



université  
PARIS-SACLAY

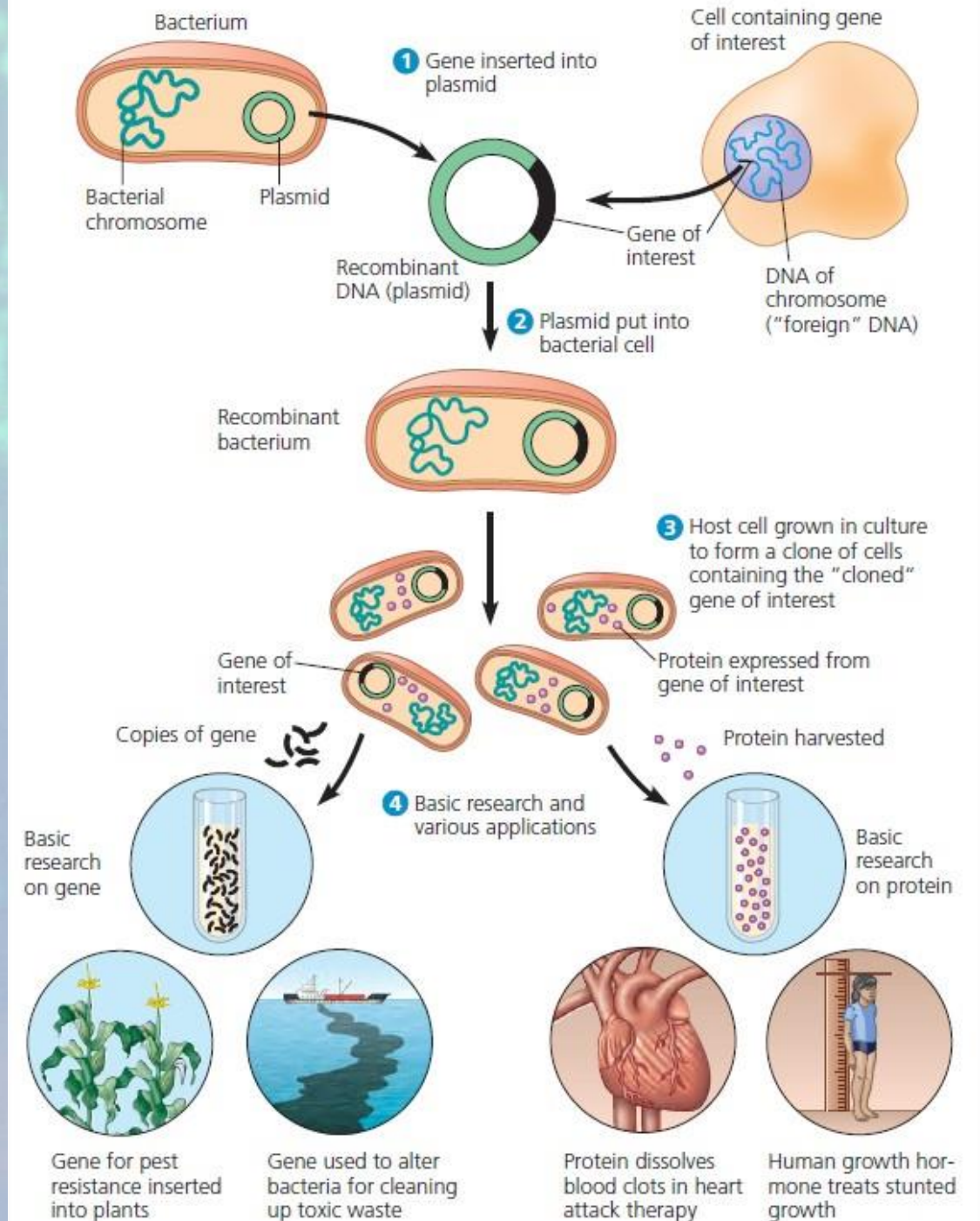
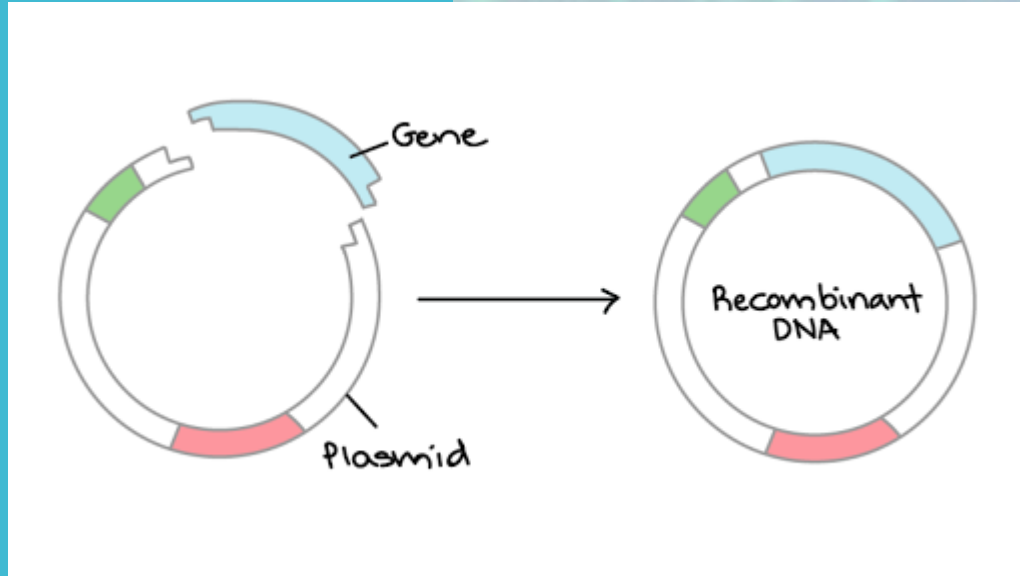
# Introduction to Gene cloning Practicals

Master Development of Drugs and Health Products

November 2024

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# Context and applications



# General objectives

- Learn the lab good practices + Work safely in a lab
- Know how to perform each experiment:
- Think about organisation to optimise working time (retro-planning, running 2 experiments in parallel, anticipation from one experiment to the other,...)
- Perform a full experimental strategy
- Write-up a protocol

