



TECHNISCHE UNIVERSITÄT  
BERGAKADEMIE FREIBERG  
Die Ressourcenuniversität. Seit 1765.

# Bioremediation and phytoremediation 2021/22

## Methods in Biohydrometallurgy



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# Sampling

- Taking samples as sterile as possible  
(sterile equipment, gloves, steril bottles,...)



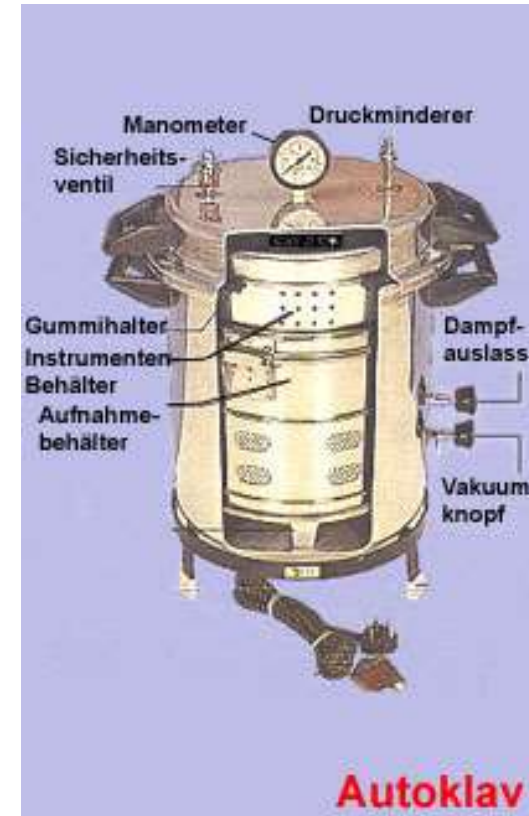
## Killing of microorganisms (sterilisation)

Basically, the following methods are used:

- moist heat
- dry heat
- filtration
- radiation
- chemicals

# 1. Moist heat and pressure

- Bacteria and fungi are killed at 60°C within 5-10 minutes
- Yeast and fungal spores only above 120 ° C (15 min)
- Clinics are recommended for 5 minutes at 134 ° C to kill even heat-resistant spores (*Bacillus stearothermophilus*)
- To reach temperatures above the boiling point of water, so-called autoclaves are used. These are devices similar to a steam pressure pot in the kitchen.





## 2. Dry heat

Dry heat spores are killed only at higher temperatures. Therefore, sterilization at least 60 minutes at 180°C or 120 minutes at 160°C. Only then proteins denature irreversibly.

Bacteria	Incubation time 150°C	Incubation time 180°C
Bacillus anthracis	60-120 min	3 min
Clostridium botulinum	25 min	5-10 min
Clostridium tetani	30 min	1 min
Soil bacteria	180 min	15 min



# Pasteurization

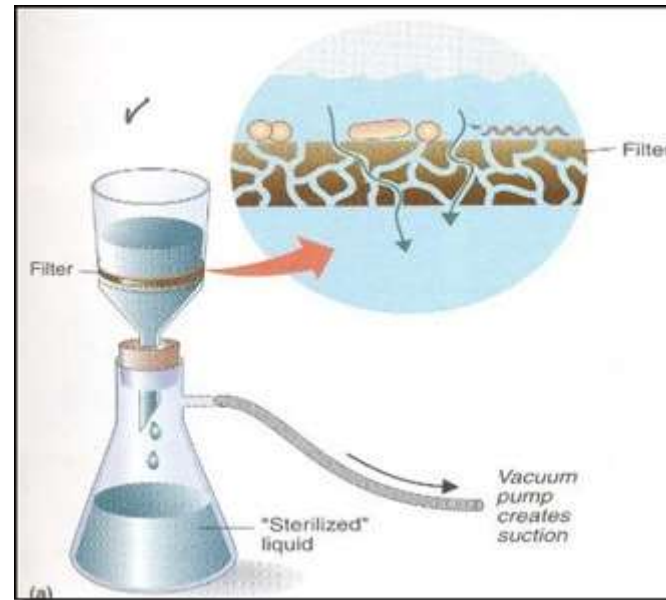
Partial germination of heat-sensitive foods is e.g. used with milk, fruit juice, sauerkraut, meat and fish salads ....



Louis Pasteur (1822-1895)

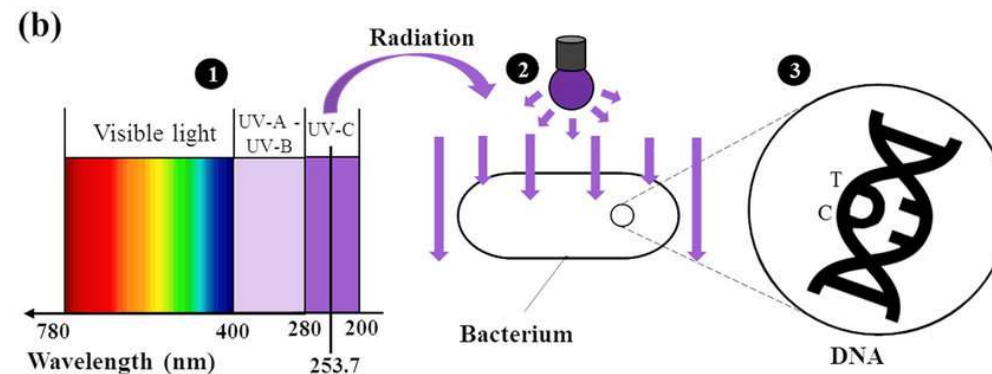
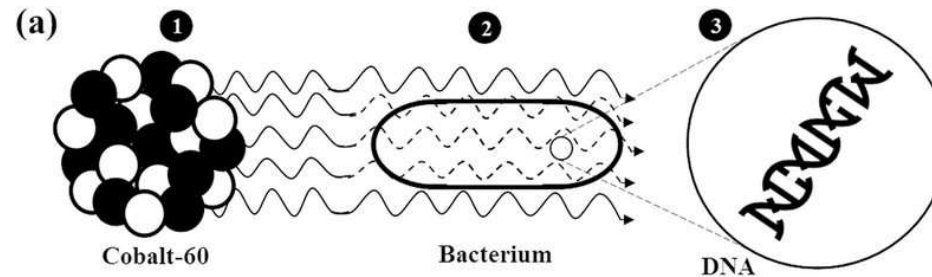
### 3. Filtration

- Heat-sensitive solutions
- sterilization by microfilter, e.g. in the drinking water disinfection filters made of diatomaceous earth or plastic with pore sizes of 0.22 to 0.45  $\mu\text{m}$
- However, bacterial toxins can not be removed that way



## 4. Radiation

- UV-, X-ray and gamma radiation are used



- generation of mutations and destruction of DNA
- UV light at 260 nm → absorbed by the DNA → death cells
- X-rays  $10^{-8}$  -  $10^{-10}$  m, gamma radiation  $10^{-10}$  -  $10^{-14}$  m (Co 60 or Cs 137)

→ ionizing effect on microbial structures, hydrogen bonds in proteins are destroyed and bases in the nucleic acids changed



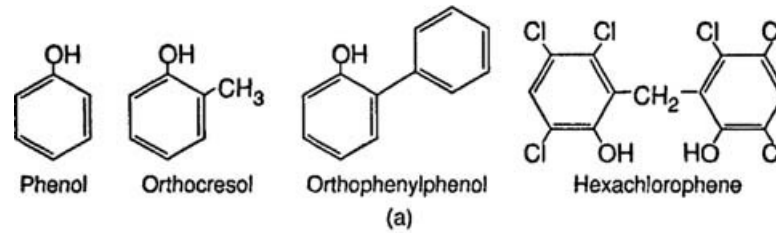


## 5. Chemicals

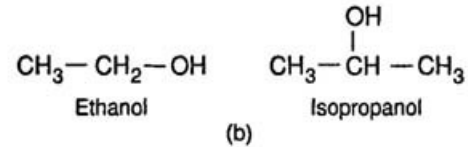
- Thermosensitive devices or areas are sterilized with chemicals such as ethylene oxide (ethene oxide), ozone or formaldehyde vapor
- Ethylene oxide only works in the presence of water (5-10%)
- Ozone is e.g. used for sterilizing plants in poultry farming (Listeria) or improving sanitation in oil platforms and disinfecting water in the fish processing industry and swimming pools

# Disinfectants

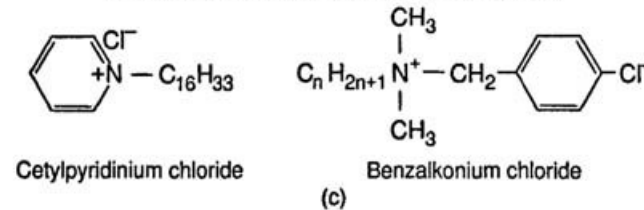
## Phenolics



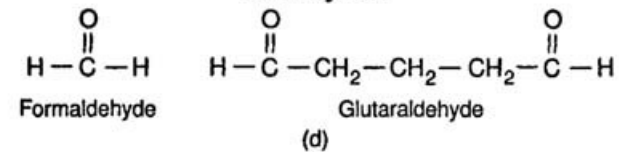
## Alcohols



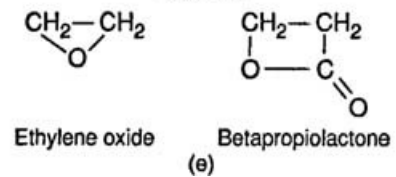
## Quaternary ammonium compounds



## Aldehydes



## Gases





# Sampling

- Taking samples as sterile as possible (sterile equipment, gloves, steril bottles,...)
- replicates
- diverse ecosystems → choice of sampling locations
- Measuring on site parameters

# Field geochemical analysis

1. Composite probes,  
e.g.<sub>1</sub> YSI multiprobe: pH/ORP/T/Ec/DO



e.g.<sub>2</sub> Hanna water tester: pH/ORP/T/Ec



2. Single measurement meters



# Field geochemical analysis

## 3. Indicator strips/paper

- narrow range pH papers



**NARROW RANGE pH INDICATOR PAPERS :**

- pH 2.0 - 4.5
- pH 3.4 - 6.0
- pH 3.8 - 5.3
- pH 5.0 - 7.5
- pH 6.5 - 9.0
- pH 8.0 - 10.5

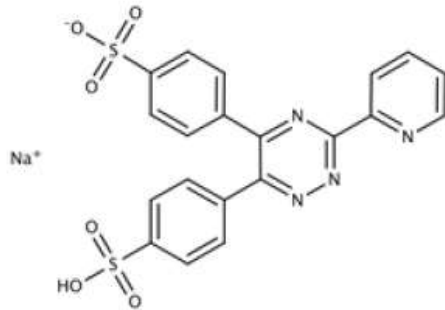
pH	pH	pH	pH	pH	pH
2.0	3.0	4.0	5.0	6.0	7.0
2.5	3.5	4.5	5.5	6.5	7.5
3.0	4.0	5.0	6.0	7.0	8.0
3.5	4.5	5.5	6.5	7.5	8.5
4.0	5.0	6.0	7.0	8.0	9.0
4.5	5.5	6.5	7.5	8.5	9.5
5.0	6.0	7.0	8.0	9.0	10.0
5.5	6.5	7.5	8.5	9.5	10.5

- Merkoquant test strips



# Field geochemical analysis

Stookey, L. (1970) *Ferrozine –a new spectrophotometric reagent for iron. Analytical Chemistry* 42:779-781



950 µL of Ferrozine reagent

+50 µL test solution  
(may require dilution;  
max 1 mM (56 mg/L) Fe<sup>2+</sup>)

The coloured complex is stable for at least 24 h.  
Samples can be taken back to the lab for spectro-photometric analysis



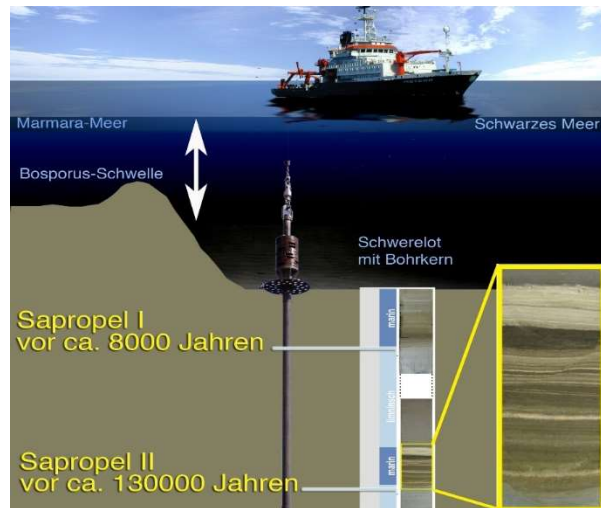
# Sampling

- Taking samples as sterile as possible (sterile equipment, gloves, steril bottles,...) replicates
- diverse ecosystems → choice of sampling locations
- Measuring on site parameters
  
- Cooling and oxygen supply of samples
- Fast sample transport
- Storage of samples depends on analyses (cold, frozen, room temperature)



# Sampling - Terrestrial sites

Terrestrial samples: drill cores, flushing, tracers,...





