

SODIUM DODECYL SULPHATE -POLYACRYLAMIDE
GEL ELECTROPHORESIS
[SDS-PAGE]

Lab#8

BCH 333

SDS-PAGE:

- **SDS-PAGE: Sodium Dodecyl Sulfate-Polyacrylamide Gel Electrophoresis.**
- The separation of macromolecules in an electric field is called **electrophoresis**.
- Separates proteins by electrophoresis using a discontinuous polyacrylamide gel.
- SDS-PAGE aims to separate and identify proteins according to their **molecular weight**.

- **Applications:**
 - Quantifying.
 - Comparing (relative abundance).
 - Characterizing proteins (e.g., determining MW of proteins).
 - Checking the purity of protein samples.

Principle of SDS-PAGE:

- In general, fractionation by gel electrophoresis is based on **differences in size, shape and net charge of macromolecules** (in molecules native conditions using native gel electrophoresis).
- **Native gel electrophoresis**, where you separate proteins under native conditions, cannot distinguish between these effects and therefore, proteins of different sizes may have the same mobility in native gels.

➔ In **SDS-PAGE**, this problem is overcome by:

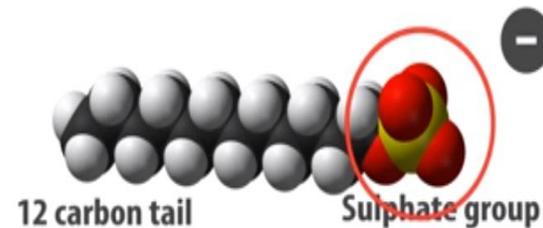
1. Using an anionic detergent, Sodium Dodecyl Sulfate (**SDS**), which binds strongly to most proteins.
2. Using disulfide reducing agent such as **β-mercaptoethanol** or **DTT** (dithiothreitol).

—————➔ **Proteins are unified in charge and shape, thus they separate based on size**

Principle of SDS-PAGE:

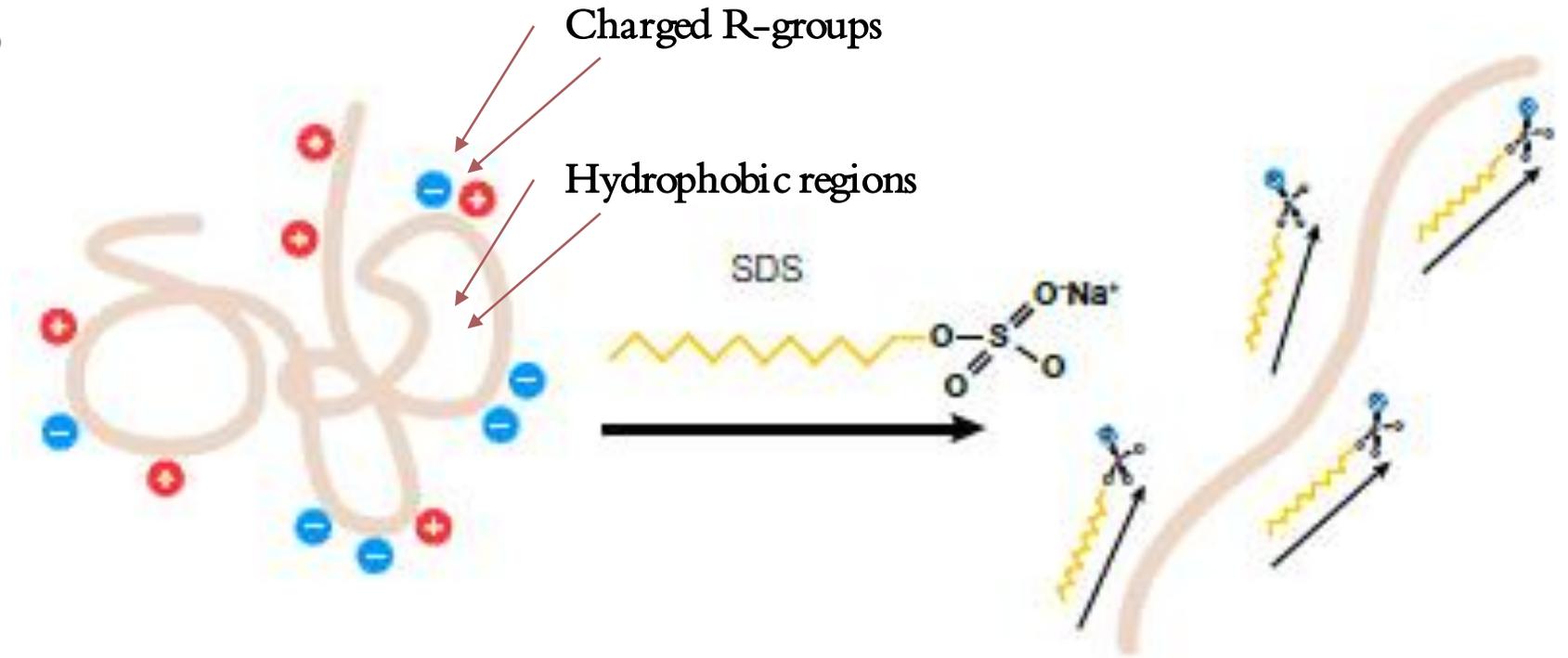
- Sodium Dodecyl Sulfate (SDS) is a **detergent** which **denature proteins** by binding to the **hydrophobic regions**, all **non-covalent bonds** will disrupt, and the proteins acquire a **negative net charge**.
 - A Concurrent treatment with a **disulfide reducing** agent such as β -mercaptoethanol or DTT (dithiothreitol) further **breaks down the strong bound disulfide**.
- ➔ So, the proteins samples are having uniformed structure and charge (-ve).

The carbon tail has a neutral charge
and the sulphate group is negatively charged



Molecular structure of SDS

Effect of SDS



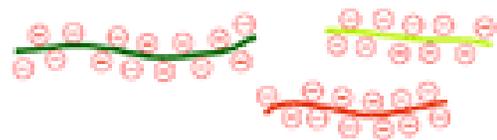
Effect of SDS on the conformation and charge of a protein. Source: bio-rad.com

The carbon tail has a neutral charge
and the sulphate group is negatively charged

