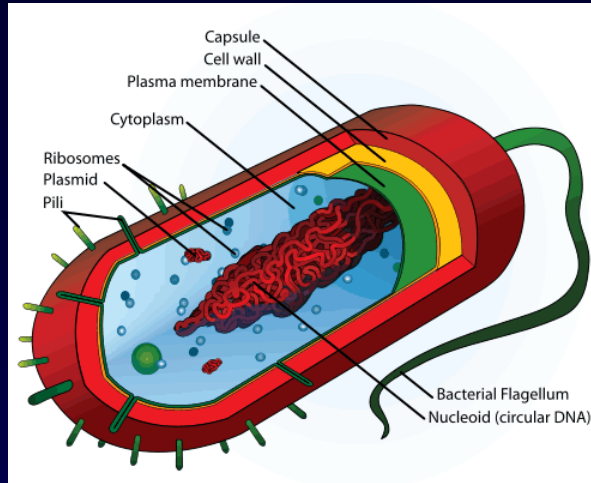


# Microbiological methods

Dr. Lidita Khandeparker



# What are Bacteria??



A typical bacterium

- ✓ Bacteria are a group of single-celled microorganisms with prokaryotic (an organism without a nucleus) cellular configuration
- ✓ The organelles of eukaryotes (mitochondria and chloroplasts) are thought to be remnants of Bacteria that invaded, or were captured by primitive eukaryotes in the evolutionary past
- ✓ Numerous types of eukaryotic cells that exist today are inhabited by endosymbiotic prokaryotes
- ✓ A typical bacterial cell is about 1 micrometer in diameter or width

## Bacteria occur in three main shapes



**Spherical**

**Rod shaped**

**Spiral**

# How they have evolved??

Modern bacteria's ancestors - single-celled microorganisms - appeared on earth about 4 billion years ago. They were the first life forms on Earth.

✓ All prokaryotes were once placed in the kingdom Monera

✓ Recently, they are divided into two different kingdoms: the Eubacteria and the Archaeobacteria

## Archaeobacteria

- Very old bacteria, evolved a long time ago
- Many live in extreme environments such as hot springs, salty water and acidic environments
- Some live in or on organisms
- Autotrophic or heterotrophic
- Can be harmful or helpful

## Eubacteria

- More recently evolved bacteria
- Live in or on organisms
- Autotrophic or heterotrophic
- Can be harmful or helpful

# How do bacteria feed themselves?



## Heterotrophic bacteria

## Autotrophic bacteria

- Photoautotrophs
- Chemoautotrophs

Nutritional Type	Energy Source	Carbon Source	Examples
Photoautotrophs	Light	CO <sub>2</sub>	Cyanobacteria, some Purple and Green Bacteria
Photoheterotrophs	Light	Organic compounds	Some Purple and Green Bacteria
Chemoautotrophs or Lithotrophs (Lithoautotrophs)	Inorganic compounds, e.g. H <sub>2</sub> , NH <sub>3</sub> , NO <sub>2</sub> , H <sub>2</sub> S	CO <sub>2</sub>	A few Bacteria and many Archaea
Chemoheterotrophs or Heterotrophs	Organic compounds	Organic compounds	Most Bacteria, some Archaea

## How do **bacteria** multiply ?

- Bacteria can reproduce at tremendous speeds
- Some bacteria can reproduce as often as once every 20 minutes!
- However, bacteria have to have certain conditions in which to reproduce
- These conditions are not often met, and that is one thing that keeps bacteria from growing out of control

### Bacteria reproduce using two basic methods

**Asexual reproduction** involves only one individual or parent. The offspring generated by asexual reproduction are exact duplicates of the parent.

**Binary fission** is the process by which a bacteria splits into two cells. Each cell gets an exact copy of the parent cell's genetic material.

**Sexual reproduction** involves joining of two parent cells and exchanging of genetic materials. The offspring will have a mixture of parent cells' traits. **Conjugation** is the process by which bacteria join and exchange genetic materials. Once genetic materials are exchanged, each bacteria cell will go through binary fission to produce an offspring with a new genetic makeup

# How they are unique?

## **Bacteria inhabit different kinds of environments**

Aerobes (aerobic bacteria)

Anaerobes (anaerobic bacteria)

Facultative anaerobes (facultative anaerobic bacteria)

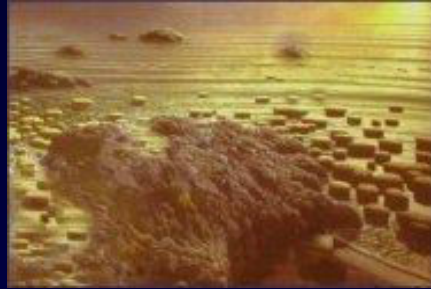
Mesophile (mesophilic bacteria)

Extremophiles (extremophilic bacteria)

- Thermophiles (thermophilic bacteria)
- Halophiles (halophilic bacteria)
- Acidophiles (acidophilic bacteria)
- Alkaliphiles (alkaliphilic bacteria)
- Psychrophiles (psychrophilic bacteria)



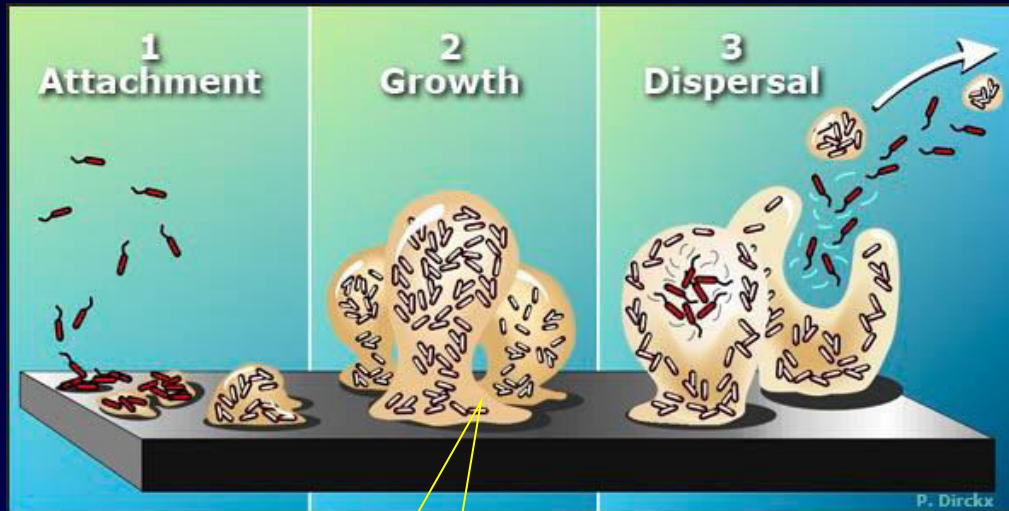
# Bacteria are found everywhere



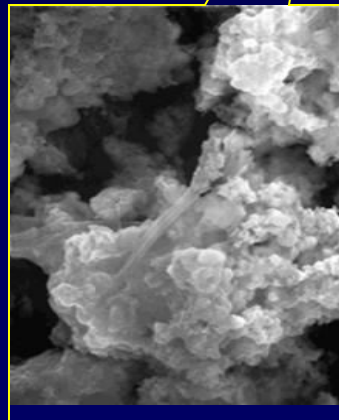
- Soil
- Radioactive waste
- Water
- Plants
- Animals
- Deep in the earth's crust
- Organic material
- Arctic ice
- Glaciers
- Hot springs
- The stratosphere (between 6 to 30 miles up in the atmosphere)
- Ocean depths - they have been found deep in ocean canyons and trenches over 32,800 feet (10,000 meters) deep.
- They live in total darkness by thermal vents at incredible pressure.



# How bacteria colonize a surface?



(Source: P. Dirckx)



Biofilm





# Phases of Bacterial growth

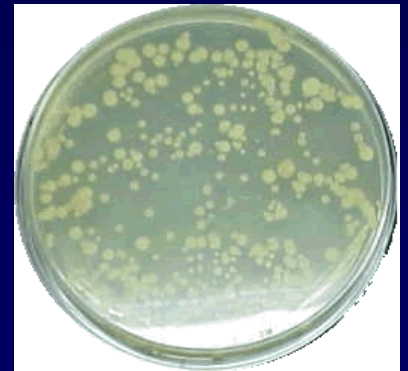
- 1) Lag phase
- 2) Log (logarithmic) phase
- 3) Stationary phase
- 4) Decline phase or death phase

## Growth in Colonies

A pure culture contains only one species or strain

A colony is a population of cells that arise from a single cell or spore or from a group of attached cells

A colony is often called as a colony-forming unit (CFU)



# Bacterial Quantification

## Direct methods

Plate counts

MPN

Direct microscopic count/Flow cytometry

Dry weight

## Indirect methods

Turbidity

Metabolic activity

Dry weight

# Methods to classify Bacteria

## Phenotypic Classification

Microscopic morphology

Macroscopic morphology

Biotyping

Serotyping

Antibiogram patterns

Phage typing

## Genotypic Classification

Guanine and cytosine ratio

DNA hybridization

Nucleic acid sequence analysis

Plasmid analysis

Ribotyping

Chromosomal DNA fragment

## Analytic Classification

Whole cell lipid analysis

Whole cell protein analysis

Multifocus locus enzyme electrophoresis

Cell wall fatty acid analysis

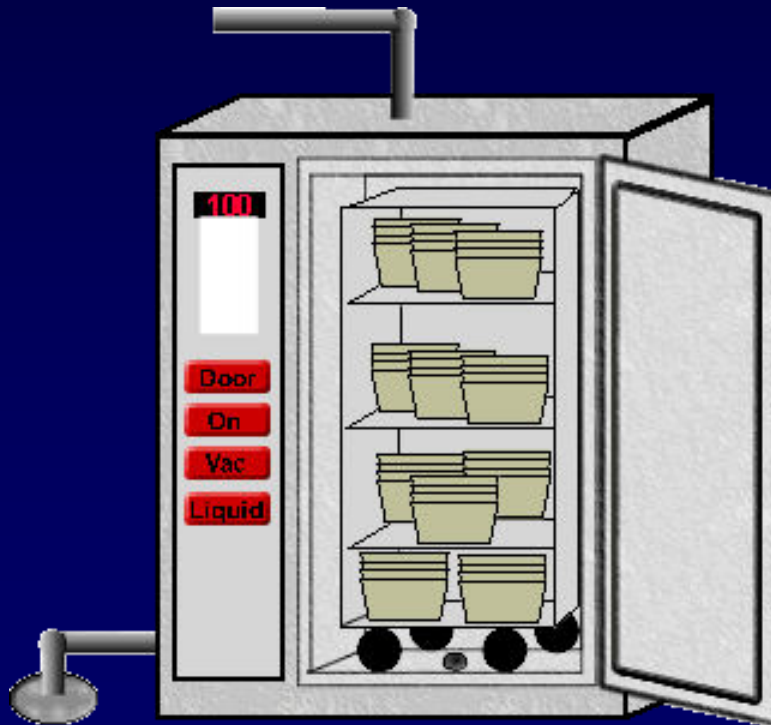
# Bacterial Identification

- The **Gram stain**, which divides most bacteria into two main groups, is the first step in bacterial identification.
- Cell shape and size
- Acid fast reaction
- Special structures

# Sterilization

Moist heat, dry heat, chemicals, and radiation

➤ Steam autoclave is primary means of sterilizing



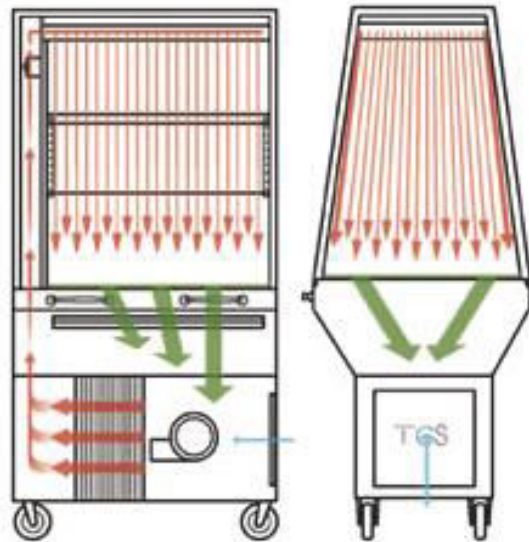
Autoclave

Steam in at top, displaces air out drain in bottom.

Load the equipments to be sterilized. To start, close door tightly and turn on timer.

When the temperature reaches  $121^{\circ}\text{C}$  and pressure reaches 15 pounds per square inch (psi), the timer begins sterilization time.

Steam is then vented.



Thoren Caging uses a different air flow as well.



(Click on the picture to view the web page.)

