

# Ion Channel Structure and Function (part 1)

# The most important properties of an ion channel

Intrinsic properties of the channel:  
**Selectivity** and **Gating (activation, inactivation)**

+

Location



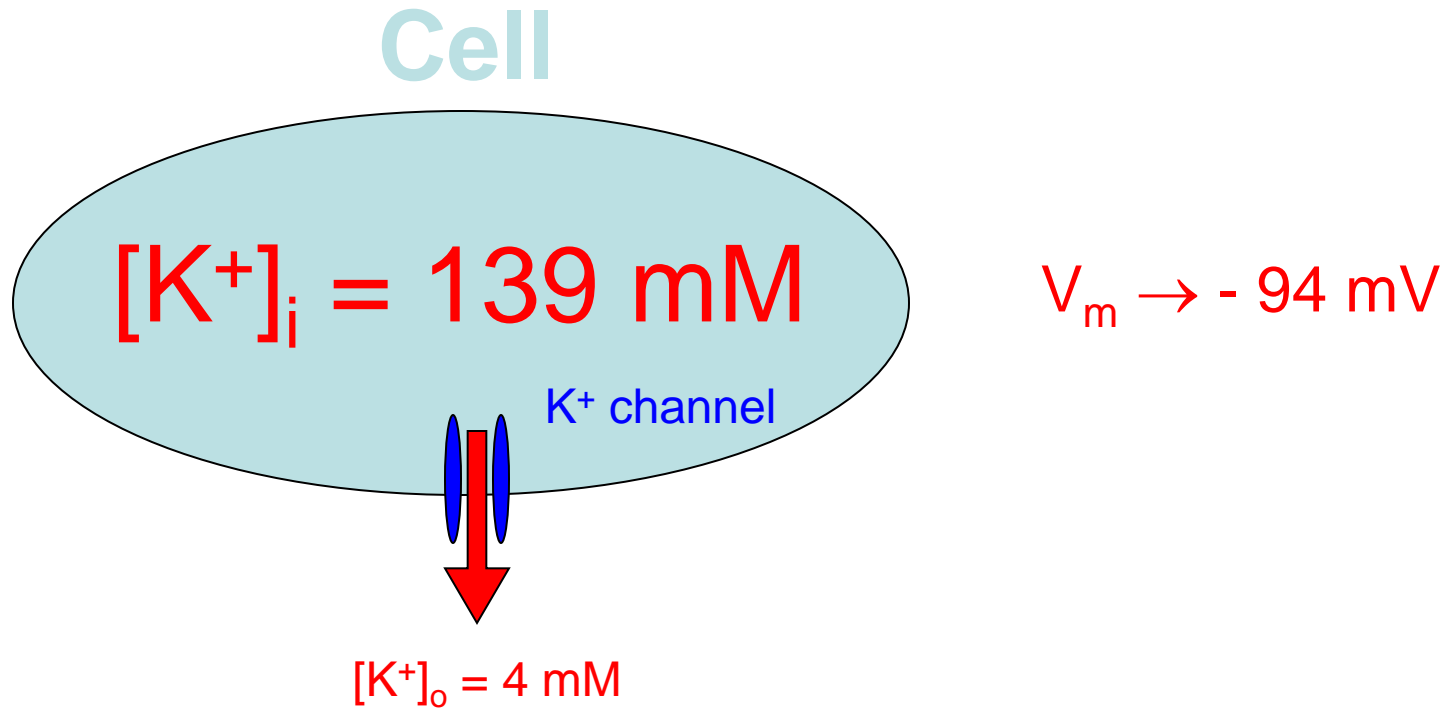
Physiological Function

# Types of ion channels by selectivity

Ion channels can be:

- Potassium ( $K^+$ )
- Sodium ( $Na^+$ )
- Calcium ( $Ca^{2+}$ )
- Nonselective Cation ( $Na^+$ ,  $K^+$ ,  $Ca^{2+}$ )
- Proton ( $H^+$ )
- Chloride ( $Cl^-$ )
- Hydroxide ( $OH^-$ )
- Nonselective Anion ( $Cl^-$ ,  $P_i$ ,  $ATP^{4-}$ , small negatively charged metabolites)
- Large-conductance nonselective ( $Na^+$ ,  $K^+$ ,  $Ca^{2+}$ ,  $Cl^-$ ,  $P_i$ ,  $ATP^{4-}$ , small metabolites)