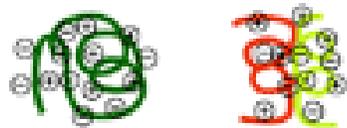


Molecular structure of SDS

(A) SDS PAGE

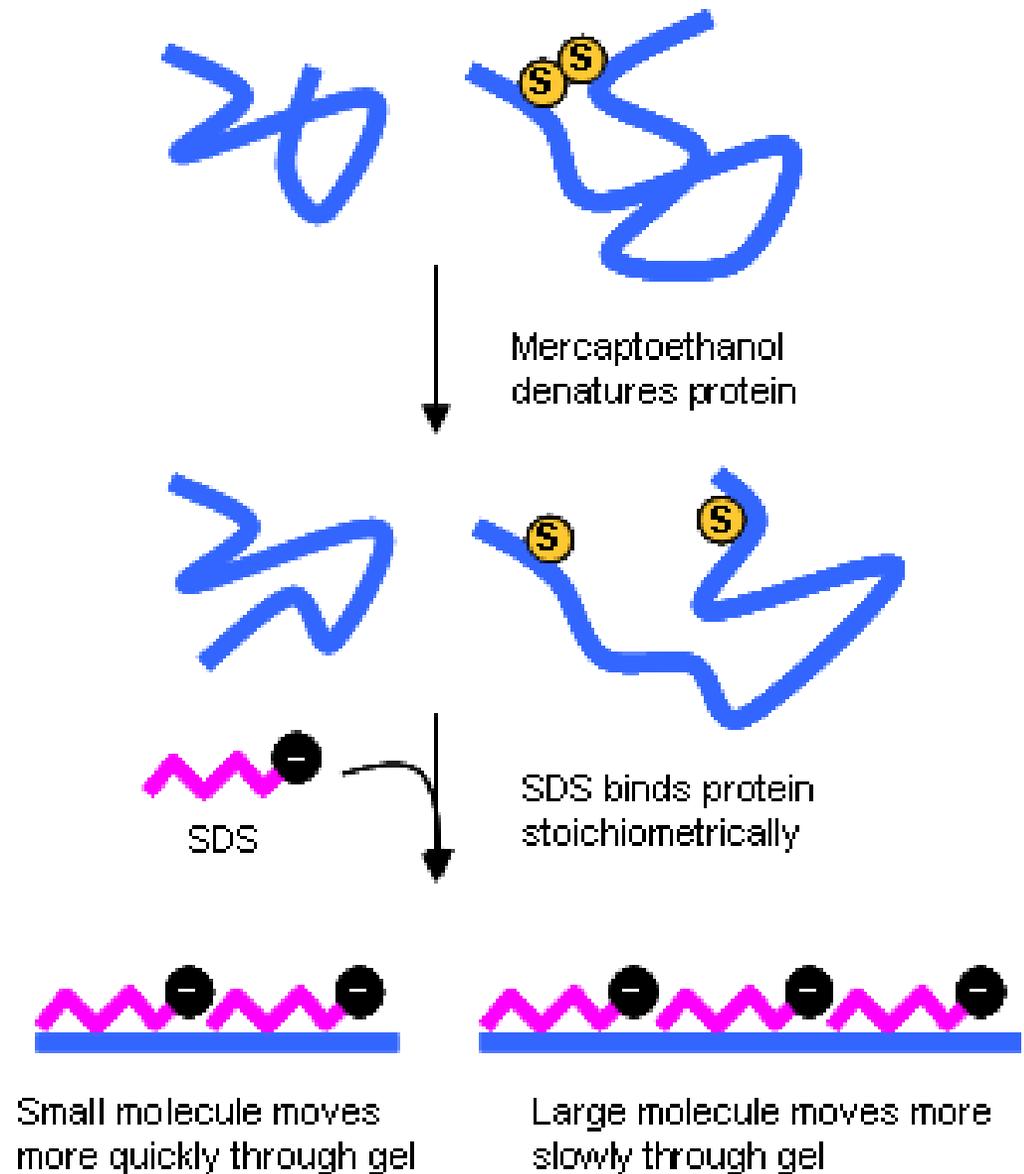


(B) Conventional native PAGE



SDS vs native PAGE

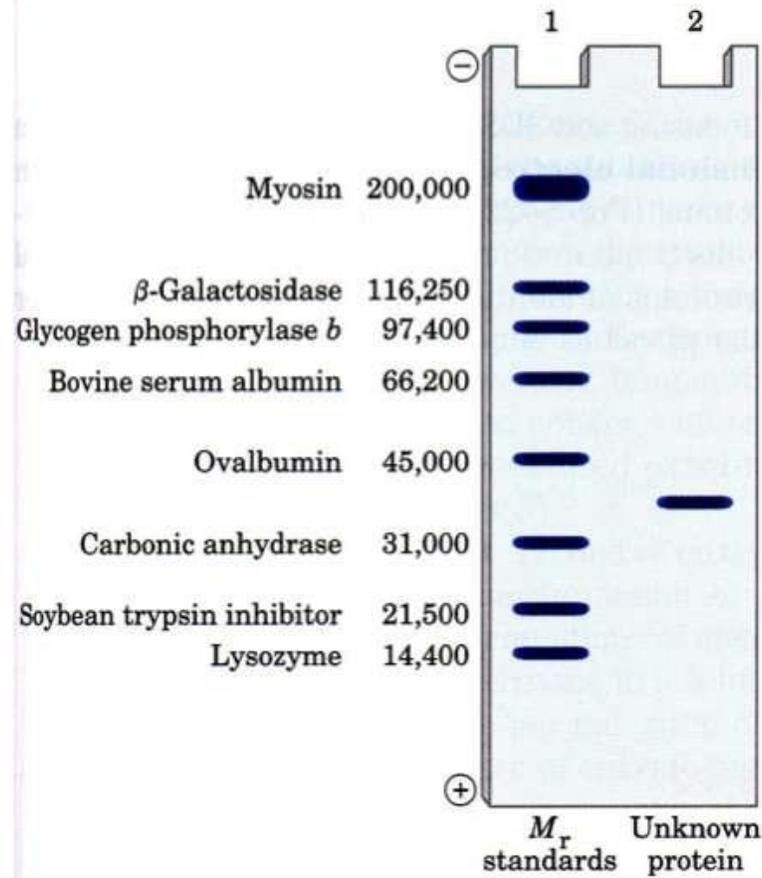
Effect of β -mercaptoethanol



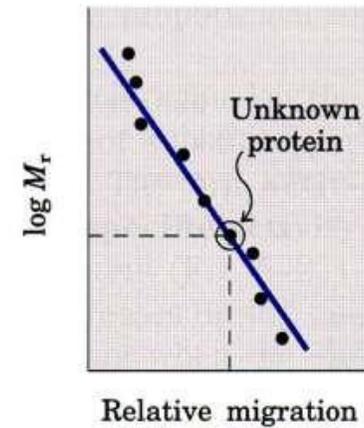
Effect of β -mercaptoethanol

Principle of SDS-PAGE:

- Because the charge-to-mass ratio is nearly the same among SDS-denatured polypeptides, the final separation of proteins is dependent almost entirely on the differences in relative molecular mass of polypeptides (separation will depend on their molecular weight only).
 - ➔ i.e The electrophoretic mobility of the molecules is now considered to be a function of their size (the protein samples are having uniformed **structure** and **charge** ➔ the separation will depend on their molecular weight only).
- Polyacrylamide gels **restrain larger** molecules from migrating **as fast as smaller molecules**.
- Thus, small proteins migrate faster through the gel under the influence of the applied electric field, whereas large proteins are successively retarded, due to the sieving effect of the gels.
 - ➔ i.e the migration of the (SDS-treated proteins), towards the anode (+) is inversely proportional to the logarithms of their molecular weights, or more simply expressed:
Small proteins migrate faster through the gel

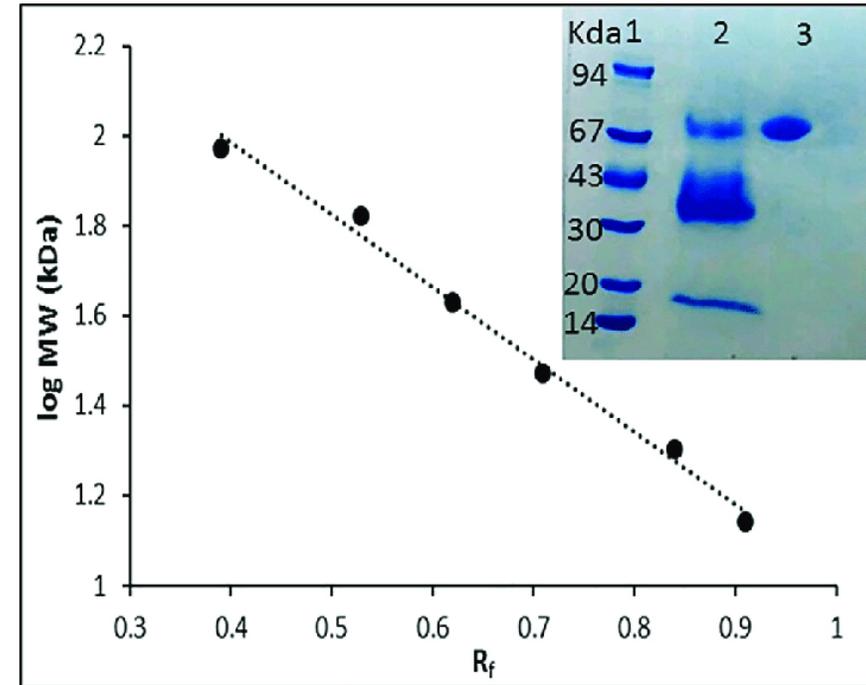
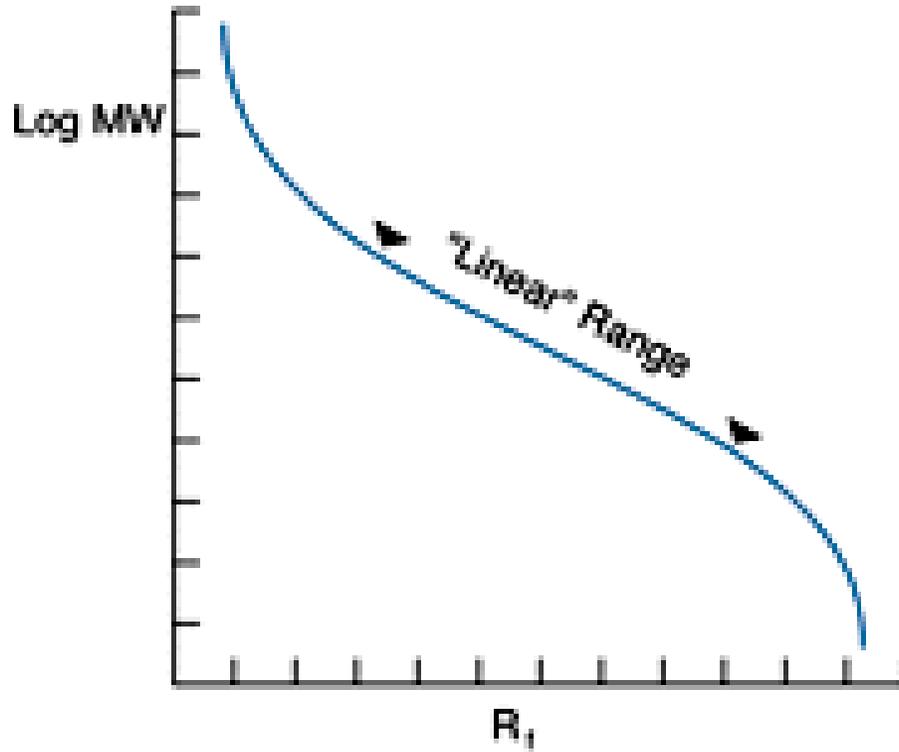