Courtesy of Thoren Caging  
([www.thoren.com](http://www.thoren.com))



Laminar flow

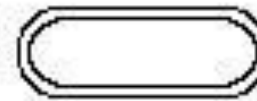
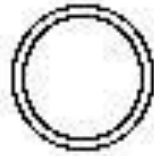


# Physiological/Biochemical Characteristics

- Traditional mainstay of bacterial identification
- Diagnostic tests for determining the presence of specific enzymes and assessing nutritional and metabolic activities
- Examples
  - Fermentation of sugars
  - Capacity to metabolize complex polymers
  - Production of gas
  - Presence of enzymes
  - Sensitivity to antimicrobial drugs

**GRAM +**

**GRAM -**



Fixation



Crystal  
Violet



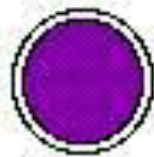
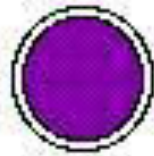
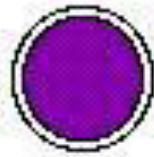
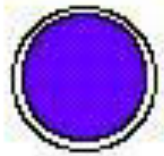
Iodine  
treatment

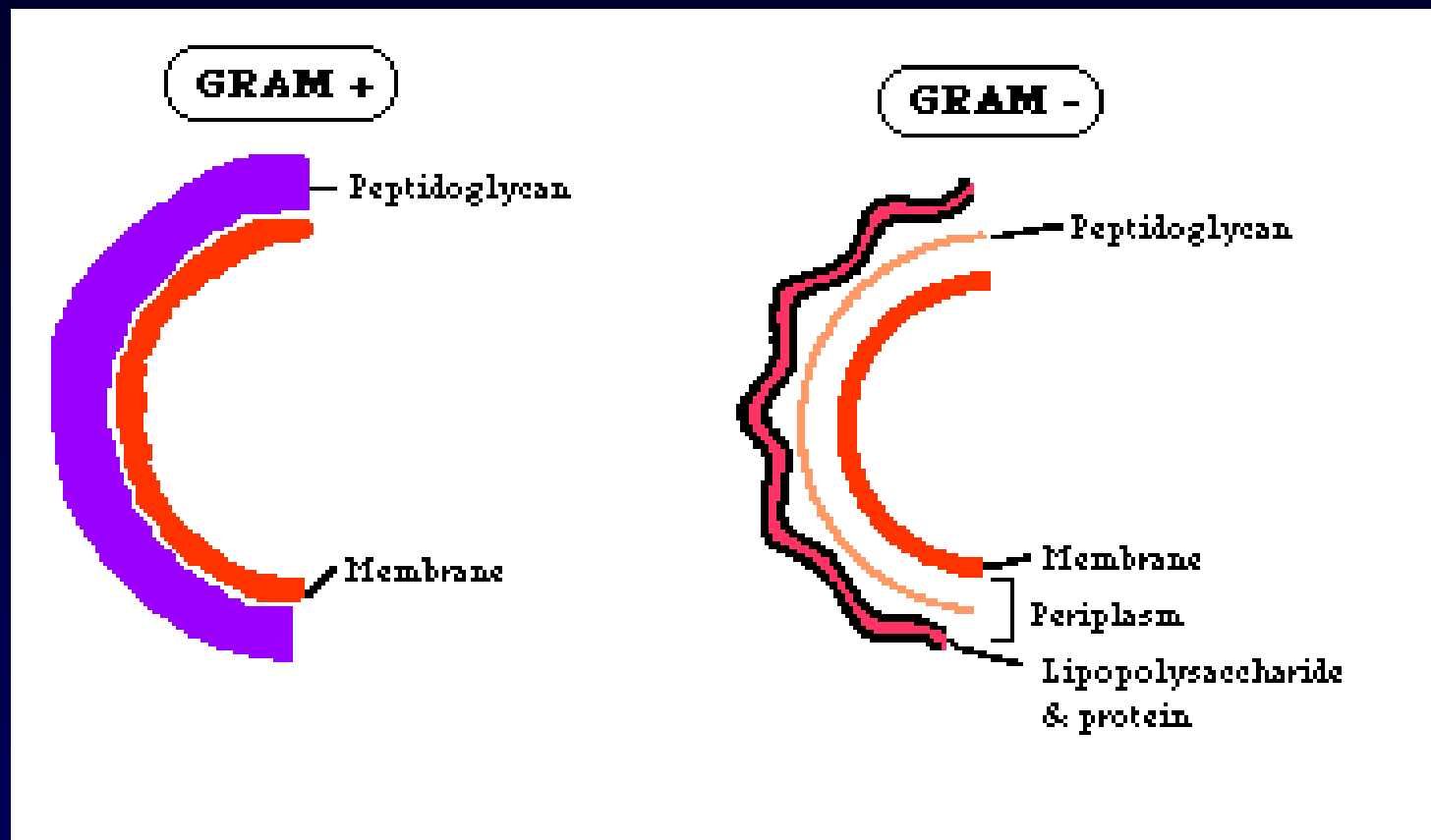


Decolorization

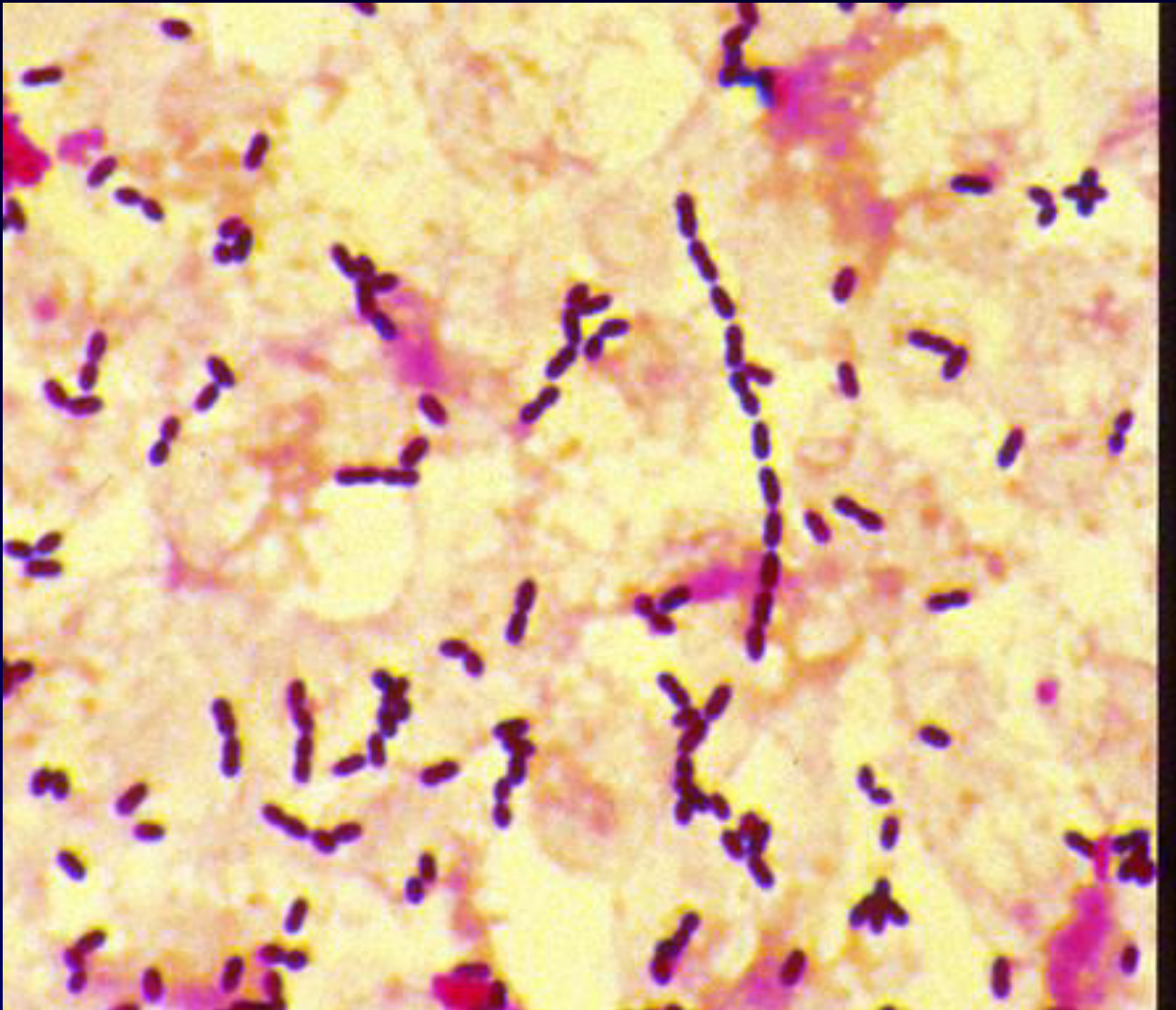


Counter stain  
(safranin)



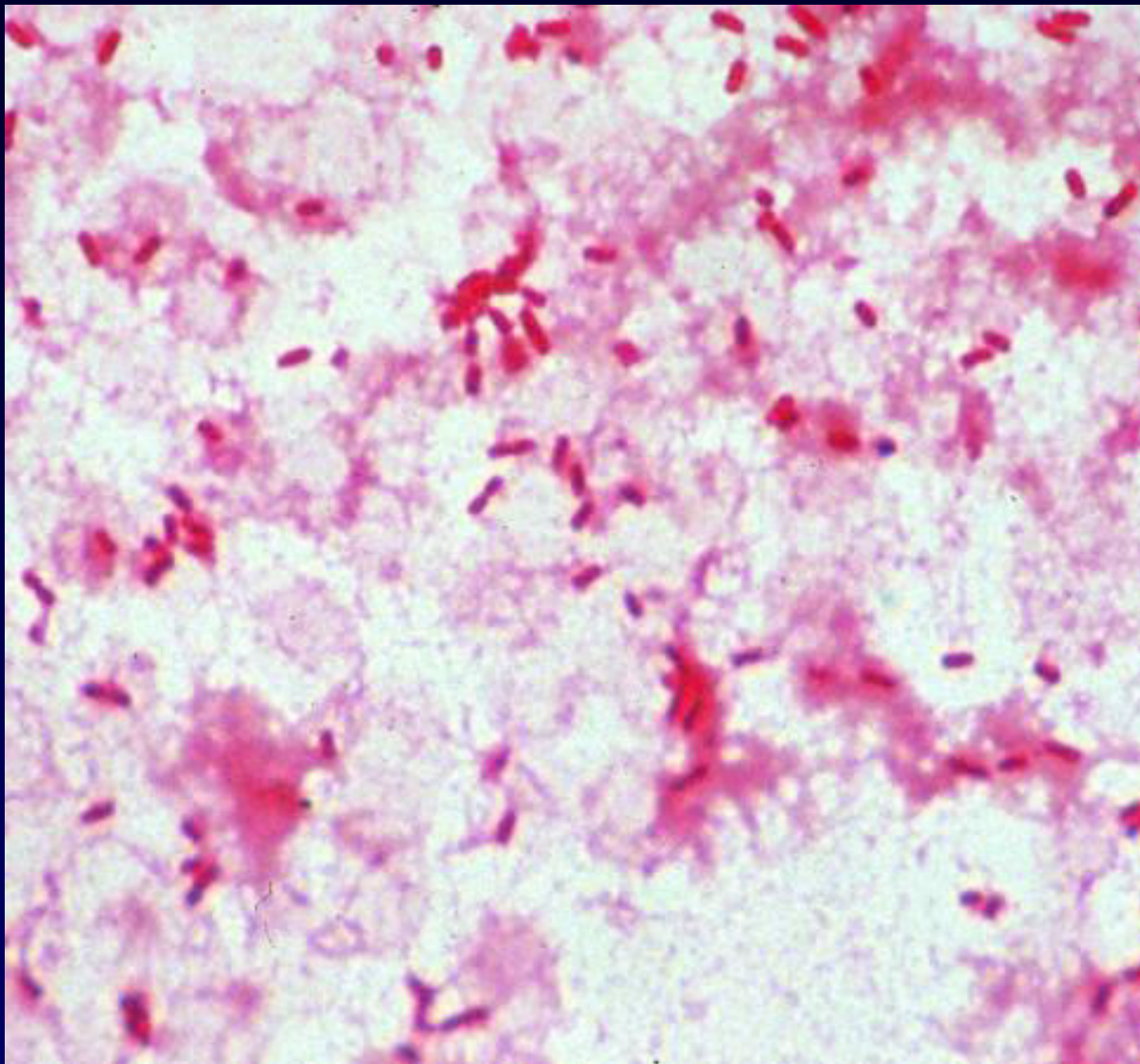


In **Gram-positive bacteria**, the purple crystal violet stain is trapped by the layer of peptidoglycan which forms the outer layer of the cell. In **Gram-negative bacteria**, the outer membrane of lipopolysaccharides prevents the stain from reaching the peptidoglycan layer. The outer membrane is then permeabilized by acetone treatment, and the **pink safranin counterstain** is trapped by the peptidoglycan layer.



Is this gram stain positive or negative?  
Identify the bacteria.



