

# Notes for Data Summary:

- If you would like to do revisions on your Data Summary
  - These are not required
- Revisions have the potential to bring a grade up 1 & 1/3 letter grade (i.e. B- to A)
- We run revisions in the same way as scientific journals
  - Write a cover letter!
  - We ask that you go one step further and show your changes with different colored font

**Data summary revision is due by Saturday, October 23rd at 10 pm** [\[edit\]](#)

**For your final report, all changes need to be in a different colored font so the improvements you made are clear.** You should also include a cover letter with your final draft that explains how you addressed the concerns raised (e.g. "paragraph x was completely rewritten to better explain...." or "Results for the agarose gel analysis were clarified by ....").

# Notes for Journal Club:

## Journal club meets in 56-614 on T/R

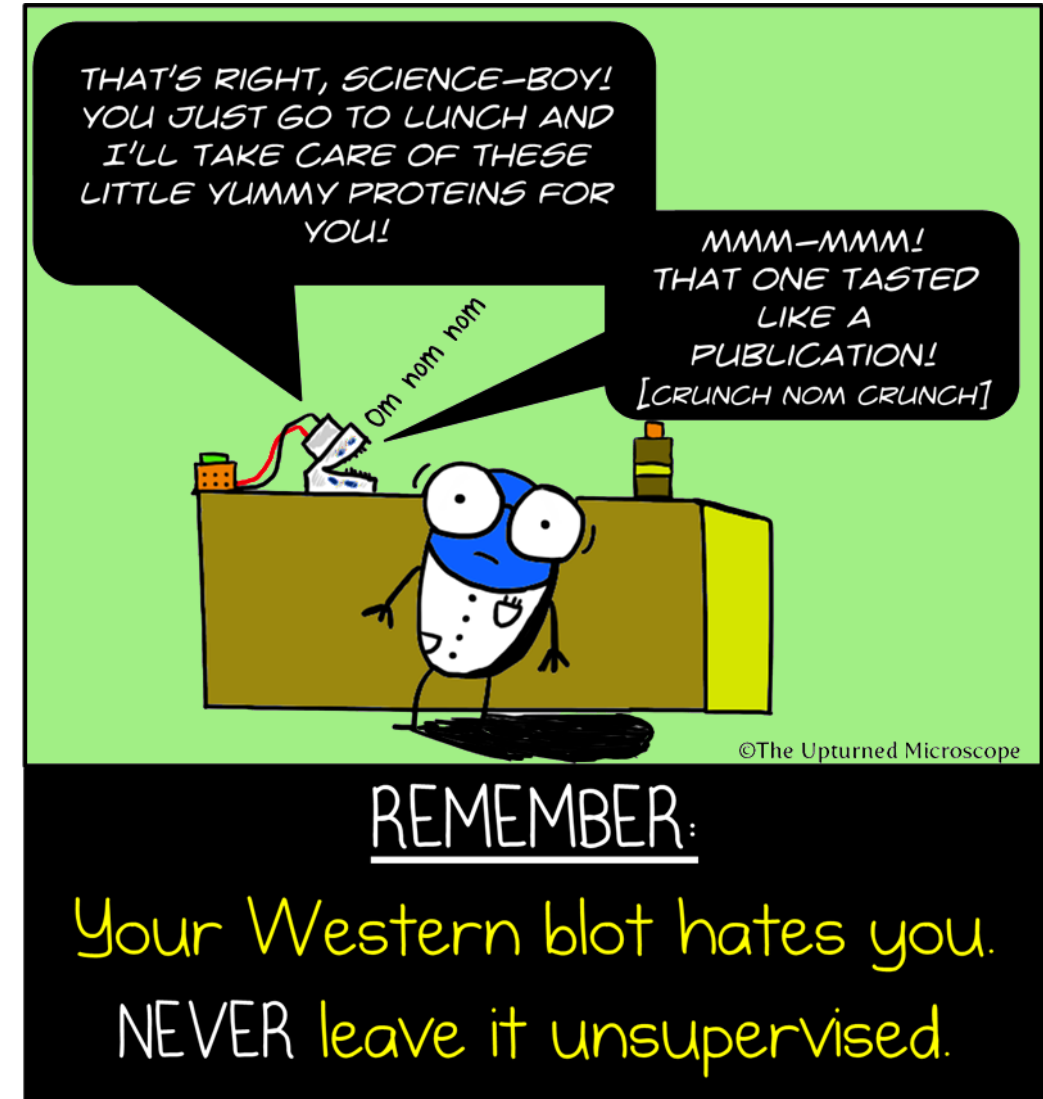
- If you want to discuss your article, please email in advance (to make sure I've read it)
- Your slides must be uploaded to Stellar 1hr prior to the class day in which you are presenting
  - The earliest slide upload gets first choice of a presentation slot on that day, and so on...
- We provide a Trader Joes smorgasbord to fortify through this experience
  - Let me know about food requests and any dietary restrictions
- No lecture on Tuesday or Thursday of next week

