

Green Fluorescent Protein I

Overview

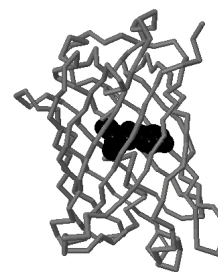
In today's lab, you will purify and study the protein "Green Fluorescent Protein" (GFP) from the *E. coli* like those you will transform with pGLO later in the semester. This lab is an illustration of protein purification and protein properties.

Background I: Green Fluorescent Protein

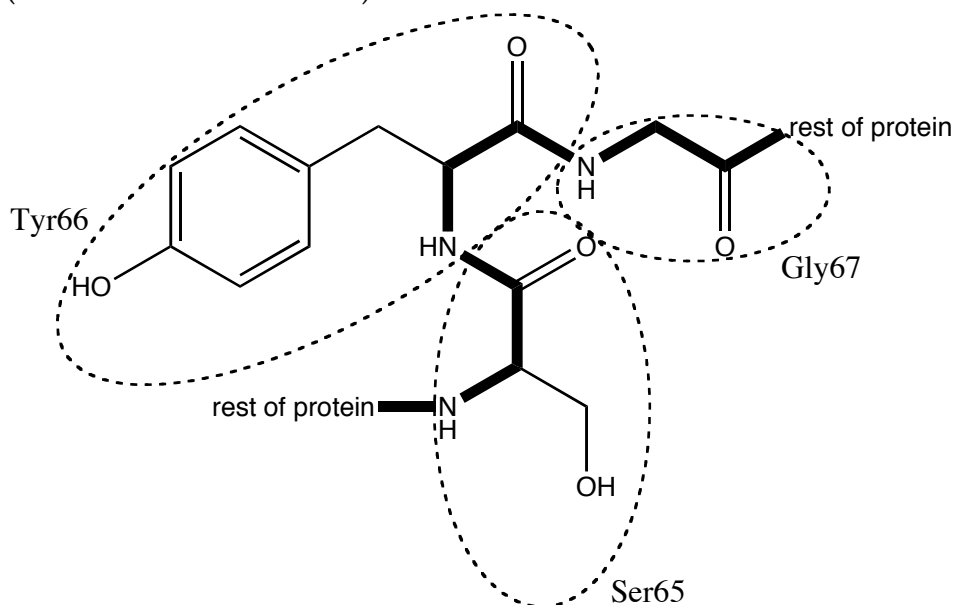
Fluorescence is a process where a molecule absorbs light at one wavelength and then gives off light at a longer wavelength. Most proteins are not fluorescent because, in order to be fluorescent, a molecule must have particular structural elements that are not normally found in proteins. However, the jellyfish *Aequoria victoria* produces a protein, called Green Fluorescent Protein (GFP), that absorbs ultraviolet (UV) light and gives off longer wavelength green light. This makes GFP an interesting protein for study. GFP has been used in a wide variety of experiments exploring protein structure and folding, gene expression, development, etc. In this lab, we will use GFP as an example of a 'typical' protein for biochemical study.

GFP is 238 amino acids in a single protein chain. It is fluorescent because it contains a *chromophore* ("color carrier"), a portion of the molecule that fluoresces. The bulk of the protein surrounds the chromophore and protects it from water, which would prevent the fluorescence.