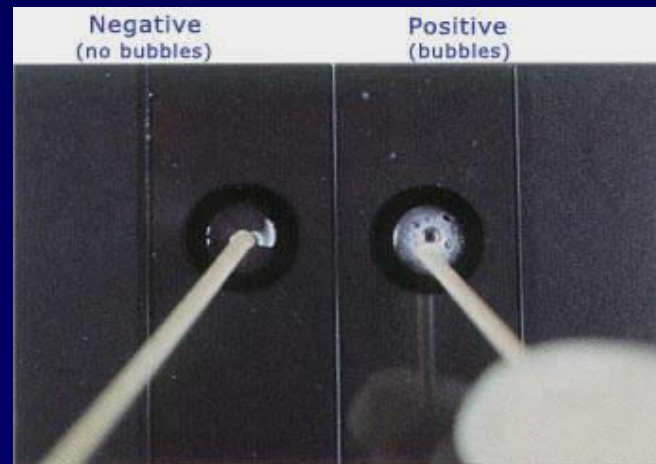


Is this gram stain positive or negative?  
Identify the bacteria.



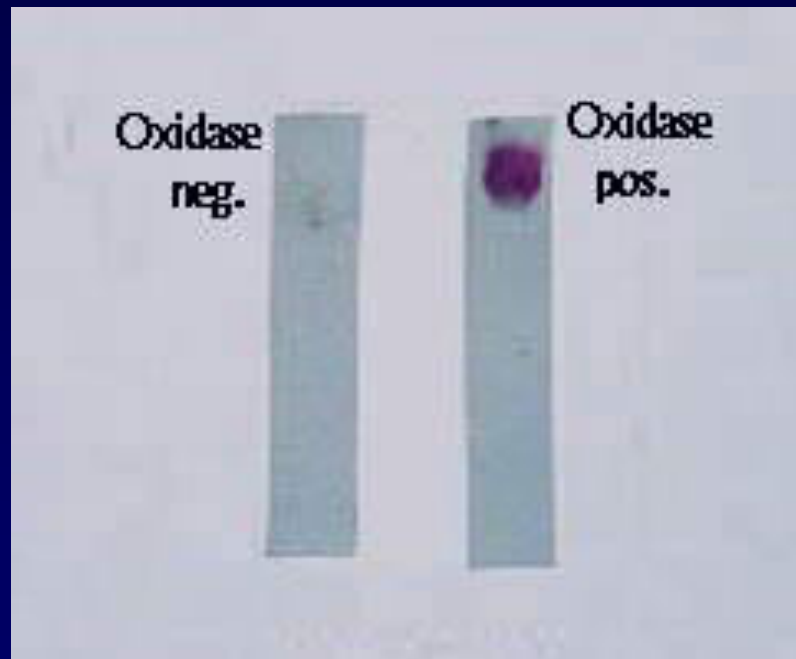
## Catalase test

- This test is used to differentiate those bacteria that produce the enzyme catalase, such as *staphylococci*, from non-catalase producing bacteria such as *streptococci*.
- This enzyme converts hydrogen peroxide into water and oxygen.



## Oxidase Test

When the organism is oxidase-producing, the phenylenediamine (oxidase reagent) will be oxidized to a deep purple color.



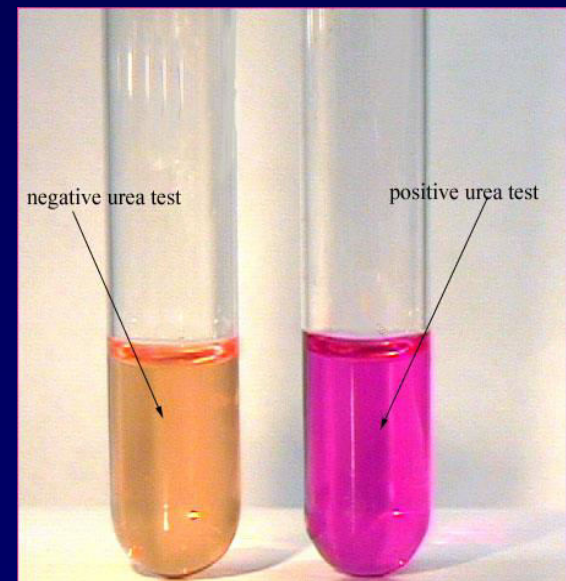
# Urease Test

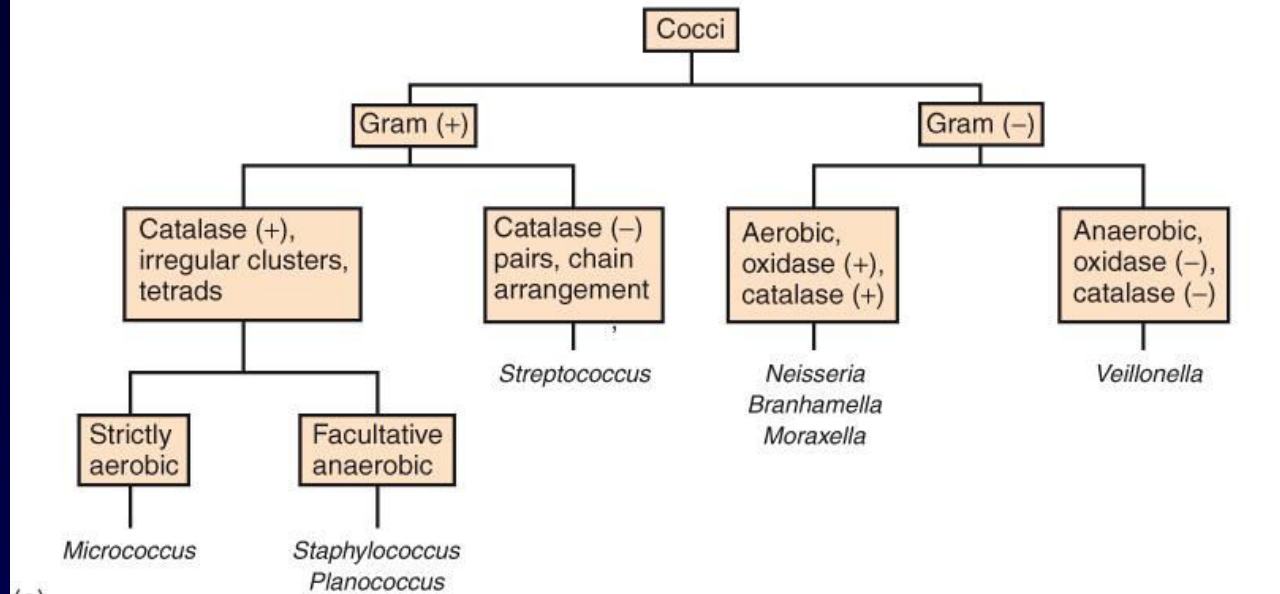
*Proteus* strains are strong urease producers.

## Principle:

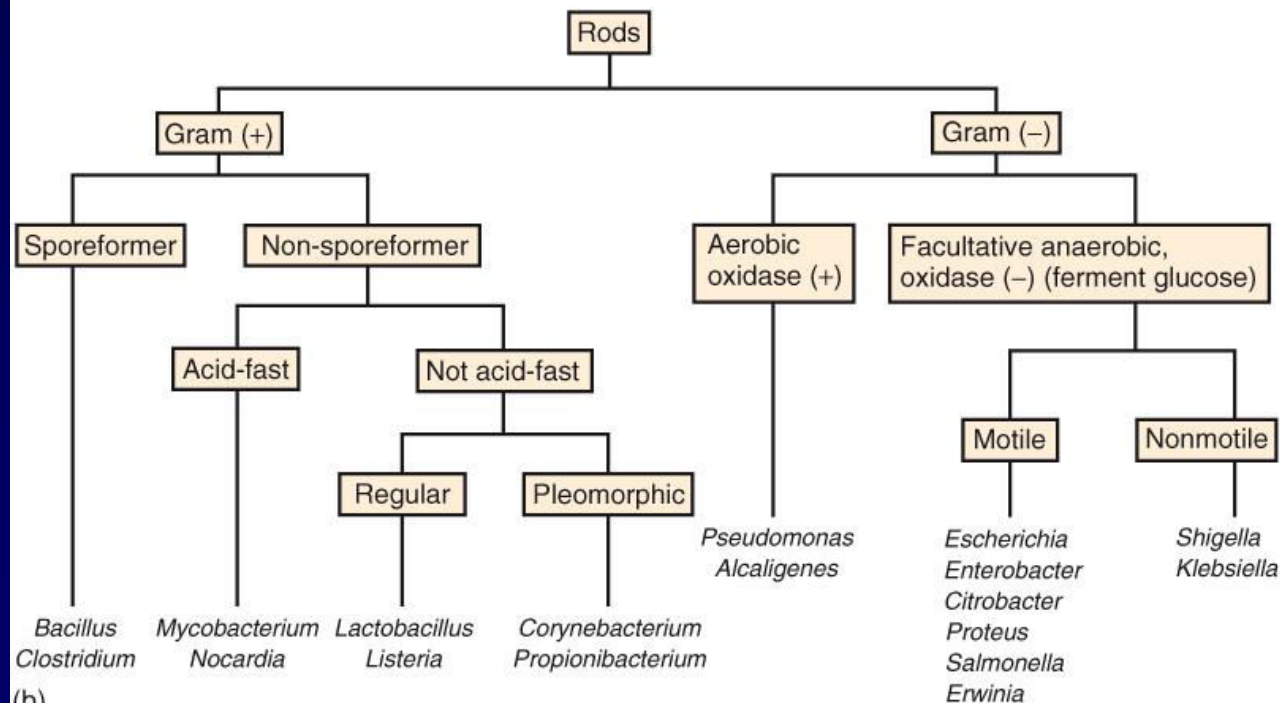
The test organism is cultured in a medium which contains urea and the indicator phenol red. When the strain is urease producing, the enzyme will break down the urea to give ammonia and carbon dioxide.

With the release of ammonia, the medium becomes alkaline as shown by a change in colour of the indicator to pink-red.





(a)



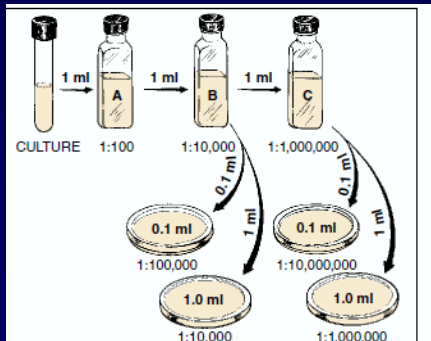
(b)

# Protocols For Viable Bacterial Enumeration



Pouring Plate using selective media

**Processing of Samples for viable bacteria**



Serial Dilutions



Spreading inoculum on surface evenly



Colonies grow

Count colonies

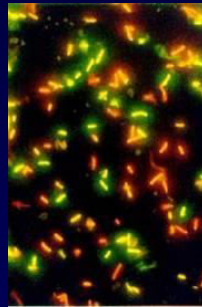


# Bacterial abundance - Total Bacterial Count (TBC) using epifluorescence microscopy

The two most widely used fluorochromes used to identify bacteria are

- Acridine Orange (AO) – 3,6-bis[dimethylamino]acridinium chloride
- DAPI - 4,6-diamidino-2-phenylindole

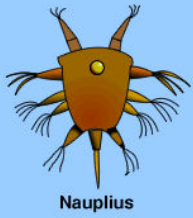
(AODC method)



- Acridine Orange binds to both DNA and RNA with an excitation maximum of approximately 470 nm
- AO-stained single-stranded nucleic acids emit orange-red fluorescence, while those that are double stranded fluoresce green in vivo

The distribution of dead, metabolically inactive but alive and living cells cannot be determined as DNA retains its staining properties even in non-viable cells.

# What is Viability of organisms



Living



Dead?

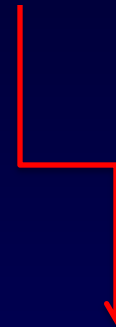


Viable?

