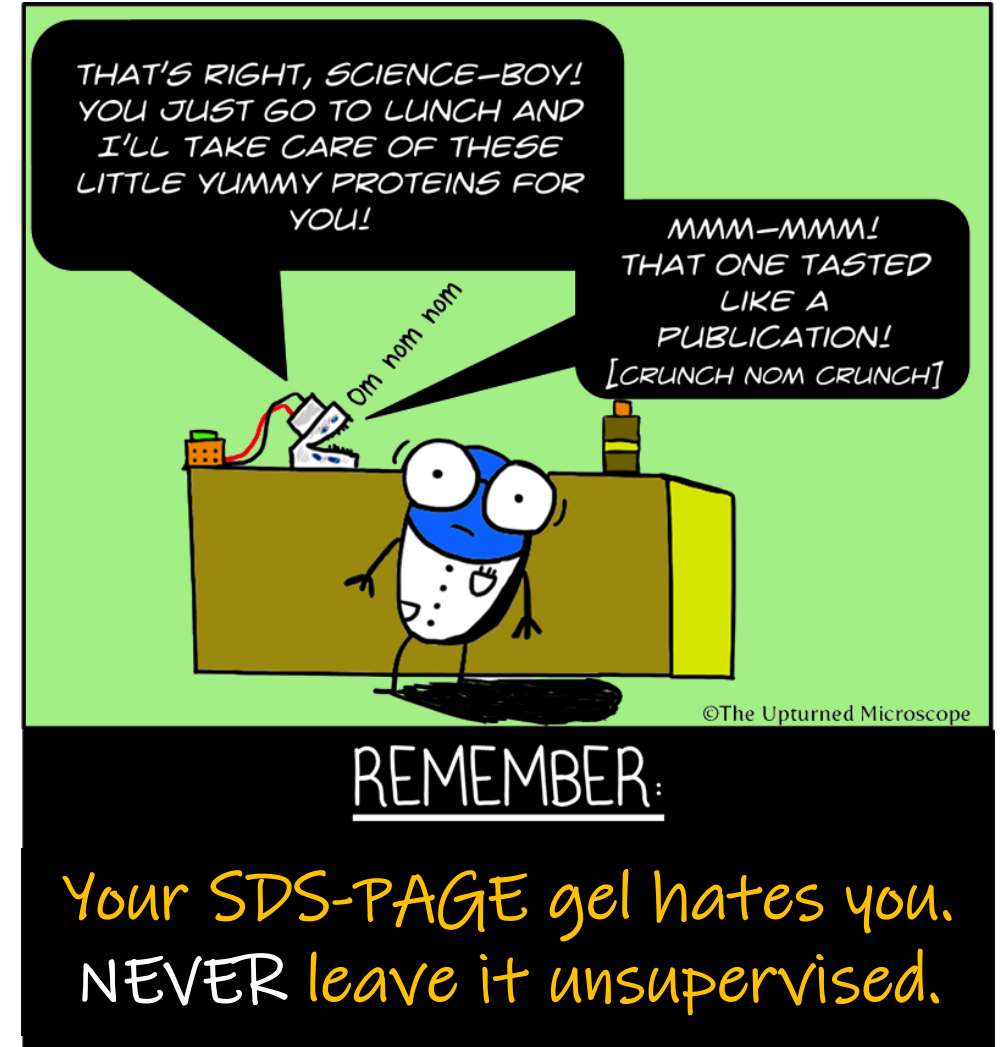


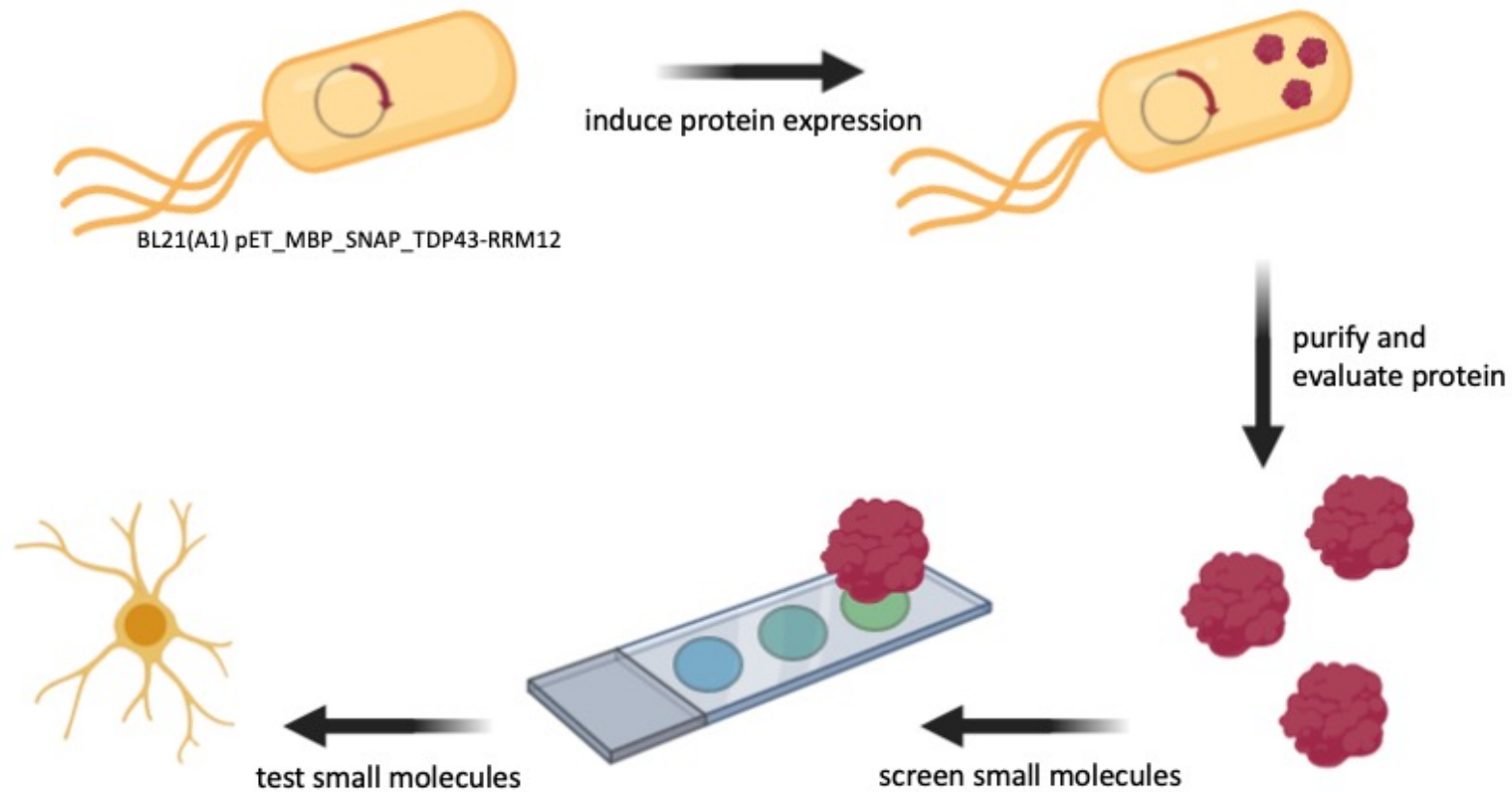
# M2D2: Assess purity and concentration of purified protein

