

# Protein Purification

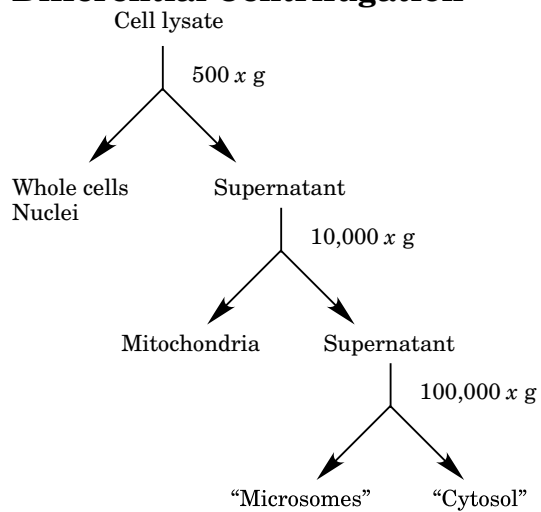
Advantages

Disadvantages

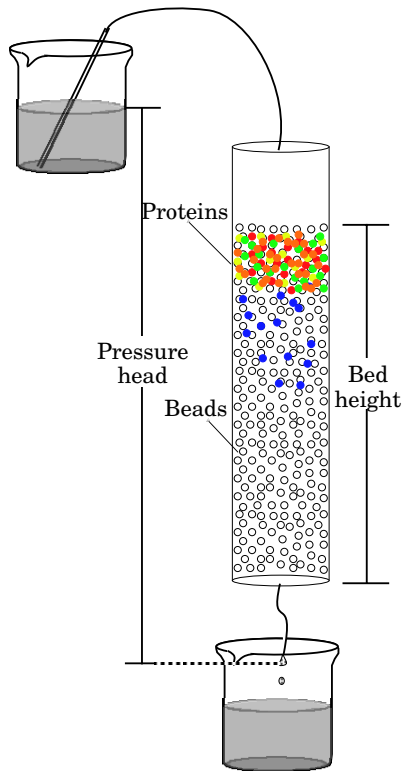
Sources

Strategy

## Differential Centrifugation



## Chromatography



## Resolution

$$R_s = \frac{\sqrt{N}}{4} \left( \frac{\alpha - 1}{\alpha} \right) \left( \frac{k_B}{1 + k_B} \right)$$

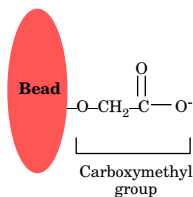
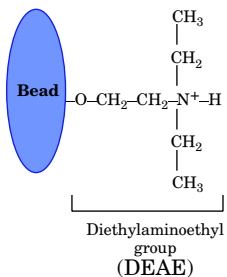
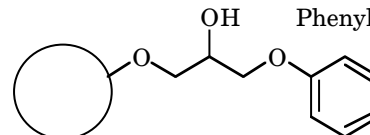
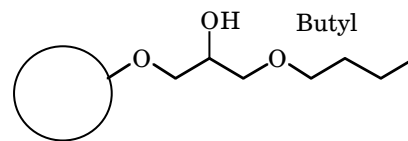
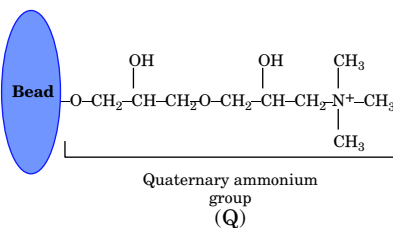
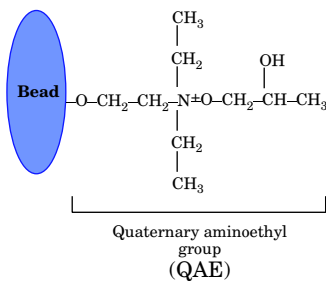
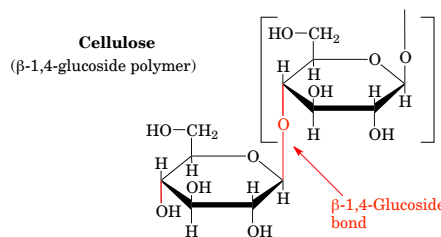
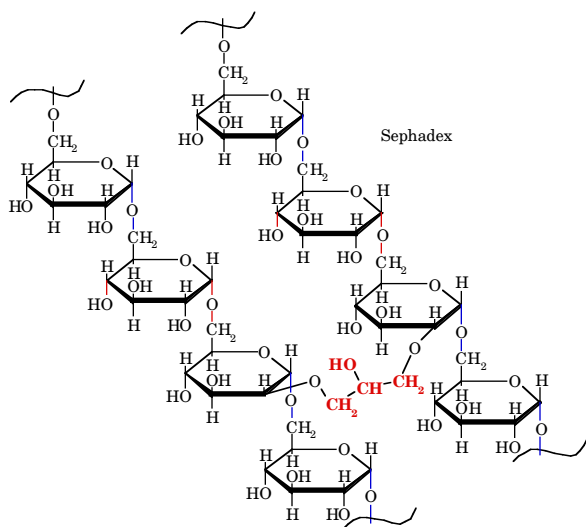
$$R_s = \frac{\sqrt{N}}{4} \left( \frac{t_{r_b} - t_{r_a}}{t_{r_b}} \right)$$

