

Chapter 12 - Membranes & Transport

- Major constituents are lipids & proteins
- Membrane structure best described by “Fluid Mosaic” model
- Membrane proteins are associated with the lipid bilayer in many ways
- Membrane fluidity is critical for many processes
- Movement of molecules across membranes is restricted
- Transport can be “passive” or “active” (energy-requiring)

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Membranes

- Define external boundaries
- Regulate “traffic” across boundary
- Segregate metabolic processes
- Bar movement of polar compounds

2

Membrane features

- Flexible
- Tough
- Self-sealing
- Selectively permeable
- Contain an array of proteins

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Molecular constituents of Membranes

Table 12-1

	Components (% by weight)				
	Protein	Phospholipid	Sterol	Sterol type	Other lipids
Human myelin sheath	30	30	19	Cholesterol	Galactolipids, plasmalogens
Mouse liver	45	27	25	Cholesterol	—
Maize leaf	47	26	7	Sitosterol	Galactolipids
Yeast	52	7	4	Ergosterol	Triacylglycerols, steryl esters
Paramecium (ciliated protist)	56	40	4	Stigmasterol	—
<i>E. coli</i>	75	25	0	—	—

Varies according to species, tissue and membrane “face”

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Membrane Structure - The Fluid Mosaic Model

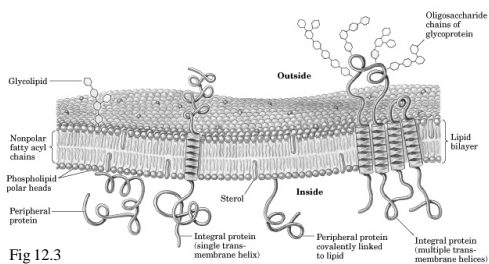
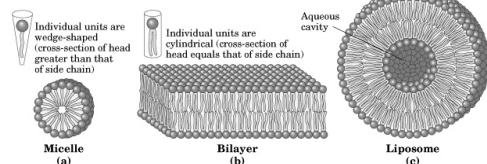


Fig 12.3

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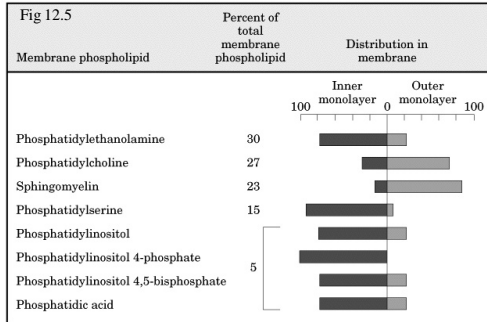
The Lipid bilayer

Fig 12.4



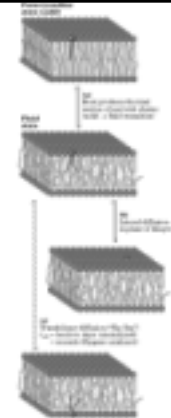
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Membrane lipids are asymmetrically distributed



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Membrane lipids are in constant motion



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Cells regulate lipid composition

table 12-2

Fatty Acid Composition of *E. coli* Cells Cultured at Different Temperatures

	Percentage of total fatty acids*			
	10 °C	20 °C	30 °C	40 °C
Myristic acid (14:0)	4	4	4	8
Palmitic acid (16:0)	18	25	29	48
Palmitoleic acid (16:1)	26	24	23	9
Oleic acid (18:1)	38	34	30	12
Hydroxymyristic acid	13	10	10	8
Ratio of unsaturated to saturated [†]	2.9	2.0	1.6	0.38

Source: Data from Marr, A.G. & Ingraham, J.L. (1962) Effect of temperature on the composition of fatty acids in *Escherichia coli*. *J. Bacteriol.* **84**, 1260.

*The exact fatty acid composition depends not only on growth temperature but on growth stage and growth medium composition.

[†]Calculated as the total percentage of 16:1 plus 18:1 divided by the total percentage of 14:0 plus 16:0. Hydroxymyristic acid was omitted from this calculation.