# Specific objective

- To produce a strain of *E. coli* expressing a His-tagged GFP protein

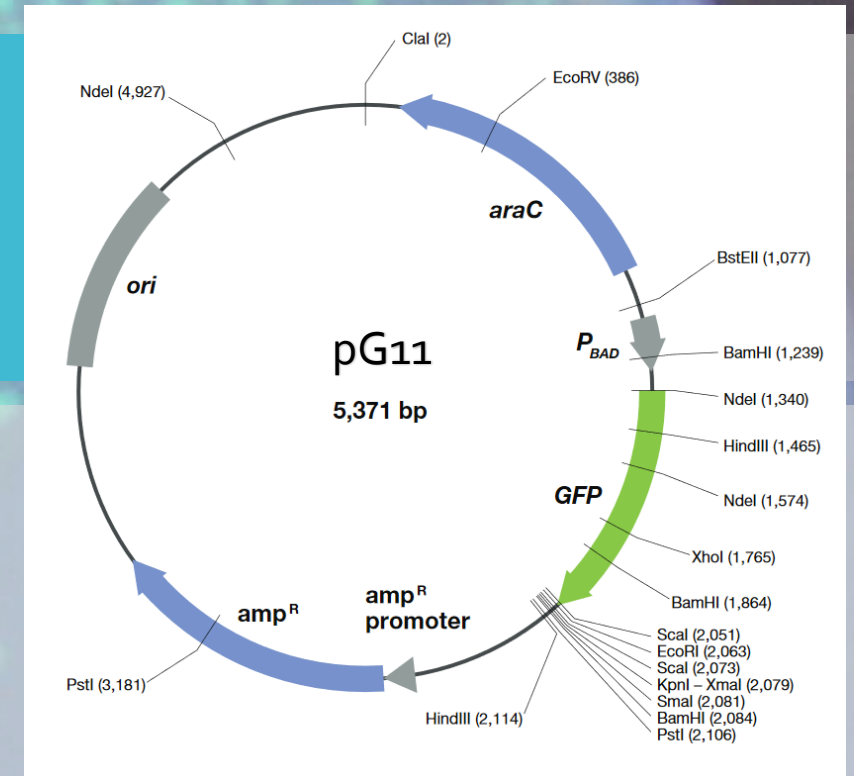


# Background



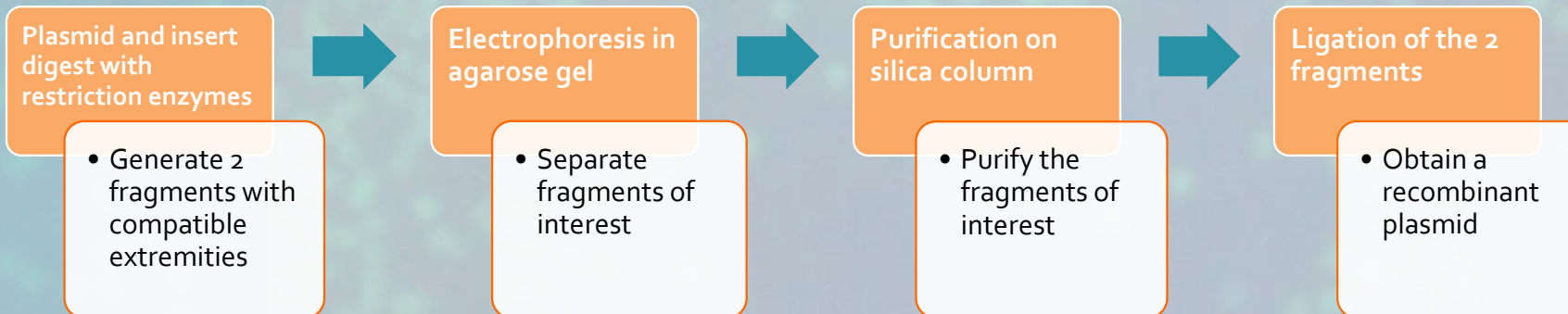
*Aequorea victoria*

- Why GFP?
- Bacterial strain: DH5- $\alpha$  (*endA recA*)
- Plasmid
- Cloning of a 6xHis encoding sequence fused to the GFP gene (! reading frame)  
=> replace the WT GFP-encoding gene by the 6xHis-tagged GFP encoding gene
- Induction

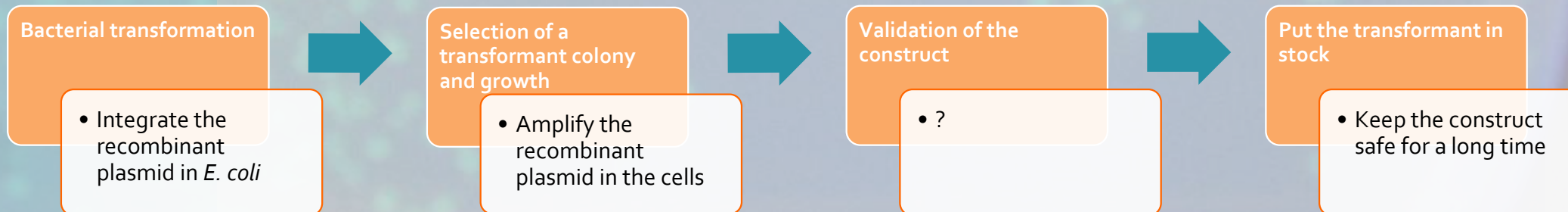


# Workflow

## Construction

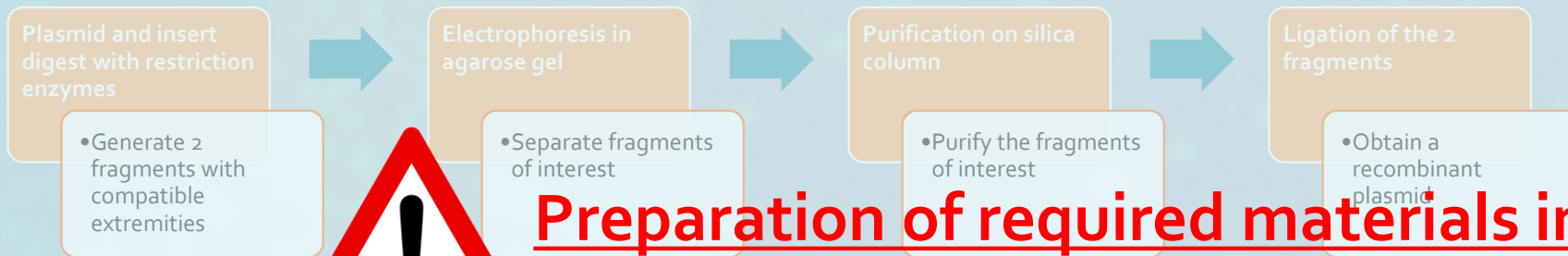


## Amplification and validation



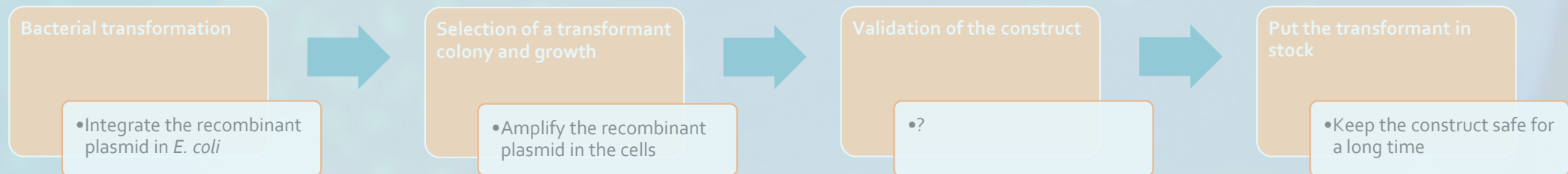
# Workflow

## Construction



**Preparation of required materials in advance**  
**=> anticipate**

## Amplification and validation





# Preparation of your experimental work

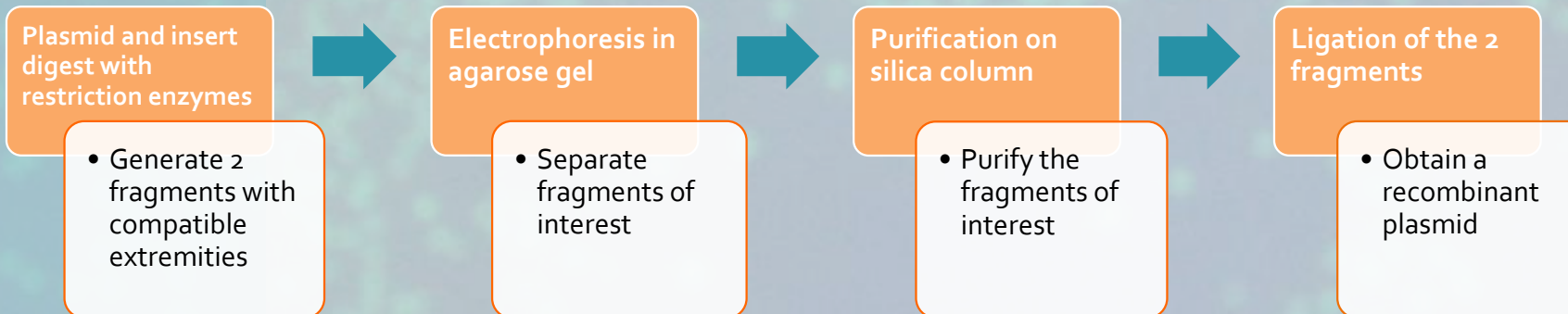
## To do before the lab course:

- Protocols: understand each step and order
  - Role of each reactive/component at each step
  - Search for and understand technical data (DNA sequence analysis, ligation conditions, agarose %, ...)
- Controls
- Calculations + DNA sequence analysis to make
- Working conditions (sterility, safety equipment, temperature, ...)
- Safety issues -> MSDS (Material Safety Data Sheet)
- Waste management
- Required materials and consumables (anticipation)
- See specific questions in your documents