Two color assays

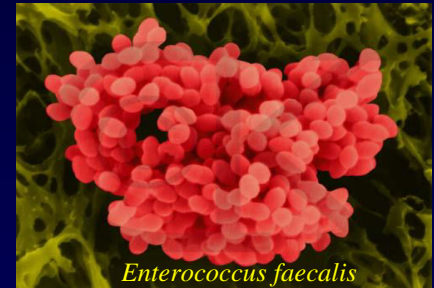
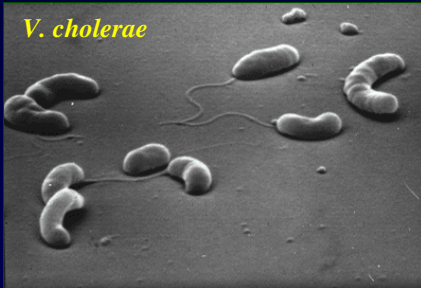


- ✓ Epifluorescence microscopy
- ✓ Confocal microscopy
- ✓ Flowcytometry



Flow cytometry allows rapid enumeration, physical and biochemical characterization of cells from a non-homogenous population

# Important Pathogenic Bacteria



# Protocols For Bacterial Enumeration

# Collection of Samples

Surface



Bottom



Zooplankton



Sediment

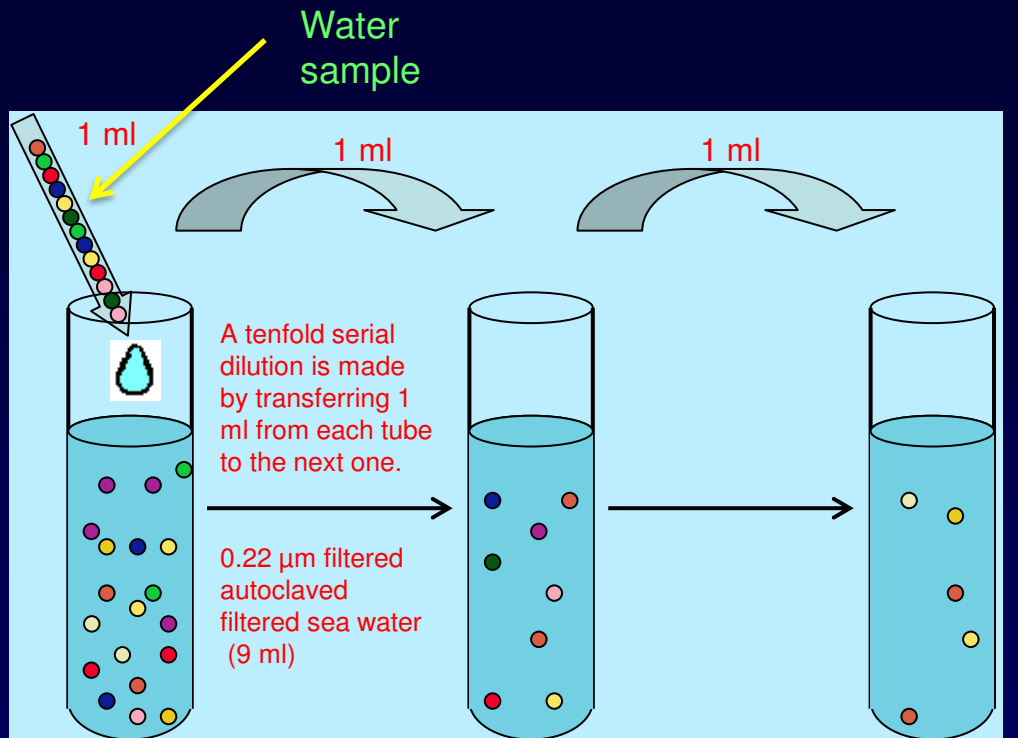


Fouling





# Protocol – water sample

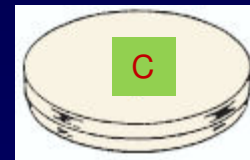
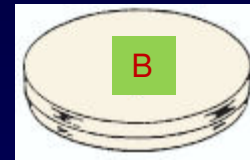
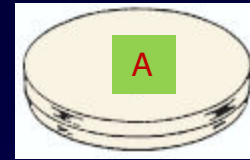


A ( $10^{-1}$ )

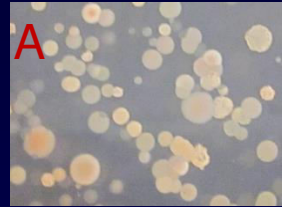
B ( $10^{-2}$ )

C ( $10^{-3}$ )

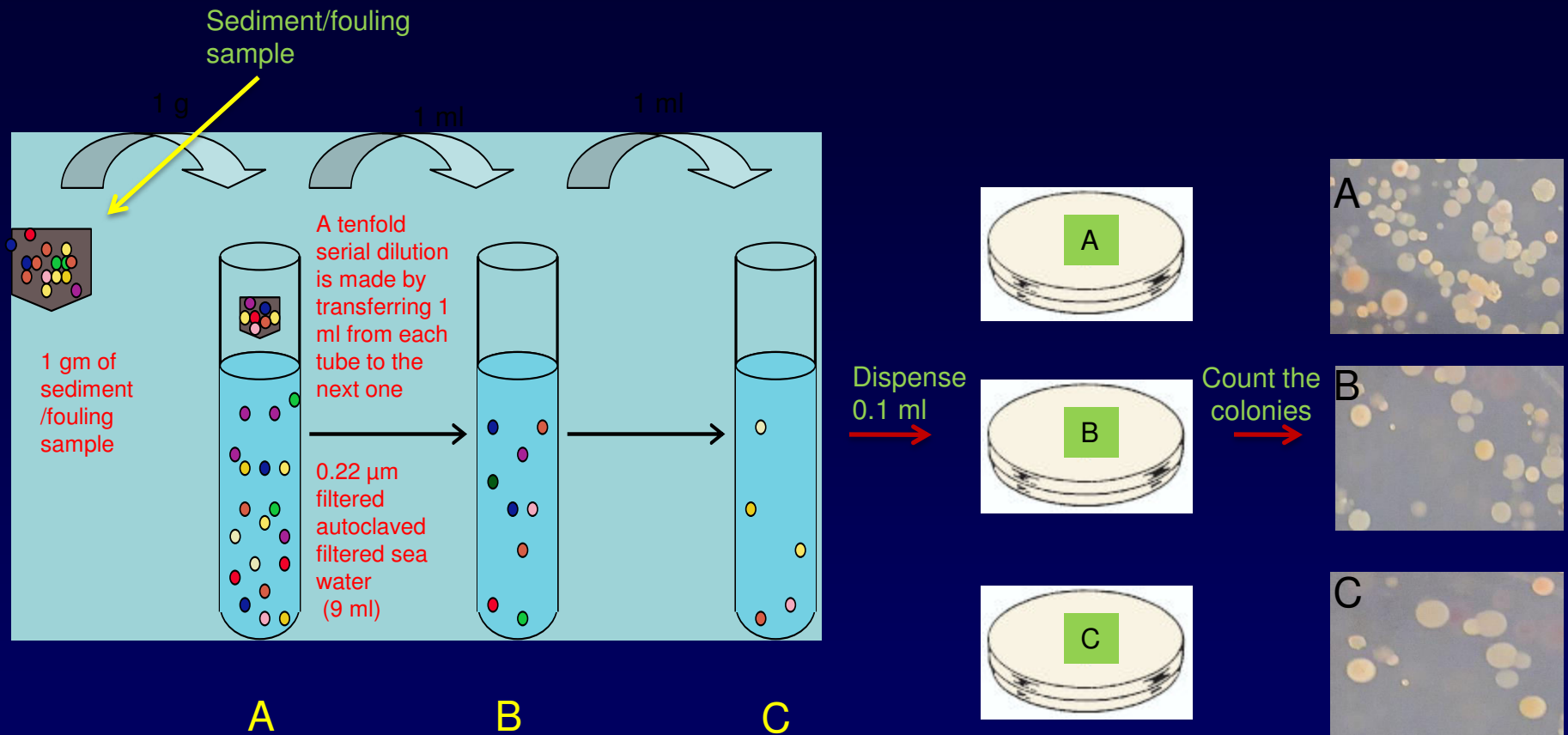
Dispense  
0.1 ml  
→



Count the colonies  
→



# Protocol – Sediment/fouling sample



# Protocol – Zooplankton sample

Sampling Net



Zooplankton

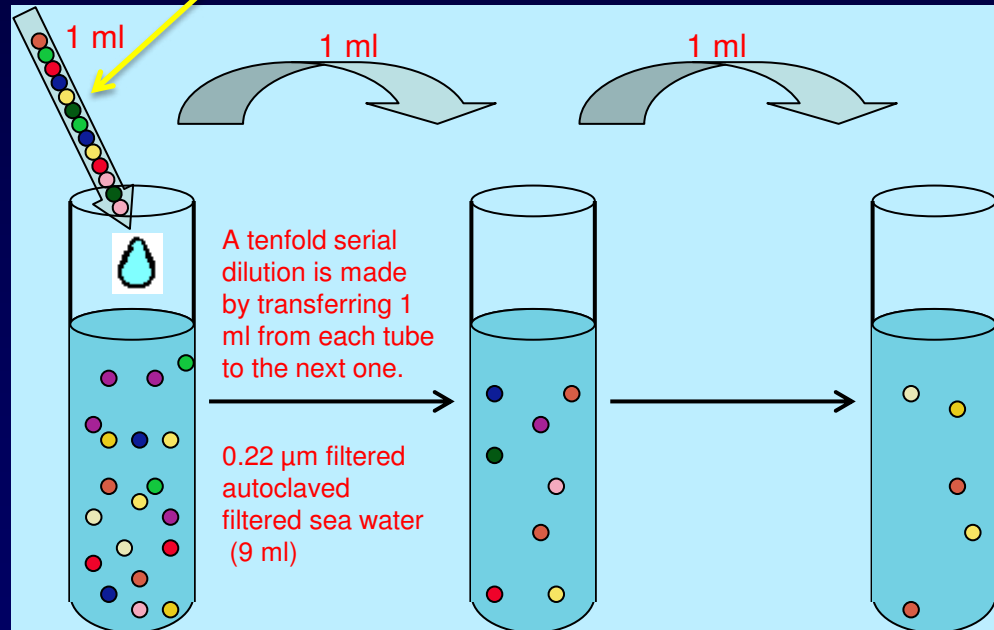


Filter through 50µm mesh

Homogenize the sample



Homogenized sample

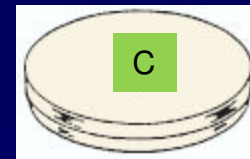
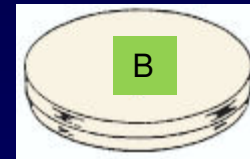
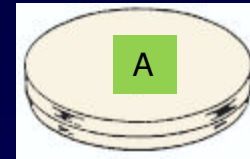


A ( $10^{-1}$ )

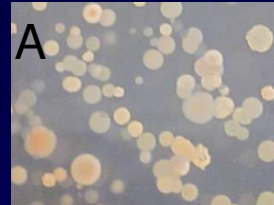
B ( $10^{-2}$ )

C ( $10^{-3}$ )

Dispense 0.1 ml



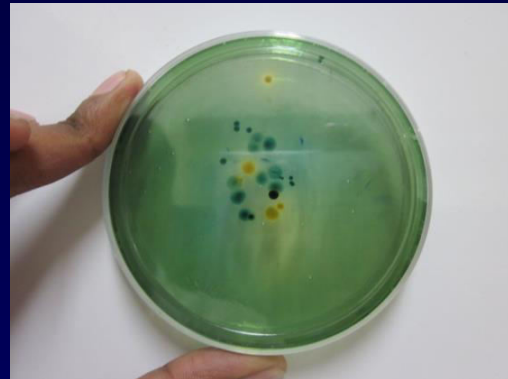
Count the colonies



# Enumeration of Total Viable Pathogenic bacterial abundance using different selective media

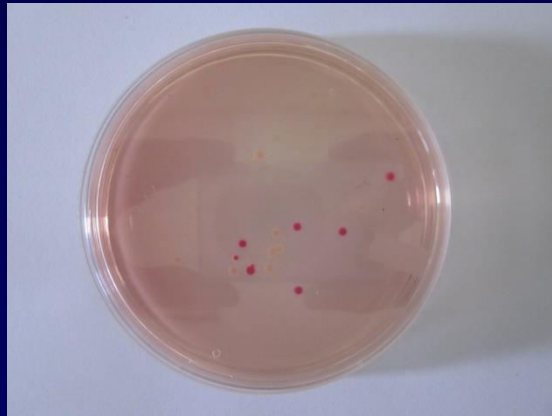
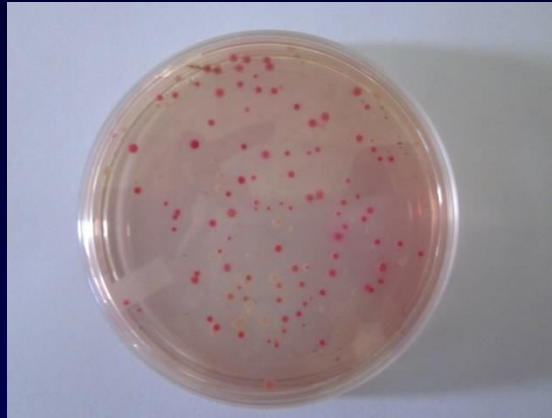
Medium	Bacteria	Observation
Thiosulfate Citrate Bile Sucrose Agar (TCBS)	<i>Vibrio cholerae</i>	Yellow colored ones, <2mm
	<i>Vibrio alginolyticus</i>	Yellow colored ones, >2mm
	<i>Vibrio parahaemolyticus</i>	Green color colonies
	<i>Aeromonas/Pseudomonas</i> spp.	Blue color colonies
MacConkey Agar	<i>Escherichia coli</i>	Pink color colonies
	<i>Salmonella/Shigella</i>	White color colonies
Enterococcus confirmatory Agar	<i>Streptococcus faecalis</i> /	Blue color colonies
	<i>Enterococcus faecalis</i>	
	General Streptococci	White color colonies
<i>E.coli</i> 0157 H:7 Agar	<i>Escherichia coli</i> 0157	Purple color colonies
	<i>Salmonella</i> serotype Enteritidis	White color colonies
Xylose Lysine deoxycholate Agar (XLD)	<i>Salmonella</i> spp.	Pink color with Black centered colonies
	<i>Shigella</i> spp.	Pink color colonies