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Membrane proteins diffuse laterally in the bilayer

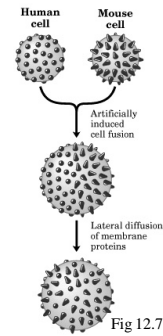
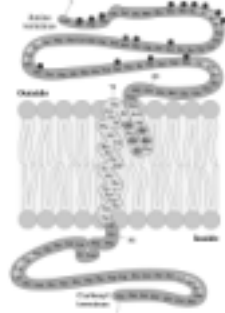


Fig 12.7

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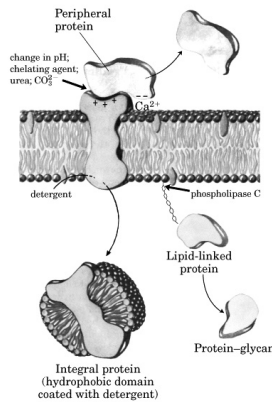
Glycophorin - Quintessential membrane protein



Outer surface, inner surface & transmembrane domains

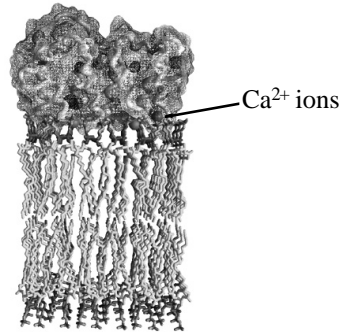
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Peripheral & integral membrane proteins



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Electrostatic interactions - 1 biological role for Ca^{2+}



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Covalently attached lipids anchor some peripheral proteins

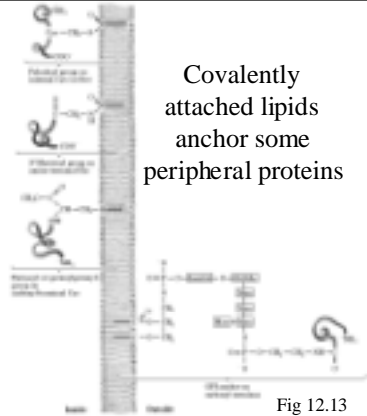


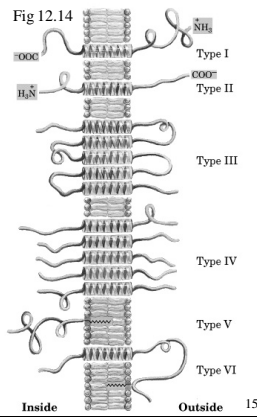
Fig 12.13

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Integral membrane proteins

Membrane-spanning α -helices (and β -sheets)

6 categories



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β -sheet-containing integral membrane proteins have " β -barrel" motifs

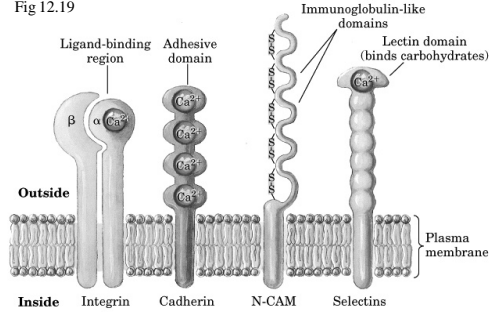


Fig 12.18

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Integral membrane proteins mediate cell-cell interactions