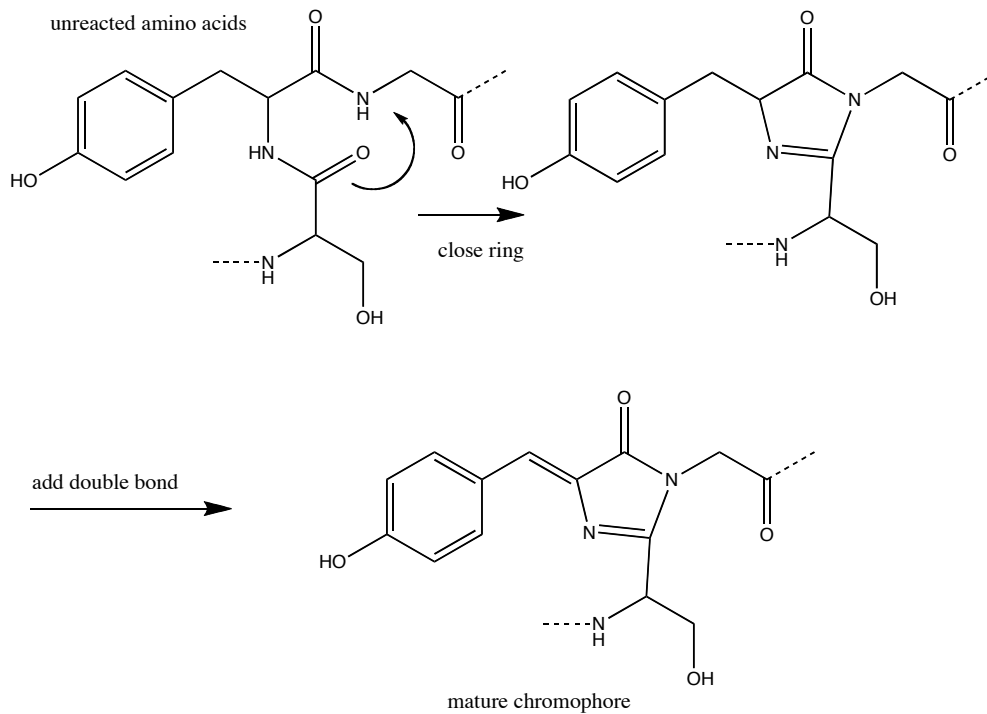
The structure of GFP is shown at right (better views are available at the OLLM for this lab). The chromophore is shown in black spacefill representation; the remaining amino acids are shown as ribbon representations of the backbone. The protein is not hollow; it was rendered this way to show how the protein protects the chromophore from water.



Interestingly, the chromophore is formed from the side chains of three amino acids as the protein folds. The side chains of Ser65, Tyr66, and Gly67 react and link via *covalent* bonds between the side chains to form the chromophore. This unusual reaction - typically side chains only interact via *non-covalent* interactions - is shown below. First, the 3 amino acids after folding, but before the reaction, look like this (backbone shown in bold):



The reaction is shown below:



Background II: Making a “protein of interest”

In general, biologists either study the properties of proteins or use proteins as reagents to accomplish various tasks. In either case, the protein must be *pure* so that researchers can be sure that their observations are due to the protein of interest and not due to various contaminants.

Unfortunately, it is not possible to synthesize large quantities of proteins chemically in a pure form; only cells can make proteins easily and cheaply. This means that producing a pure protein involves three steps:

- 1) **Get the cells to make or “express” the protein.** Any given cell will not necessarily make the desired protein. In our case, it is difficult to obtain jellyfish that make GFP, so we will use the bacterium *E. coli*, which are easy to grow in large quantities. Since normal *E. coli* don't contain the gene for GFP, we will use a strain of *E. coli* to which you have added the GFP gene by *transformation* with the pGLO plasmid.
- 2) **Break open the cells to let the protein out.** Bacterial cells are enclosed by a tough cell wall and a cell membrane. Both must be broken to let the protein out. We will digest away the cell wall with the enzyme lysozyme. We will then break the cell membrane with repeated cycles of freezing and thawing - the ice crystals formed rupture the membrane. These techniques are fierce enough to release the GFP without damaging it.
- 3) **Purify the protein from the other cell contents.** Cells contain thousands of other proteins, DNA, RNA, small molecules, etc. We will separate GFP from all this other 'gunk' in two steps. First, we will discard any material that is not water soluble. We will then separate the GFP from the water-soluble material using *chromatography* that takes advantage of its hydrophobic properties (see next section).