# Thiosulfate Citrate Bile Sucrose Agar (TCBS)



Thiosulfate Citrate Bile Sucrose Agar (TCBS)	<i>Vibrio cholerae</i>	Yellow colored ones, <2mm
	<i>Vibrio alginolyticus</i>	Yellow colored ones, >2mm
	<i>Vibrio parahaemolyticus</i>	Green color colonies
	<i>Aeromonas/Pseudomonas</i> spp	Blue color colonies

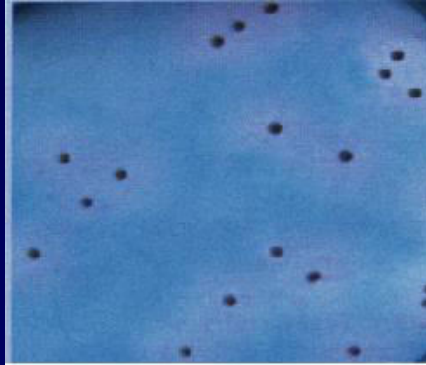
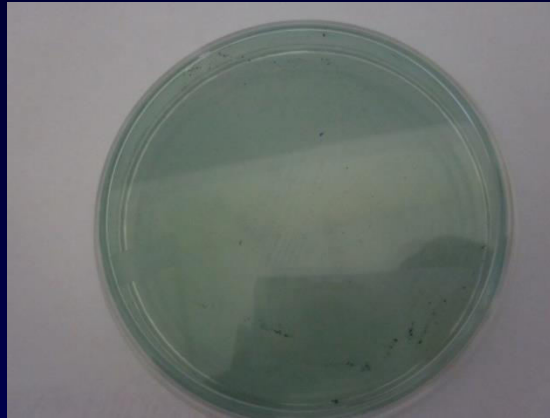
## MacConkey Agar



MacConkey Agar	<i>Escherichia coli</i>	Pink color colonies
	<i>Salmonella/Shigella</i>	White color colonies

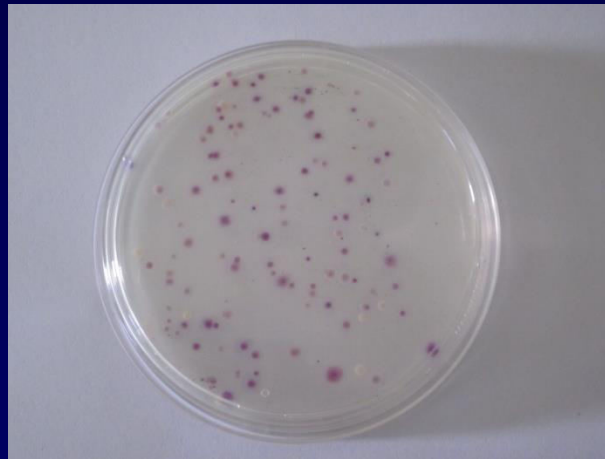
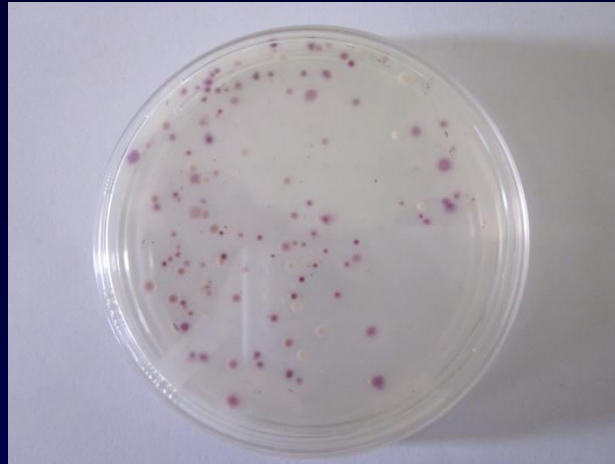


## *Enterococcus* confirmatory Agar



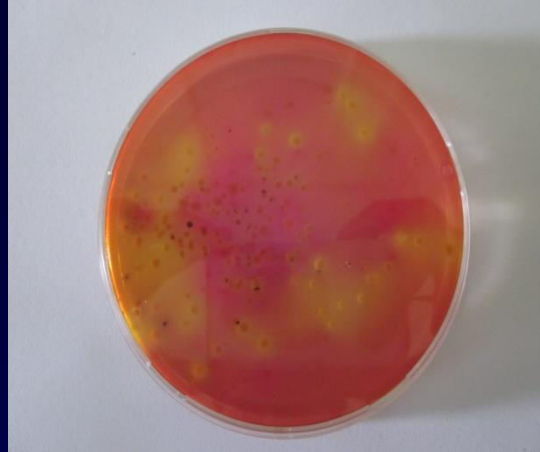
<b><i>Enterococcus</i> confirmatory Agar</b>	<b><i>Streptococcus faecalis</i>/ <i>Enterococcus faecalis</i></b>	<b>Blue color colonies</b>
	<b><i>General Streptococci</i></b>	<b>White color colonies</b>

## *E.coli* 0157 H:7 Agar



<i>E. coli</i> 0157 H:7 Agar	<i>Escherichia coli</i> 0157	Purple color colonies
	<i>Salmonella</i> serotype	White color colonies
	Enteritidis	

## Xylose Lysine deoxycholate Agar (XLD)



Xylose Lysine deoxycholate  
Agar (XLD)

*Salmonella* spp.

Pink color with Black  
centered colonies

*Shigella* spp.

Pink color colonies

# Protocol for Total Bacterial Count (TBC) using epifluorescence microscopy

Samples are immediately fixed with 0.22 $\mu$ m filtered Formalin or were stored in ice until fixed (250  $\mu$ l of formalin is added for 5 ml of sample)

↓  
1 ml of formalin fixed sample is stained with 20  $\mu$ l of Acridine orange

↓  
Incubate the sample for 10 min in dark

↓  
Filter the sample on 0.22  $\mu$ m Black Polycarbonate filter paper

↓  
Place the filter paper on a clean glass slide

↓  
Place a drop of oil on top of filter paper

↓  
Cover this with a clean glass cover slip

↓  
Observe the slide under Epi-fluorescence microscope under oil immersion

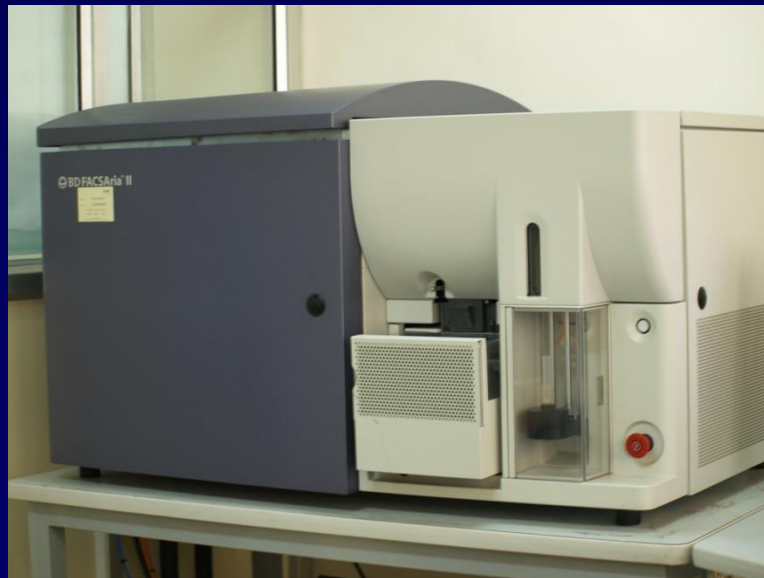
↓  
Count the number of cells in each field

↓  
Take a note of 10 different fields

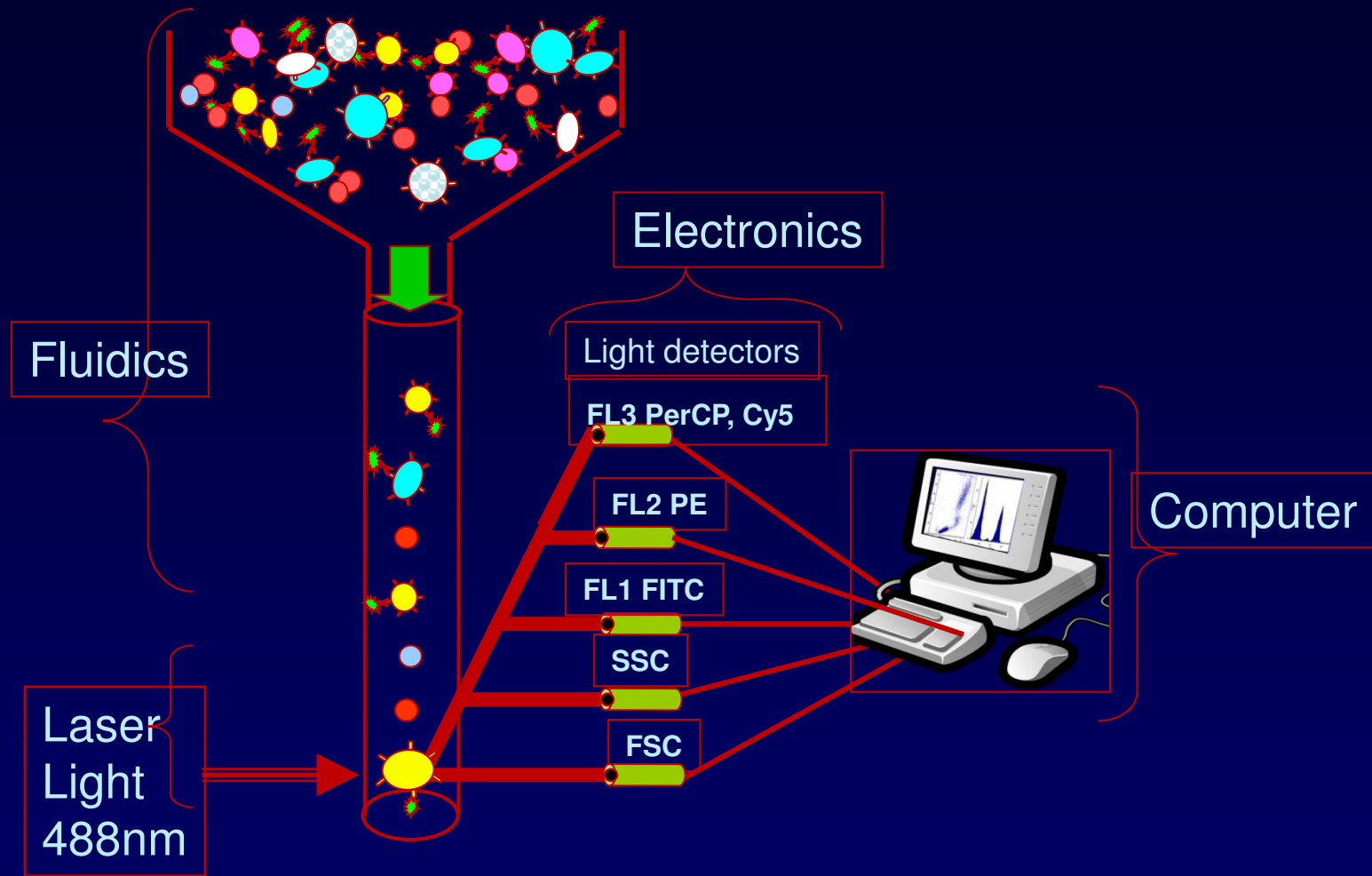


# Flow cytometric assay to quantify bacteria

- ✓ Flow cytometry is a useful tool for counting microorganisms, assessing their viability and cell sorting
- ✓ Cells are usually stained or labelled with fluorescent tags that enable them to be electronically identified when passing through a beam of laser light
- ✓ When compared to conventional culturing techniques there is also the advantage of being able to detect viable but non-culturable cells in seawater and ship's ballast water which represent a major fraction of marine bacterial species



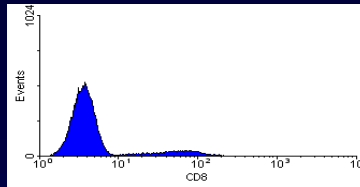
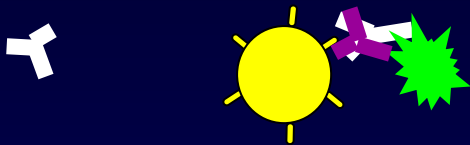
# Flow cytometer consists of



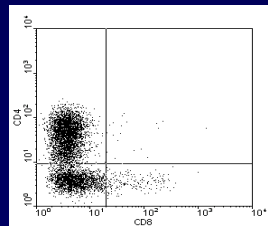
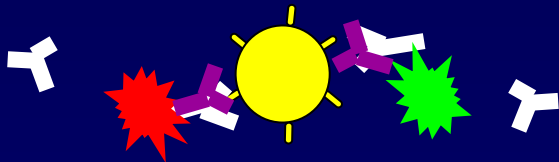