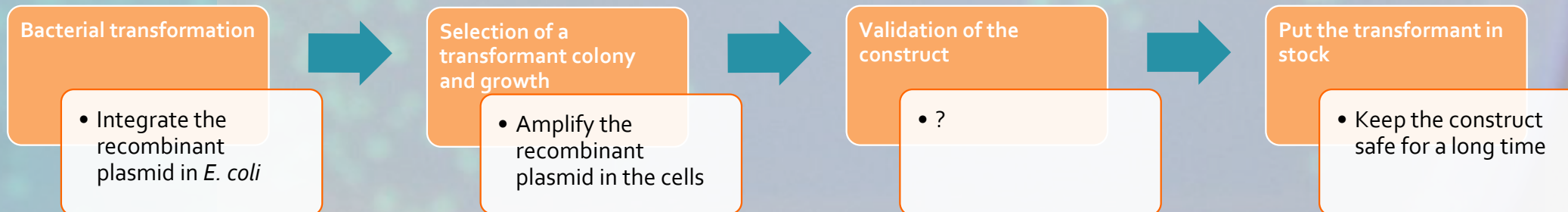


# Workflow

## Construction



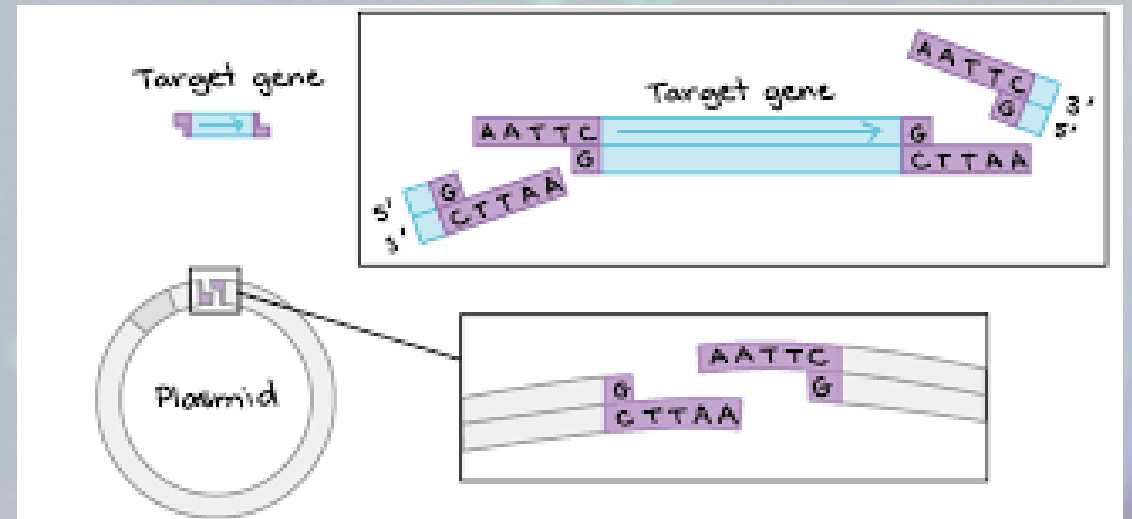
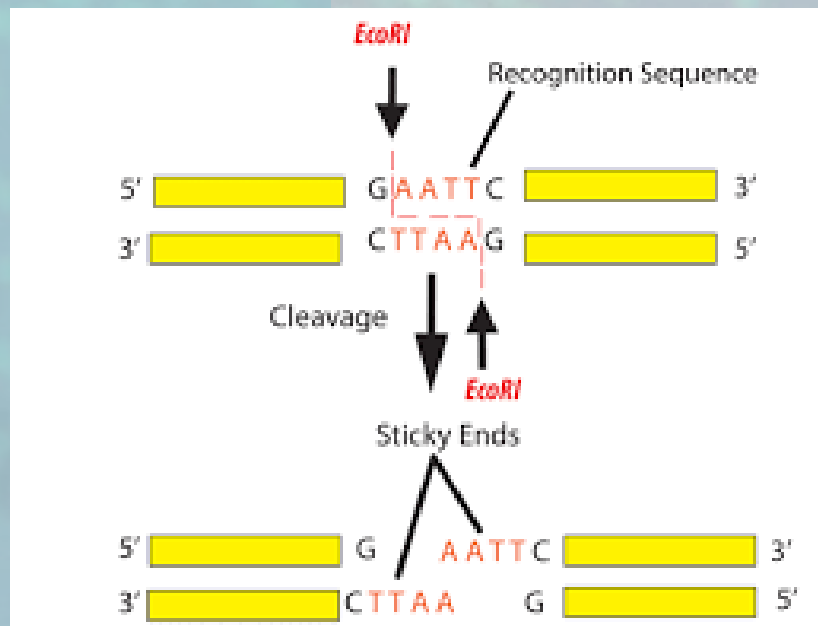
## Amplification and validation



# Plasmid and insert digest

## Principle

- Restriction enzymes



# Plasmid and insert digest Reaction

## Restriction Enzyme Single/Double Digestion

EcoRI-HF

Select 2nd Enzyme

✕ clear 2nd selection

Digest in rCutSmart Buffer

Show Detailed Protocol

Name	Cat #	Temp °C	Supplied Buffer	Add SAM	% Activity in NEBuffer™			
					r1.1	r2.1	r3.1	rCutSmart
EcoRI-HF®	R3101	37	rCutSmart Buffer	No	10	100	10	100

	pG11 plasmid	Insert
Ultra-pure H <sub>2</sub> O	qsf 20 µl	qsf 20 µl
Restriction buffer (10 X): <b>YOU MUST DETERMINE WHICH ONE IN ADVANCE</b>	1 X	1 X
DNA (pG11 @250 ng/µL; Insert @25 ng/µL)	500 ng	150 ng
EcoRI-HF (5,000 U/ml)	5 U	5 U
NheI-HF (5,000 U/ml)	5 U	5 U

- ! Order of the reagents for the mix + temperature
- Which buffer? Sequence & length of produced fragments?
- Volume of each reagent ?

→ NEB cutter tools :

<https://nebcloner.neb.com/#!/redigest>

<http://nc2.neb.com/NEBcutter2/>

Enter a DNA sequence, or select from other options, to identify cut sites. Once you submit a sequence, you may choose to customize your digest.

### 1. Input or choose sequence. ⓘ

Text File GenBank Plasmid Vector Viral & Phage

Type or paste sequence

### 2. Set preferences. ⓘ

Circular  Additional Preferences (enzymes, oligos, etc)

