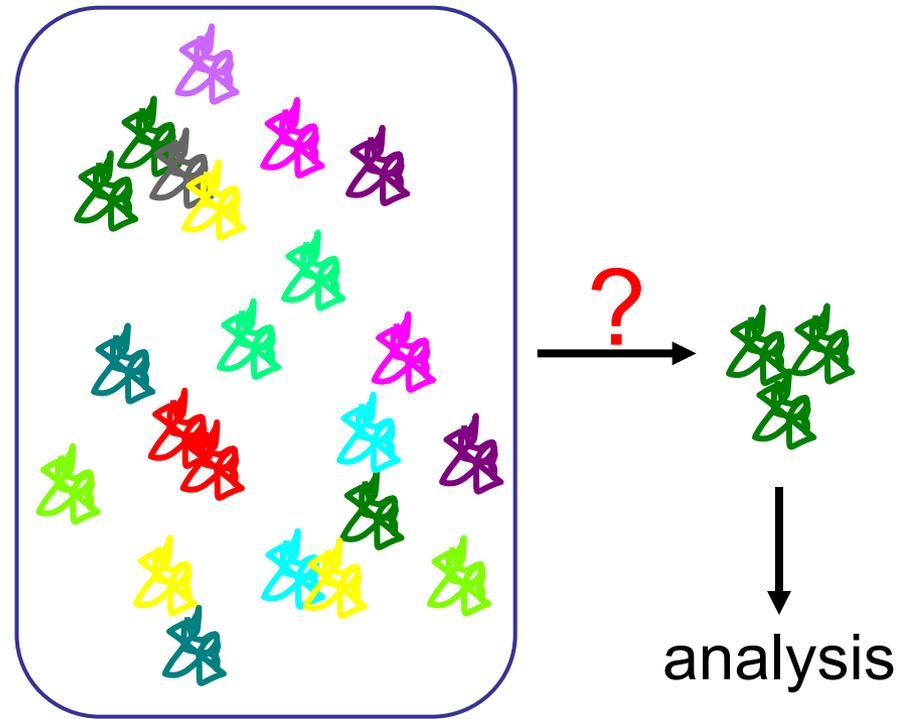


# Protein Purification

In order to carry out detailed studies of a protein (structure, enzyme activity, amino acid sequence, etc) \_\_\_\_\_

How is a single protein isolated \_\_\_\_\_?



# Protein Purification

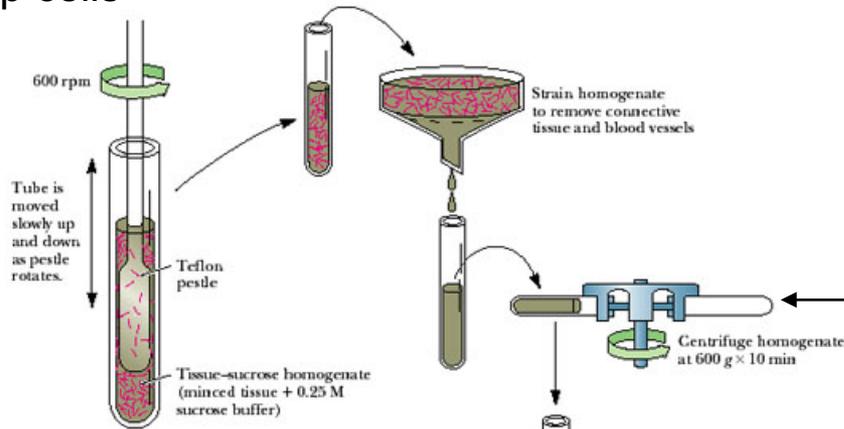
1. Extract \_\_\_\_\_
2. Separate extract contents by \_\_\_\_\_  
\_\_\_\_\_
3. Separate protein extract by \_\_\_\_\_
  - \_\_\_\_\_
  - \_\_\_\_\_
  - \_\_\_\_\_
4. Analyze purity of isolated protein \_\_\_\_\_

# Protein Purification

## Making protein extract

### 1. Grind tissue to break up cells

mortar & pestle



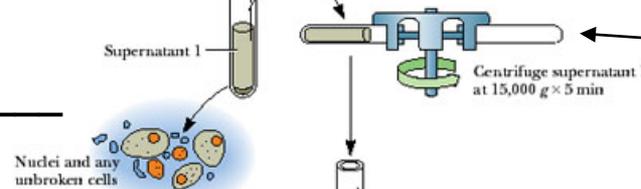
### 2. Centrifuge to

\_\_\_\_\_

\_\_\_\_\_

Low speed centrifugation to remove large contents, nuclei, cell wall debris (plants), etc.

Separate liquid sample into \_\_\_\_\_ and \_\_\_\_\_.



