

# Protein Purification and Gel Filtration



by Peter Van Haarhoven Nov 9, 2007



#### What is Protein Purification?



• Protein purification is a multi-step process for separating our target protein from a complex mixture of cell membranes, organelles, non-target proteins and other impurities.

### Procedure

- This is for a 10g bacterial pellet in which target protein expression has (hopefully) been induced.
- To a 50ml conical vial, add:

CLA

- 50ml native wash buffer
- 500µl Protease Inhibitor Complex (PIC)
- 20µl DNAse I
- 20mg Lysozyme
- 7.4µl BME
- Resuspend the pellet in the solution.





#### French Press



CLA



- What does the French Press do for us?
  - It forces the cells individually through a tiny opening at high pressure, causing the cells to burst, releasing the contents.
- Steps
  - First clean the French Press with 3 H<sub>2</sub>O rinses.
  - Run the sample through the machine twice to ensure full cell lysis.
  - Clean with 2 more  $H_2O$  rinses and a final rinse with ethanol.



- Spin down the lysate in the low temp. ultra-centrifuge at 15,000rpm for 30 min.
  - Make sure that everything is balanced, and that you are using the correct plastic centrifuge tubes (small rotor only)
  - Start the centrifuge and run like hell
  - All of the insoluble bodies will be in the pellet, such as lipids, organelles and large or hydrophobic proteins
  - The target protein should be in the supernatant
  - Remove 9µl aliquots of the supernatant and pellet for analysis.



### Ni-NTA Beads



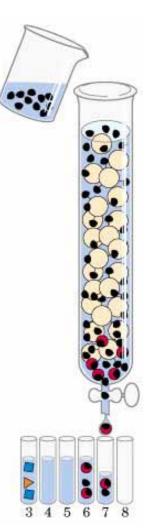
- Next we add Ni-NTA Beads to the supernatant, which are made of a resin infused with nickel.
- Our target protein has been engineered to contain a polyhistidine sequence (6xHis). This region binds tightly to the nickel beads, removing the protein from the solution.
- The vial is nutated for an hour at 4 ° C to allow as much of the protein to bind to the beads as possible.

## Affinity Chromatography

- Next the contents of the flask are poured into a column, and the flowthrough is run over the beads 4 times. Save the F/T.
  - Add 10ml wash buffer 2x, collect fractions
  - Add 10ml hi-salt wash, collect fraction
  - Add 10ml low-conc. imidizol, collect fraction
  - Elute target protein with 20 1ml washes containing a high concentration of imidizol
- Never allow the column to run dry.

CLA

• Hopefully elutions 1-20 contain our purified target protein.





To the first gel add:

<u>UC'LA</u>

- Broad Standard
- Supernatant
- Pellet
- F/T
- Wash Buffer 1
- Wash Buffer 2
- Hi-salt Rinse
- Imidizol Rinse

- To the second gel add:
  - Elution 1
  - Broad Standard
  - Elution 2
  - Elution 3
  - Elution 4
  - Elution 5
  - Elution 6
  - Elution 7

## **Bio-Rad Gel Filtration**

Commercial size exclusion chromatography system

CLA

- The purified protein is loaded into the column, and the target protein, along with any additional impurites is eluted at different times based on protein size
- A mixture of known proteins is loaded in a separate step for size comparisons
- Smaller proteins must flow through pores in the gel beads, causing more collisions and longer elution times
- Larger proteins bypass the pores and simply flow through the spaces between beads, leading to fast elution
- The absorption of the flow-through at 280nm is plotted and the elution time of the target protein is compared to the standards giving a rough estimate of protein size.
- The fraction which contains the target protein should hopefully be pure





## Acknowledgements

- All of the PETC Staff
  - Annie
  - Irina
  - Sum
  - Tung
  - Mark