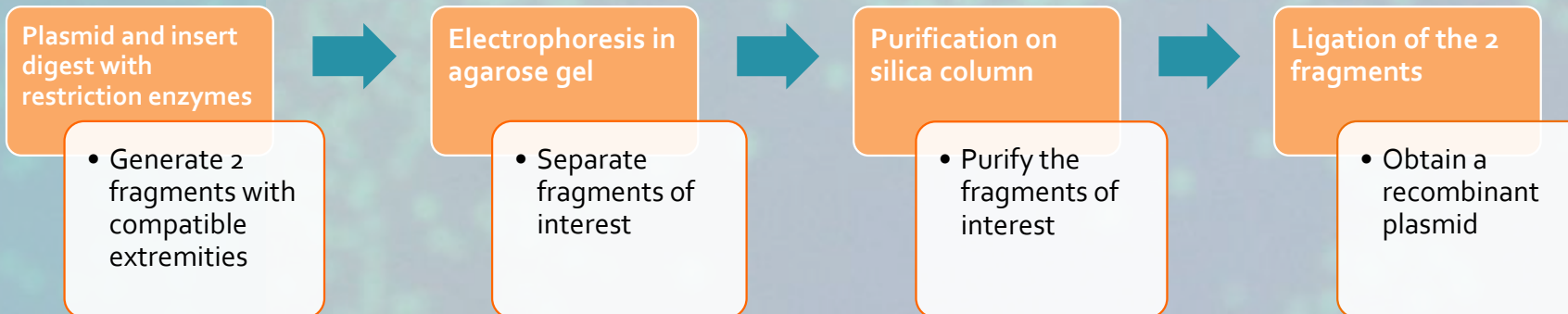
### 3. Name project (optional). ⓘ

Enter project name

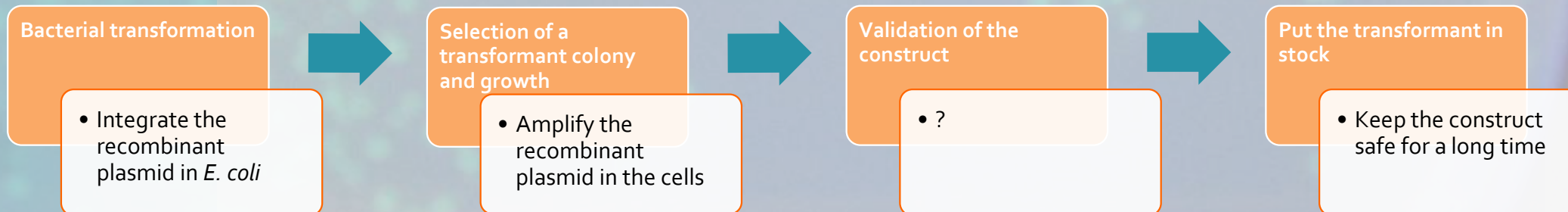
Submit

# Workflow

## Construction



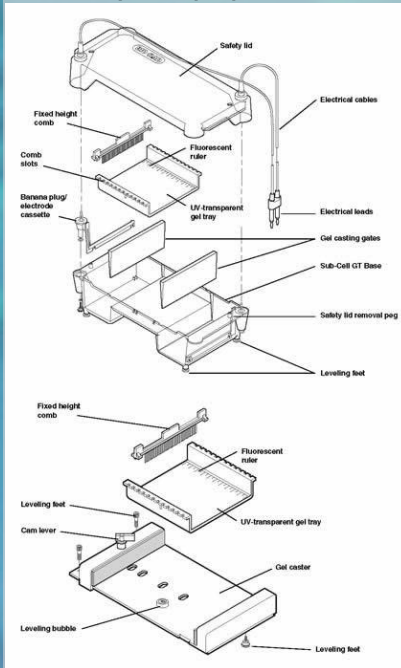
## Amplification and validation



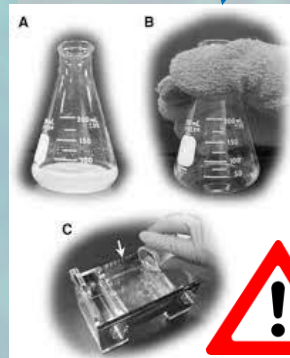


# Electrophoresis in agarose gel

## 1. Gel casting system preparation

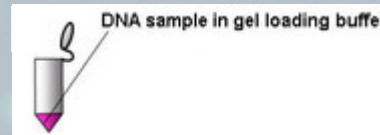


## 2. Agarose gel preparation



EtBr  
= CMR

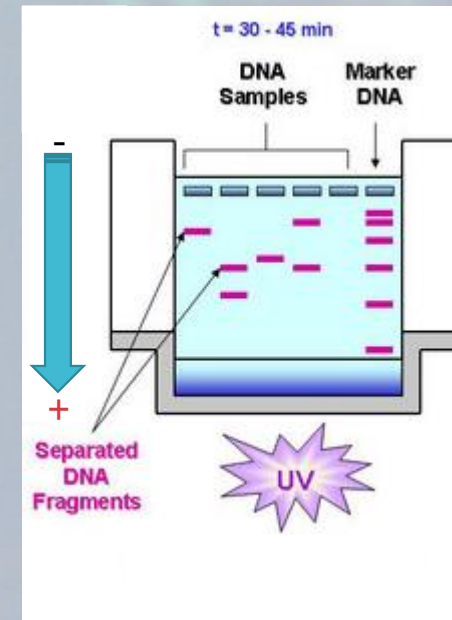
## 3. Addition of loading buffer & sample loading



## 4. Setting up connection & start migration



## 5. Results observation



# Electrophoresis in agarose gel

sample preparation for loading

<b>Sample of digested plasmid or insert</b>	500 ng pG11 or 150 ng insert
<b>Loading buffer (6 X or 10 X)</b>	1 X final
<b>H<sub>2</sub>O UP (only if necessary)</b>	qsf 24 µl

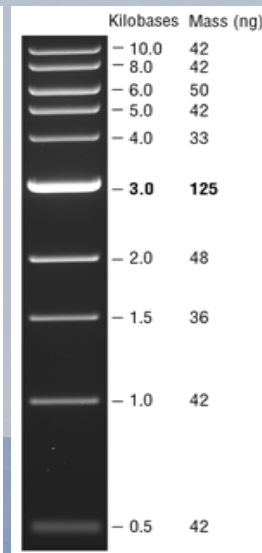
1. Which volume of loading buffer and H<sub>2</sub>O?

2. Which DNA ladder?

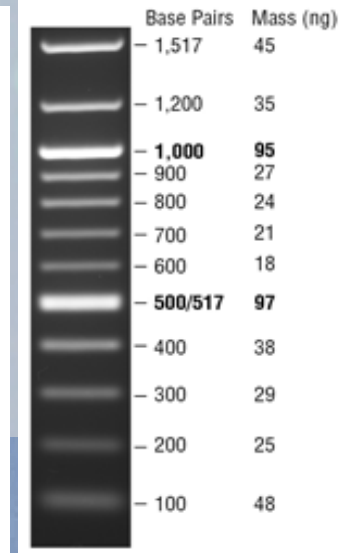
1 kb Plus DNA Ladder



1 kb DNA Ladder

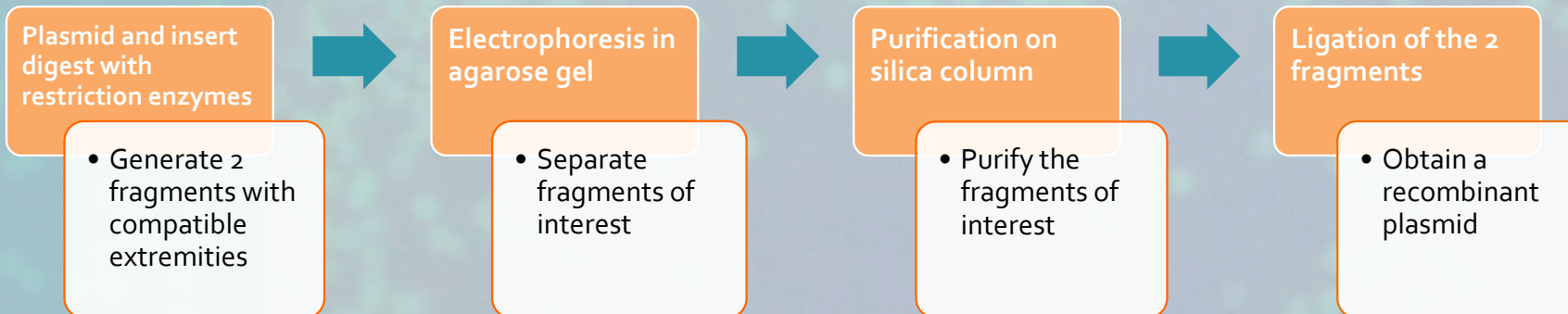


100 bp DNA Ladder

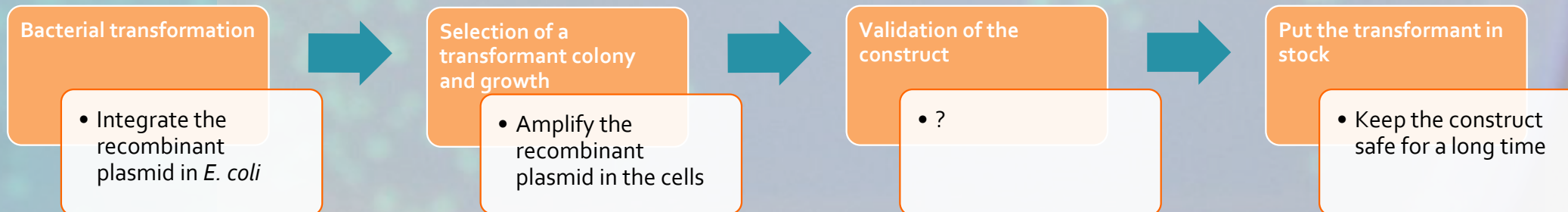


# Workflow

## Construction



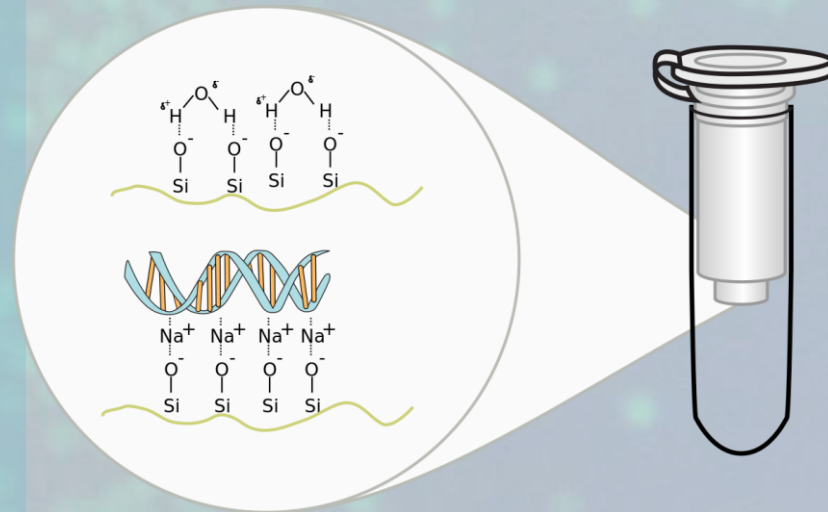
## Amplification and validation





# Purification on silica column

## Materials



DNA adsorbs to silica matrices in high-salt buffer, at optimal pH.



