

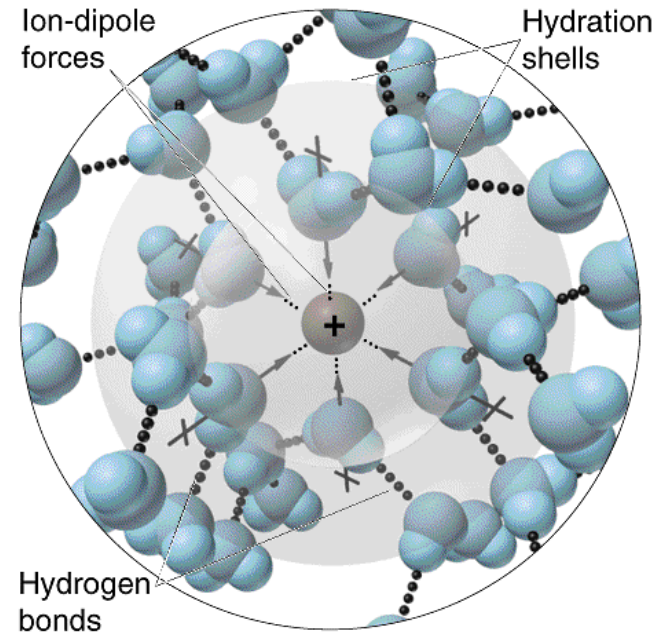
Hydrophobic barrier

Born-Formula

✚ There are about 7 water molecules in the first hydration shell of potassium ion.

✚ Each water molecule stabilizes the ion by approximately 24 kT.

$$\Delta E = \frac{1}{2} \frac{Q^2}{r} \left(\frac{1}{\epsilon_m} - \frac{1}{\epsilon_w} \right)$$

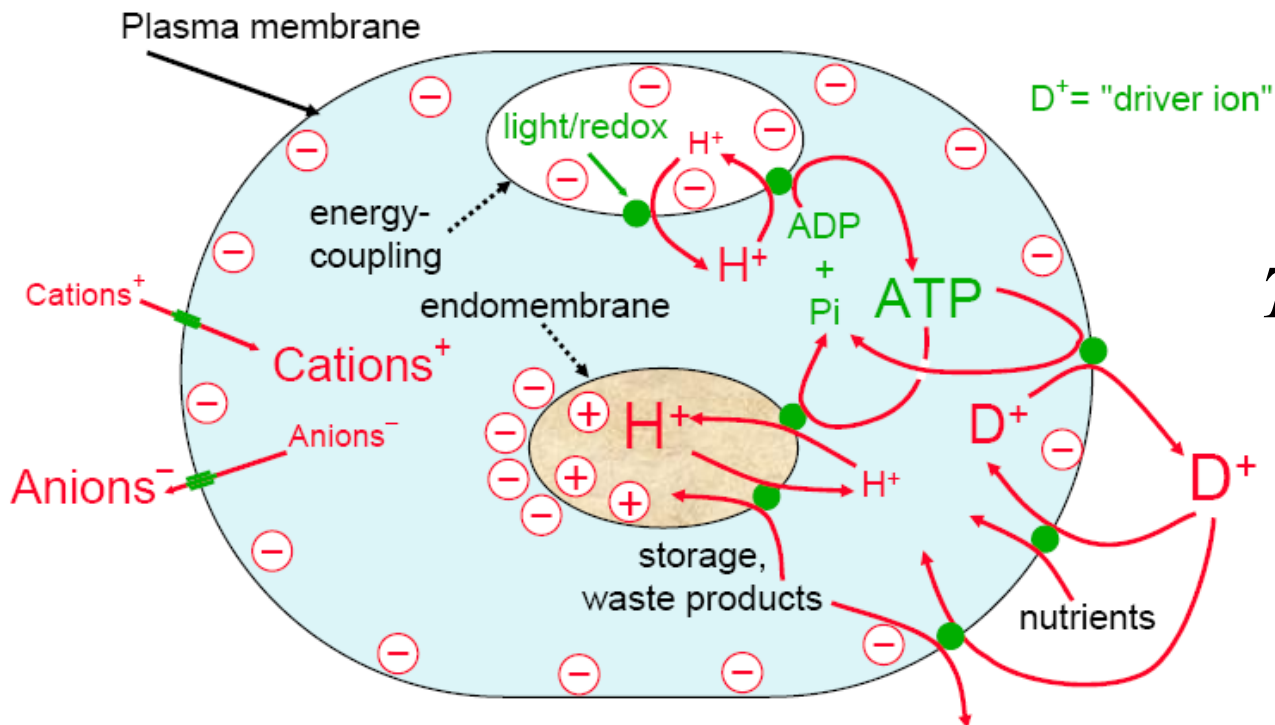


Completion and analysis of various genomes revealed that about 10% of **all** proteins function in **transport** (in E.coli – 427 transporters)

In eucaryotic cells, 2/3 of cellular energy at rest is used to transport ions (H^+ , K^+ , Na^+ , Ca^{++})

About 200 families of transporters are recognized

The largest family: ABC-transporters



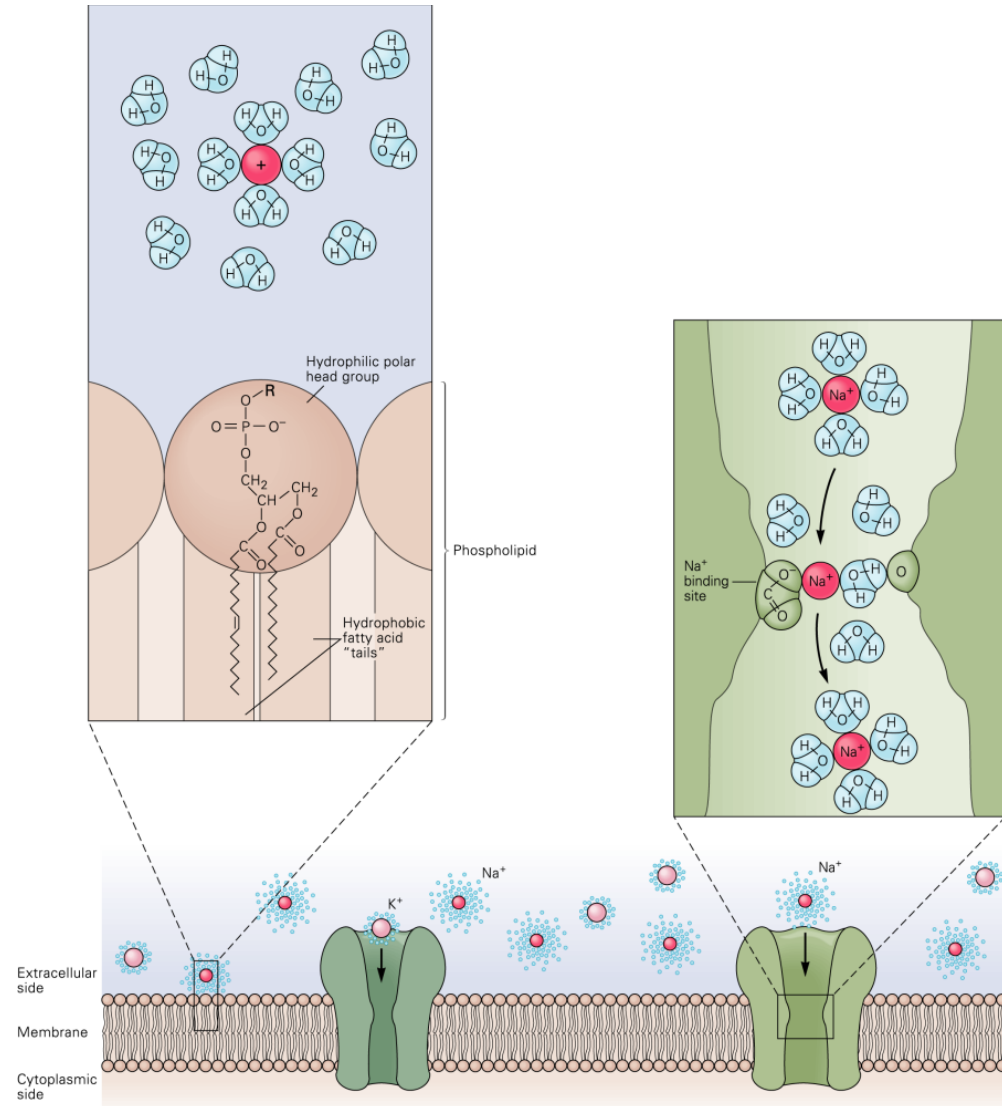
Transport Processes in an Idealized Eukaryotic Cell

Ion channels are enzymes that catalyze the flow of ions across cell membranes causing picoamp current.

