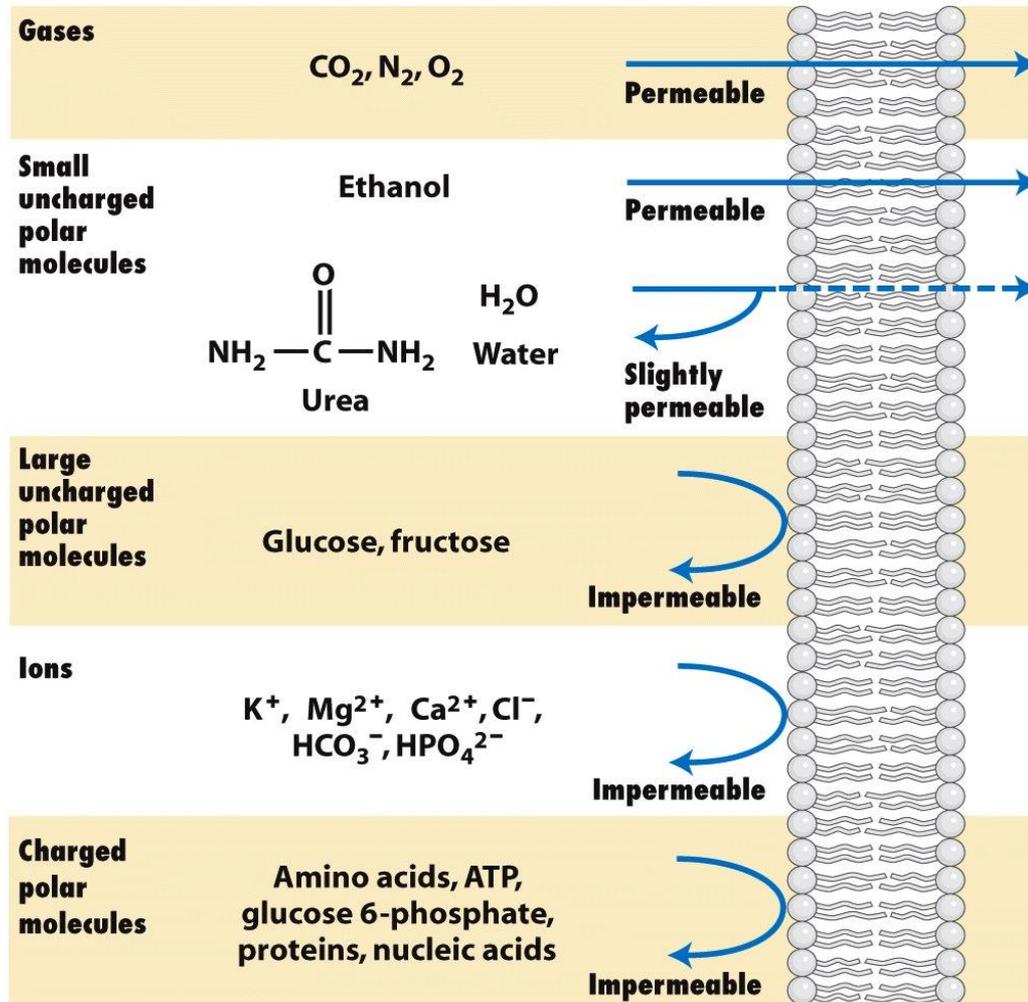


Chapter 5. Ion Channels



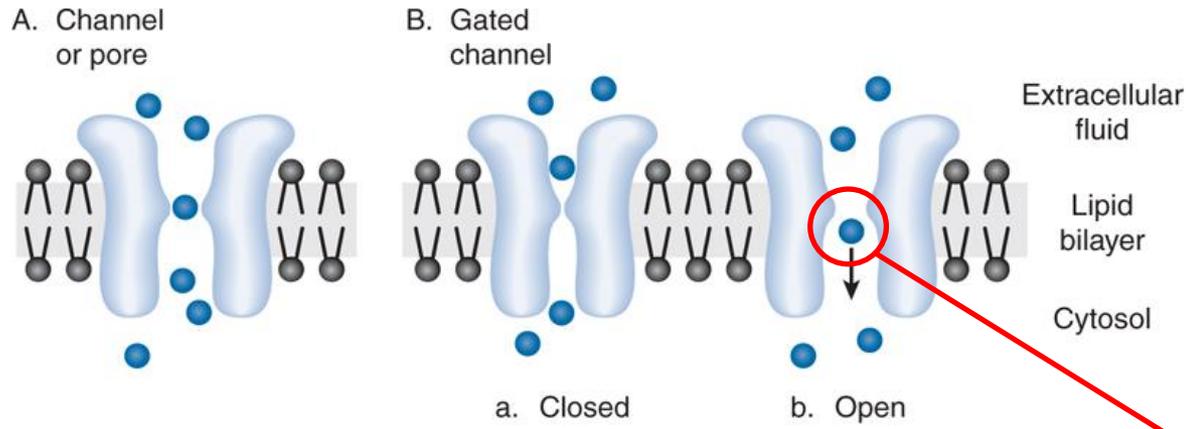
A pure artificial phospholipid bilayer is permeable to small hydrophobic molecules and small uncharged polar molecules.

Ion channels

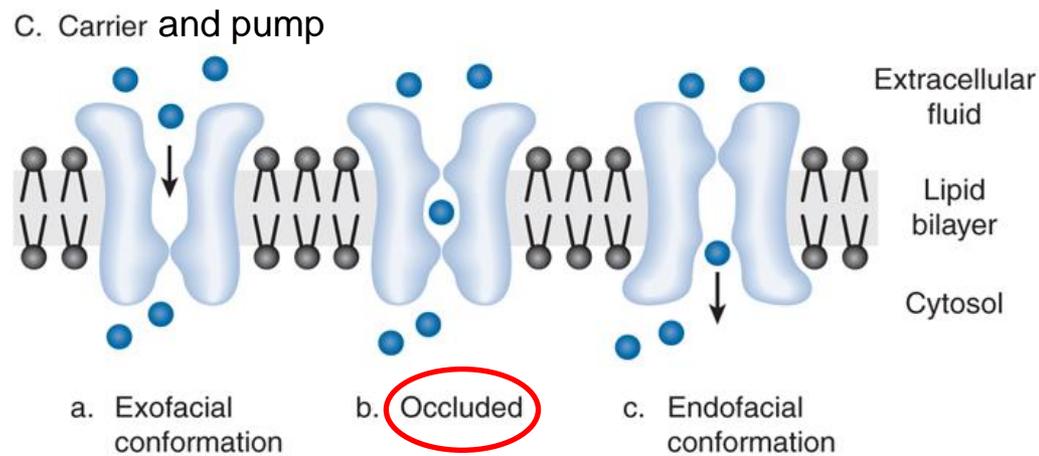
- Integral membrane protein
- select specific ions (selectivity)
- open and close (gating) by electrical, mechanical, or chemical signals
- conduct ions across the membrane

TRANSPORTER	TURNOVER NUMBER* (PER SEC)
K ⁺ channel	30,000,000
Valinomycin (carrier)	30,000
Glucose carrier (GLUT-1)	3,000
Na ⁺ /Ca ²⁺ exchanger	2,000
Ca ²⁺ pump (SERCA)	200
Na ⁺ pump	150
enzymes	10-1000

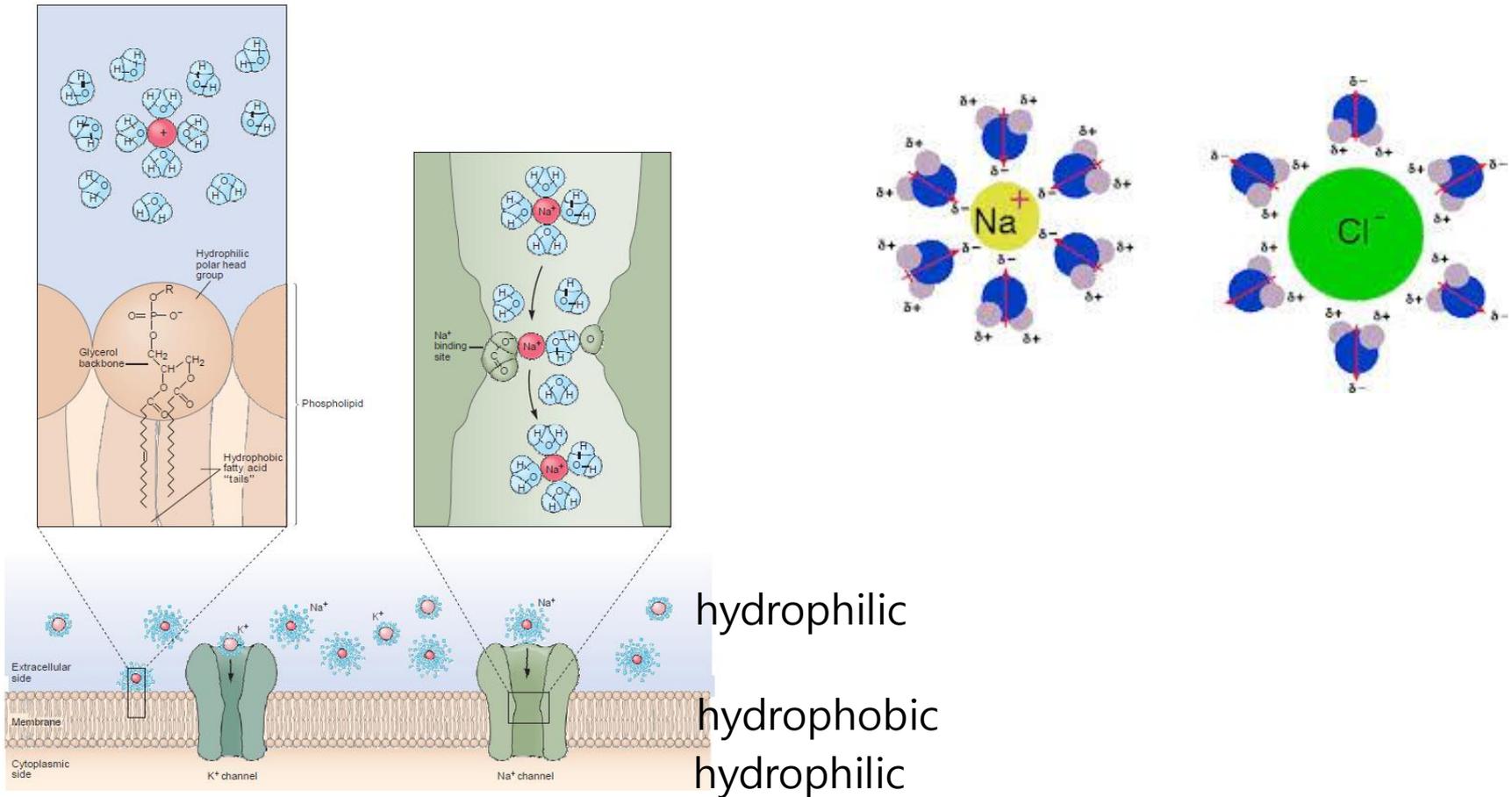
Transport across biological membrane through channel, carrier, and pump protein



Continuous pathway

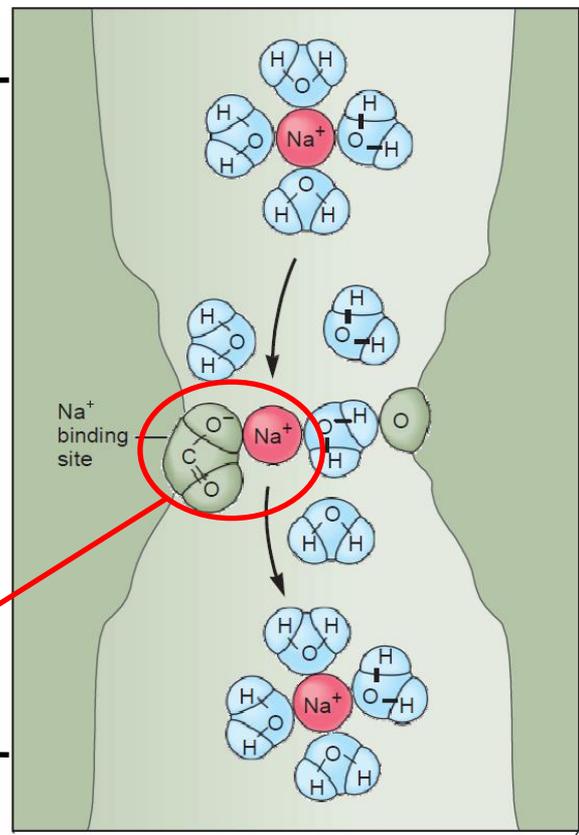


Ion channels provide polar pathways for ions to cross the membrane



- Relatively fast transport
- Channel density controls membrane permeability to a substance
- Rate of transport through open channels depends on the net driving force
- Transport through some channels is controlled by gating

