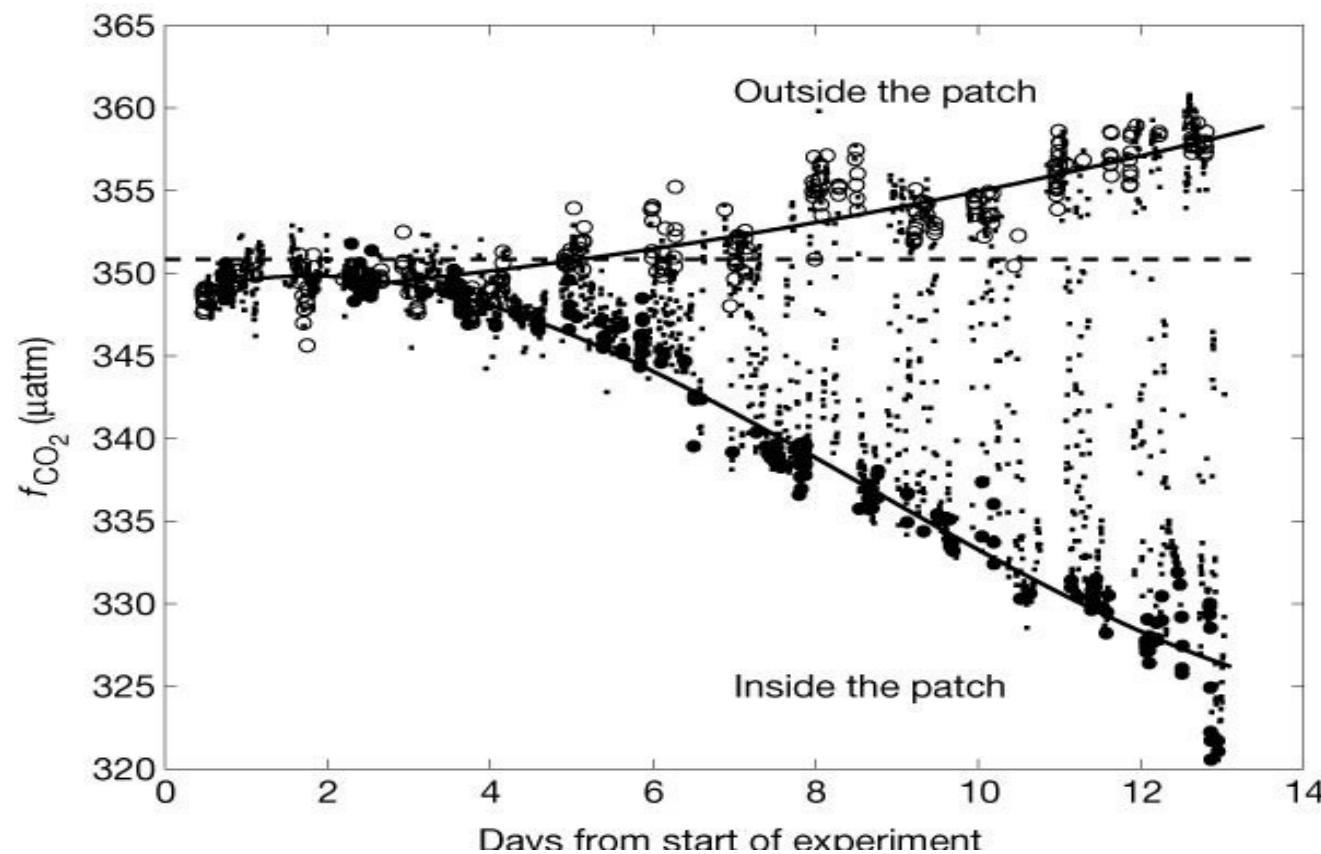
The Fear:
Iron leads to the depletion of deep-water oxygen, alters food chain, and promotes toxic species; CO₂ soon resurfaces.



Iron Fertilization



Quantifiable reduction in atmospheric CO₂ in "patches" of iron fertilization. SOFEX Experiment



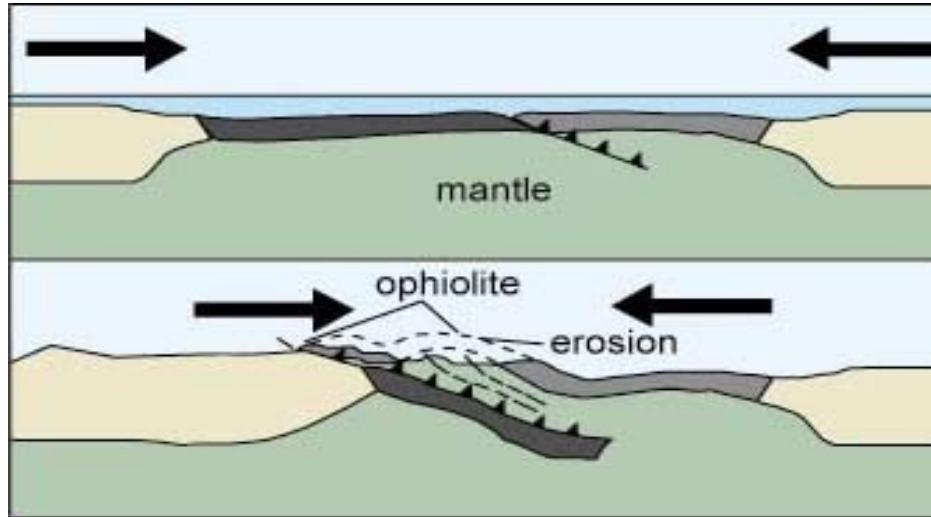


Carbon Sequestration: Underground

Natural Carbon Sequestration

An **ophiolite** is a section of the Earth's oceanic crust and the underlying upper mantle that has been uplifted and exposed above sea level and often emplaced onto continental crustal rocks

Made of mantle rocks (peridotite) rich in iron and magnesium, they react with water and CO₂, sequestering carbon into carbonate





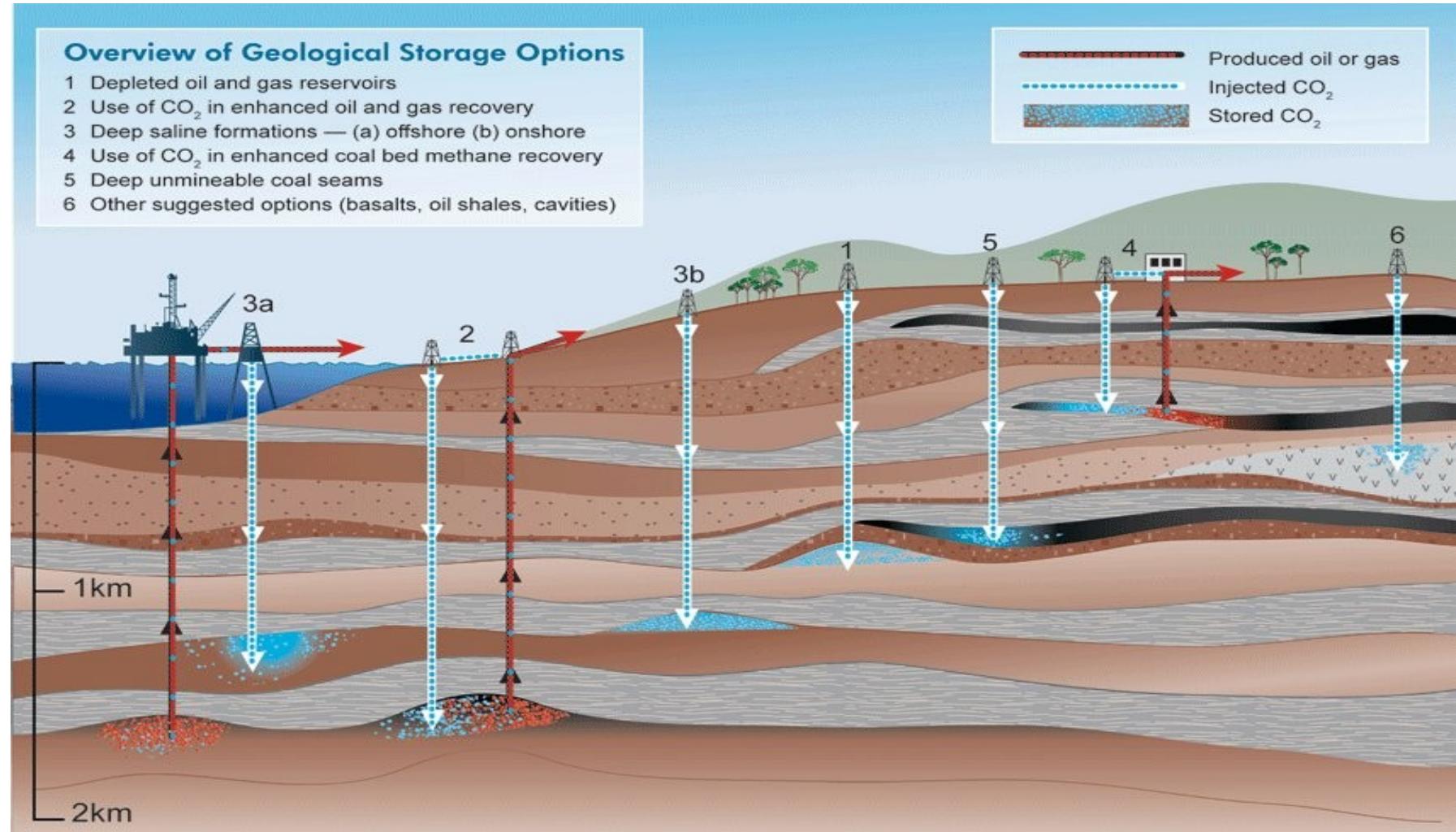
Manmade Carbon Sequestration

Geological sequestration refers to the storage of CO₂ underground in depleted oil and gas reservoirs, saline formations, or deep, un-minable coal beds

Once CO₂ is captured from a gas or coal-fired power plant, it would be compressed to ≈ 100 bar so that it would be a supercritical fluid. In this fluid form, the CO₂ would be easy to transport via pipeline to the place of storage.

The CO₂ would then be injected deep underground, typically around 1 km, where it would be stable for hundreds to millions of years.