For every ion that crosses the membrane through the lipid bilayer, 10^{13} do so by traversing an ion channel.

The downhill diffusion of ions through biological membranes occurs through ion channels;

- an ion channel has a resistance of $\sim 10^{10} \Omega$,
- a patch of lipid bilayer of similar cross section has a resistance of $\sim 10^{23} \Omega$.



Physical Characteristics of Ion Channels

Natural Nanodevices

Ion channels have **Selectivity**.

K channel selects K^+ over Na^+ by $\sim 10^4$.

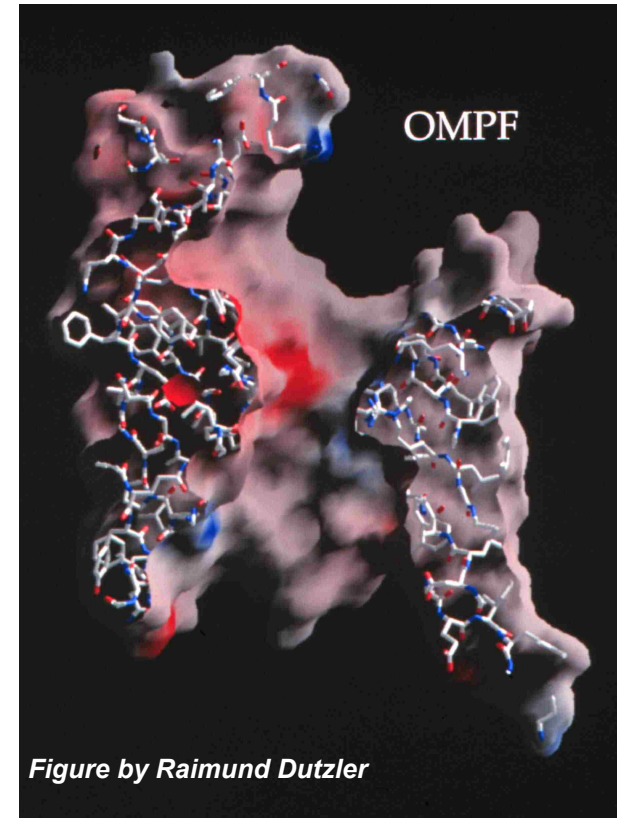
Ion channels **Gate/Switch** in response to pH, voltage, chemical species and mechanical force from conducting to nonconducting state.

Ion channels have **VERY Large Charge Densities** critical to I-V characteristics and selectivity (and gating?)

Ion channels allow **Mutations** that modify conductance, selectivity, and function.

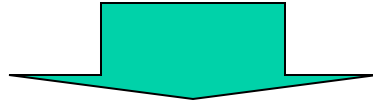
Ion channels are **Device Elements** that self-assemble into perfectly reproducible arrays.

Ion channels form **Templates** for design of bio-devices and biosensors.

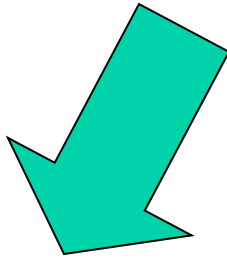


←→
~30Å

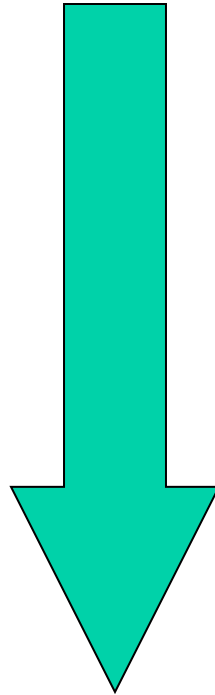
Ion channel



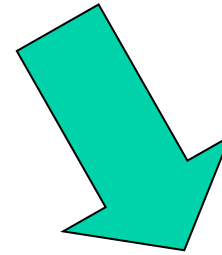
ion-permeation pathway through the membrane



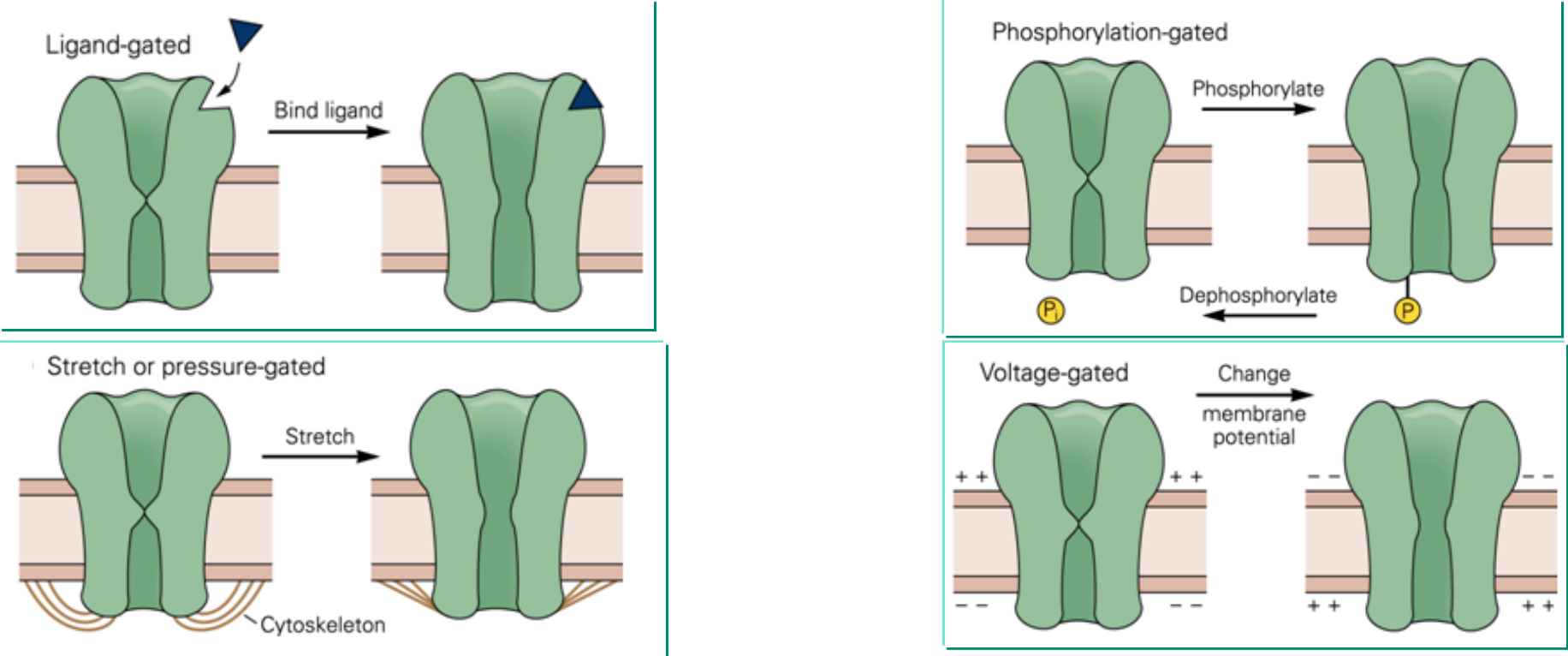
Selectivity filter
(narrowest constriction
in the 'open'
conformation)



Gate
(narrowest constriction
in the 'closed' conformation)



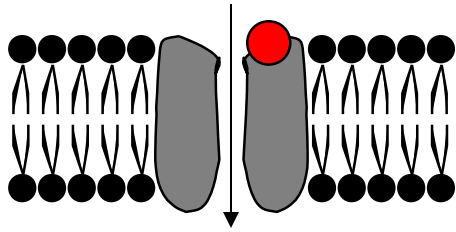
***Elements that
control the gate***
(ligand-binding sites,
voltage-sensor, pH-sensor,
temperature-
sensor, mechanical-
deformation sensor)



Depending on the type of the channel, this gating process may be driven by:

- ligand binding (ligand-gated channels)
- changes in electrical potential across cell membrane (voltage-gated channels)
- mechanical forces acting on cellular components (mechanosensitive channels)