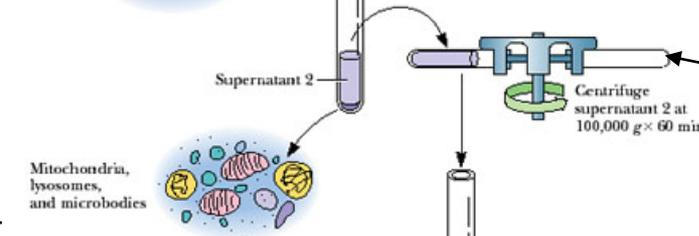
Higher speed centrifugation

\_\_\_\_\_

\_\_\_\_\_

**Supernatant:** liquid above the \_\_\_\_\_

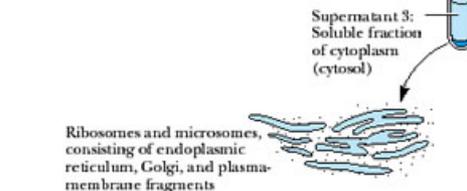
\_\_\_\_\_



Very high speed centrifugation to remove \_\_\_\_\_

\_\_\_\_\_

\_\_\_\_\_



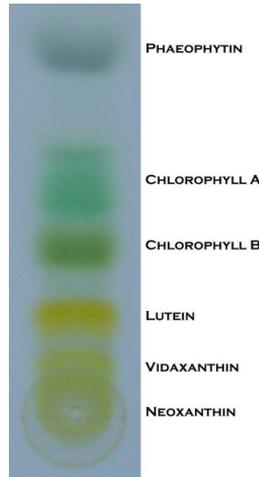
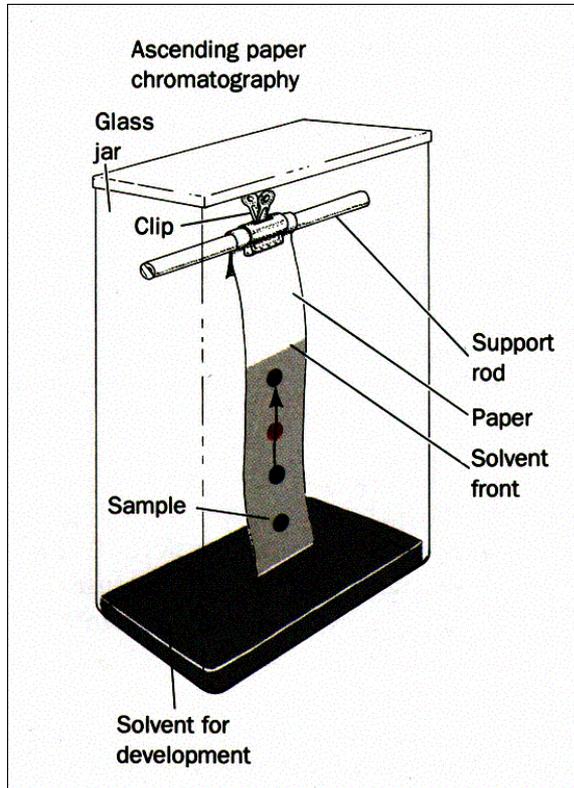
# Protein Purification

## Chromatography

*chroma* - "color"  
*graphien* - "to write"

**Theory:** separation of \_\_\_\_\_ within a sample based on their \_\_\_\_\_ as they pass through a medium.

First used to separate plant pigments



Phases:

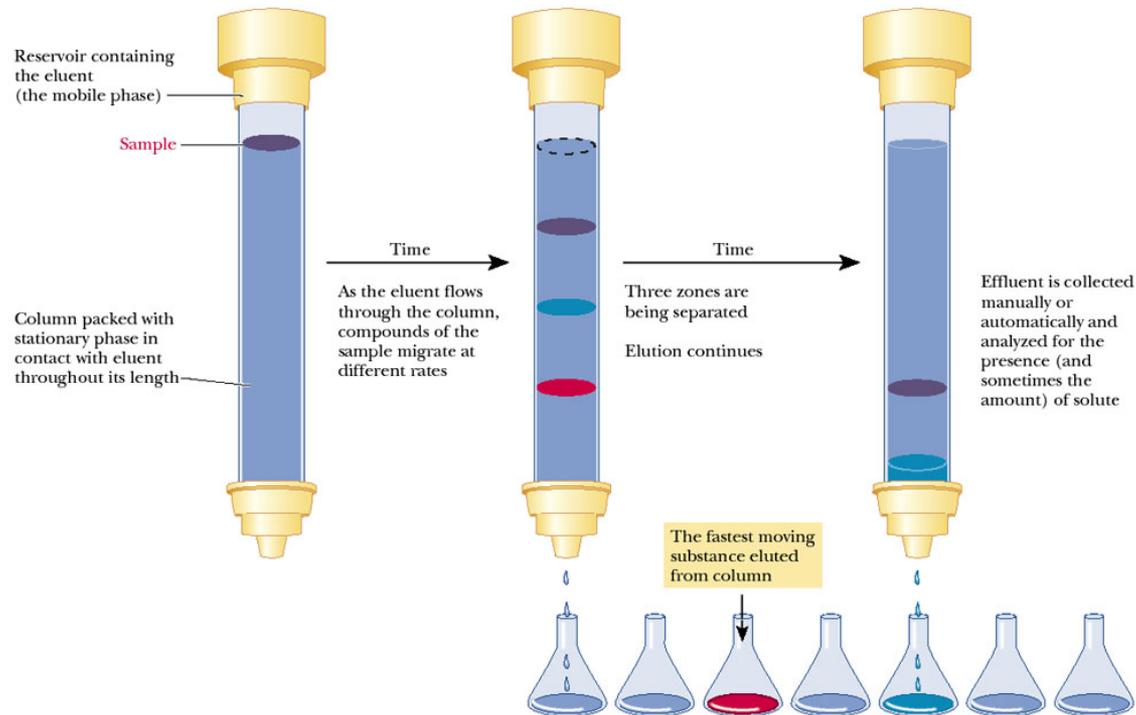
**stationary phase:** a solid material over which \_\_\_\_\_. Imparts separation of compounds.