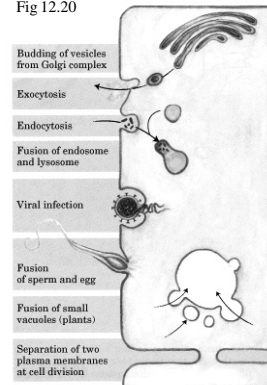
Fig 12.19



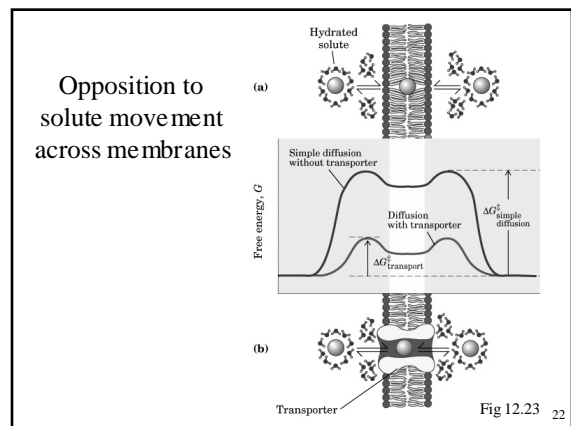
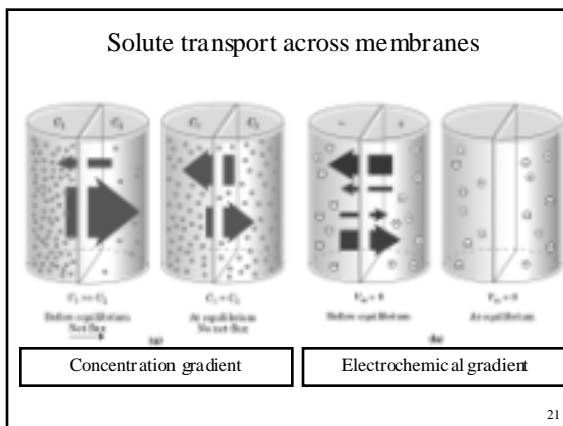
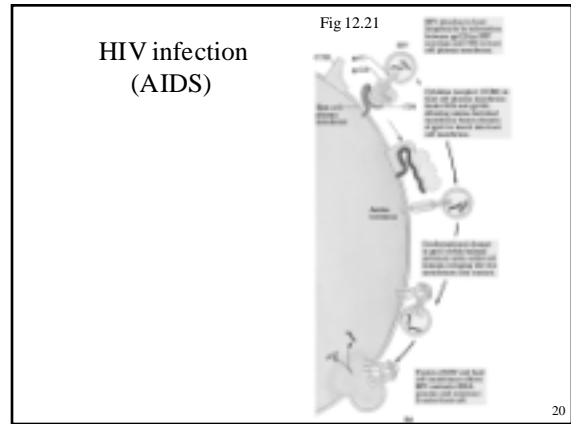
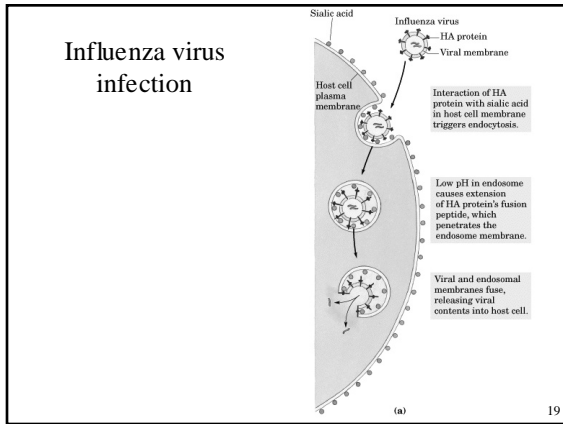
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Membrane fusion central to many biological processes

Fig 12.20



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Transmembrane passage of polar compounds

- Integral membrane proteins (transporters)
- “Passive” or “active” transport
- Action analogous to enzymes
 - Bind substrates via multiple weak interactions
 - Negative free energy of binding counterbalances positive free energy of H_2O removal
 - Span of lipid bilayer by transporter provides alternative path for substrate movement

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3 examples of Passive transport (RBC's)

- Aquaporins (H_2O)
- Glucose T transporter
- Chloride-bicarbonate exchanger

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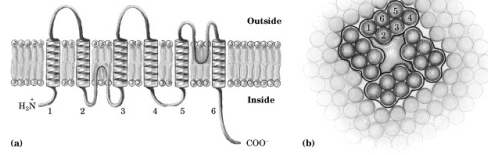
Passive transport - Hallmarks

- High rates of diffusion down concentration or electrochemical gradients (high \rightarrow low)
- Saturability of transporter
- Specificity of transporter

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Aquaporins

Fig 12.24



Channel for rapid ($5 \times 10^8 \text{ s}^{-1}$) movement of water

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Glucose transporter (GluT1)

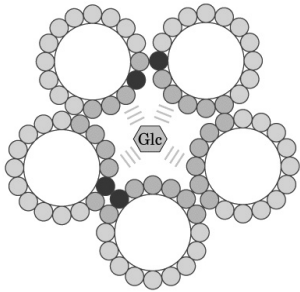
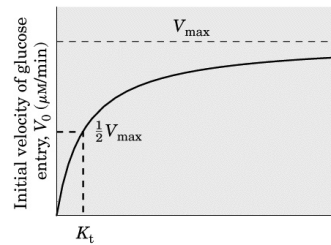


Fig 12.25 (c)

50,000 x more glucose transported

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Transport analogous to enzymatic reaction



Extracellular glucose concentration, $[S]_{\text{out}}$ (mM)

