

Module 2: Western Blot

BMES Cell Team

Fall 2020



Outline

- Protein Basics
- What is a Western Blot?
- Western Blot Protocol
 - BCA Assay
 - SDS Gel Electrophoresis
 - Immunoblotting
- Western Blot Video
- Pipetting basics



Proteins

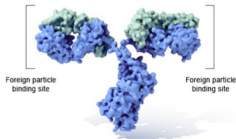
- **Definition:** **Proteins** are macromolecules made of amino acids.

→ Chains of amino acids make up proteins

→ Proteins have a wide structural range → large functional range

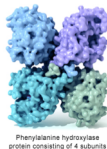
→ Key players in organism's metabolic & regulatory activity

Immunoglobulin G (IgG)



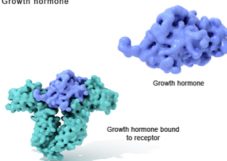
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Phenylalanine hydroxylase



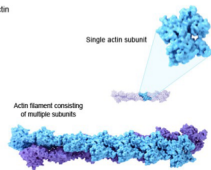
U.S. National Library of Medicine

Growth hormone



U.S. National Library of Medicine

Actin

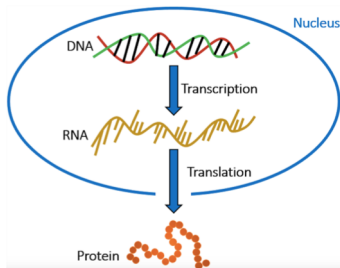


U.S. National Library of Medicine

What is Western Blotting?

- **Definition:** A **Western Blot** separates and identifies target proteins.

- Two stage procedure confirms protein presence and quantifies target
 - Proteins separated by size through gel electrophoresis
 - Target proteins visualized through immunostaining



SN❄W

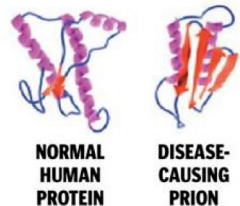
DR💧P

S	= SOUTHERN	= DNA	= D
N	= NORTHERN	= RNA	= R
O	= OOOOOOOO	= OOOO	= O
W	= WESTERN	= PROTEIN	= P

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Dyeing techniques mnemonic

Western Blot Utility

- Understand mechanisms of cell behavior
 - Protein's primary function in organism
- Probe for a specific disease
 - **Viral:**
 - HIV
 - **Bacterial:**
 - Meliodosis
 - **Prion:**
 - Creutzfeldt-Jakob disease



Basic Protocol for a Western Blot

1. Lyse cells and collect proteins
2. BCA Assay
 - Calculate sample protein concentration
3. SDS Gel Electrophoresis
 - Separate proteins by size
4. Transfer proteins to membrane
5. Immunoblot target proteins
 - Antibody Binding
6. Image membrane

Western Blot Protocol



Sample extraction



Run Gel



Transfer proteins to membrane



Antibodies incubations and washes



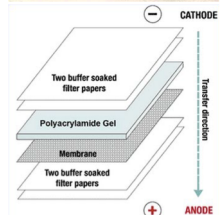
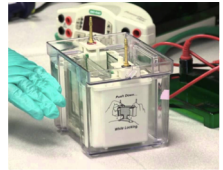
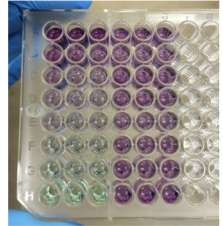
Film to visualize protein bands



Cry

Basic Protocol for a Western Blot

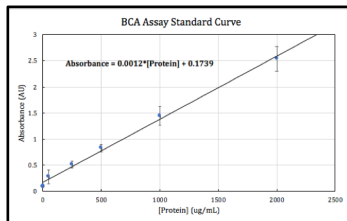
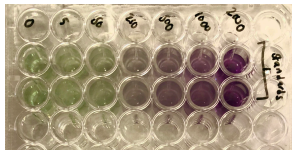
1. Lyse cells and collect proteins
2. **BCA Assay**
 - Calculate sample protein concentration
3. **SDS Gel Electrophoresis**
 - Separate proteins by size
4. Transfer proteins from gel to membrane
5. **Immunoblot target proteins**
 - Antibody Binding
6. Image membrane



BCA Assay

- **Definition:** A bicinchoninic acid assay (**BCA Assay**) uses absorbance readings to determine total protein concentration in a sample.