### Method of submission [\[edit\]](#)

Please submit your completed Journal club slides **1 hr prior to your scheduled laboratory session time** to Stellar [🔗](#), with filename **Name\_LabSection\_JC.pptx** (for example, ImaStudent\_TR\_JC.pptx).

# M2D5: Assess purity and concentration of purified protein

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



# Module 2 Roadmap

Determine putative PF3D7\_20109-F21 binders via high throughput screening (SMM)



Create plasmid of PF3D7\_20109-F21 to use in validation assays



Express PF3D7\_20109-F21 (from plasmid) in bacteria and purify protein

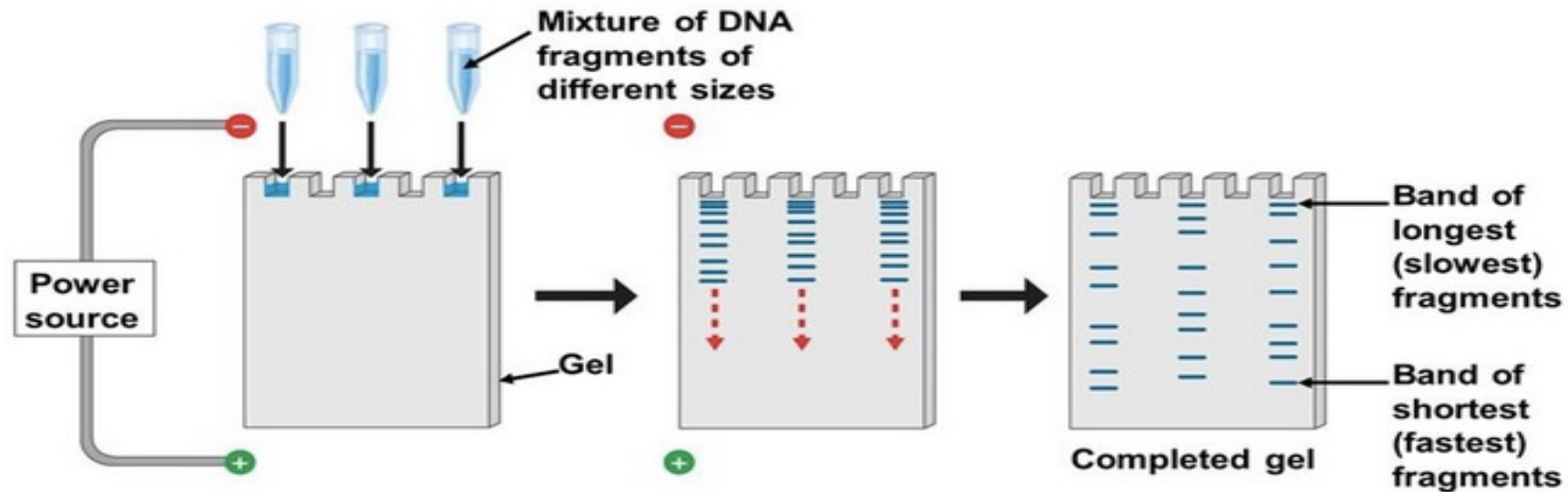


Assess purity and concentration of purified protein



Use purified protein to validate binding of small molecules identified in SMM

# DNA electrophoresis review



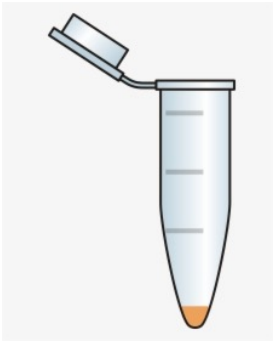
How do you visualize the migration through the gel?