Selectivity filter



Hydration (water shell)

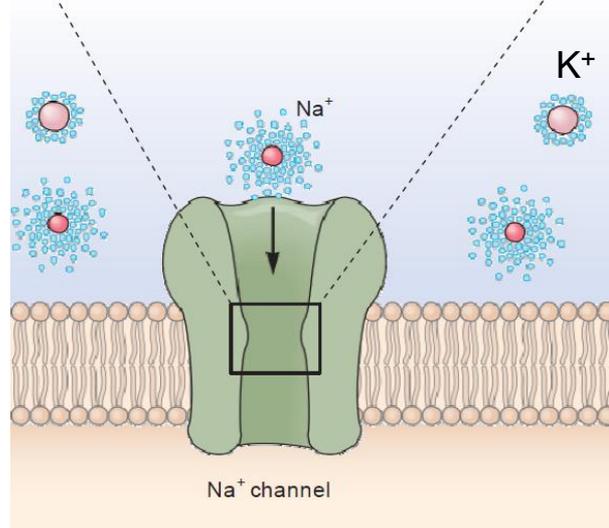
Dehydration

Transfer

Rehydration (water shell)

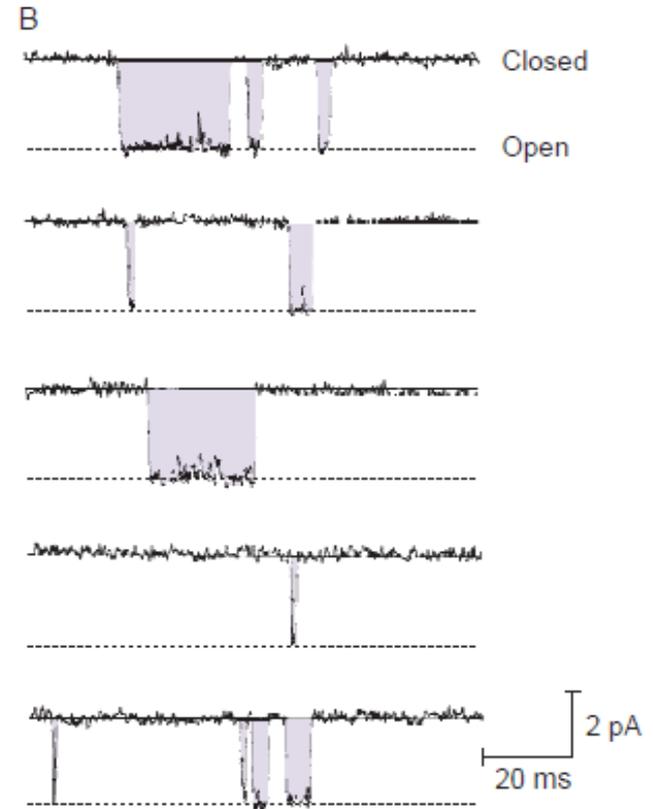
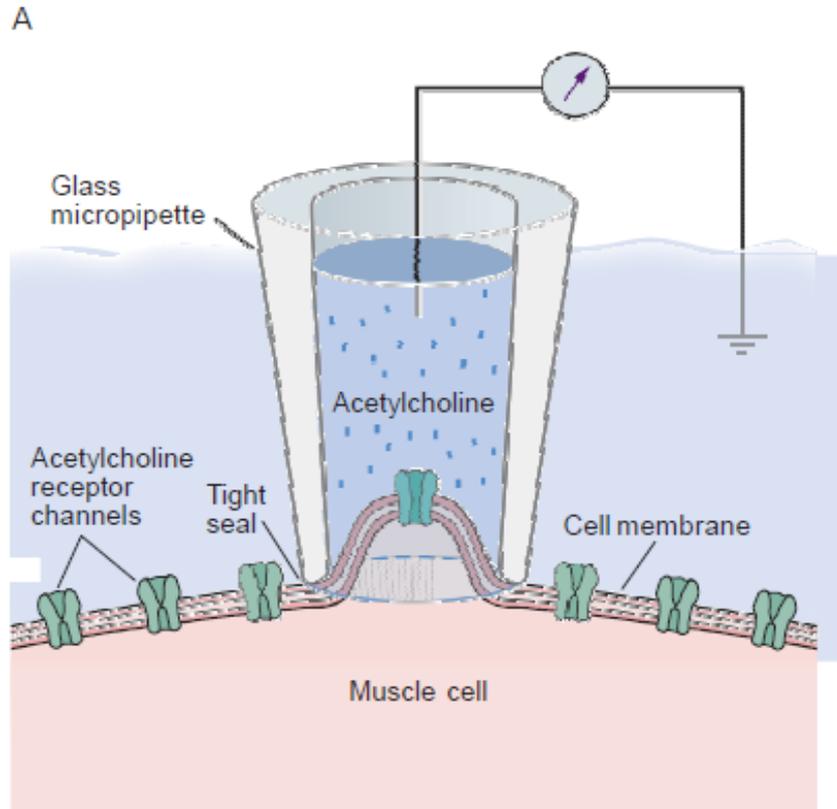
Na^+ binding site

electrostatic interaction
(< 1 μs)

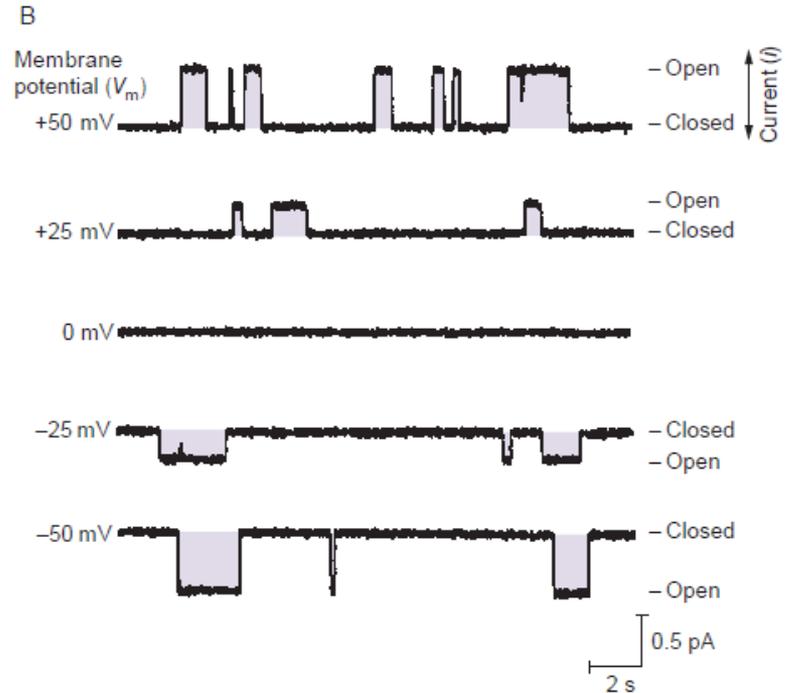
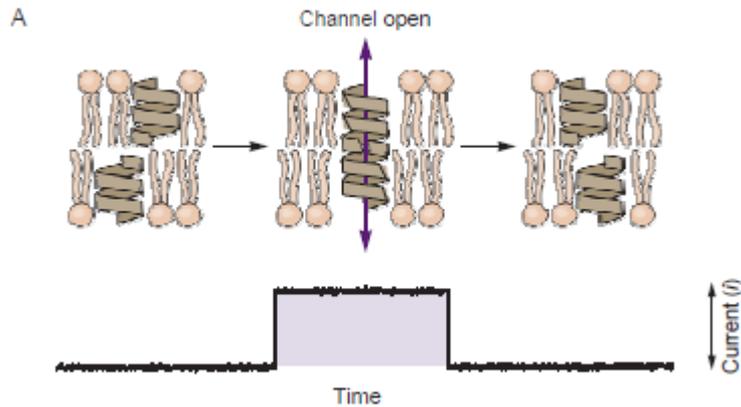


Na^+ channel

Recording Current in Single Ion Channels: The Patch Clamp



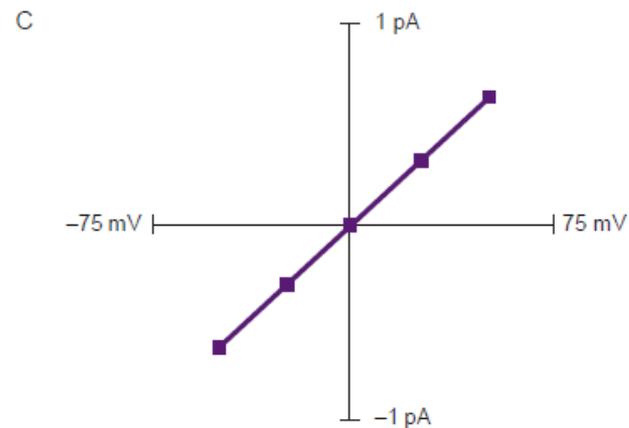
Characteristics of the current in a single ion channel



Ohm's law

$$i = V/R \text{ or } i = \gamma \times V$$

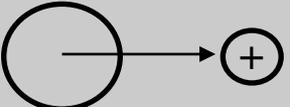
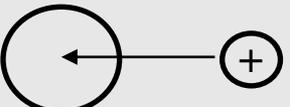
(Conductance; $\gamma = 1/R$)



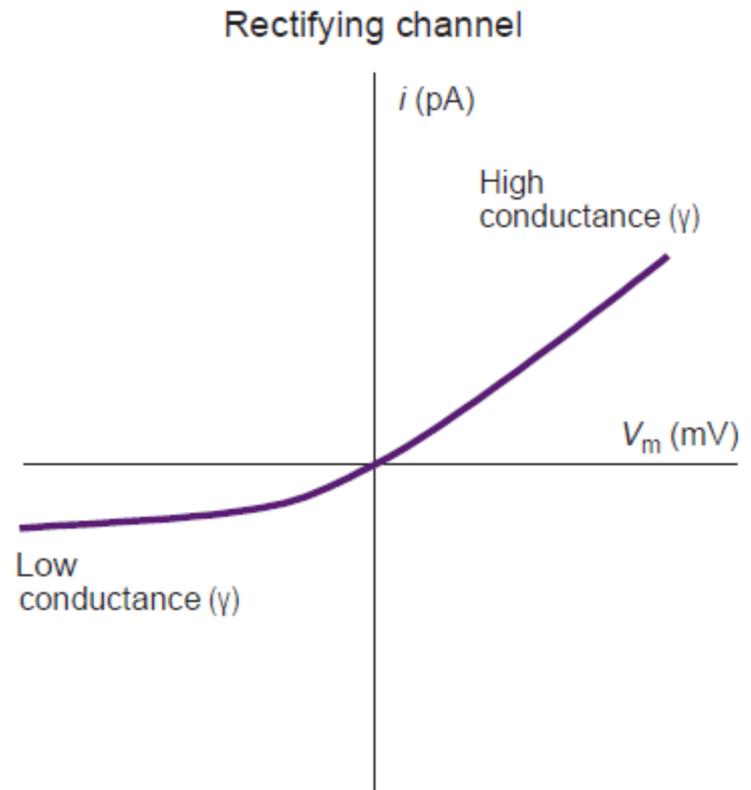
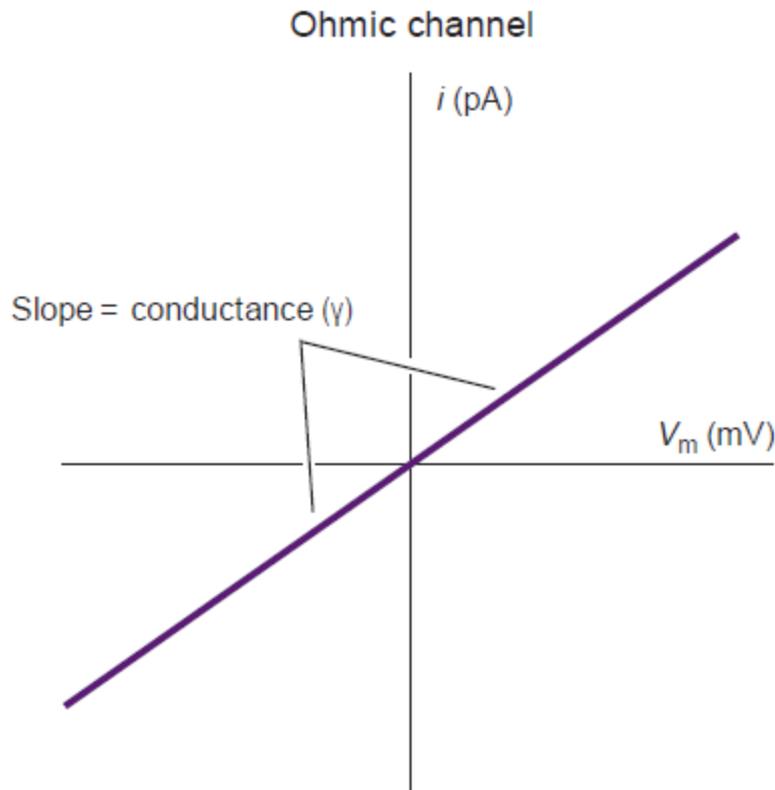
Ionic fluxes and ionic currents