**mobile phase:** a liquid solvent that carries \_\_\_\_\_

# Protein Purification

## Column Chromatography

1. Pack column with \_\_\_\_\_ mobile phase
2. Add sample to \_\_\_\_\_
3. Use pump or gravity to pass mobile phase over stationary phase thus \_\_\_\_\_
4. Collect samples as they \_\_\_\_\_



# Protein Purification

## Column Chromatography

### Types of Stationary Phases

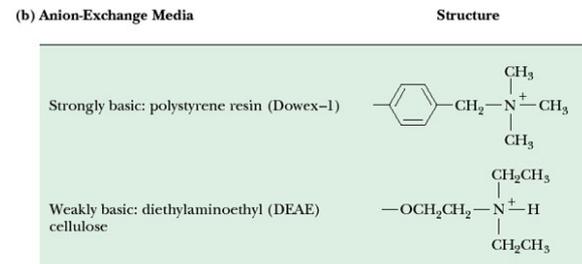
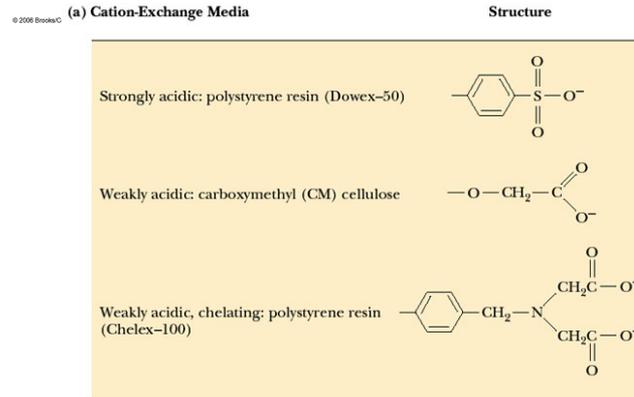
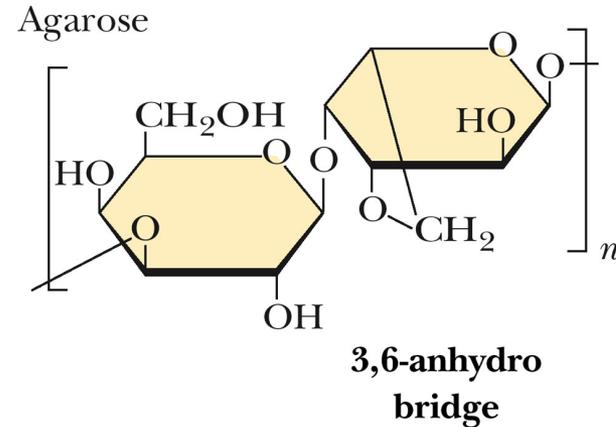
Size-exclusion/gel-filtration:

- dextran (Sephadex) - \_\_\_\_\_  
 \_\_\_\_\_ from  
 certain bacteria.

- agarose (Sepharose) - a  
 polysaccharide of \_\_\_\_\_  
 \_\_\_\_\_.