(a)



(b)

SDS-treated proteins have very similar charge-to-mass ratios and similar shapes. During SDS-PAGE, the rate of migration of SDS-treated proteins is effectively determined by **molecular weight**.



Graph of log MW vs. R_f is sigmoidal; it is nearly linear for a range of molecular weights excluding very small and very large MW.

Practical part



OBJECTIVES:

-
- To separate and calculate the molecular size of proteins by comparing the separated bands with a known standard molecular weight marker.



SDS-Polyacrylamide Gel Electrophoresis preparation



1. Sample Preparation:

40 μl of protein sample + 10 μl of disruption buffer → boil the mixture 3 min at 99 °C, why?

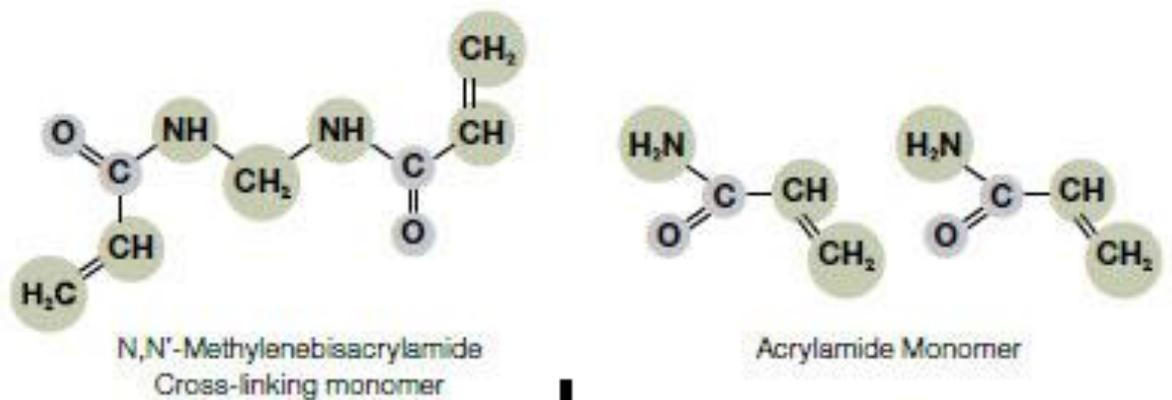
• Disruption buffer [loading buffer] contains:

- 10% (w/v) SDS [?]
- 1M Tris/HCl, pH 6.8
- Glycerol [?]
- β -Mercaptoethanol [?]
- Bromophenol blue [?]

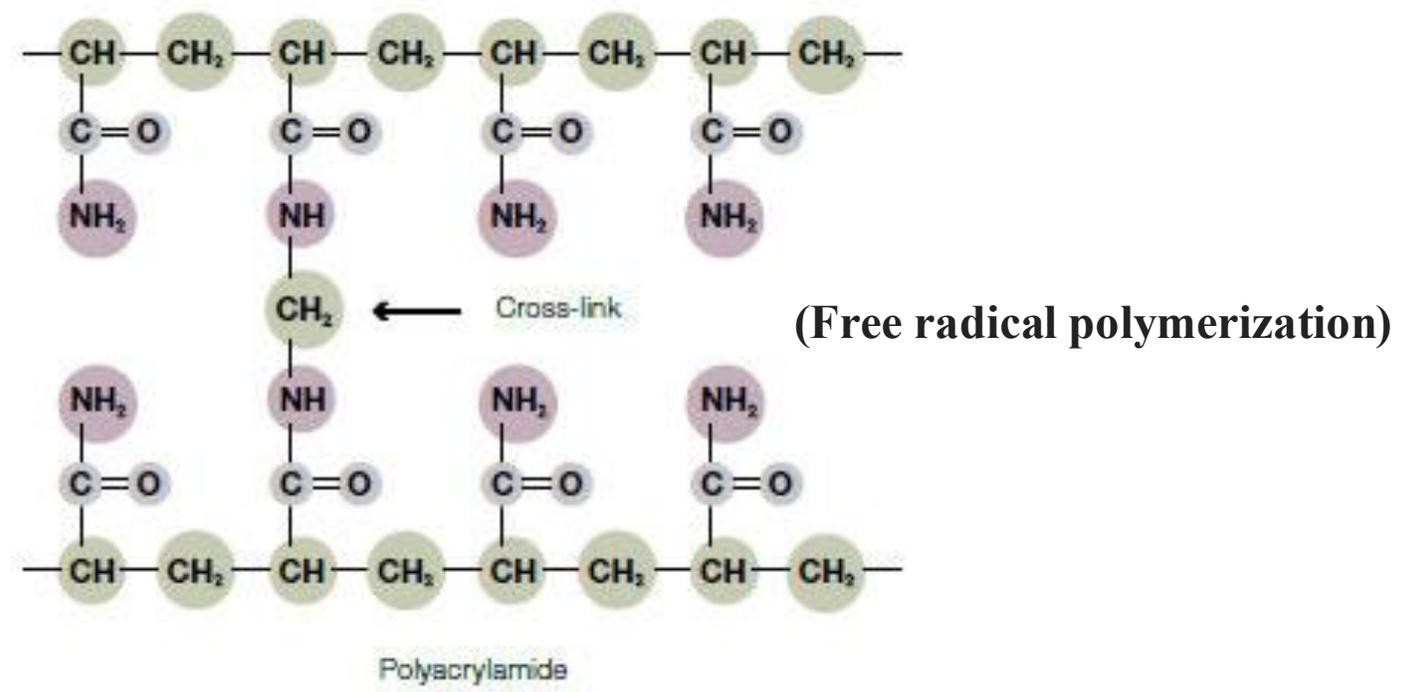
2. Polyacrylamide Gel (Acrylamide stock):

- The polyacrylamide gel is formed by co-polymerization of acrylamide and **cross-linking** by N, N'-methylene-bis-acrylamide “bis-acrylamide”.
- Polyacrylamide = acrylamide + bis-acrylamide.

- **To polymerize the gel a system:**
- 1- Ammonium PerSulfate (APS) [initiator].
- 2-TetraMethylene Ethylene Diamine (TEMED) [catalyst].