- First, create a **standard curve**
 - Concentration of protein in each well is known
- Then, add experimental samples
 - Concentration of protein is unknown
- Use standard curve absorbance readings to correlate protein concentration to absorbance
- Use standard curve to calculate experimental protein concentration



How does a BCA Assay work?

- Step 1: Biuret Reaction

- Green Cu^{2+} in BCA reagent binds to sample protein

→ reduction of cuprous Cu^{1+}

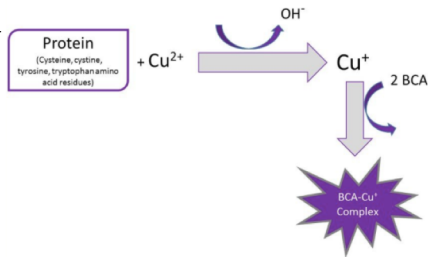
- Step 2: BCA and Copper Chelation

- 2 BCA molecules bind to Cu^{1+}

→ purple chelated complex

- Step 3: Measure Absorbance

- Purple complex absorbs maximally at 562 nm
 - Absorbance \propto purple complexes \propto peptide

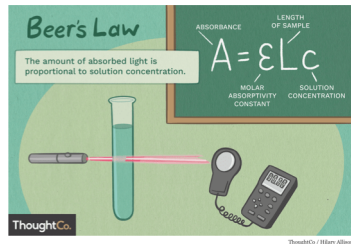


Beer's Law

- **Definition:** **Beer's Law** relates a sample's absorbance reading to total protein concentration.

$$A = \epsilon \cdot L \cdot C$$

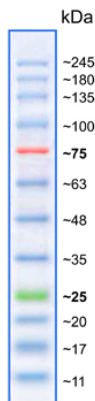
- A = absorbance reading from plate reader
- ϵ = molar absorptivity constant
- L = path length
- C = protein concentration
 - As ϵ and L are constant, there is a linear relationship between absorbance and protein concentration



SDS-PAGE

- **Definition:** Sodium dodecyl sulfate-polyacrylamide gel electrophoresis (**SDS-PAGE**) separates proteins by size.

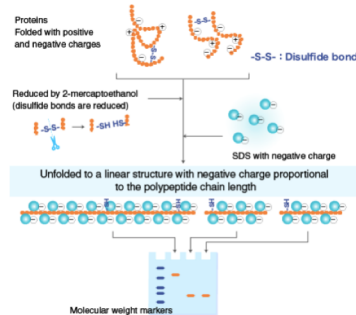
- First, load the protein ladder at the two ends of the well
 - Set of standards that allow us to estimate protein size
 - Dalton (Da) = atomic mass unit
 - kDa = 1000 Da
- Then, load your sample into the central wells
- Run the gel and use the ladder to estimate protein size



How does SDS-PAGE work?

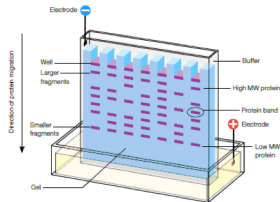
Key Reagents

- Sodium Dodecyl Sulfate (**SDS**)
 - Anionic detergent that binds to protein side chains → disruption noncovalent bonds → protein denaturation
 - Coats denatured protein in uniform negative charge → charge of protein \propto length of protein
- Beta-mercaptoethanol (β ME)
 - Reduces disulfide bonds in protein → disruption covalent bonds → protein linearization



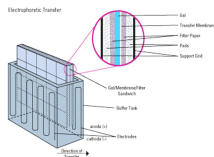
How does SDS-PAGE work?