# Potassium (K<sup>+</sup>) channels



**Equilibrium (Nernst) potential for K<sup>+</sup>:**

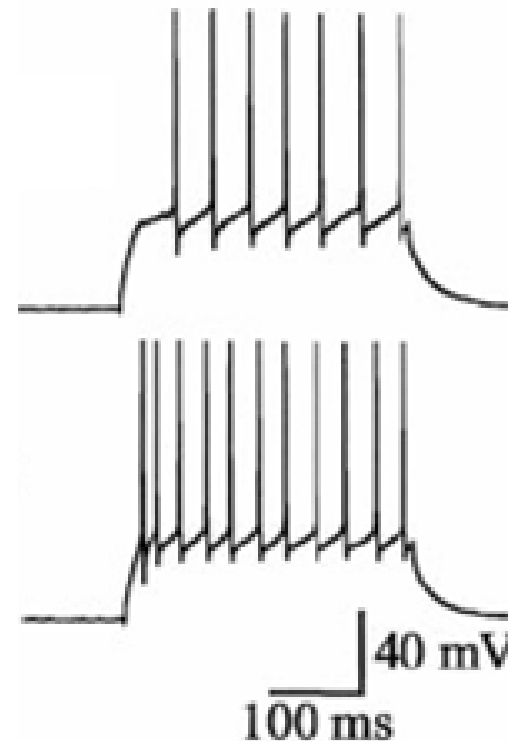
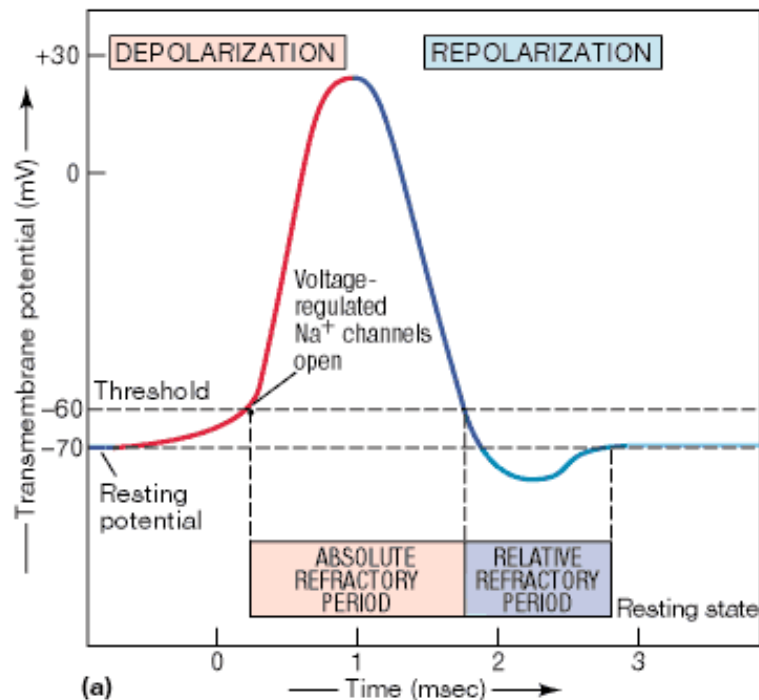
$$E_K = RT/zF \{\ln[K^+]_o/[K^+]_i\} = 61 \{\log_{10}[K^+]_o/[K^+]_i\} = -94 \text{ mV}$$

# $K_v$ , voltage-gated $K^+$ channels

Gating: opened by membrane depolarization; there are fast inactivating (A-type, ms) and slow inactivating (delayed-rectifier type, s)  $K_v$  channels

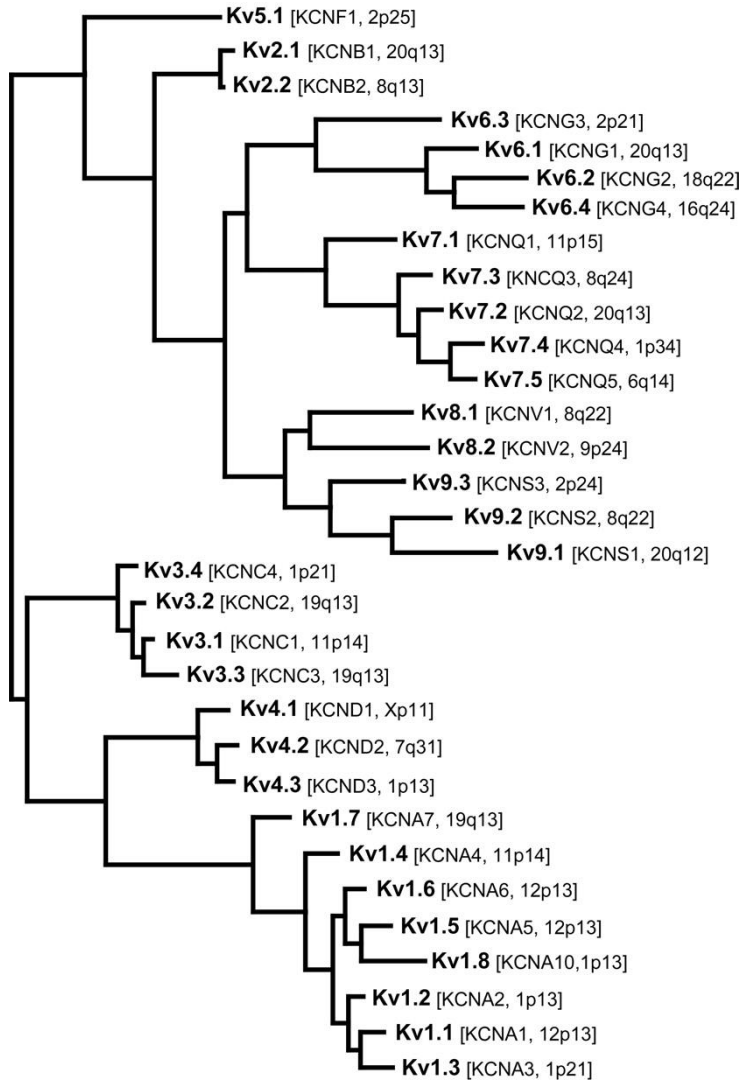
Location: plasma membrane of neurons, muscle cells, and many non-excitable cells