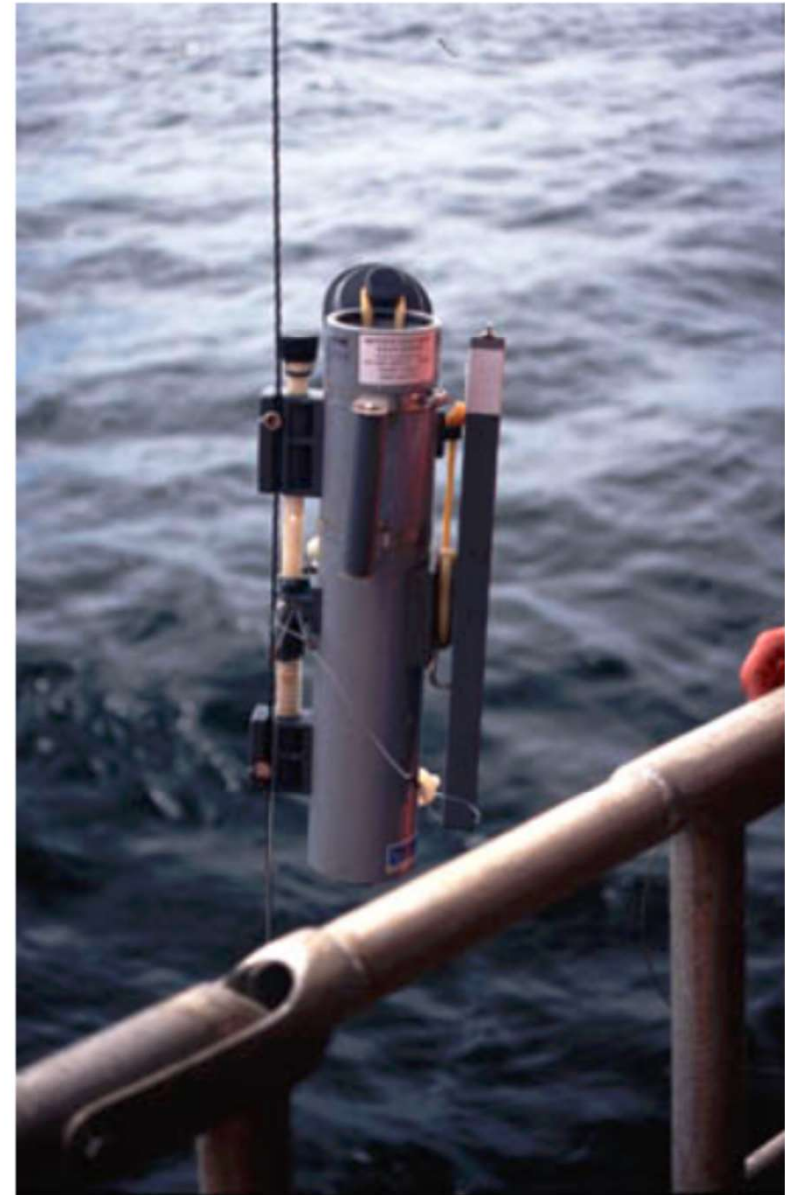


# Water samples



# Water sampling

- Niskin Bottle
  - used for greater depths.
  - lids open or 'cocked' as the instrument is put into the seawater
  - At the desired depth the lids are triggered to close
  - Bottles are sealed airtight to avoid contamination
  - Water analysis conducted on board





## Sample transport/ storage

(i) Solids



→ sealed polythene bags

(ii) Liquids → sterile containers (for enumeration/isolation): keep COOL (not frozen!)

→ filter through 0.2  $\mu\text{m}$  (pore size) membranes

→ biomass (for DNA; can be preserved by freezing)

→ sterile water sample (for lab analyses; DOC sulfate etc))



→ add small amount of conc.  $\text{HNO}_3$  to some sterile water sample to maintain metals in solution for lab analyses



# Microbial characterization methods

## ▶ *In situ* methods

- light and electron microscopy

## ▶ Cultivation-dependent methods

- growth on nutrient medium
- conditions often unknown
- only 1% of all organisms can be cultivated

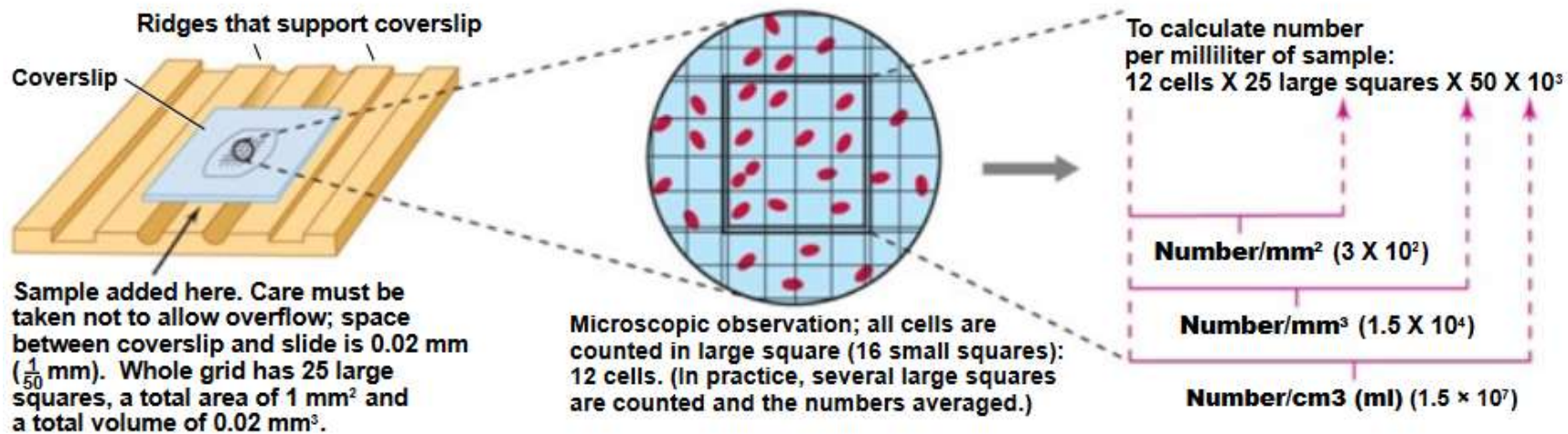
## ▶ Culture-independent methods

- extraction of nucleic acids, proteins, etc.



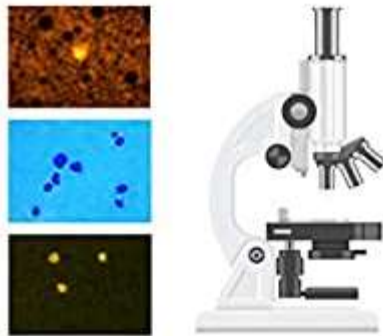
# Total cell counts using microscopy

- Microbial cells are enumerated by microscopic observations (Figure 5.15)



## Direct cell counts under the microscope

- Highest expected cell number
- Minimal equipment
- Quick and easy
  
- No discrimination between dead and living cells
- Background interference
- Pretreatment to remove e.g. iron minerals
- Suitable for protozoa, eukaryots.... Difficult for bacteria



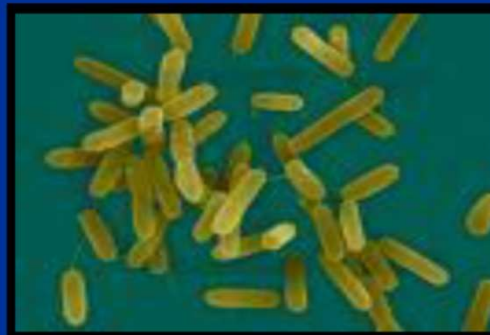


## Why is it so difficult to distinguish bacteria by microscopy?

- Exist on earth since many years (much longer than higher organisms) → higher diversity
- Limited morphology (cocci, rods, spirilli, vibri)



*Escherichia coli*



*Pseudomonas aeruginosa*





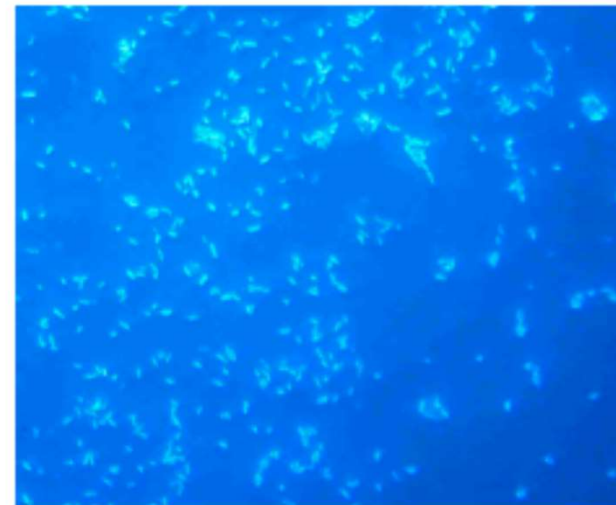
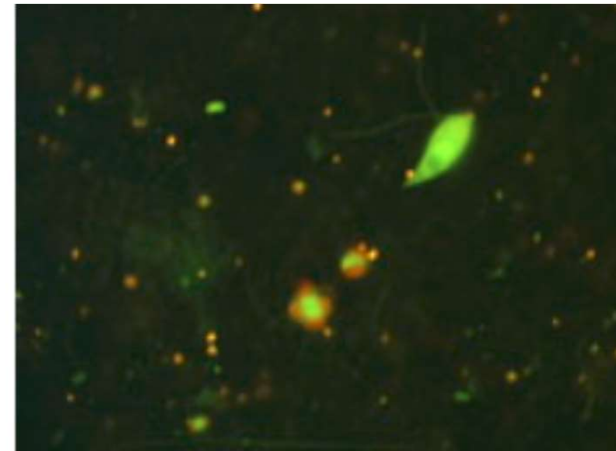
# Fluorescence microscopy