- Polyacrylamide (PA)
 - Water soluble polymer
 - 3D networks of polyacrylamide → porous gel
 - Smaller proteins can travel faster through the porous gel
- Gel Electrophoresis (GE)
 - When placed in an electric field, the negatively charged proteins will migrate toward the positive electrode
 - Since smaller proteins can travel faster through a porous gel, loading protein samples into a gel then creating an electric field around the gel **separates proteins by size**



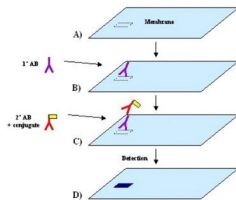
Protein Transfer

- Need to transfer proteins from gel to nitrocellulose membrane
 - Antibodies cannot bind to proteins when they are on the gel
 - Must transfer proteins onto a nitrocellulose membrane and retain the gel electrophoresis size sorting
- Transfer proteins using **electroblotting**
 - Align the gel and the membrane
 - Use an electric current to pull negatively charged proteins toward a positively charged anode and onto the membrane

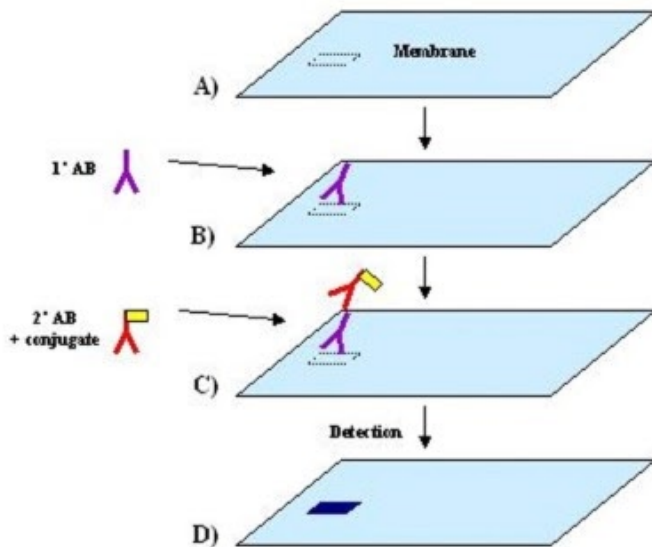


Immunoblotting (What Makes it a Western Blot!)

- **Definition:** Immunoblotting uses antibodies to identify proteins.
- Antibodies are proteins in the immune system that target specific antigens
- Primary Antibody: binds to target protein
 - Loading Control: Actin (constitutively expressed in all cells)
- Secondary antibody: binds to primary antibody and amplifies the signal
 - Primary and secondary antibodies must be from a different species than the target protein
 - If not, will have non-specific binding

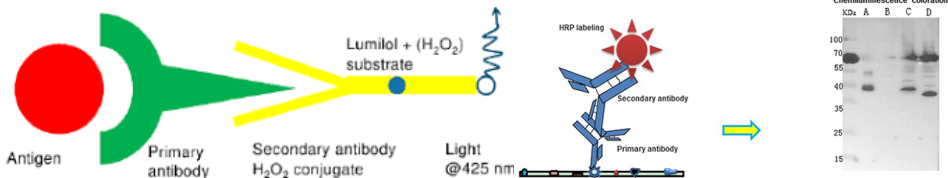


How does Immunoblotting work?



Imaging

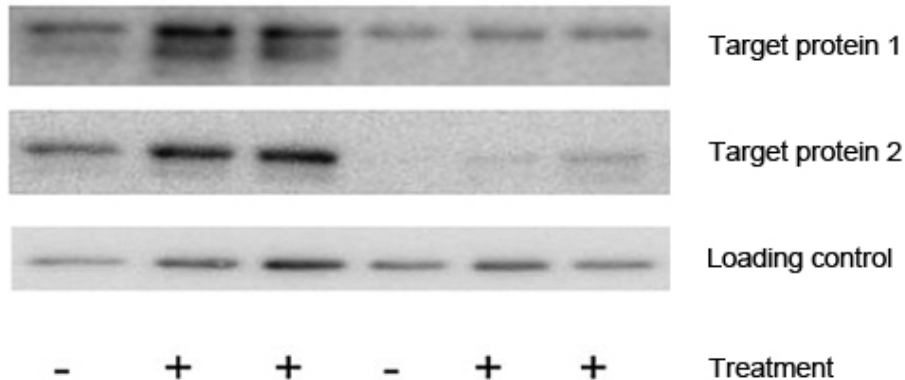
- Chemiluminescence
 - Chemical reaction between ECL substrate and HRP enzyme in secondary antibody
 - Releases energy as light
 - One of the easiest ways to understand internal cell behavior and controls