Function: maintaining membrane potential; repolarization of action potential and shaping its waveform; modulating firing pattern and electrical excitability in neurons and muscle.

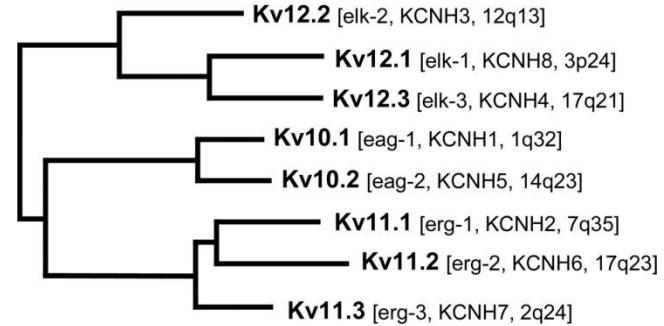


# K<sub>v</sub>, voltage-gated K<sup>+</sup> channels

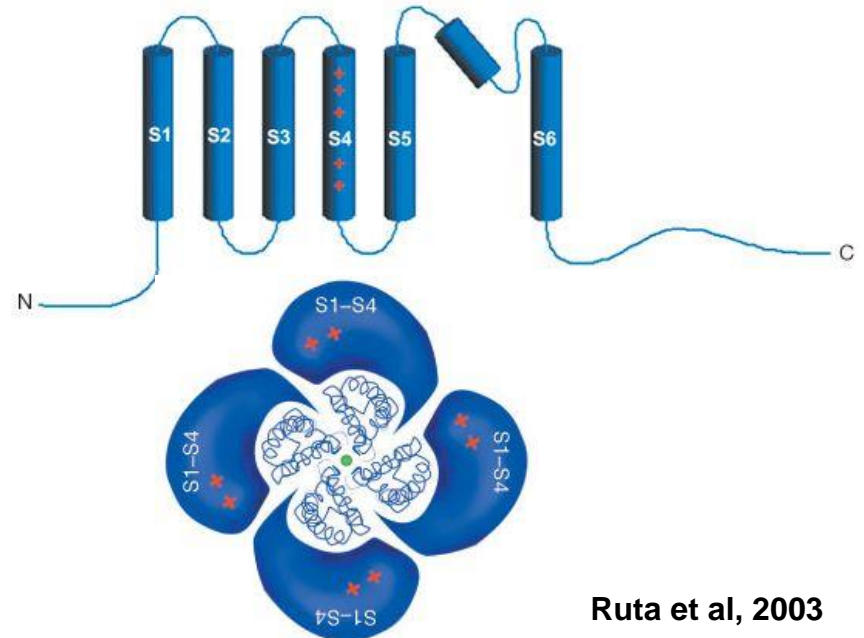
Phylogenetic Tree, Kv1-9 Families



Phylogenetic Tree, Kv10-12 Families



K<sub>v</sub> α subunits make homo- and hetero-tetrameric complexes

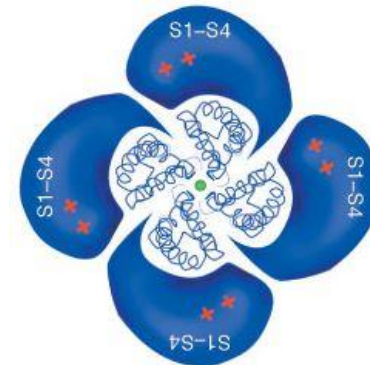
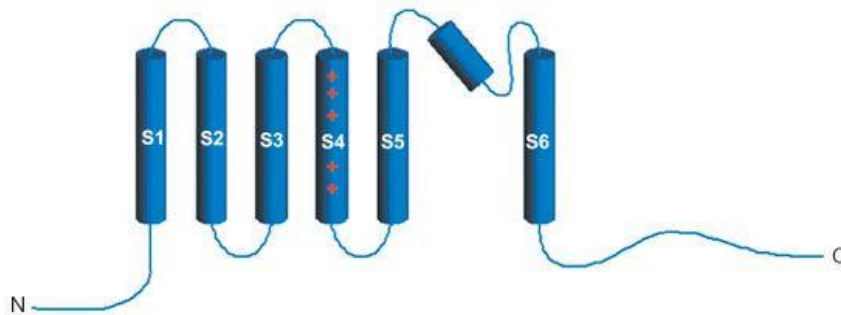