**Direct counts:**

**Fluorescent dyes &  
epifluorescence microscopy**

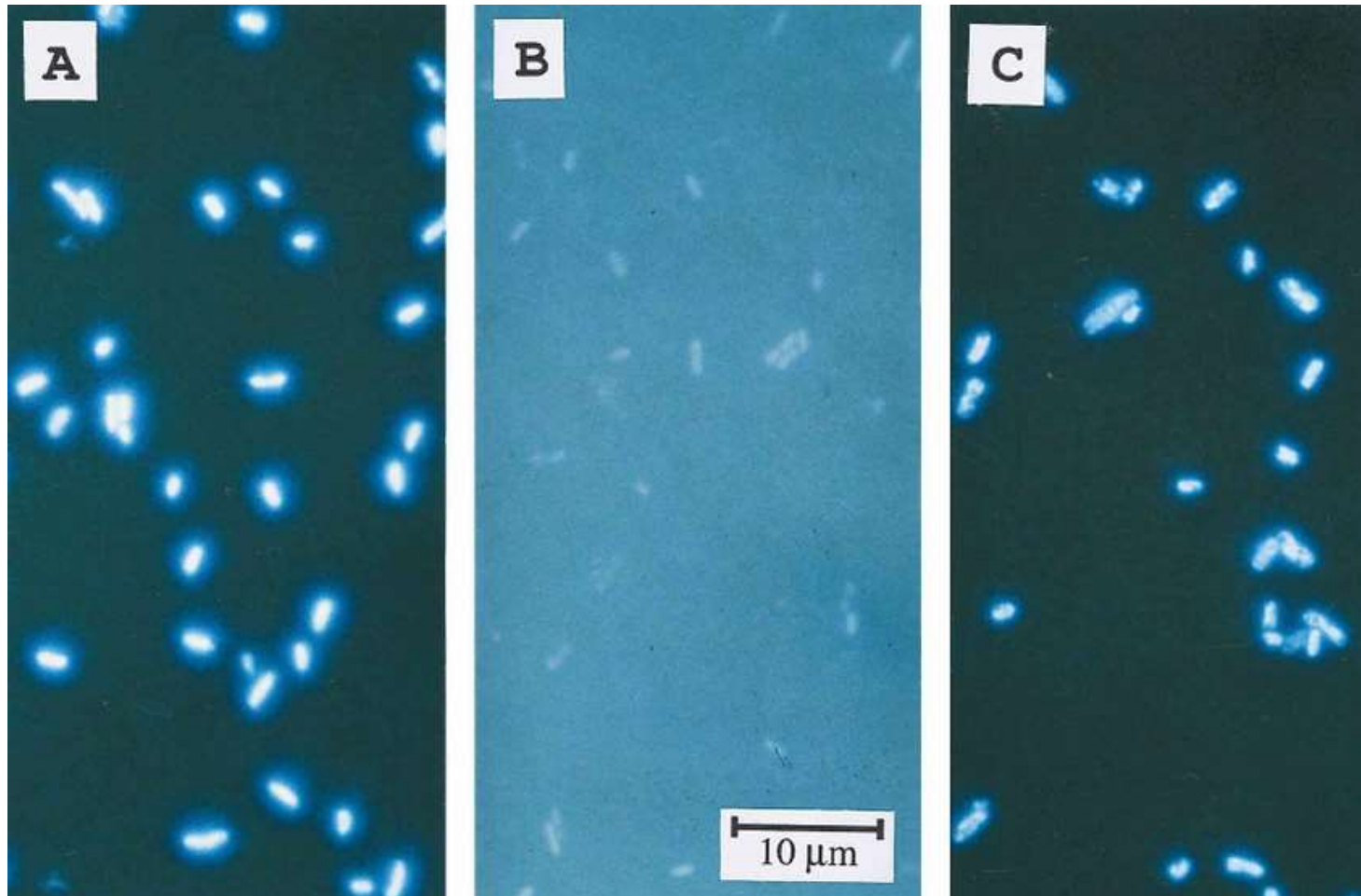
-Acridine Orange Direct  
Counts (AODC) general  
nucleic acid stain,  
fluoresces green or red

-DAPI – 4,6-diamidino-2-  
phenylindole AT-specific  
DNA stain, fluoresces  
blue or white





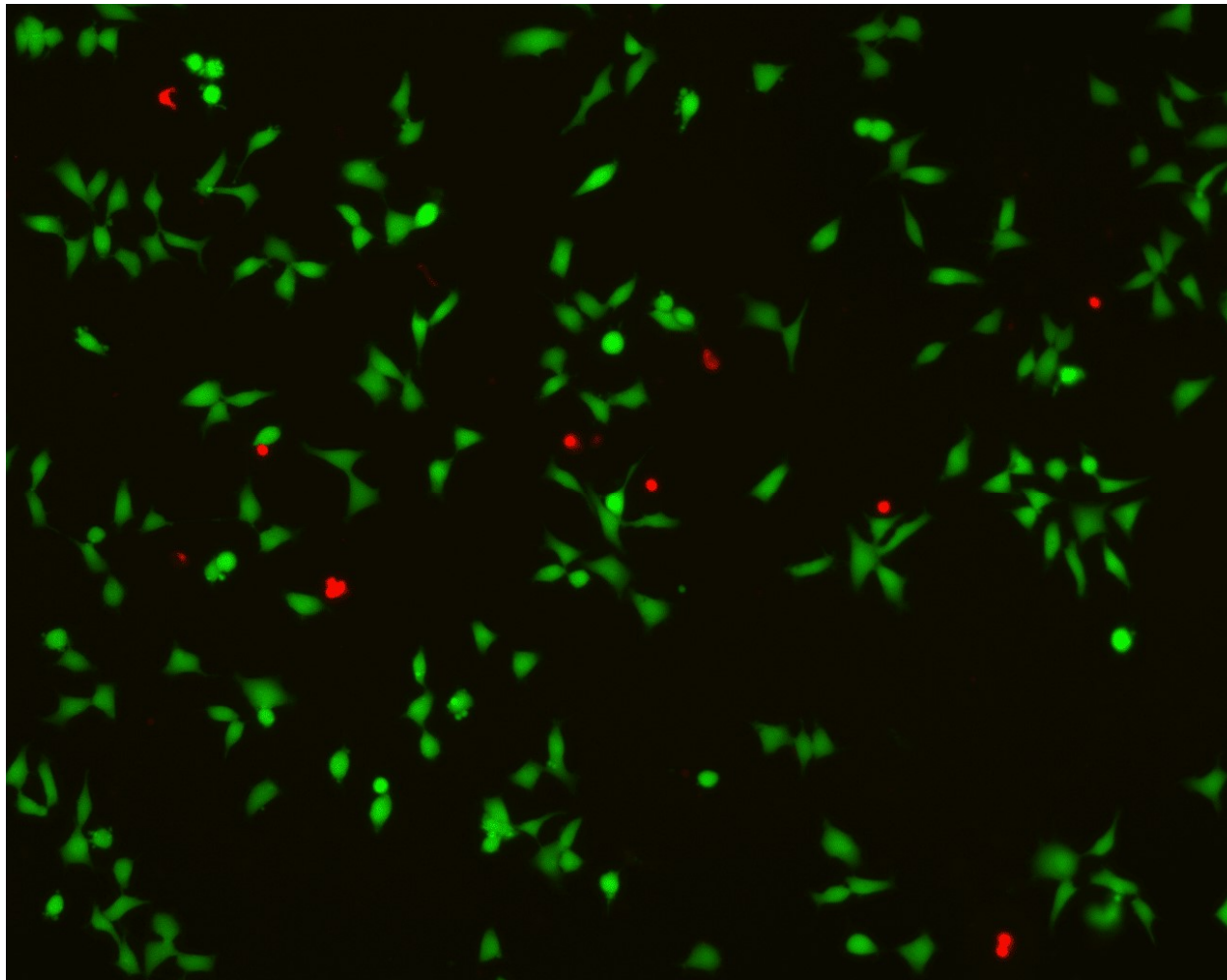
## Fluorescence microscopy



Cells of *E. coli* stained with DNA stain (DAPI) to determine total cell number



## Live/dead staining



Viability staining to distinguish living (green) and dead (red) cells using the LIVE/DEAD Bac Light™ kit



# Microcalorimetry

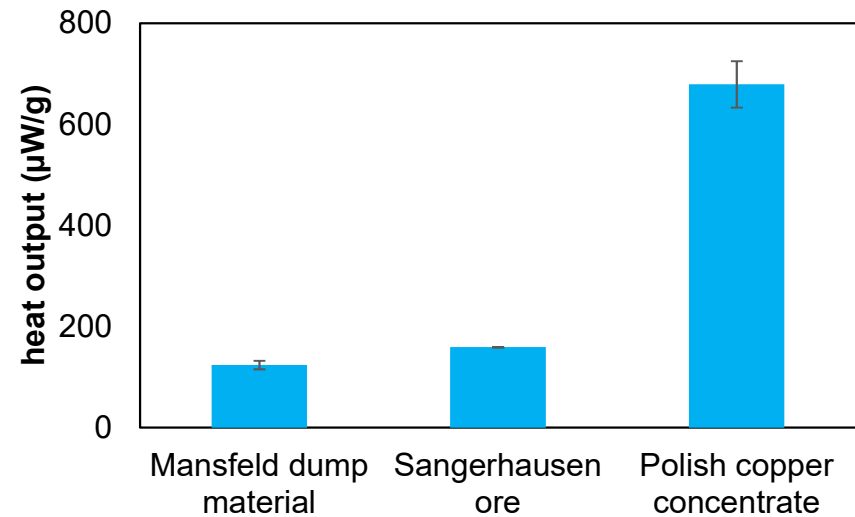
- performance of active culture on substrate (ore)
- determination of cell activity by heat release

## 48 Channel Microcalorimeter



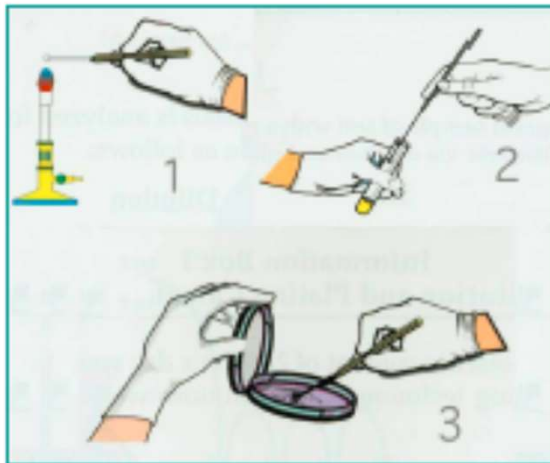
Specimen ampoule goes here:

Heat flow measured relative to a reference:



## Cultivation-dependent techniques

- Enumeration and isolation using selective media
- Target specific organisms
- Gives indication of activity
- Gives model organisms for laboratory studies
- But will underestimate diversity