Substrate = Glucose_{outside}
Product = Glucose_{inside}

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Blood [glucose] = 5 mM
GluT1 $K_T = 1.5 \text{ mM}$

GluT1 saturated -
operates at V_{max}

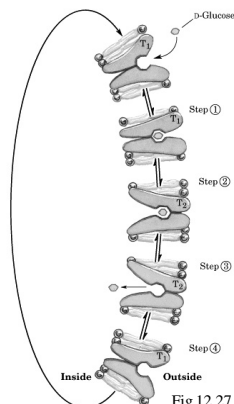
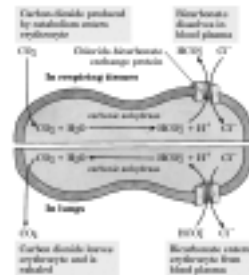


Fig 12.27

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Chloride-bicarbonate exchanger



- Cotransport system: simultaneous & opposite movement of solutes
- Essential for CO_2 transport; \uparrow permeability by 10^6

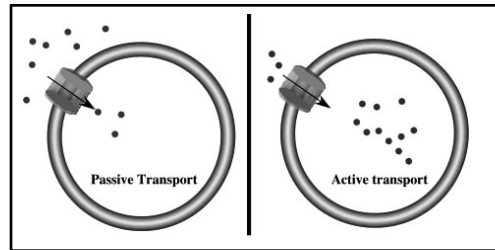
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Cotransport systems

- Antiport (Chloride-bicarbonate exchanger): 2 substrates in opposite direction
- Symport: 2 substrates in same direction
- Uniport (Glucose transporter): one substrate

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Active transport moves solutes against their concentration or electrochemical gradient



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Primary & Secondary Active Transport

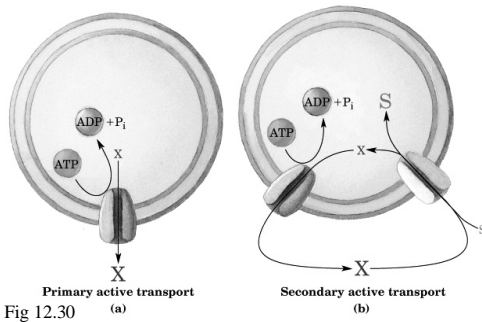


Fig 12.30

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Active Transport - ATPase

Table 12-4

Four Classes of Transport ATPases			
	Organism or tissue	Type of membrane	Role of ATPase
P-type ATPases			
Na ⁺ /K ⁺	Animal tissues	Plasma	Maintains low [Na ⁺], high [K ⁺] inside cell, creates transmembrane electrical potential
H ⁺ /K ⁺	Acid-secreting (parietal) cells of mammals	Plasma	Acidifies contents of stomach
H ⁺	Fungi (<i>Neurospora</i>)	Plasma	Create H ⁺ gradient to drive secondary transport of extracellular solutes into cell
H ⁺	Higher plants	Plasma	
Ca ²⁺	Animal tissues	Plasma	Maintains low [Ca ²⁺] in cytosol
Ca ²⁺	Myocytes of animals	Sarcoplasmic reticulum (endoplasmic reticulum)	Sequesters intracellular Ca ²⁺ , keeping cytosolic [Ca ²⁺] low
Cd ²⁺ , Hg ²⁺ , Cu ²⁺	Bacteria	Plasma	Pumps heavy metal ions out of cell
V-type ATPases			
H ⁺	Animals	Lysosomal, endosomal, secretory vesicles	Create low pH in compartment, activating proteases and other hydrolytic enzymes
H ⁺	Higher plants	Vacuolar	
H ⁺	Fungi	Vacuolar	
F-type ATPases			
H ⁺	Eukaryotes	Inner mitochondrial	Catalyze formation of ATP from ADP + P _i
H ⁺	Higher plants	Thylakoid	
H ⁺	Prokaryotes	Plasma	
Multidrug transporter			
	Animal tumor cells	Plasma	Removes a wide variety of hydrophobic natural products and synthetic drugs from cytosol, including vinblastine, doxorubicin, actinomycin D, mitomycin, taxol, colchicine, and paclitaxel