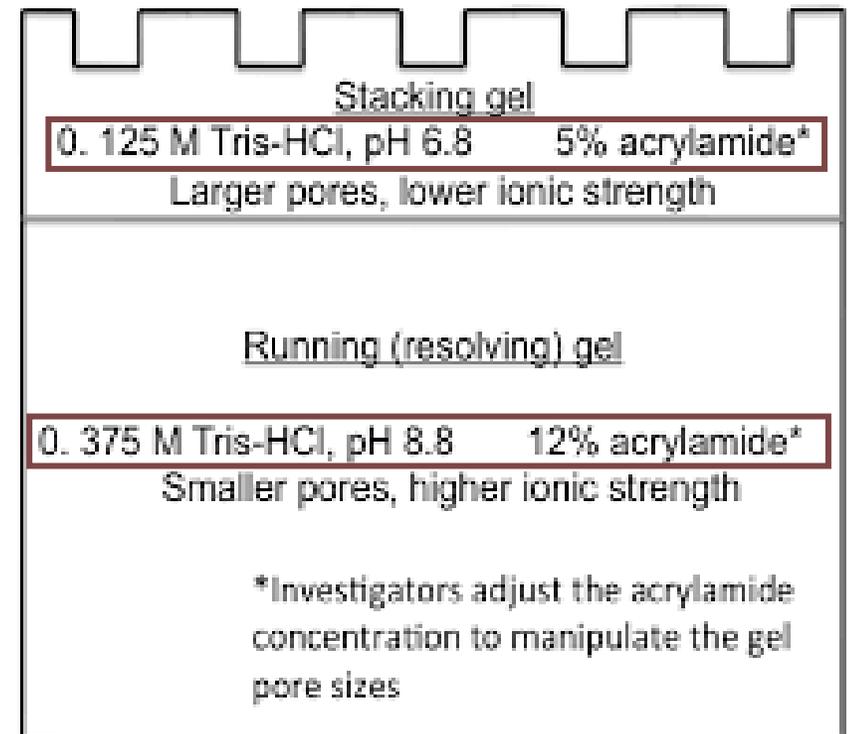
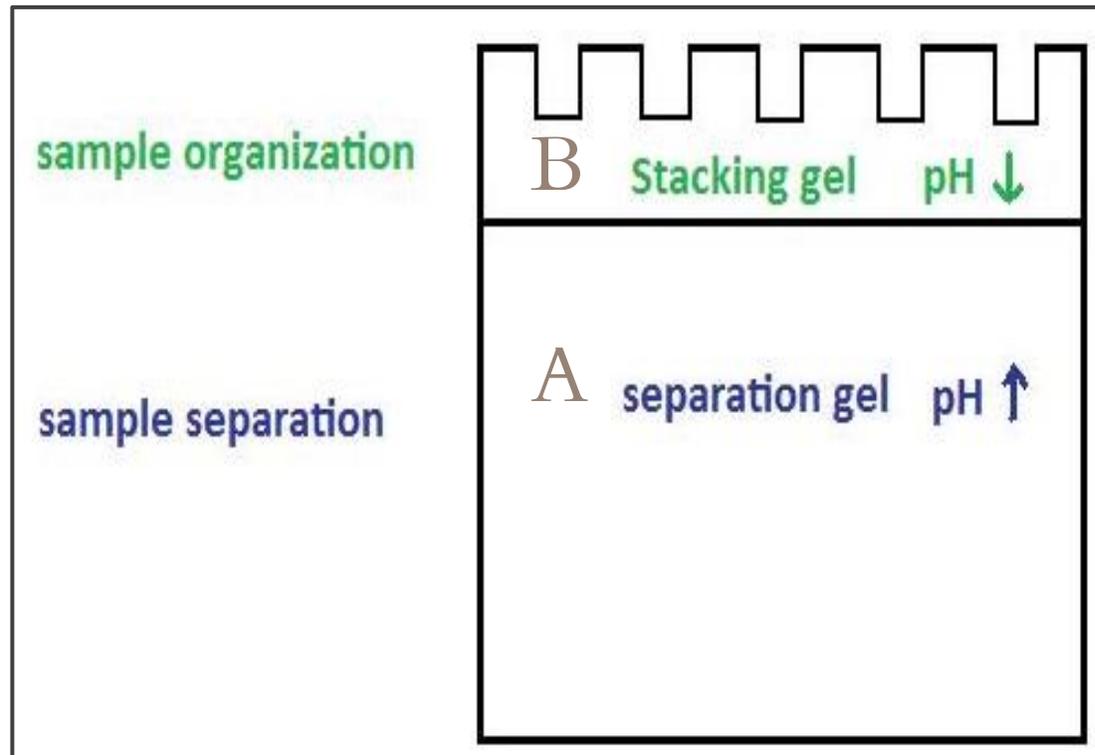
ammonium persulfate (APS) **↓** Tetramethylethylenediamine (TEMED)



2. Polyacrylamide Gel:

- It is composed of **two** gels layers:
 - A. Separation gel:
 - **HIGH** concentration of the acrylamide stock and pH value.
 - B. Stacking gel:
 - **LOW** concentration of the acrylamide stock and pH value.

The gel's two layers:



Discontinuous polyacrylamide gel

A. Separation Gel Contents:

- For 10 ml of separation gel:

Acrylamide%	6%	8%	10%	12%	15%
H ₂ O	5.2 ml	4.6 ml	3.8 ml	3.2 ml	2.2 ml
Acrylamide/Bis-acrylamide (30% / 0.8% w/v)	2 ml	2.6 ml	3.4 ml	4 ml	5 ml
1.5 M Tris (pH=8.8)	2.6 ml				
10% w/v SDS	0.1 ml				
10% w/v Ammonium persulfate (APS)	100 µl				
TEMED	10 µl				

*Note: APS and TEMED must be added right before each use.

- The acrylamide percentage in SDS PAGE gel depends on the **size of the target protein in the sample.**

Acrylamide %	M.W. Range
7%	50 kDa - 500 kDa
10%	20 kDa - 300 kDa
12%	10 kDa - 200 kDa
15%	3 kDa - 100 kDa

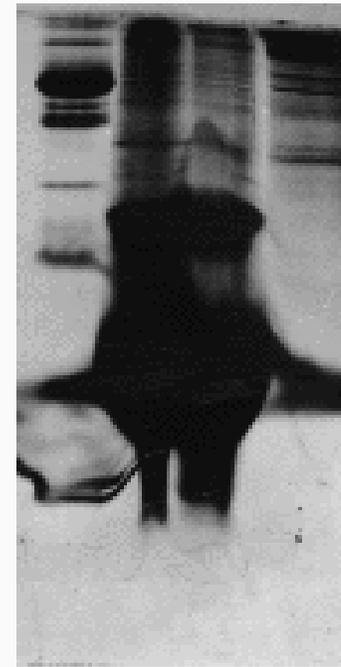
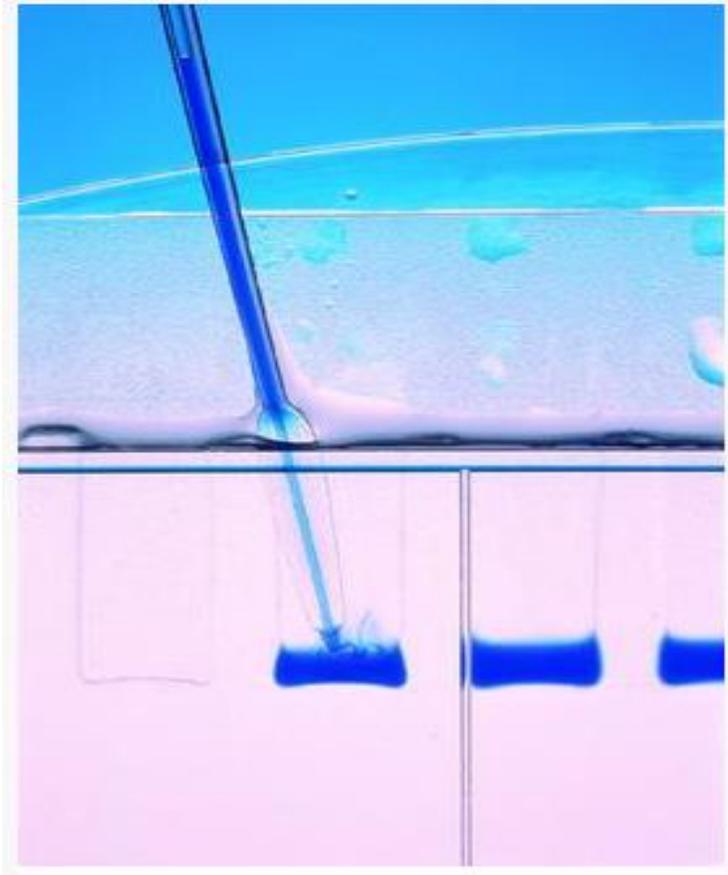
B. Stacking Gel Contents:

- For 5 ml of stacking gel:

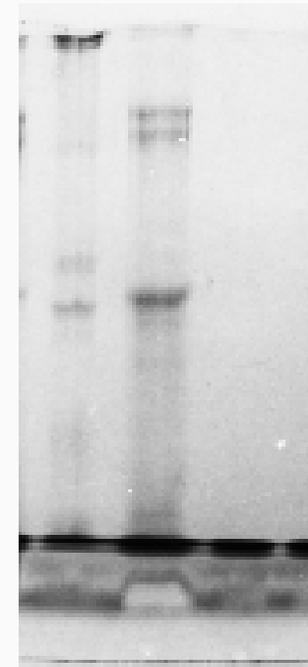
	Volume
H ₂ O	2.975 ml
Acrylamide/Bis-acrylamide (30% / 0.8% w/v)	0.67 ml
0.5 M Tris-HCl (pH=6.8)	1.25 ml
10% w/v SDS	0.05 ml
10% w/v Ammonium persulfate (APS)	0.05 ml
TEMED	0.005 ml