Ionic flux (J)	Ionic current (I)
number of moles of ions moving through a unit area of membrane per unit time	Movement of charges per unit time
[mol/cm ²]/sec	Coulombs/sec, A

$$I = zF \times J \times A_{\text{mem}}, \text{ where } F=96,485 \text{ coulombs/mol}$$

Flow	Direction and Sign of Flux (J)	Direction and Sign of current (I)
	Outward, negative ($J < 0$)	Outward, positive ($I > 0$)
	Inward, positive ($J > 0$)	Inward, negative ($I < 0$)
	Inward, positive ($J > 0$)	Outward, positive ($I > 0$)
	Outward, negative ($J < 0$)	Inward, negative ($I < 0$)

Current–voltage relations

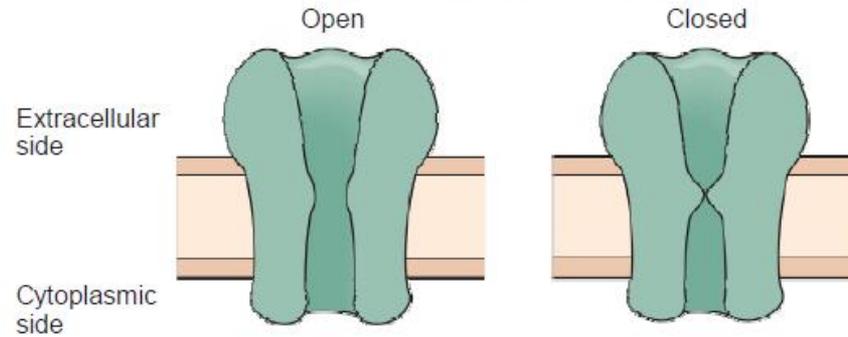


Ohm's law

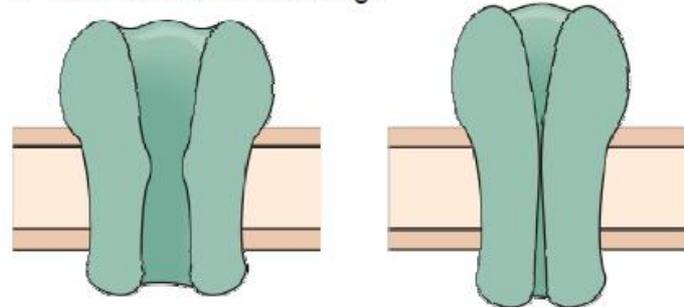
$$i = V/R \text{ or } i = \gamma \times V$$

Models for the opening and closing of ion channels

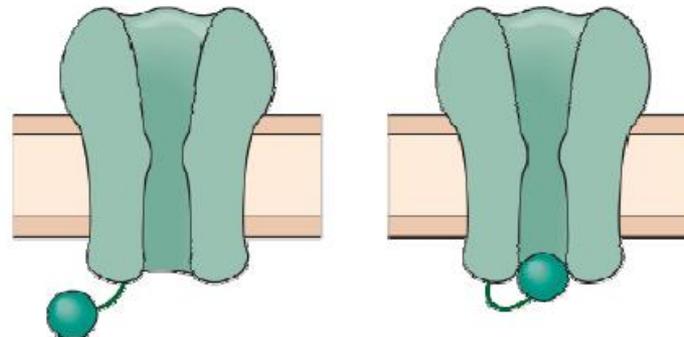
A Conformational change in one region



B General structural change

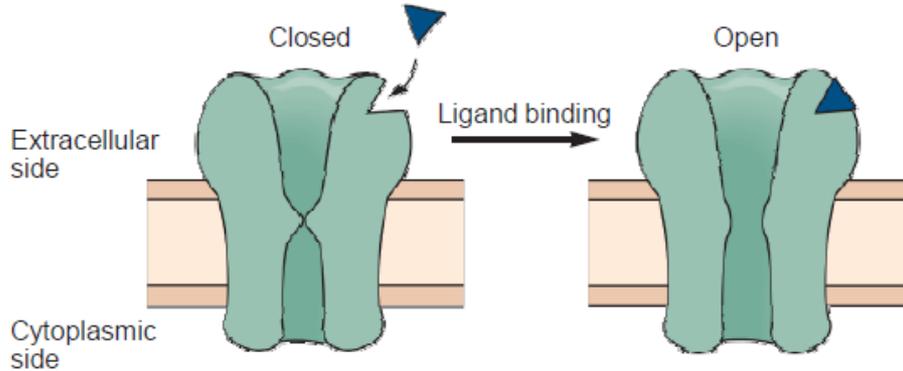


C Blocking particle

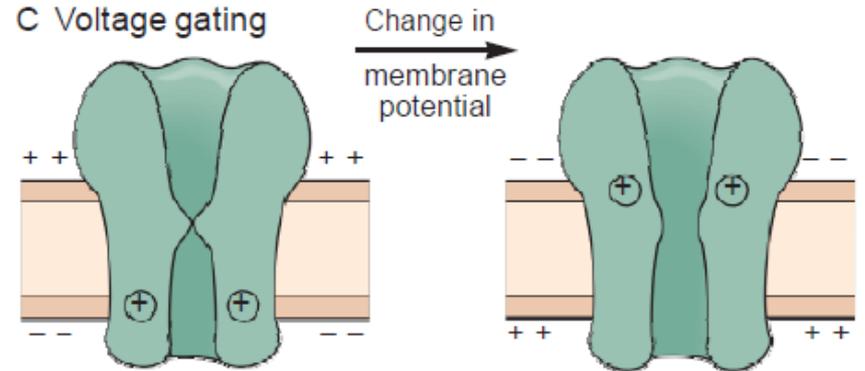


Classification by gating of ion channel

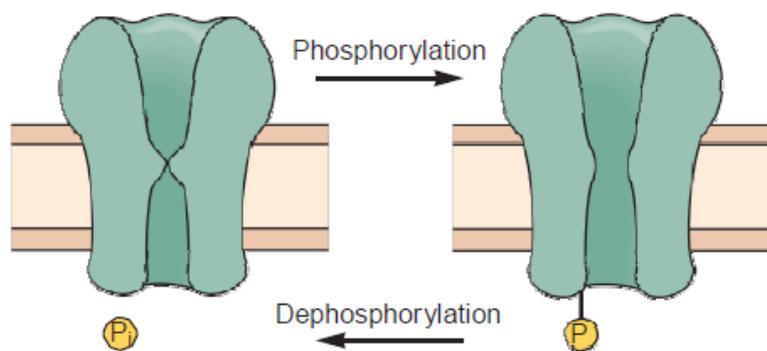
A Ligand gating



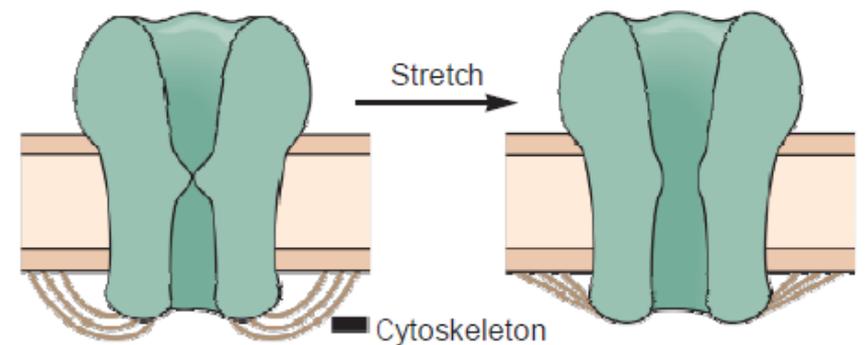
C Voltage gating



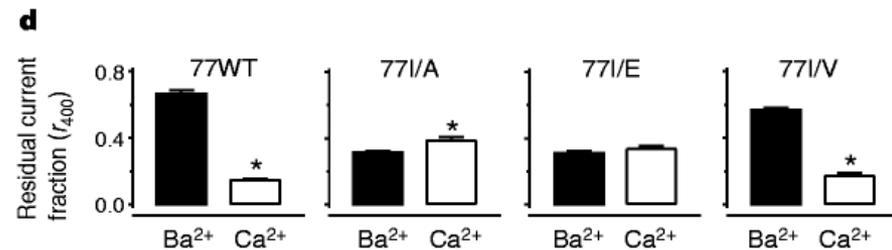
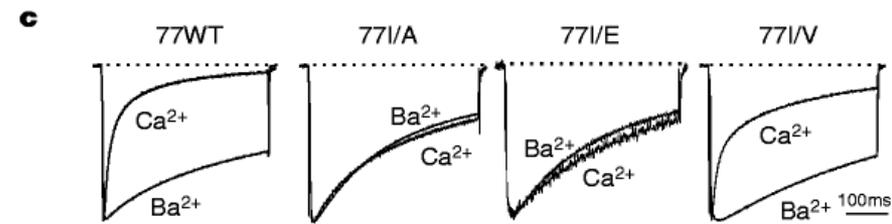
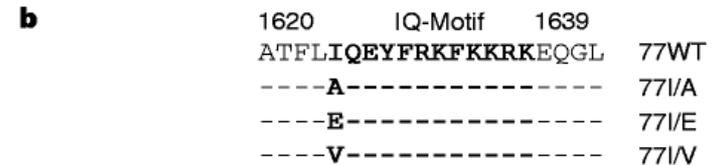
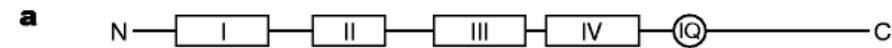
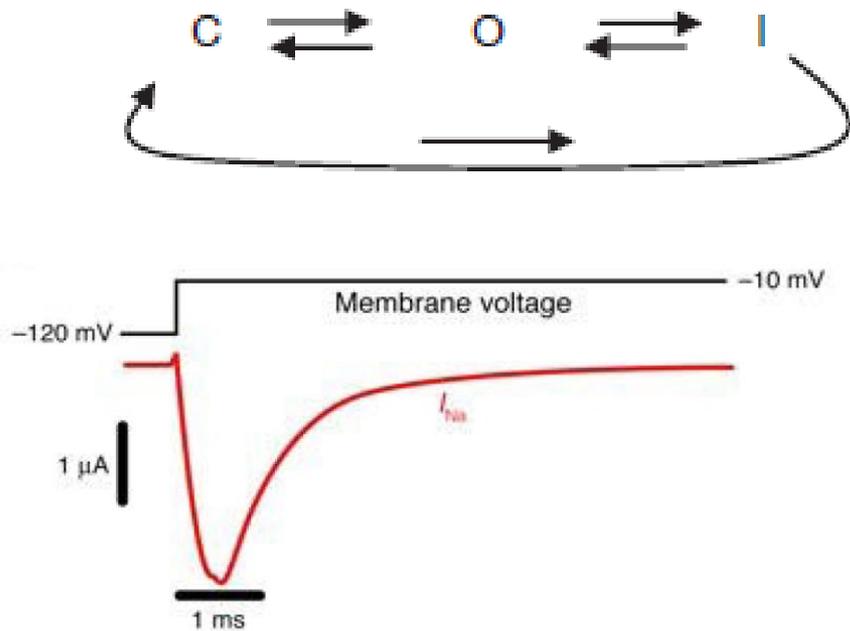
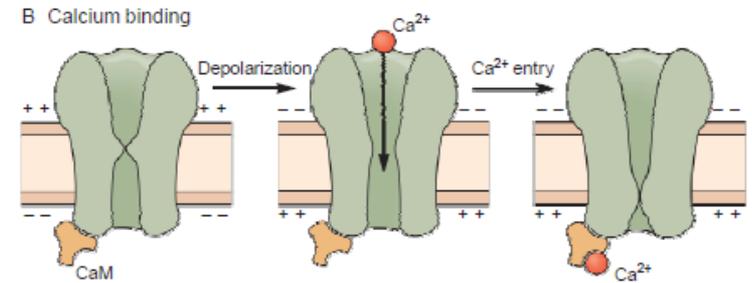
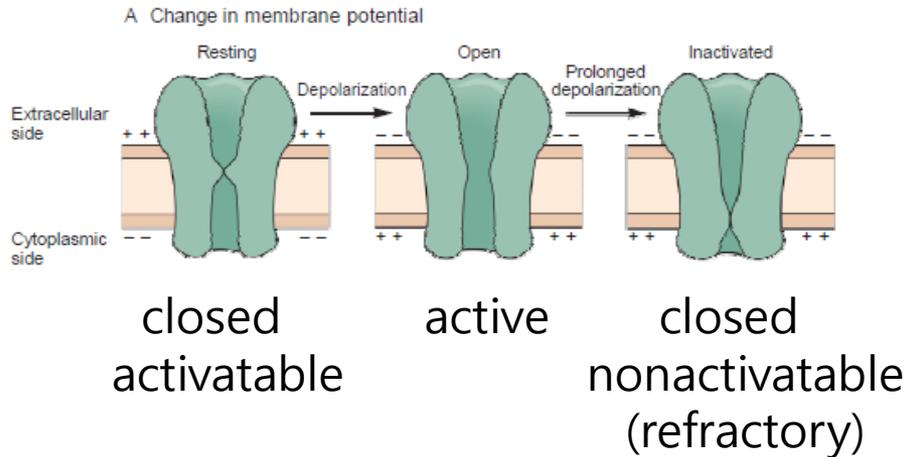
B Phosphorylation gating