1. Prelab discussion
2. Visualize protein purity using SDS-PAGE
3. Measure protein concentration using BCA assay



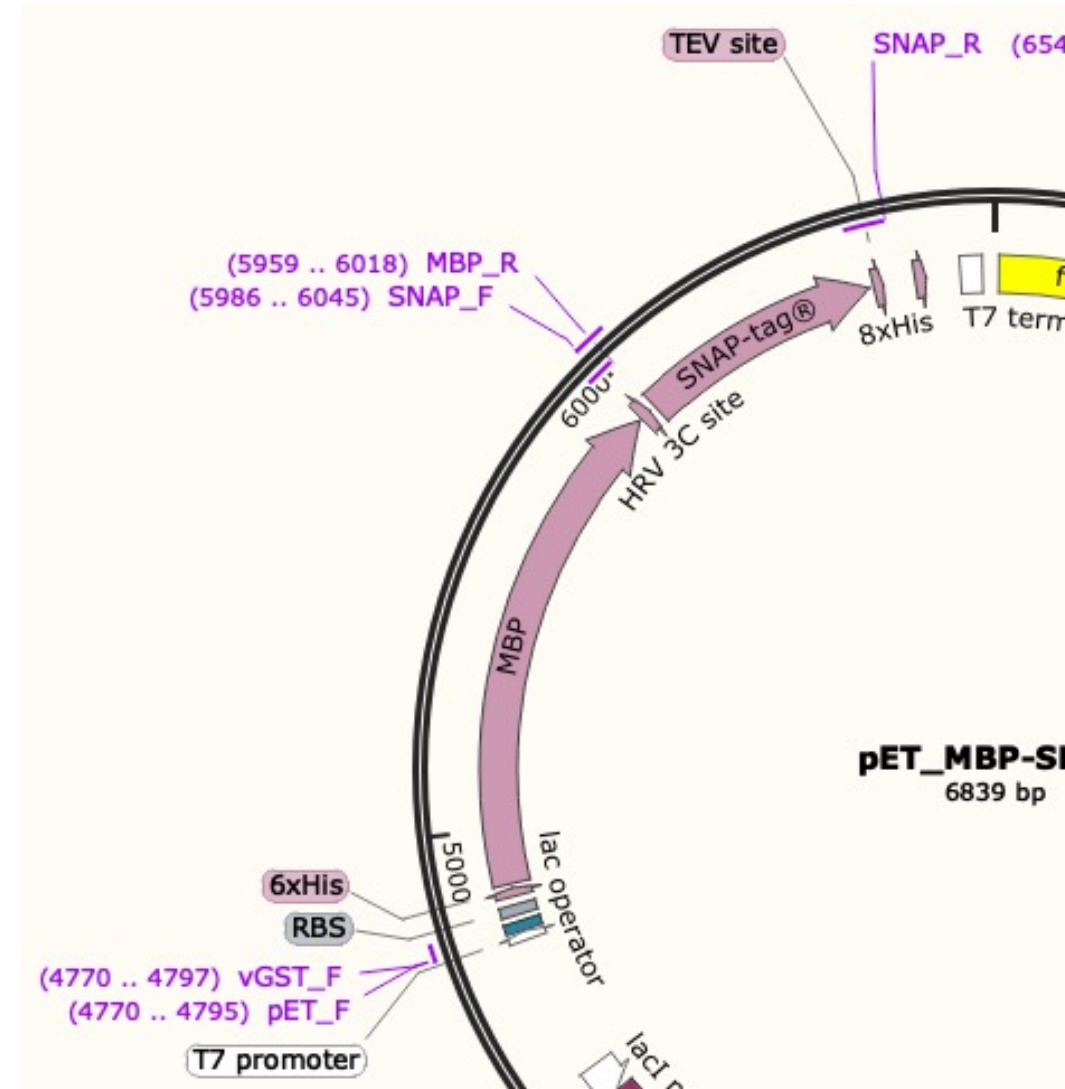
# Overview of M2

**Research goal:** Identify and characterize small molecule binders to a protein drug target.

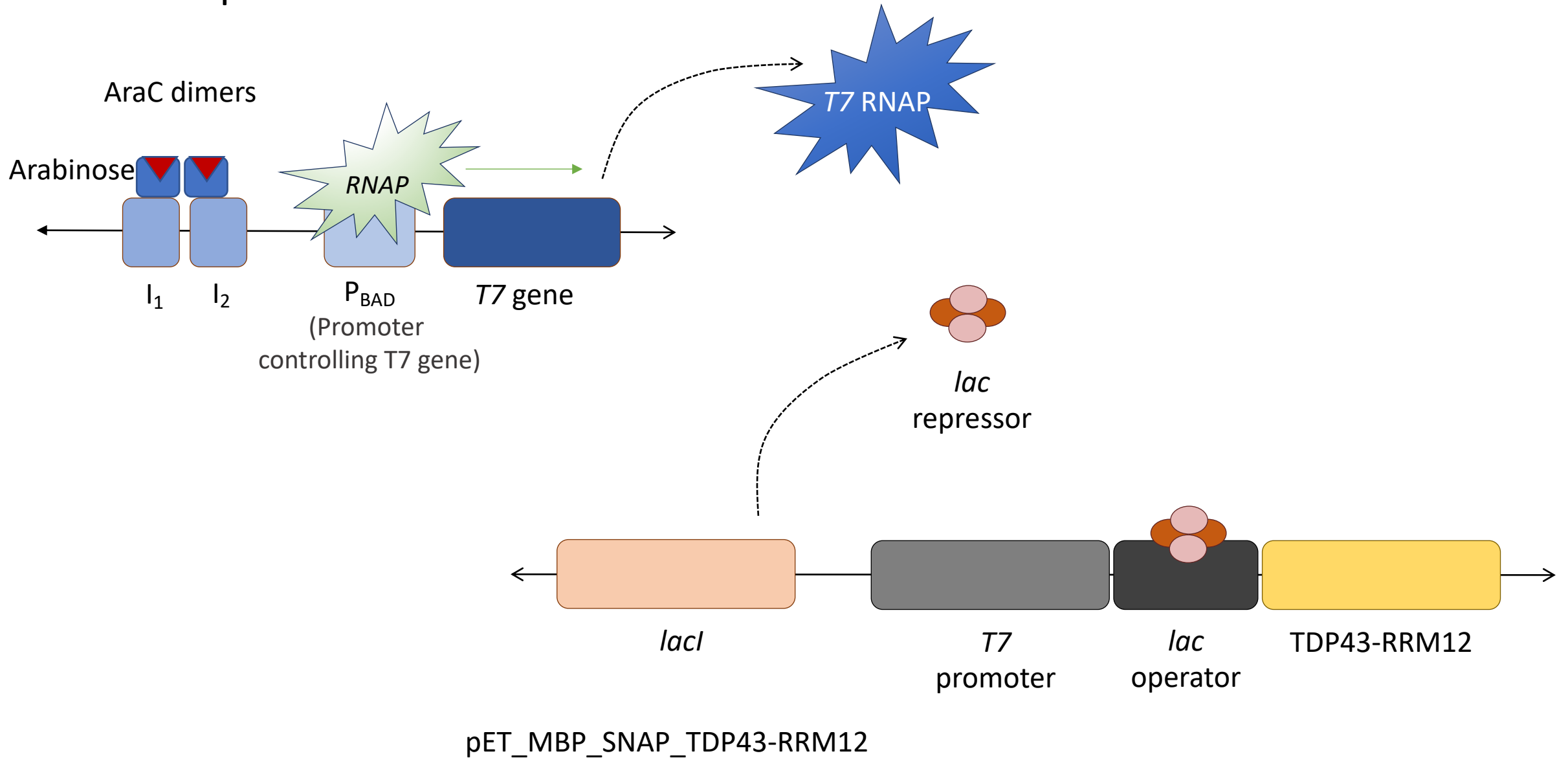