Background III: Hydrophobic Interaction Chromatography (HIC)

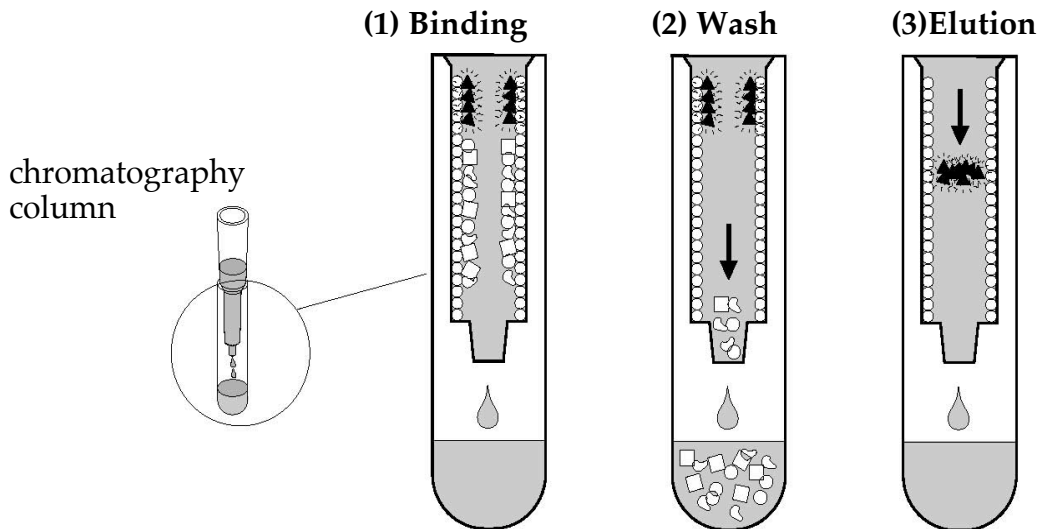
Chromatography is a term that describes a huge and diverse set of methods for separating molecules based on their different properties. In its simplest form, it consists of three steps:

- 1) **Binding.** A mixture of molecules is exposed to a solid material. The molecule of interest binds to the solid via some non-covalent interaction; the others do not.
- 2) **Wash.** The solid is rinsed with a solution (wash buffer) which allows the molecule of interest to remain bound to the solid while washing off the unwanted molecules.
- 3) **Elution.** The solid is rinsed with a different solution (elution buffer) which releases the molecules of interest from the solid. The resulting solution contains (ideally) only the molecule of interest.

In this lab, we will use the fact that GFP has significantly more hydrophobic amino acids on its surface than most proteins. Therefore, it will bind - via the hydrophobic effect - to a solid that also has a hydrophobic surface more tightly than most other proteins.