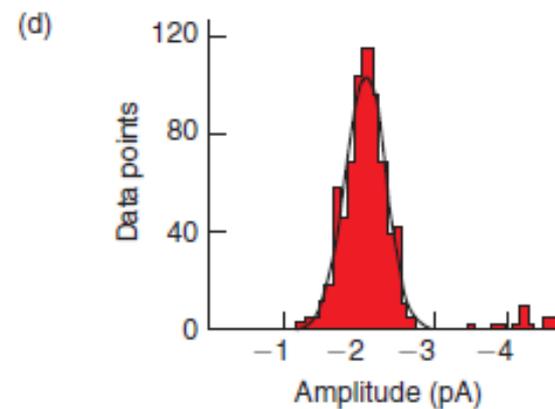
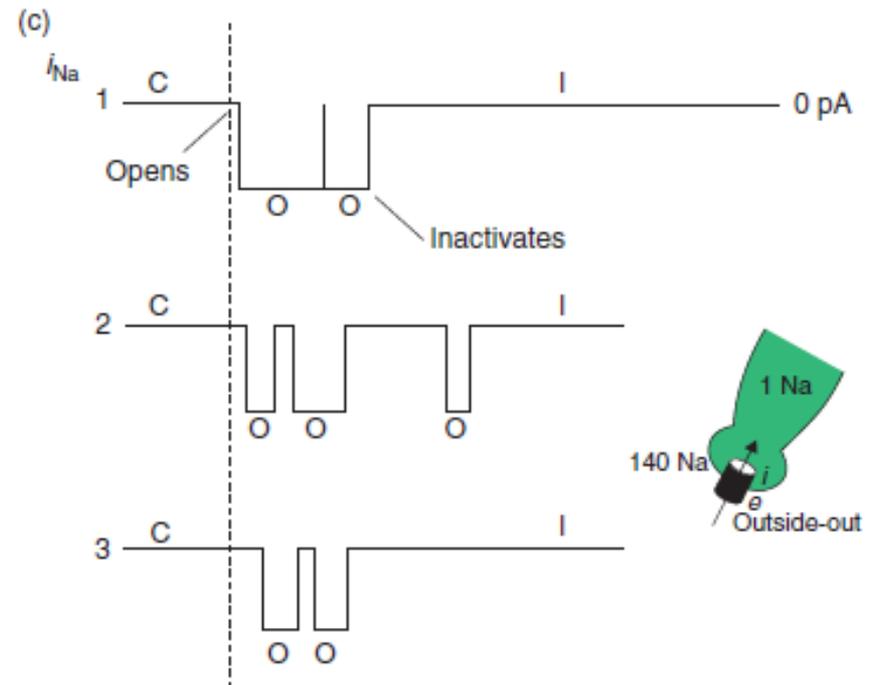
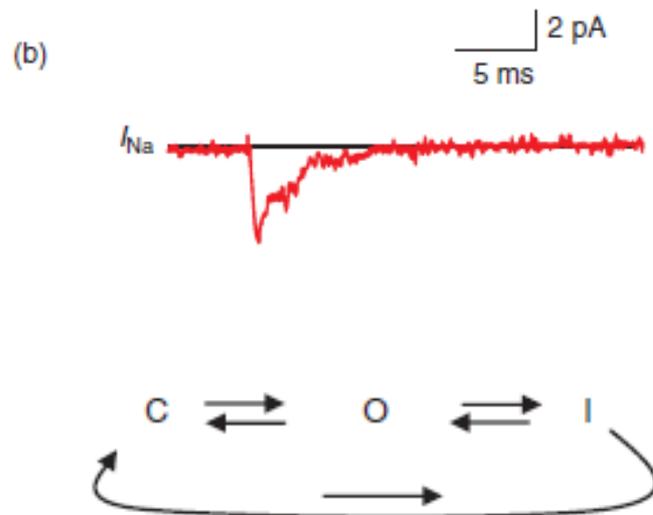
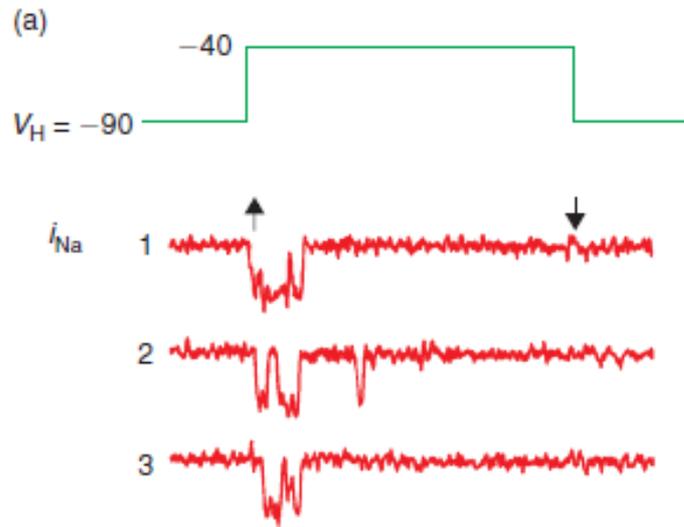
D Stretch or pressure gating



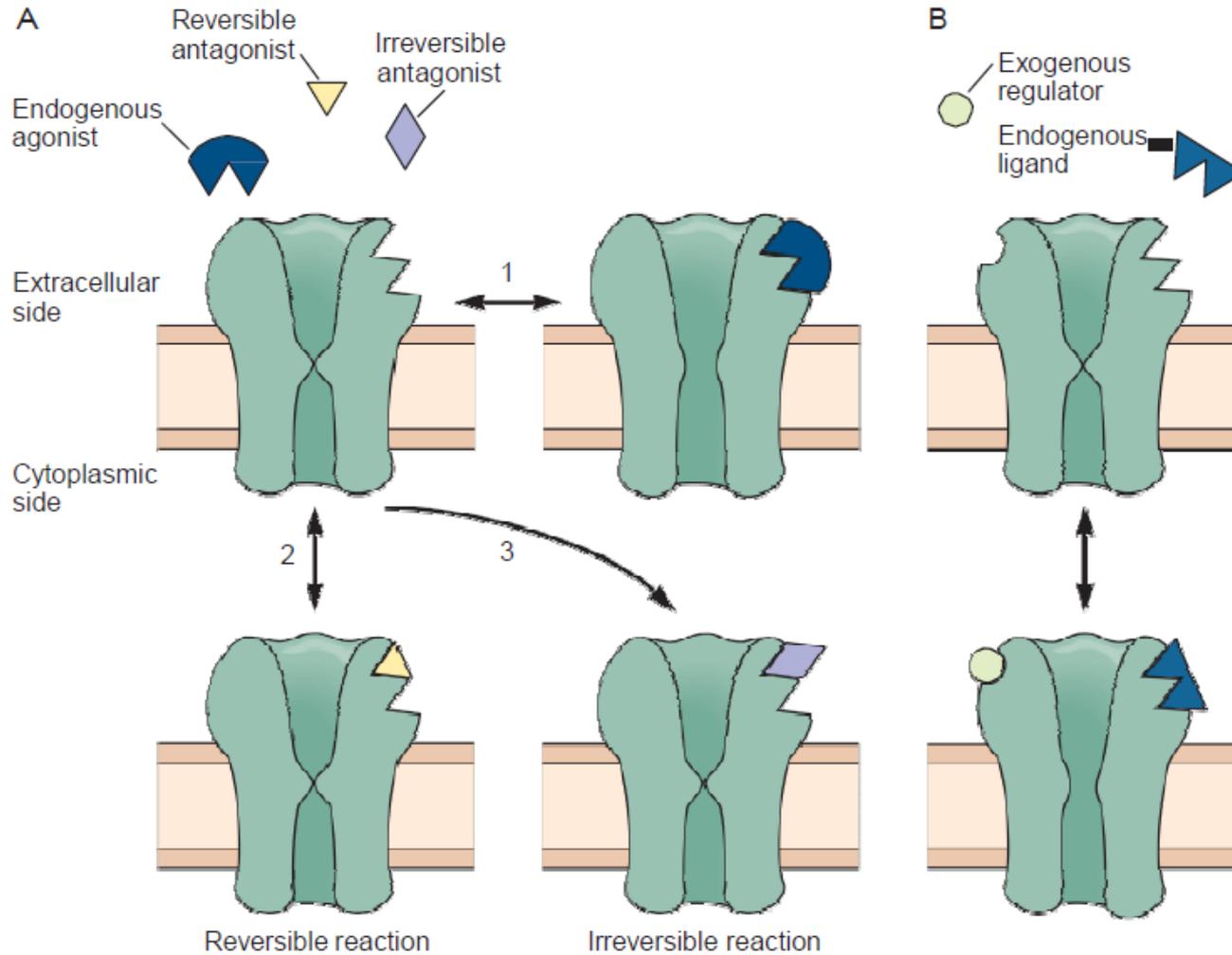
Inactivation mechanisms of voltage-gated ion channels