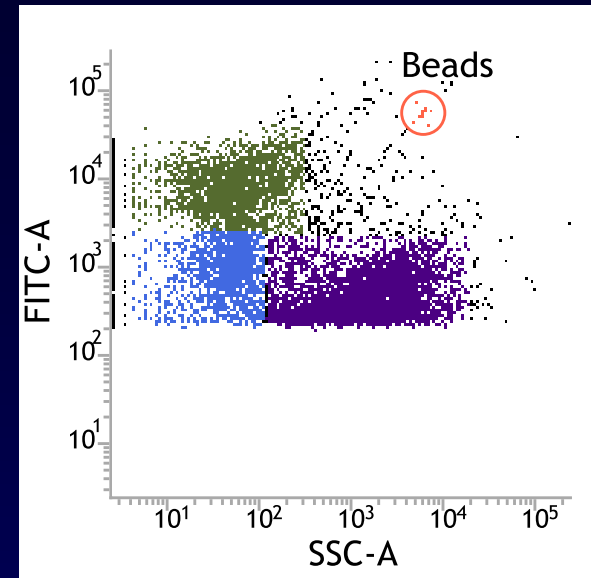
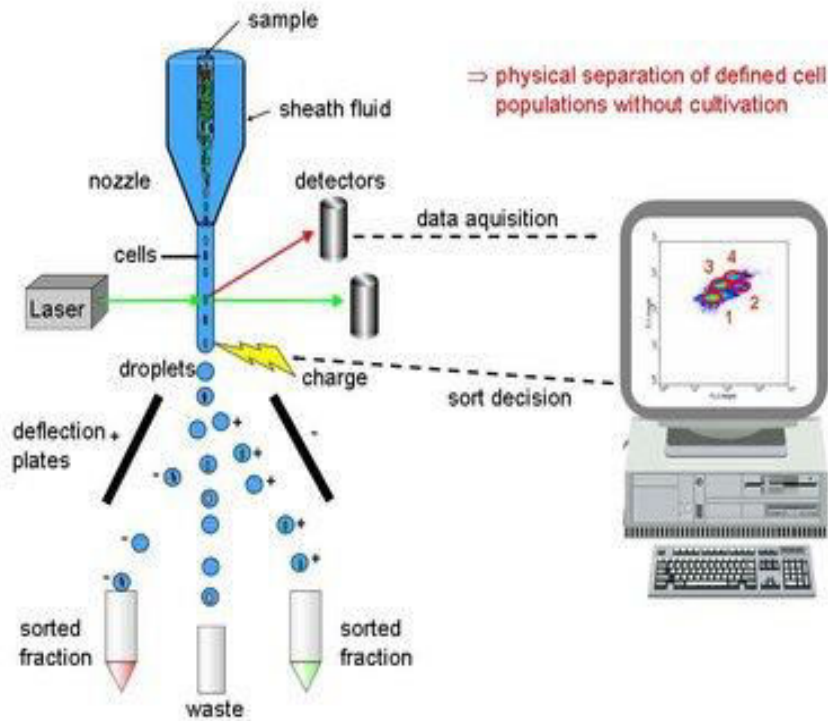
## Presenting and Interpreting Data

Single parameters can be displayed as a histogram.

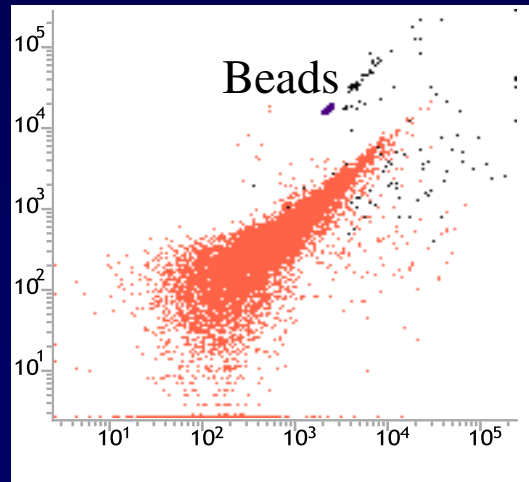


Dual parameter data can be displayed in two dimensions using dot, density or contour plots.

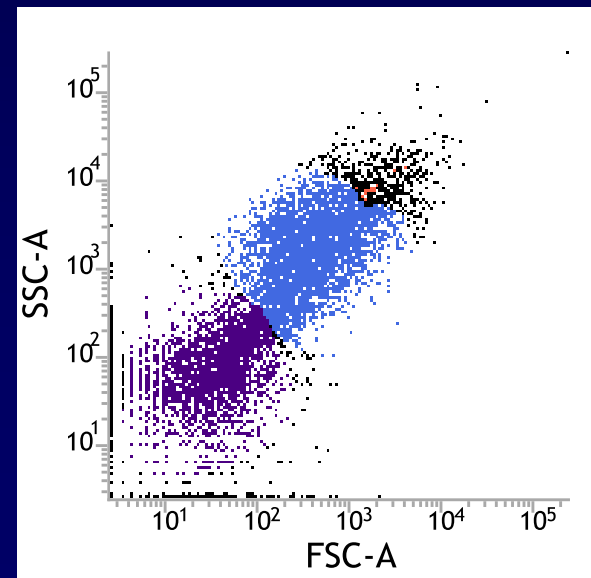




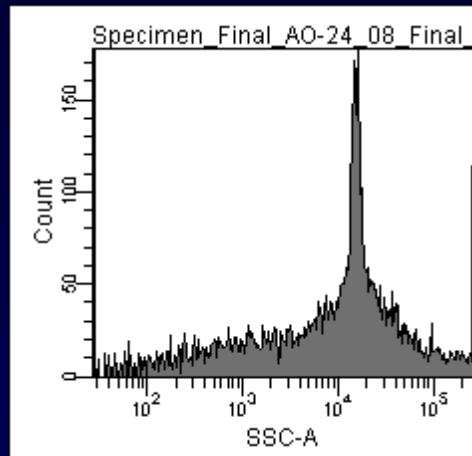
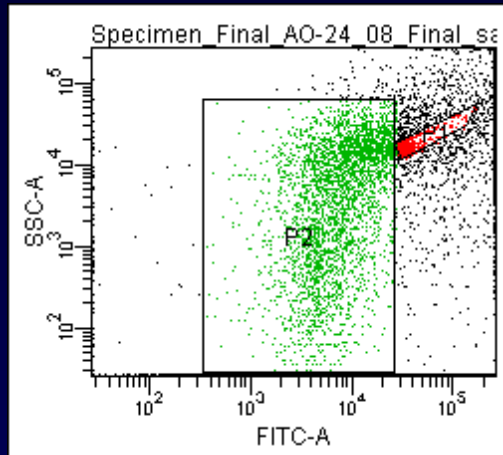
Metabolically active  
cells



Side scatter

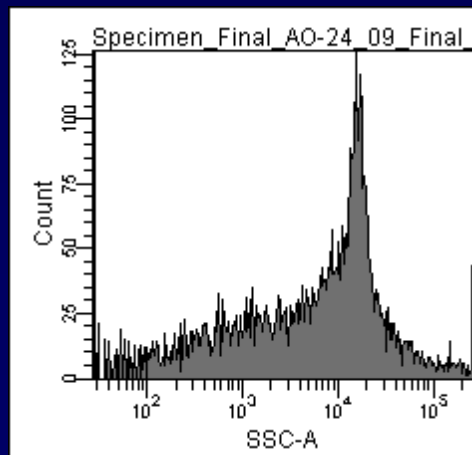
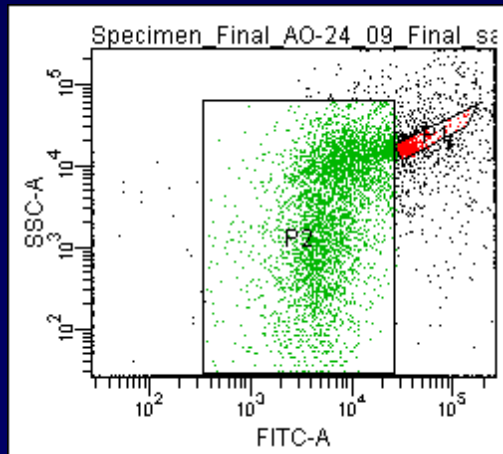


# Quantification of bacteria using Acridine Orange



Tube: 24\_08\_Final\_sampe\_BACTERIA-AO-20μL

Population	#Events	%Parent	%Total
All Events	20,000	###	100.0
P1	766	3.8	3.8
P2	3,403	17.0	17.0



Tube: 24\_09\_Final\_sampe\_BACTERIA-AO-20μL

Population	#Events	%Parent	%Total
All Events	20,000	###	100.0
P1	496	2.5	2.5
P2	3,925	19.6	19.6

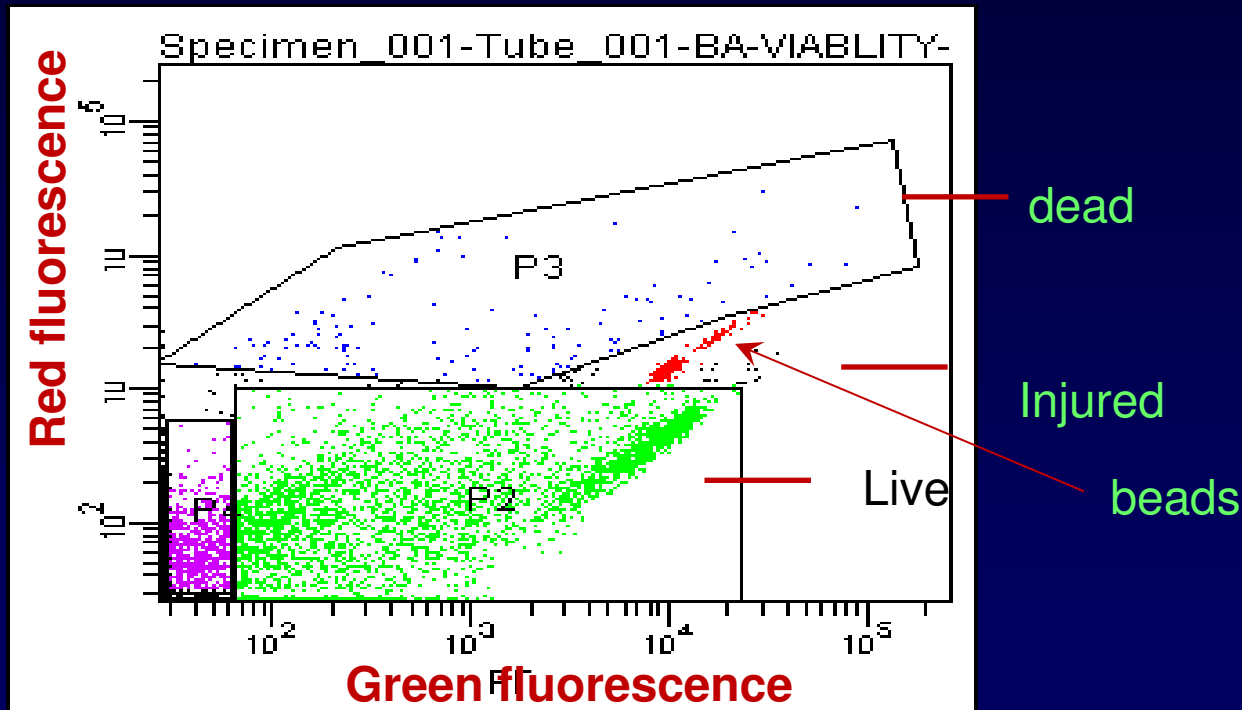
# Flow cytometric assay to assess viability