How do you visualize DNA bands in the gel?

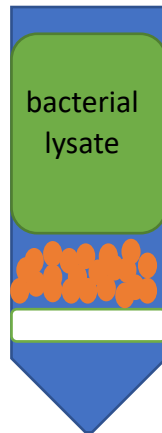
# Protein purification review

- Why this step?

## Pellet



## Lysate



## Flowthrough



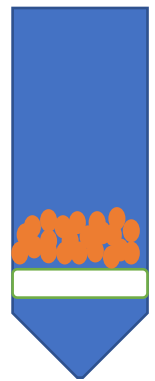
## Wash



## Elution



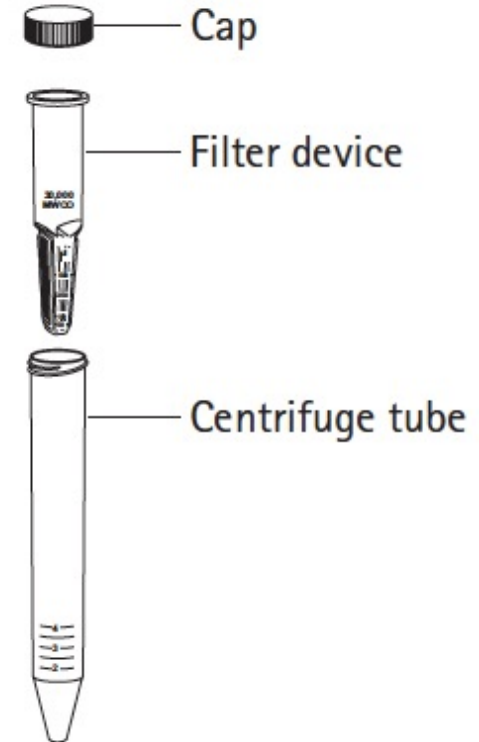
## Slurry



- What's on the resin?
- What's in the expelled liquid?

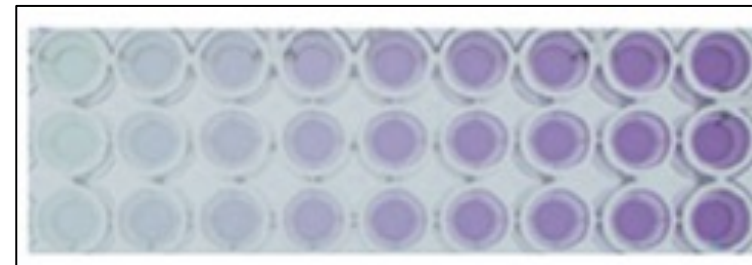
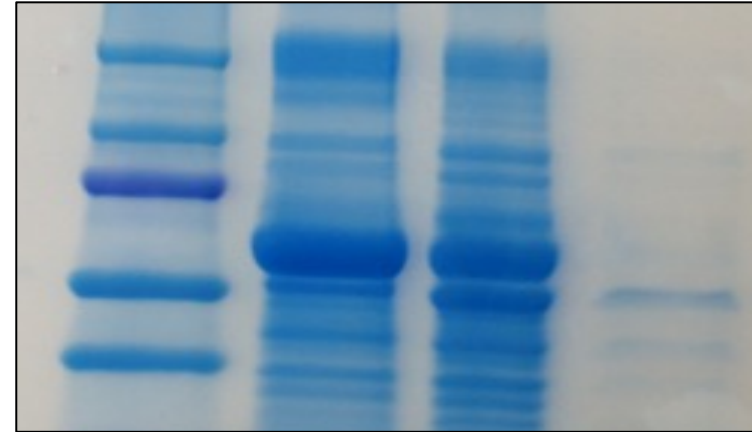
# Concentrate protein before testing

- Filter device sits within centrifuge tube...**add protein to filter device** for centrifugation
- Filter device has MW cutoff of 3 kDa ...**protein is retained in the filter device** during centrifugation
  - **PF3D7\_1351100 = 23kDa**
  - **Strep-tag = 3kDa**
- How does this concentrate the protein?
- How does this remove excess biotin?