Ion-exchange:

- \_\_\_\_\_-exchange  
 - \_\_\_\_\_-exchange

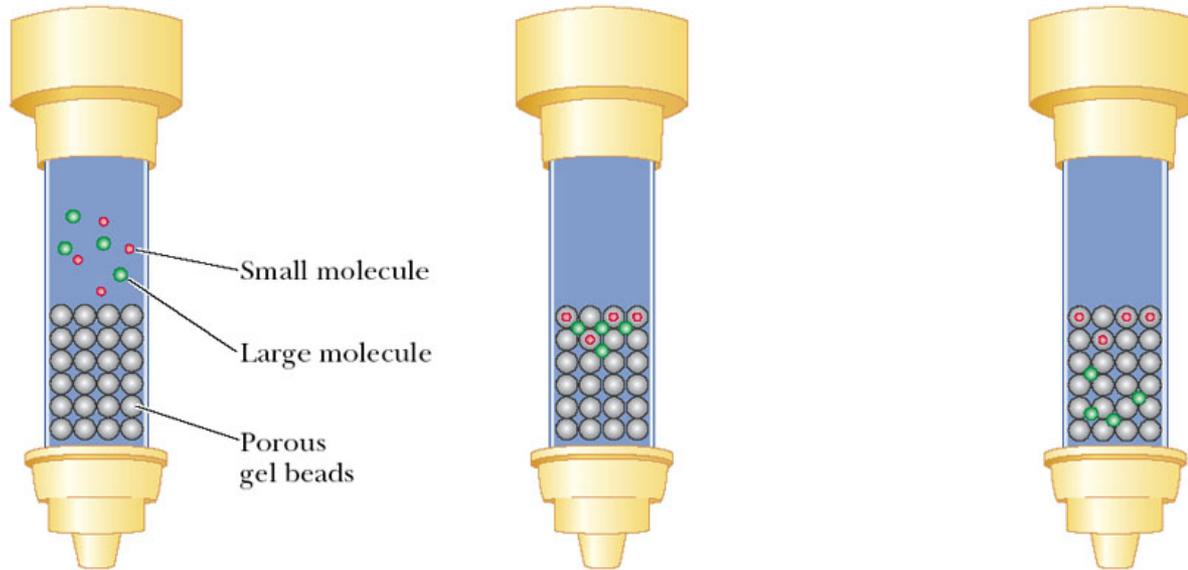


# Protein Purification

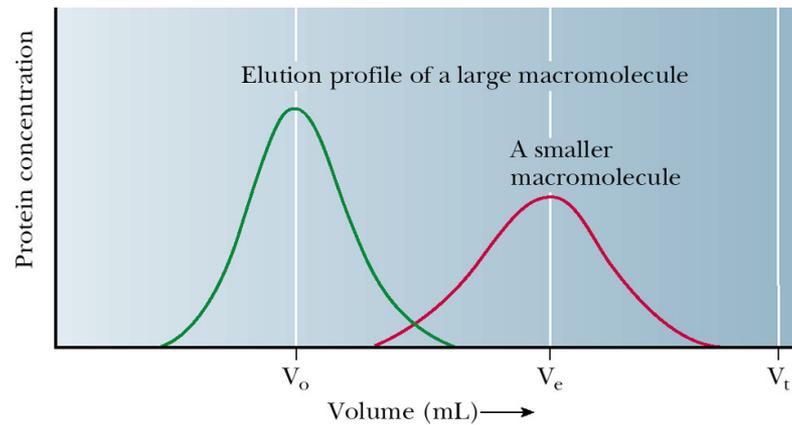
## Gel-filtration Chromatography

Principle: to separate \_\_\_\_\_

(a)



© 2006 Brooks/Cole - Thomson

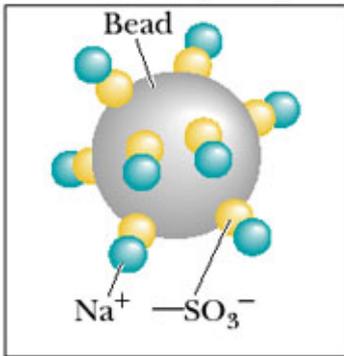


# Protein Purification

## Ion-exchange Chromatography

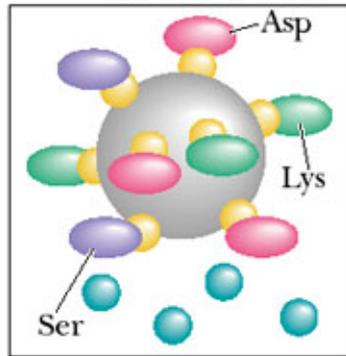
Principle: a charged ion bound to the column is displaced by \_\_\_\_\_ . The protein is eluted from the column using \_\_\_\_\_ .

Cation exchange bead before adding sample



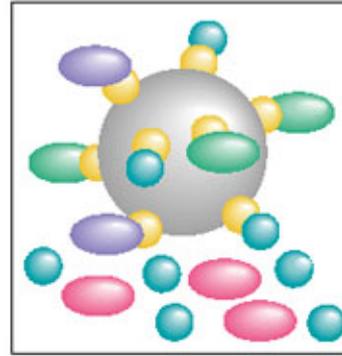
(a)

Add mixture of Asp, Ser, Lys



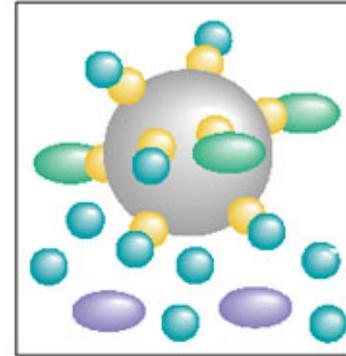
(b)