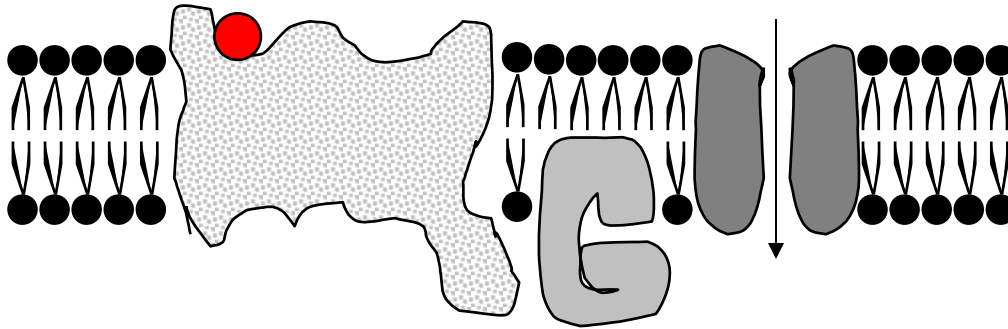
Receptor with integral ion channel



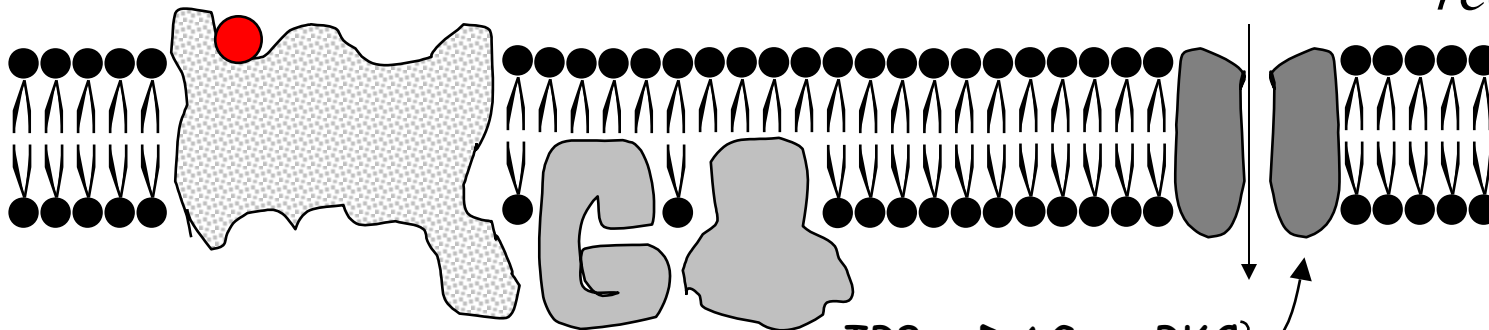
(Nicotinic ACh receptors)

Ligand-gated ion channels

Receptor linked to ion channel via g-protein



Receptor linked via g-protein & 2nd messenger



IP3 + DAG $\Rightarrow$ PKC
or cAMP $\Rightarrow$ PKA

K⁺ channel

*(GABA_B
receptors)*

Methods for Studying Ion Channels

Biochemistry

- agonist, antagonist or drug binding
- isolation and purification
- reconstitution
- radioactive ion flux

Structural biology

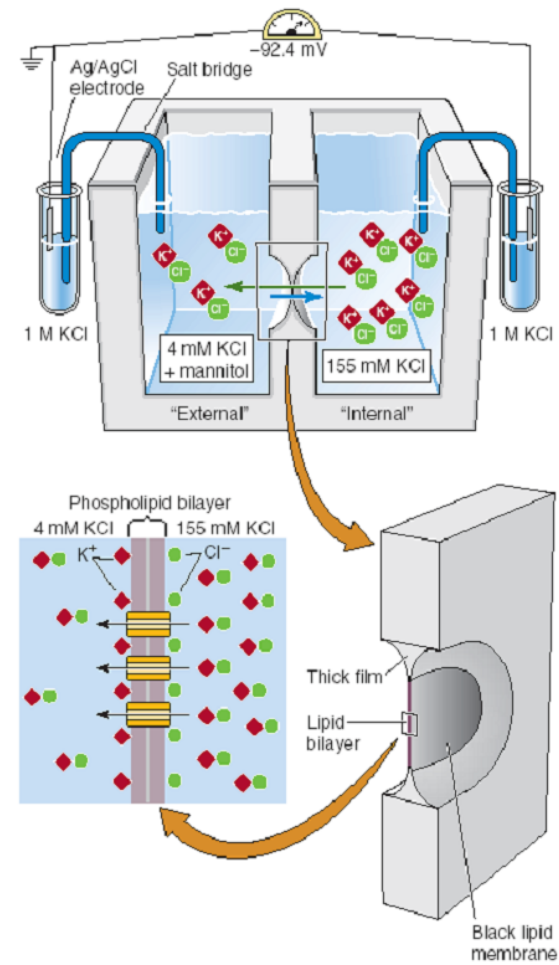
- microscopy, crystallography, NMR, ...