# Protein induction review

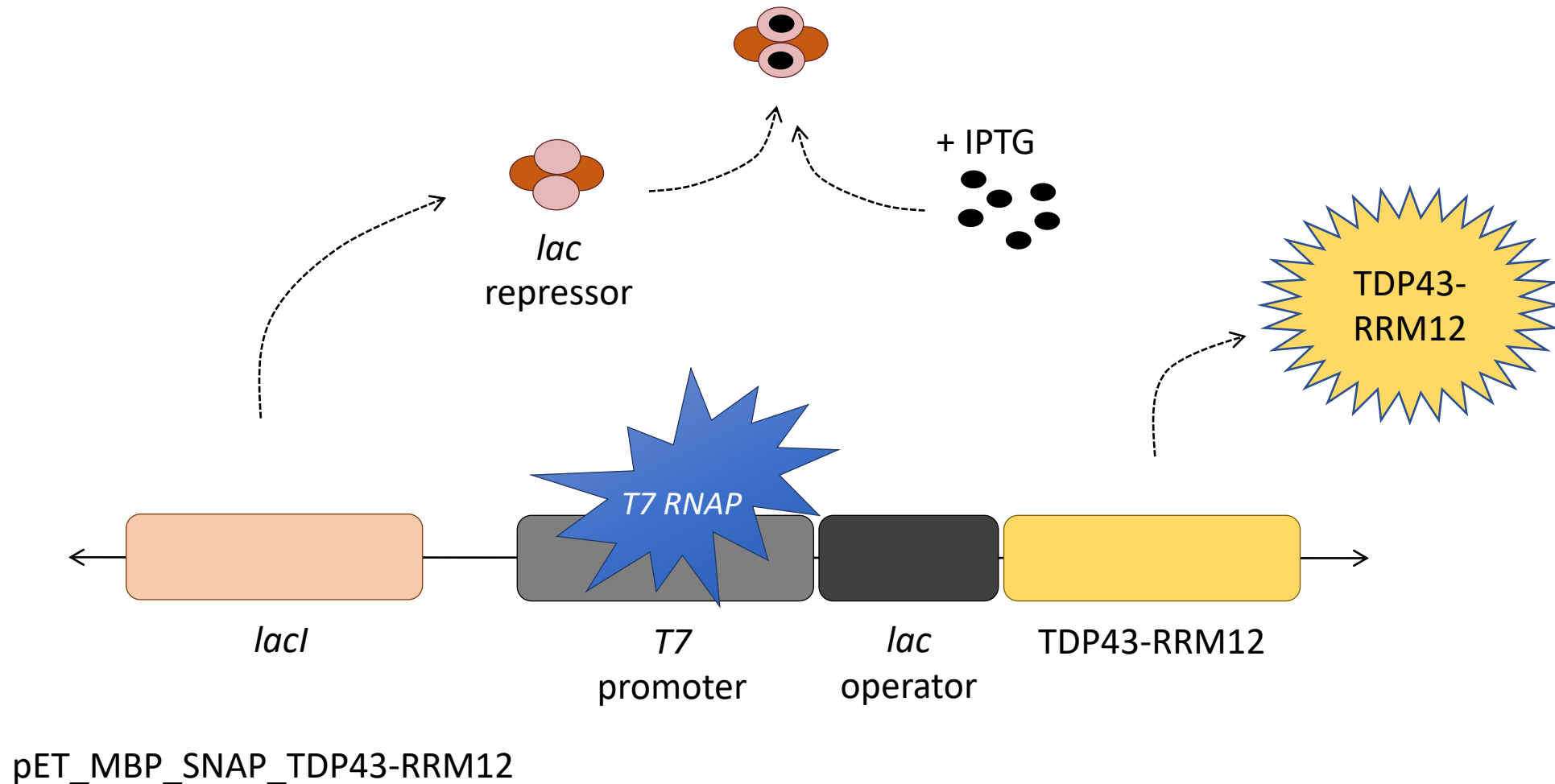
- What were the two chemicals used to induce TDP43\_RRM12 expression?
- What do they allow to be expressed/how?



# Arabinose controls T7 expression while LacI repressor blocks transcription

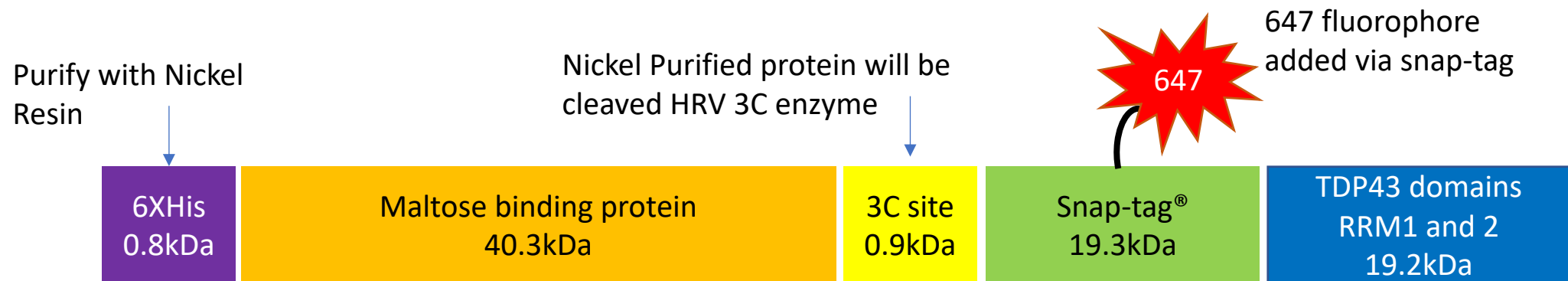


# IPTG 'induces' protein expression by preventing LacI repression



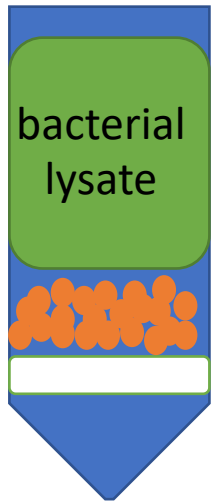
# What is protein expressed in our system

## Our protein for this module:



# Protein purification review

- Why this step?



## Flowthrough

On beads:



## Wash

On beads:



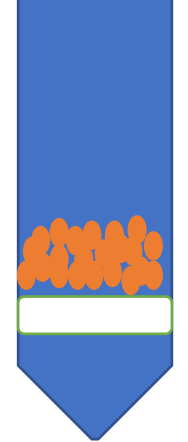
## Elution

On beads:



## Slurry

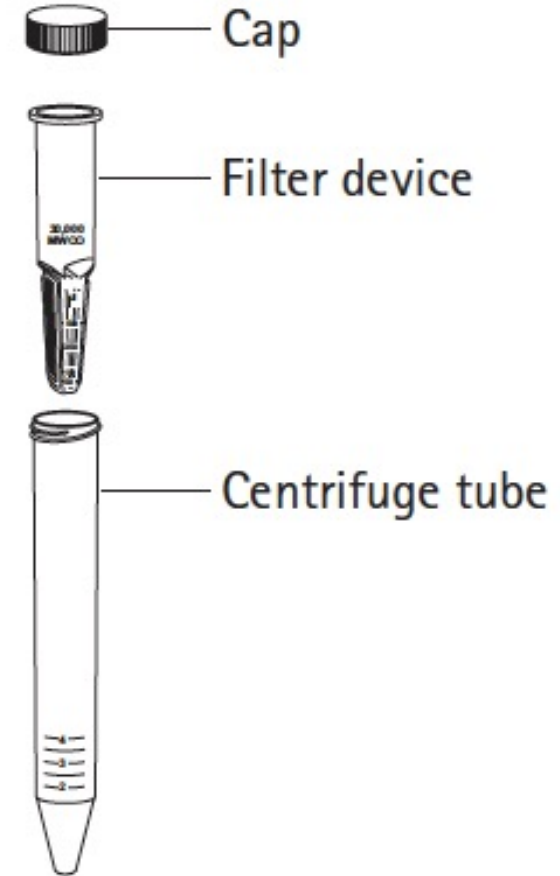
On beads:



- What's on the  $\text{Ni}^{2+}$  beads?
- What's in the expelled liquid?

# Protein is concentrated after purification

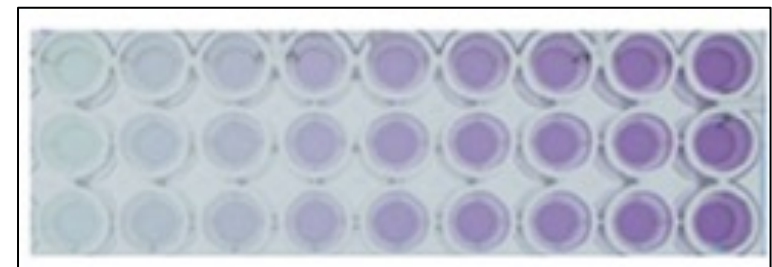
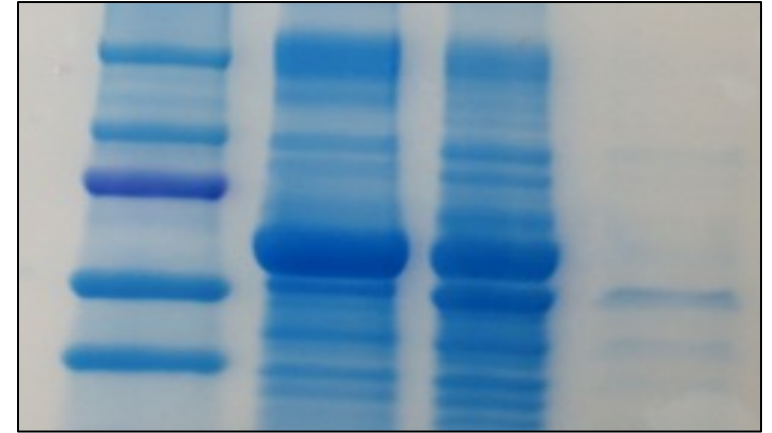
- Filter device sits within centrifuge tube
  - Protein added to filter device before centrifugation
- Filter has MW cutoff of 3 kDa
  - protein retained in the filter device during centrifugation
  - TDP43-RRM12 + Snap-tag = ?
  - 6x His tag = 2.5 kDa
- How does this concentrate the protein?



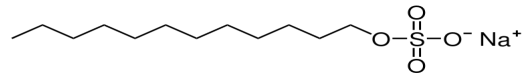


# How will you assess purity and concentration?

- Check purity using SDS-PAGE
  - Identifies presence of protein during purification procedure
  - Visual detection of other proteins in sample
- Measure concentration using BCA assay
  - Colorimetric assay
  - Calculate concentration from standard curve

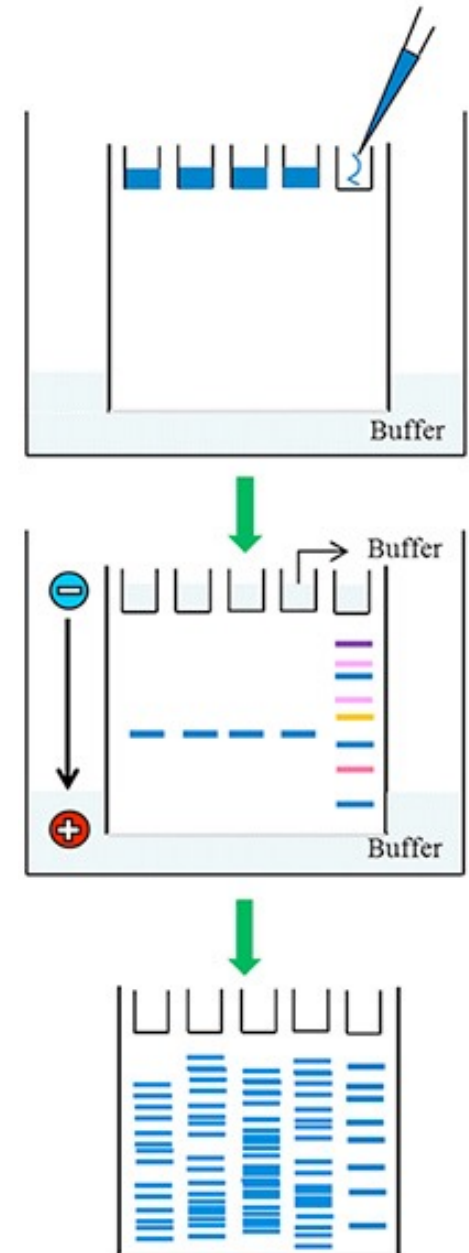


Purity: Sodium dodecyl sulfate polyacrylamide gel electrophoresis (SDS-PAGE)