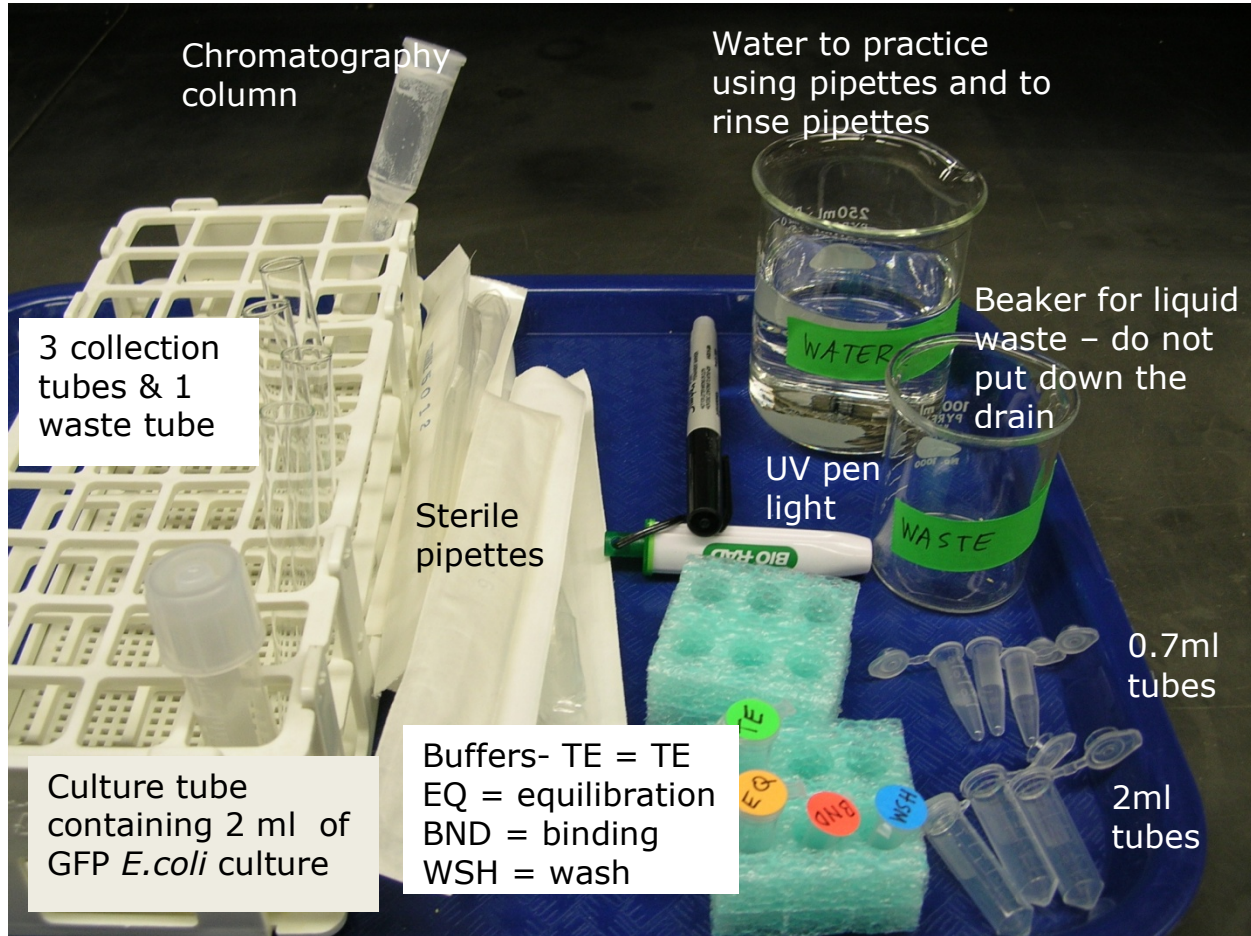
We will also use another important property of the hydrophobic effect to selectively bind and elute GFP. In solutions with a *high* salt concentration, the hydrophobic effect is *strengthened*; in solutions with *low* salt concentrations, the hydrophobic effect is *weakened*. It turns out that it doesn't matter what the salt is; we will use ammonium sulfate ((NH₄)₂SO₄) because it is very soluble in water and does not harm most proteins.

To make the process easier to carry out, the solid material called the "resin" will be packed into a tube called a *chromatography column*. Here's how it will work (the hydrophobic solid resin is indicated by circles; GFP is indicated by black triangles; other molecules are indicated by the other shapes):



- 0) **Equilibration** – a solution containing the same buffer as Step (1) is run through the column to prepare it for chromatography (not shown above).
- 1) **Binding** - a solution containing the cell contents, including GFP, is run through the column in *high salt*. GFP binds via the hydrophobic effect; most other molecules don't. Those that don't stick *flow through* the column and are discarded.
- 2) **Wash** - a solution of *medium salt* is run through the column. GFP remains bound because it has more hydrophobic amino acids on its surface while any remaining other molecules wash out. The molecules in the wash are discarded.
- 3) **Elution** - a solution of *low salt* is run through the column. This disrupts the hydrophobic interaction between the column and GFP so GFP comes off the column in relatively pure form for study.