Add Na<sup>+</sup> (NaCl)



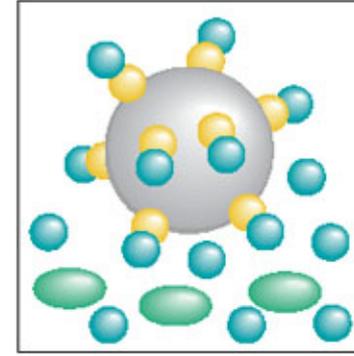
(c) Asp, the least positively charged amino acid, is eluted first

Increase [Na<sup>+</sup>]



(d) Serine is eluted next

Increase [Na<sup>+</sup>]

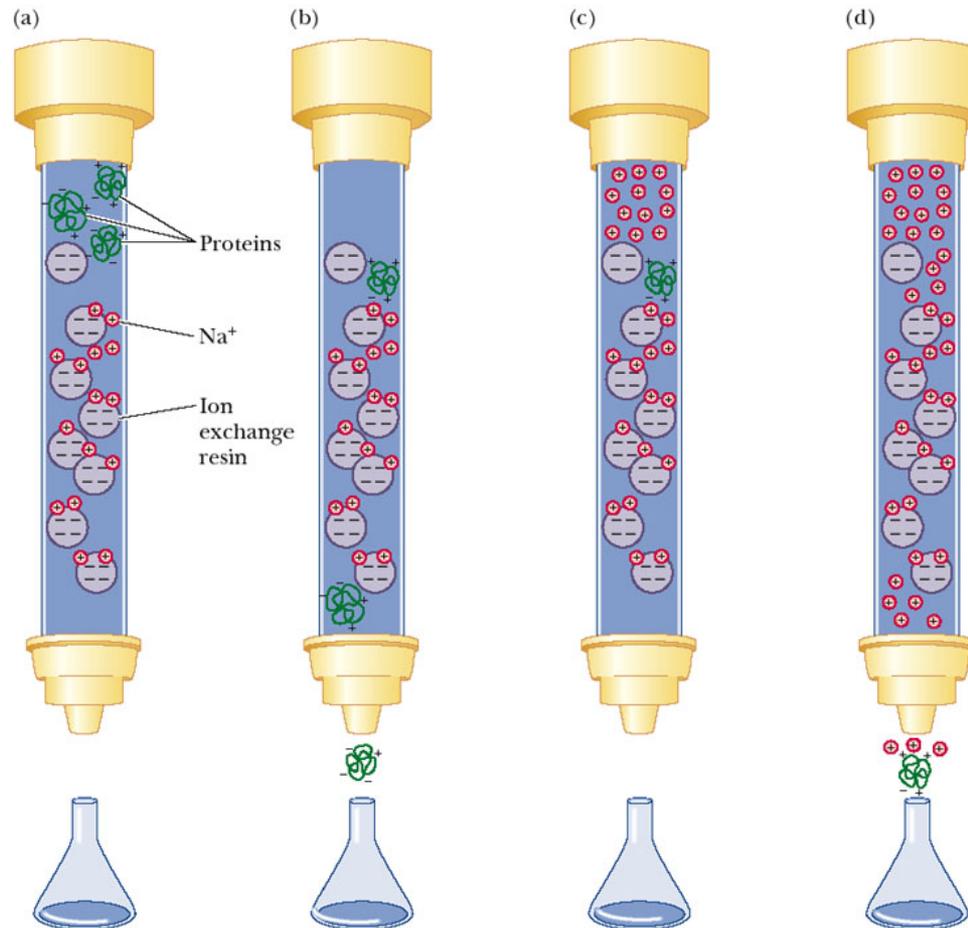


(e) Lysine, the most positively charged amino acid, is eluted last

# Protein Purification

## Ion-exchange Chromatography

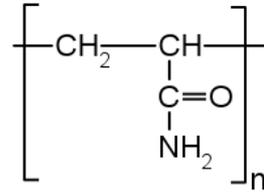
Principle: a charged ion bound to the column is displaced by the charged side groups on a protein. The protein is eluted from the column using increasing amounts of the ion.