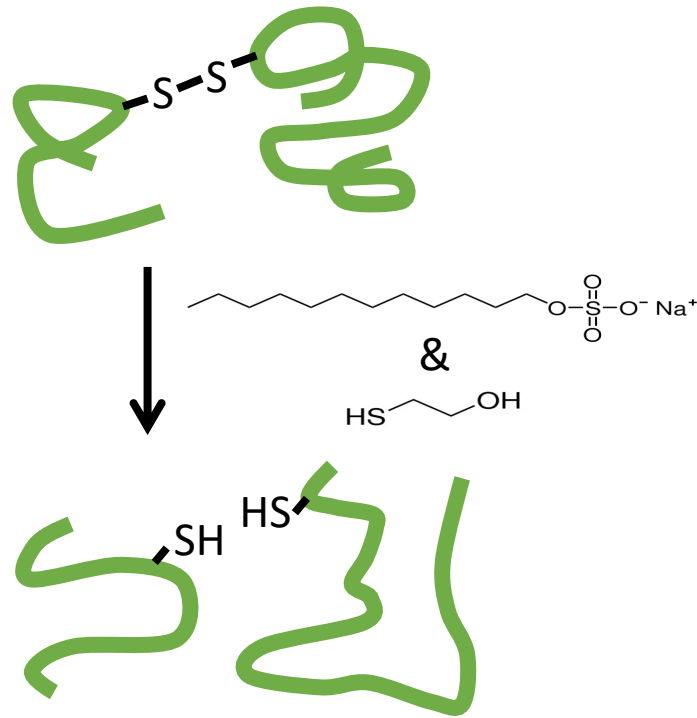
# How will you assess purity and concentration?

- Check **purity** using **SDS-PAGE**
  - Visual detection of other proteins in sample
  - Identifies purity of sample at multiple stages of purification
- Measure **concentration** using **BCA assay**
  - Colorimetric assay
  - Calculate concentration from standard curve





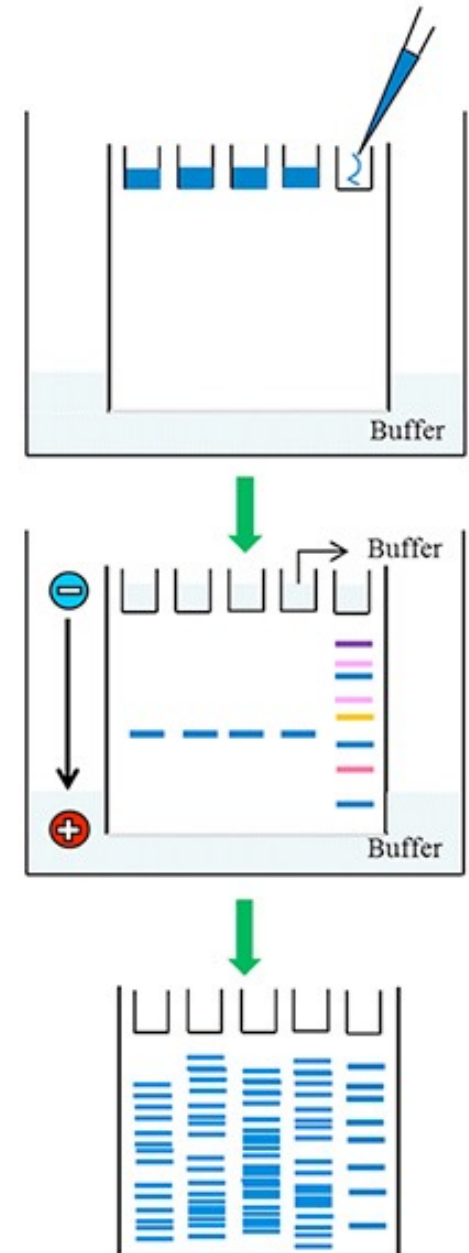
# Purity: Sodium dodecyl sulfate polyacrylamide gel electrophoresis (SDS-PAGE)



- Laemmli sample buffer / loading dye:
  - SDS
  - $\beta$ -mercaptoethanol (BME)
  - bromophenol blue
  - glycerol
- Boiling:

# How are proteins separated?

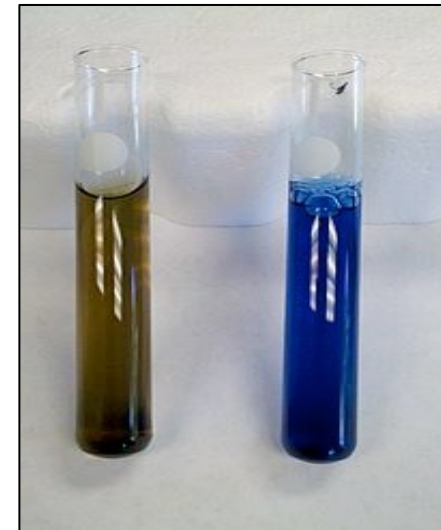
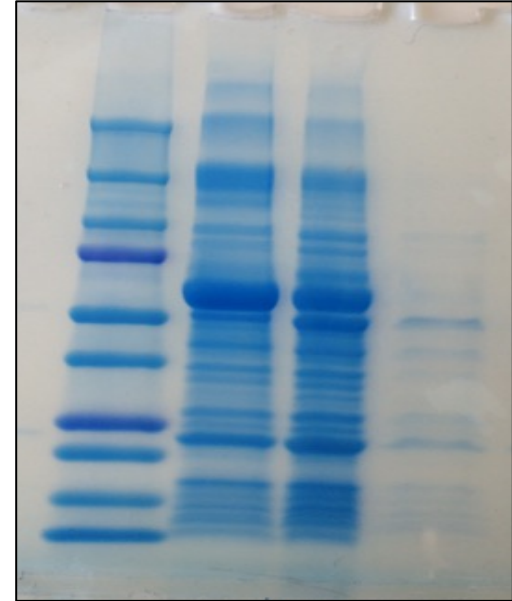
- Laemmli buffer and boiling results in \_\_\_\_\_ and \_\_\_\_\_ charged proteins
- SDS-PAGE separates proteins by \_\_\_\_\_
- Electrophoresis completed in TGS buffer
  - Tris-HCl
  - SDS
  - Glycine



# 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)



# Be mindful when assessing SDS-PAGE protein samples

Consider the order of your samples:

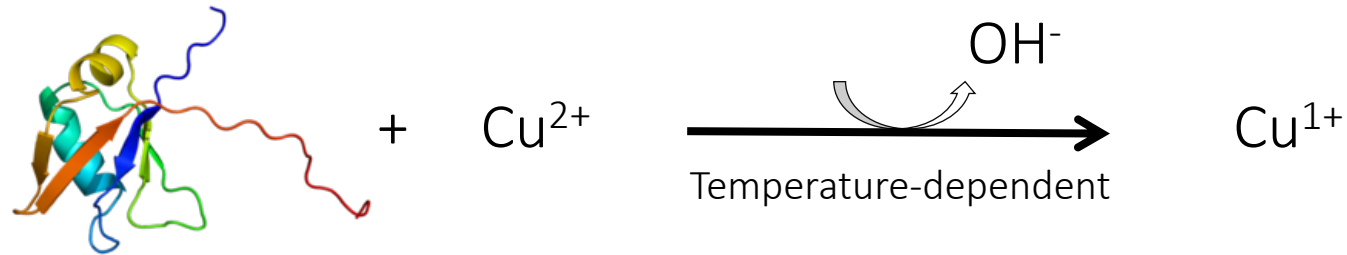
1. molecular weight ladder
2. pellet
3. lysate
4. flow-through
5. wash
6. elution
7. resin
8. concentrated protein.