Kit of Parts:



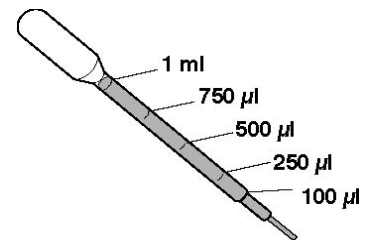
Procedure I: Pipette Practice

In order that your experiment work, it is very important to measure the quantities of various reagents carefully. Although the pipettes we will be using look easy to use, you will need to learn several tricks to using them properly. For this reason, we will start with a little practice.

The pipette you will be using is shown at right. It measures volumes from 100 μ l (micro-liter; 1 millionth of a liter) to 1000 μ l or 1ml (milli-liter; 1 thousandth of a liter). It is calibrated with marks on the side.

To use one, you **first** squeeze out the air **before** you put it in the liquid (otherwise you make bubbles, mix your sample, and cause chaos). You then put it in the liquid and release the bulb **slowly** to draw up the desired amount of liquid. You then **gently** squeeze the bulb to release the measured volume of liquid. You should do the following until all the members of your group can do this with ease.

- One person puts 500 μ l of water in a microtube.
- Another person takes out 250 μ l, discards it, and takes the remaining 250 μ l. There should be no bubbles in the pipette (indicating that less than 250 μ l remained) nor should there be any liquid left (indicating that more than 250 μ l remained).



Procedure II: Look at the Results of your Transformation

Your TA will give you the plates of colonies of *E. coli* bacteria carrying the pGLO plasmid. You should look at them in room lighting and under UV. Do the results match your predictions on the pre-lab?

Procedure III: Purification of GFP

WARNINGS:

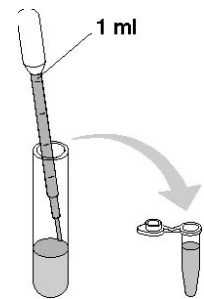
1. In general, the lab is unforgiving of mistakes like using the wrong solution or taking the wrong amount. The construction folks at "This Old House", say "Measure twice; cut once." We'll adapt this to "Check twice; pipette once".
2. Although the *E. coli* strain we use is non-pathogenic (it is not known to cause disease in healthy individuals), you should be careful with it. *Always wear gloves, don't eat or drink in lab, and wash your hands thoroughly when you are all done.*
3. We will be using ultraviolet light in the lab. *Never shine the UV light directly into anyone's eyes.*
4. We will be using liquid nitrogen in this lab. It is very cold (-321° F, -196° C).
 - *Always use insulating gloves when working with liquid nitrogen.*
 - *Always use liquid nitrogen with your TA's supervision.*
 - *Do not get liquid nitrogen on your hands or other body parts.*

Last night, the set up staff started 2ml cultures of *E. coli* that contain the GFP gene. The staff used colonies from plates like those you will make in a later lab. They have been growing and producing GFP until there are roughly 10 billion cells in the culture tube. You will start with this culture.

• Concentrate the bacterial cells in a suitable buffer

1) You will be given a microtube with a liquid culture of *E. coli* containing pGLO and expressing GFP. Vortex the tube briefly to resuspend the cells. Observe it in normal room lighting and then with the UV light. The GFP in the bacteria should show green fluorescence.

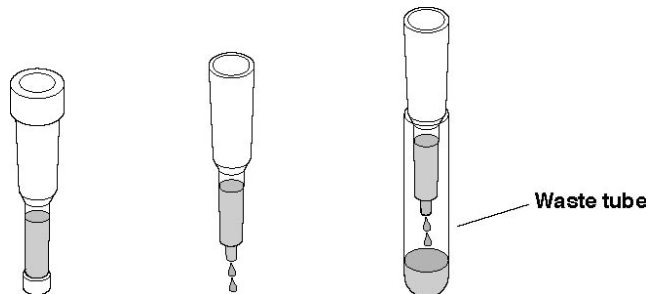
2) Using a marker, label one new microtube with your group initials. Put 1 ml of the resuspended culture in the tube.