# Important questions about molecular architecture of $K_v$ channels

1. How high permeability and amazing selectivity for  $K^+$  are simultaneously achieved?

- $K_v$  channels enable extremely fast ion flow,  $\sim 10^8 K^+$  per second
- $K^+$  is at least 10,000 times more permeant than  $Na^+$ , a feature that is essential to the function of  $K^+$  channels

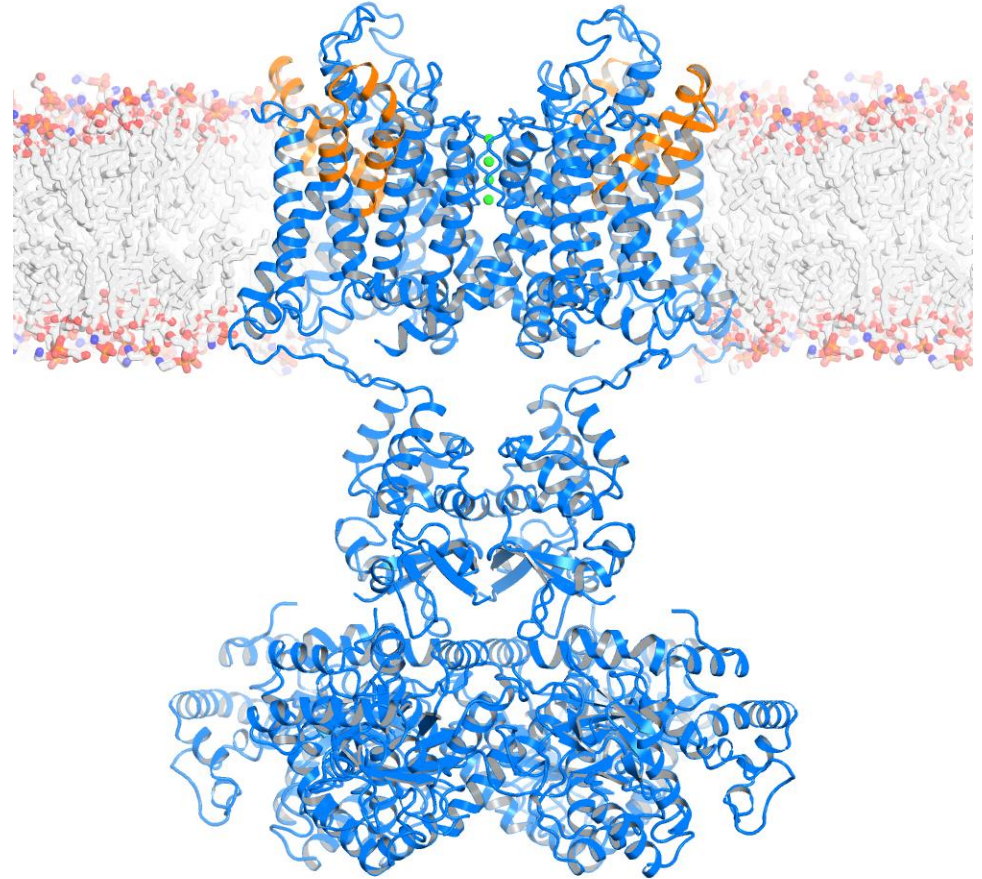
2. How changes in membrane voltage are coupled to channel opening?



# Roderick MacKinnon

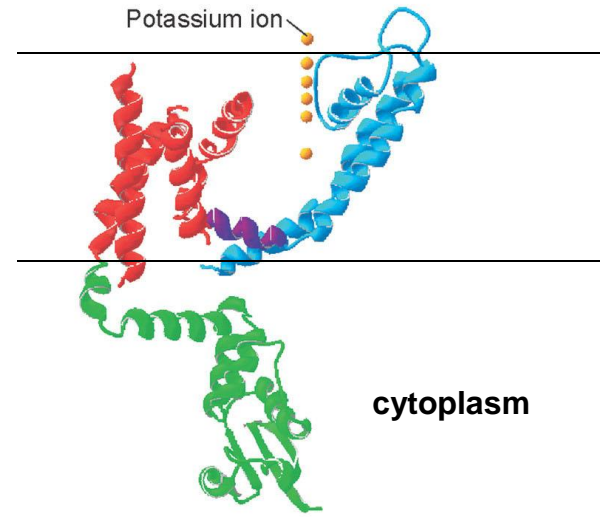
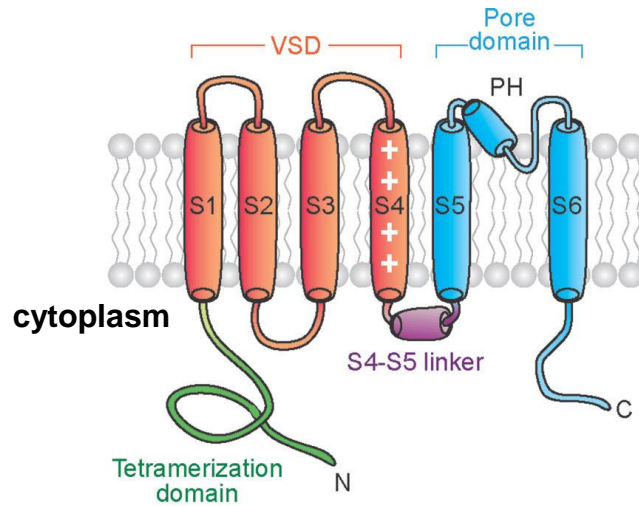