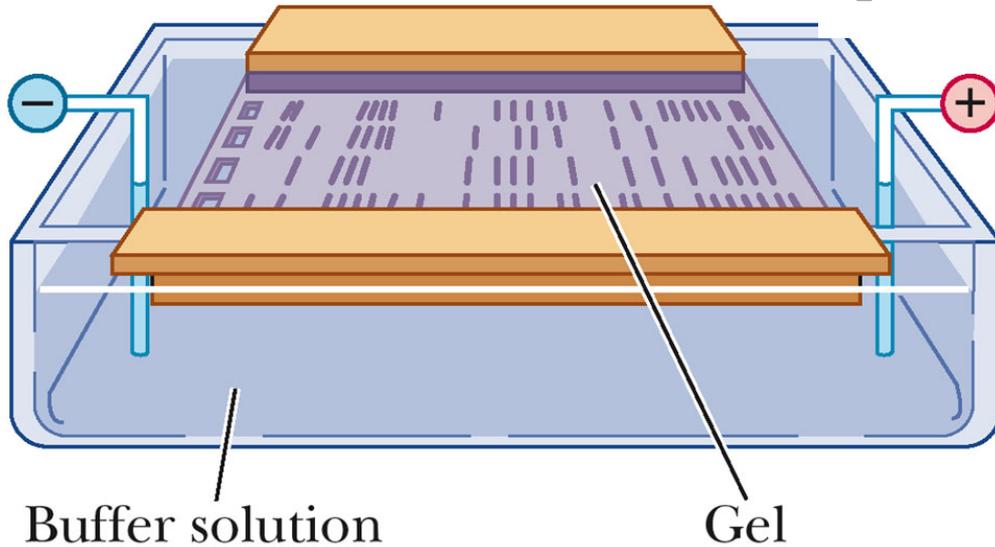
# Protein Purification

## Electrophoresis

**Principle:** protein samples are separated through \_\_\_\_\_. Proteins are “pulled” through the gel by an electric field.

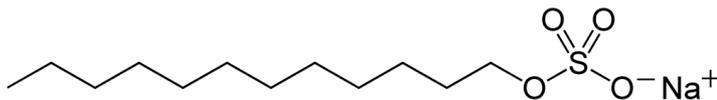


**Gel:** polymer of \_\_\_\_\_



**Sodium dodecyl sulfate (SDS):** a negatively charged \_\_\_\_\_ proteins and give a uniform “-” charge.

In the presence of \_\_\_\_\_ will travel towards anode, “+”



**SDS-PAGE:** \_\_\_\_\_

# Protein Purification

## Electrophoresis

### Cathode:

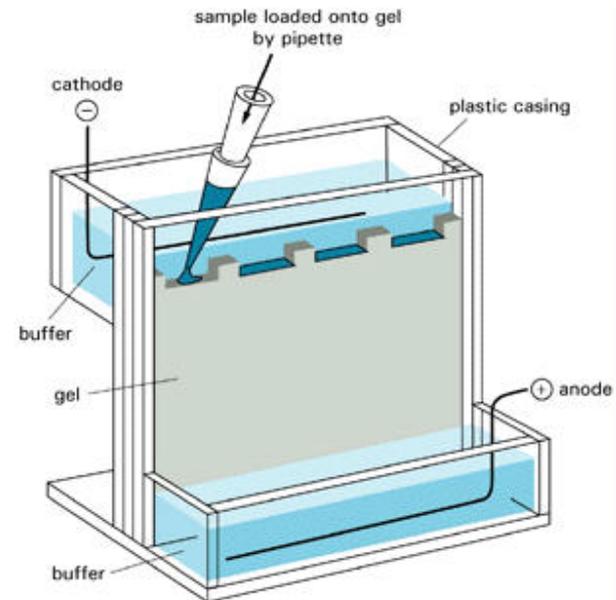
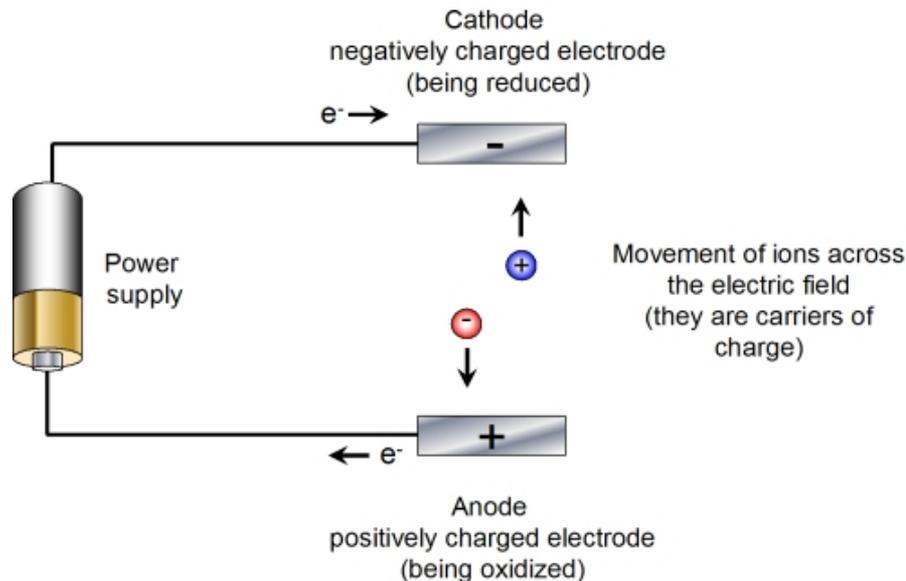
- The negatively charged electrode by which electrons enter an electrical device.
- The positively charged electrode of an electrical device that supplies current.