Single channel activity of a voltage-gated Na⁺ channel

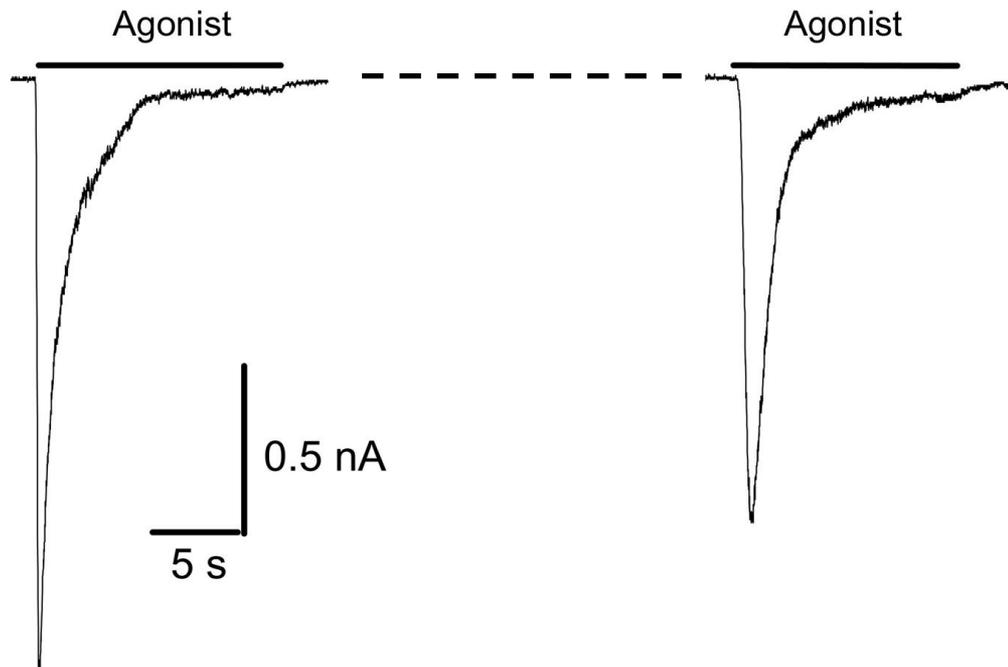


Exogenous ligands affect gating of ion channel

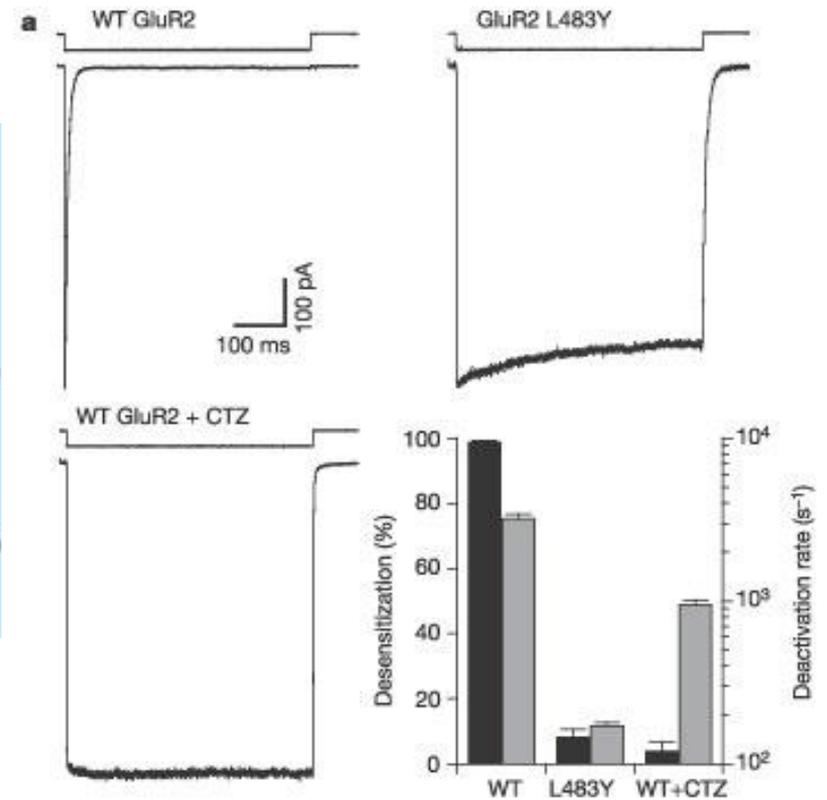
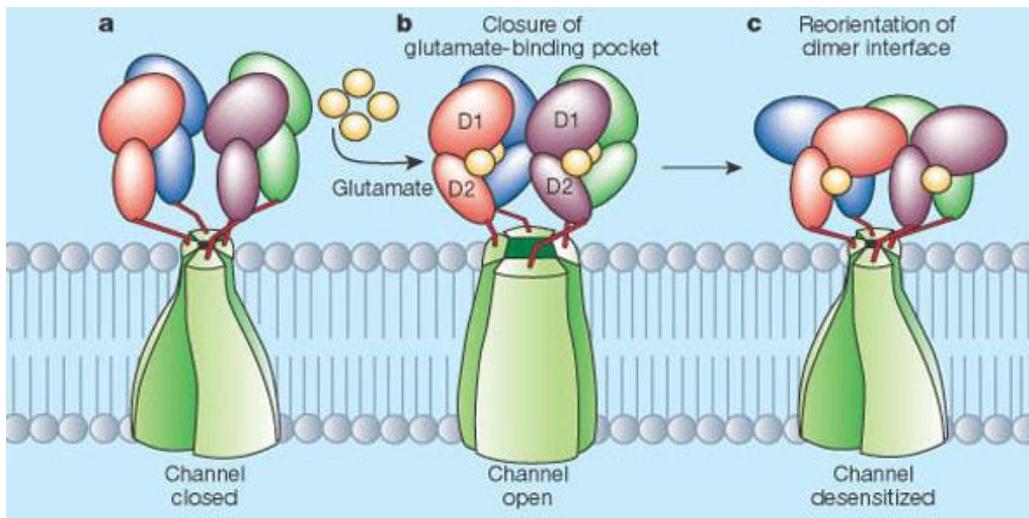


Desensitization

- loss of responsiveness to a continuing dose of agonist
- refractory state
- mechanisms
 - intrinsic property of the interaction between ligand and channel
 - phosphorylation by protein kinase



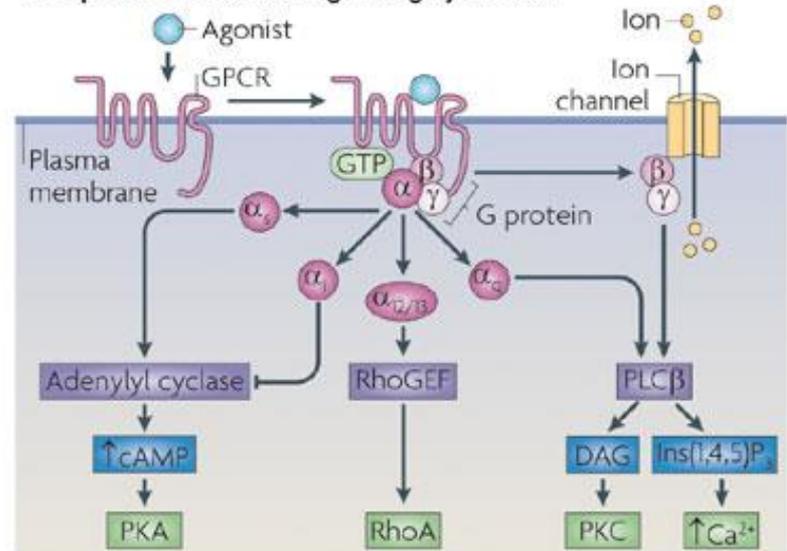
AMPA receptor desensitization



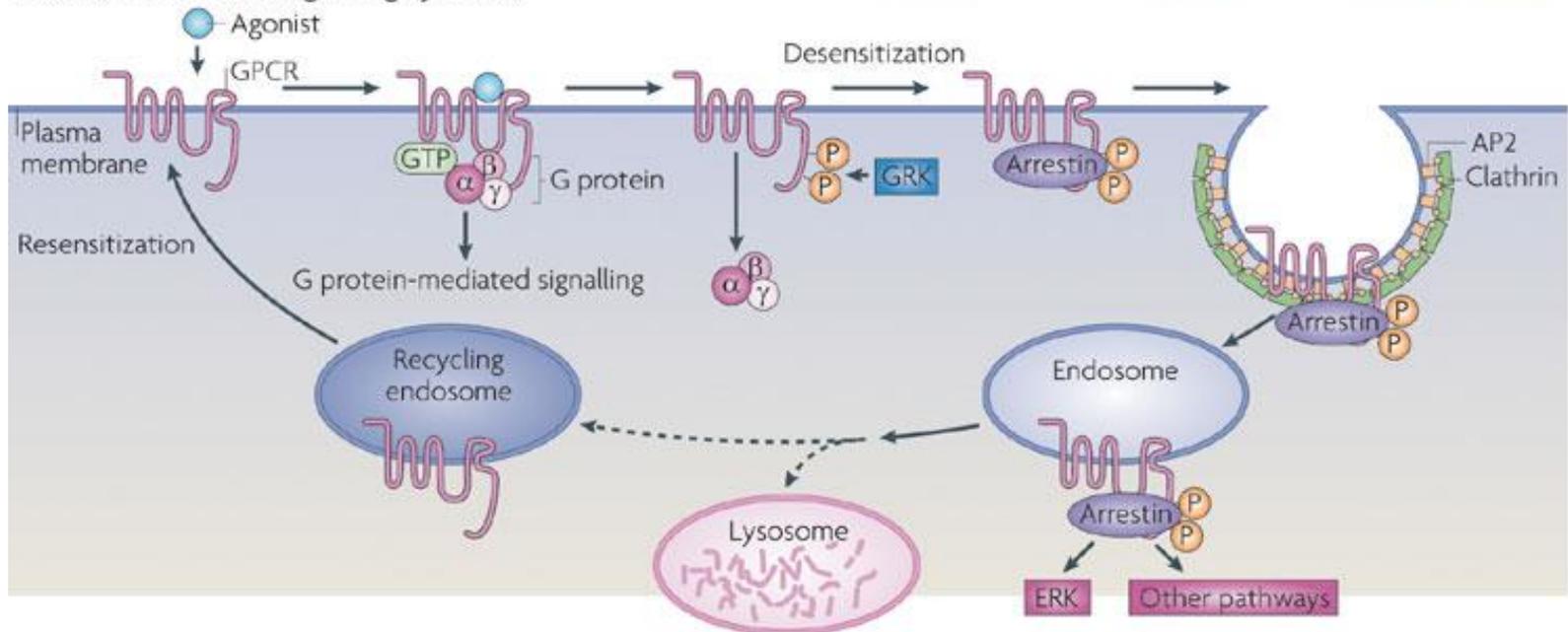
-intrinsic property of the interaction between ligand and channel