Thiazole Orange

→ To discriminate live and dead cells

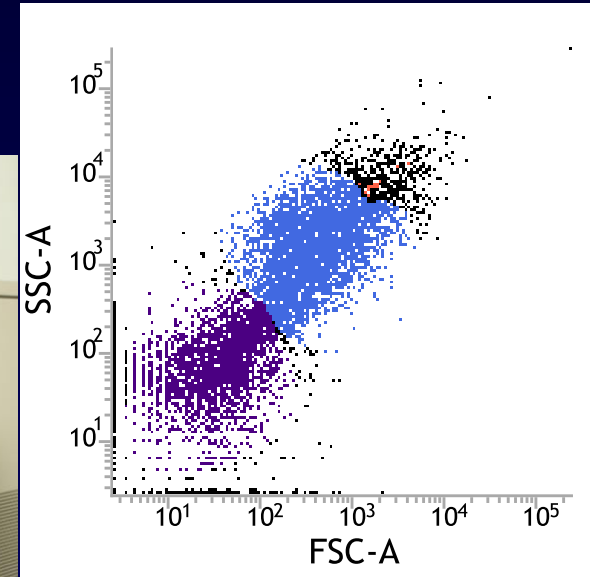
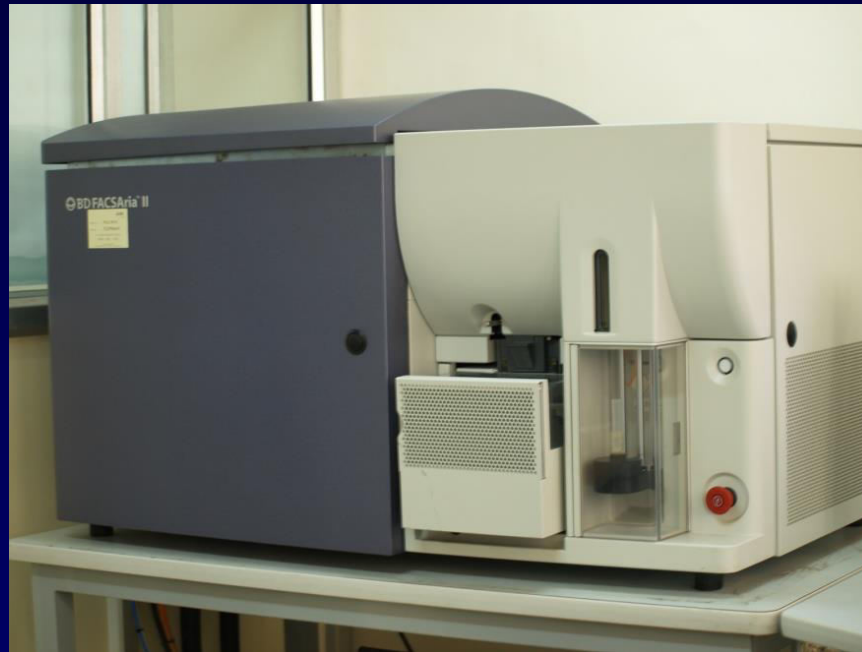
Propidium Iodide

Flow cytometry which offers the ability to rapidly distinguish between active and compromised or dead cells is a valuable tool to study the effect of environmental variables



Nucleic acid double staining (with SYTO9/ Propidium iodide) which differentiates active, live cells from inactive, dead cells

# FISH-FCM method for rapid estimation of bacterial communities with special reference to *Vibrio cholerae*



# Flow cytometry with FISH

- ✓ Flow cytometry combined with FISH is also an increasingly popular method of enumerating cells in environmental samples
- ✓ The oligonucleotide probes used for FISH are dependant upon the micro-organism which is to be identified.

Specific probe for *Vibrio* spp.

*Vibrio* spp.

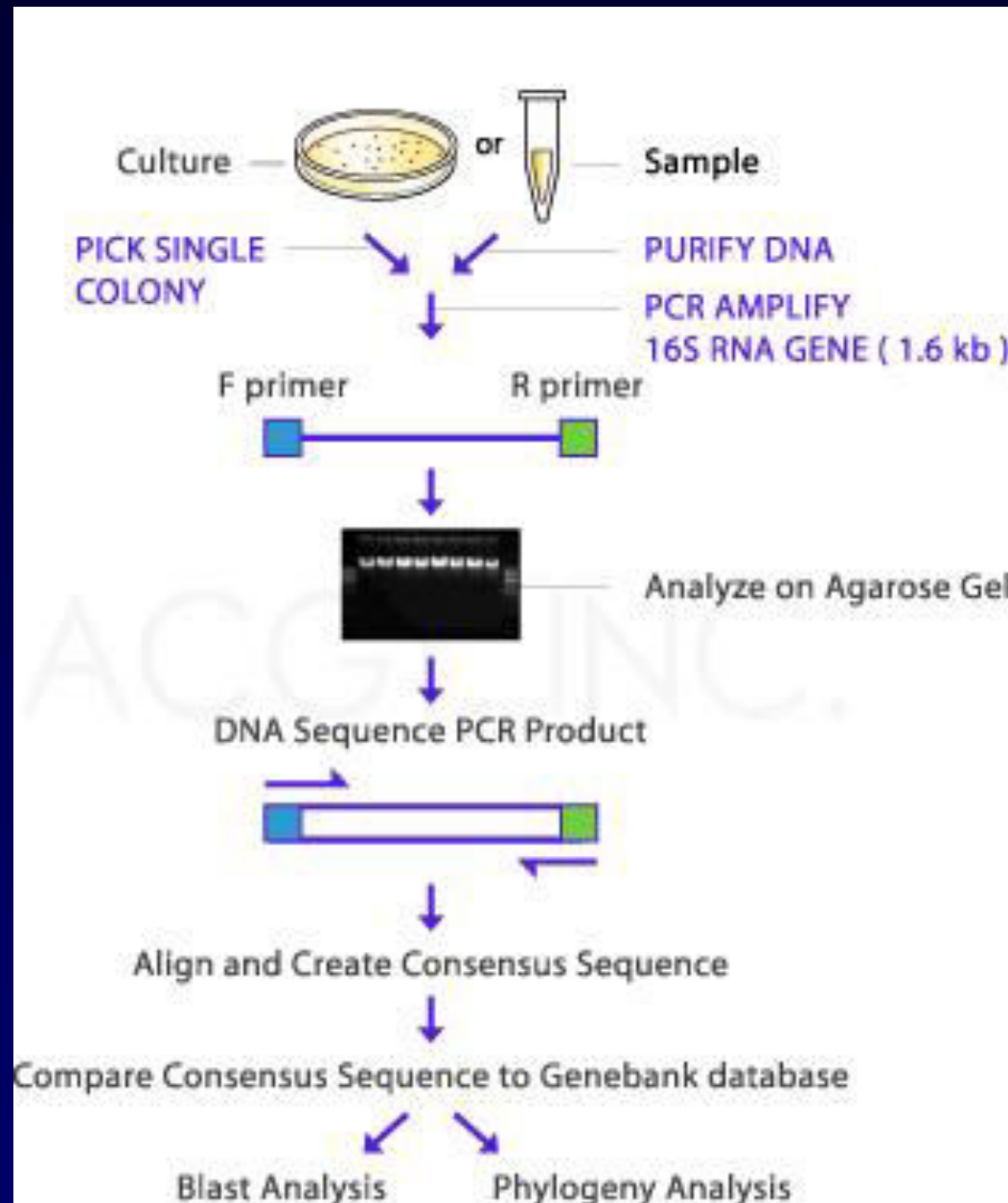
AGGCCACAACCTCCAAGTAG

Cy3-AGGCCACAACCTCCAAGTAG

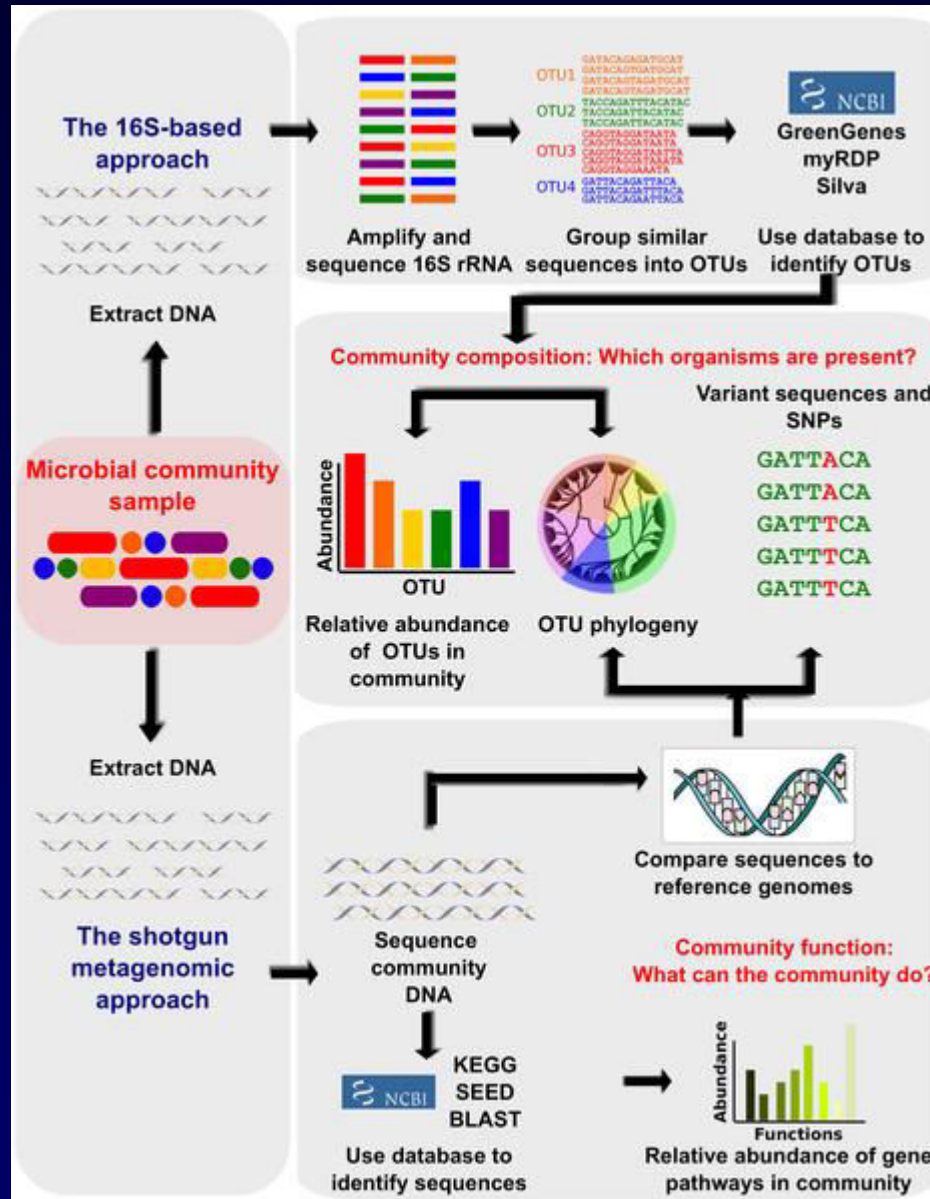
T<sub>m</sub> (°C) 59.4 Eilers et al. (2000b)



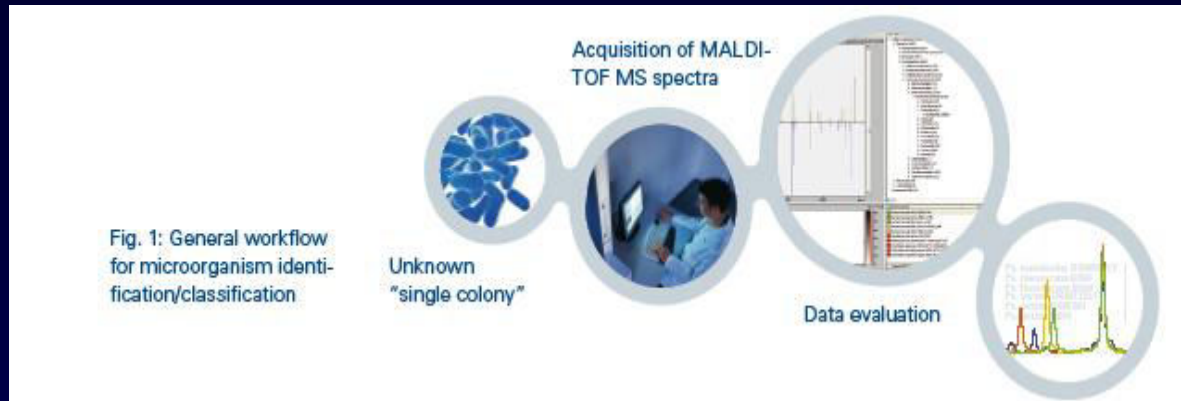
# Phylogenetic Analysis



# Metagenomics Approach



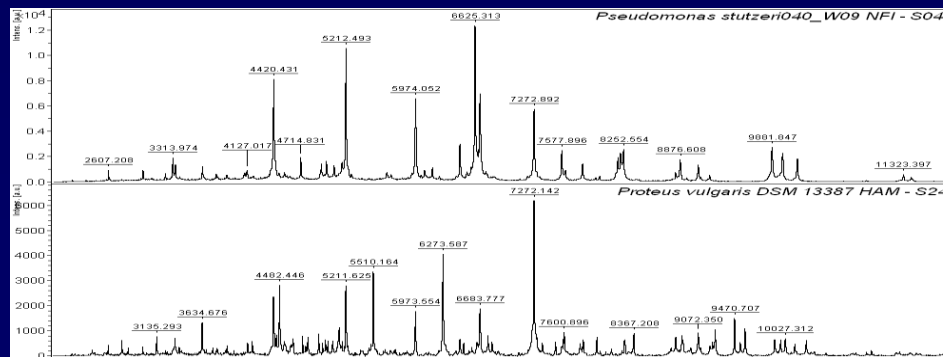
# Microorganism Identification and Classification Based on MALDI-TOF MS Fingerprinting with MALDI Biotyper