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P-type ATPases

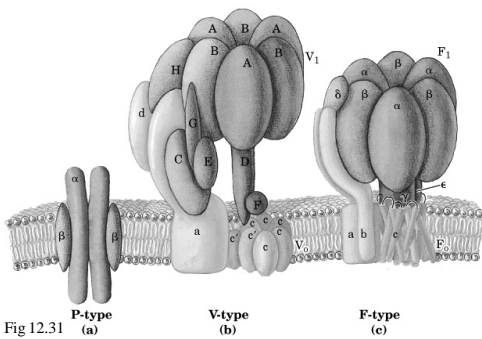


Fig 12.31

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P-type ATPase: The Na⁺K⁺ATPase

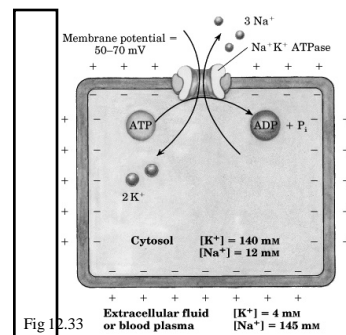
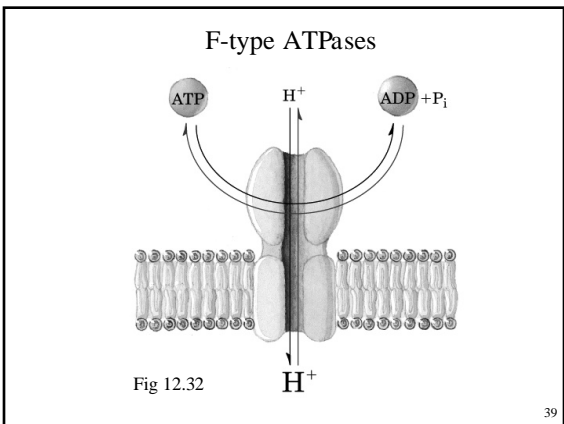
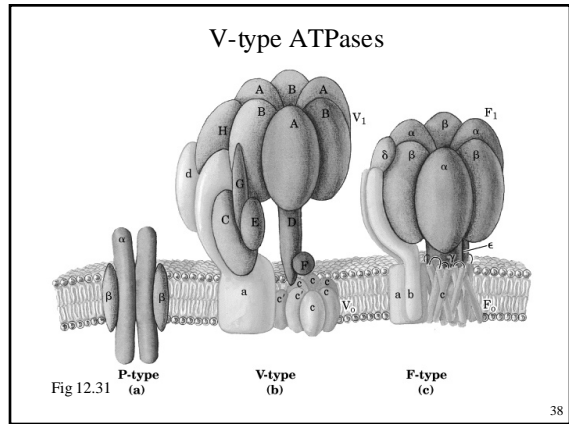
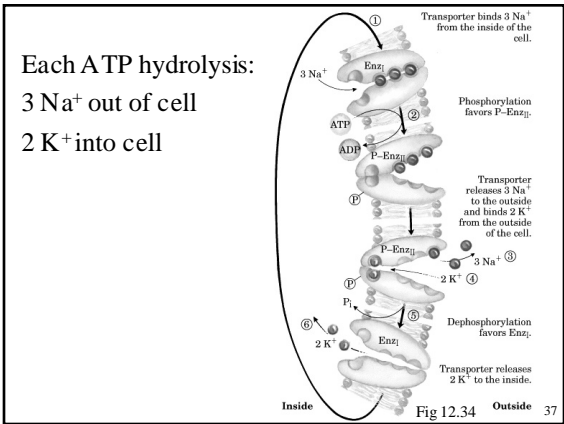


Fig 12.33

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“Bad ATPase” - The multidrug transporter

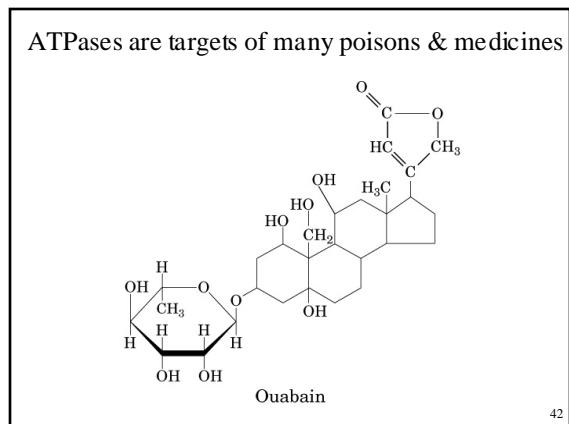
- Found in plasma membrane of some tumor cells
- ATP transporter
- Pumps many hydrophobic drugs out of tumor cells