Carbon Sequestration

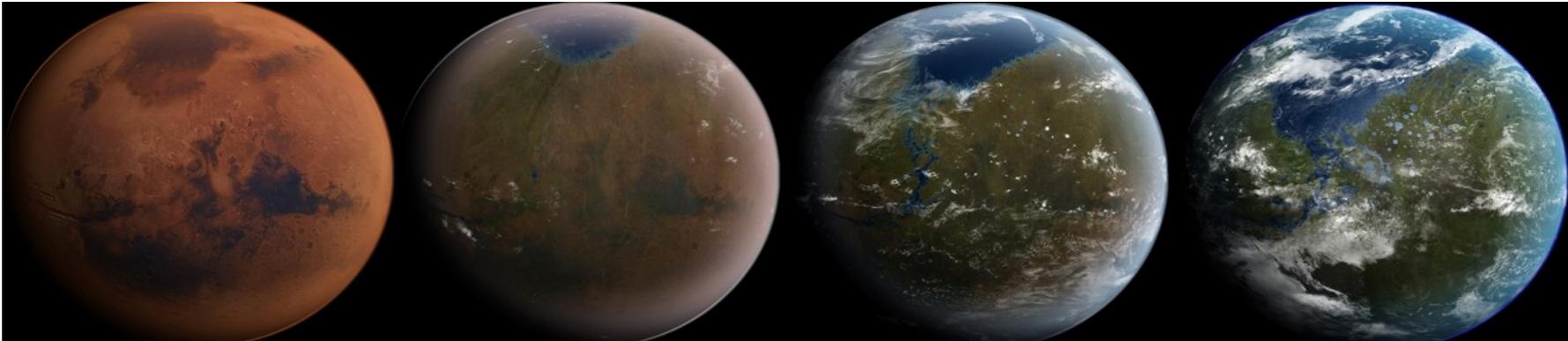


Geoengineering: not just this planet

Geoengineering concept can also be applied to **extraterrestrial planets**

In this case they are often collectively referred to as **Terraforming**

A number of project exists for example on the possible terraforming of Mars





UNDERGROUND HYDROGEN STORAGE

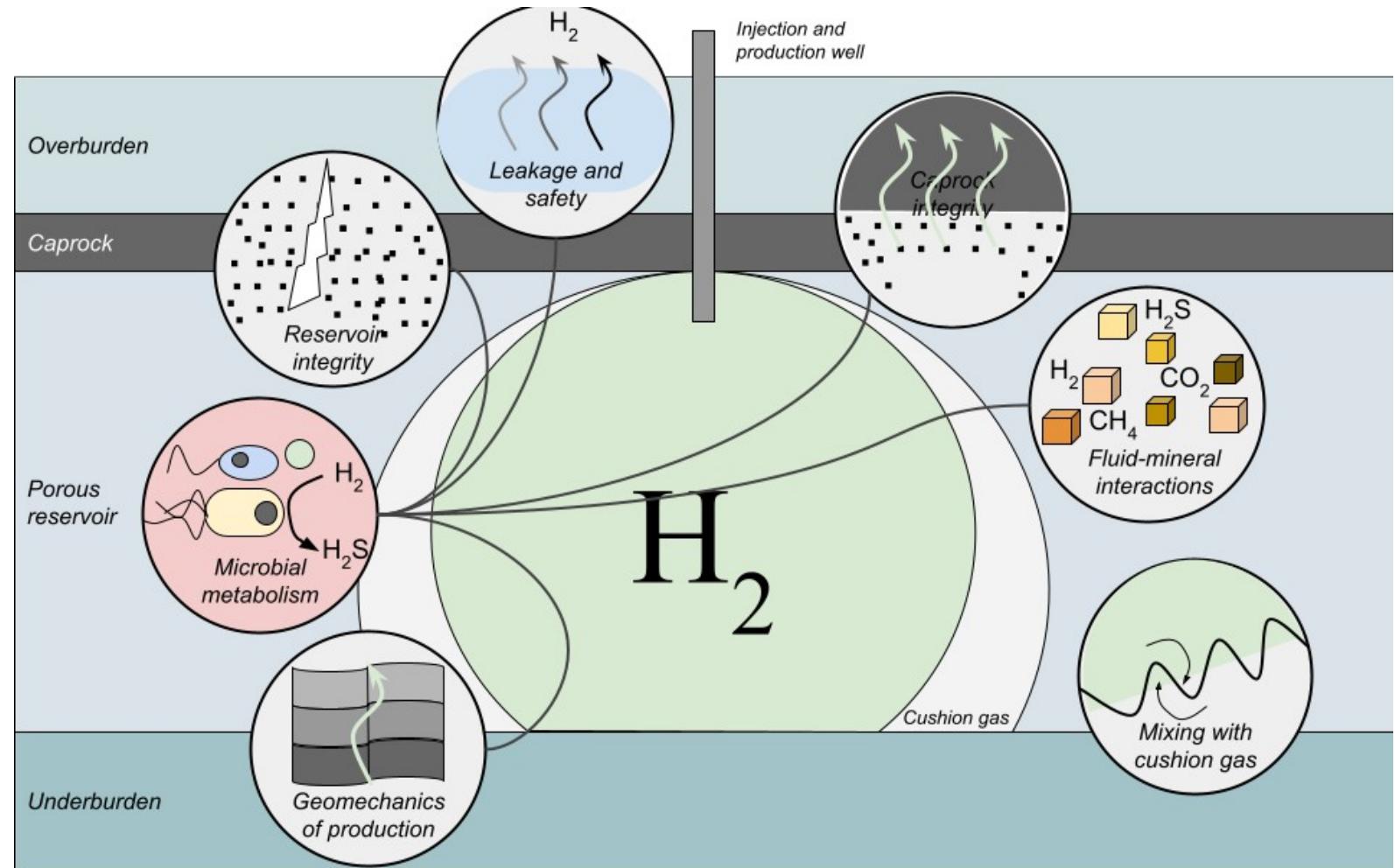
Underground Hydrogen Storage

Renewable energies (RE) will play a key role in decarbonization of our society. However they are for their nature intermittent and not uniformly distributed.

Converting excess energy from RE into hydrogen is a key technology that will allow storage and transportation to meet the societal demands. For safety reason hydrogen is better stored underground



Underground Hydrogen Storage





BIOLEACHING AND BIOMINING



Bioleaching and Biomining

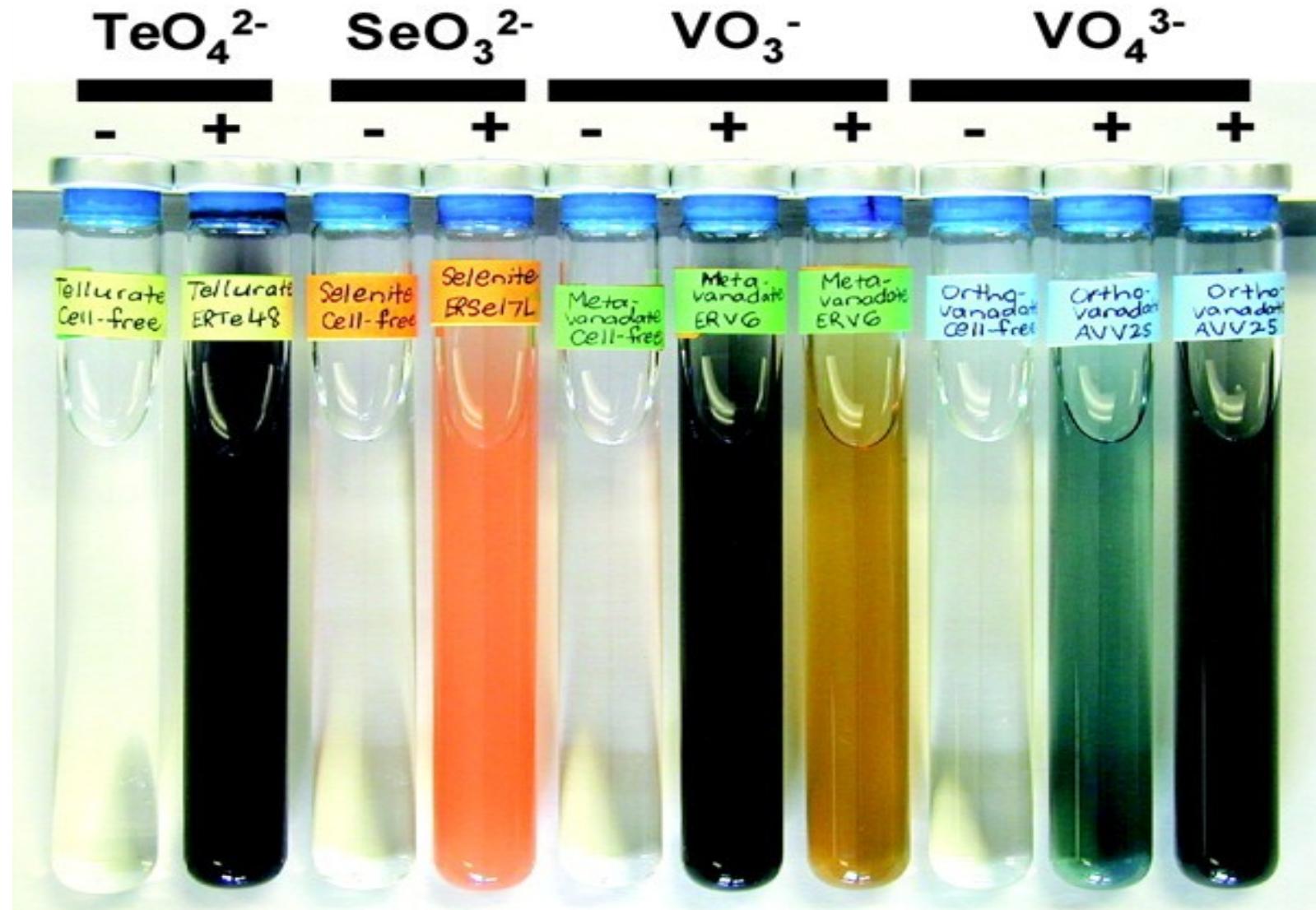
Bioleaching and Biomining are subdivision of **Biohydrometallurgy**, a branch of hydrometallurgy that deals with the employment of biotechnology in hydrometallurgy.

Biomining is the generic term that refers to the use of **biotechnology in mining operations**. The most used form of biomining is bioleaching

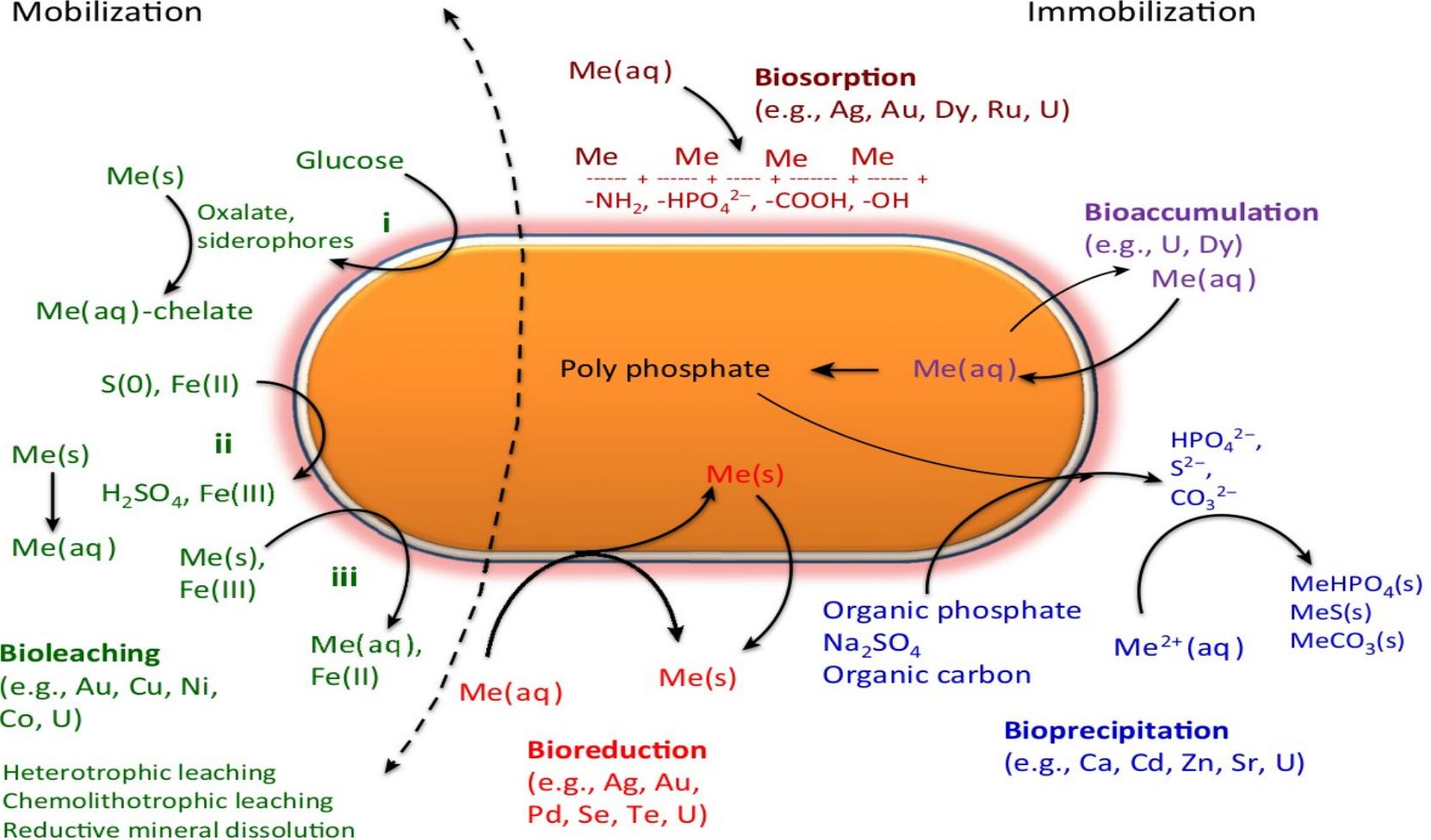
Bioleaching is the **solubilization** of metals from their ores through the use of living organisms.

Bioleaching is currently used to recover **copper, zinc, lead, arsenic, antimony, nickel, molybdenum, gold, silver, and cobalt**.





Mobilization





Bioleaching for Biomining

Until 50 years ago, nobody was aware that bacteria played an active part in ore extraction process. Despite the lack of knowledge, bioleaching had been **unknowingly used in medieval times** to extract and recover metals.

The modern era of biomining began within the discovery of a bacterium able to catalyse the oxidation of Fe^{3+} to Fe^{2+} in low pH (*Thiobacillus ferrooxidans*, renamed ***Acidithiobacillus ferrooxidans***) and accelerate the oxidative dissolution of pyrite (FeS_2), the most abundant sulfide mineral in the lithosphere.

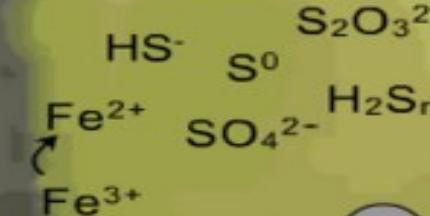
By the **mid-1960s** the **bioleaching** of run-of-mine copper waste rocks in mines operated by the (then) Noranda corporation in the USA started.

Today microbe-aware mining turn billions of tons of low-value copper ore into pure copper. United States, Canada, Chile, Australia, and South Africa produce **>25% of the total copper** production through bioleaching. More than **10% of gold** and **3% of cobalt** and **nickel** are biotechnologically produced.

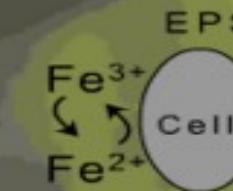
Sulfide mineral Leachate

Non-contact leaching

Fe³⁺ or Fe³⁺-proton attack

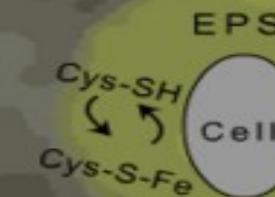


Contact leaching



Cooperative leaching

Sulfur-colloids, sulfur intermediates and mineral fragments for planctonic iron- and sulfur-oxidizers



Sulfur-oxidizers

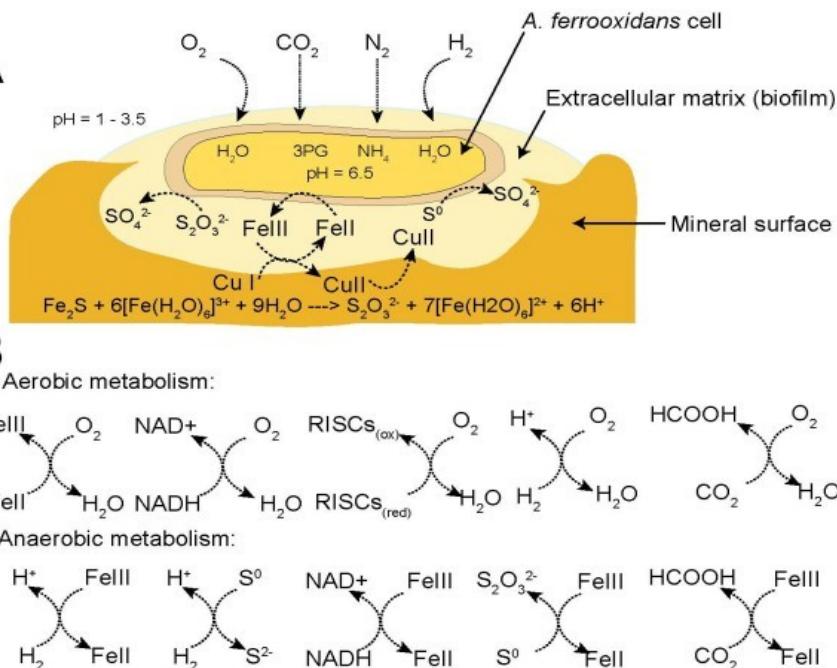
Acidithiobacillus ferrooxidans

Acidithiobacillus ferrooxidans (formerly *Thiobacillus*) is a Gram-negative, **Gammaproteobacterium** that thrives optimally at 30°C and pH 2, but can grow at pH 1 or lower.