The cathode is negative in a device that consumes or uses power, and the cathode is positive in a device that provides power

### Anode:

- The positively charged electrode by which electrons leave an electrical device.
- The negatively charged electrode of a device that supplies current.

the anode is positive in a device that consumes power, and the anode is negative in a device that provides power

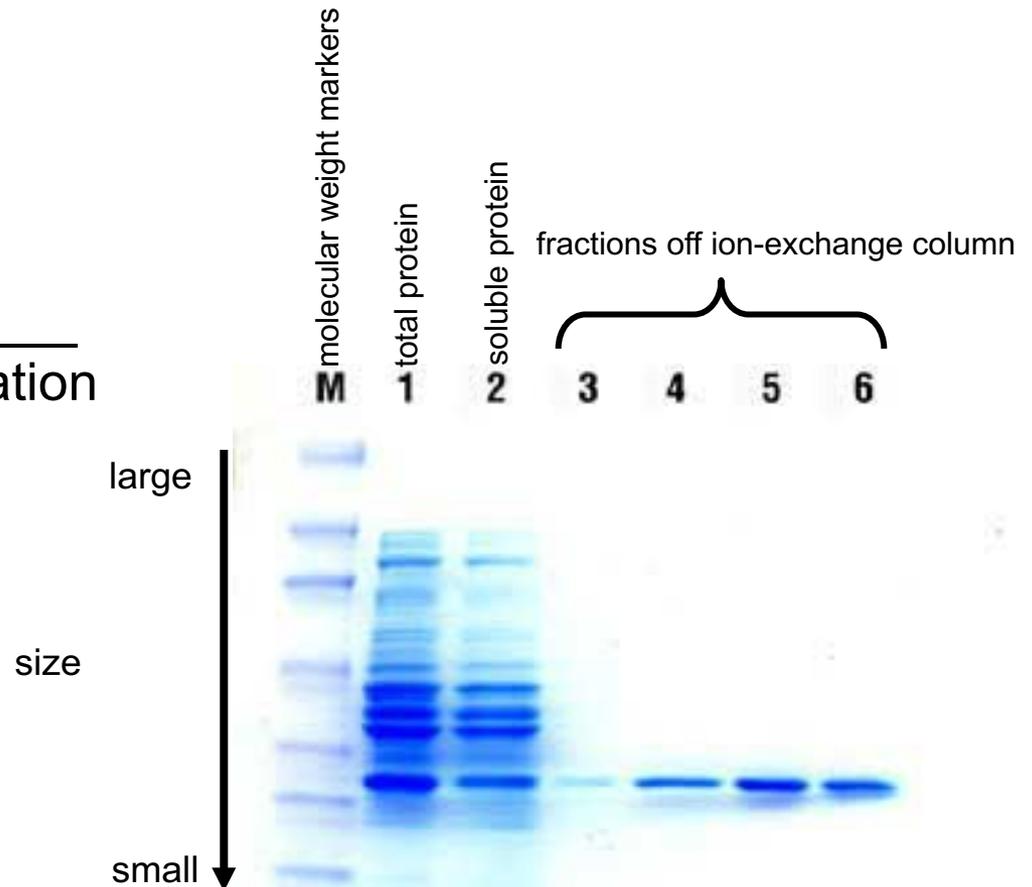


# Protein Purification

## Electrophoresis

**SDS-PAGE:** separates proteins within a sample \_\_\_\_\_.  
**large proteins** migrate \_\_\_\_\_  
**small proteins** migrate \_\_\_\_\_

Stain gel with \_\_\_\_\_  
\_\_\_\_\_ for visualization



[SDS-PAGE](http://www.youtube.com/watch?v=pjEHJIPrtIU)

(<http://www.youtube.com/watch?v=pjEHJIPrtIU>)



# Protein Purification

Once a protein has been purified how is the primary structure (sequence) determined?

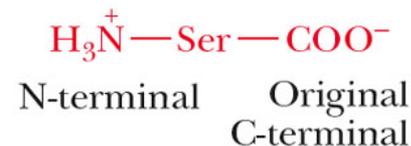
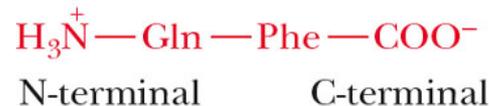
**Chymotrypsin:** enzyme that \_\_\_\_\_

---

Original protein



Chymotrypsin  
digestion



# Protein Purification

Once a protein has been purified how is the primary structure (sequence) determined?