3) Spin the microtube for 5 minutes in the centrifuge at maximum speed. Be sure to balance the tubes in the machine (your TA will help with this). If you do not know how to balance the tubes, do not operate the centrifuge.

4) While you are waiting for the centrifuge, prepare the chromatography column. Before performing the chromatography, shake the column vigorously to resuspend the beads of resin. Then shake the column down one final time to bring the beads to the bottom.

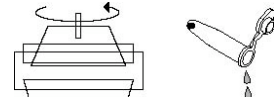
Tapping the column on the table-top will also help settle the beads at the bottom. Remove the top cap and snap off the tab bottom of the chromatography column. Allow all of the liquid buffer to drain from the column into the waste tube (this will take ~3–5 minutes).



You can create a “paper crutch” by folding a small piece of paper, about the size of a match stick, and wedging it between the column and the collection tube. This crutch makes it impossible for an air tight seal to form, and insures that the column will flow.

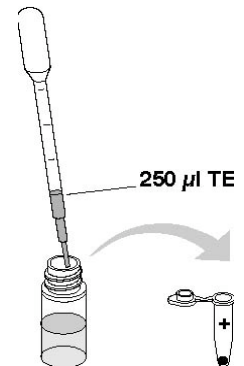
5) While you are waiting for the centrifuge. Prepare the column by adding 2 milliliters of **Equilibration Buffer** (check twice; pipette once) to the top of the column, 1 milliliter at a time using a well rinsed pipette. Drain the buffer from the column into the waste tube.

6) After the bacterial liquid culture has been centrifuged, open the tube and slowly pour off the liquid supernatant above the pellet. After the supernatant has been discarded to the waste beaker, there should be a large bacterial pellet remaining in the tube.



7) Observe the pellet under UV light. Note your observations.

8) Using a new pipette, add 250 μl of **TE Solution** to the tube (check twice; pipette once). Resuspend the bacterial pellet thoroughly by vortexing the tube (your TA will demonstrate). You are done when the pellet is no longer visible on the side of the tube.



- Break open (“lyse”) the bacterial cells

9) Using a rinsed pipette, your TA will add 1 drop of lysozyme solution to the resuspended bacterial pellet. Cap and mix the contents by flicking the tube with your index finger. The lysozyme will start digesting the bacterial cell wall. Observe the tube under the UV light.

10) Let tube sit 5 minutes at room temperature.

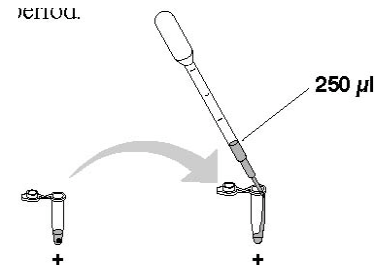
11) Freeze and thaw the cells. Put your microtube into the tube rack and lower it into the LN_2 container by the knob on the top. Leave it in the LN_2 for 10 seconds. Remove the rack by using the knob - **BE CAREFUL NOT TO DIP YOUR FINGERS!** Hold it there until it freezes - about 10 seconds. Take the tube out and thaw it in your hands until it is fully melted. Freeze and thaw one more time to be sure the bacteria are fully lysed.

- Purification Part 1: Remove insoluble material

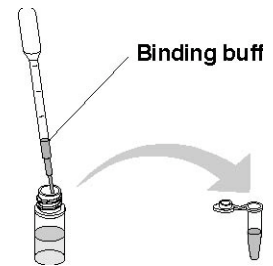
12) After making sure that the suspension is completely melted, place the tube in the centrifuge, have your TA balance the tubes, and pellet the insoluble bacterial debris by spinning for 10 minutes at maximum speed. Label a new microtube with your team’s initials.