It is **abundant in natural environments** associated with pyritic ore bodies, coal deposits, and their acidified drainages.

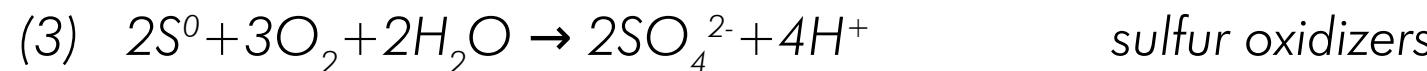
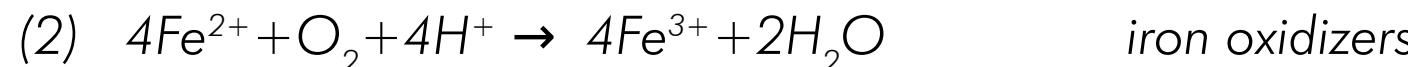
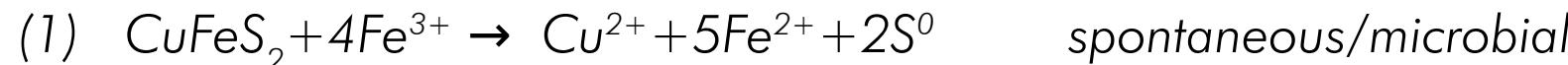
It is capable of coupling **Fe²⁺, elemental sulfur** and **thiosulfate oxidation** to oxygen respiration.

It is an important member of **microbial consortia** used to recover copper via a process known as bioleaching or biomining.



Bioleaching of Chalcopyrite

Chalcopyrite bioleaching is a multistep process, involving both direct and indirect bioleaching



In general, sulfides are first oxidized to elemental sulfur, whereas disulfides are oxidized to give thiosulfate, and the processes above can be applied to other sulfidic ores.

Bioleaching of non-sulfidic ores such as pitchblende also uses ferric iron as an oxidant (e.g., $\text{UO}_2 + 2\text{Fe}^{3+} \rightarrow \text{UO}_2^{2+} + 2\text{Fe}^{2+}$). In this case, the sole purpose of the bacterial step is the regeneration of Fe^{3+} .



Copper mine aerial view



Environmental issues: Mine Drainage Fluids

Aside from the commercial use of microbiology to facilitate mining operations, microbiology also interacts with mining wastes, often in unwanted manners.

The most abundant mining waste in many cases are the **Mine Drainage Fluids**. These are composed by high amounts of **sulfide**, **heavy metals** and **very low pH**.

Mine Drainage Fluids are responsible for the vast majority of the mining accidents and spills threatening neighboring ecosystems

The microbiology of Mine Drainage Fluids has been extensively studied, and the body of knowledge collected can help both **biomining** operations as well as **bioremediation** efforts.

Mine Drainage Fluids



Mine Drainage Fluids





Recovery of e-wastes

Electronic waste or e-waste describes discarded electrical or electronic devices.

Up to **60 elements** can be found in complex electronics. In the United States alone, an estimated **70% of heavy metals in landfills** comes from discarded electronics.

The short lifespan and rapid technology turnover makes electronics one of the **fast growing special waste** of the future.

Additionally many of the **elements are considered endangered**, and their recovery will become of primary importance for the sustainability of our society.

Informal processing of e-waste in **developing countries** can lead to **adverse human health effects** and **environmental pollution**.

ELEMENTS OF A SMARTPHONE

ELEMENTS COLOUR KEY: ● ALKALI METAL ● ALKALINE EARTH METAL ● TRANSITION METAL ● GROUP 13 ● GROUP 14 ● GROUP 15 ● GROUP 16 ● HALOGEN ● LANTHANIDE

SCREEN

49 In Indium	8 O Oxygen
50 Sn Tin	

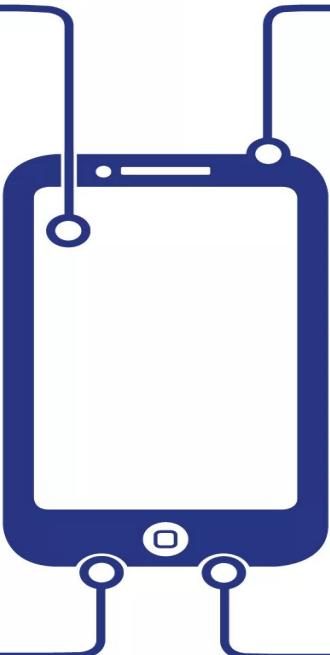
Indium tin oxide is a mixture of indium oxide and tin oxide, used in a transparent film in the screen that conducts electricity. This allows the screen to function as a touch screen.

13 Al Aluminium	14 Si Silicon
8 O Oxygen	19 K Potassium

The glass used on the majority of smartphones is an aluminosilicate glass, composed of a mix of alumina (Al_2O_3) and silica (SiO_2). This glass also contains potassium ions, which help to strengthen it.

39 Y Yttrium	57 La Lanthanum	65 Tb Terbium
59 Pr Praseodymium	63 Eu Europium	66 Dy Dysprosium
64 Gd Gadolinium		

A variety of Rare Earth Element compounds are used in small quantities to produce the colours in the smartphone's screen. Some compounds are also used to reduce UV light penetration into the phone.



ELECTRONICS

Copper is used for wiring in the phone, whilst copper, gold and silver are the major metals from which microelectrical components are fashioned. Tantalum is the major component of micro-capacitors.

29 Cu Copper	47 Ag Silver
79 Au Gold	73 Ta Tantalum

Nickel is used in the microphone as well as for other electrical connections. Alloys including the elements praseodymium, gadolinium and neodymium are used in the magnets in the speaker and microphone. Neodymium, terbium and dysprosium are used in the vibration unit.

28 Ni Nickel	66 Dy Dysprosium	59 Pr Praseodymium
65 Tb Terbium	60 Nd Neodymium	64 Gd Gadolinium

Pure silicon is used to manufacture the chip in the phone. It is oxidised to produce non-conducting regions, then other elements are added in order to allow the chip to conduct electricity.

14 Si Silicon	8 O Oxygen	51 Sb Antimony
33 As Arsenic	15 P Phosphorus	31 Ga Gallium

Tin & lead are used to solder electronics in the phone. Newer lead-free solders use a mix of tin, copper and silver.

50 Sn Tin	82 Pb Lead
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BATTERY

3 Li Lithium	27 Co Cobalt	8 O Oxygen
6 C Carbon	13 Al Aluminium	

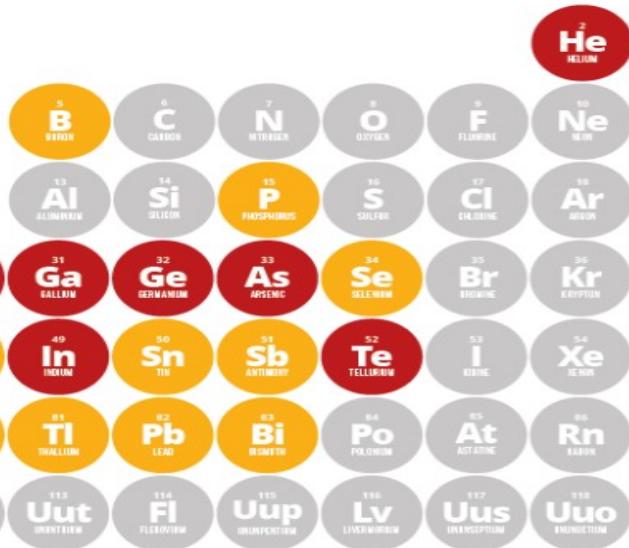
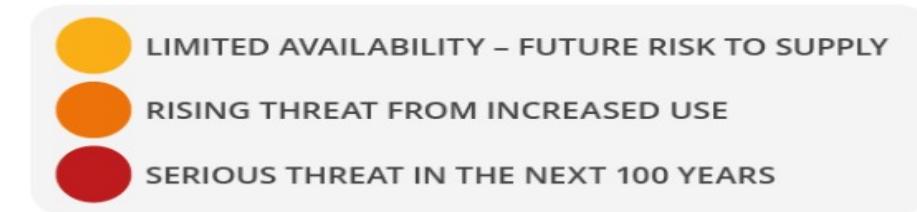
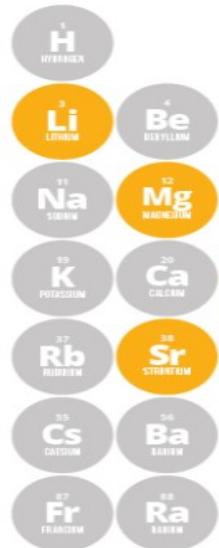
The majority of phones use lithium ion batteries, which are composed of lithium cobalt oxide as a positive electrode and graphite (carbon) as the negative electrode. Some batteries use other metals, such as manganese, in place of cobalt. The battery's casing is made of aluminium.

Magnesium compounds are alloyed to make some phone cases, whilst many are made of plastics. Plastics will also include flame retardant compounds, some of which contain bromine, whilst nickel can be included to reduce electromagnetic interference.

6 C Carbon	12 Mg Magnesium
35 Br Bromine	28 Ni Nickel



THE PERIODIC TABLE'S ENDANGERED ELEMENTS



SOURCE: CHEMISTRY INNOVATION KNOWLEDGE TRANSFER NETWORK



BIOENGINEERING



A timeline of CRISPR discovery

1987 – Ishino **accidentally cloned part of a CRISPR** together with the *E. coli* gene. The function of the clustered repeats was unknown at the time

1993 – Van Soolingen and colleagues reported in two papers clusters of interrupted direct repeats in *Mycobacterium tuberculosis* used their diversity to devise a named **spoligotyping** (still in use) to investigate *M. tuberculosis* diversity

2001 – Mojica and Jansen, working on *Haloferax*, reported the presence of the cluster in many microbes and **proposed the acronym CRISPR**

2002 – Tang showed that CRISPR repeat from *Archaeoglobus fulgidus* were **transcribed into long RNA** subsequently processed into small RNAs. At the same time Jansen described the **CRISPR-associated systems genes**



2005 – Three independent research groups showed that some CRISPR spacers are **derived from phage DNA** and extrachromosomal DNA such as plasmids and Mojica proposed a role of CRISPR-Cas in microbial immunity. All were rejected from high profile journals

2010 – The first experimental evidence that CRISPR was an **adaptive immune system** was published

2012 – Doudna and Charpentier (UC Berkeley, USA) **re-engineered the Cas9** endonuclease manageable two-component system. By manipulating the guide RNA, the artificial Cas9 system could be programmed to target any DNA. Two more teams, Šikšnys (Vilnius University, Lithuania) and Zhang (MIT, USA)

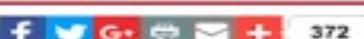
2013 – Two groups simultaneously described **genome editing in human cell cultures** using CRISPR-Cas9 for the first time

By the end of **2014** some **1,000 research papers** had been published that mentioned CRISPR



New Scientist Live Last chance to book

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SHORT SHARP SCIENCE 25 January 2017

Gene editing has saved the lives of two children with leukaemia



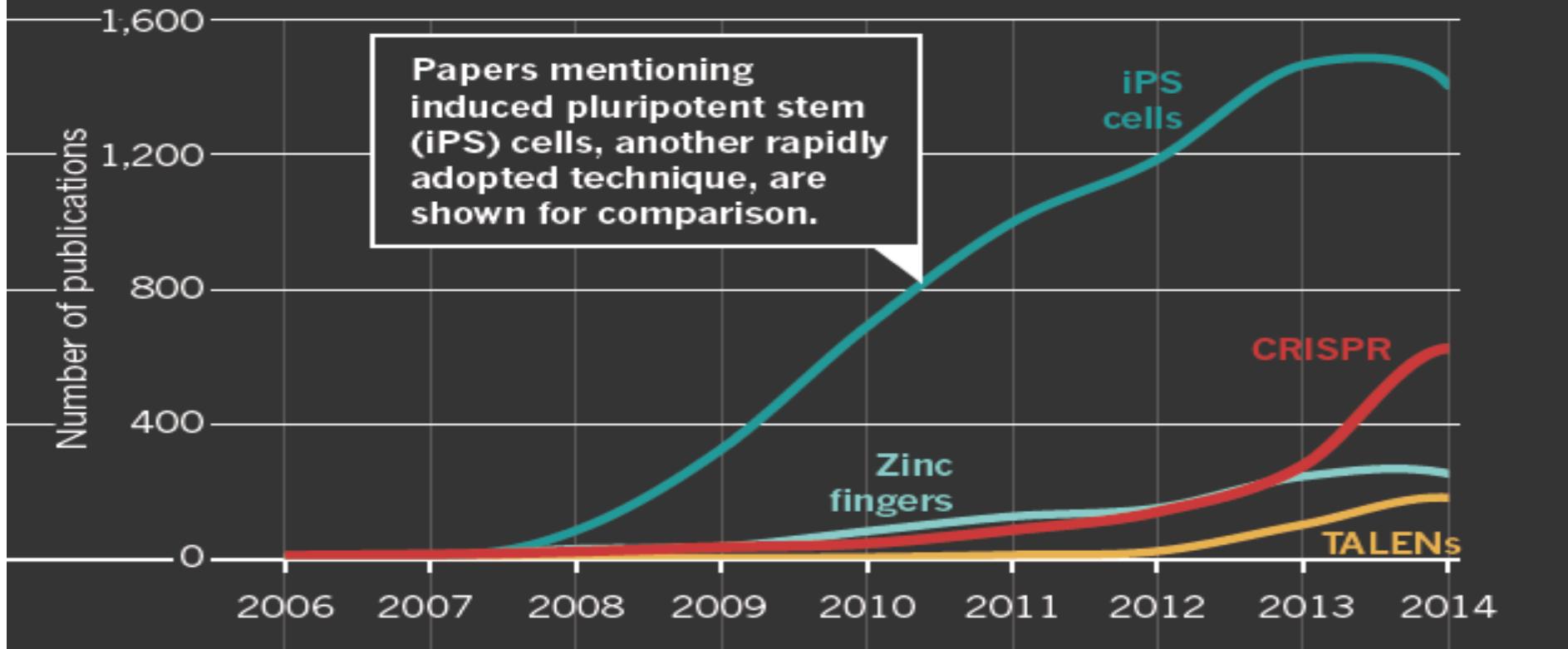


CRISPR-Cas is a *vertically transmitted adaptive immune system* against phage infection present in a large number of Bacteria and Archaea