$$\alpha = \frac{k_b}{k_a} = \frac{t_{r_b} - t_m}{t_{r_a} - t_m} = \frac{K_b}{K_a}$$

$\alpha \geq 1$  (although if  $\alpha = 1$  then the analytes are not separating)

# Protein Chromatography Resins

Resin	Carbohydrate	Cross-linking agent
Cellulose	Cellulose	none
Biogel	Polyacrylamide	Bisacrylamide
Sephacel	Cellulose	Epichlorhydrin
Sephacryl	Dextran	Bisacrylamide
Sephadex	Dextran	Epichlorhydrin
Sepharose	Agarose	none
Superdex	Dextran, Agarose	Highly cross-linked
Superose	Agarose	Highly cross-linked



## **Chromatography types**

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**Type**

**Capacity**

**Resolution**

**Expense**

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**Ion exchange**

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**Hydrophobic  
interaction**

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**Gel filtration**

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**Affinity**

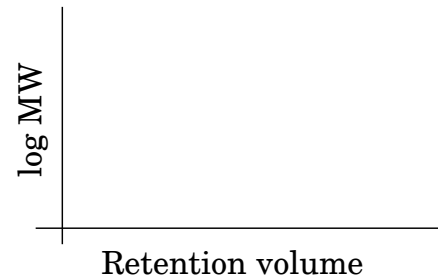
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## Gel Filtration Chromatography

Preparative

Analytical

Stokes radius



## Hofmeister series (effect on hydrophobic interactions)

Anions:  $\text{PO}_4^{3-} > \text{SO}_4^{2-} > \text{CH}_3\text{COO}^- > \text{Cl}^- > \text{Br}^- > \text{NO}_3^- > \text{ClO}_4^- > \text{I}^- > \text{SCN}^-$

Cations:  $\text{NH}_4^+ > \text{Rb}^+ > \text{K}^+ > \text{Na}^+ > \text{Cs}^+ > \text{Li}^+ > \text{Mg}^{2+} > \text{Ca}^{2+} > \text{Ba}^{2+}$

## Effect on water surface tension

$\text{Na}_2\text{SO}_4 > \text{K}_2\text{SO}_4 > (\text{NH}_4)_2\text{SO}_4 > \text{Na}_2\text{HPO}_4 > \text{NaCl} > \text{LiCl} > \text{others} > \text{KSCN}$