Interpreting a Western Blot

- Loading control band:
 - should be the same in all samples
 - If loading control is not the same, result is invalid
- Band position on gel:
 - Different sized proteins show up at different heights
 - Larger proteins show up closer to the original well position
- Band intensity:
 - The darker the band, the more protein is present

Interpreting a Western Blot



- Loading control band is the same in all trials
- Greater target protein intensity in wells 2 and 3

Western Blot Video

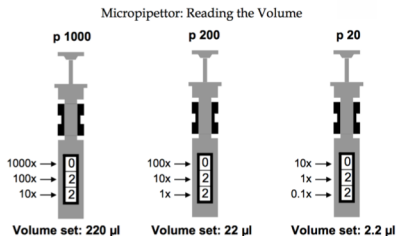


<https://youtu.be/yUstng0npaY>

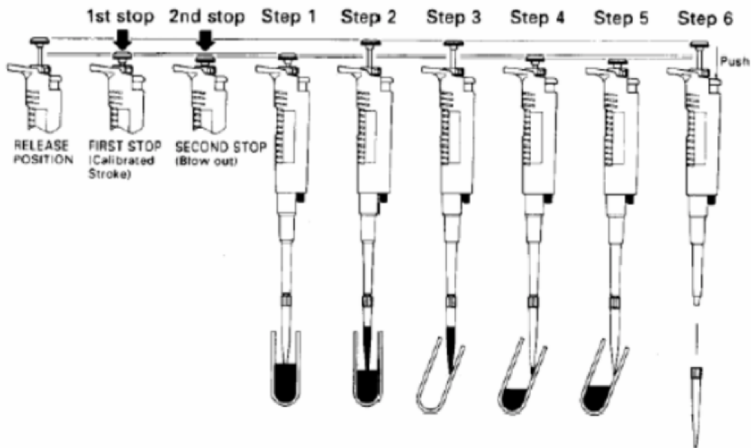
Micropipettes

- Definition:** A **micropipette** is a laboratory instrument used to measure small volumes (on the order of microliters).

- Micropipette sizes
 - P20: 2 – 20 μ L
 - P200: 20-200 μ L
 - P1000: 100-1000 μ L
- Be mindful of which size you are using



How to use a Micropipette



<https://www.youtube.com/watch?v=TMFeV9h6zEA>

Module 2 Worksheet

BMES Cell Team

Fall 2020



Overview of Module 2

1. Name one protein function:

2. What is the purpose of a Western blot?

3. What is the purpose of a BCA Assay? Which metal is present in BCA reagent?

4. What is the purpose of SDS-PAGE?

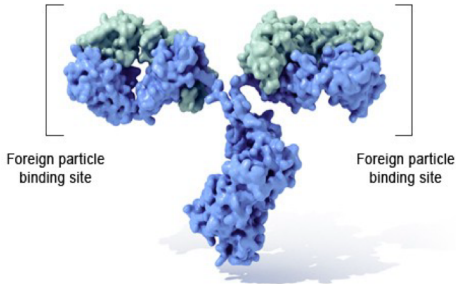
5. Why do we need to transfer proteins from a gel to a membrane?

Overview of Module 2

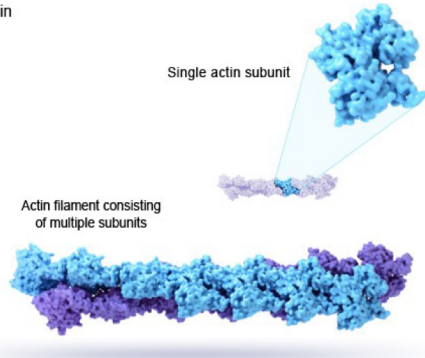
1. Name one protein function:

antibody, structural component, messenger, enzyme, regulatory

Immunoglobulin G (IgG)



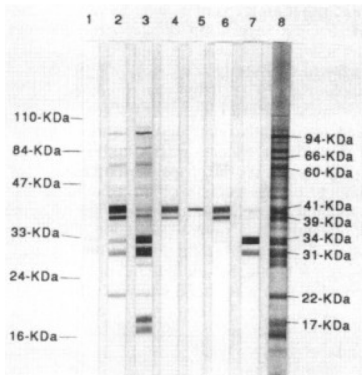
Actin



Overview of Module 2

2. What is the purpose of a Western blot?

separate and identify target protein to better understand the protein's role and how protein mutations can result in disease state