CRISPR-Cas is a *tool for targeted genetic engineering* (or gene editing) extremely diffused in modifying eukaryotic genomes

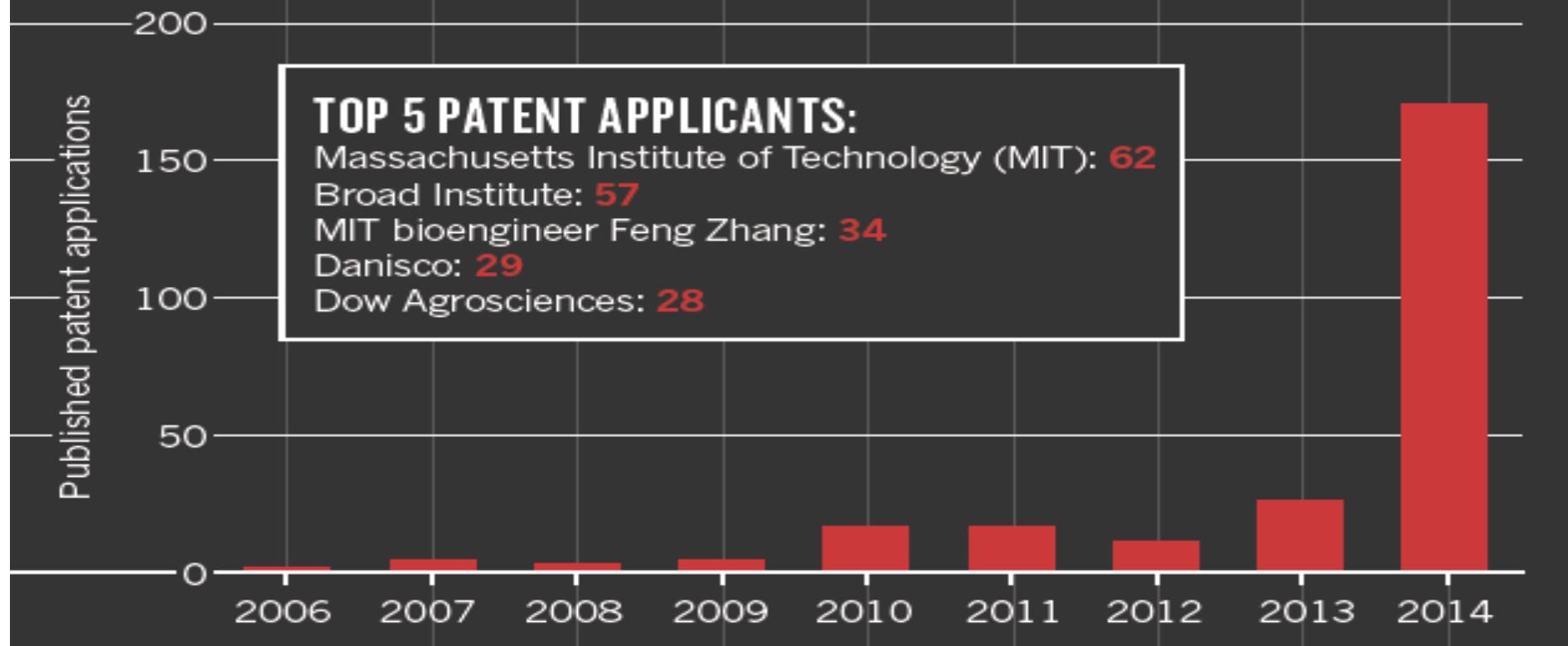
PUBLICATIONS

The number of papers about CRISPR has outstripped the numbers mentioning the gene-editing technologies known as TALENs and zinc fingers.



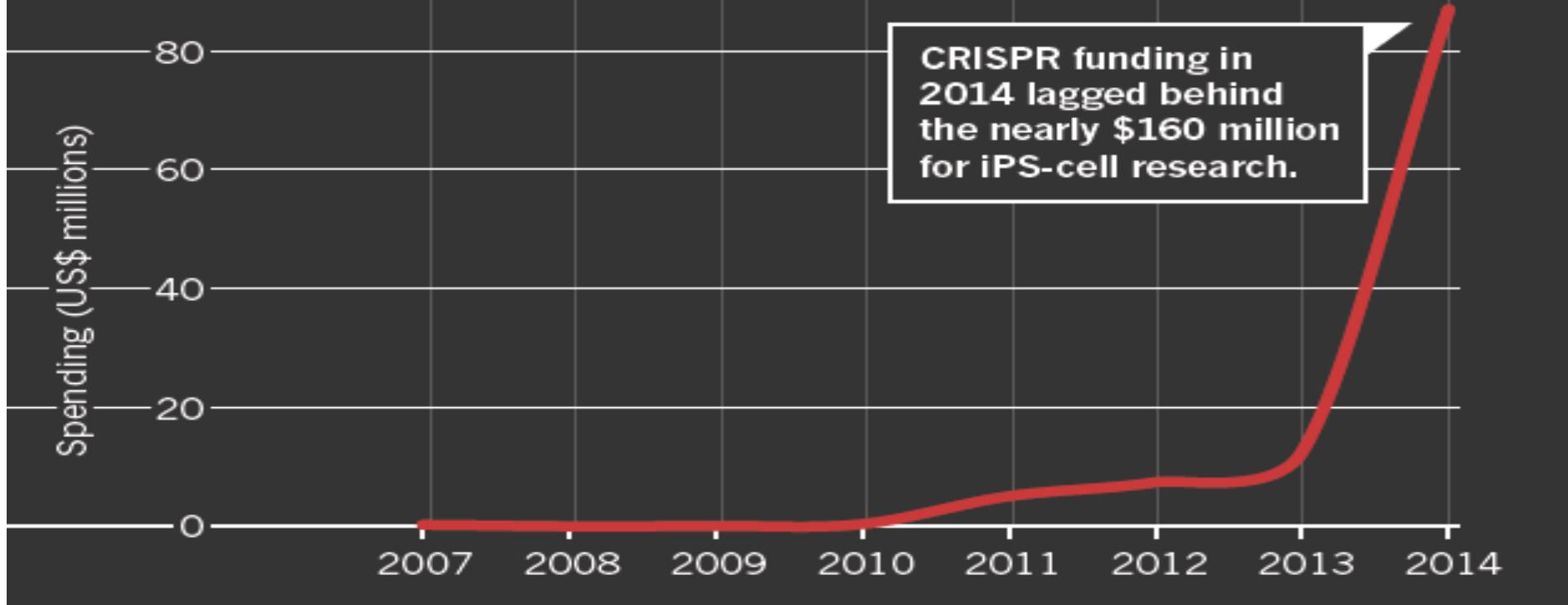
PATENTS

In 2014, worldwide patent applications that mention CRISPR leapt and a patent battle intensified.



FUNDING

A sharp jump in US National Institutes of Health funding for projects involving CRISPR is a harbinger of future advances.





"It seems appropriate to remember here that the great motor of molecular biology during the 20th century, molecular cloning, was made possible by restriction endonucleases and plasmid vectors, both derived from serendipitous discoveries of prokaryotic cell biology features involved in self/nonself discrimination as well. The lesson here to scientists, science policy makers and mankind at large is that ***the only way forward is enlarging evenly the sphere of knowledge supporting fundamental research.*** In words of Louis Pasteur: 'There does not exist a category of science to which one can give the name applied science. There are science and the applications of science, bound together as *the fruit of the tree which bears it*'."

Mojica and Rodriguez-Valera, 2016 FEBS Journal

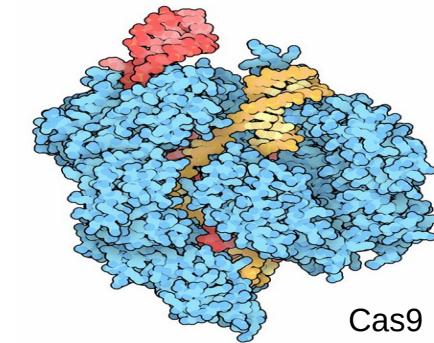
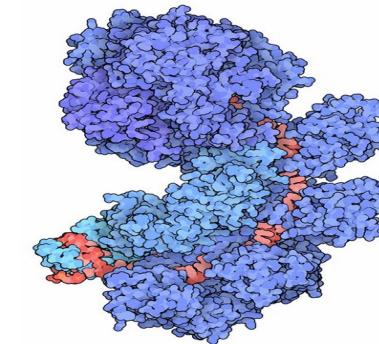
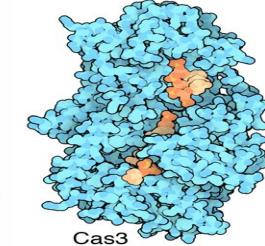
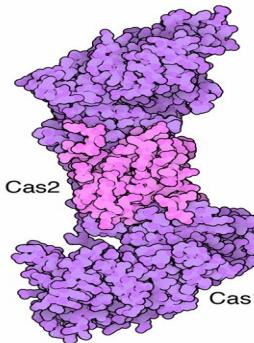
CRISPR-associated system

CRISPR repeat-spacer arrays are generally associated to protein coding genes called **cas genes**. A complete CRISPR-Cas locus has at least one gene belonging to the cas core.

Collectively the **93 cas genes** are grouped into **35 families** based on sequence similarity of the encoded proteins. 11 of the 35 families form the cas core.

CRISPR-Cas systems fall into two classes. **Class 1** systems use a complex of **multiple Cas proteins** to degrade foreign nucleic acids. **Class 2** systems use a **single large Cas protein** for the same purpose (see www.rcsb.org)

Cascade



Cas9

Coevolution

CRISPR sequences revealed **coevolution** of host and viral genomes.

The basic model of CRISPR evolution is newly incorporated spacers driving phages to mutate their genomes to avoid the bacterial immune response, creating diversity in both the phage and host populations.

To fight off a phage infection, the sequence of the CRISPR spacer **must correspond perfectly** to the sequence of the target phage gene.



"Now, here, you see, it takes all the running you can do, to keep in the same place." - Alice:
Through the Looking Glass



BIOACTIVE COMPOUNDS

Thermus aquaticus gen. n. and sp. n., a Non-sporulating Extreme Thermophile

THOMAS D. BROCK AND HUDSON FREEZE

Department of Microbiology, Indiana University, Bloomington, Indiana 47401

Received for publication 15 January 1969

The isolation of a new thermophilic bacterium, *Thermus aquaticus* gen. n. and sp. n., is described. Successful enrichment requires incubation at 70 to 75 C, and the use of nutrient media relatively dilute with respect to the organic components. Strains of *T. aquaticus* have been isolated from a variety of thermal springs in Yellowstone National Park and from a thermal spring in California. The organism has also been isolated from man-made thermal habitats, such as hot tap water, in geographical locations quite distant from thermal springs. Isolates of *T. aquaticus* are gram-negative nonsporulating nonmotile rods which frequently form long filaments at supraoptimal temperatures or in the stationary phase. All isolates form a yellow cellular pigment, probably a carotenoid. A characteristic structure formed by all isolates is a large sphere, considerably larger than a sphaeroplast. These large spheres, as well as lysozyme-induced sphaeroplasts, are resistant to osmotic lysis. Deoxyribonucleic acid base compositions of four strains were determined by CsCl density gradient ultracentrifugation and found to be between 65.4 and 67.4 moles per cent guanine plus cytosine. The growth of all isolates tested is inhibited by fairly low concentrations of cycloserine, streptomycin, penicillin, novobiocin, tetracycline, and chloramphenicol. Nutritional studies on one strain showed that it did not require vitamins or amino acids, although growth was considerably faster in enriched than in synthetic medium. Several sugars and organic acids served as carbon sources, and either NH₄⁺ or glutamate could serve as nitrogen source. The organism is an obligate aerobe and has a pH optimum of 7.5 to 7.8. The optimum temperature for growth is 70 C, the maximum 79 C, and the minimum about 40 C. The generation time at the optimum is about 50 min. The possible relationships of this new genus to the myxobacteria, flexibacteria, and flavobacteria are discussed.



1967 Brooks explained: "Bacteria are able to grow [...] at any temperature at which there is liquid water, even in pools which are above the boiling point."

In 1976 the thermostable enzyme DNA polymerase was first isolated from *Thermus aquaticus*

In 1983 the Taq enzyme became the cornerstone of Kary Mullis invention of the Polymerase Chain Reaction (PCR) for which he won the Noble Prize in 1993

Thermostable polymerase enzymes derived from Taq are now an industry worth ca. 400 Million euro/year

PCR and polymerases are key technologies for biological, genetics, biomedicine and biotechnology research, a global business of hundred billions of euro

Bioprospecting is defined as a systematic and organized search for useful products derived from bioresources including plants, microorganisms, animals, etc., that can be developed further for commercialization and overall benefits of the society.