GPCR signaling and desensitization

a G protein-mediated signalling by GPCRs



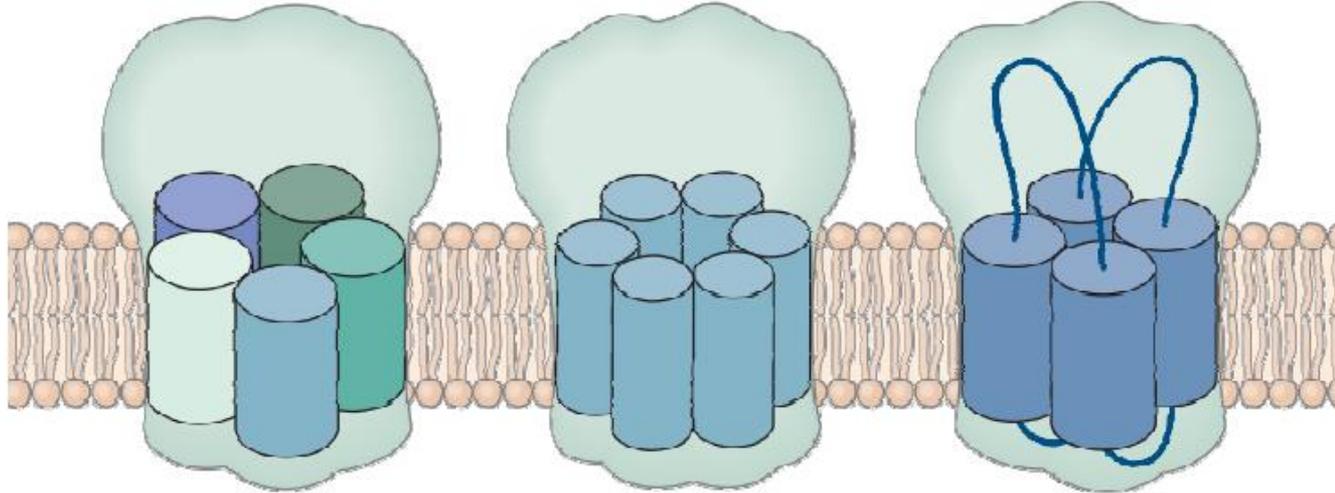
b Arrestin-mediated signalling by GPCRs



-phosphorylation by protein kinase

Ion channels are integral membrane proteins composed of several subunits

A

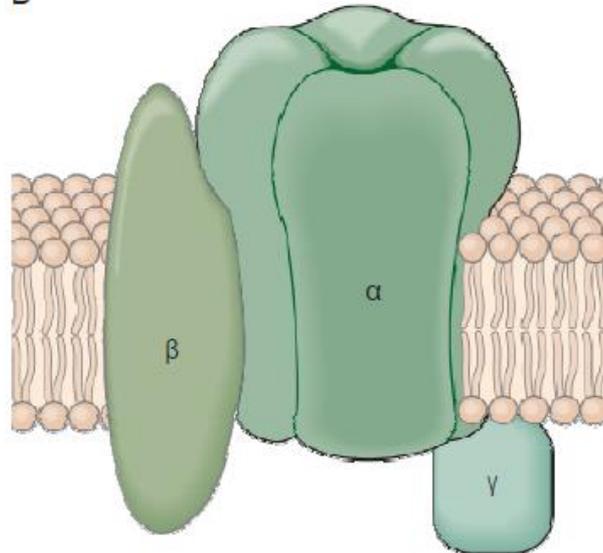


Hetero-oligomer

Homo-oligomer

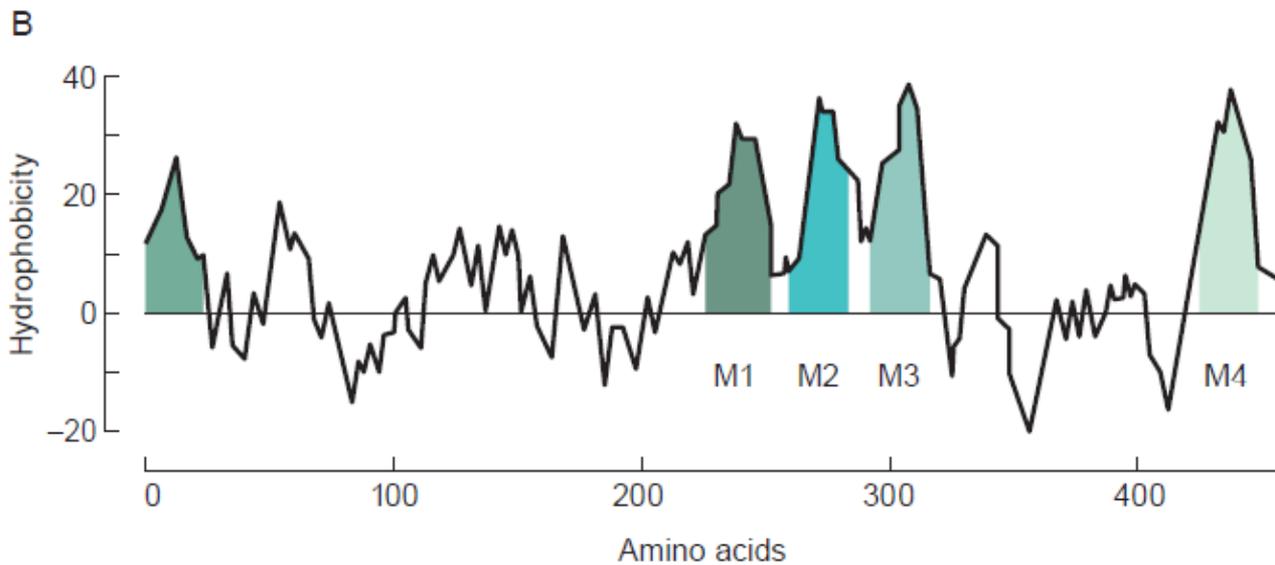
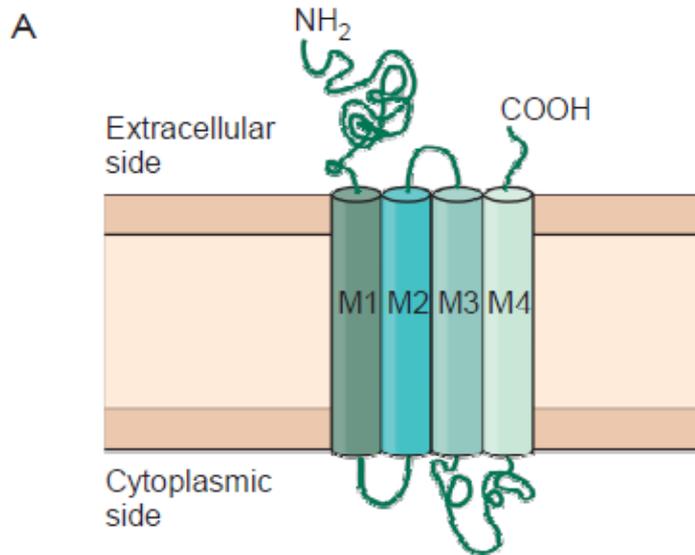
Single peptide chain
with repeating motifs

B



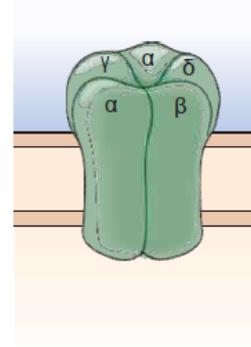
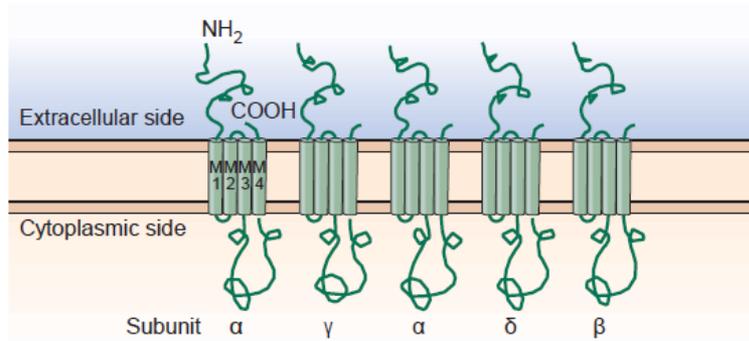
Pore forming subunit (α)
+ auxiliary subunits (β or γ)

The secondary structure of membrane-spanning proteins

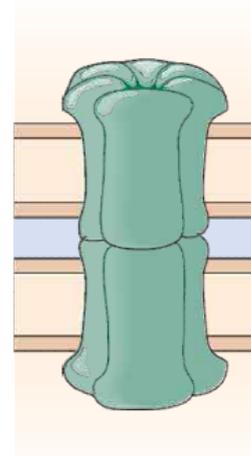
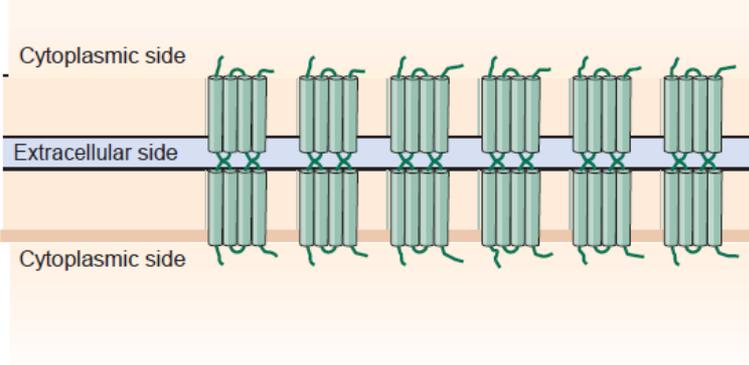


Three superfamilies of ion channel

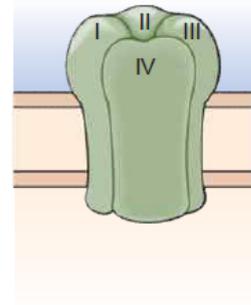
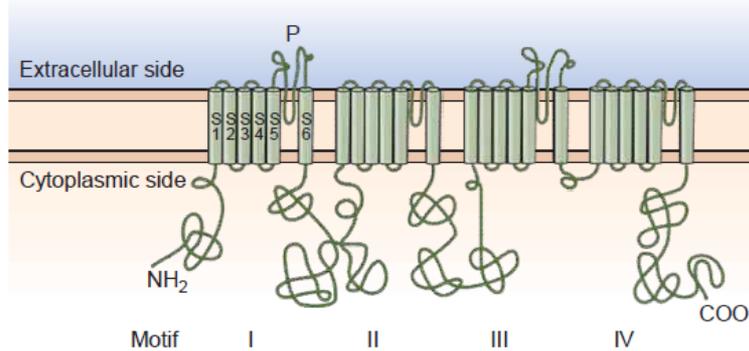
A Ligand-gated channel (ACh receptor)



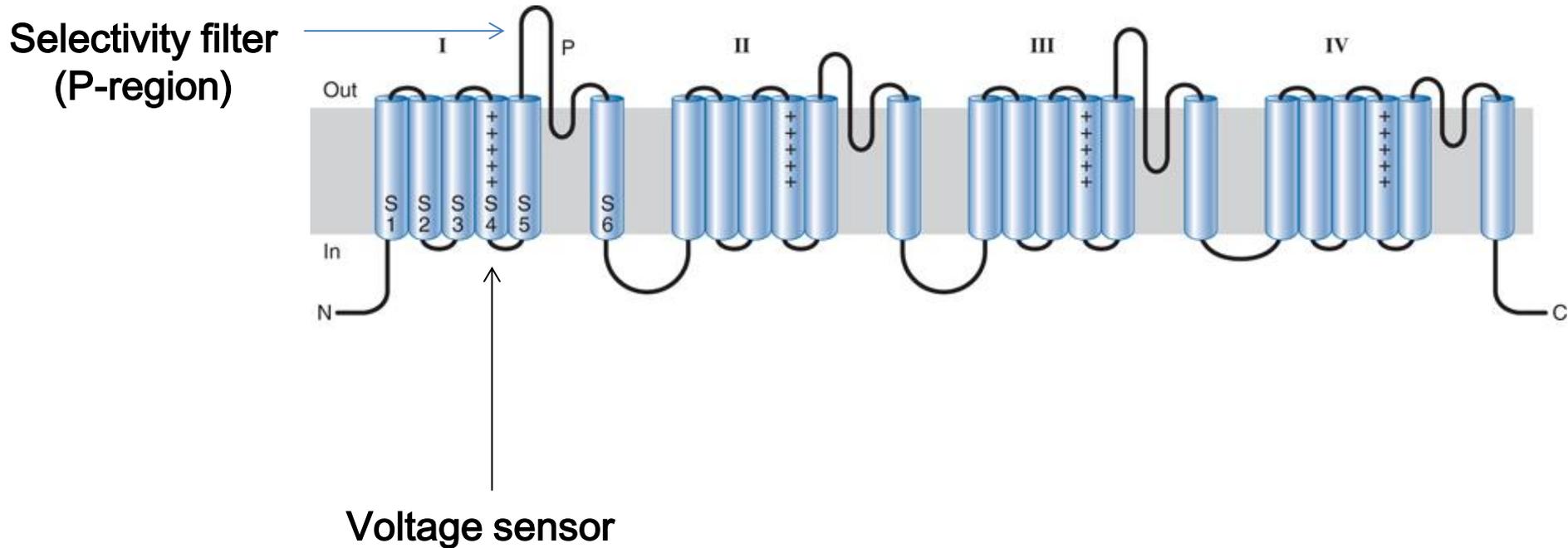
B Gap-junction channel



C Voltage-gated channel (Na⁺ channel)



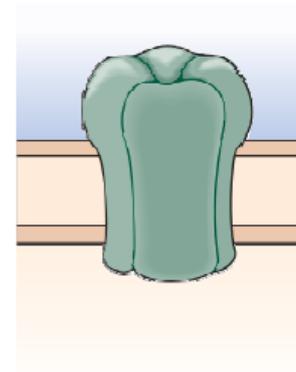
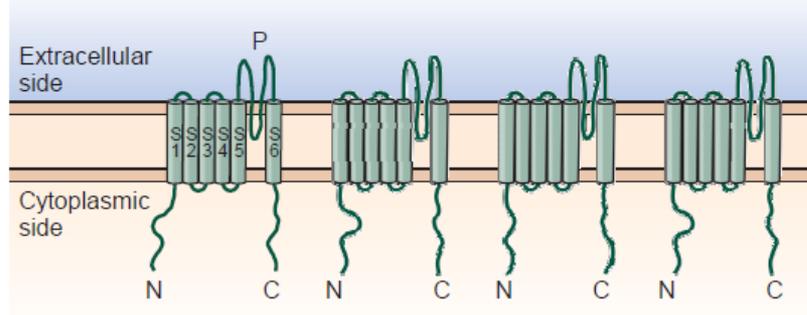
Voltage-gated Na⁺ channel (α subunit)



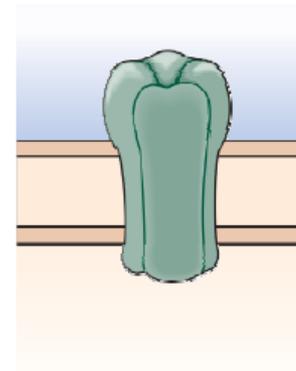
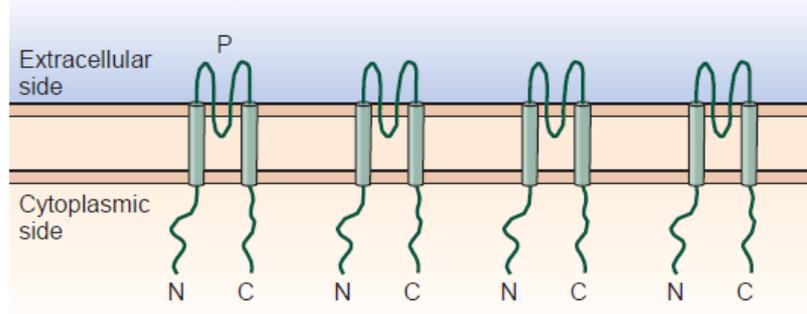
-positive amino acids located at every fourth position