Nobel Prize in Chemistry 2003





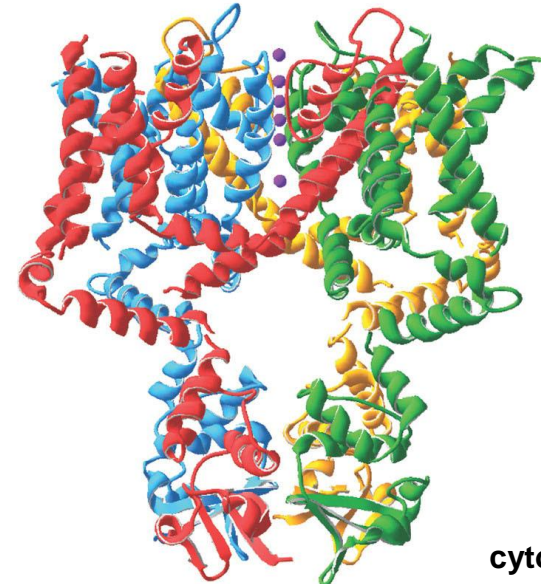
# Overview of the K<sub>v</sub> channel structure



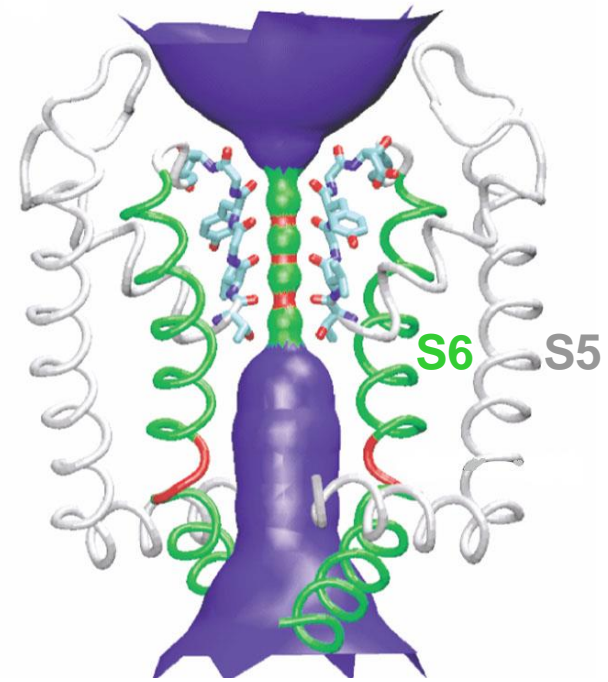
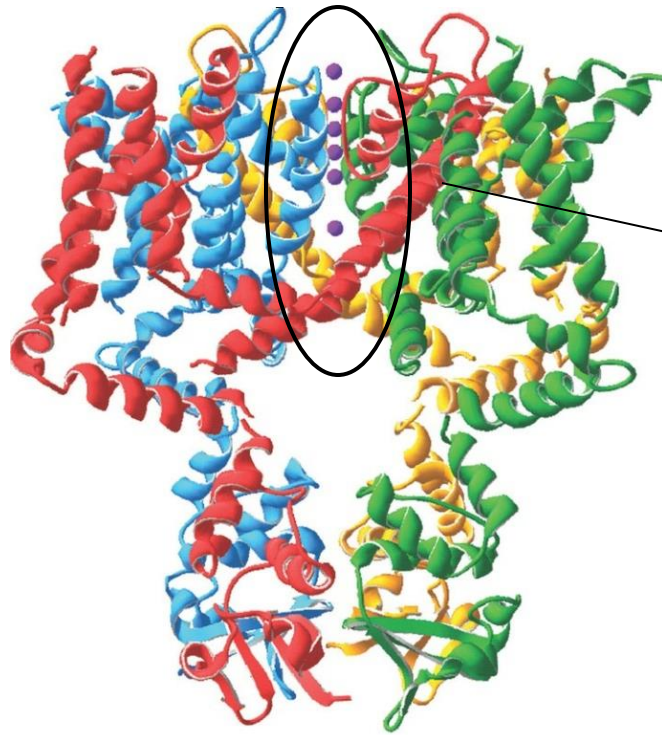
Top view