**You know what goes great
with the Corona virus?**



Lyme Disease

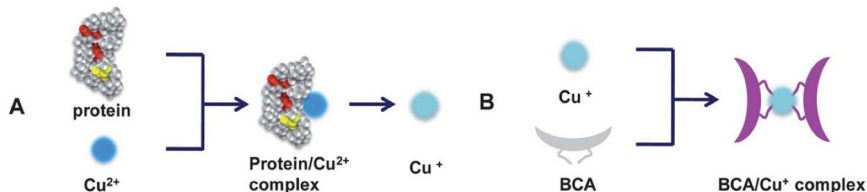


Overview of Module 2

3. What is the purpose of a BCA Assay? Which metal is present in BCA reagent?

A BCA Assay allows us to calculate sample protein concentration using absorbance readings.

The copper ion (Cu^{2+}) is reduced during the Biuret Reaction in a BCA Assay

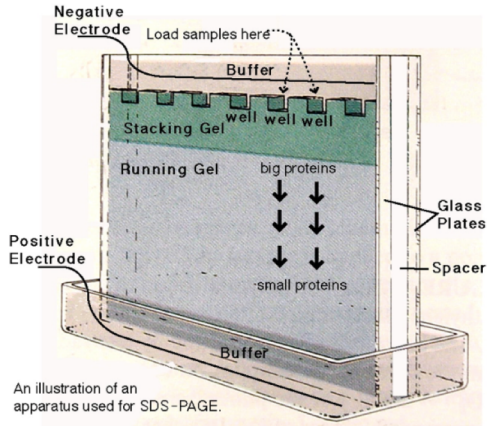


Overview of Module 2

4. What is the purpose of SDS-PAGE?

Gel electrophoresis separates protein fragments by size.

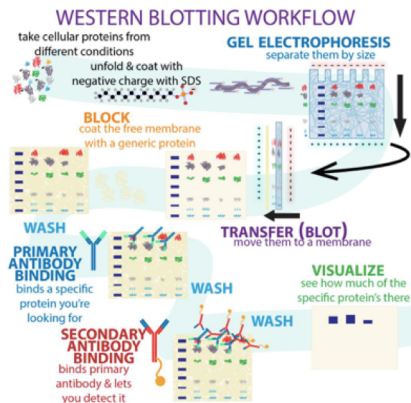
1D SDS-PAGE



Overview of Module 2

5. Why do we need to transfer proteins from a gel to a membrane?

The antibodies utilized in immunoblotting cannot bind to the protein when the protein is on the gel.



Practice Preparing a BCA Assay

1. Calculate serial dilution volumes.

We will start with a 20 ug/mL BSA solution.

- Need 25 uL of standard for each well x 3 replicates = 75 uL per standard
- Always make a little extra: _____ uL per standard

BSA Concentration (ug/mL)	BSA Solution Added (uL)	Water Added (uL)	Total Volume (uL)	Final Volume (uL)
2000 ug/mL	160 uL of 2000 ug/mL	0	160	160 - 80 = 80
1000 ug/mL	80 uL of 2000 ug/mL	80	160	160 - ____ = ____
500 ug/mL	____ uL of 1000 ug/mL	____	160	160 - ____ = ____
250 ug/mL	____ uL of 500 ug/mL	____	160	160 - ____ = ____
50 ug/mL	____ uL of 250 ug/mL	____	160	160 - ____ = ____
5 ug/mL	____ uL of 50 ug/mL	____	160	160 - ____ = ____
0 ug/mL	0	160	160	160

Practice Preparing a BCA Assay

1. Calculate serial dilution volumes.

We will start with a 20 ug/mL BSA solution.