**Cyanogen bromide:** \_\_\_\_\_

1. \_\_\_\_\_ by several different methods, individually
2. Produces \_\_\_\_\_
3. Sequence peptides using \_\_\_\_\_
4. Arrange peptides to \_\_\_\_\_

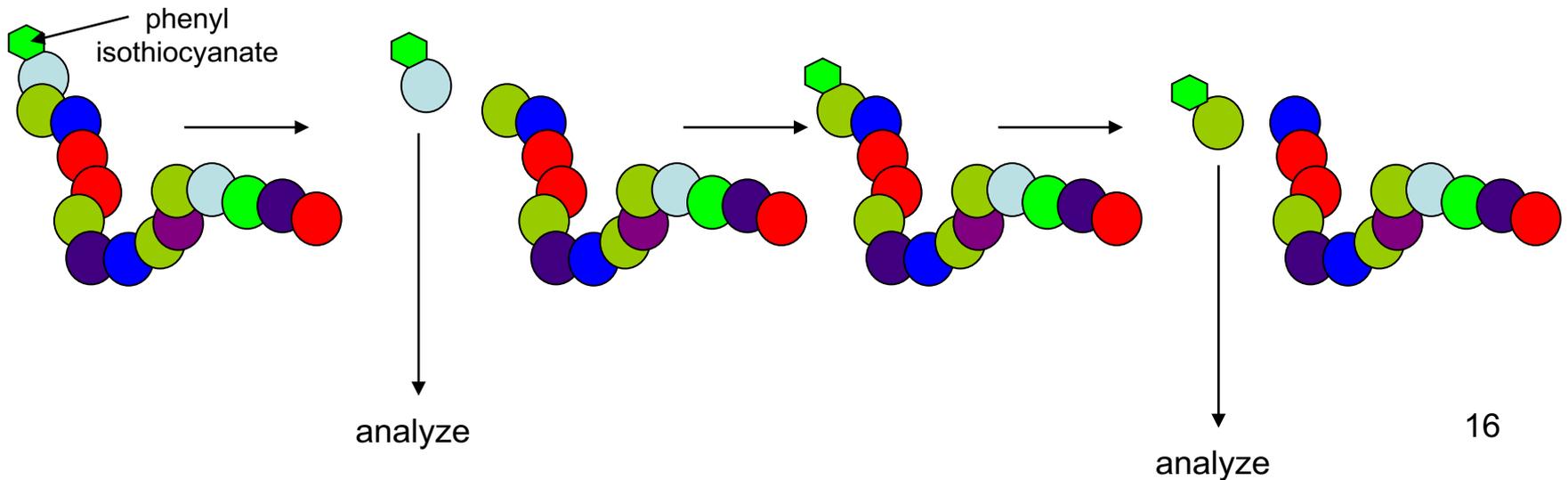
Chymotrypsin	H <sub>3</sub> N <sup>+</sup> —Leu—Asn—Asp—Phe
Cyanogen bromide	H <sub>3</sub> N <sup>+</sup> —Leu—Asn—Asp—Phe—His—Met
Chymotrypsin	His—Met—Thr—Met—Ala—Trp
Cyanogen bromide	Thr—Met
Cyanogen bromide	Ala—Trp—Val—Lys—COO <sup>-</sup>
Chymotrypsin	Val—Lys—COO <sup>-</sup>
Overall sequence	H <sub>3</sub> N <sup>+</sup> —Leu—Asn—Asp—Phe—His—Met—Thr—Met—Ala—Trp—Val—Lys—COO <sup>-</sup>

# Protein Purification

Once a protein has been purified how is the primary structure (sequence) determined?

**Edman degradation:** a method used to determine \_\_\_\_\_

1. Peptide is reacted with \_\_\_\_\_, bonds to N-terminal aa.
2. The modified aa is \_\_\_\_\_
3. Repeated until all \_\_\_\_\_.



# Protein Purification

Once a protein has been purified how is the primary structure (sequence) determined?

Example: a protein sample is split into two equal samples. One is treated with trypsin and the other with chymotrypsin. Given the peptides found, determine the protein sequence.

Trypsin peptides: \_\_\_\_\_  
and  
\_\_\_\_\_

Chymotrypsin peptides: \_\_\_\_\_  
and  
\_\_\_\_\_

# Protein Purification

Once a protein has been purified how is the primary structure (sequence) determined?

Trypsin peptides: Leu - Ser - Tyr - Ala - Ile - Arg

and

Asp - Gly - Met - Phe - Val - Lys

Possibilities for protein sequence:

1. \_\_\_\_\_

or

2. \_\_\_\_\_

# Protein Purification

Once a protein has been purified how is the primary structure (sequence) determined?

Chymotrypsin peptides: Val - Lys - Leu - Ser - Tyr  
and  
Ala - Ile - Arg  
and  
Asp - Gly - Met - Phe

1. Leu - Ser - Tyr - Ala - Ile - Arg - Asp - Gly - Met - Phe - Val - Lys

2. Asp - Gly - Met - Phe - Val - Lys - Leu - Ser - Tyr - Ala - Ile - Arg