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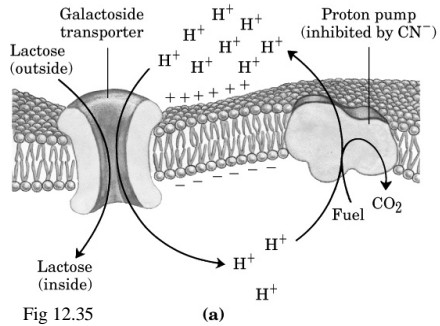
ATP-driven Ca^{2+} pumps (P-type)

- Ca^{2+} far lower inside vs. outside cells
- Ca^{2+} forms insoluble compounds with phosphate
- Ca^{2+} pumped out of cells by specific pumps
- Other ATPases (SERCA's) move Ca^{2+} into the ER and sarcoplasmic reticulum (SR) of myocytes
- Similar to Na^+K^+ ATPase - cycling between phosphorylated & dephosphorylated forms

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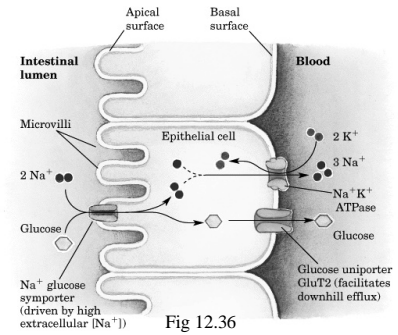


Ion gradients provide the energy for 2^o active transport



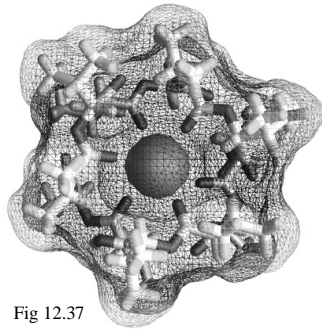
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Glucose transport in intestinal epithelial cells



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Ionophores "short circuit" ion gradients



Valinomycin - potent antibiotic

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Ion channels allow rapid movement of ions across membranes

Ion channels distinguished from transporters

- Flux much higher than channels ($\uparrow 10^8 \text{ ions} \cdot \text{s}^{-1}$)
- Ion channels are not saturable
 - No V_{max} , linear increase with concentration
- Ion channels are gated
 - Ligand-gated channels
 - Voltage-gated channels

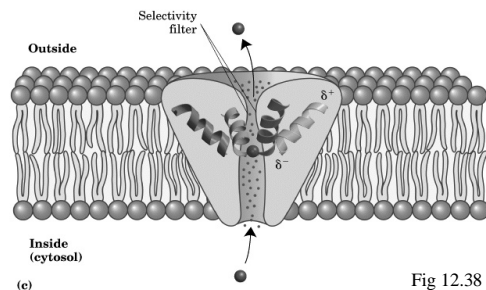
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3 examples

- Bacterial K^+ channel
- Acetylcholine receptor ion channel
- Voltage-dependent Na^+ channel (neurons)