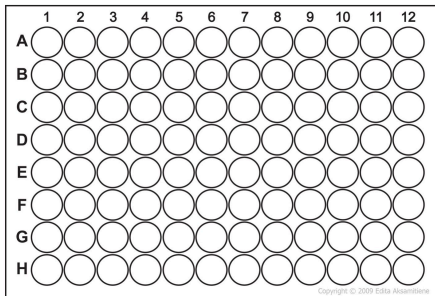
- Need 25 uL of standard for each well x 3 replicates = 75 uL per standard
- Always make a little extra: 80 uL per standard

BSA Concentration (ug/mL)	BSA Solution Added (uL)	Water Added (uL)	Total Volume (uL)	Final Volume (uL)
2000 ug/mL	160 uL of 2000 ug/mL	0	160	160 - 80 = 80
1000 ug/mL	80 uL of 2000 ug/mL	80	160	160 - 80 = <u>80</u>
500 ug/mL	<u>80</u> uL of 1000 ug/mL	<u>80</u>	160	160 - 80 = <u>80</u>
250 ug/mL	<u>80</u> uL of 500 ug/mL	<u>80</u>	160	160 - 80 = <u>80</u>
50 ug/mL	<u>32</u> uL of 250 ug/mL	<u>128</u>	160	160 - 16 = <u>144</u>
5 ug/mL	<u>16</u> uL of 50 ug/mL	<u>144</u>	160	160 - 0 = <u>160</u>
0 ug/mL	0	160	160	160

Practice Preparing a BCA Assay

2. Design layout for 96 well plate:

- a. Number of standards _____ x Number of repeats _____ = _____ wells
- b. Number of samples 2 x Number of repeats 2 = 4 wells
- c. Total number of wells = Standard wells _____ + Sample wells 4 = _____ wells
- d. Label the well plate below with standard and sample placement



Practice Preparing a BCA Assay

2. Design layout for 96 well plate:

- a. Number of standards 7 x Number of repeats 3 = 21 wells
- b. Number of samples 2 x Number of repeats 2 = 4 wells
- c. Total number of wells = Standard wells 21 + Sample wells = 4 = 25 wells
- d. Label the well plate below with standard and sample placement:

	standards					samples							
	1	2	3	4	5	6	7	8	9	10	11	12	
A	2000 $\mu\text{g/ml}$	2000 $\mu\text{g/ml}$	2000 $\mu\text{g/ml}$										
B	1000 $\mu\text{g/ml}$	1000 $\mu\text{g/ml}$	1000 $\mu\text{g/ml}$			PC9	PC9						
C	500 $\mu\text{g/ml}$	500 $\mu\text{g/ml}$	500 $\mu\text{g/ml}$										
D	250 $\mu\text{g/ml}$	250 $\mu\text{g/ml}$	250 $\mu\text{g/ml}$			PC9 AHL-RQ	PC9 AHL-RQ						
E	50 $\mu\text{g/ml}$	50 $\mu\text{g/ml}$	50 $\mu\text{g/ml}$										
F	10 $\mu\text{g/ml}$	10 $\mu\text{g/ml}$	10 $\mu\text{g/ml}$										
G	0 $\mu\text{g/ml}$	0 $\mu\text{g/ml}$	0 $\mu\text{g/ml}$										
H													

Practice Preparing a BCA Assay

3. Prepare reaction buffer:

- a. Total number of wells _____ x 200 uL/well = _____ uL reaction buffer
- b. Always make a little extra: _____ uL
- c. Combine reagents A, B, and C in ratio 25:24:1 to make reaction buffer
 - i. Reagent A = (total volume of buffer _____ uL)(25/50) = _____ uL
 - ii. Reagent B = (total volume of buffer _____ uL)(24/50) = _____ uL
 - iii. Reagent C = (total volume of buffer _____ uL)(1/50) = _____ uL

3. Prepare reaction buffer:

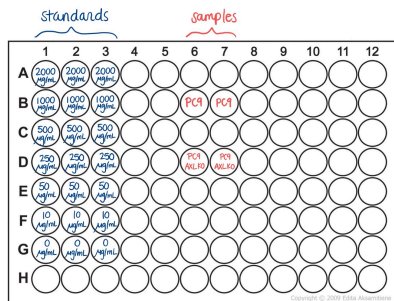
- a. Total number of wells 25 x 200 uL/well = 5000 uL reaction buffer
- b. Always make a little extra: 5500 uL
- c. Combine reagents A, B, and C in ratio 25:24:1 to make reaction buffer
 - i. Reagent A = (total volume of buffer 5500 uL)(25/50) = 2750 uL
 - ii. Reagent B = (total volume of buffer 5500 uL)(24/50) = 2640 uL
 - iii. Reagent C = (total volume of buffer 5500 uL)(1/50) = 110 uL