From: Medicinal Spices and Vegetables from Africa, 2017

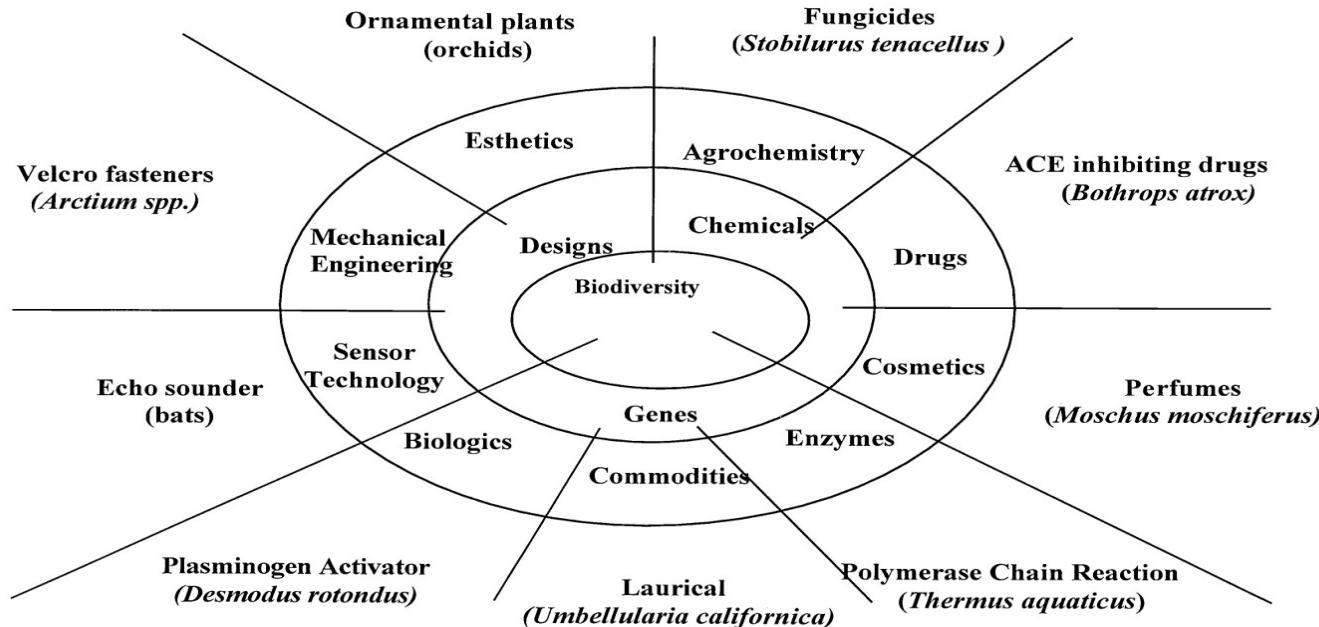
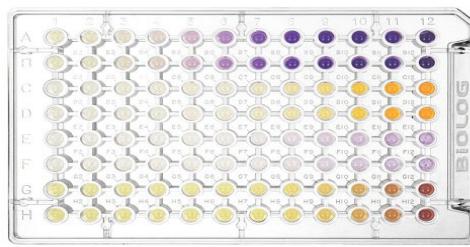


FIGURE 1 The three principal sources of inspirations from biodiversity and their applications. Product examples are described in the text (with variations derived from Nader and Hill, 1999).

Bioprospecting program	Global health issues	Sources	Parameters of species selection	Complementary approaches	Products to obtain in a bioprospecting program
Terrestrial	Antimicrobial resistance	Plants	Ethnopharmacology	Metagenomics	Pharmaceuticals
Freshwater origin	Cardiovascular diseases	Fungi	Chemotaxonomy	Proteomics	Chemical scaffolds
Marine	Cancer	Animals	Ecological chemistry	Metabolomics	Bioprocesses
Polar regions	Malnutrition	Algae	Random search	Combinatorial chemistry	Formulations
Extreme environments	Mental diseases	Insects	Bioassay-guided isolation	Medicinal chemistry	Genetic modified organisms
	Pollution	Ciliates		Organic synthesis	
	Emerging infectious diseases	Bacteria			
		Archaea			
		Microbial endophytes			
		Marine organisms			

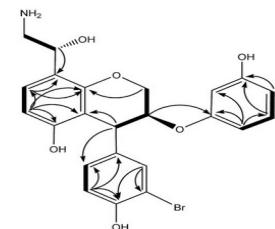
Table 1: Parameters for to be considered in a rational bioprospecting program.

Juan, 2017 J Microb Biochem Technol

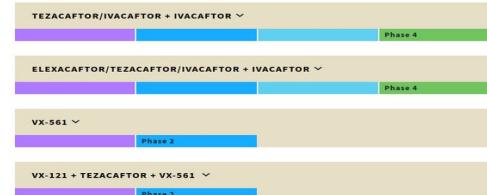


Access Biological Resources

Screen for Activity



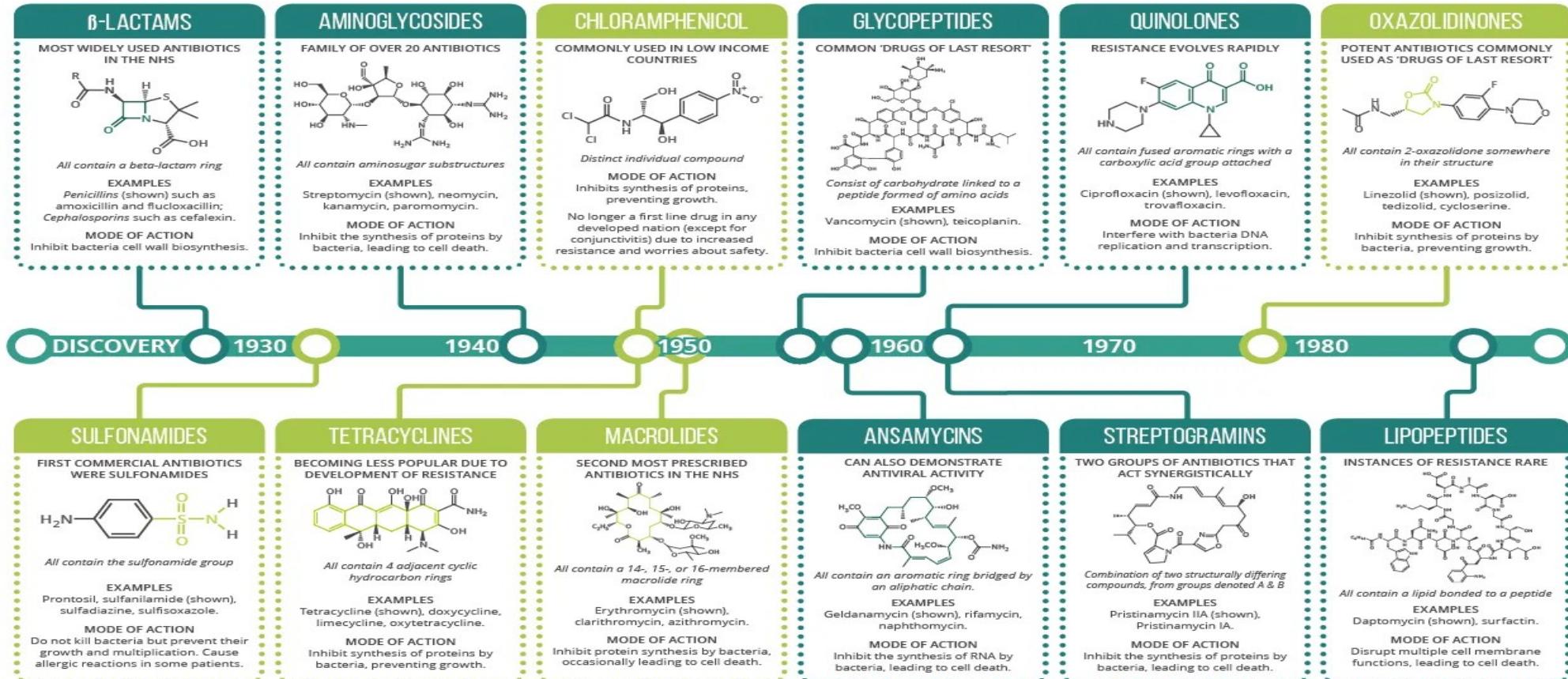
Isolate and Characterize



Test in Controlled Trials

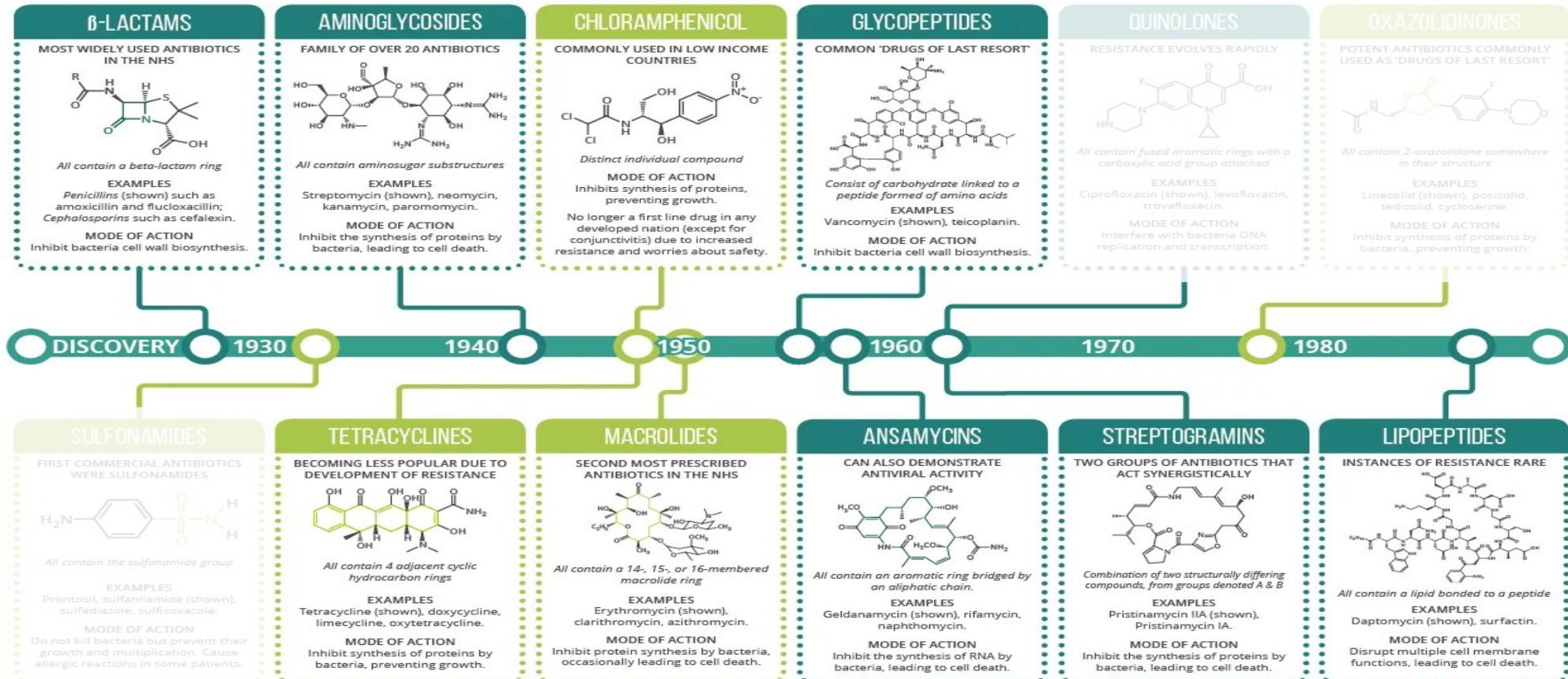
DIFFERENT CLASSES OF ANTIBIOTICS - AN OVERVIEW

Key: ● COMMONLY ACT AS BACTERIOSTATIC AGENTS, RESTRICTING GROWTH & REPRODUCTION ● COMMONLY ACT AS BACTERICIDAL AGENTS, CAUSING BACTERIAL CELL DEATH