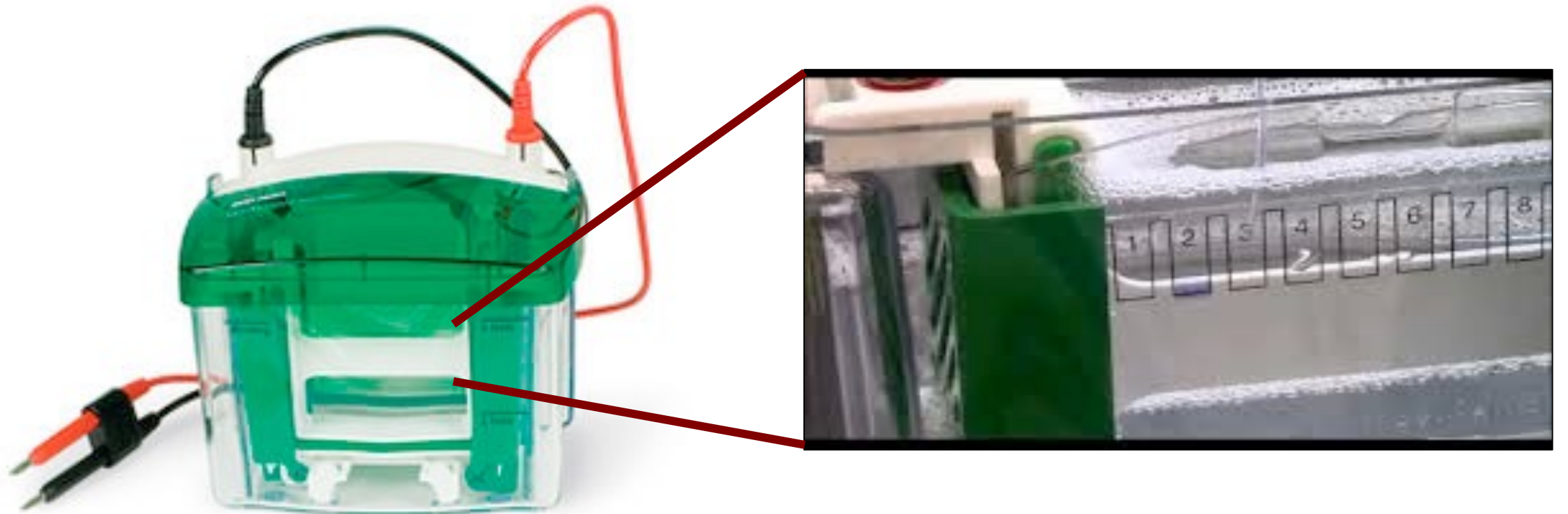


# How are proteins separated?

- Laemmli buffer and boiling results in denatured and \_\_\_\_\_ charged proteins
- SDS-PAGE separates proteins by \_\_\_\_\_
- Electrophoresis completed in TGS buffer
  - Tris-HCl
  - SDS
  - Glycine



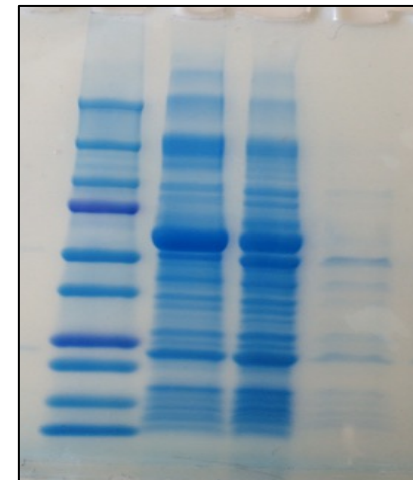
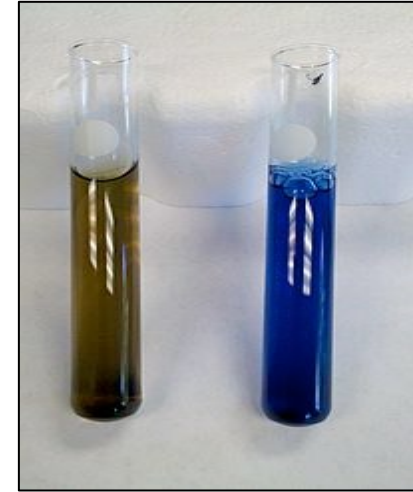
# Demonstration of SDS-PAGE