Isolation techniques



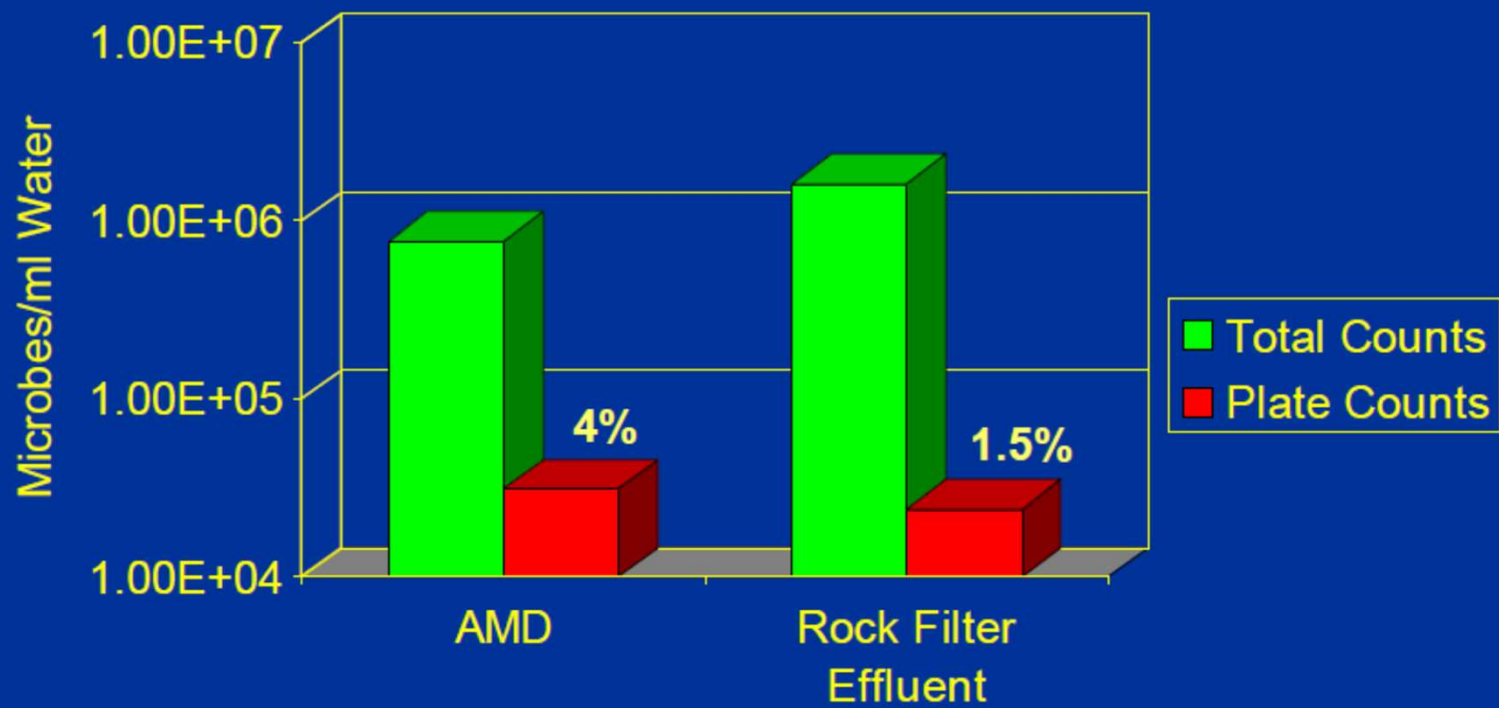
Agar plate + colonies



Anaerobic cabinet



## Total microbes vs cultivatable microbes at Wheal Jane



## Selection by variation of growth medium:

- salts
- pH- value
- C-source

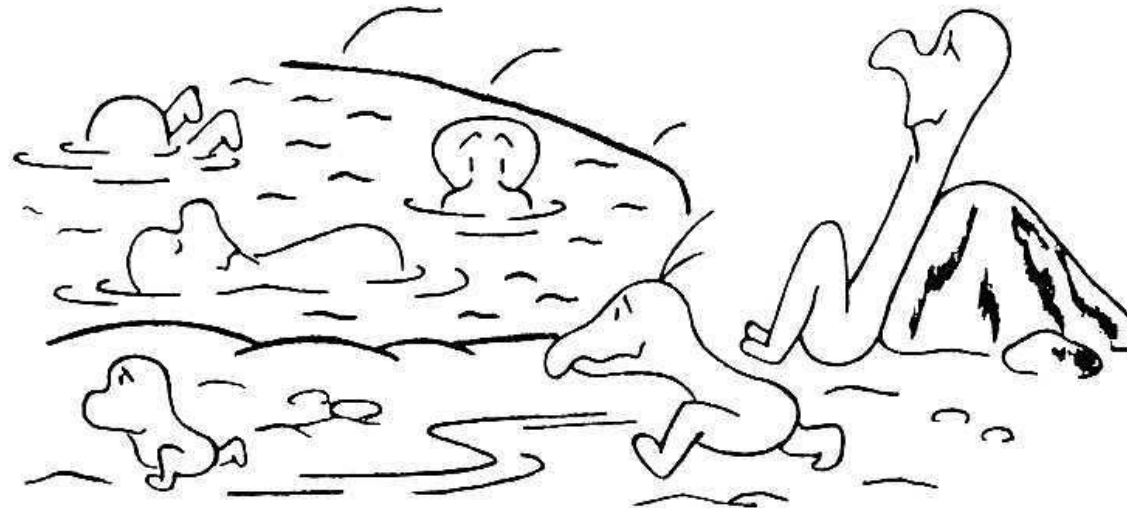
Enrichment culture



*Die Anzucht unter speziellen Laborbedingungen wird nur von wenigen überstanden.*

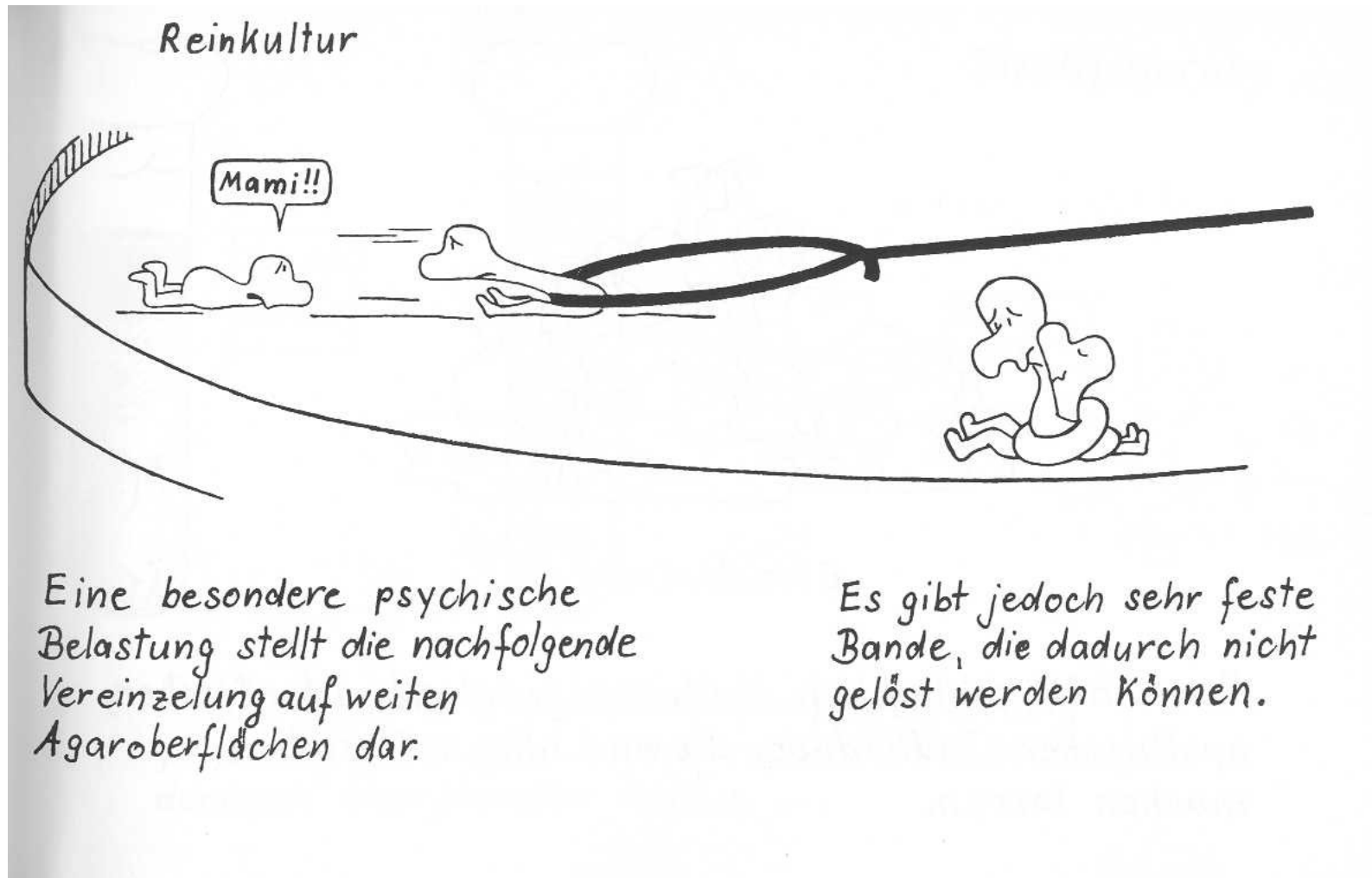


## Mixed culture



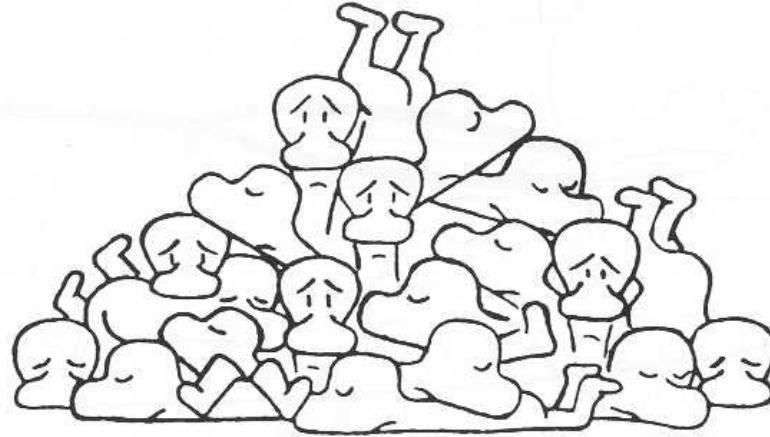
*leben die Bakterien in  
Mischkultur...*

## Isolation using agar plates





## Pure culture

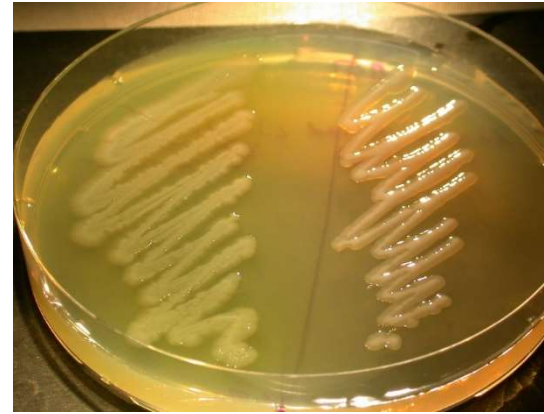


Einzelkolonie

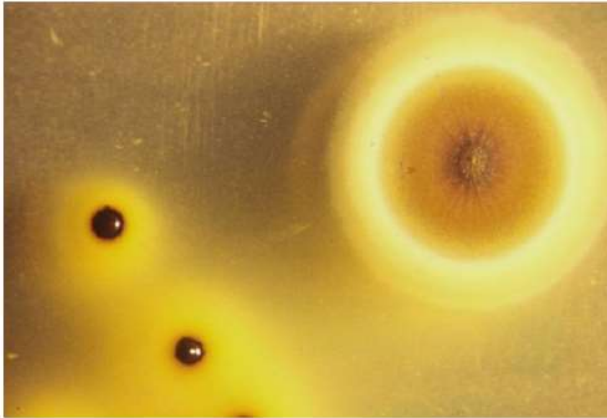
Man erhält schließlich Kulturen genetisch identischer, apathischer Individuen, die nun alles mit sich machen lassen.



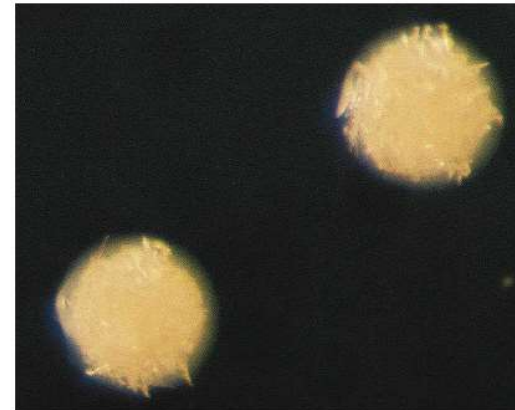
# Isolation of pure cultures on agar plates