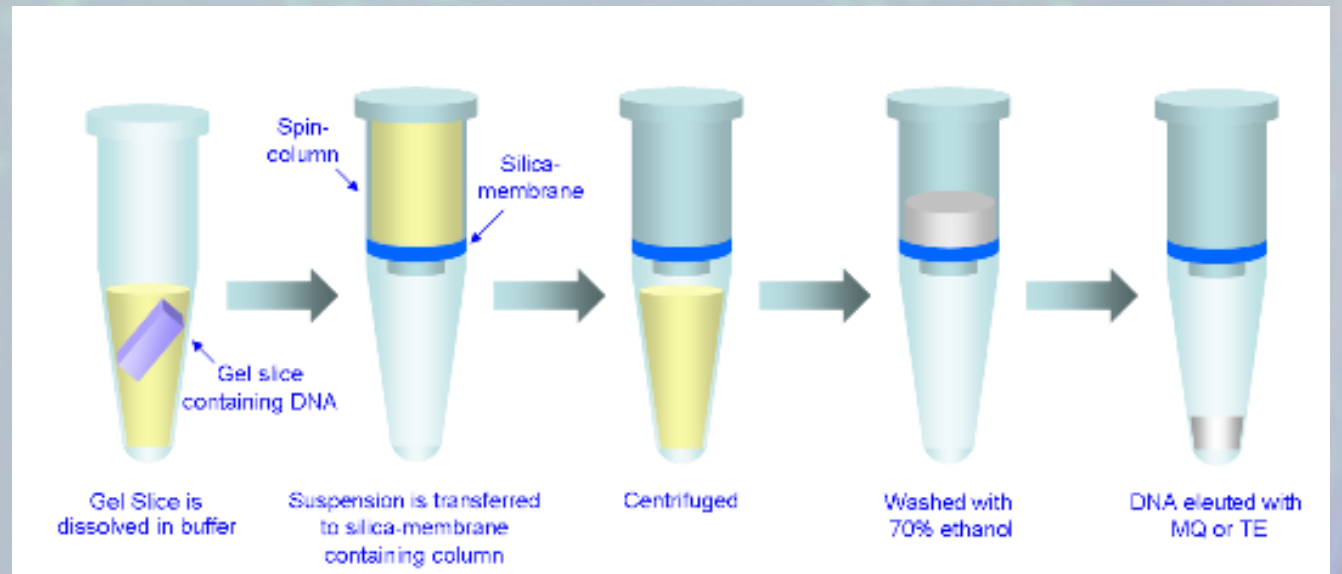


# Purification on silica column

## Procedure overview



Eye protection against burn  
= wearing an anti-UV mask



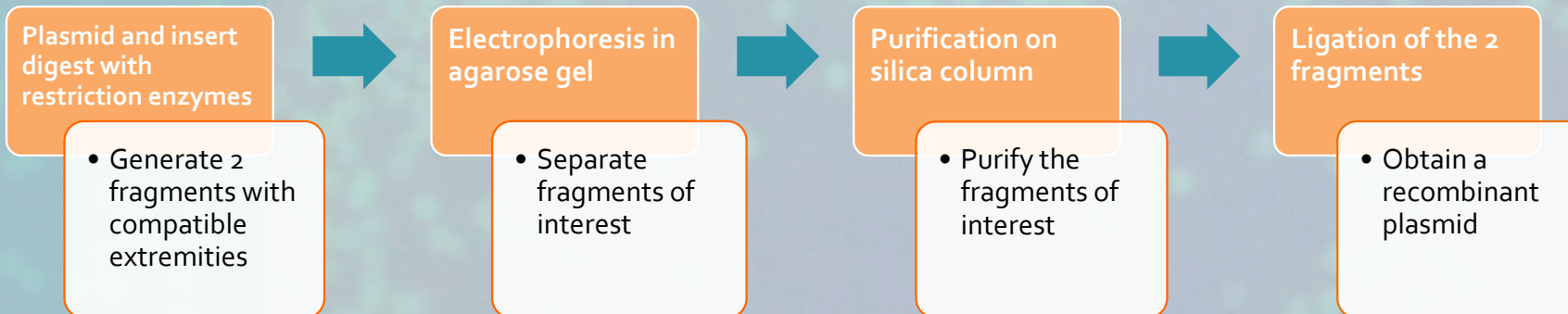
# Purification on silica column

Anticipate expected results

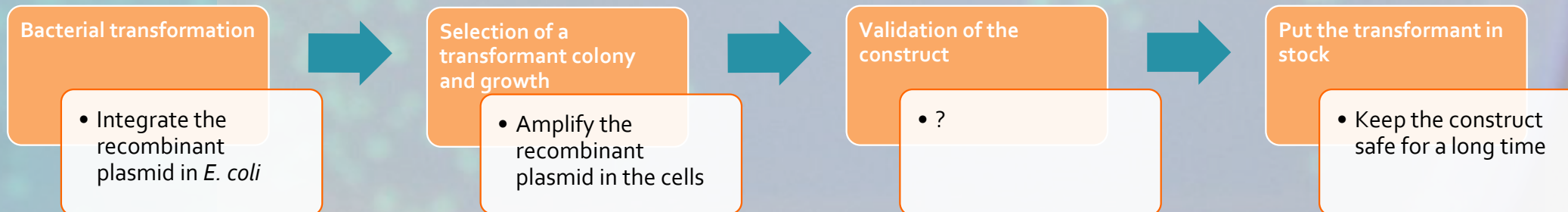
1. What is the expected size (length) and sequence of the 2 DNA fragments obtained after digest?
2. Why and how are they able to ligate to each other?
  - Tool recommended for sequence analyses: APE free software
  - !You must consider all the possibilities of ligation that can happen