Molecular biology

- sequencing, cloning, mutagenesis



Chris Miller
Brandeis University



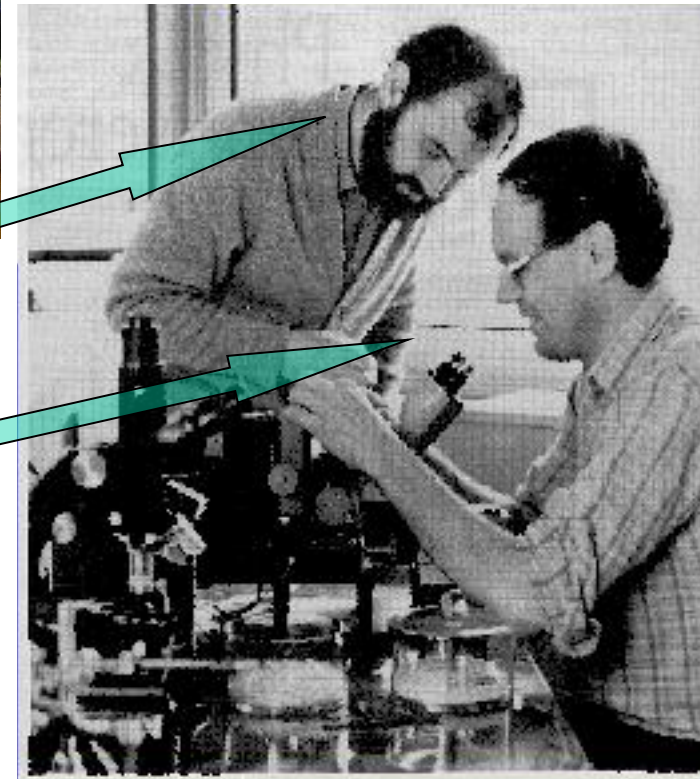
Patch clamp technique

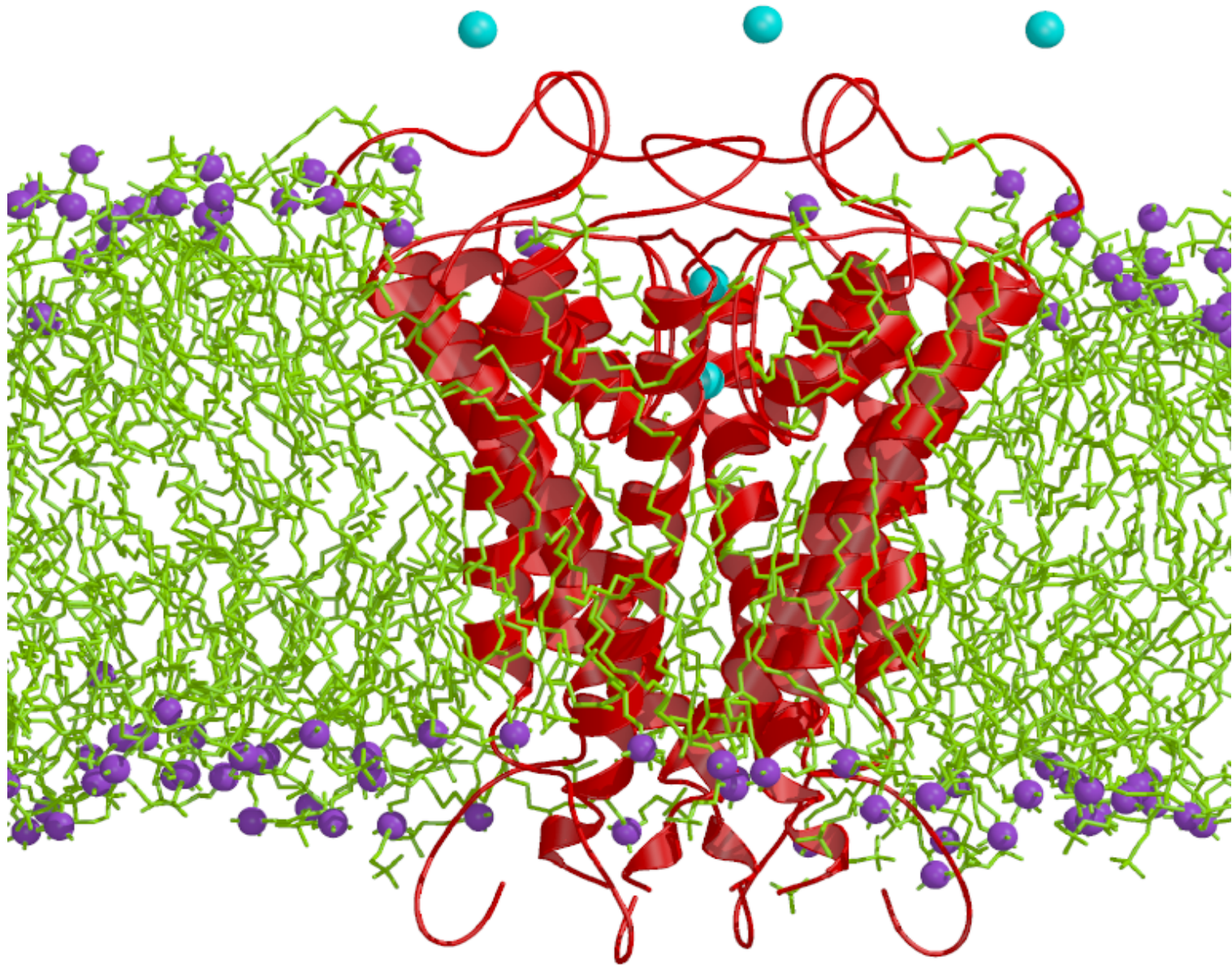


Erwin Neher

Bert Sakmann

*Germany
(1991 Nobel Laureates)*



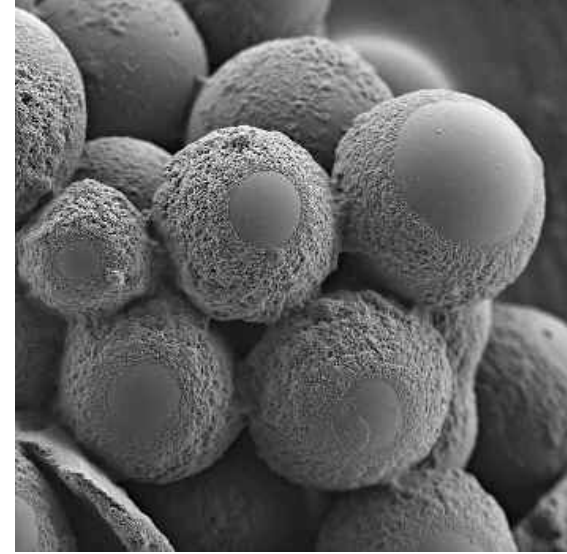


Potassium channel

Potassium channel KcsA

The 3D-structure of KcsA from Streptomyces lividans was the first X-ray structure of an ion channel.

For the Atomic Basis of Selective Ion Conduction in Potassium Channels



Streptomyces lividans



Advanced information on the Nobel Prize in Chemistry, 8 October 2003



KUNGL.
VETENSKAPSAKADEMIEN
THE ROYAL SWEDISH ACADEMY OF SCIENCES



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Membrane channels

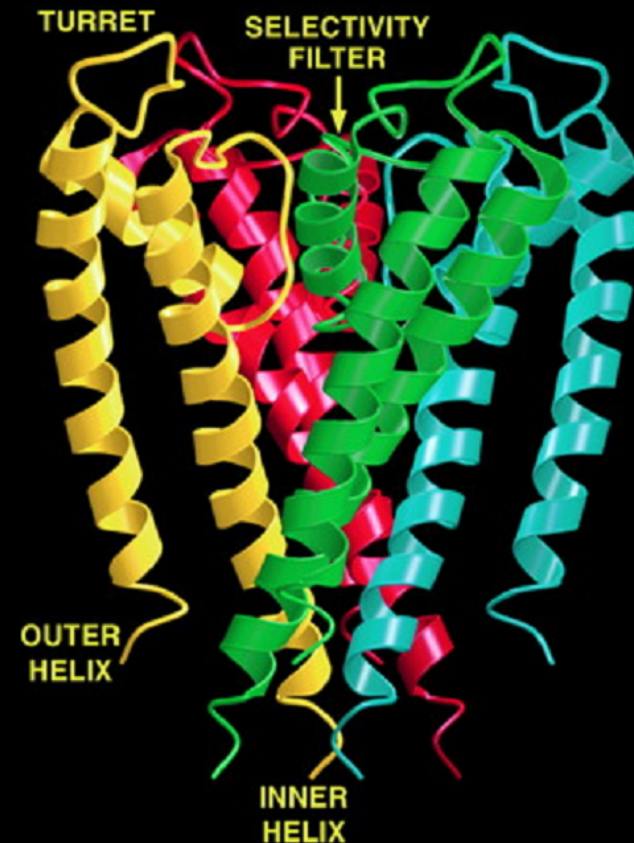
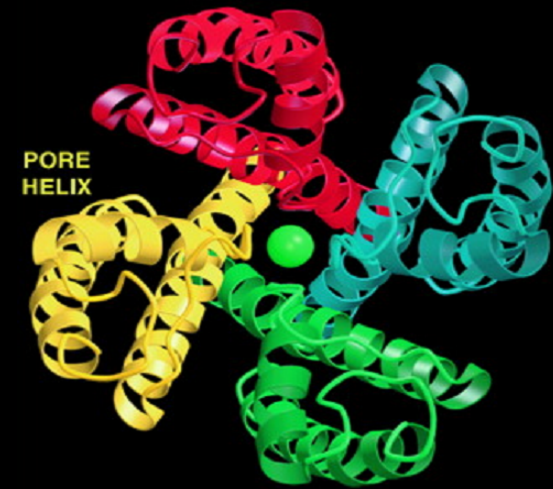
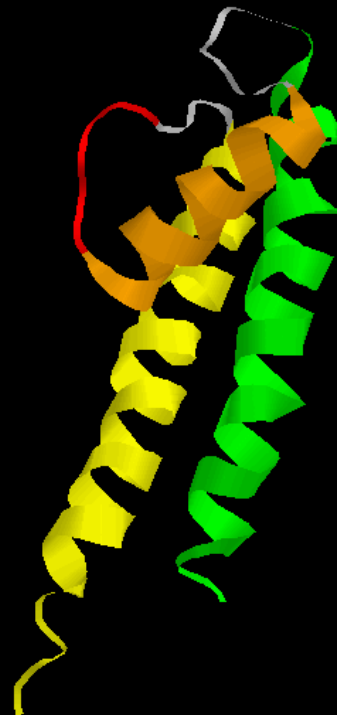
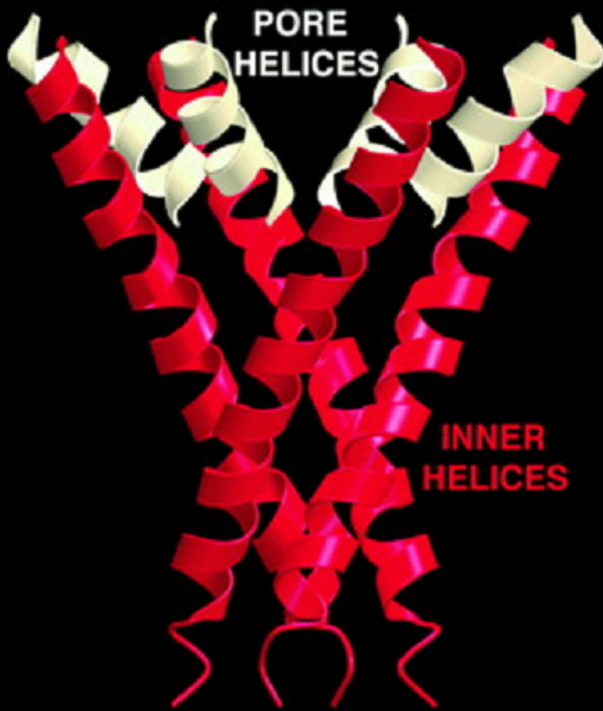
Introduction

The Nobel Prize in Chemistry for 2003 is shared between two scientists who have made fundamental discoveries concerning how water and ions move through cell membranes. Peter Agre discovered and characterized the first water channel protein and Roderick MacKinnon has elucidated the structural and mechanistic basis for ion channel function.

The Potassium Channel (KcsA)

It is a **tetramer**.

Each monomer is 96 amino acids long and consist of 3 helices.