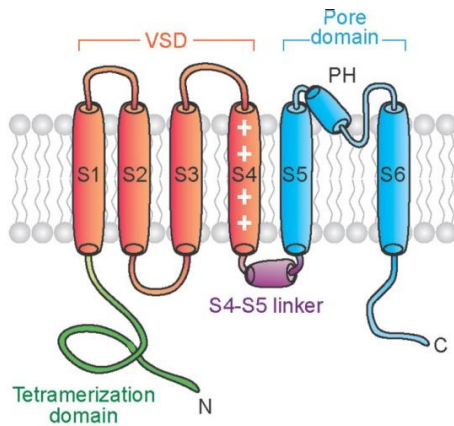
Side view



# Pore Region



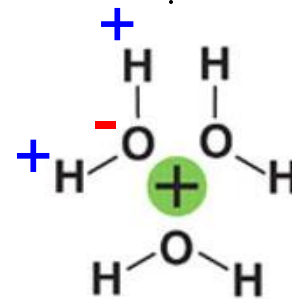
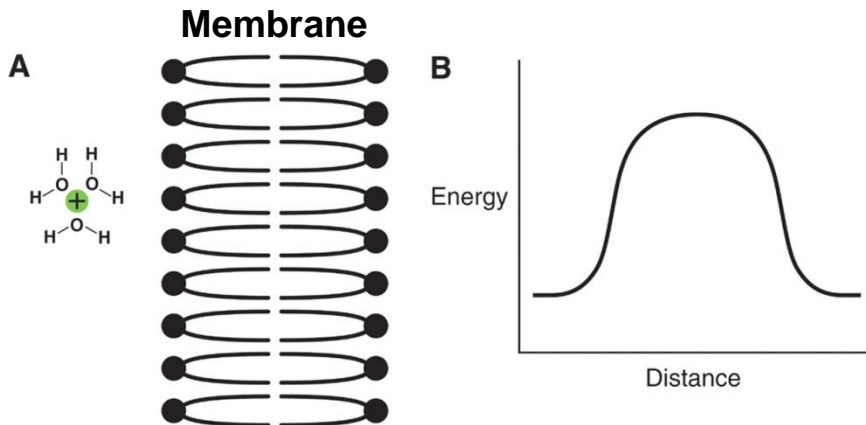
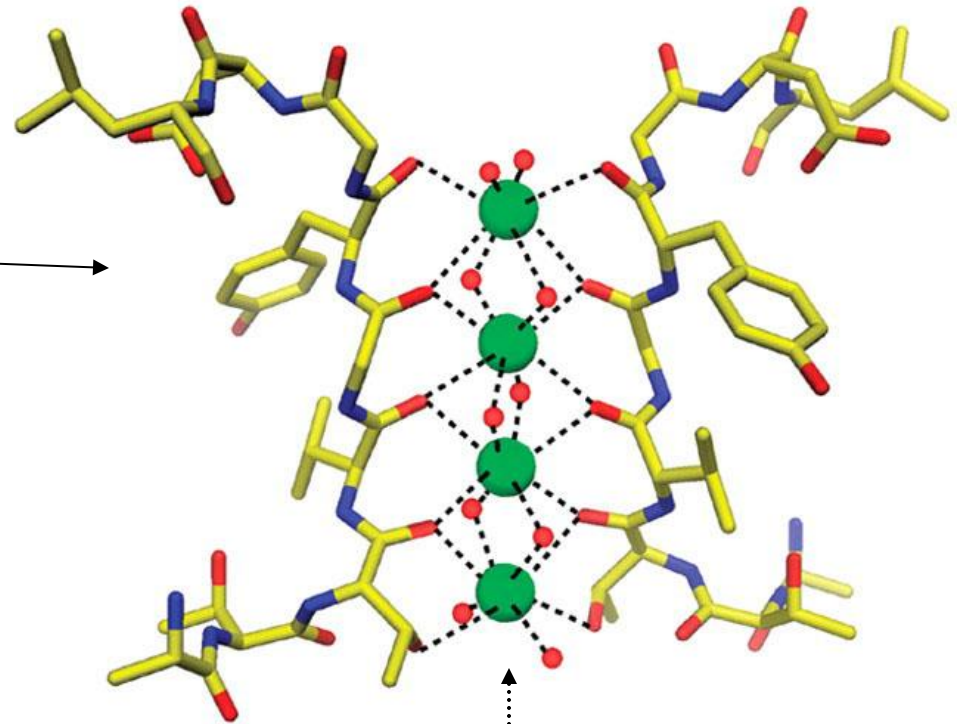
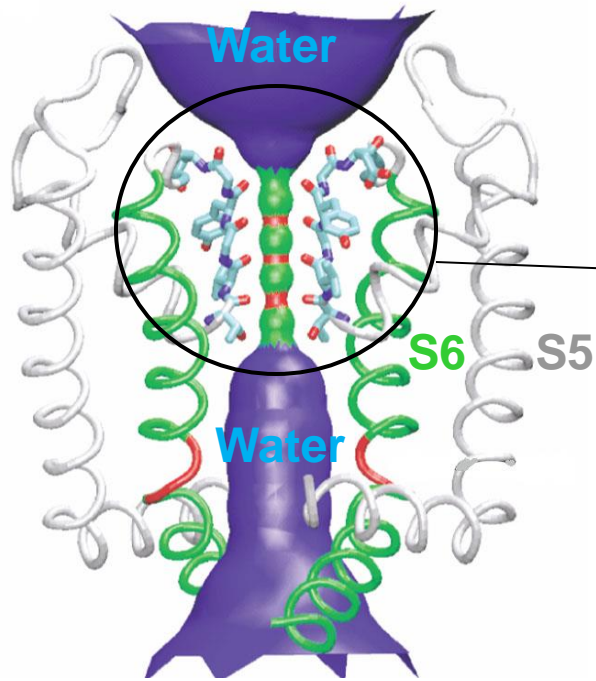
cytoplasm