13) Obtain 3 collection tubes and label them 1, 2, and 3. Place the tubes in a rack. When the last of the buffer has reached the surface of the column bed, gently place the column on collection tube 1.

14) After the 10 minute centrifugation, immediately remove the microtube from the centrifuge. Examine the tube with the UV light. The bacterial debris should be visible as a pellet at the bottom of the tube. The liquid that is present above the pellet is called the supernatant. Note the color of the pellet and the supernatant. Using a new pipette, **VERY CAREFULLY** transfer 250 μ l of the supernatant into the new microtube. Again, rinse the pipette well for the rest of the steps of this lab.



15) Using the well-rinsed pipette, transfer 250 μ l of **Binding Buffer** (check twice; pipette once) to the microtube containing the supernatant.

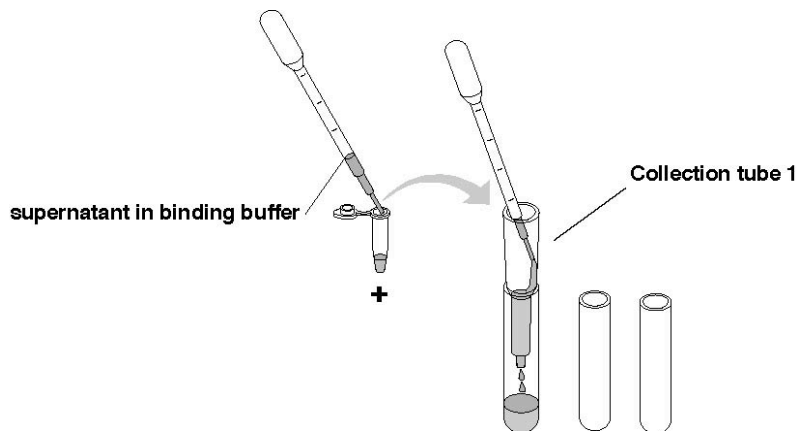


• Purification Part 2: Hydrophobic Interaction Chromatography

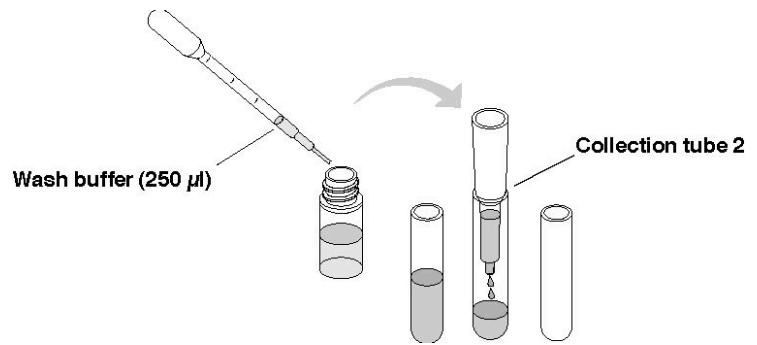
16) Based on the description in the background section of the lab manual, predict which tube the GFP will end up in. Explain your reasoning.

1 2 3

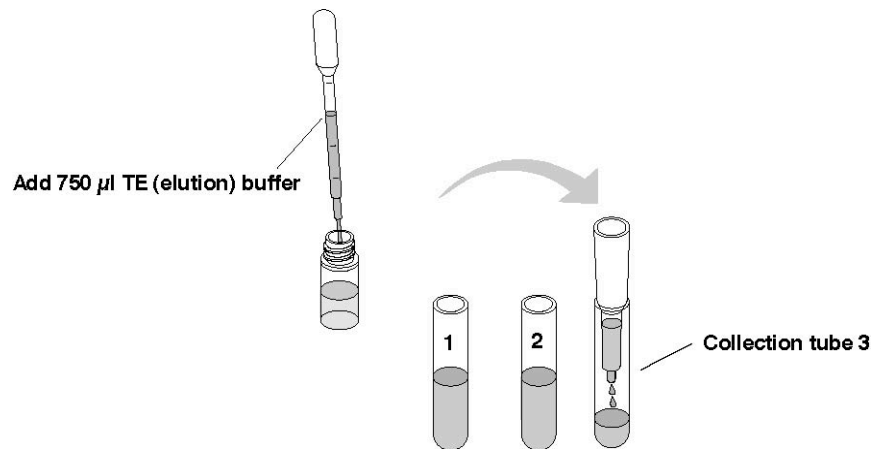
17) Using a new pipette, carefully load **all** of the supernatant into the top of the column by resting the pipette tip against the side of the column and letting the supernatant drip down the side of the column wall. Let the entire volume of supernatant flow into tube 1. Examine the column using the UV light. Where is the GFP?