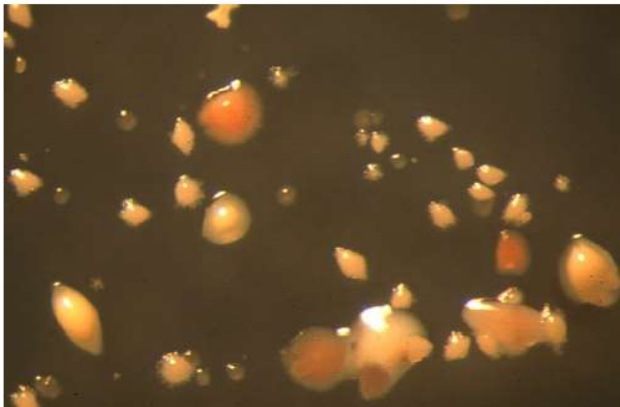
# Single colonies on agar plates



Colonies of Fe-oxidising  
bacteria on iFe<sub>0</sub> medium



Colonies of S-oxidising bacteria  
on FeS<sub>0</sub> medium

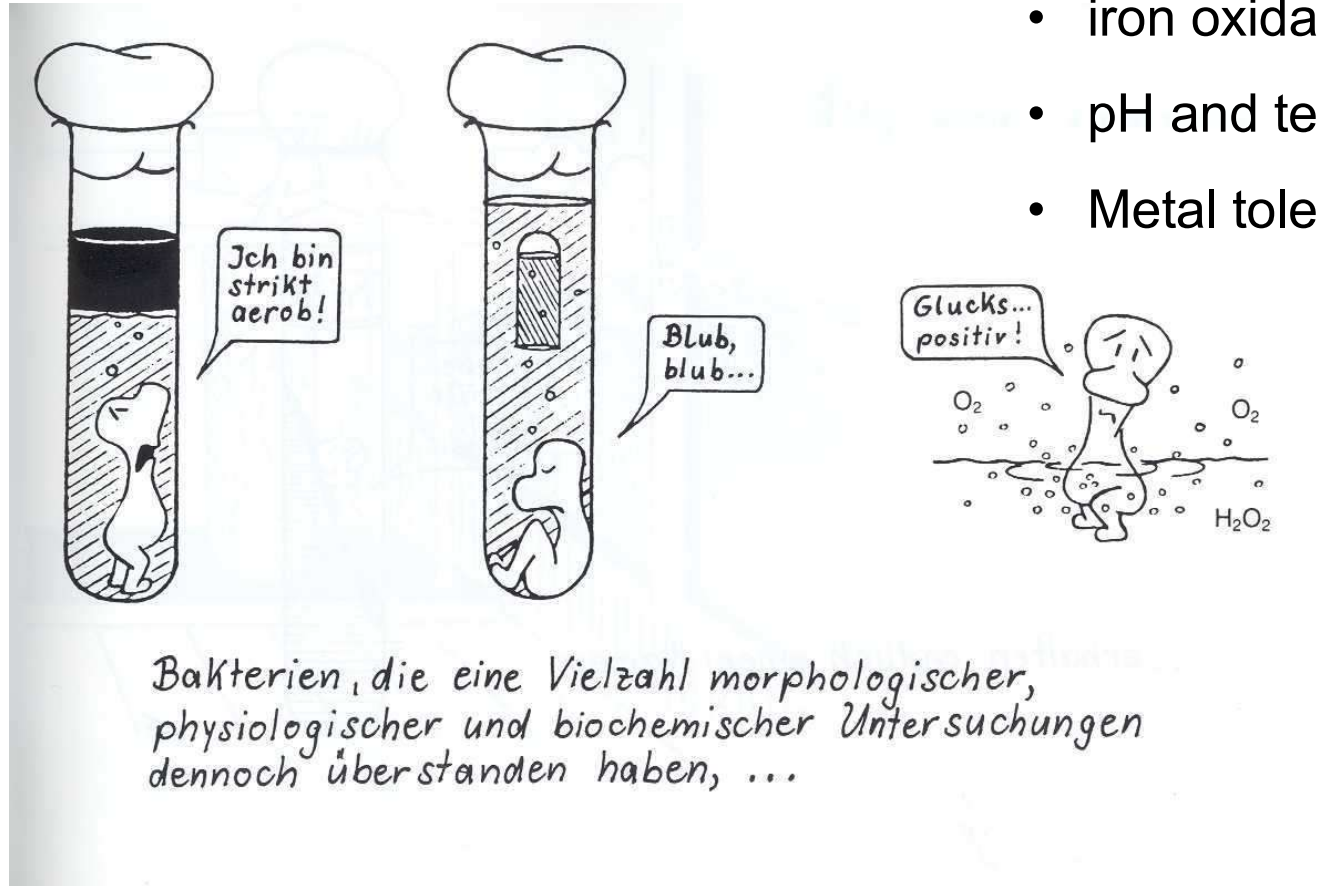


Colonies of heterotrophic  
acidophiles on YE3<sub>0</sub> medium

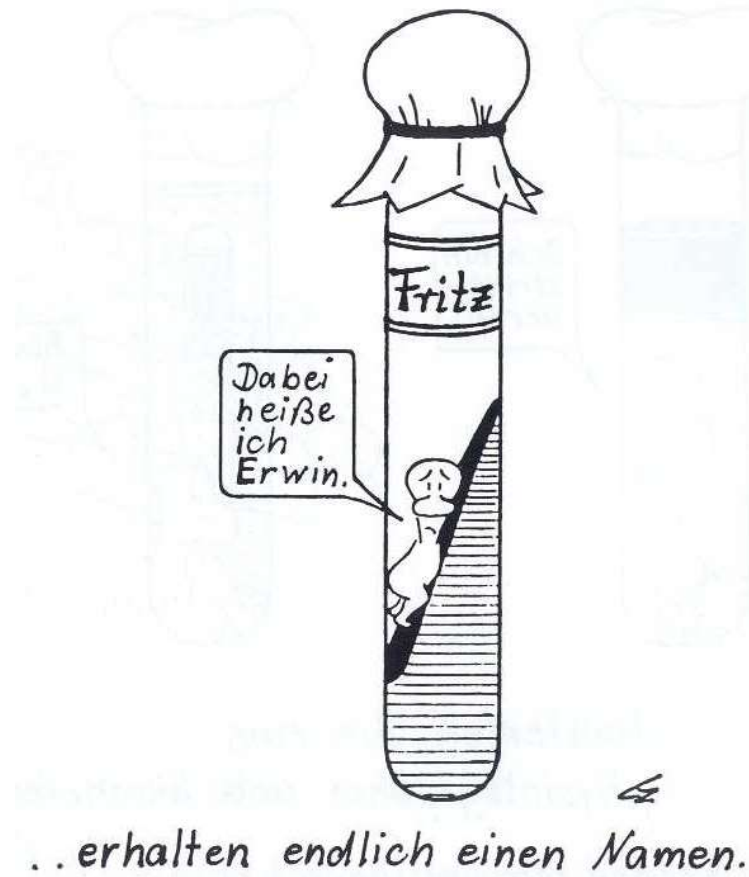
# Naming a novel microorganisms

Test physiological properties:

- iron oxidation
- pH and temperature limits
- Metal tolerance



# Naming a novel microorganisms

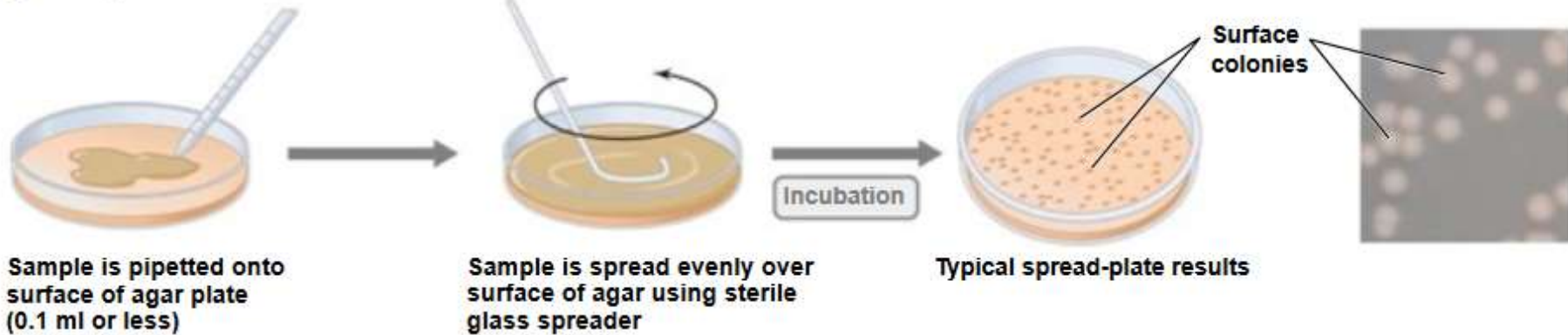




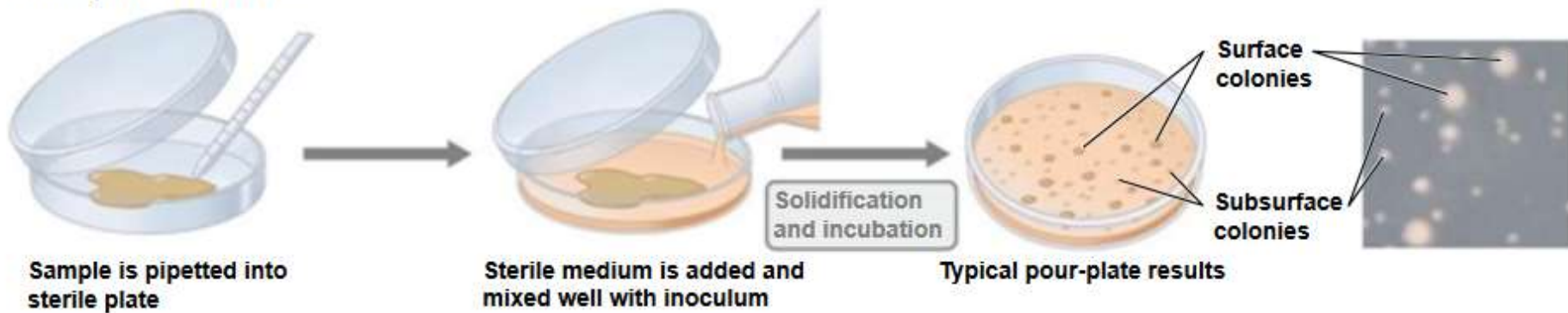
# Viable cell counts (plate counts)