The stacking gel has a lower concentration of acrylamide

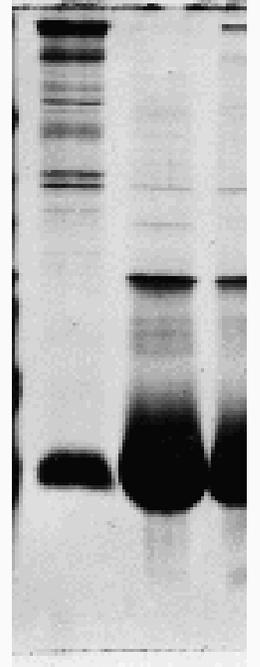
3. Sample Loading:



too much



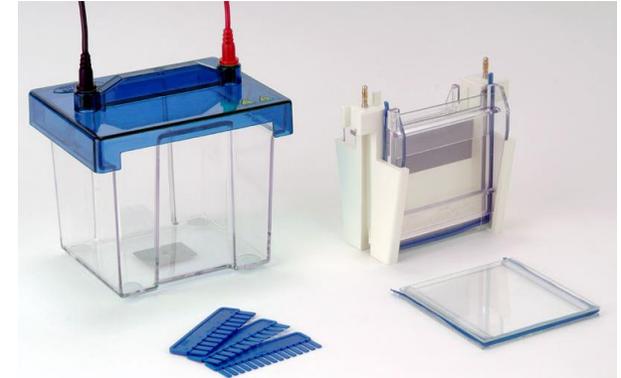
too little



good

4. Running the gel using Running buffer (pH 8.3): →

- It helps to deliver the electric current through the gel.
- It contains:
 - Tris-HCl.
 - Glycine.
 - SDS.



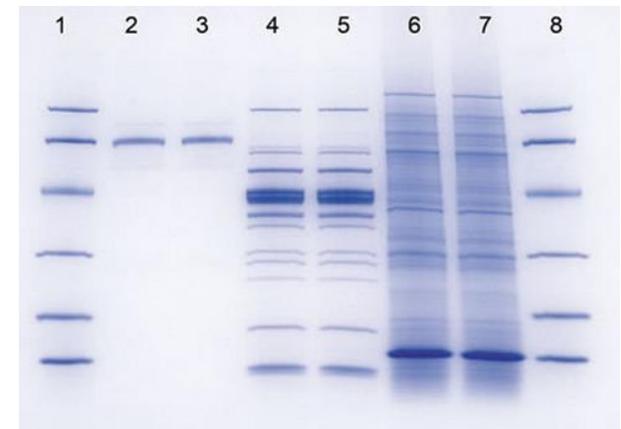
5. Stain the gel using staining buffer : →

- It contains:
 - Glacial acetic acid
 - Methanol
 - Coomassie brilliant blue 250-R.

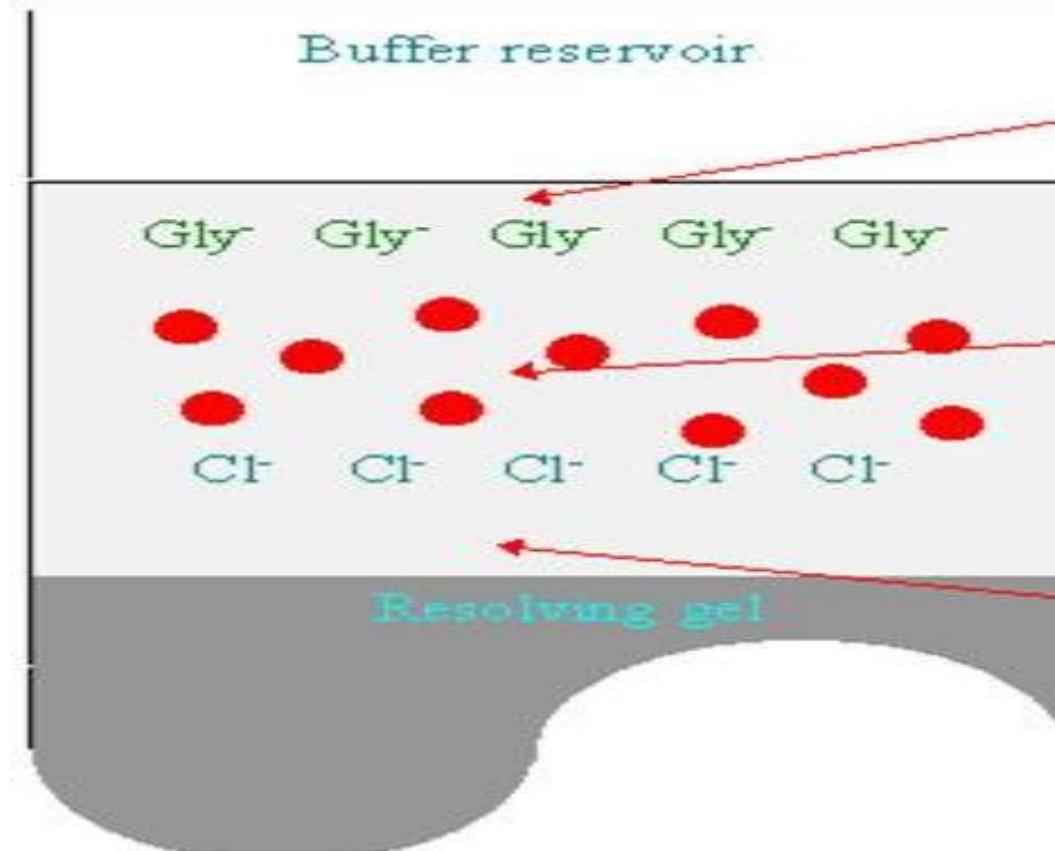


6. De-stain the gel using De-staining buffer: →

- It contains:
 - Glacial acetic acid
 - Methanol



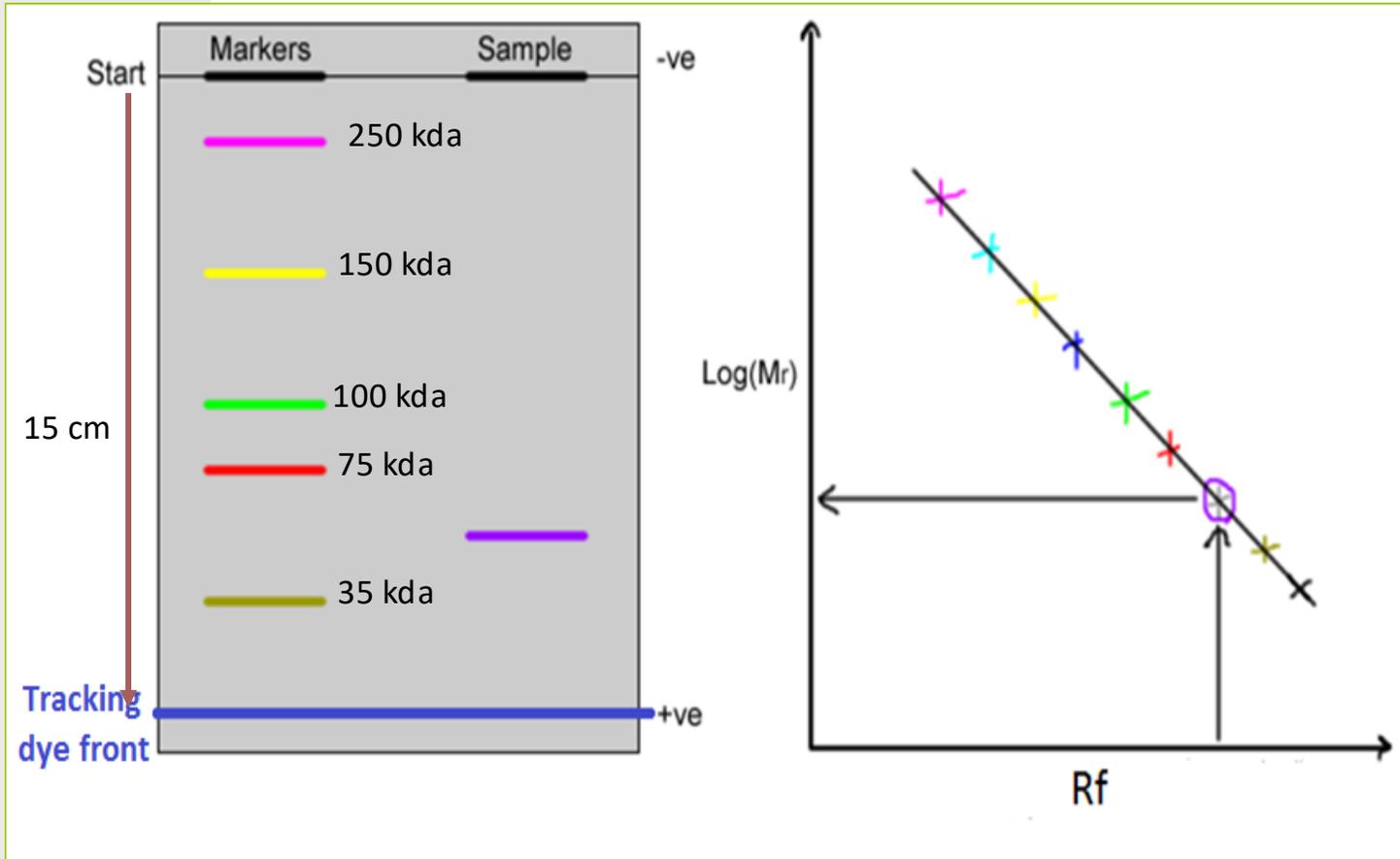
Proteins are sandwiched between the Cl^- ions and the glycine molecules.