18) Transfer the column to collection tube 2. Using the rinsed pipette and the same loading technique described above, add **250 μ l of Wash Buffer** and let the entire volume flow into the column. Examine the column using the UV light. Where is the GFP?



19) Transfer the column to tube 3. Using the rinsed pipette, add **750 μ l of TE buffer (Elution Buffer)** and let the entire volume flow into the column. Examine the column using the UV light. Where is the GFP?



20) Examine all of the collection tubes using the UV lamp and note any differences in color between the tubes. Do these results match your prediction from step (16)?

21) You now have purified GFP in one of your tubes. You can then study it in the remainder of the lab.

Procedure IV: Studies of Purified GFP

- Temperature denaturation of GFP

If you heat GFP (or any protein), eventually you reach a temperature where the protein *denatures* - where the energy of the molecules is sufficient to disrupt the *non-covalent* interactions that give the protein its shape. The protein is then said to be *denatured*. Different proteins denature at different temperatures depending on the particular non-covalent bonds that give them their shape. When GFP denatures, the chromophore is exposed to the surrounding water and it is no longer fluorescent.

22) Take three small samples of your purified GFP - about 100 μ l each - and put each in a new glass test tube. Label each tube with your initials.

23) Put one tube at in the 50°C water bath, one in the 75°C water bath, and one in the boiling (100°C) water bath. Let them sit for 5 minutes. Remove the tubes with a metal tube holder or a paper towel. Be careful, they're HOT!

24) Observe the tubes under UV light. Which are fluorescent; which are not? What does this tell you about the temperature required to denature GFP?

- Denaturation by pH

pH measures the relative acidity of a solution. The charges on the side chains of certain amino acids (Arg, Lys, His, Glu, and Asp) depend on the pH as follows:

- At **neutral** pH (7.0)
 - (-)-charged side chains (Asp and Glu) are *mostly* charged
 - (+)-charged side chains (Arg, Lys, and His) are *mostly* charged.
- At **higher** pHs (> 7.0) (more alkaline; less acidic),
 - (-)-charged side chains (Asp and Glu) are *more* likely to be charged
 - (+)-charged side chains (Arg, Lys, and His) are *less* likely to be charged.
- At **lower** pHs (< 7.0) (more acidic; less alkaline),
 - (-)-charged side chains (Asp and Glu) are *less* likely to be charged
 - (+)-charged side chains (Arg, Lys, and His) are *more* likely to be charged.

As a result, the strength of ionic interactions involving these amino acids will change as the pH changes. The buffers we use have a pH of roughly 8.0, which is close to neutral. When GFP denatures, the chromophore is exposed to the surrounding water and it is no longer fluorescent.

25) Each group will test a different pH. Add about 100 μ l of your pH solution to the remainder of your GFP using a new pipet, wait a few minutes, and observe it under UV light. Is the GFP still native at this pH?

26) Clean up; you are done. Specifically:

- All *E. coli* waste should go into the *E. coli* waste jar including the liquid in the waste beaker.
- All used plastic pipettes and used 2ml tubes should be put in the orange autoclave bags.
- We re-use the 15 ml tubes and the chromatography columns; place them in the labeled bin.
- Gloves and pipet wrappers go in the regular trash.

Lab report

There is no lab report for this lab.