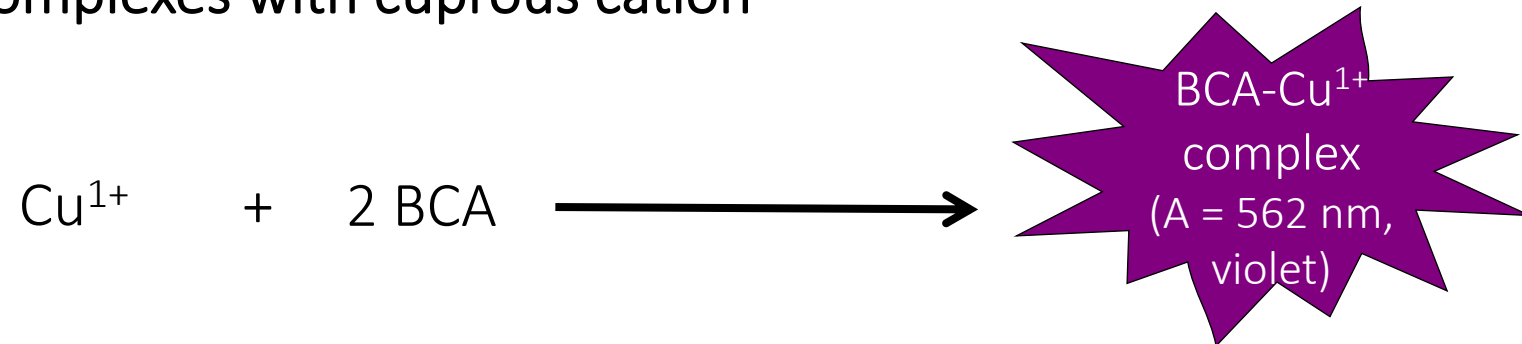
- Figure will be included in your Research Article!

# Concentration: Bicinchoninic acid (BCA) protein assay

Step 1: Chelation of copper with protein, reduction of copper sulfate to copper ion



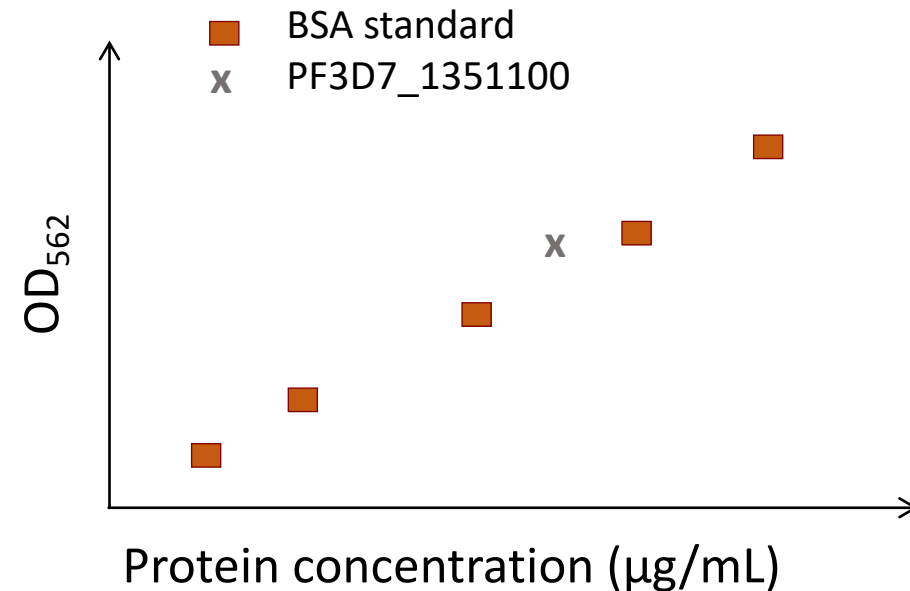
Step 2: BCA complexes with cuprous cation



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

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

- Use fresh tips between tubes
- Mix well between dilutions
- Be mindful of volumes



# For today...

- Complete the purity and concentration assessments
  - It's good to divide the work load here!
  - Start immediately by putting your Elution into the concentration column to spin!

# For M2D5...

- Draft methods for M2D2-M2D4
  - For your research article
  - Individual assignment

# Methods gift

Confirmation digest of pET-28b(+)\_PF3D7\_20109-F21

Topic sentence! (Include mention that construct was a gift from Dr. Khan Osman)

To confirm that PF3D7\_20109-F21 was cloned into pET-28b(+) expression vector, a digest was completed. Restriction enzymes Abcl and DefIII were used to digest X ng of pET-28b(+)\_PF3D7\_20109-F21 in single digests (only one enzyme added) and in a double digest (both enzymes added) using Y U / uL of each enzyme and CutSmart buffer (NEB). Digests were incubated at 37C for Z hrs. ...