Practice Making a Standard Curve

→ Consider: You run this plate ...
and get the below absorbance readings

→ Create a standard curve

→ Use the standard curve to determine
the sample concentration of protein



1	2	3	4	5	6	7
A	2.799	2.482	2.343	0.047	0.049	0.048
B	1.644	1.313	1.368	0.048	0.253	0.246
C	0.904	0.806	0.774	0.049	0.047	0.049
D	0.566	0.529	0.452	0.047	0.305	0.26
E	0.206	0.201	0.438	0.048	0.048	0.047
F	0.106	0.119	0.101	0.047	0.048	0.048
G	0.097	0.096	0.096	0.048	0.048	0.048

Practice Making a Standard Curve

Step 1: Determine the average and standard deviation of replicates

1	2	3	4	5	6	7
A	2.799	2.482	2.343	0.047	0.049	0.048
B	1.644	1.313	1.368	0.048	0.253	0.246
C	0.904	0.806	0.774	0.049	0.047	0.049
D	0.566	0.529	0.452	0.047	0.305	0.26
E	0.206	0.201	0.438	0.048	0.048	0.047
F	0.106	0.119	0.101	0.047	0.048	0.048
G	0.097	0.096	0.096	0.048	0.048	0.048



[Protein] (ug/mL)	AVG Absorbance (AU)	STD Absorbance (AU)
2000	2.541333333	0.233718492
1000	1.441666667	0.17737061
500	0.828	0.067734777
250	0.515666667	0.058157831
50	0.281666667	0.135411718
5	0.108666667	0.009291573
0	0.096333333	0.00057735

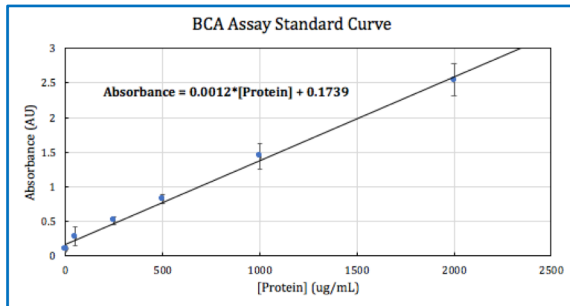
Practice Making a Standard Curve

Step 2: Plot the average and standard deviation of replicates

[Protein] (ug/mL)	AVG Absorbance (AU)	STD Absorbance (AU)
2000	2.541333333	0.233718492
1000	1.441666667	0.17737061
500	0.828	0.067734777
250	0.515666667	0.058157831
50	0.281666667	0.135411718
5	0.108666667	0.009291573
0	0.096333333	0.00057735

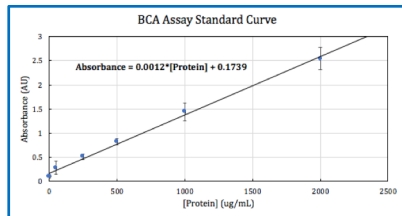
→ Display equation of trendline

→ Note y-intercept



Practice Making a Standard Curve

Step 3: Use standard curve equation and absorbance data to calculate sample protein concentration



1	2	3	4	5	6	7
A	2.799	2.482	2.343	0.048	0.048	0.048
B	1.644	1.313	1.368	0.048	0.253	0.246
C	0.904	0.806	0.774	0.048	0.047	0.049
D	0.566	0.529	0.452	0.047	0.305	0.26
E	0.206	0.201	0.438	0.048	0.048	0.047
F	0.106	0.119	0.101	0.047	0.048	0.048
G	0.097	0.096	0.096	0.048	0.048	0.048



$$Absorbance = 0.0012 * [Protein] + 0.1739 \rightarrow [Protein] = \frac{Absorbance - 0.1739}{0.0012}$$

Sample	Absorbance 1	Absorbance 2
PC9	0.253	0.246
PC9 AXO KO	0.305	0.26

Sample	[Protein] 1	[Protein] 2
PC9	65.91666667	60.08333333
PC9 AXO KO	109.25	71.75

Sample	AVG [Protein] (ug/mL)	STD [Protein] (ug/mL)
PC9	63	4.124789557
PC9 AXO KO	90.5	26.51650429