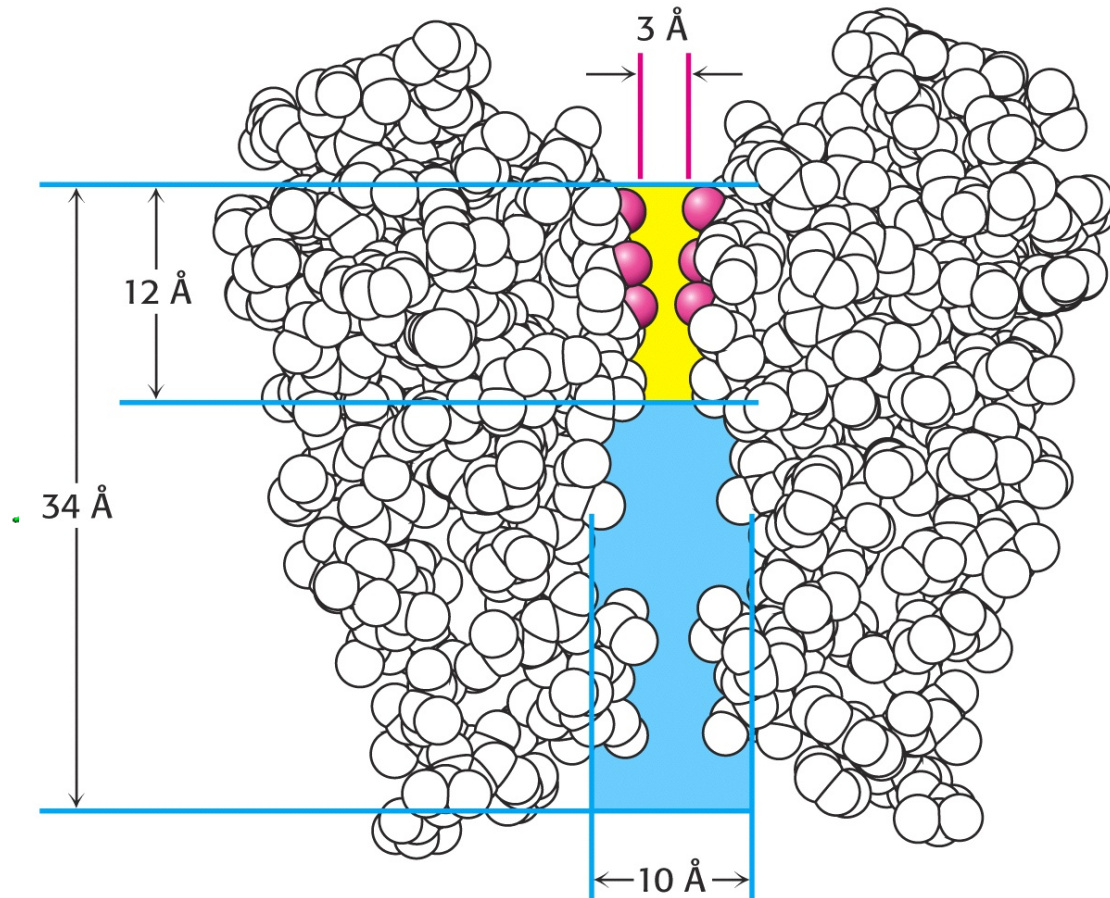


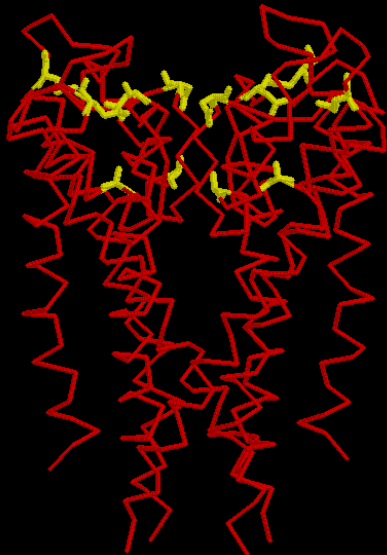
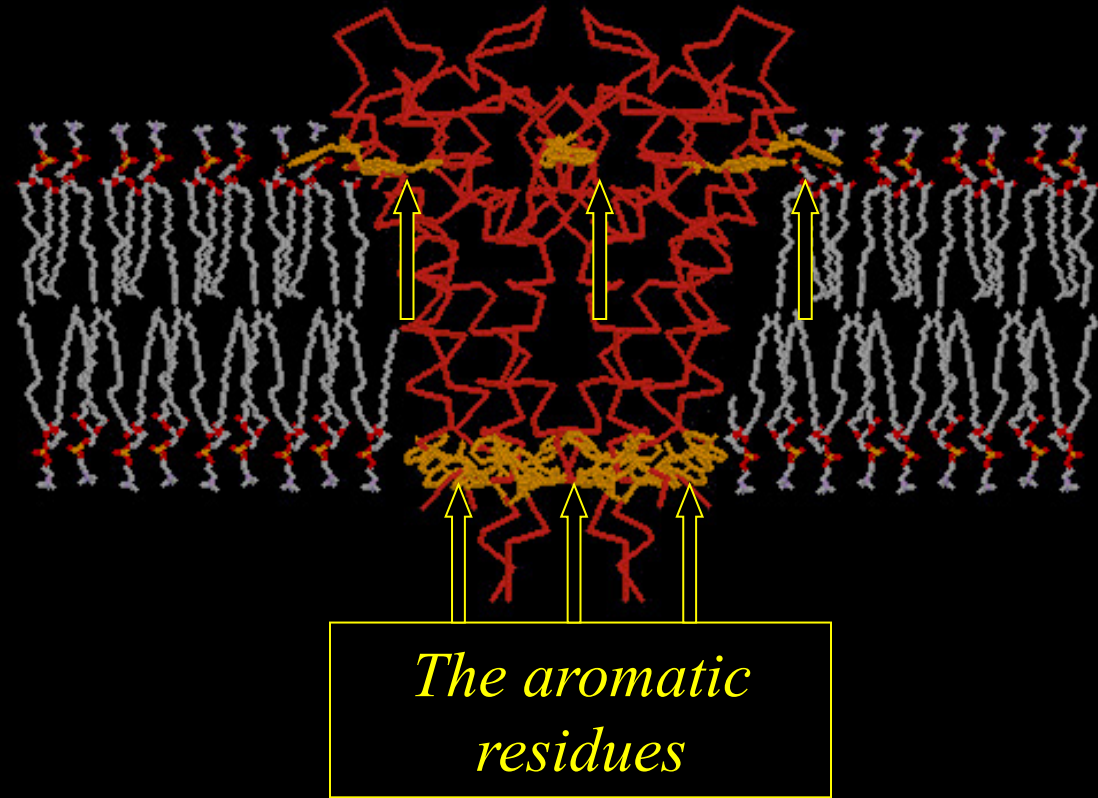
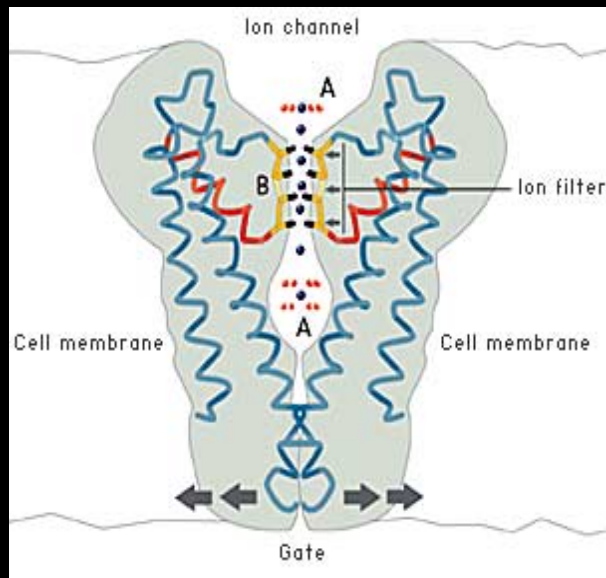
Potassium channel

Potassium channels are a class of ion channels, managing to distinguish between K^+ and Na^+ ions while maintaining a very high throughput of K^+ ions when gated.

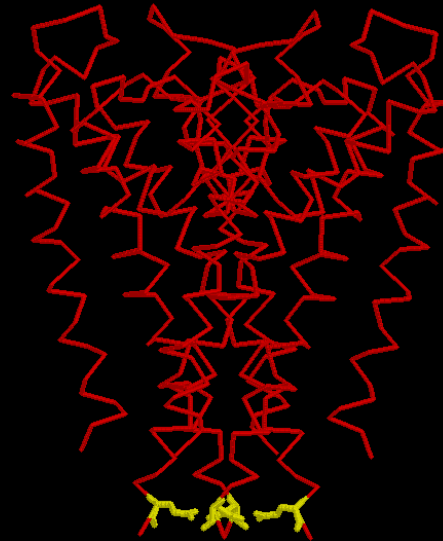
It has a diffusion rate of 10^8 ions per second.

One K^+ ion is dehydrated, transferred, and rehydrated every 10 ns.