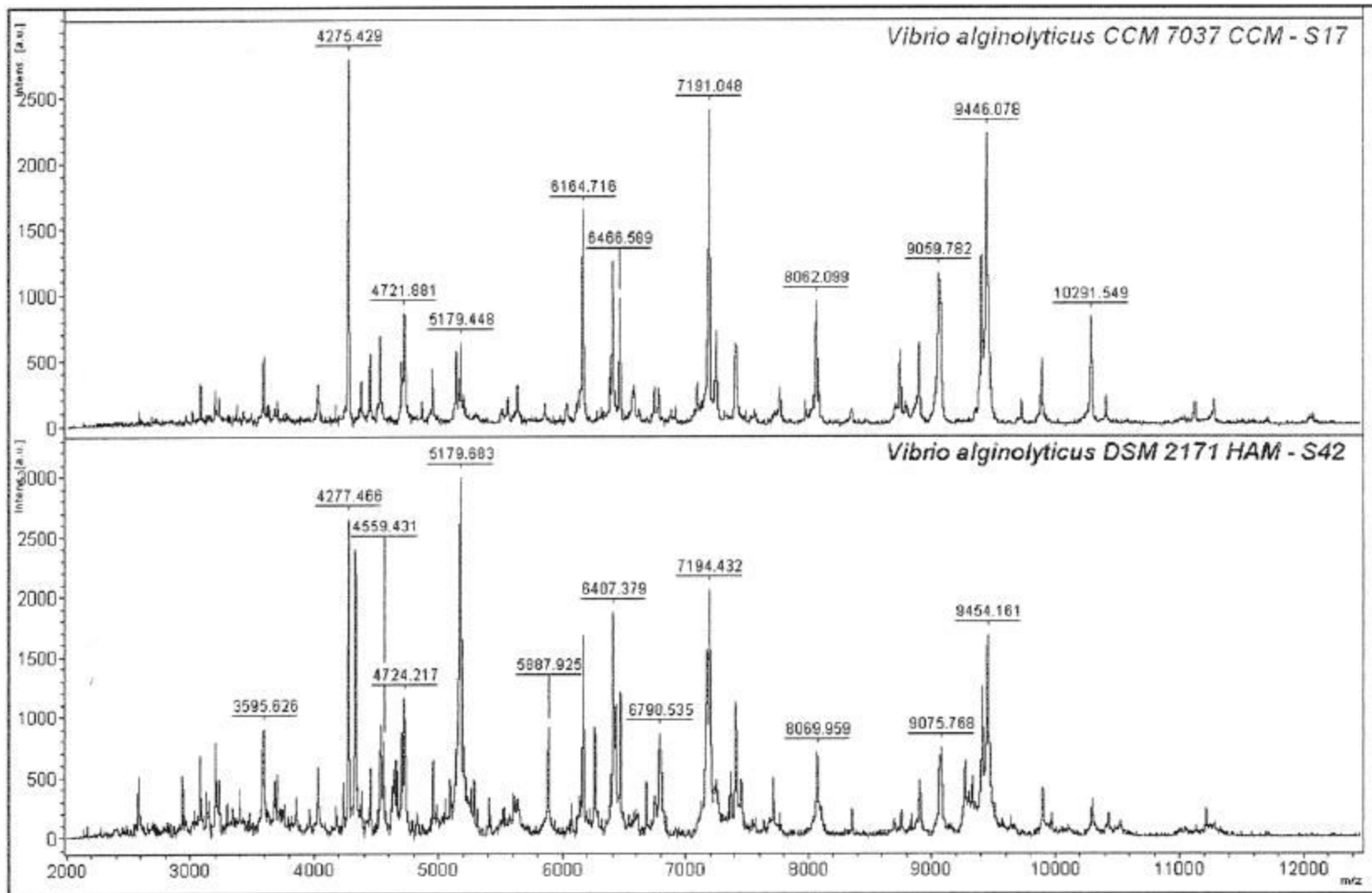
MALDI-TOF MS



Spotting of bacterial proteins on MALDI plate



MALDI-TOF MS spectra



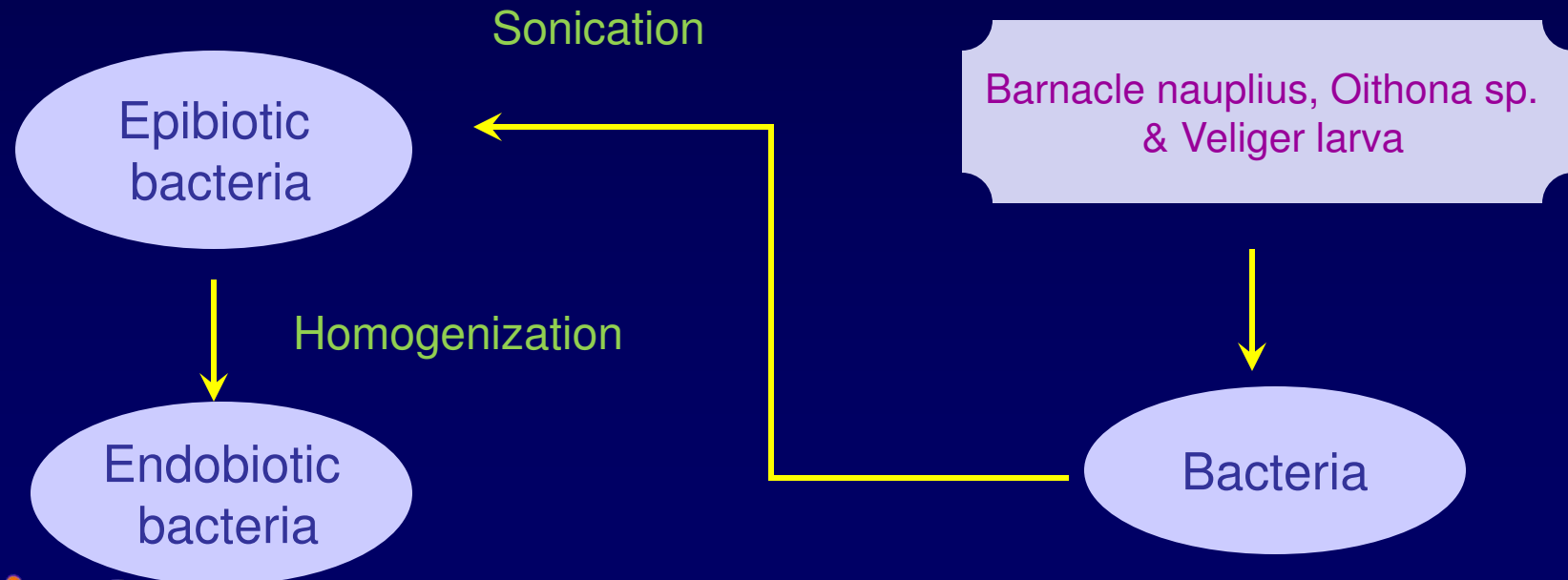
MALDI-TOF mass spectra of *V. alginolyticus* distinguishable by their protein fingerprints

Source: Kaveh E, Vahid A, Matthias U, Khwajah M, AC Anil, Lidita Khandeparker, Burgess JG, Mesbahi E, Plos One ( 2012)

# Larvae as hotspots of bacterial diversity – a case study

- ❖ Marine invertebrate larvae offer extensive body surfaces for attachment of bacteria, their decomposition would contribute to increased bacterial production in the marine environment
- ❖ Microbial decomposition of larval carcasses serves as an alternate mechanism for nutrient regeneration, elemental cycling and microbial production.

# Protocol





# Conclusions

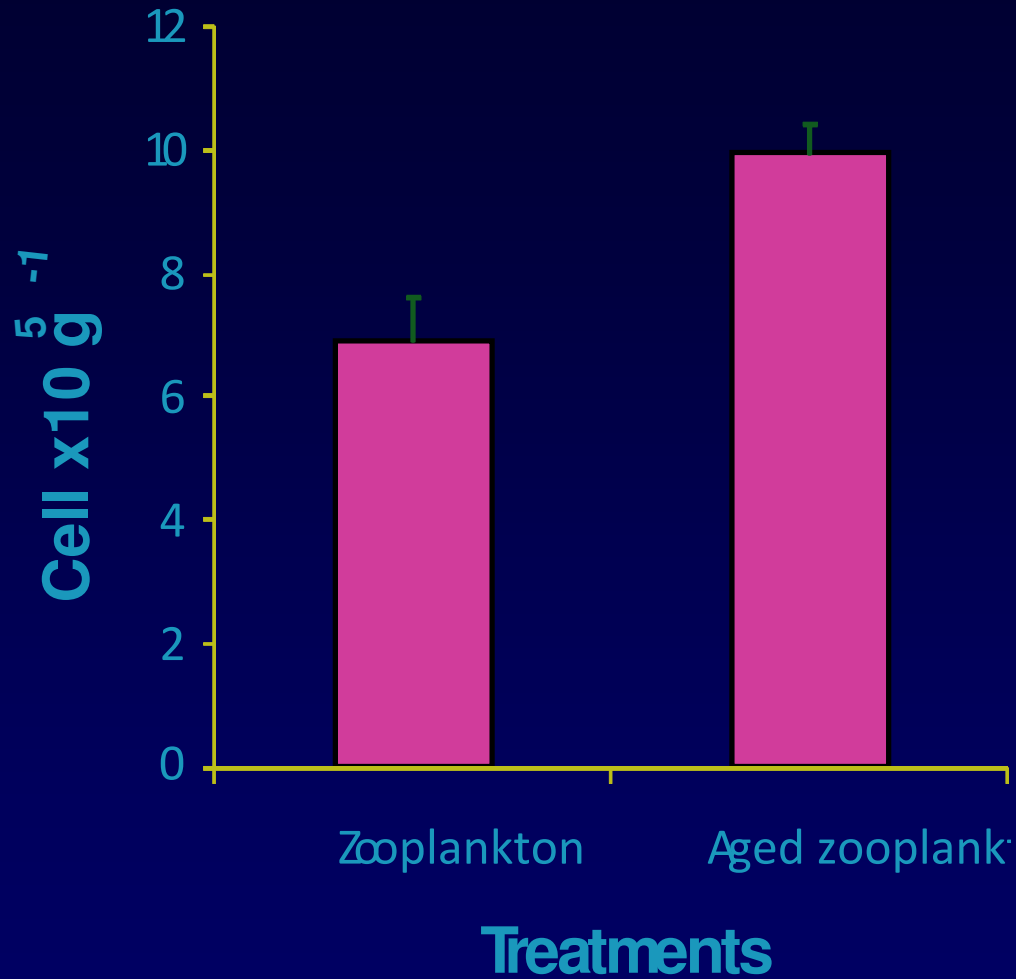
- ✓ Barnacle nauplius and veliger larva harbored  $\sim 4.4 \times 10^5$  cells ind<sup>-1</sup> whereas *Oithona* sp. had  $8.8 \times 10^5$  cells ind<sup>-1</sup>
- ✓ Computation of bacterial contribution based on bio-volume indicated that veliger larva, smallest form of the lot harbored highest numbers of bacteria and barnacle nauplius with largest bio-volume contributed the least

# List of epibiotic and endobiotic bacteria specific to different invertebrate forms

Type of organisms	Epibiotic bacteria	Endobiotic bacteria
Barnacle nauplii	<i>Aeromonas salmonicida salmonicida</i>	<i>Bacillus brevis</i>
	<i>Aeromonas schubertii</i>	
	<i>Photobacterium angustum</i>	
Copepod	<i>Aeromonas schubertii</i>	<i>Actinobacillus capsulatus</i>
		<i>Bacillus brevis</i>
Bivalve larva	<i>Vibrio fischeri</i>	<i>Aeromonas schubertii</i>
	<i>Photobacterium angustum</i>	

Certain degree of form specificity was observed in case of epibiotic and endobiotic bacteria of each of the tested invertebrate forms

# Bacterial Abundance on aging



# Pathogenic bacteria associated with the zooplankton

Pathogens	Initial (CFU g <sup>-1</sup> )	After pulverization (CFU g <sup>-1</sup> )
<i>Streptococcus faecalis</i>	$2.1 \times 10^2 \pm 4.9 \times 10^1$	$2.5 \times 10^5 \pm 3.8 \times 10^4$
<i>Vibrio cholerae</i>	$3.5 \times 10^3 \pm 1.4 \times 10^3$	$5.4 \times 10^5 \pm 3.1 \times 10^5$
<i>Escherichia coli</i>	$5.0 \times 10^2 \pm 2.8 \times 10^2$	$1.3 \times 10^4 \pm 7.6 \times 10^3$

- ✓ The numbers of the pathogenic bacteria increased enormously subsequent to zooplankton pulverization
- ✓ *V. cholerae* which was initially  $3.5 \times 10^3$  increased to  $5.4 \times 10^5$  CFU g<sup>-1</sup> following pulverization of zooplankton. Similar increase was also observed in *E. coli* and *S. faecalis*

# Conclusions

- ✓ After aging of pulverized zooplankton, *Chromobacterium violaceum*, an opportunistic pathogen in animals and humans emerged
- ✓ Association of pathogenic bacteria in vast numbers with the zooplankton is a cause of concern once they are destroyed or dead

# Conclusions

Bacterial numbers were highest in the pre-monsoon season at the Fishing harbour in surface and bottom water and sediments

Lowest bacterial abundance was observed in the sediments of outer port stations in the monsoon and offshore station in the post-monsoon season

# Thank you