Data Analysis



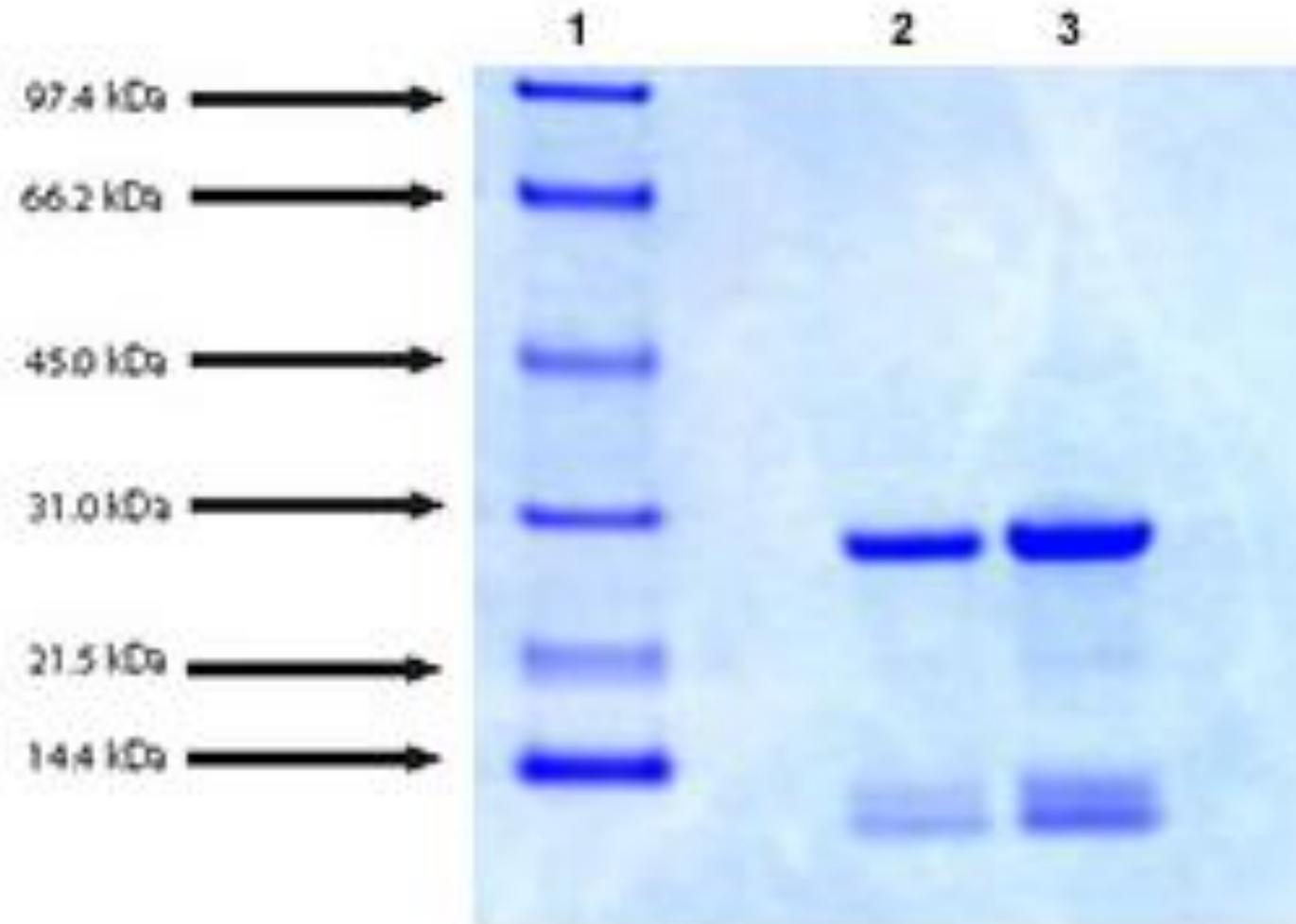
For Molecular Weight Determination:

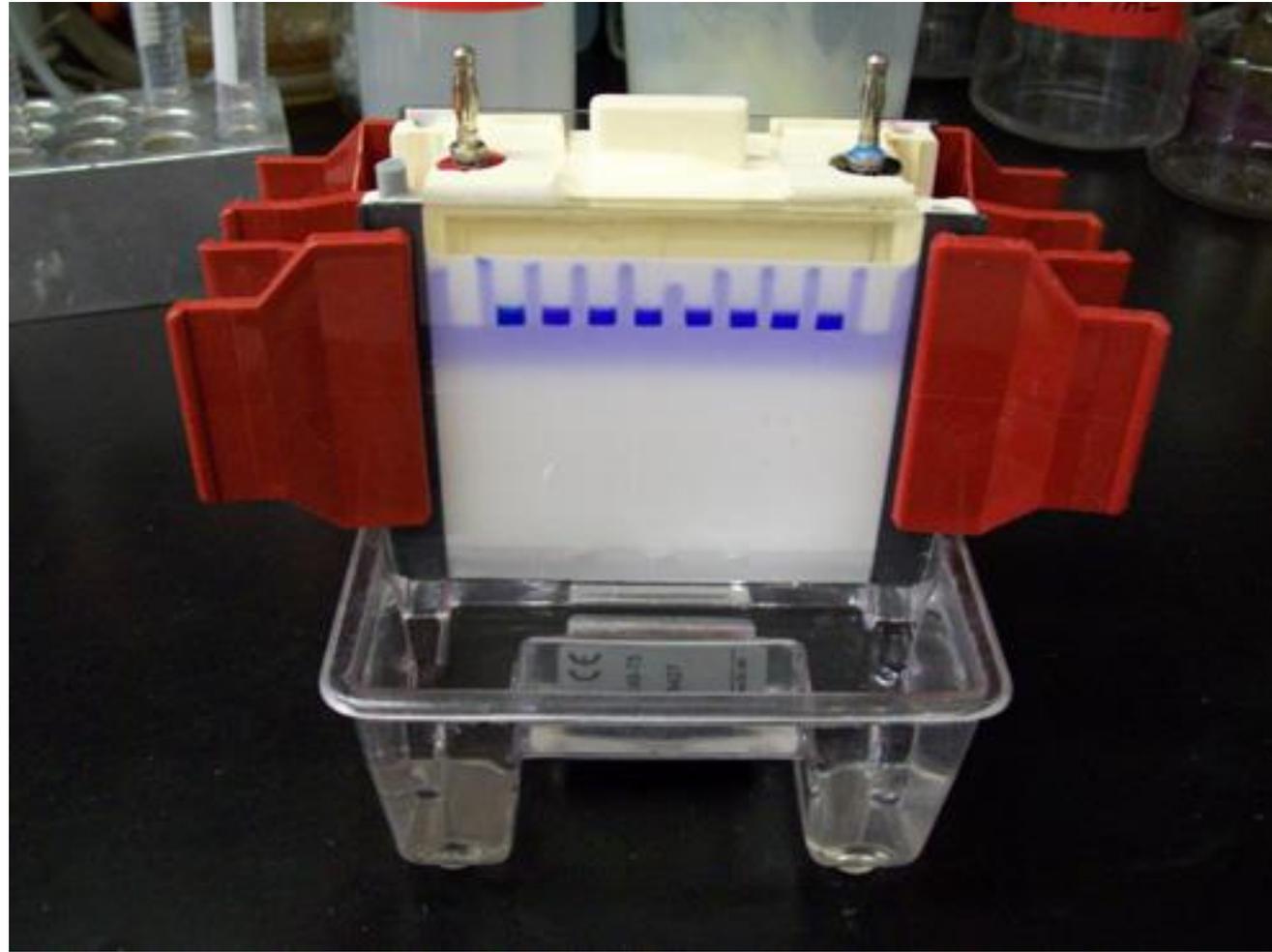


$$- R_f = \frac{\text{Distance of migration of sample}}{\text{Distance moved by tracking dye}}$$