# How are proteins visualized?

Coomassie brilliant blue G-250 dye used to stain gel after electrophoresis

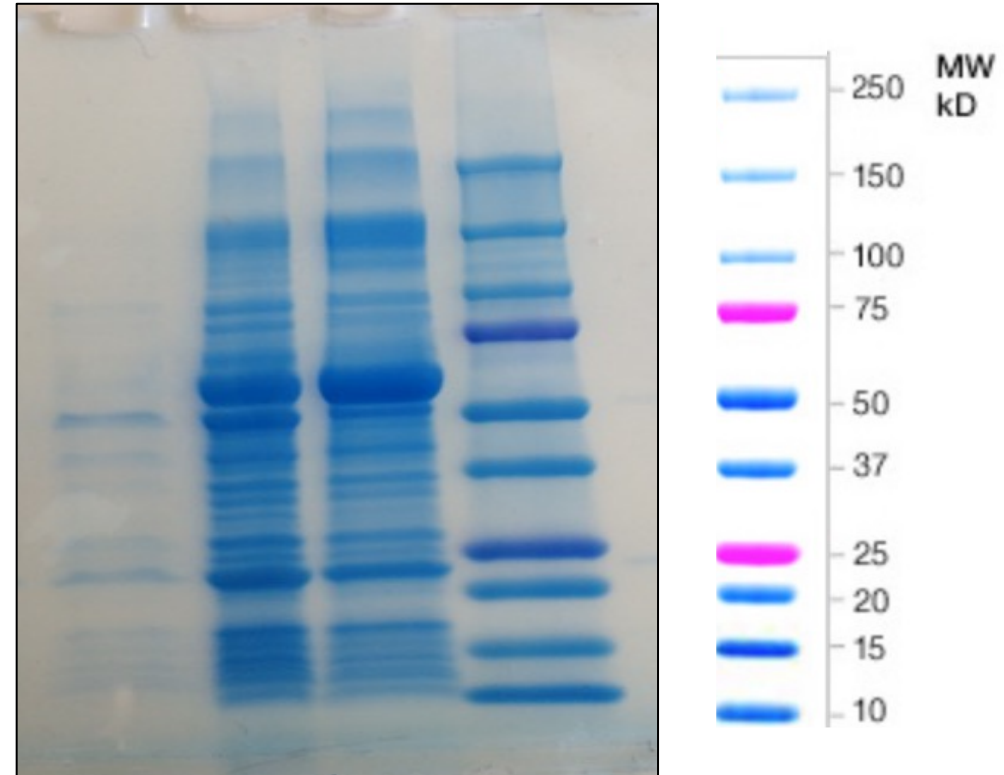
- Red if unbound (cationic form)
- Blue if bound to protein (anionic form)
- Hydrophobic and electrostatic interactions with basic residues
- Arg (also His, Lys, Phe, Trp)



# What are the expected results of SDS-PAGE?

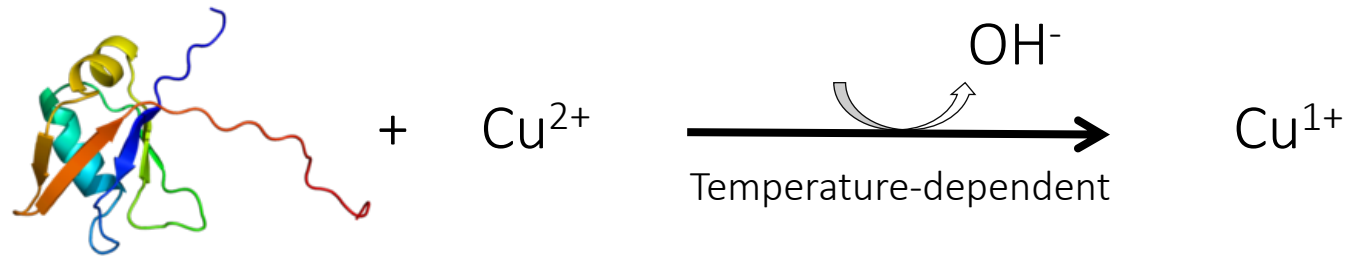
Each lane of the gel should be explained in the results

- What bands are expected? Do you see the bands you expected?
- Do you see any unexpected bands?
- What do the bands tell you about the purity of your protein?
- What does might this tell you about the protein concentration calculated in the next step?