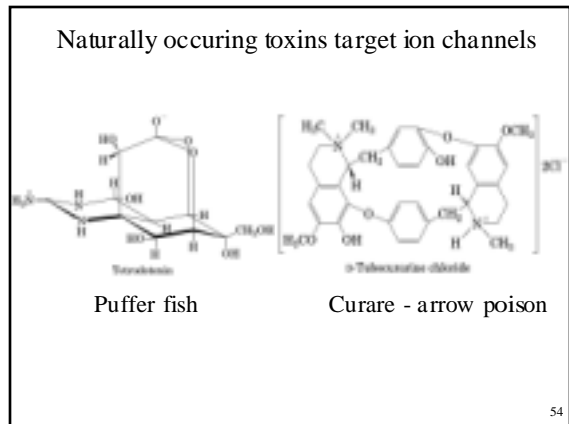
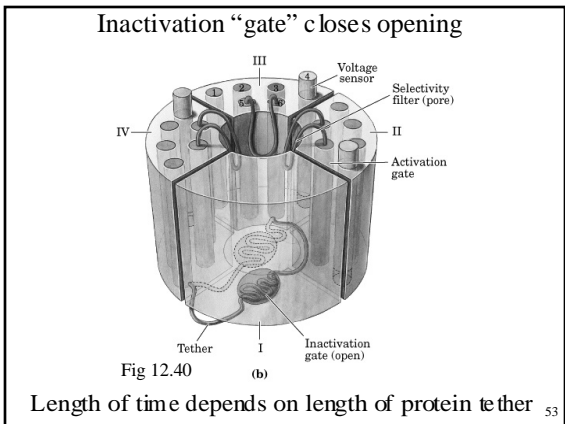
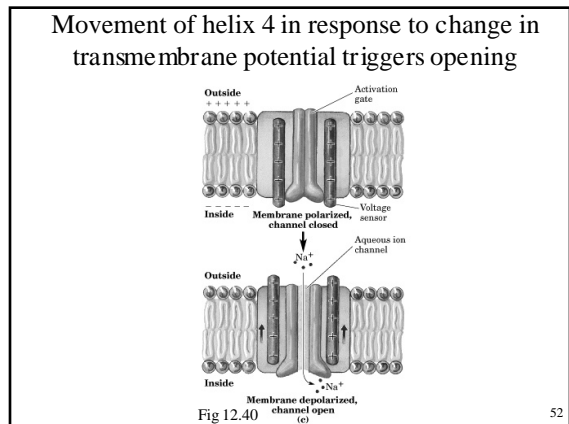
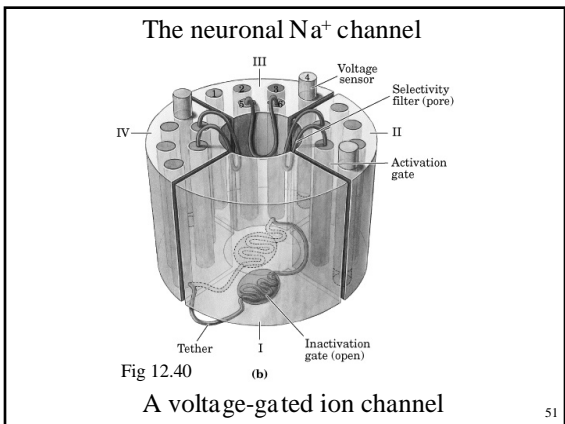
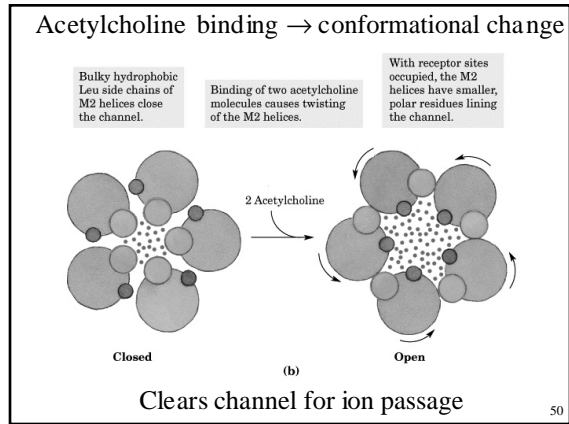
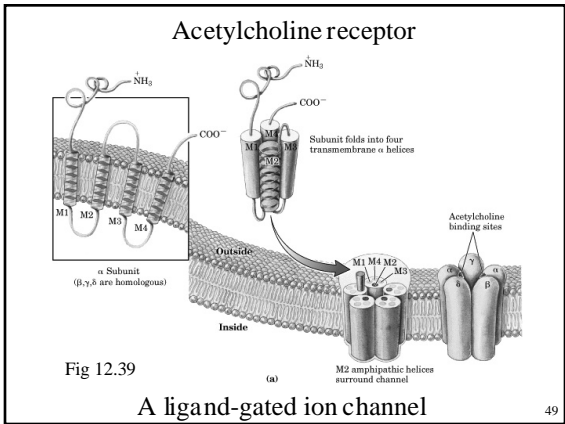
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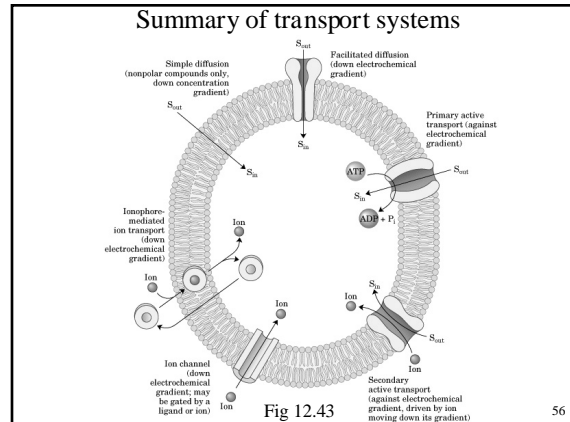
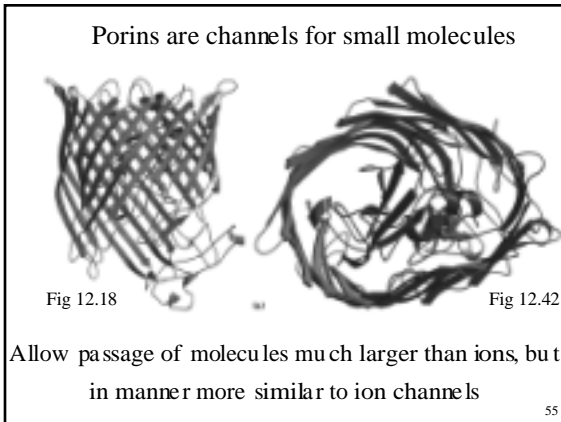
Bacterial K^+ channel