# Workflow

## Construction



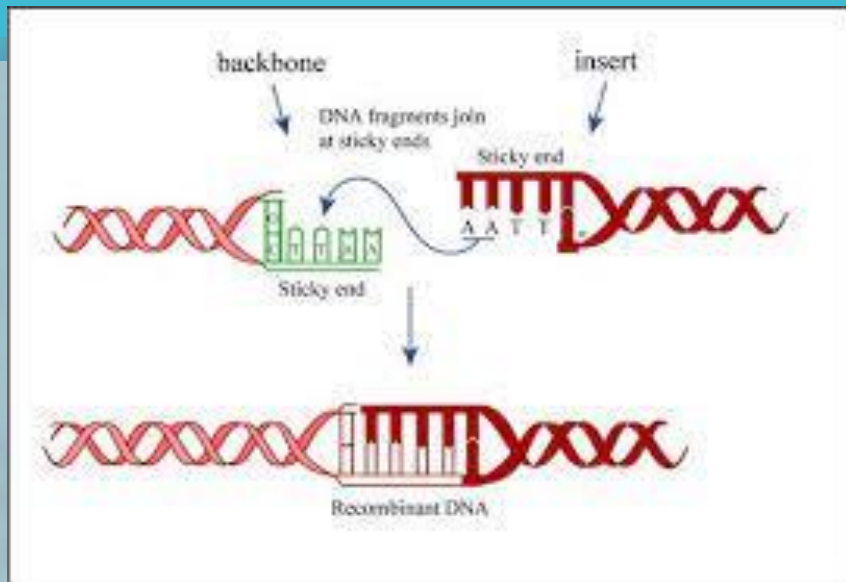
## Amplification and validation



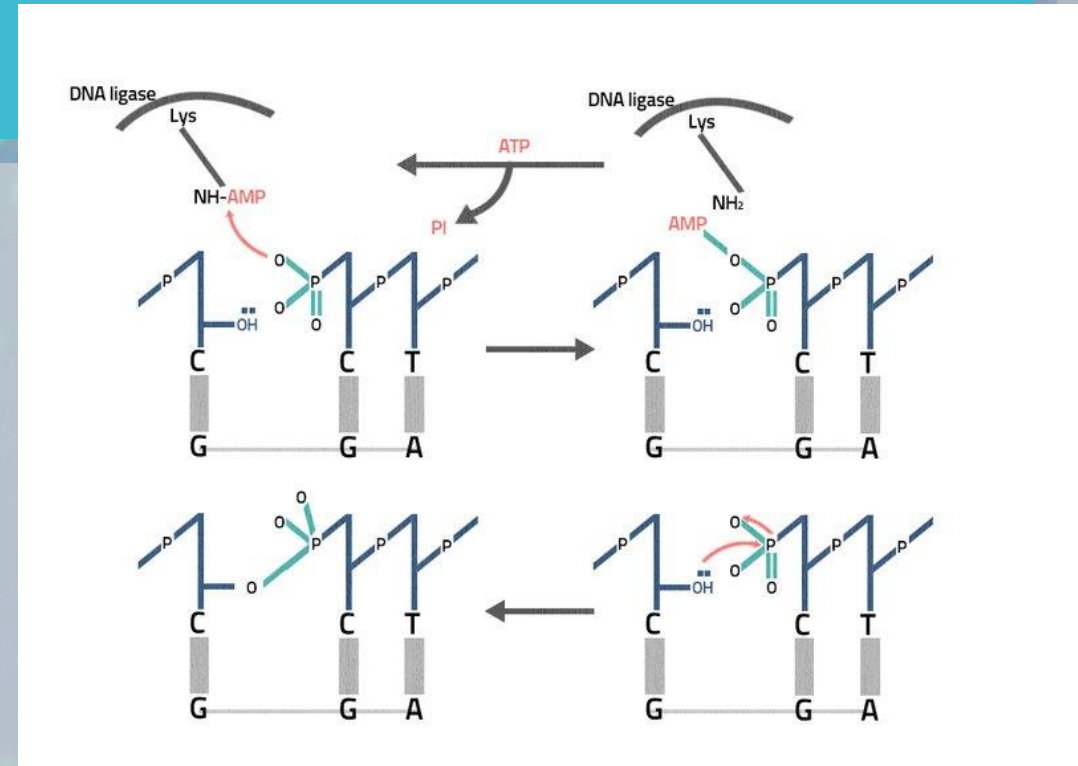


# Ligation

## Principle



- ! Molar ratio and not quantity ratio





# Ligation Reaction

	Tube Lig A (ratio 1:2)	Tube Lig B (ratio 1:3)	Tube Lig C (ratio idem A or B)	Tube Lig D
H <sub>2</sub> O	qsf 20 µl	qsf 20 µl	qsf 20 µl	qsf 20 µl
Ligase buffer (10 X)				
<b>! Make sure <u>you</u> vortex it before pipetting it</b>	1 X	1 X	1 X	1 X
Cloning vector (previously digested and purified)	50 ng	50 ng	50 ng	50 ng
Insert (previously digested and purified)	x ng <b>(YOU MUST CALCULATE*)</b>	x ng <b>(YOU MUST CALCULATE*)</b>	x ng <b>(YOU MUST CALCULATE*)</b>	-
Ligase	1 µl	1 µl	-	1 µl

You need to calculate the **required amount of insert**  
**! Molar ratio (and not mass ratio)**

→ Tool recommended for calculation: <http://nebiocalculator.neb.com>

Or simply apply the following calculation:

$$\frac{\text{ng of vector} \times \text{kb size of insert}}{\text{kb size of vector}} \times \text{insert:vector molar ratio} = \text{ng of insert}$$

# Ligation

## Ligation Calculator

This tool will calculate the mass of insert required at several molar insert:vector ratios in the range needed for typical ligation reactions.

Ligation

Tutorials

### Insert DNA length

kb



### Vector DNA length

kb



### Vector DNA mass

ng



### Required insert DNA mass

--- (1:1)

--- (2:1)

--- (3:1)

--- (5:1)

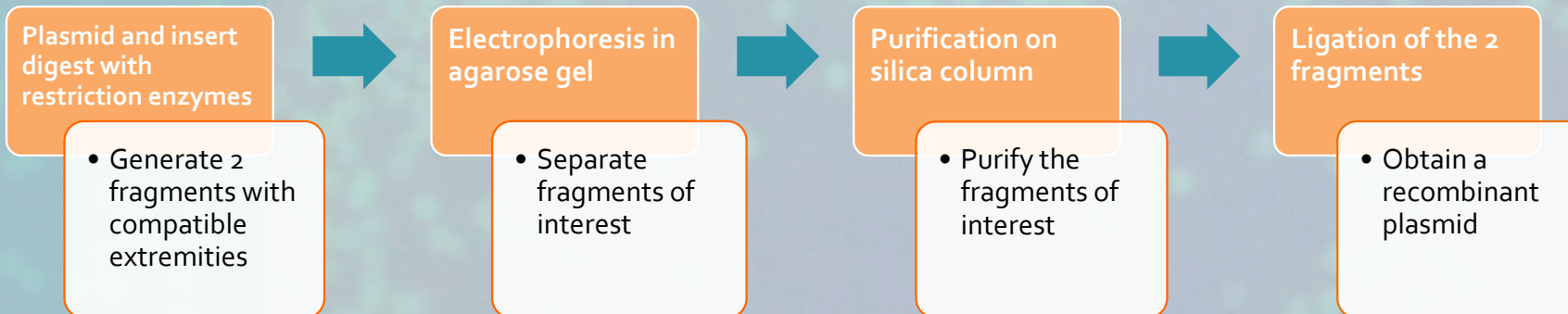
--- (7:1)

### Formula

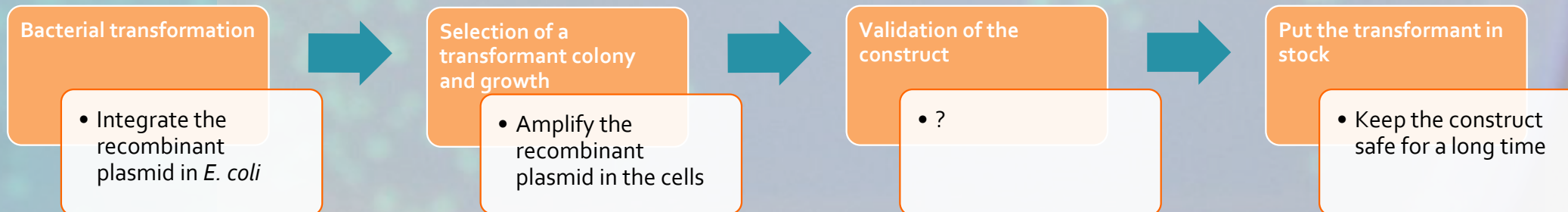
required mass insert (g) = desired insert/vector molar ratio x mass of vector (g) x ratio of insert to vector lengths

# Workflow

## Construction

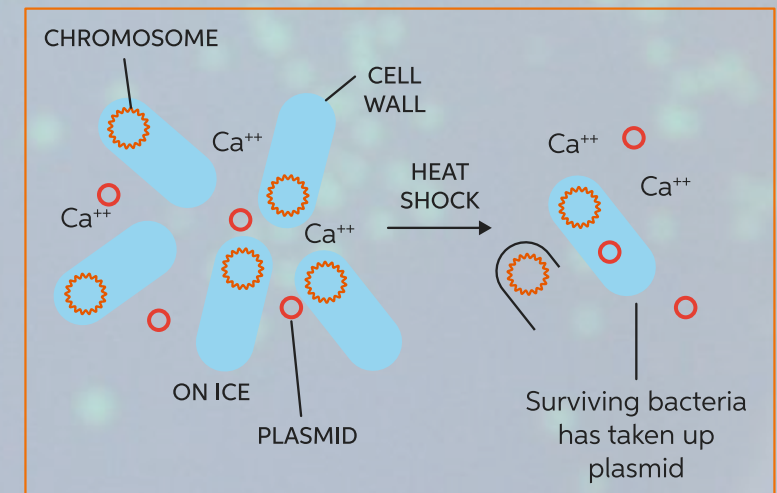
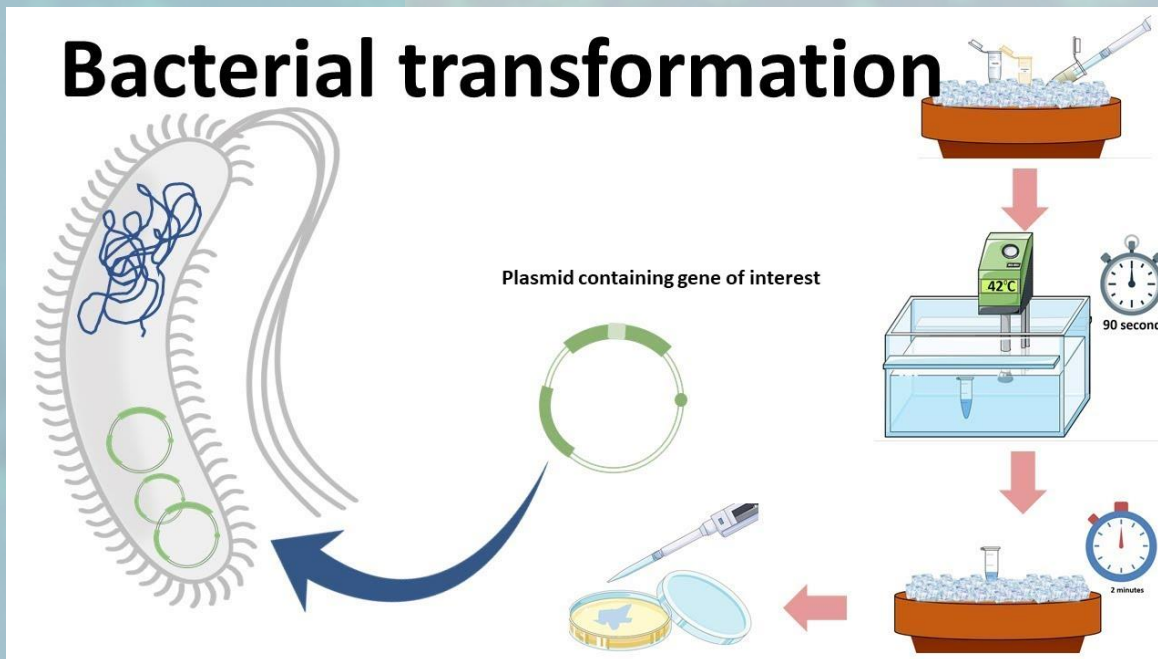