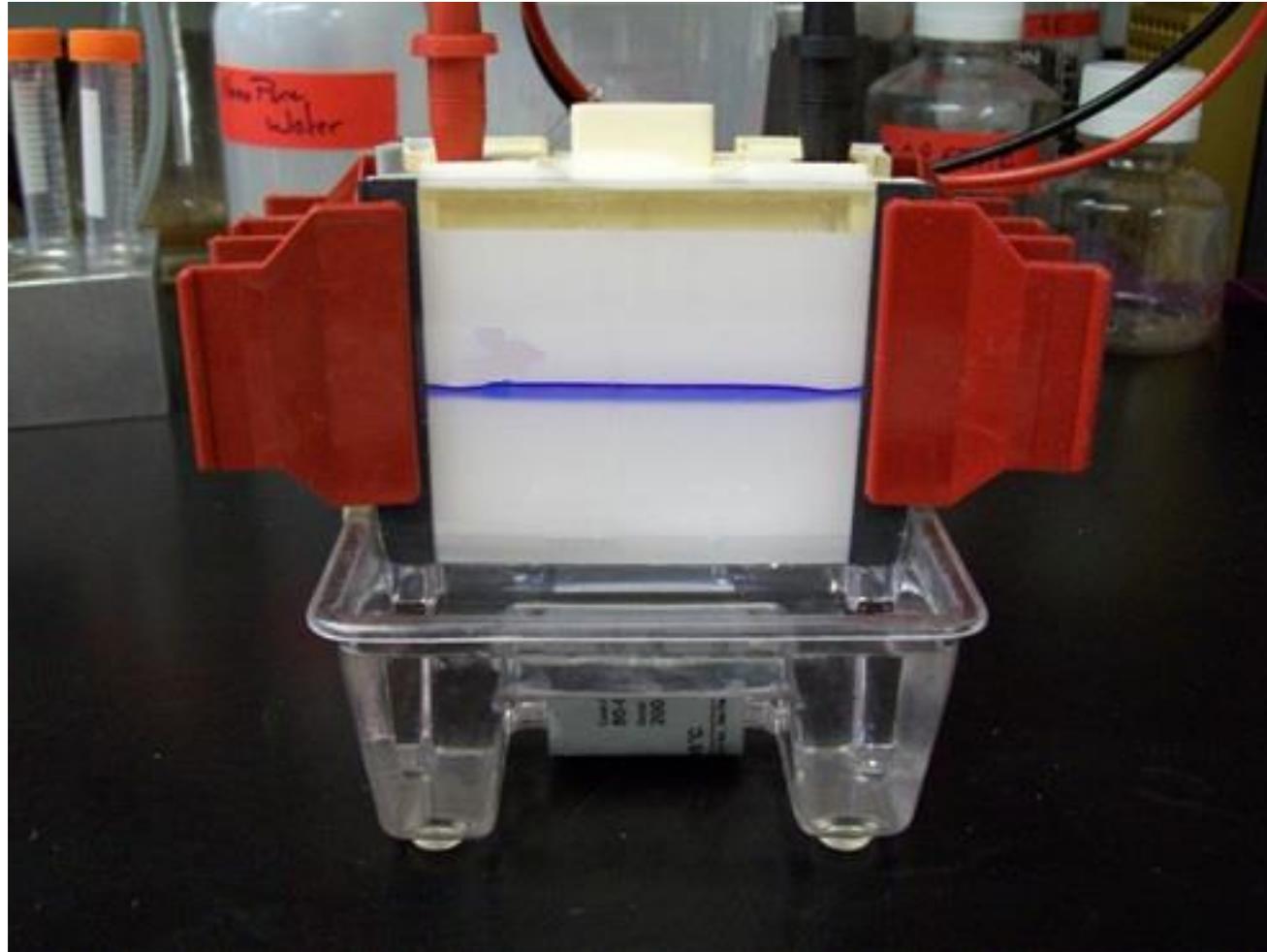
	Dye front (cm)	Distance (cm)	R_f	M.W (Da)	Log M.W (Da)
std ₁	15 cm	1.5	0.1	250000	5.4
std ₂		4	0.27	150000	5.18
std ₃		5.5	0.37	100000	5
std ₄		6.4	0.43	75000	4.88
std ₅		12.3	0.82	35000	4.54
Un-known M.W sample		9.3	0.623	?	?

For Molecular Weight Determination cont':