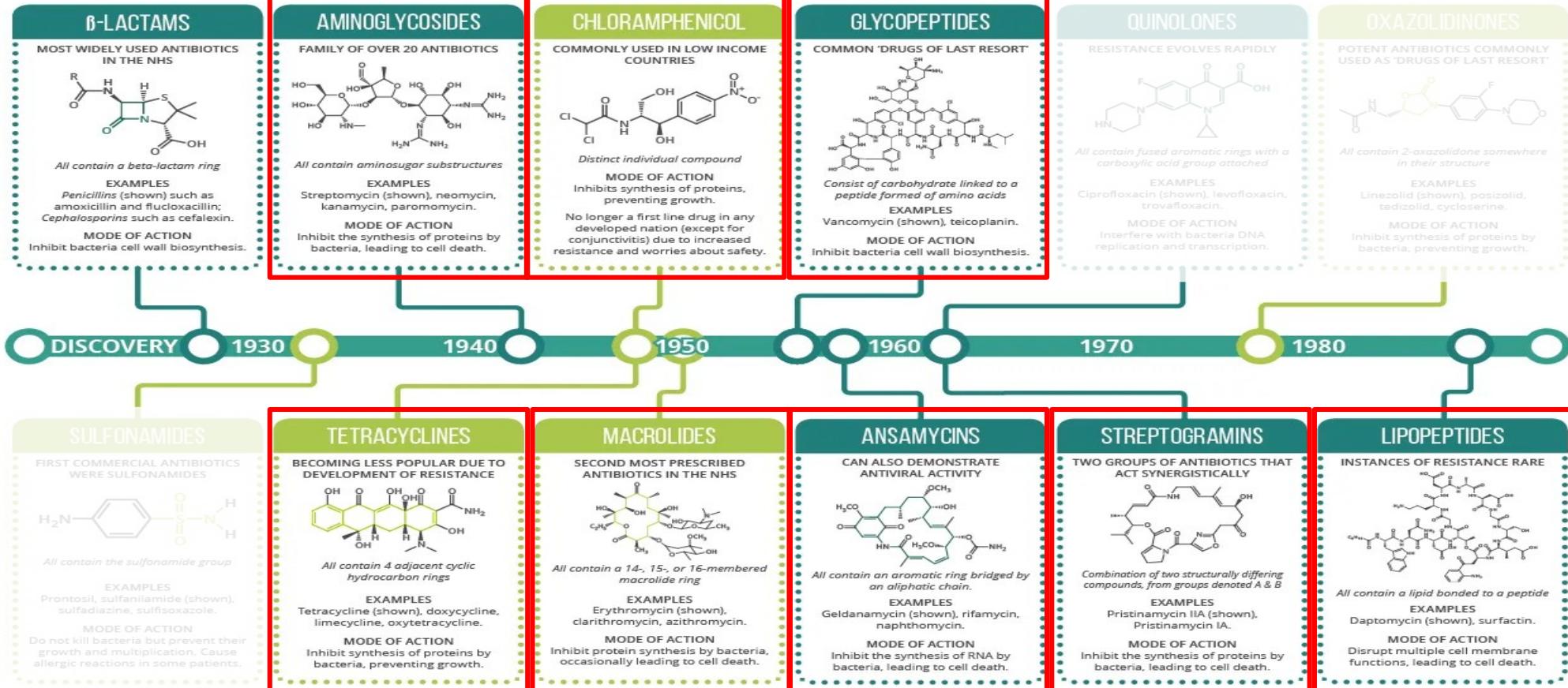
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DIFFERENT CLASSES OF ANTIBIOTICS - AN OVERVIEW

Key: ● COMMONLY ACT AS BACTERIOSTATIC AGENTS, RESTRICTING GROWTH & REPRODUCTION ● COMMONLY ACT AS BACTERICIDAL AGENTS, CAUSING BACTERIAL CELL DEATH





EXPLO(R/IT)ATION OF BIOLOGICAL RESOURCES

The Search for Private Profit in the Nation's Public Parks



Yellowstone National Park has more than 200 continuing experiments by outside scientists.
Anne Sherwood for The New York Times

By **Jim Robbins**

Nov. 28, 2006



YELLOWSTONE NATIONAL PARK, Wyo. — In the 1980s, a researcher at the Cetus Corporation named Kary Mullis was searching for a way to speed the replication of DNA. Enzymes used to amplify the genetic code broke down when they were heated. Dr. Mullis had the idea of using a heat-resistant enzyme called Taq

Taq, isolated from *Thermus aquaticus* in the Yellowstone National Park, made billions of dollars in profit, of which none went to the public park services. It is referred to as “**the great Taq rip-off**”.

Researchers working in National Parks are now required to sign “benefits sharing” agreements that would send a portion of later profits back to the Park Service.

Who owns biological diversity?

"Until the 1970s, biodiversity was considered to be part of the "common heritage of humankind." Under this regime, biological resources are treated as belonging to the public domain and are not owned by any individual, group, or state."

"The last three decades have seen a significant change in the regime governing access to biodiversity. From a common heritage of mankind, biodiversity is evolving into a resource under the sovereignty of nation states and is subject to intellectual property rights (IPRs)."

Illegal Bioprospecting and Biopiracy



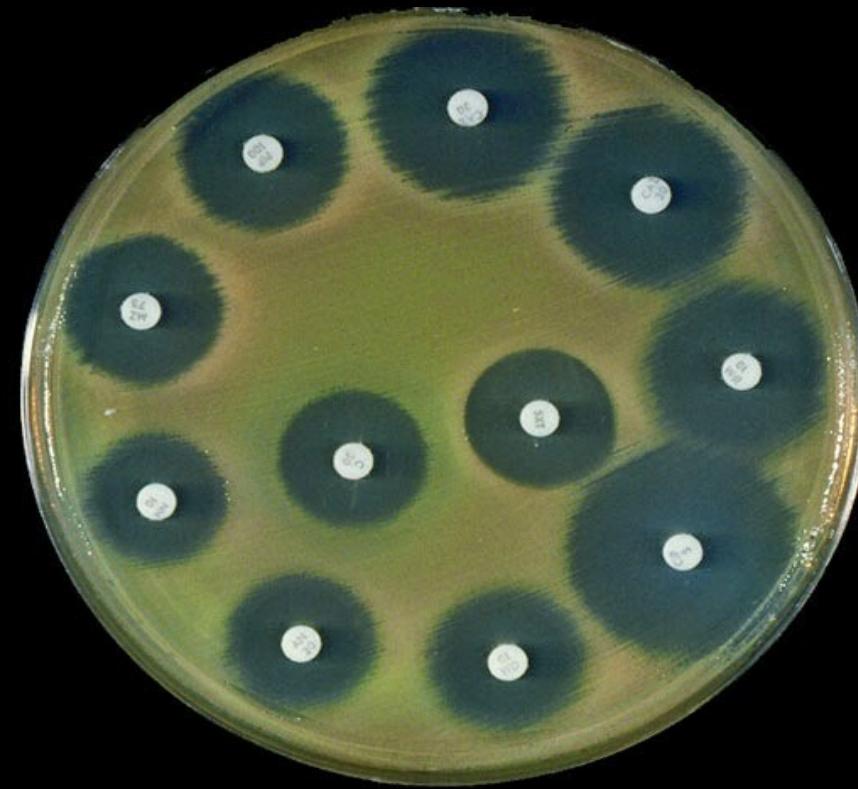
**Convention on
Biological Diversity**

The **Convention on Biological Diversity** (CBD; <http://www.biodiv.org>) was signed in 1992 and came into force in 1993

The **Nagoya Protocol on Access and Benefit-sharing** (Nagoya Protocol, <https://www.cbd.int/abs>) entered into force on 12 October 2014

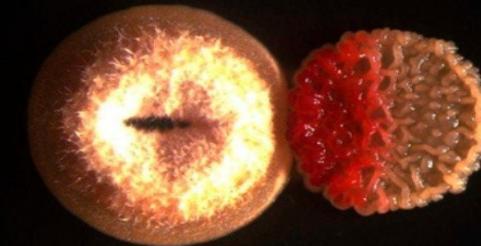


THE HIDDEN ROLE OF SOCIAL INTERACTION IN MICROORGANISMS





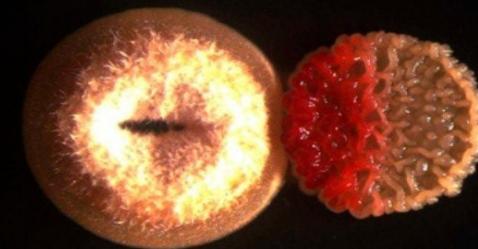
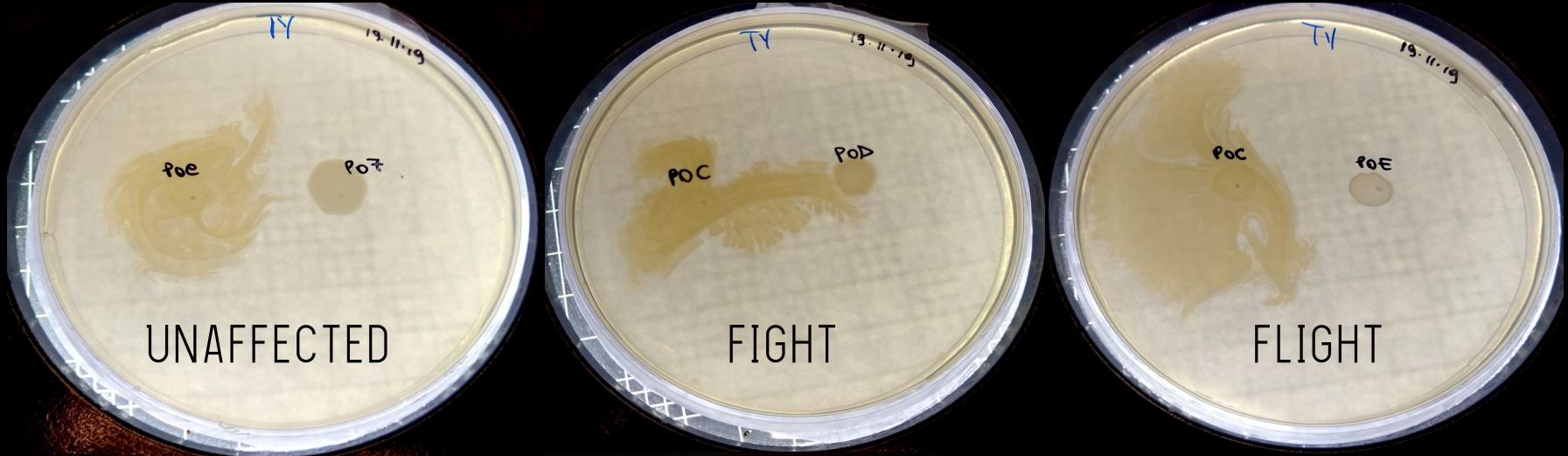
Competitive cocultures to stimulate alternative phenotypes in wild-type microorganisms: fight or flight



Competitive interaction alter the expression of secondary metabolites



Competitive cocultures to stimulate alternative phenotypes in wild-type microorganisms: fight or flight



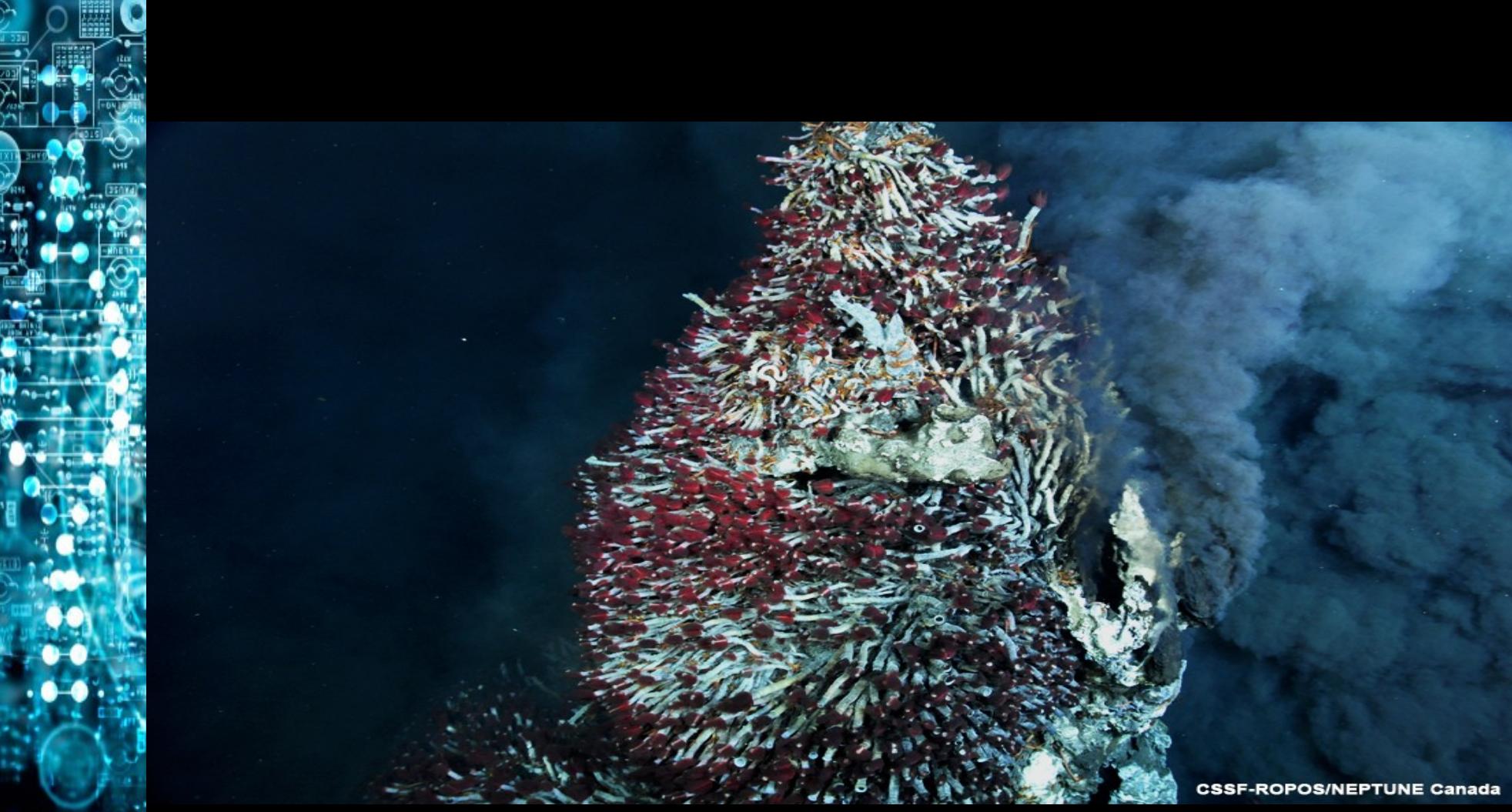
Competitive interaction alter the
expression of secondary metabolites