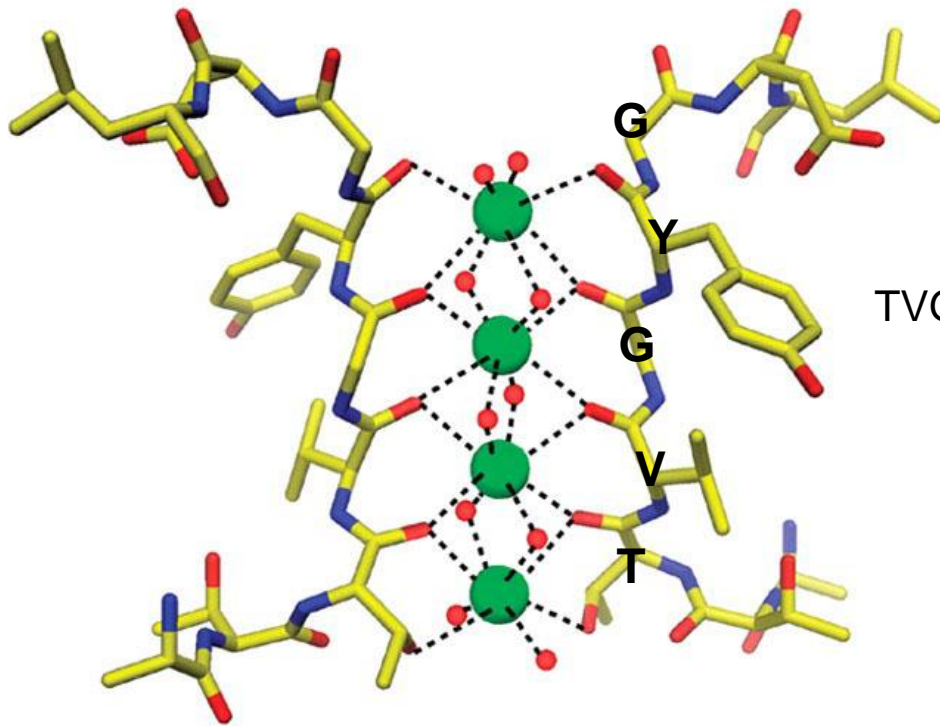
Tombola et al. 2006

Grottesi et al. 2005

# Pore Region - Selectivity Filter

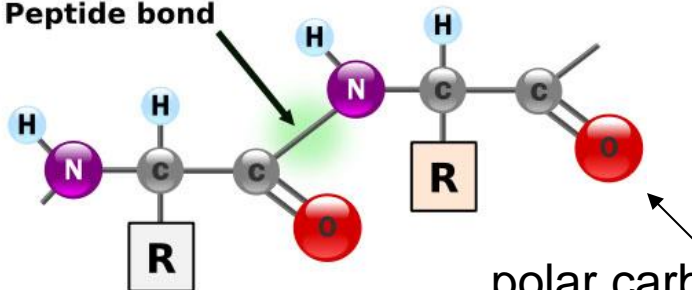


# Pore Region - Selectivity Filter

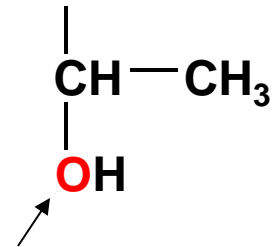


TVGYG (threonine-valine-glycine-tyrosine-glycine)  
– signature motif of K<sup>+</sup> channels