→ Measurement of living, reproducing population

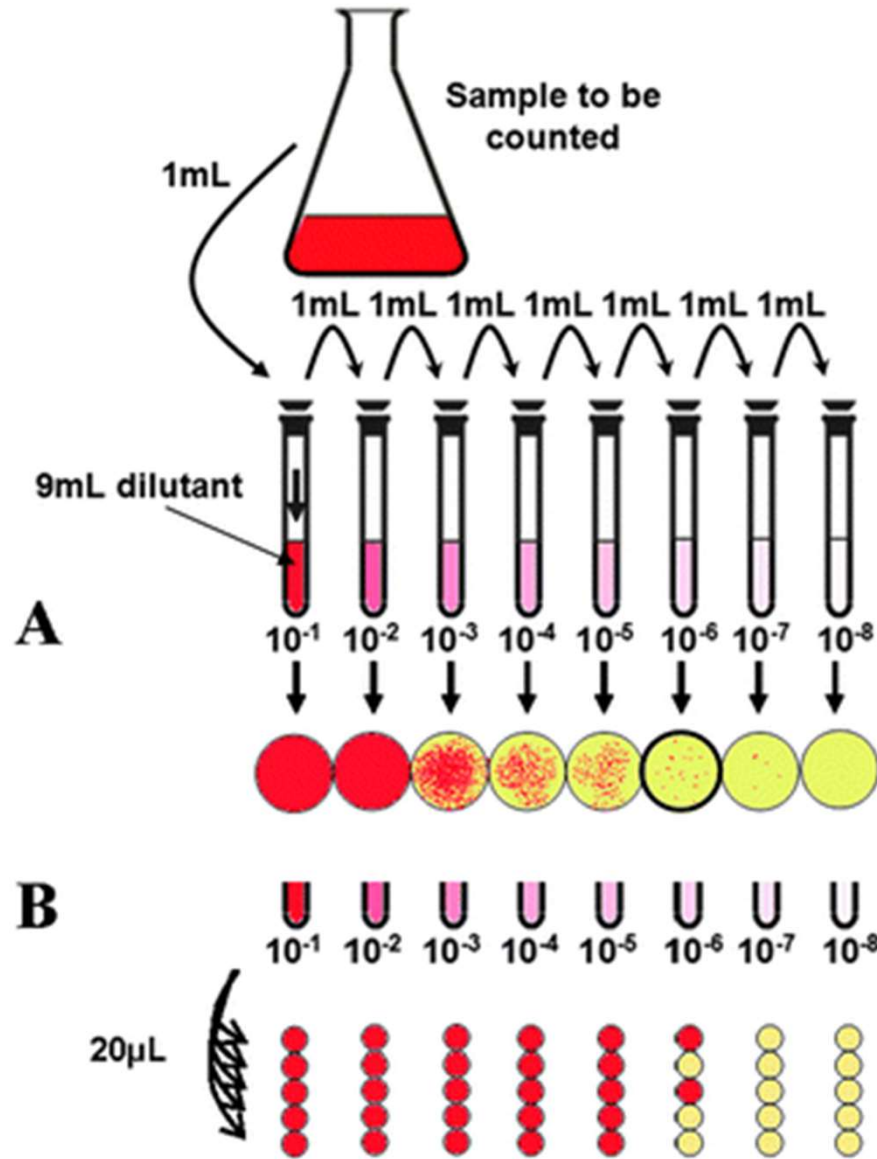
## Spread-plate method



## Pour-plate method



# MPN (most probable number) counts



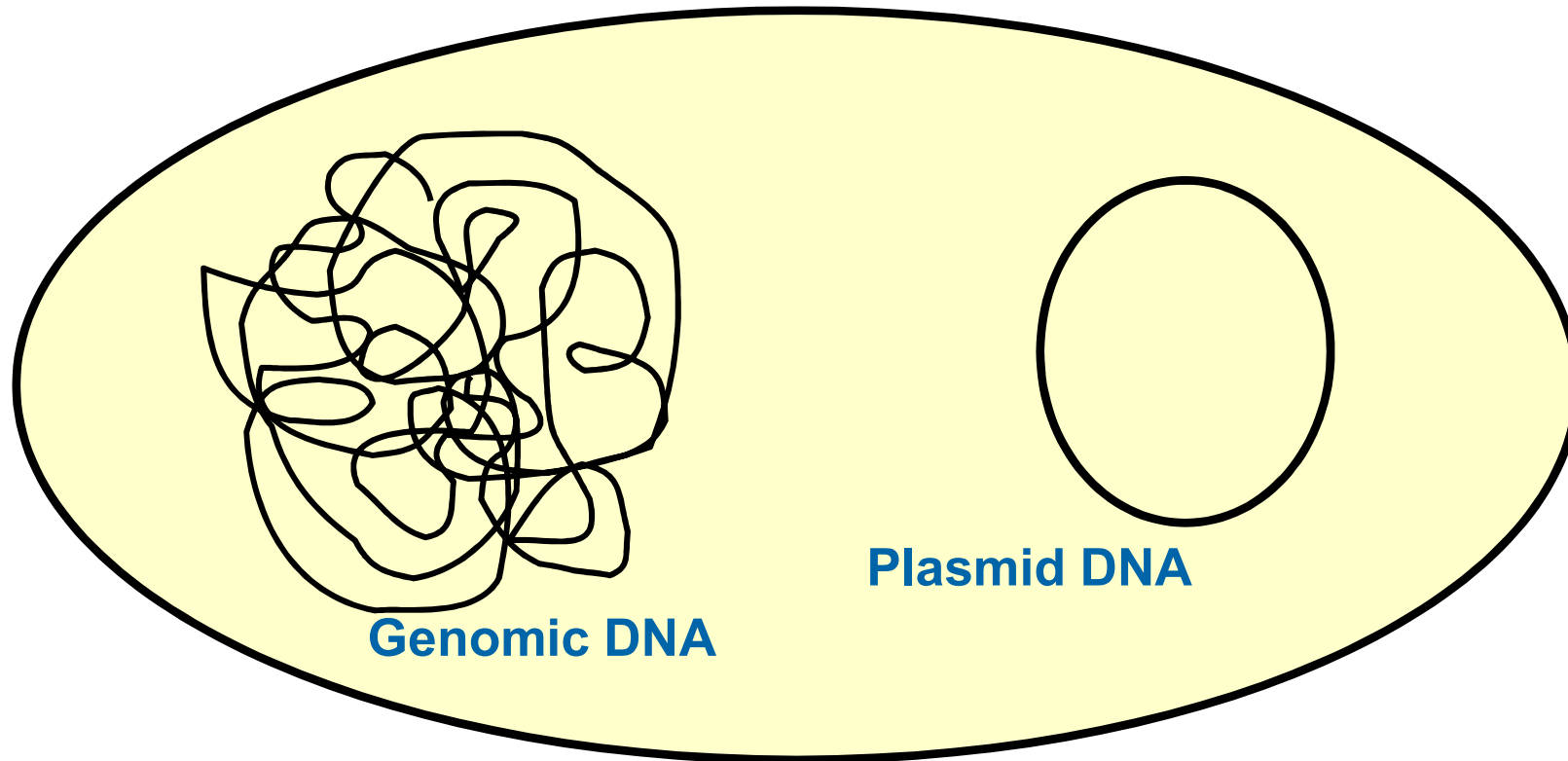
# Molecular methods





# Molecular methods

## Bacterial cell





# Molecular methods

**Aim:** Quantification of different groups of microorganisms in different geosystems

- Conventional cultivation methods are time consuming and capture only a small portion of the microorganisms
- Molecular biological methods are based on the specific detection of nucleic acids (DNA, RNA) and detect all microorganisms



## The 16S rRNA gene (16S rRNA)

- a subunit of the ribosomal RNA responsible for translation of a gene into a protein
- present in all organisms and highly conserved since it performs the same function in all organisms (16S rRNA in prokaryotes, 18S rRNA in eukaryotes)
- easily isolated and detected since it is a relatively abundant molecule in cell
- most importantly, the sequence of 16S rRNA gene has been determined from over 30,000 microbes



## Role of rRNA in the cell

- All functions of a cell are encoded as DNA as **genes** (including the rRNA)
- Genes do nothing but store information
- The information in a gene is transcribed into **mRNA** (messenger RNA) based on the nucleotide sequence of the gene
- mRNA transfers the information in the gene to ribosomes
- **Ribosomes** produce proteins that carry out cellular functions (or synthesize molecules that do) from amino acids based on the nucleotide sequence in the mRNA



## Genes for specific metabolic groups

Gene	Enzyme activity	Microbe function
N-cycle		
<i>amo</i>	ammonia monooxygenase	ammonia oxidation
<i>nirK</i>	nitrite reductase	nitrite reduction
<i>narG</i>	nitrate reductase	nitrate reduction
S-cycle		
<i>apsA, dsrAB</i>	sulfate reduction	sulfate reduction
<i>dsrAB*</i>	sulfur oxidation	sulfur oxidation
C-cycle		
<i>nahA</i>	naphthalene dioxygenase	aromatic degradation
<i>tfdA</i>	2,4-D dioxygenase	herbicide degradation



# DNA extraction

## DNA- enrichment:

- Cultivation of organisms
- Concentration of biomass by e.g. centrifugation, filtration

## DNA- extraction:

- cell lysis (enzymatically, mechanically, chemically)
- protein precipitation
- washing of DNA