Four related families of ion channels with P-regions

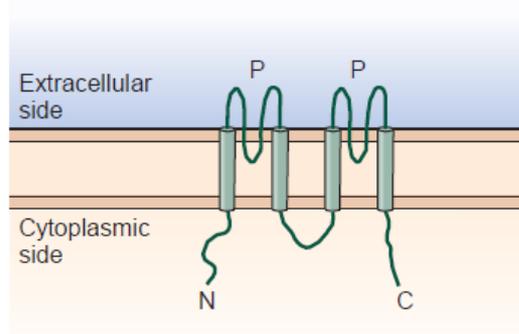
A Voltage-gated K⁺ channel



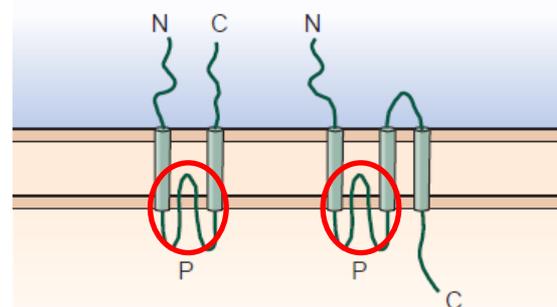
B Inward-rectifying K⁺ channel



C K⁺ channel subunit with two P regions



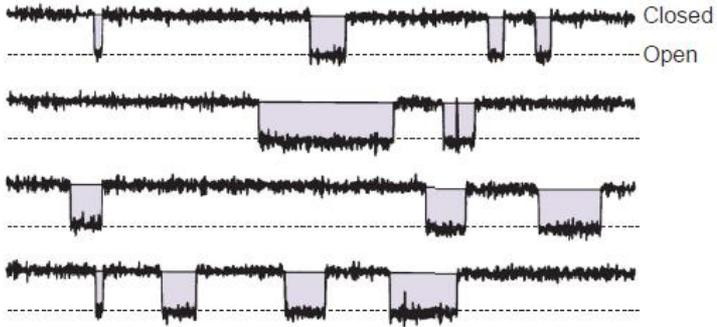
D Glutamate-gated channel subunits



Bacteria Higher organism

nACh receptor channel

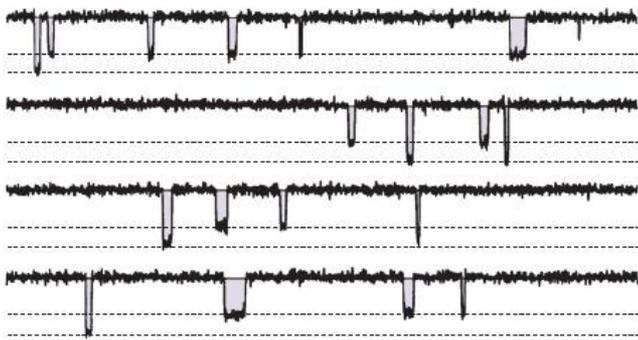
1.1 days



Subunit composition

$2 \alpha, \beta, \gamma, \delta$

2.4 days

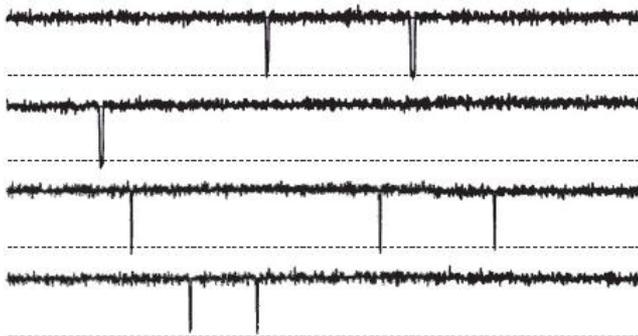


$2 \alpha, \beta, \gamma, \delta$

+

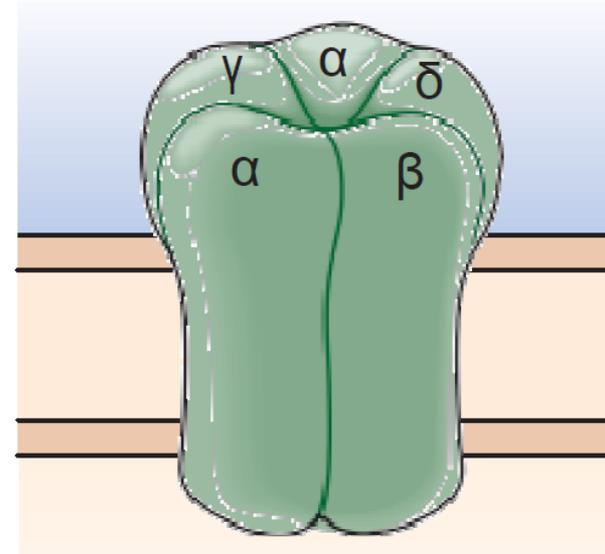
$2 \alpha, \beta, \gamma, \epsilon$

48 days



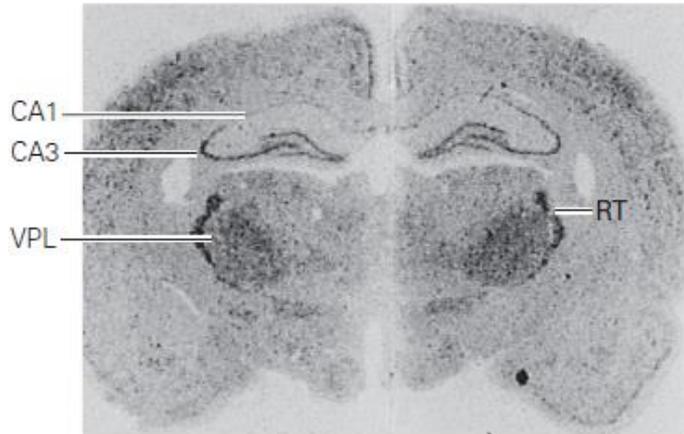
$2 \alpha, \beta, \gamma, \epsilon$

5 pA
10 ms

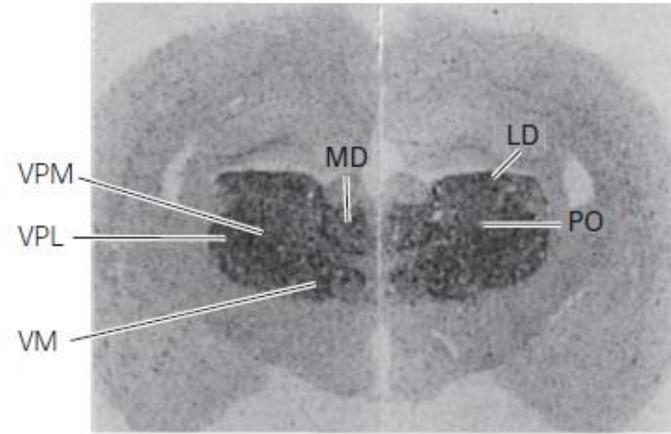


Brain expression patterns of Kv3 channel variants

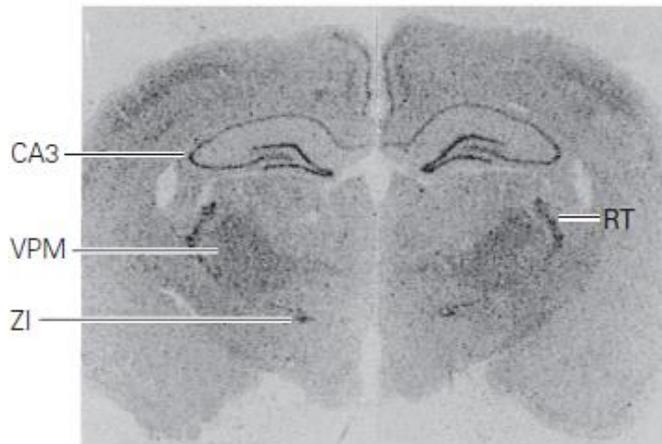
A Kv3.1



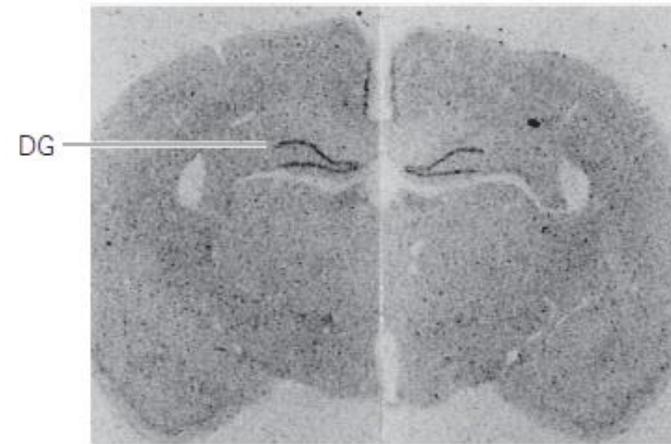
B Kv3.2



C Kv3.3

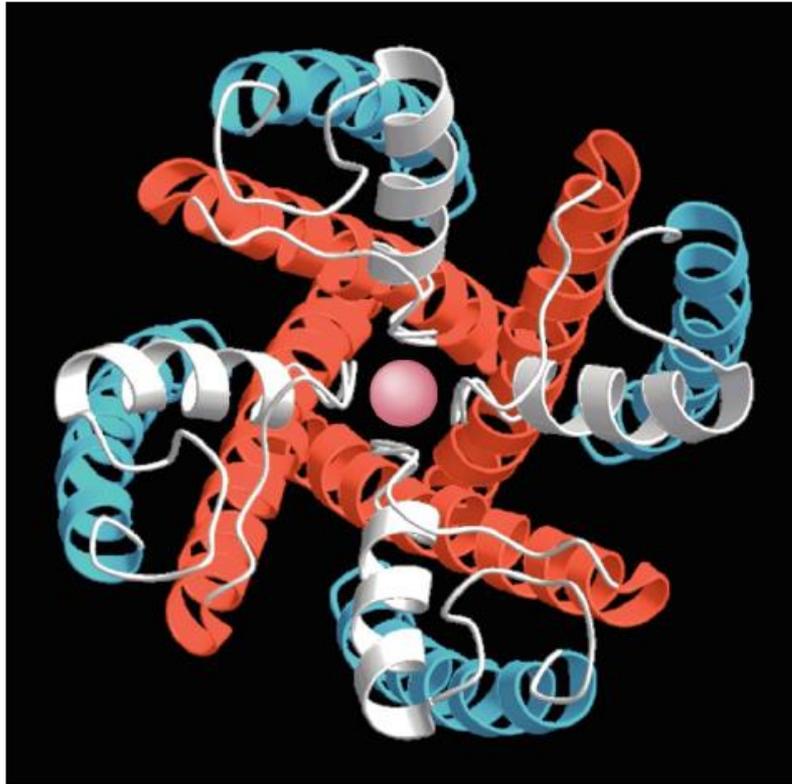


D Kv3.4

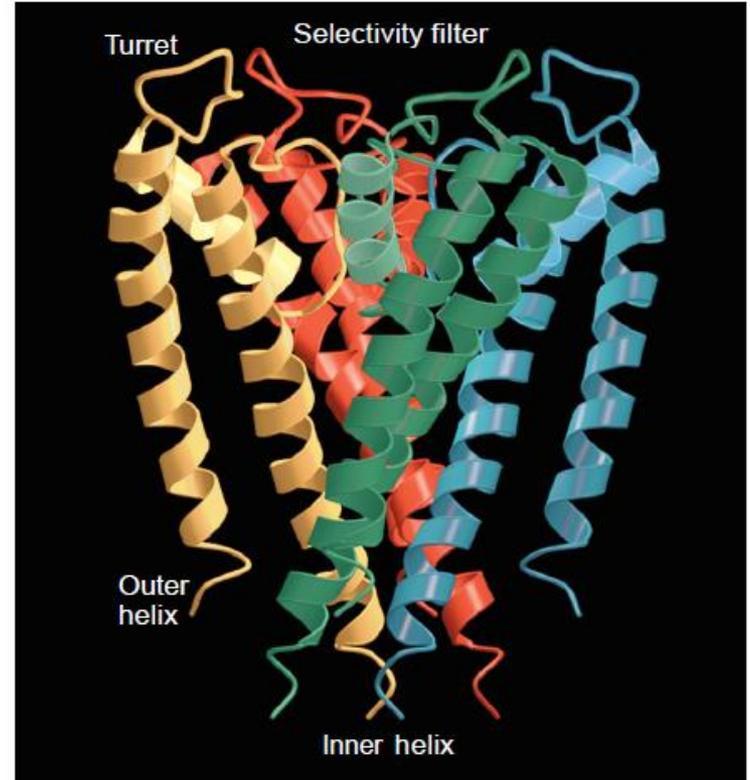