## Affinity Chromatography

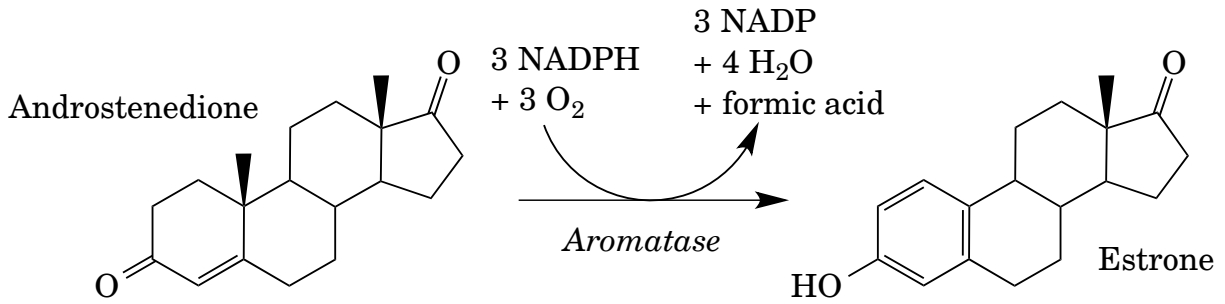
His-tag

Maltose-binding protein

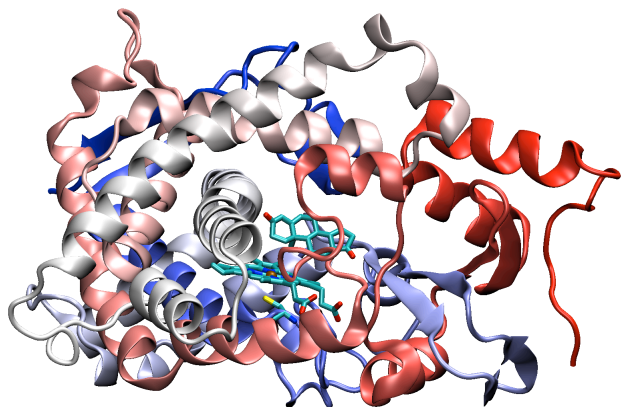
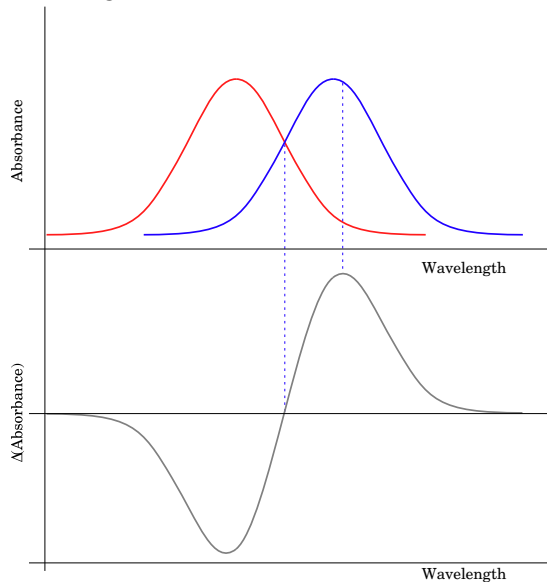
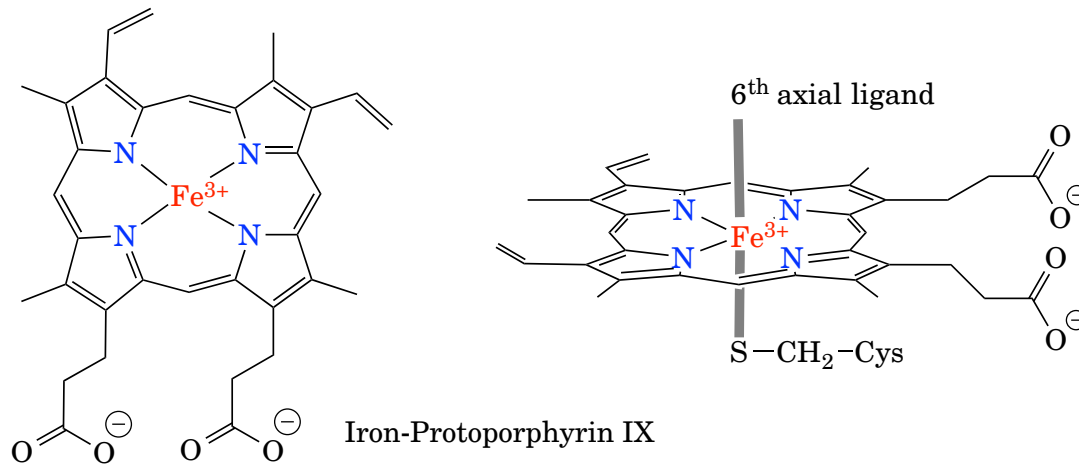
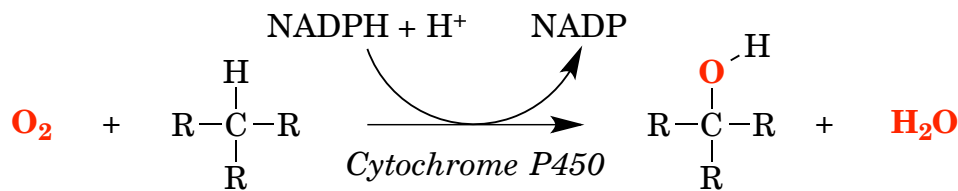
Ligands

Antibodies

## Aromatase



## Cytochrome P450 Enzymes



P450<sub>arom</sub> Aromatase bound to androstenedione (PDB ID 3EQM)