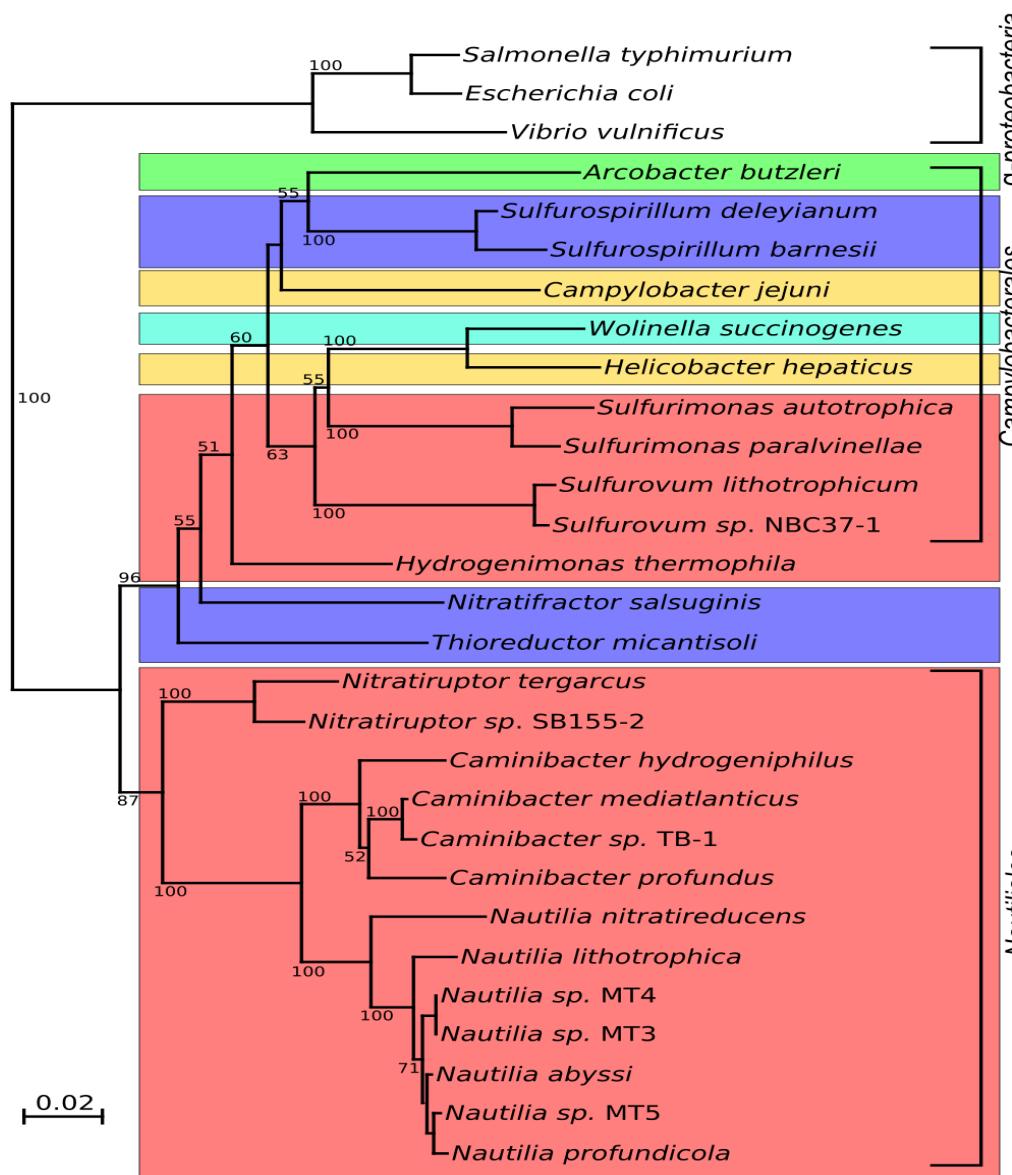
Martina
Cascone



BEYOND THE OBVIOUS: WHERE DO PATHOGENS COME FROM?



CSSF-ROPOS/NEPTUNE Canada



Extremes

From extreme environments to human pathogens: an evolutionary journey

**Donato Giovannelli and
Costantino Vetriani** (Earth-
Life Science Institute, Japan
and Rutgers University, USA)

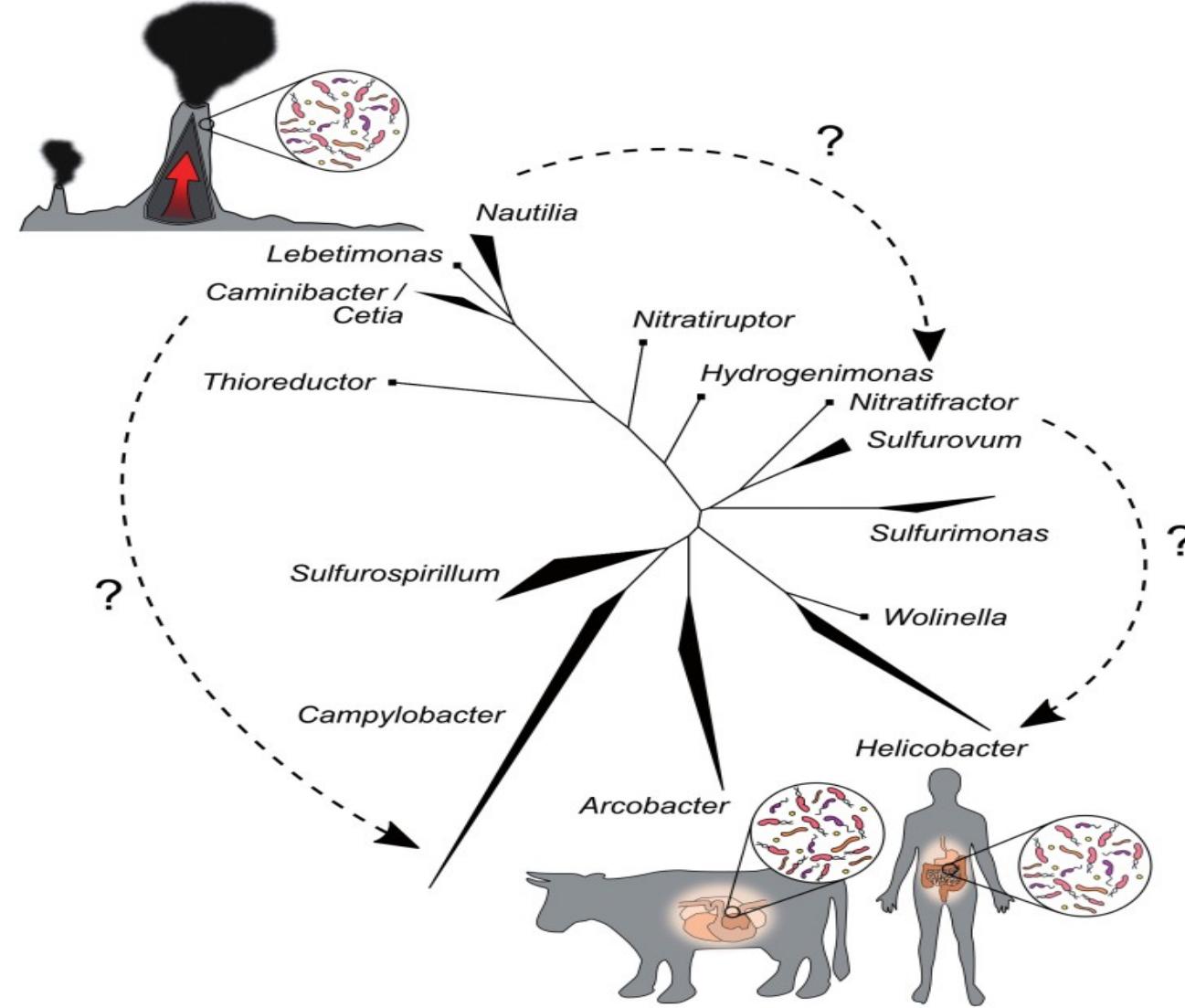
The history of our planet is underpinned by roughly 4 billion years of microbial evolution. From its emergence in a (probably) hot and anoxic environment, microbial life has evolved to colonize every available niche on our planet, including the inside and outside of other organisms. Yet, the emergence and evolution of microbial metabolism remains a major unsolved problem. How have microbes adapted to colonize every available environmental niche, including other organisms? How did they evolve to colonize mammals and our human ancestors? Answers to these questions will allow us to understand the emergence and evolution of life on our planet, inform the search for life elsewhere and, in the making, reveal important insight that will help us fight infections.

A microbial world

There is little doubt today that prokaryotes were the earliest forms of life on our planet. Life as we know it is currently based on energy-conserving reactions that allow the exploitation of naturally occurring redox gradients to perform chemical, mechanical and transport work. Over 30 years of research on the origin of life has led to recognition

overlooked. Only in the last decade have we begun to appreciate how the geosphere and biosphere have co-evolved, ultimately resulting in the complex network of metabolic reactions we see today.

Besides controlling biogeochemical cycles, microbes also affect human and animal health. Microorganisms that comprise both beneficial strains and potential pathogens colonize the exposed areas of our bodies and our gastrointestinal tract. Recent studies have



RESEARCH ARTICLE

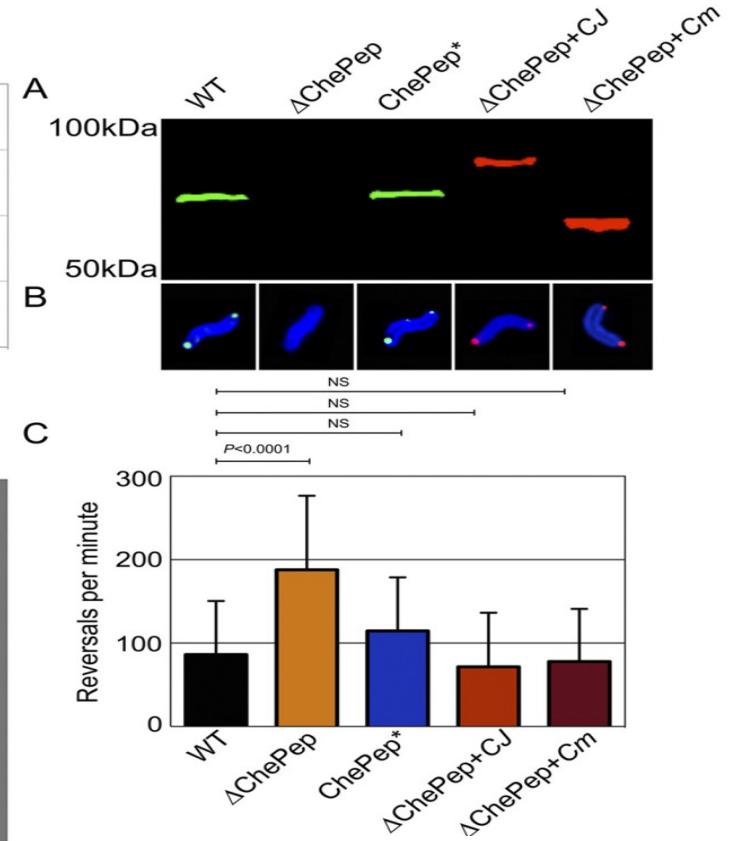
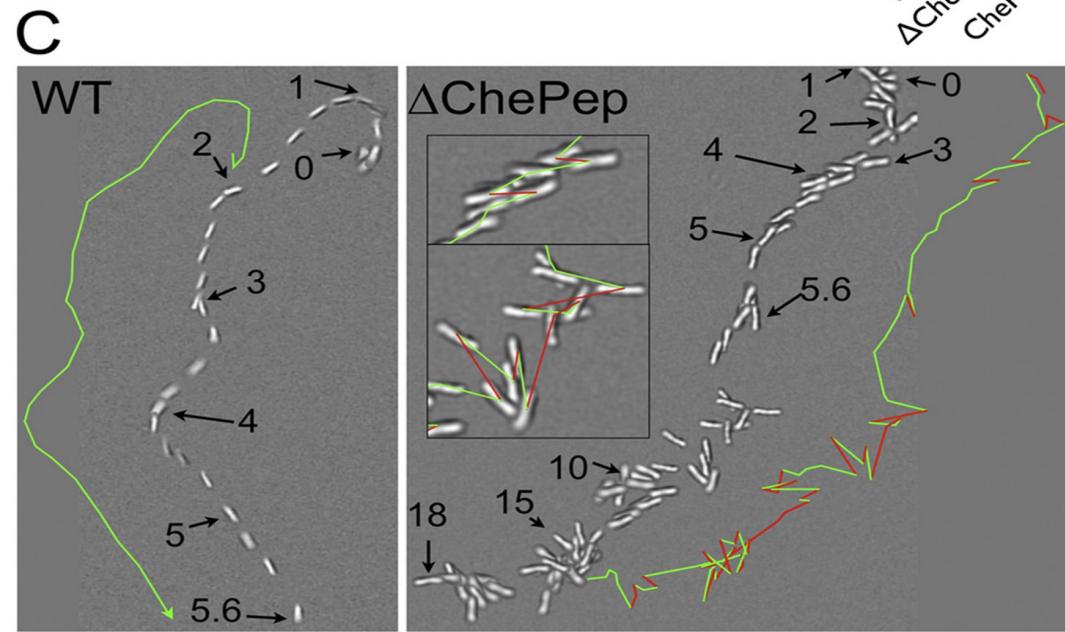
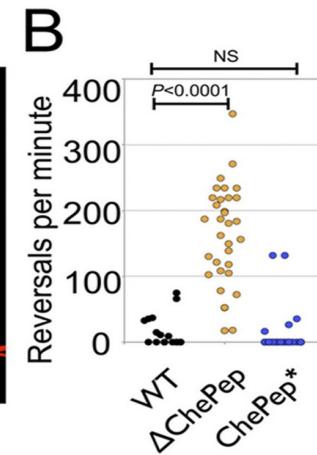
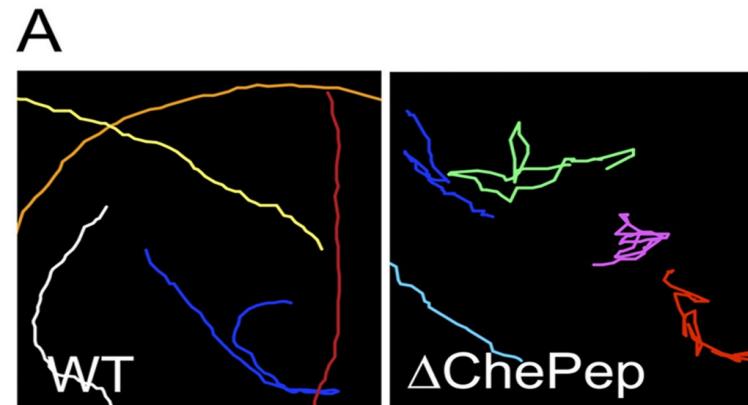
ChePep Controls *Helicobacter pylori* Infection of the Gastric Glands and Chemotaxis in the *Epsilonproteobacteria*