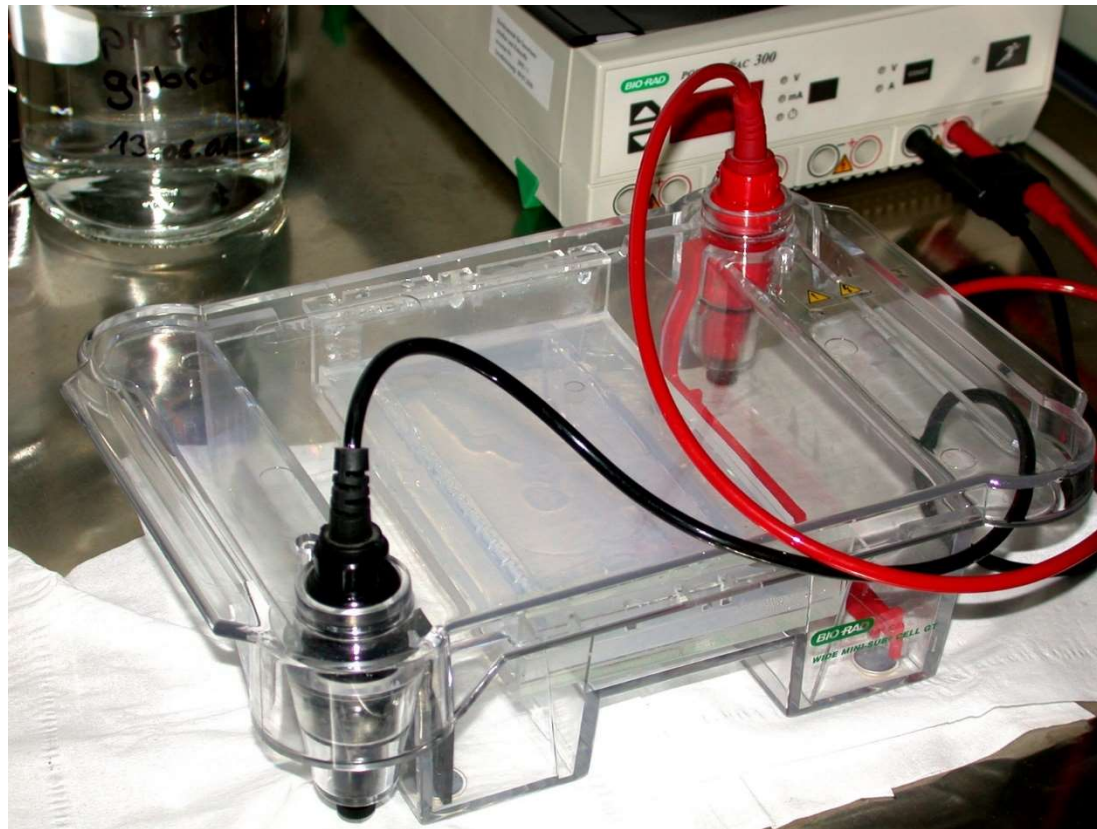
# Polymerase chain reaction







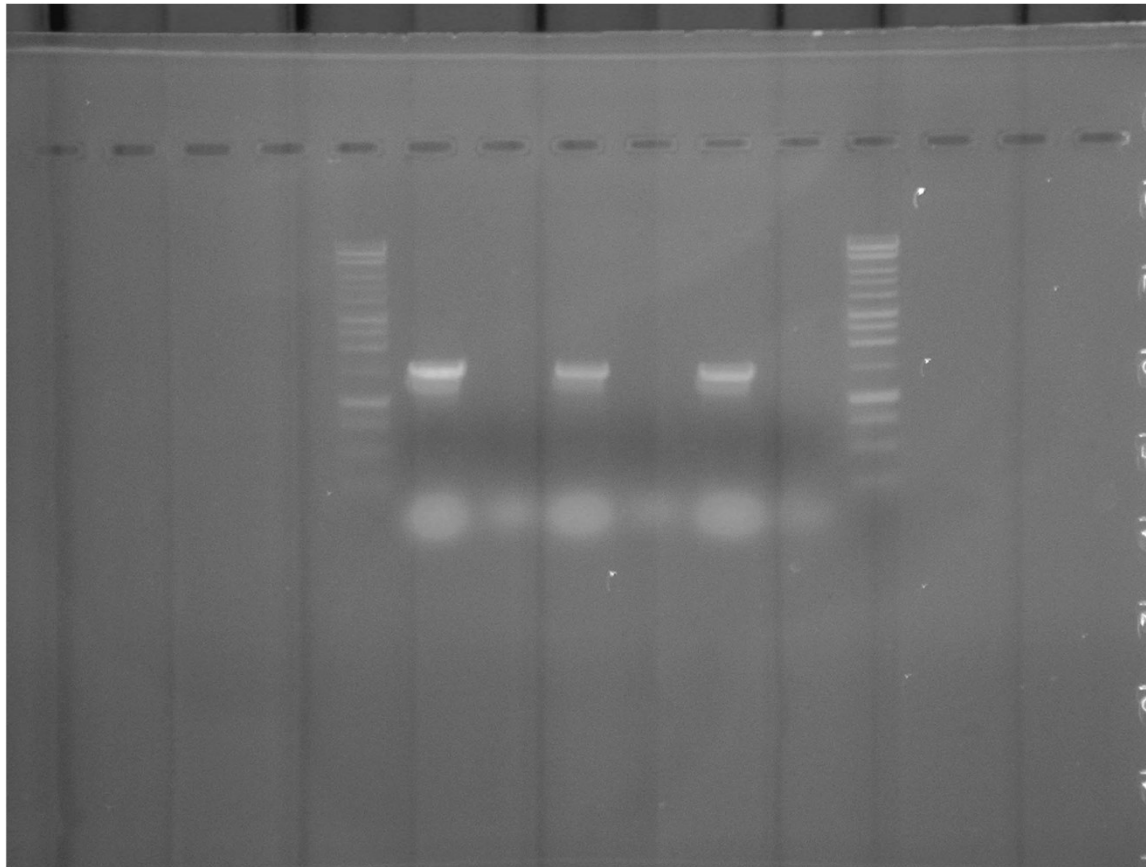
# Gel electrophoreses







## Agarose gel with DNA bands



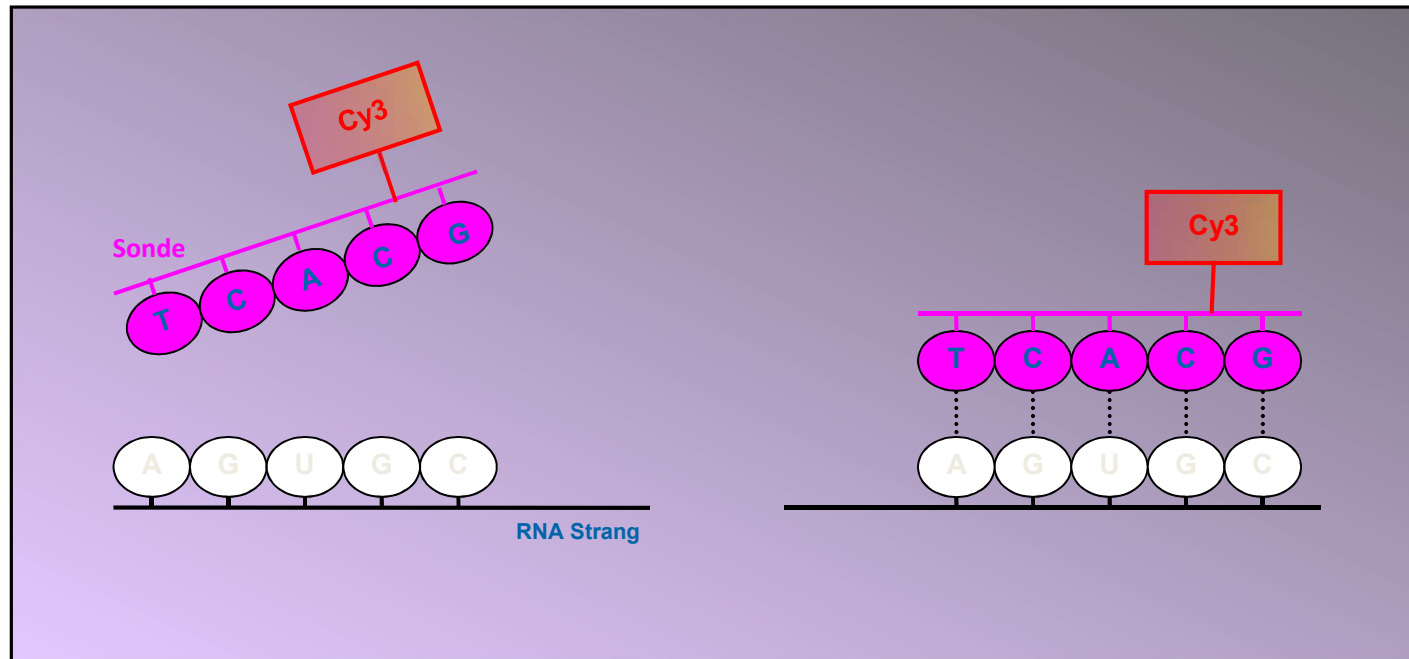


## Fluorescence In Situ Hybridization (FISH oder CARD-FISH)

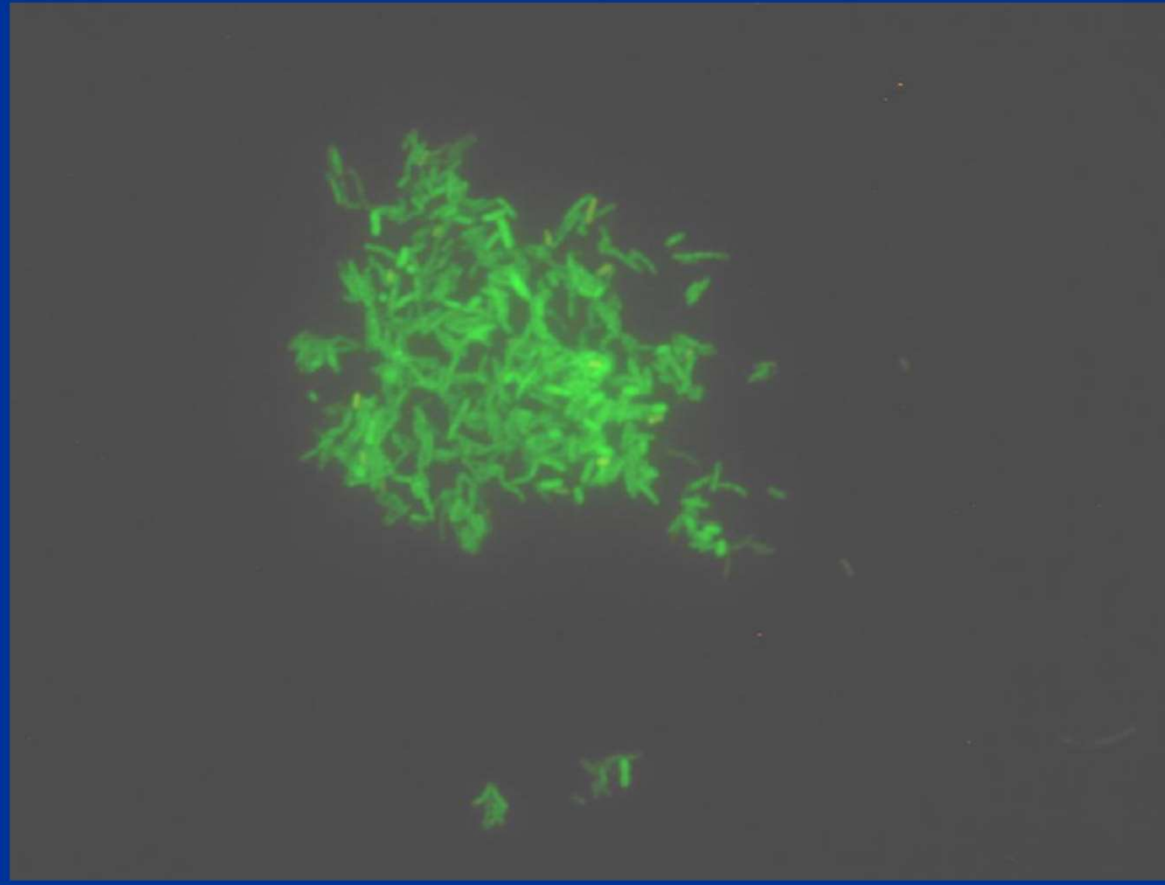
- Samples are chemically fixed to formaldehyde.
- Genetically relevant groups of microorganism-specific gene probes hybridize to a target sequence of ribosomal RNA found only in living cells.
- The gene probes emit a fluorescent signal, causing the microorganisms to shine.
- By counting on the fluorescence microscope, the number of living microorganisms in the sample is determined.
- Analysis time: 2 days

# FISH

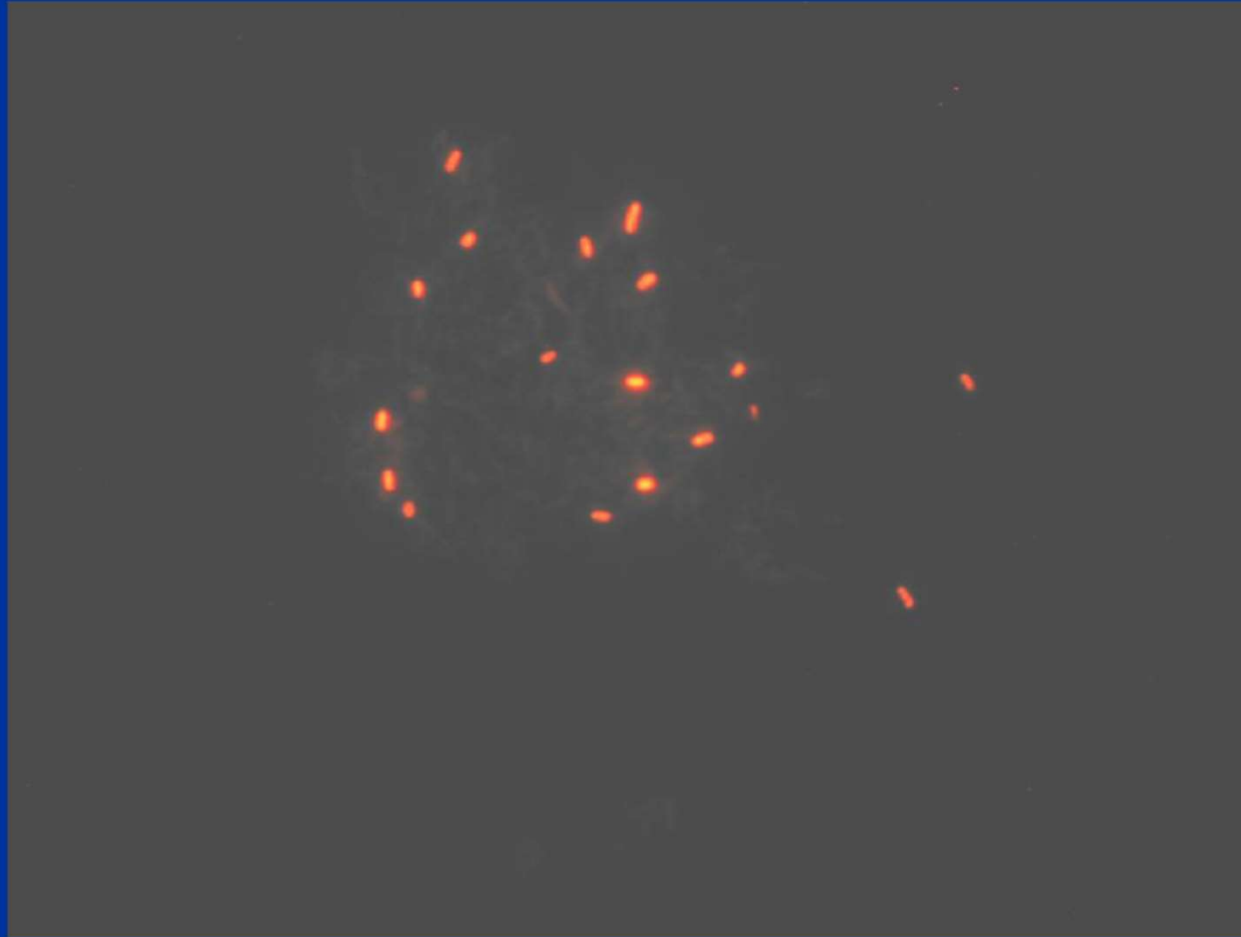
(Fluorescence In Situ Hybridization)



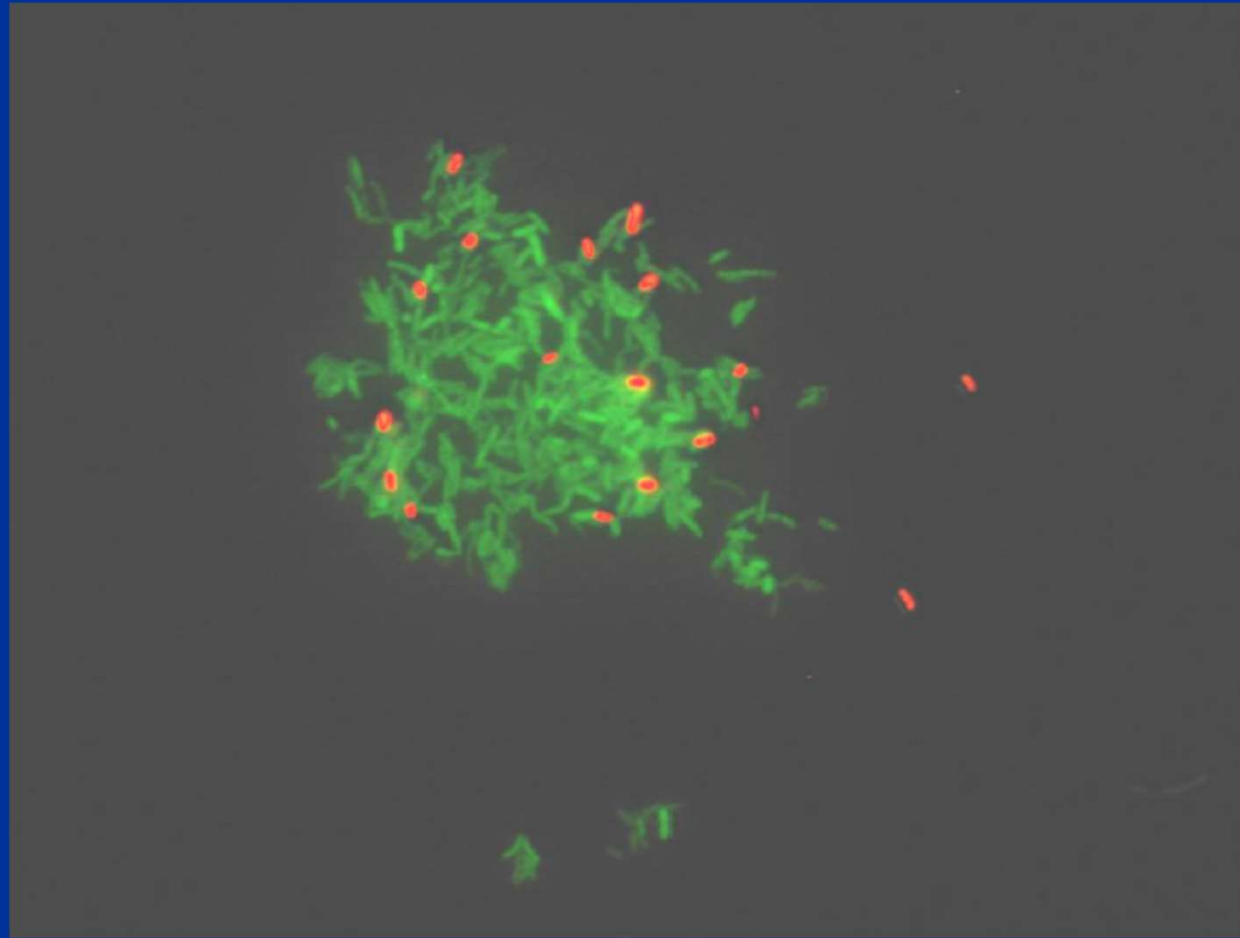
## Biofilm probed with bacterial specific oligonucleotide