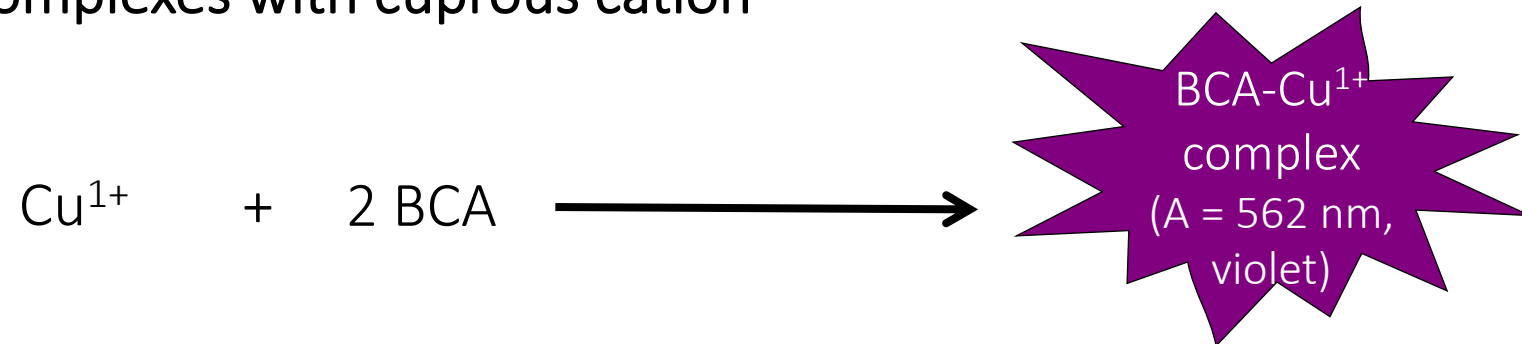


# Concentration: Bicinchoninic acid (BCA) protein assay

Step 1: Biuret reaction; chelation of copper with protein, reduction of copper



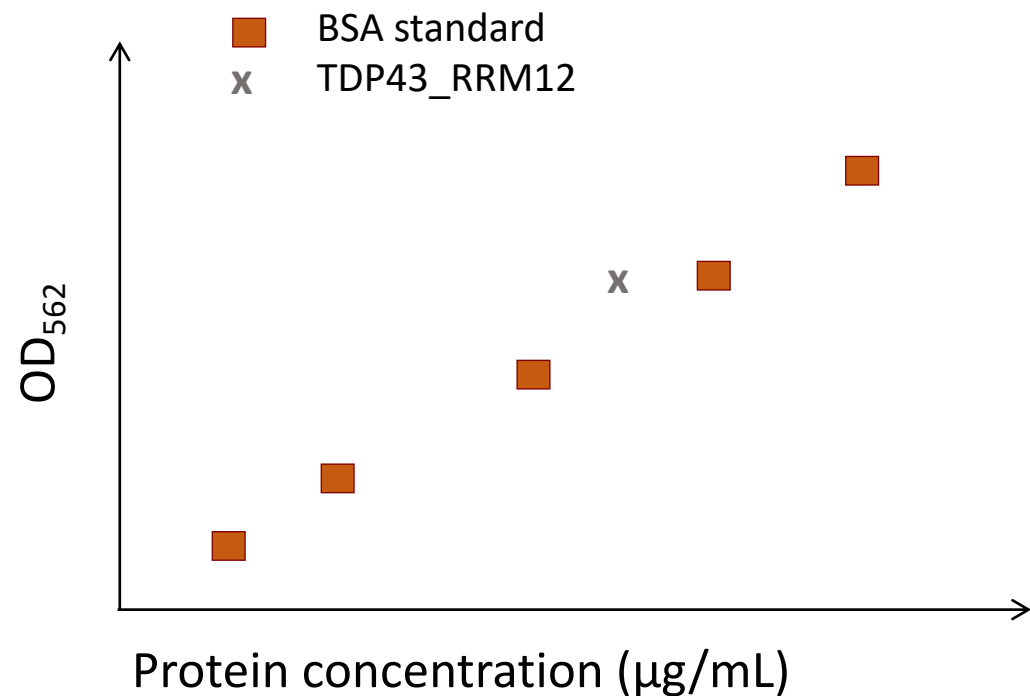
Step 2: BCA complexes with cuprous cation



# BCA/Cu<sup>1+</sup> absorbance proportional to protein concentration

Standard curve generated using serial dilutions of bovine serum albumin (BSA)

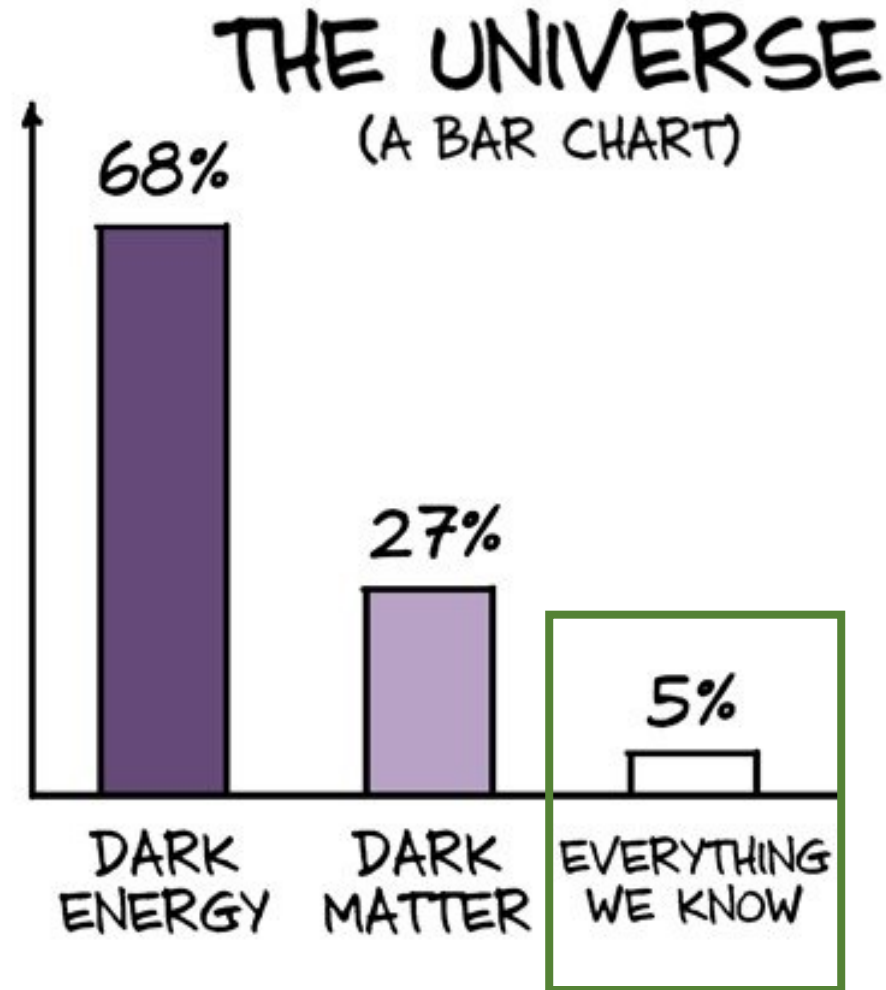
- Equation of the line used to calculate protein concentration
- What does the R<sup>2</sup> value tell you about the standard curve?  
What does this tell you about the calculated concentration?





We know a small fraction of what the universe has to offer (a take-home message)

- Minimal text included to understand the figure
- Oooh, 5%. That's better than expected.
- But, what does that even mean?



## For today...

- Work through M2D2 laboratory exercises with partner
- Work on Mini-presentation!

## For M2D3...

- Outline the Introduction section for Research article
- Review paper for in-class discussion with partner,
  - Draft slide, script for presenting Figure 1 from that paper