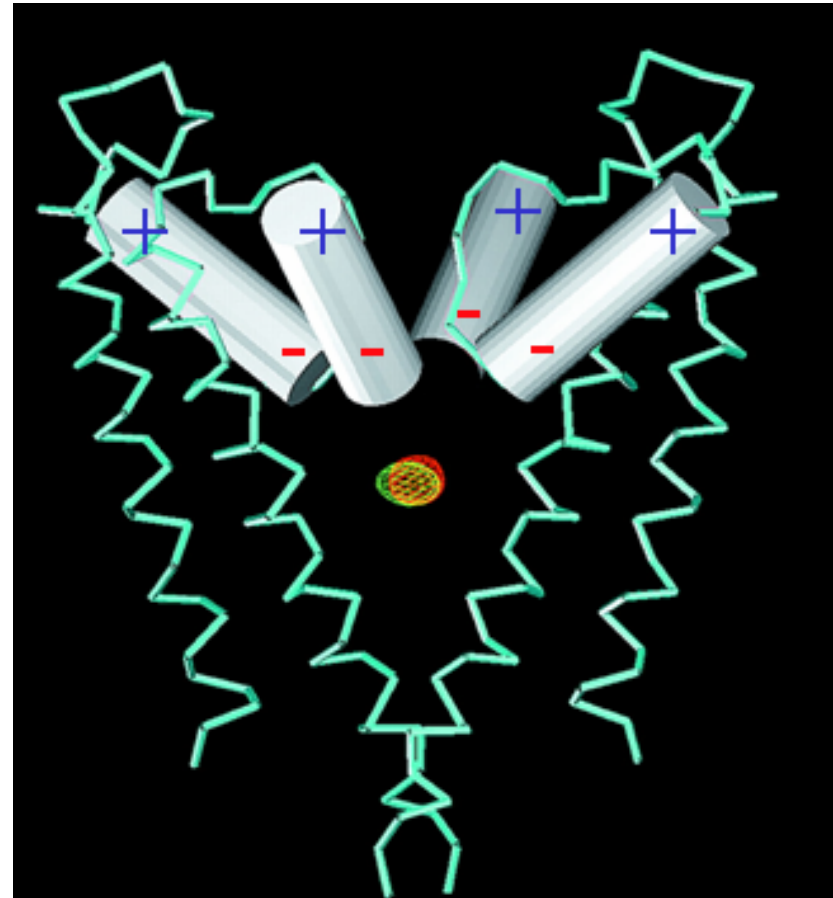
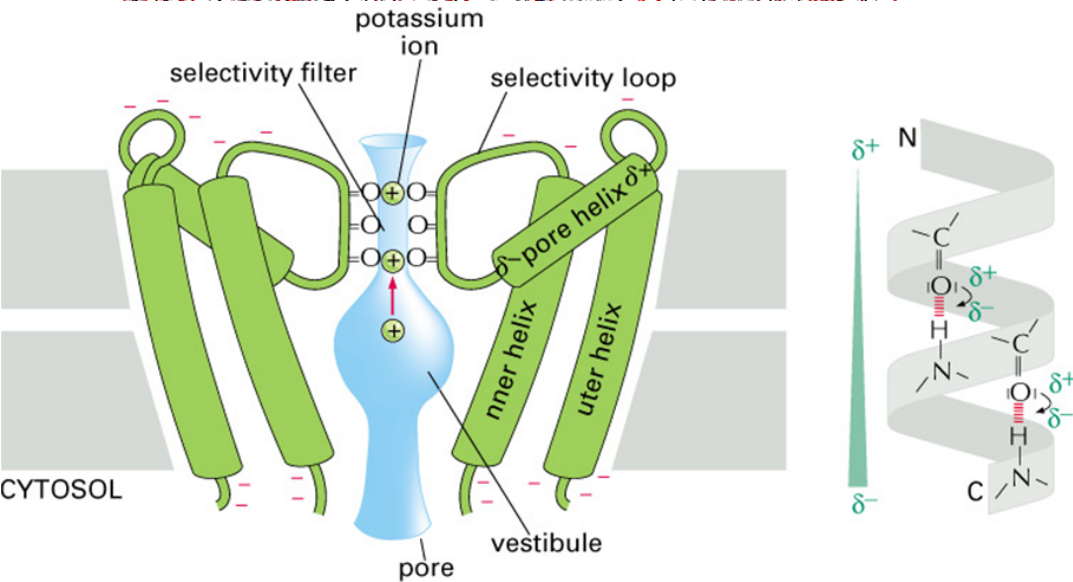
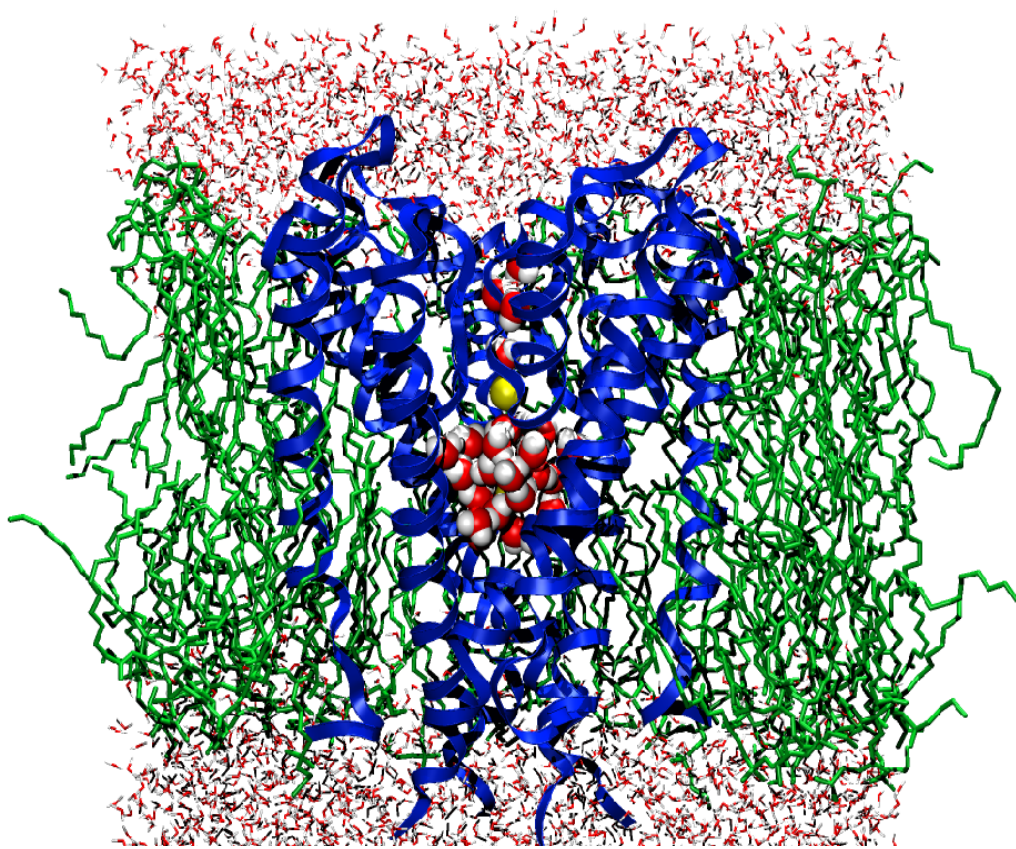


Negatively charged acidic residues are on top attract K^+



The negative charges on the bottom help to pull the K^+ ion through the pore.

The central cavity is lined with hydrophobic residues.