**Same sample with *Acidithiobacillus ferrooxidans* probe**

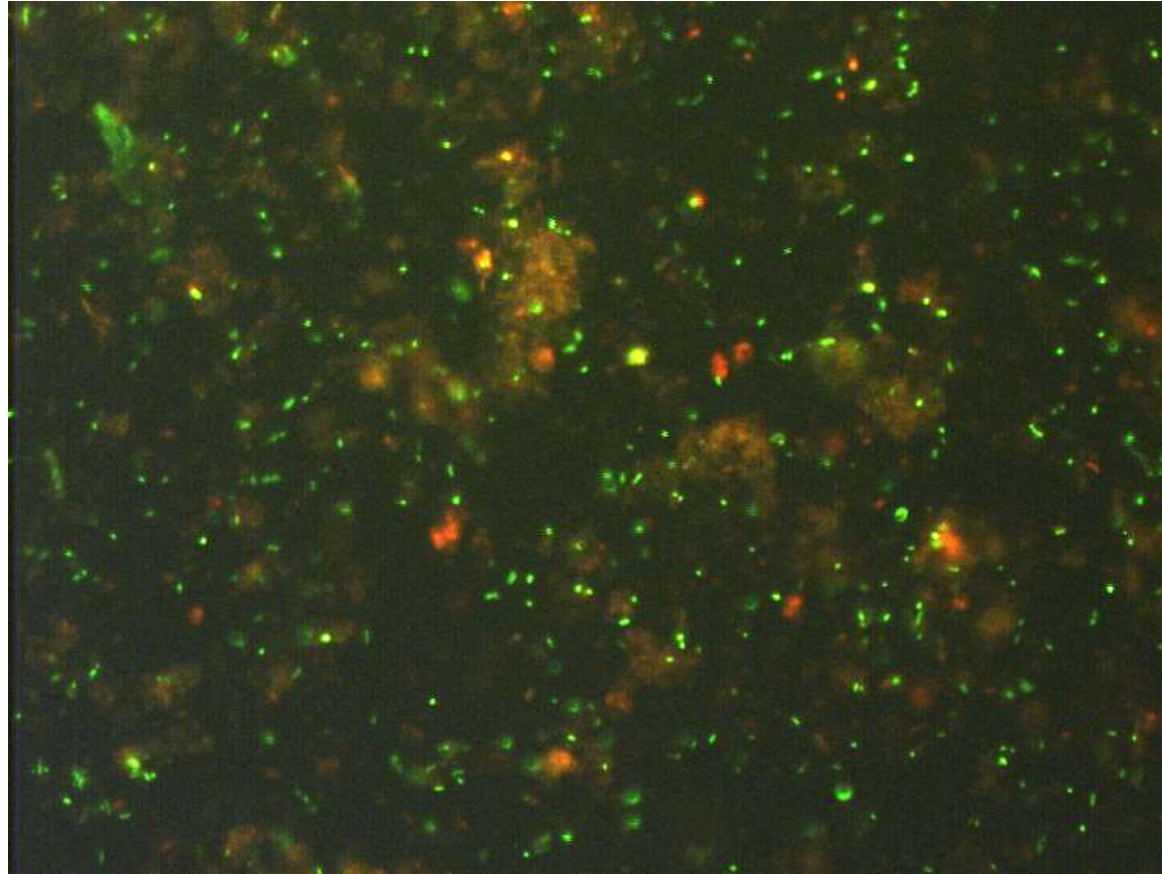


## Combined images

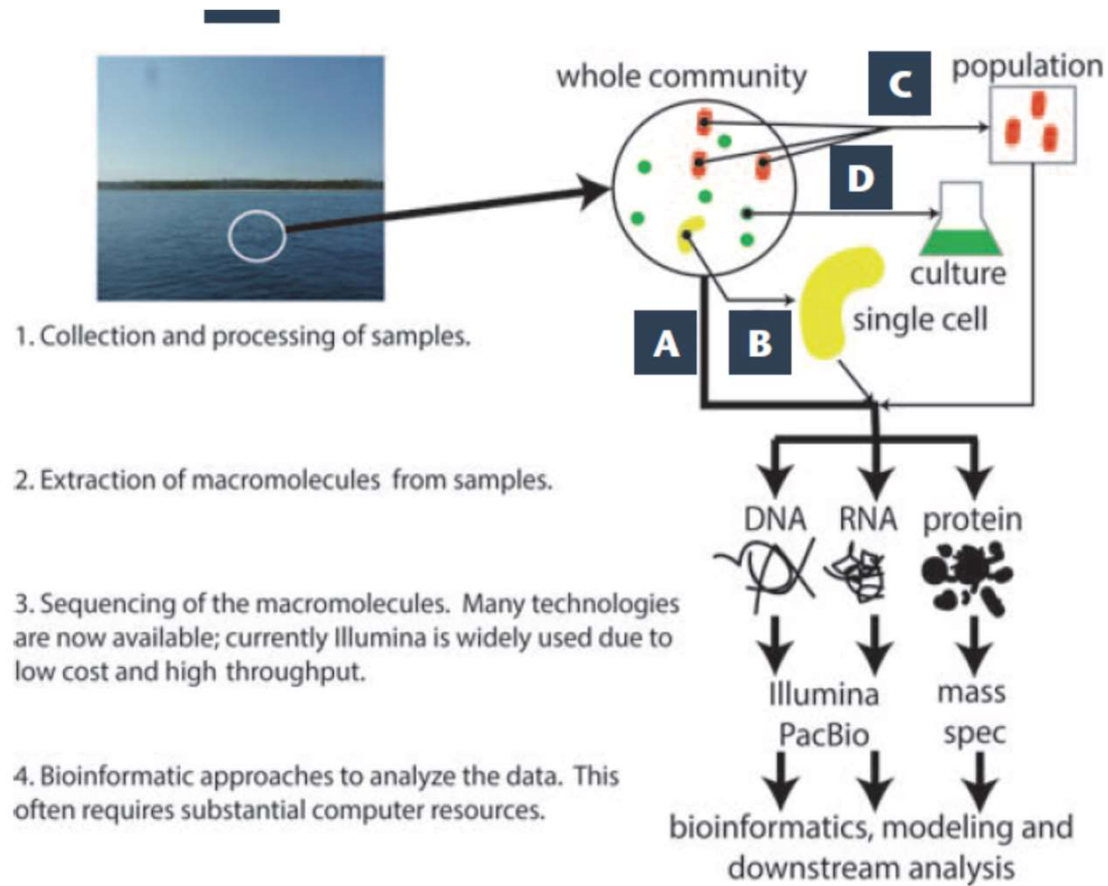




## Microbial cells in tailing sample



# Omics studies



1. Collection and processing of samples.

2. Extraction of macromolecules from samples.

3. Sequencing of the macromolecules. Many technologies are now available; currently Illumina is widely used due to low cost and high throughput.

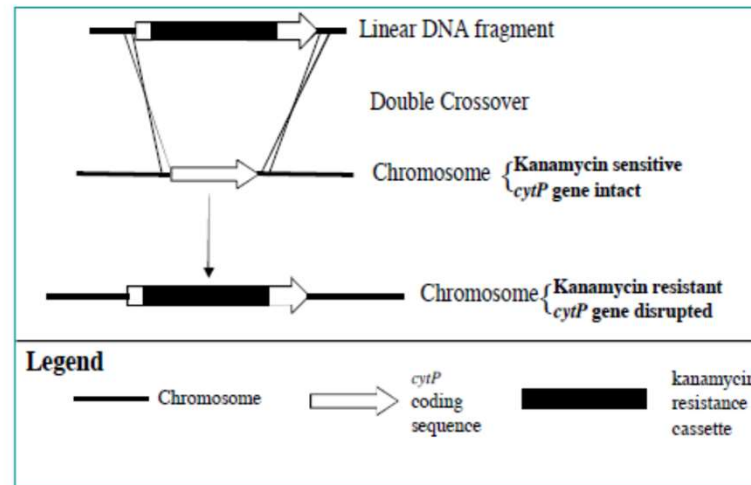
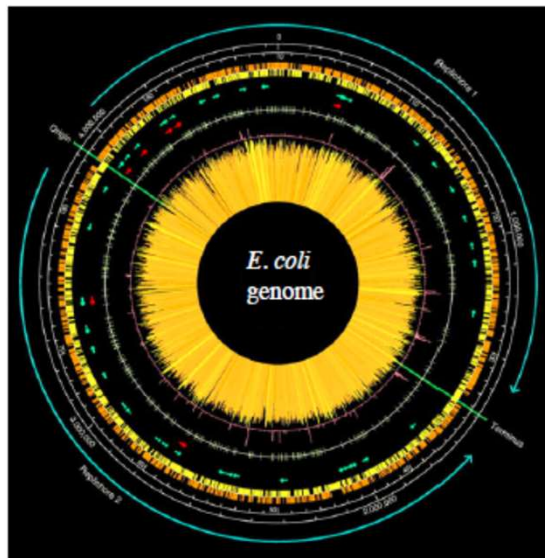
4. Bioinformatic approaches to analyze the data. This often requires substantial computer resources.

**FIGURE 2** Processing samples for omics approaches. Samples may be from the field, (e.g. seawater, soil, or rock), experiments, or laboratory cultures, and can be analyzed directly in four ways: **(A)** as whole communities (“metagenomics”); **(B)** processed to select specific cells (single cell genomics); **(C)** as populations; **(D)** as cultures.



# The genomics revolution

- Genomes sequenced and available for many subsurface microbes
- Tools available for genetic analysis via WWW
- Genetic systems available for many subsurface microorganisms
- Gene arrays and proteomics tools





## Omics studies

- Computational power for analysis, assembly, annotation, and comparative studies
- Data storage and accessibility
- Adoption of worldwide standards (standardized genome, metagenome reports)
  - The Genomic Standards Consortium
  - Standards in Genomic Sciences journal



## Molecular methods to determine diversity and function of microorganisms

- Statement about type of microorganisms available
- Selective quantification of different microorganism groups
- Determination (and quantification) of existing metabolic processes
- Depending on the efficiency of the extraction of nucleic acids / proteins
- often not enough comparison data available
- more complex methods have lower resolution
- expensive



## Summary

- total microscopic cell count good for overview,
- BUT difficult to define bacteria types and physiology
- Cultivation-dependent methods easy to handle,
- BUT difficult to cultivate all organisms
- Overall picture about microbiology of a sample achieved only with molecular biological methods
- Molecular biological methods use DNA, RNA or proteins
- Each method has its drawbacks, so more than one method should be used to characterize a sample
- (Cultivation-dependent methods supplement molecular biological data)



**Thank you for your attention!**

**Glück auf!**