## Amplification and validation



# Bacterial transformation

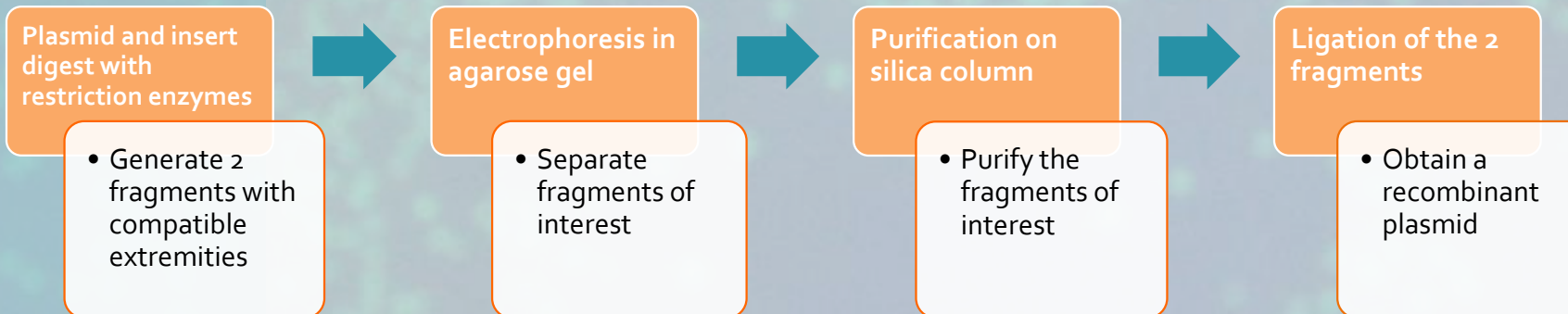
## Principle



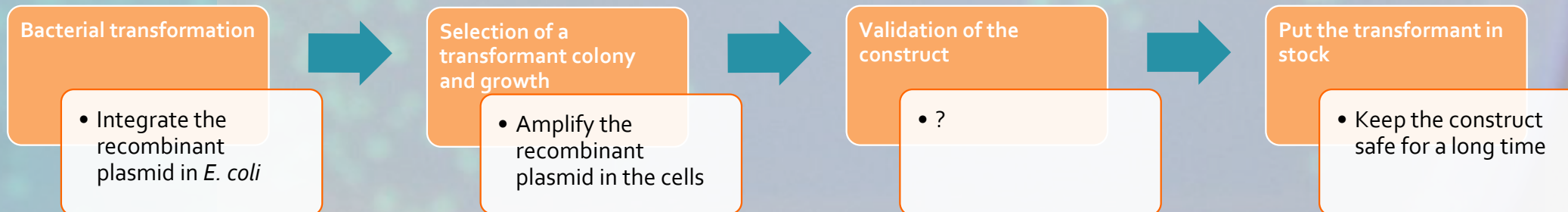


# Workflow

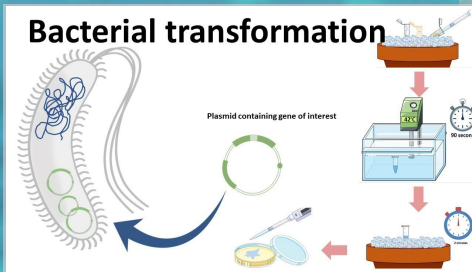
## Construction



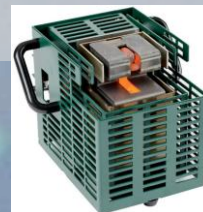
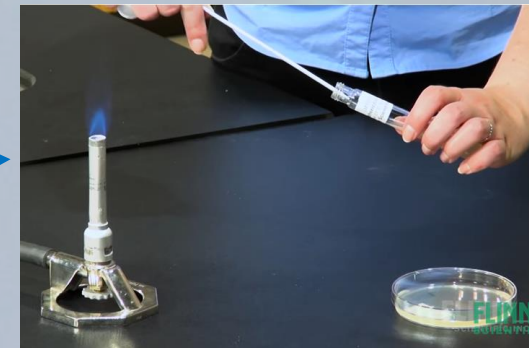
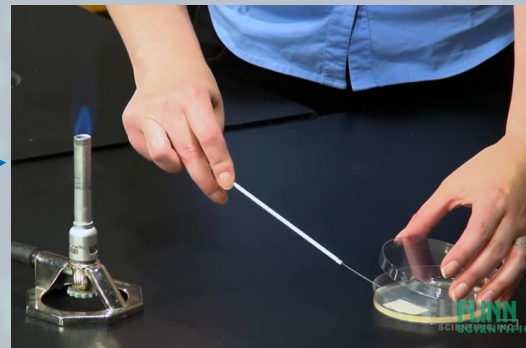
## Amplification and validation



# Selection of transformant colonies and growth in liquid nutritive medium

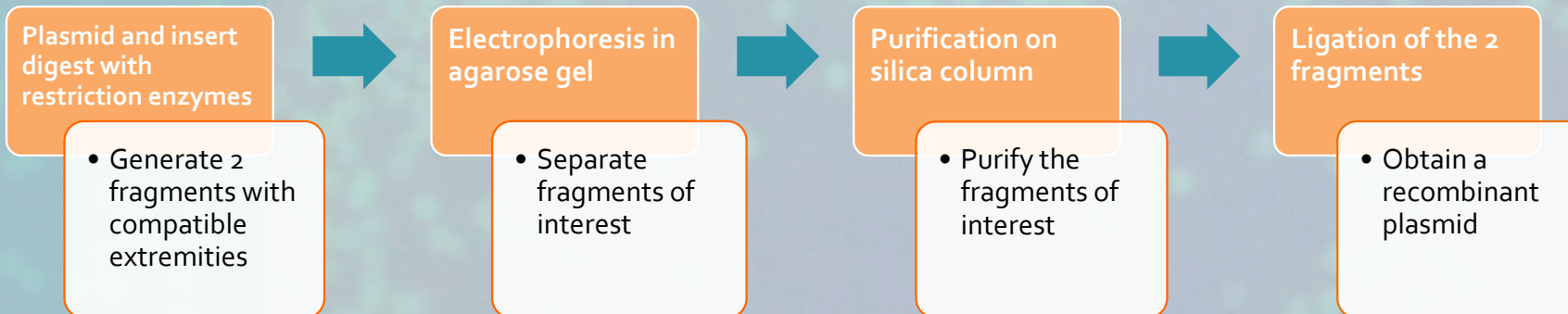


Medium = ?  
Controls = ?