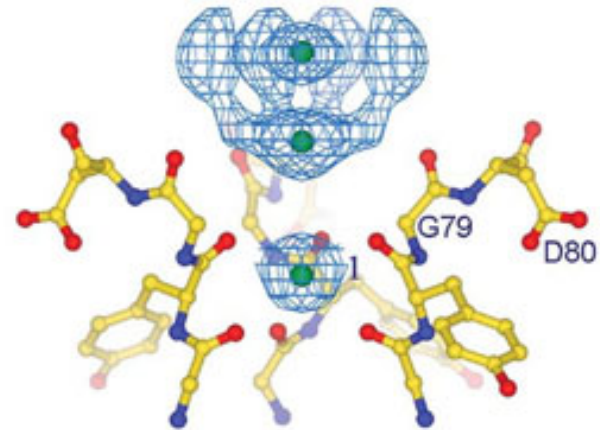
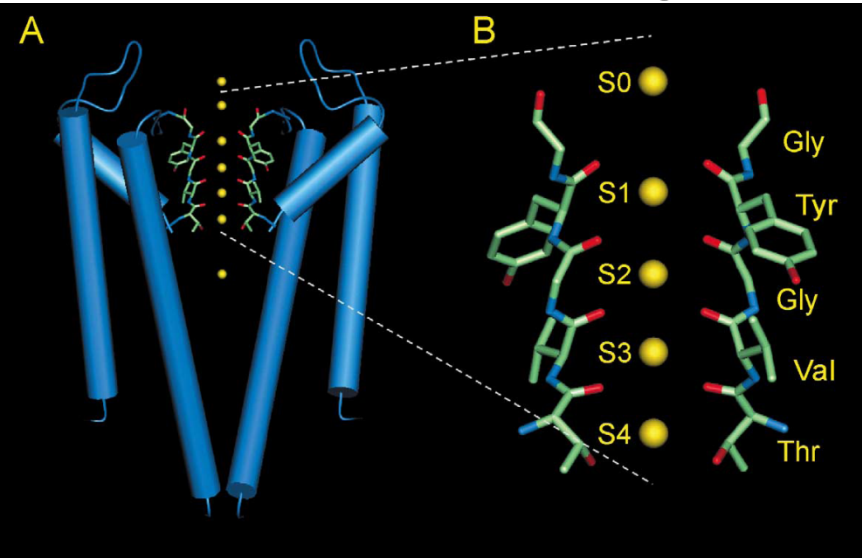
Selectivity filter

$C=O$ atoms of the protein backbone form selectivity filter ($4 \times$ *Tyr-Val-Gly-Tyr-Gly*).

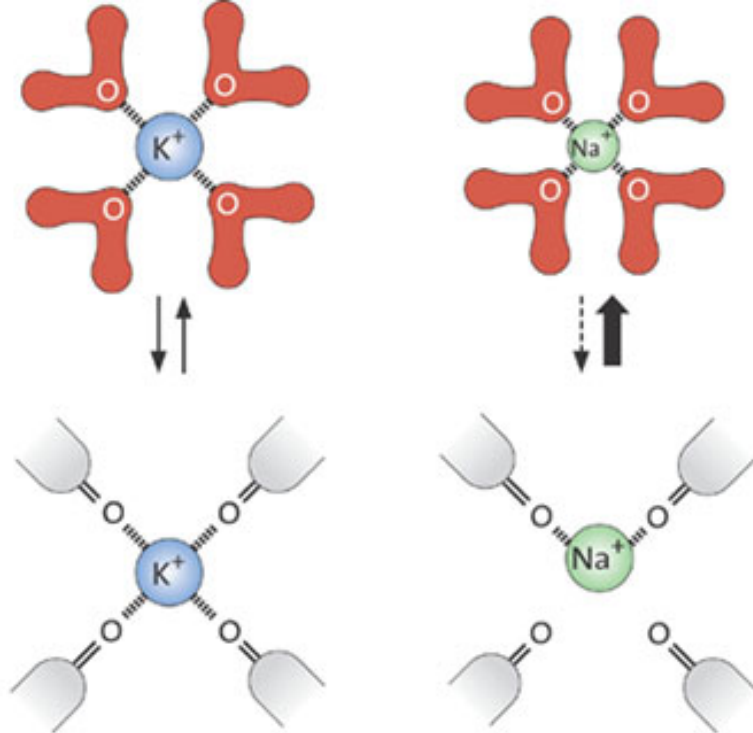
The sequence is **conserved** in all K^+ -channels.

❑ *Four K^+ ion binding sites.*

	- Filter	Inner Helix	
MthK	: YWTFVTIATVGYGDYS--PSTPLGMYFTVTLIVL	SIGTFFAVAVERLLEFLINRE	
KCSA	: WWSVETATTVGYGDLY--PVTLWGRIVAVVMVA	ITSEGLVTAALATWFGVRE	
Dradio	: YWAVVTVTTVGYGDIS--PKTGLGKEFIATLAML	SYAIIAVPTGIVTVGLQQAQ	
Ecoli	: YFSIETMSTVGYGDIV--PVSESARLETISVII	SITVFATSMTSIFGPLIRGG	
Shaker	: WWAVVTMTTVGYGDMT--PVGFWGKIVGSLCVVA	VLTIALPVPVIVSNFNIFY	
hDRK1	: WWATITMTTVGYGDIY--PKTLLGKIVGGLCCIA	VLVIALPIPIIVNNFSEFY	
hBK	: YLLMVTMSTVGYGDVY--AKTTLGRLFMVFFILG	LAMFASYVPEIIEELIGNRK	
hSK3	: WLISITFLSIGYGDMV--PHTYCGKGVCLLTGIM	AGCTALVVAVVARKLELTK	
hERG2	: YFTFSSLTSVGFNVN--PNTNSEKIFSICVMLI	SLMYASIFGNVSAIIQRLY	
hGIRK2	: LPSIETETTIGYGYRVITDKCPEGIILLIIQSVL	SVINAFMVGCMFVKISQPK	
hIRK1	: LPSIETQTITIGYGERCVTDECPIAVFMVVFQSI	VCIIDAFIIGAVMAKMAKPK	
bcNG1	: YWSTLTTLTIG--ETPP-PVRDSEYFFVVADELI	VLIFATIVGNIGSMISNMN	

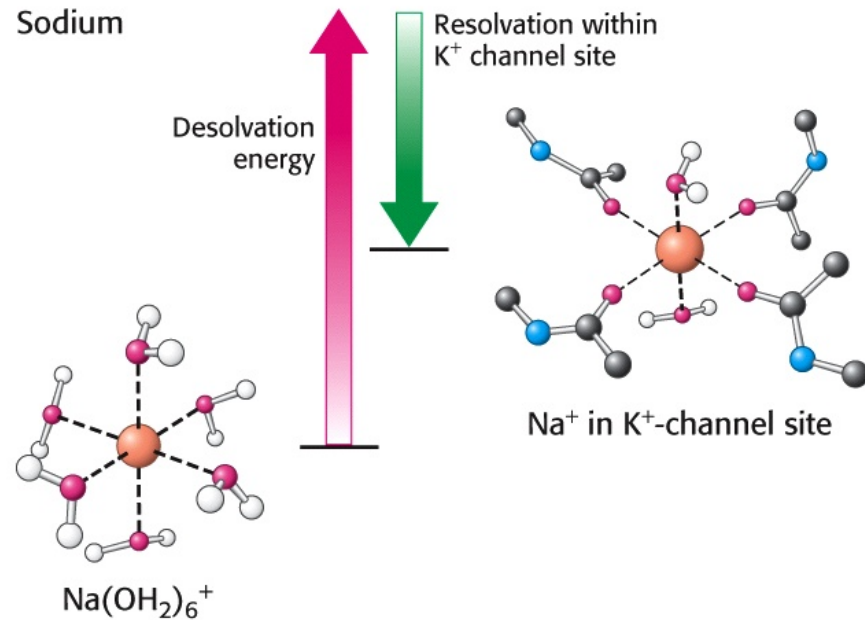
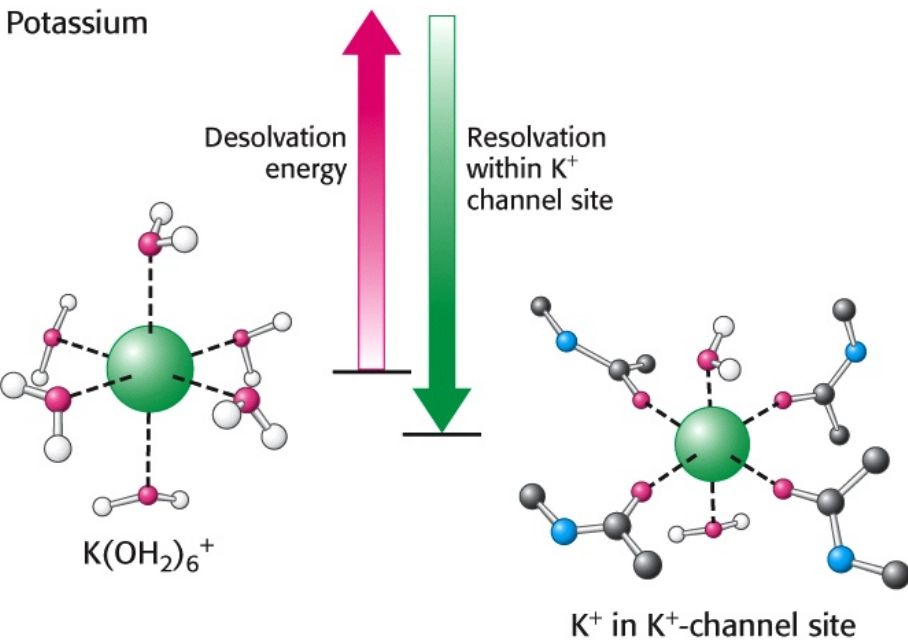


- ❑ *K^+ is surrounded by eight oxygen atoms from the protein*
- four 'above' and four 'below'.
 - very similar to water molecules around hydrated K^+ .