The X-ray crystal structure of a bacterial potassium (KcsA K⁺) channel

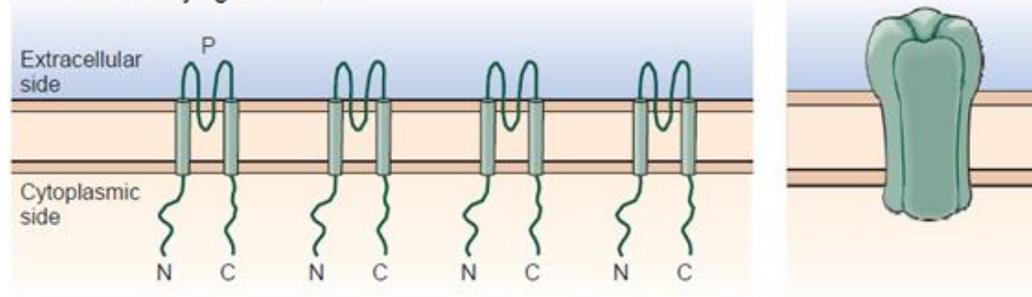
A Looking down the channel



B Cross section

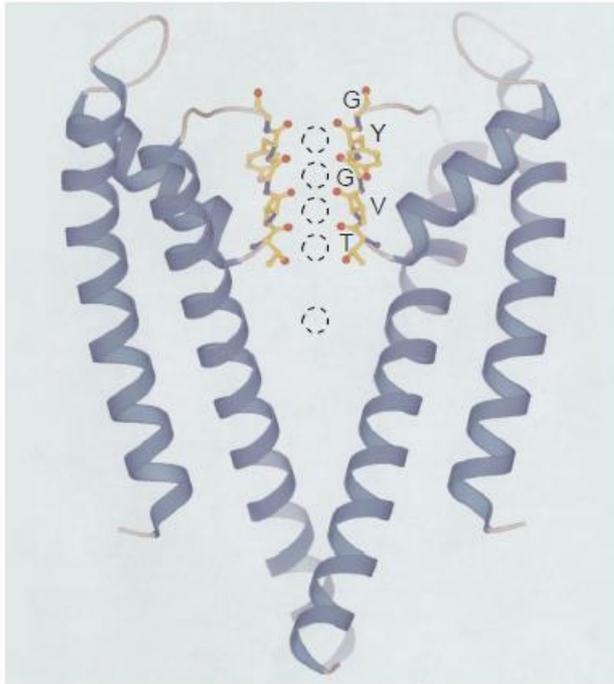


B Inward-rectifying K⁺ channel

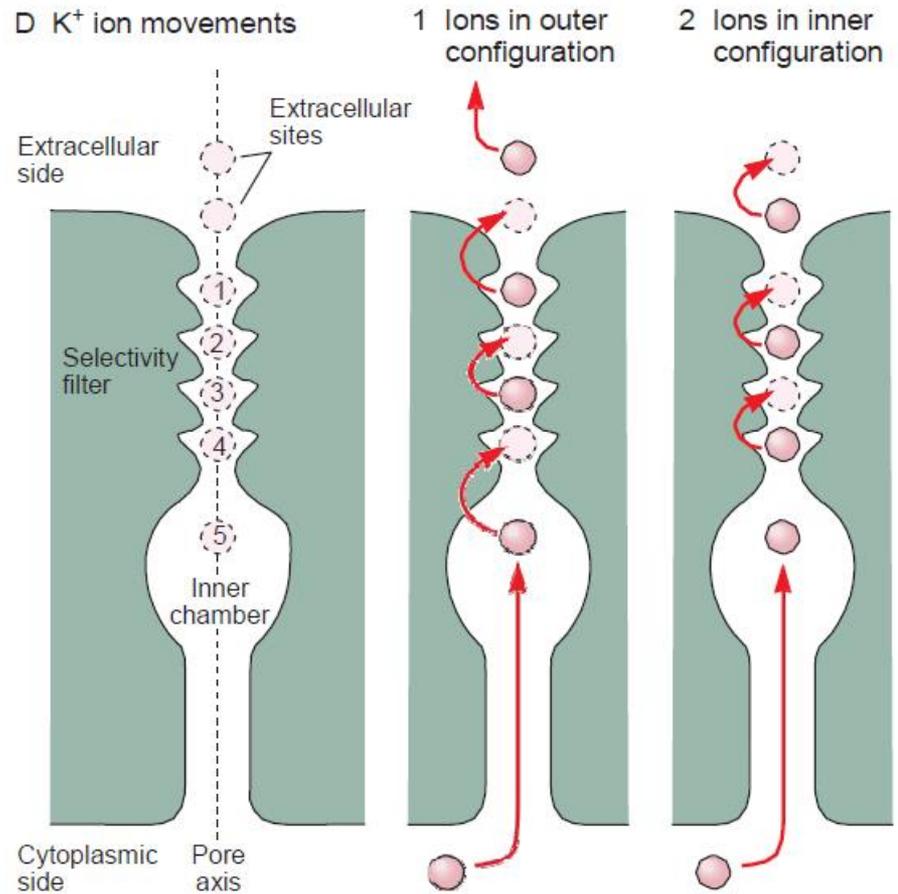


Ion permeation through the KcsA K⁺ channel

C K⁺ ion binding sites

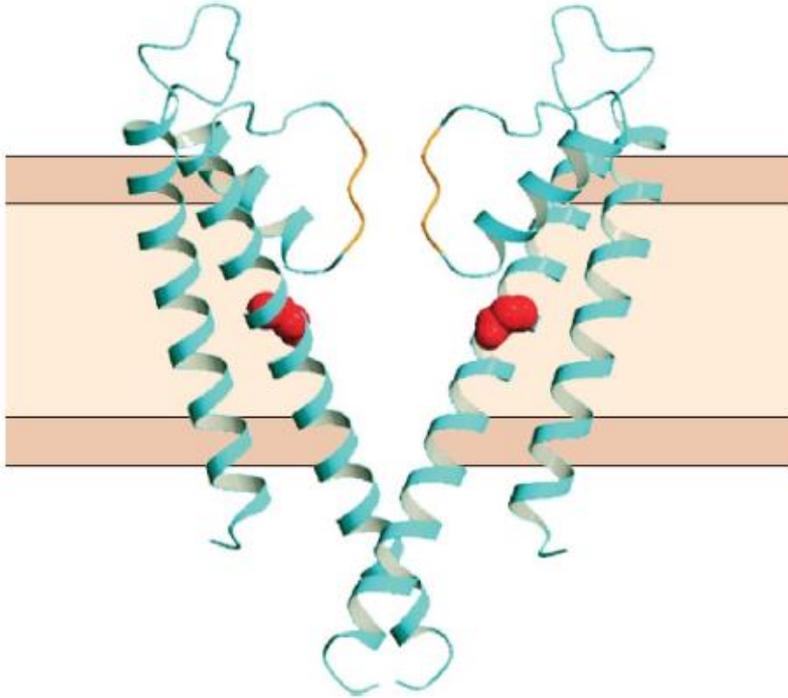


D K⁺ ion movements

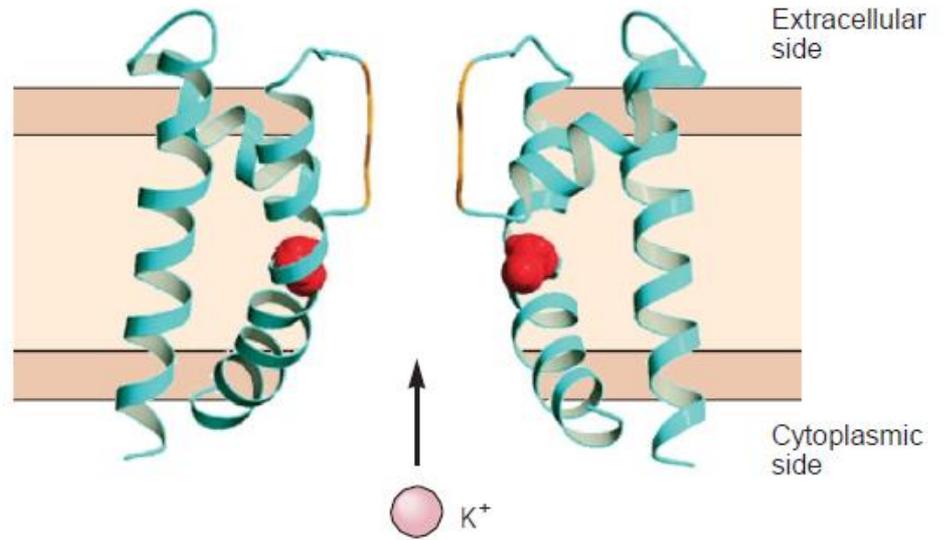


Gating of bacterial potassium (MthK) channels

A Closed state



B Open state



Double-barrel channels: CIC family chloride channels and transporters

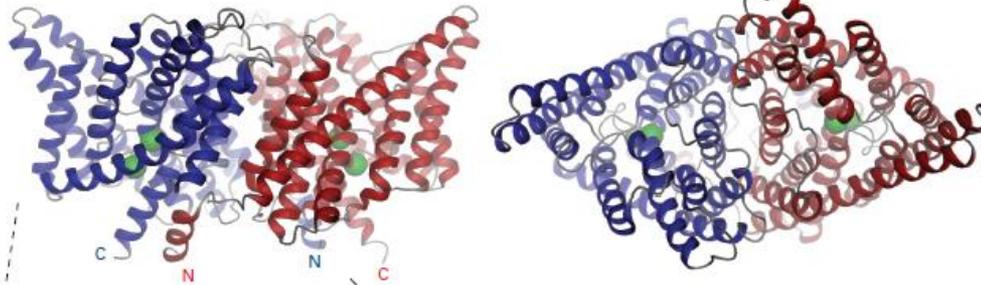
A 척추동물의 단일 Cl⁻ 채널을 통한 전류



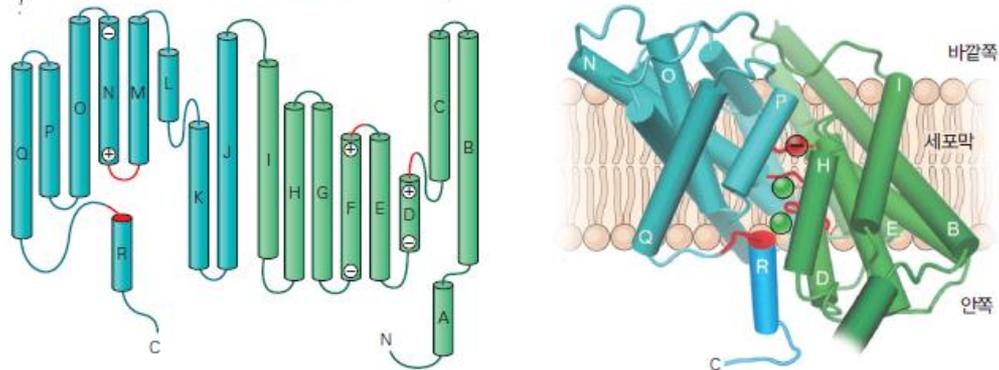
B CIC 채널의 모델



C 대장균 CIC 수송체

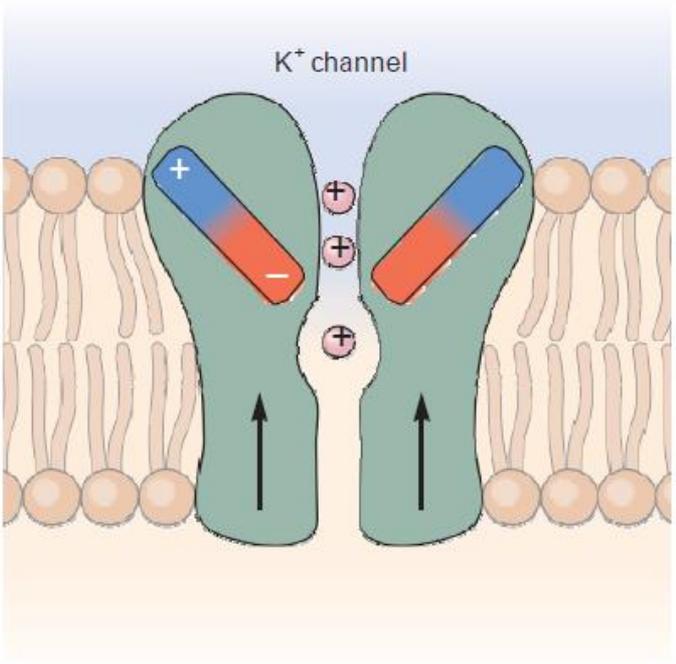


D 대장균 CIC 수송체의 단일 하위단위

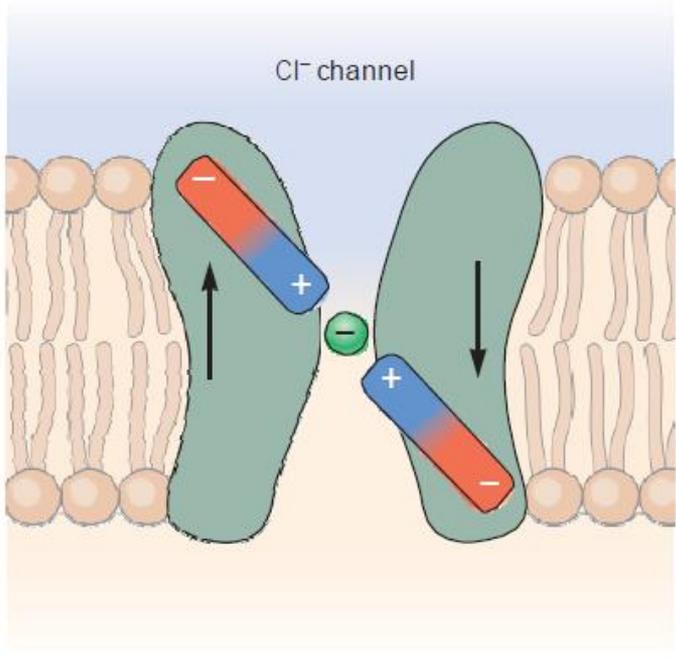


Comparison of general architectures of potassium and chloride channels

A Parallel (barrel stave)



B Anti-parallel



The functional difference between ion channels and pumps