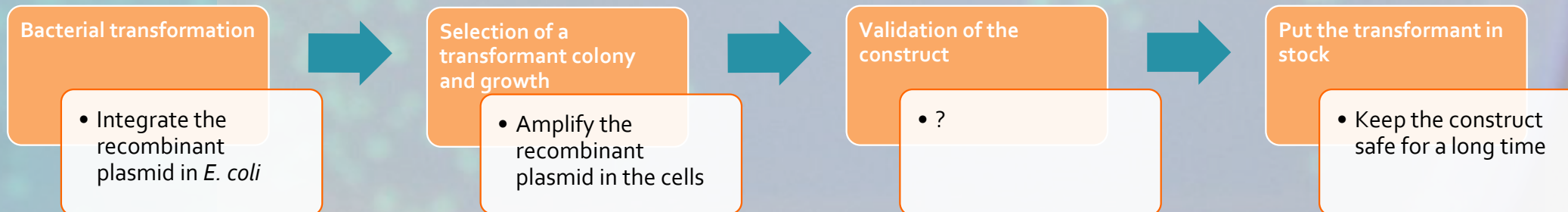


# Workflow

## Construction



## Amplification and validation





# Validation of the construct

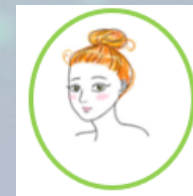
- Has the insert been integrated in the plasmid? In the correct orientation?
- → You must **plan the experimental steps you want to perform to prove it**
- → To do so, you must **make a scheme of all the possibilities of ligation reactions that can take place** (some of them are more likely to happen than others)

# Time schedule

- 4 lab sessions:
  - Day 1 = plasmid and insert digest, agarose gel preparation, electrophoresis and purification, ligation
  - Day 2 = bacterial transformation and discussion about validation protocol
  - Day 3 = analysis of transformation results and culture of transformants
  - Day 4 = cloning validation, finishing report writing-up

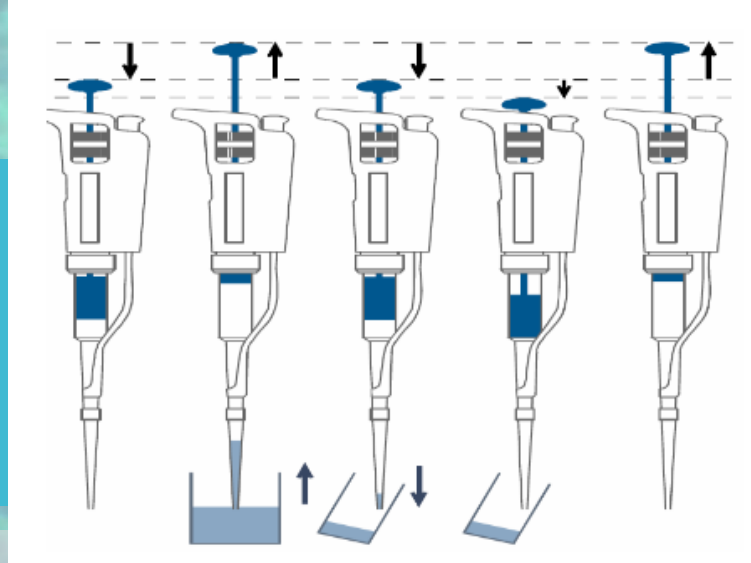
# Requirements for lab work

- Basic hygiene and security rules in the lab:
  - !!! Work with alkaline/acidic/CMR reactivities
  - Wear a lab coat in cotton
  - Be careful when and how to use lab gloves
  - No eating and drinking
  - Not use your mobile phone
  - Keep long hair attached





# Pipeting technique



- Pipeting small volumes of reactives → use of micropipettes = precision tools
  - Handle with care and precision
  - Effect of different parameters on pipeting: temperature of sample, viscosity...
  - Most simple technique: direct pipeting
  - Always use pipette tips
  - Keep the end of the pipette tip just below the level of the taken liquid
  - Always look at the volume you take up and transfer



Push button and volume setting

Tip ejector

Volume display

Piston assembly (inside)

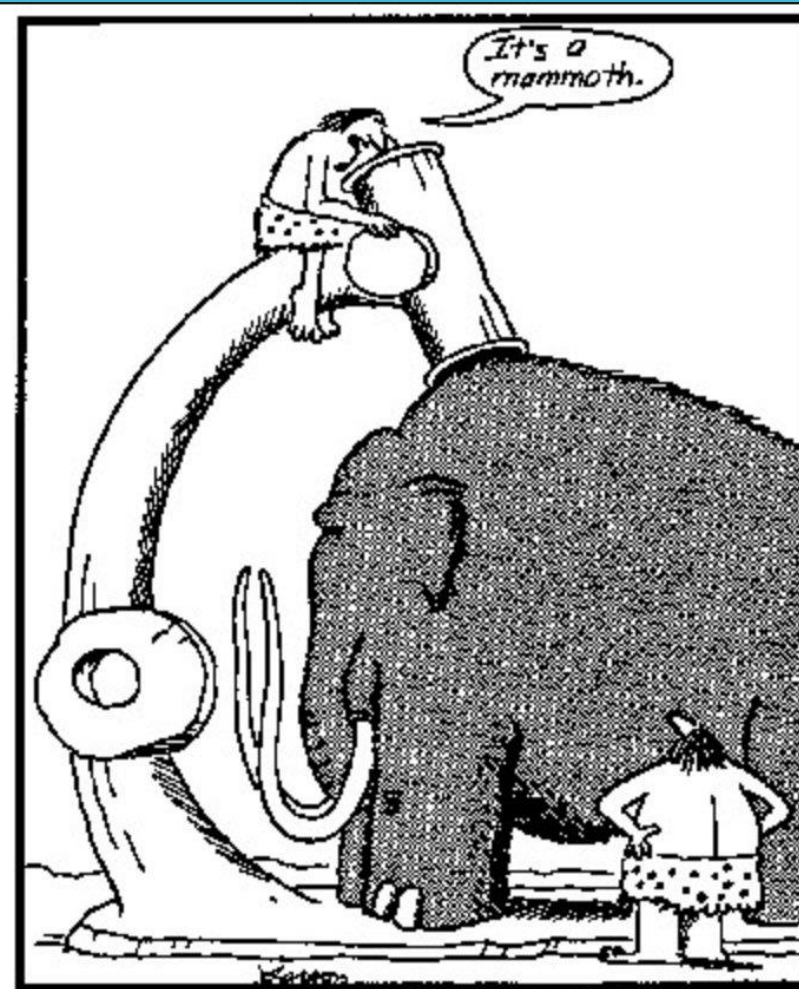
Ejector collar

Tip holder

Tip

Thank you for you attention!

Hope you enjoy the lab class



Early microscope

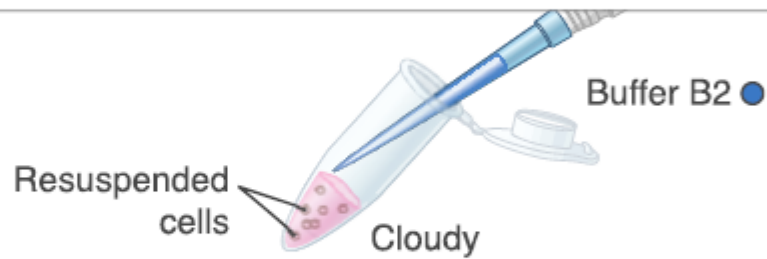
# Appendix - Purification of plasmid

## Procedure overview

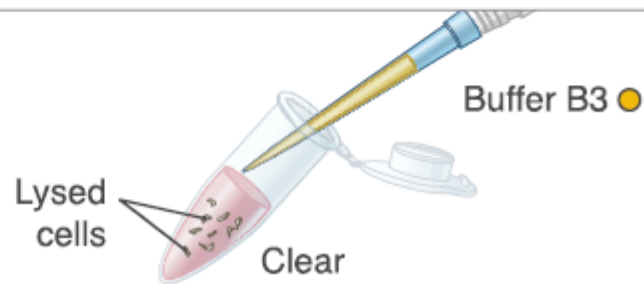
### 1. Resuspend Cells



### 2. Lyse Cells



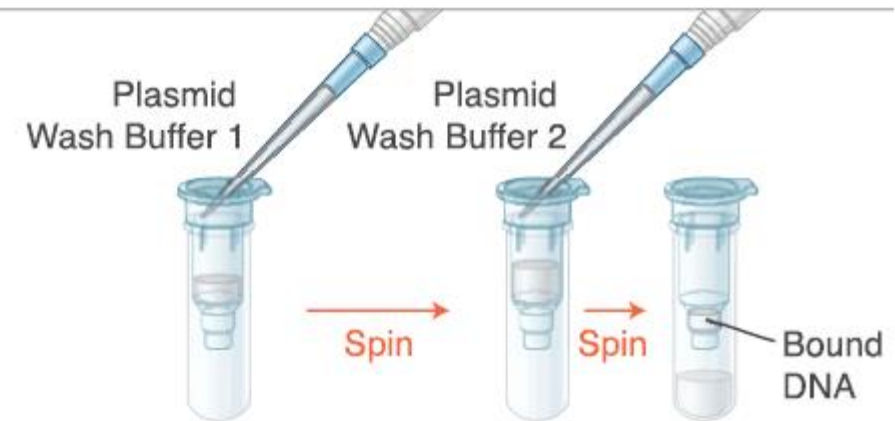
### 3. Neutralize Lysate



### 4. Bind DNA from Supernatant to Matrix



### 5. Wash Matrix



### 6. Elute DNA