Potassium

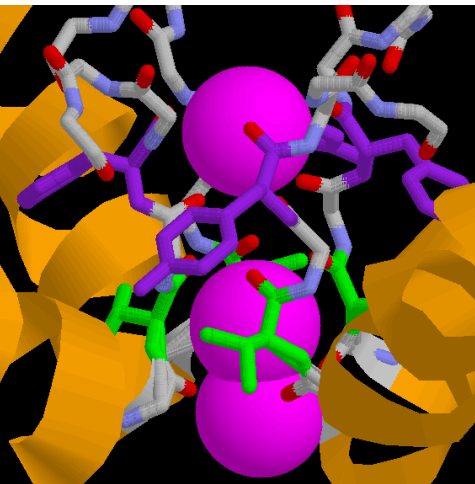
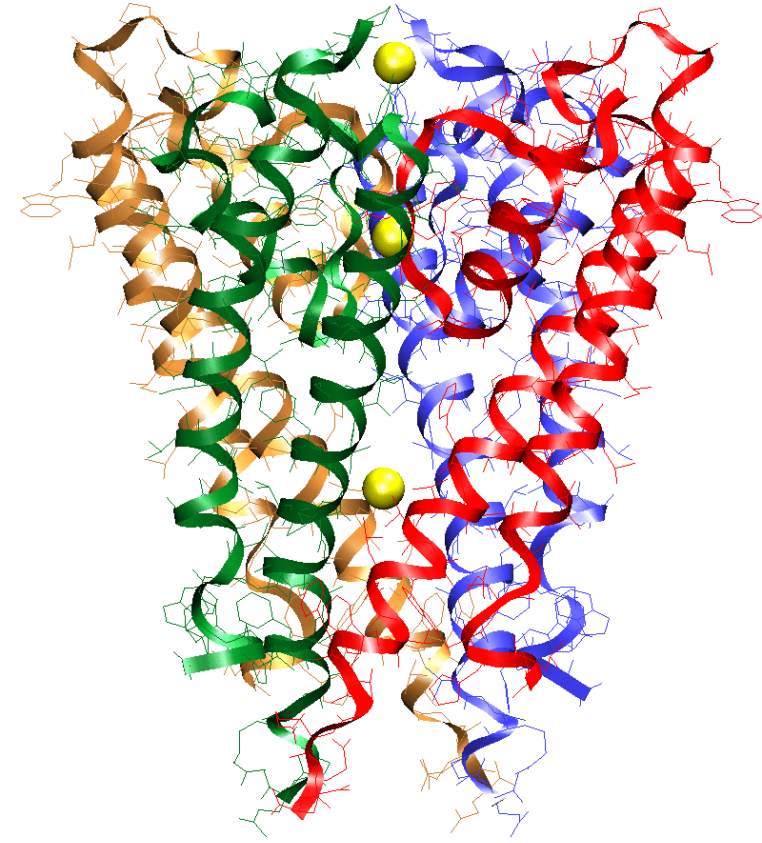
Sodium



The selectivity filter needs to be rigid

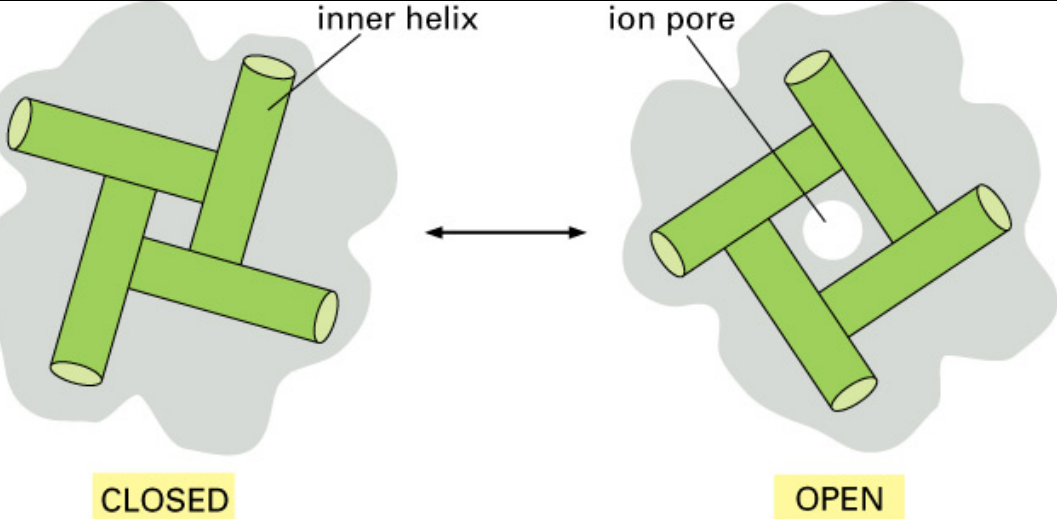
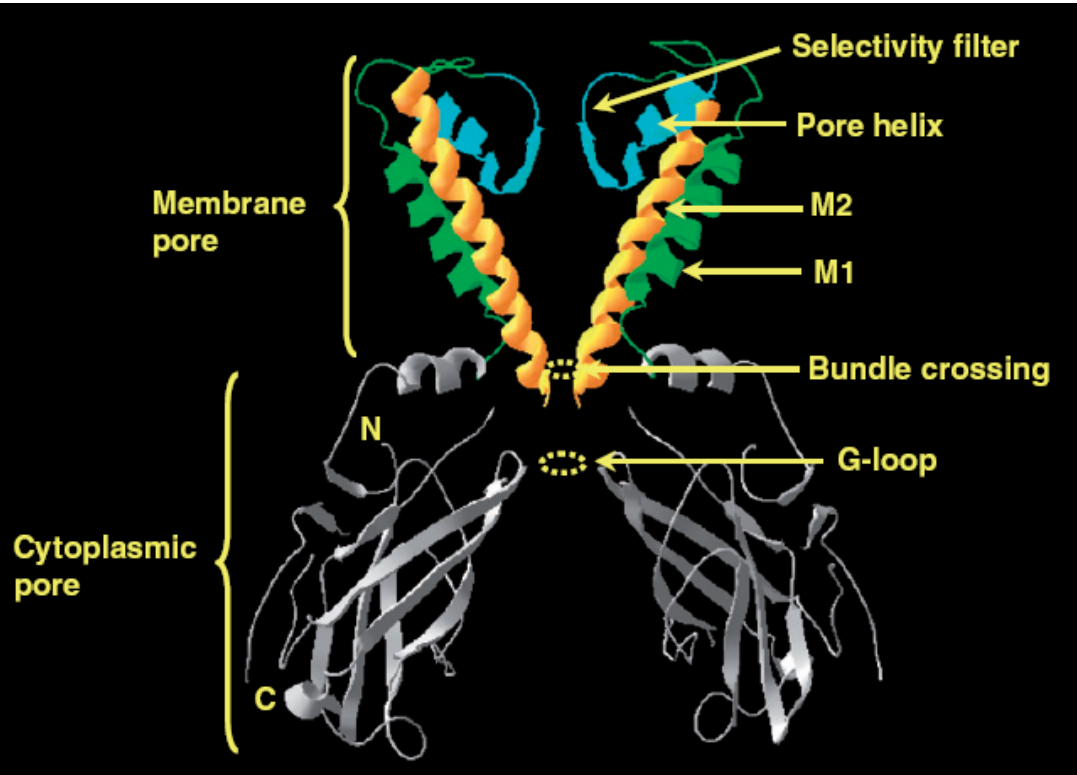
- ❑ The Val and Tyr hold the selectivity filter at a certain diameter by hydrogen bonding with the inner helix.
- ❑ They form hydrogen bonds which acts as tight springs that will not allow the pore to collapse.

**KcsA Potassium Channel from
*Streptomyces lividans***

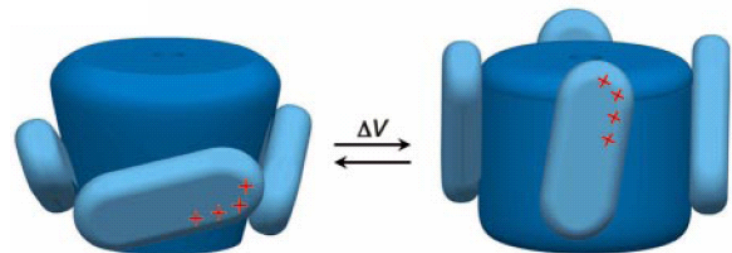


- ❑ The "springs" prevent the selectivity filter from interacting with cations smaller than K^+ .

Topology of a Kir subunit.



The bacterial voltage-gated K⁺ channel KvAP



The gate