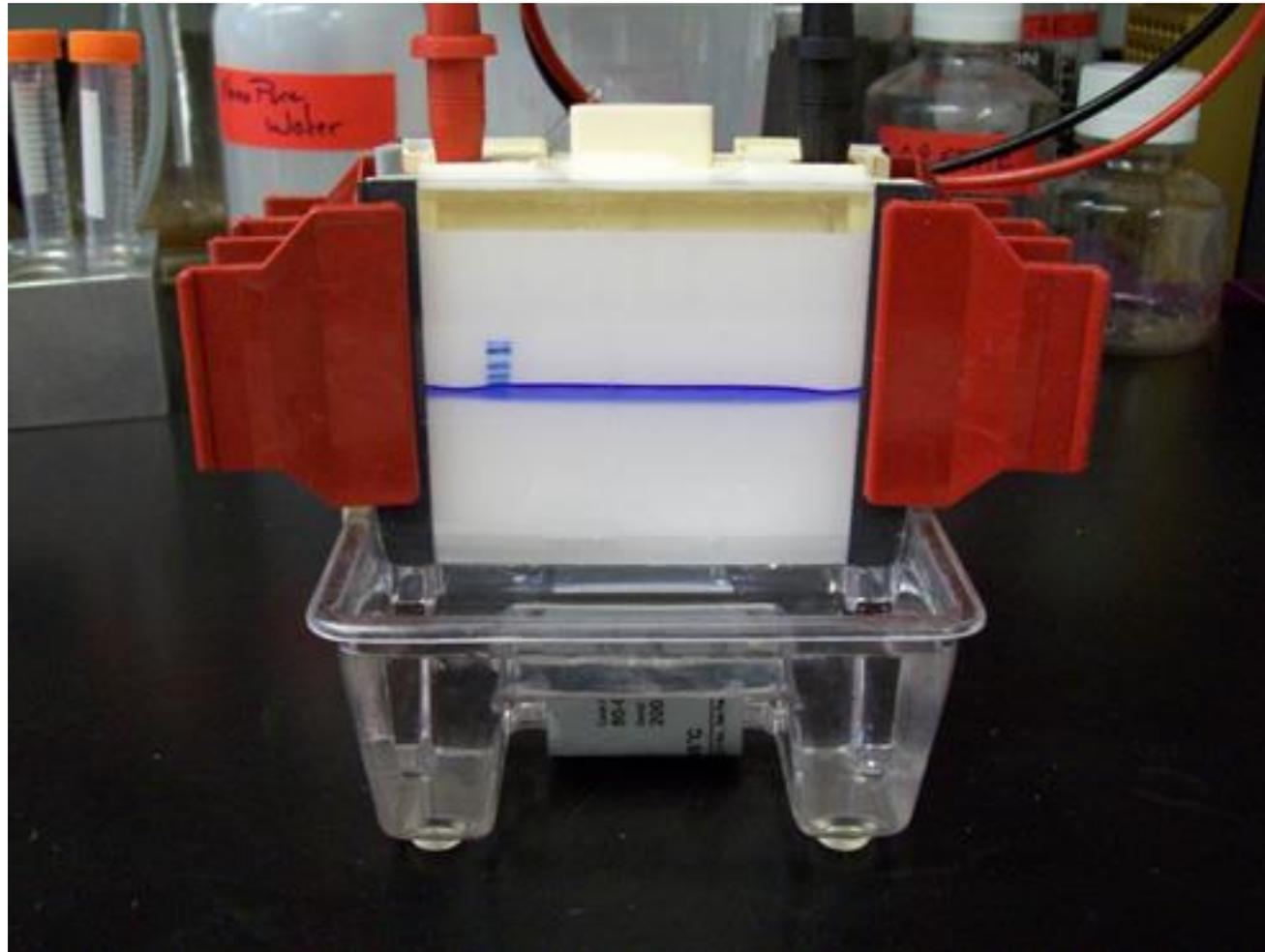




At beginning

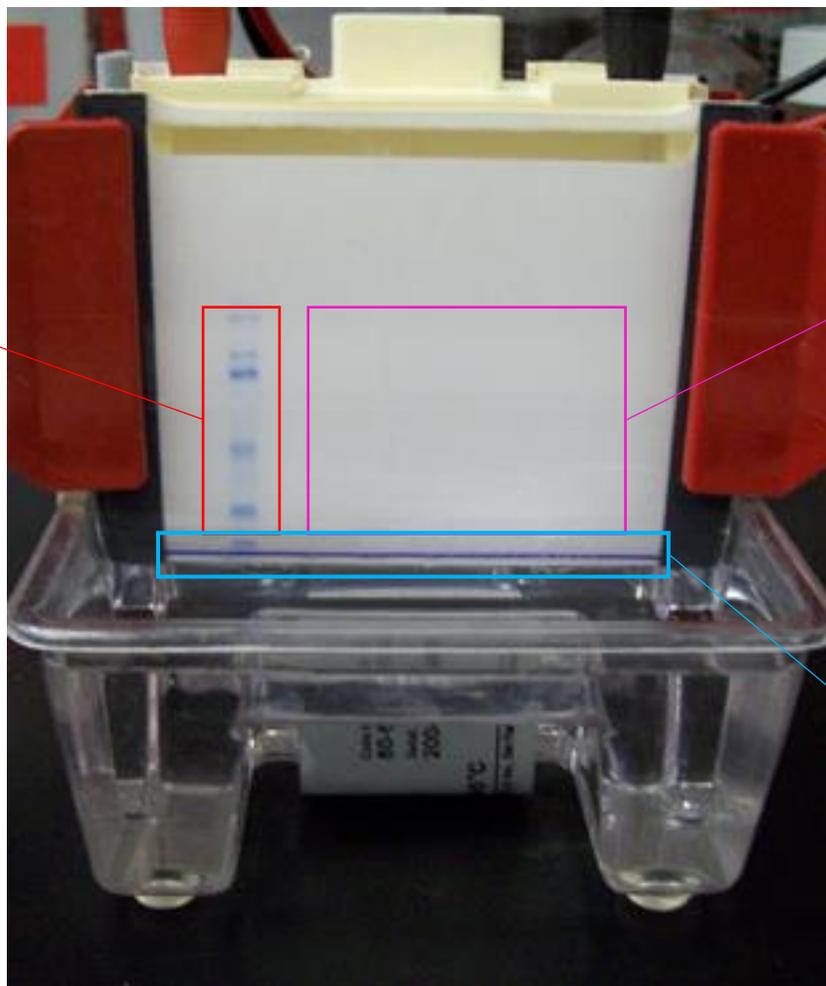


After a while



After more mins.

Marker
Ladder)



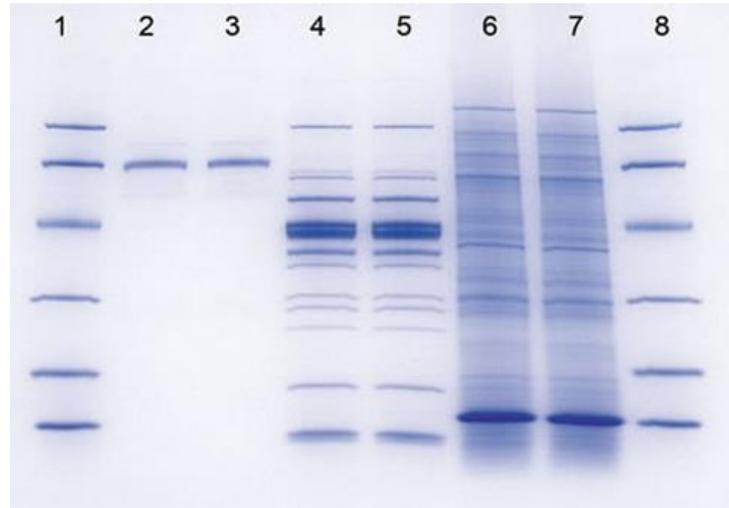
Separated Sample

Dye front

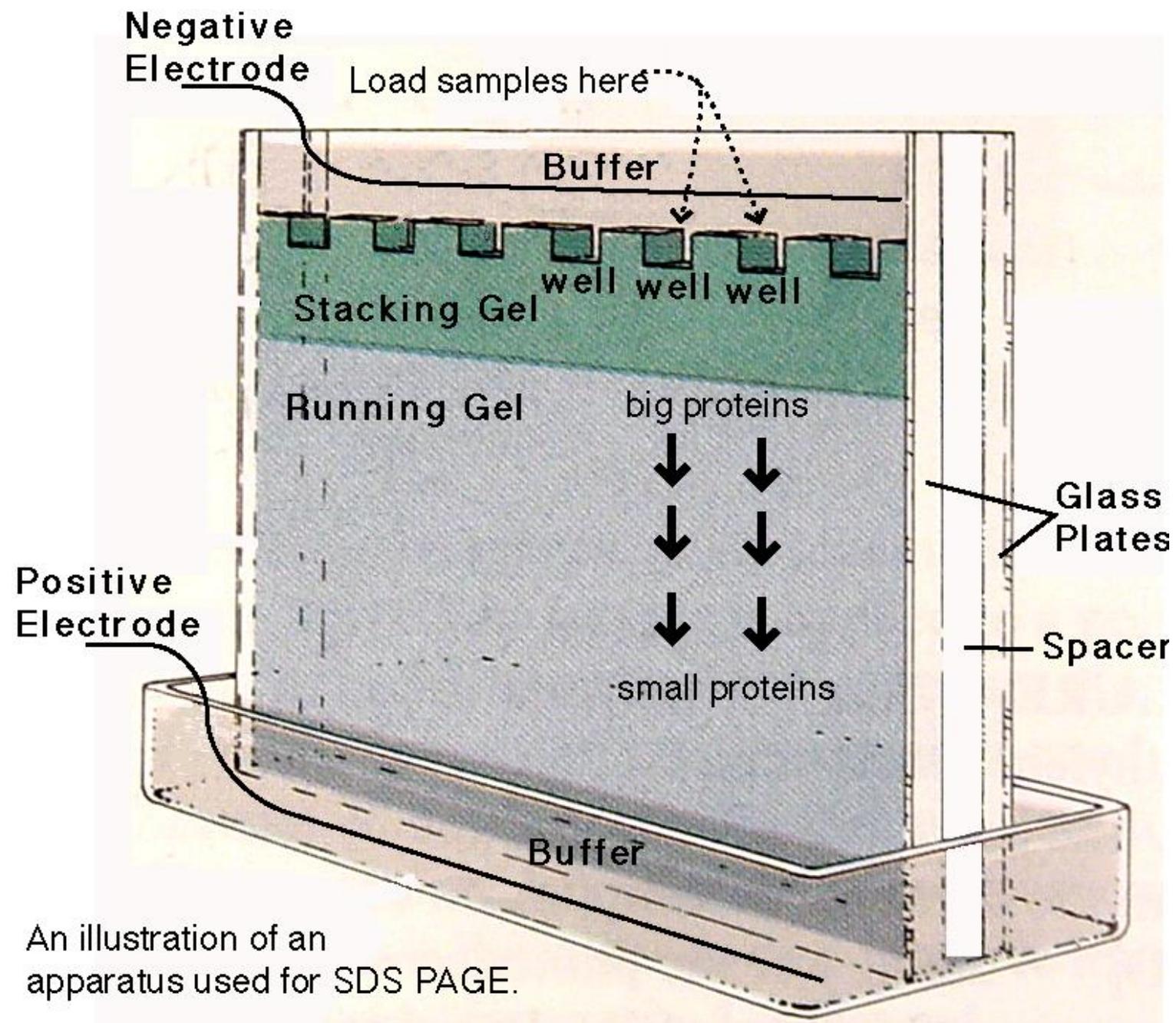
At the end



Staining



De Staining



An illustration of an apparatus used for SDS PAGE.

Supporting Materials:

- SDS PAGE principles - simple animated tutorial:

<https://www.youtube.com/watch?v=3CrzY7jb9fQ>

https://www.youtube.com/watch?v=i_6y6Z5UvwE