Michael R. Howitt,^a Josephine Y. Lee,^a Paphavee Lertsethtakarn,^b Roger Vogelmann,^{a*} Lydia-Marie Joubert,^c Karen M. Ottemann,^b and Manuel R. Amieva^{a,d}

Department of Microbiology and Immunology, Stanford University School of Medicine, Stanford, California, USA^a; Department of Microbiology and Environmental Toxicology, University of California, Santa Cruz, Santa Cruz, California, USA^b; Cell Sciences Imaging Facility, Stanford University School of Medicine, Stanford, California, USA^c; and Department of Pediatrics, Stanford University School of Medicine, Stanford, California, USA^d

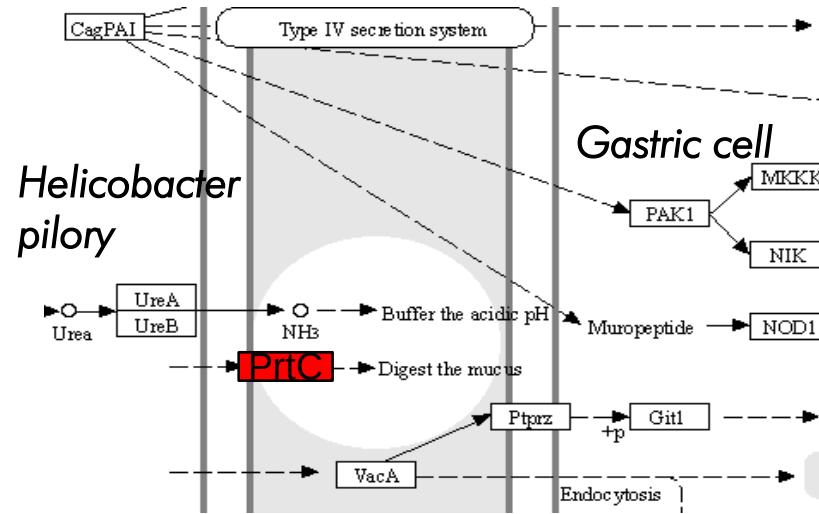
* Present address: II, Medizinische Klinik Universitätsmedizin Mannheim, Mannheim, Germany

ABSTRACT Microbes use directed motility to colonize harsh and dynamic environments. We discovered that *Helicobacter pylori* strains establish bacterial colonies deep in the gastric glands and identified a novel protein, ChePep, necessary to colonize this niche. ChePep is preferentially localized to the flagellar pole. Although mutants lacking ChePep have normal flagellar ultrastructure and are motile, they have a slight defect in swarming ability. By tracking the movement of single bacteria, we found that Δ ChePep mutants cannot control the rotation of their flagella and swim with abnormally frequent reversals. These mutants even sustain bursts of movement backwards with the flagella pulling the bacteria. Genetic analysis of the chemotaxis signaling pathway shows that ChePep regulates flagellar rotation through the chemotaxis system. By examining *H. pylori* within a microscopic pH gradient, we determined that ChePep is critical for regulating chemotactic behavior. The *chePep* gene is unique to the *Epsilonproteobacteria* but is found throughout this diverse group. We expressed ChePep from other members of the *Epsilonproteobacteria*, including the zoonotic pathogen *Campylobacter jejuni* and the deep sea hydrothermal vent inhabitant *Caminibacter mediatlanticus*, in *H. pylori* and found that ChePep is functionally conserved across this class. ChePep represents a new family of chemotaxis regulators unique to the *Epsilonproteobacteria* and illustrates the different strategies that microbes have evolved to



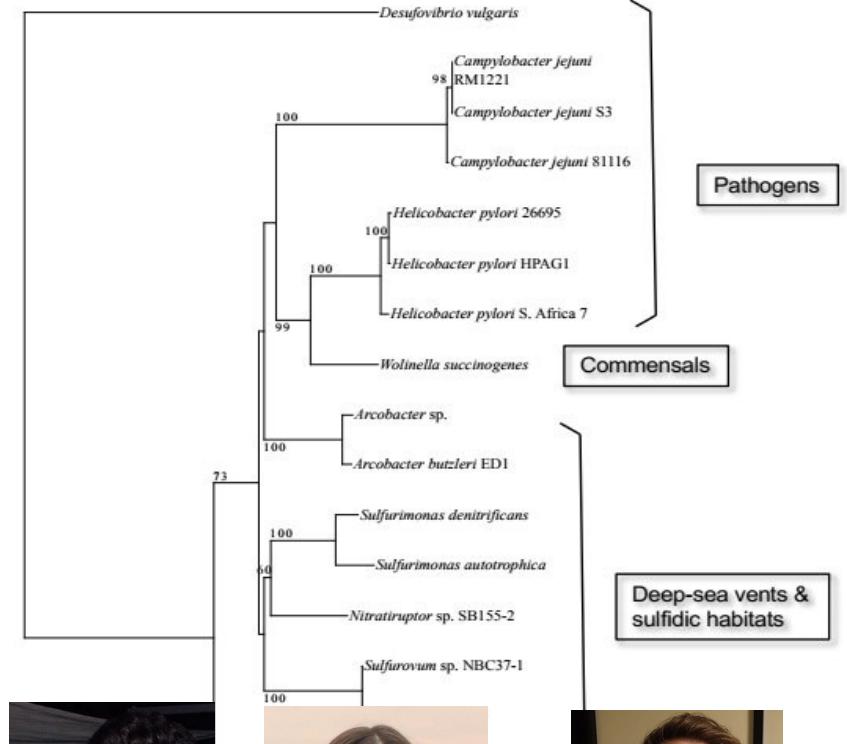
$\text{ChePep}_{H_p}:\text{ChePep}_{C_m}$ 35% aa sim

Conserved function of PrtC in environmental and pathogenic Epsilonproteobacteria



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QC128907.1	RGLTDAYLVKRPYERDDTNQLETSLTHGEWQVSGVVTQEGDSFMCKYKTFPGDVVEILLP
AFV41388.1	RGFTDGYLMMRPFERLDTQNHQTAISEGDFQVNGEITEDGRFFACKFTTTNTAYEIAP
CAE11120.1	RGFTDGYLIRHPRFQLDTQNHETAISEGSFQGVSEVEDGRTFFCKFTIKPSSEEHEIVP
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WP_041354064.1	RGFTDGYLVSRRPYEKSSSTQKLDSSESEGTHQVKAEMEDGLHFTKDNCQGDVLEIVAP
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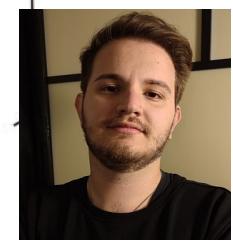
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AFV41388.1	KNAAITPVNEIGKLYTF-EKRSYLVLYKILLENTTELETIHSGNVNLVRPAPLPAFSF
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WP_041354064.1	EGAKLEPCNSEIGSVFEN-ERRWLKLKNKIE-ANKEYECIHSGNVNPVVKLPCPLPAYTF
KYJ86685.1	VGASVELVDNEYGKIDEV-AGQYWLTFKKIVTKEGKELECVHSGNLNALLPAKLPGYTI



Alessandro
Coppola



Angelica
Severino



Alessandro
Masi



WHERE THEM ARCHAEA AT?

Bacteria: medical and environmental relevance

About $>16,000$ bacterial species are known today
($>150,000$ if we include uncultured)

538 are known pathogens or facultative pathogens
($\sim 0.36\%$)

Archaea: any medical relevance?

If we play the number game we should have identified 16-20 Archaeal pathogens

And yet there are no known Archaea pathogens, and the etiologic agents of all major disease are known



PEARLS

Archaea in and on the Human Body: Health Implications and Future Directions

Mor N. Lurie-Weinberger, Uri Gophna*

SPECIAL VIEWPOINTS

The Search for Unrecognized Pathogens

David A. Relman

* See all authors and affiliations

Science 21 May 1999;
Vol. 284, Issue 5418, pp. 1308-1310
DOI: 10.1126/science.284.5418.1308



Infection and Immunity

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Archaea and Their Potential Role in Human Disease

Paul B. Eckburg, Paul W. Lepp, David A. Relman

DOI: 10.1128/IAI.71.2.591-596.2003



Problems and paradigms

Pathogenic archaea: do they exist?

Ricardo Cavicchioli, Paul M.G. Curmi, Neil Saunders, Torsten Thom

First published: 17 October 2003 | <https://doi.org/10.1002/bies.10354> | Citations: 50

Hypotheses

The proportional lack of archaeal pathogens: Do viruses/phages hold the key?

Erin E. Gill and Fiona S. L. Brinkman*

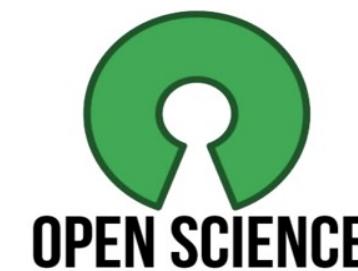


THE ROLE OF OPEN SCIENCE

Open Science: what is it?

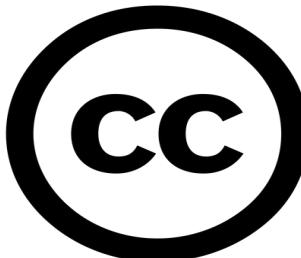
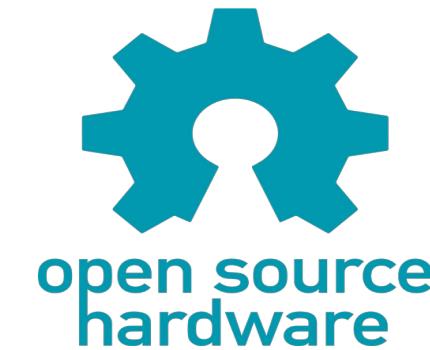
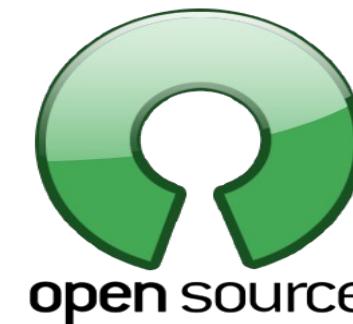
Open science is the movement to make scientific research (including publications, data, physical samples, and software) and its dissemination accessible to all levels of an inquiring society, amateur or professional.

Open science is transparent and accessible knowledge that is shared and developed through collaborative networks.



Open Science: Why?

OpenScience is part of a large movement, started with Open Source Software many years ago.



see <https://www.gnu.org/philosophy/free-sw.html> for more info



Reproducible

Raphael Ritson-Williams

FAST





IMMORTAL



The power of sharing

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Great power comes from comparison!

Biologist as Early Innovators in Big Data Resources

'70s-'80s

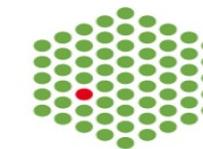


'90s



National
Center for
Biotechnology
Information

EMBL



2000s



MG-RAST
metagenomics analysis server



AddGene

Addgene is a global, nonprofit repository that was created to help scientists share plasmids. Plasmids are DNA-based research reagents commonly used in the life sciences. When scientists publish research papers, they deposit their associated plasmids at Addgene.

It played an important role in the spread of CRISPR-cas technology allowing quick access to reagents.



Sharing Speeds Science



If it takes **5** hours to make a plasmid...
Addgene has saved researchers
5,000,000 hours.

This is the same as **570** years saved!



Deposits and Requests



We've now shipped
1,000,000+
plasmids.



Our repository
contains **67,000+**
plasmids available
from **4,000+**
depositing labs.

Fun Addgene Facts



Since we were
founded in 2004,
Addgenies have
formed Boston
Triathlon teams
for **9** years.



We've also gone
on **317** outreach
trips to reach
scientists around
the world.



OpenSourceMalaria

The Open Source Malaria project is an open science initiative designed to accelerate the pre-phase one testing of anti-malaria compounds by leveraging the open science principle in 2011.

All the experiments go on the web (including the ones that did not turn out well). All the data are available. Anyone can do anything they wish with the compounds (CC-BY licence). Anyone can take part in the research.

The screenshot shows the main homepage of the Open Source Malaria project. It features the OSM logo at the top left. Below it is the title "OPEN SOURCE MALARIA" in large red letters, followed by the subtitle "Looking for New Medicines". To the right of the title is a section titled "Catch us on the Daily Show?", which includes a link to "Click here!". Below this is a paragraph about the project's approach to curing malaria, followed by a "Read More" link. At the bottom of this section is a "How would you Improve GitHub?" form. On the left side, there are links to "Lab Notebook", "Project Wiki", and "Molecule Database". A "Recent Contributors" chart shows activity levels for users matodd, Mferrinflower, drc007, and holeung. At the bottom, there are buttons for "Activity", "Join the Team", and "Meet Us".

The screenshot shows the "Story so far" page from the OpenSourceMalaria wiki. The page has a header with the OSM logo and the title "OpenSourceMalaria:Story so far". Below the header is a navigation bar with links for "Malaria Home", "OSM So Far", "Compound Series", and "Links". The main content area contains a summary of the project's history and a table of contents for the story. The table of contents includes sections such as "The story so far in the open source malaria project", "History", "Synthesis and Evaluation Completed to Date", "What Are the Compounds Doing?", "Obtaining Other Compounds", "Related Series", "Comments", and "Licence".



Readings

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