Selectivity filter screens out alternative ions

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- Chapter 12 - Summary
- Biological membranes
 - Define cellular boundaries
 - Divide cells into discrete compartments
 - Organize reaction sequences (pathways, etc.)
 - Roles in signal reception, transduction
 - Membrane composition
 - Lipids (phospholipids, sphingolipids, glycolipids)
 - Protein (peripheral & integral)
 - Varies in composition according to species, cell type, organelle & even "face" of membrane
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- Fluid Mosaic Model
 - Lipid bilayer
 - Fatty acyl chains of phospholipids & sterols oriented towards interior
 - Lipids & proteins free to diffuse laterally
 - Fluidity affected by temperature, fatty acid composition (saturation) & sterol content
 - Cells regulate their lipid composition to reflect changes in their environment
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- Membrane proteins
- Peripheral
 - Loose association with membrane
 - Electrostatic interactions, H-bonds or covalent attachment to lipids
 - Integral
 - Firm association with membrane
 - Hydrophobic interactions with lipid bilayer
 - Both α -helical and β -sheet-type structures
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- Membranes are structurally & functionally Asymmetric
- Two sides differ in lipid/protein composition
 - Different functional roles (adhesion, signal reception vs. cytoskeleton attachment, exocytosis, etc.)
 - Variety of proteins mediate protein fusion (SNARE's)
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Lipid bilayer is impermeable to polar substances

- Exception = water (concentration v. high)
- Other polar molecules cross via transporters or ion channels

Transporters

- Similar to enzymes - show saturation & substrate specificity
- Passive or Active transport
 - Passive - down an electrochemical gradient
 - Active - against an electrochemical gradient, requires energy (ATP, oxidation, light, 2^o transport)

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Transport

- Uniport: single species transport
- Symport: simultaneous transport, same direction
- Antiport: simultaneous transport, opposite direction

Differences in $[Na^+]$ & $[K^+]$ between inside/outside of Cell maintained by active transport via Na^+K^+ ATPase

- Resulting gradient used as energy source for 2^o transport

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4 general types of ATPases

- P-type
 - Reversible phosphorylation
 - Vanadate inhibition (P_i analog)
 - Ex's: Na^+K^+ ATPase, SERCA (Ca^{2+} transporter)
- V-type
 - Produce proton gradients across intracellular membranes
- F-type
 - Produce proton gradients
 - Reverse reaction (Proton flow) drives ATP synthesis
 - Mitochondria & chloroplasts

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• Multidrug transporter

- Tumor cells
- Energy of ATP to export hydrophobic drugs

• Ionophores

- Lipid soluble
- Bind & transport ions across membranes
- Short-circuit electrochemical gradients

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Ion channels

- Provide hydrophilic pores for movement of specific ions
- Movement down electrochemical gradient
- Differ from transporters
 - Unsaturable
 - V. high flux rate
- Gated by ligand binding or voltage potentials
- K^+ channel prototype
 - Ion movement restricted by "selectivity filter"
 - Protein domain "gate" obstructs channel when not activated

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Porins

- Integral membrane β -barrel proteins
- Substrate binding opens pore, allowing movement of molecule to interior

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