Peptide bond

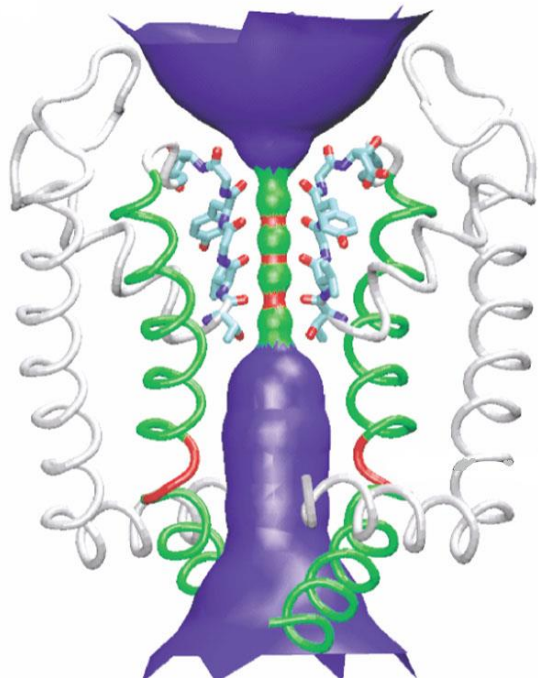


polar carbonyl oxygen  
(main chain)



polar hydroxyl oxygen of threonine  
(side-chain)

# What types of gating motions might one expect?



A. Trap door motion



B. Collapsing motion



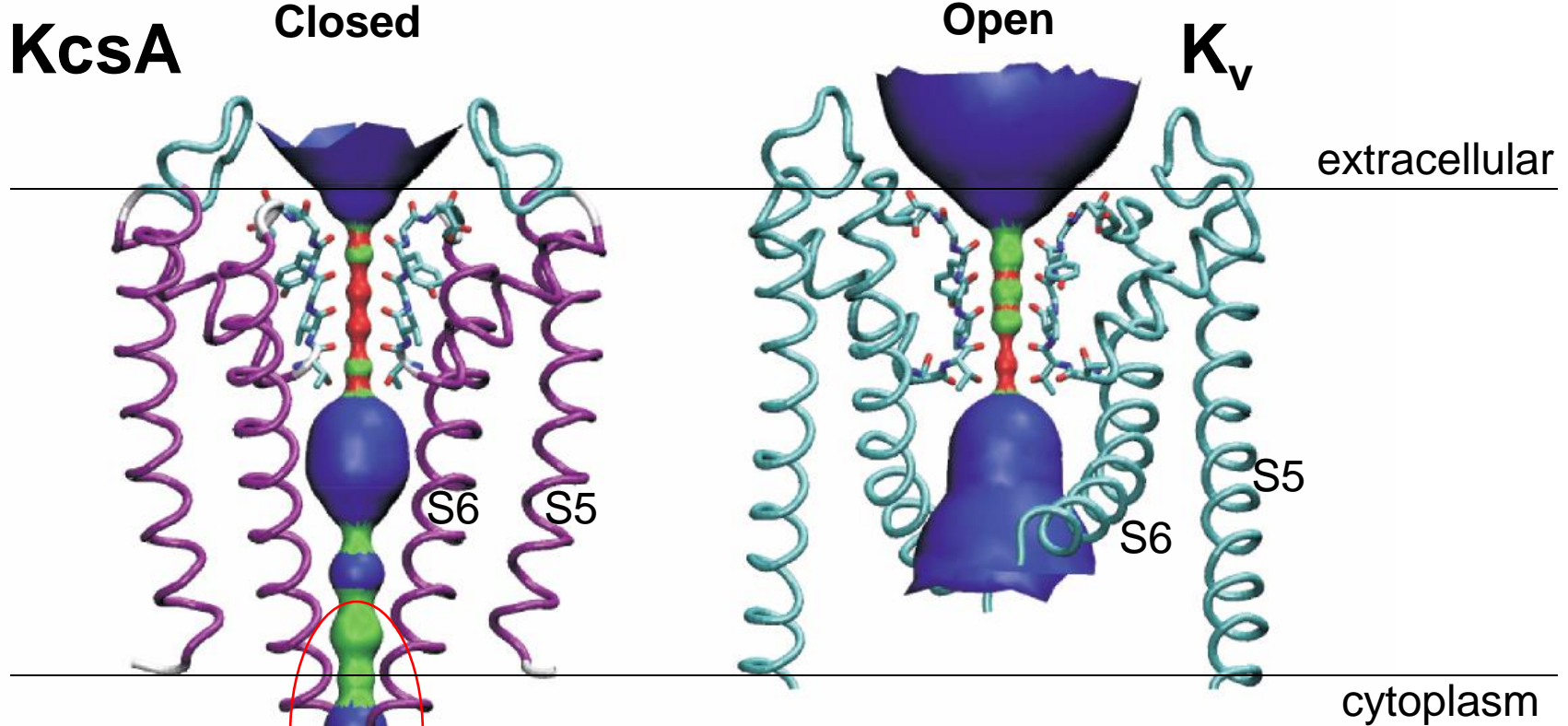
C. Pinching of selectivity filter



Closed

Open

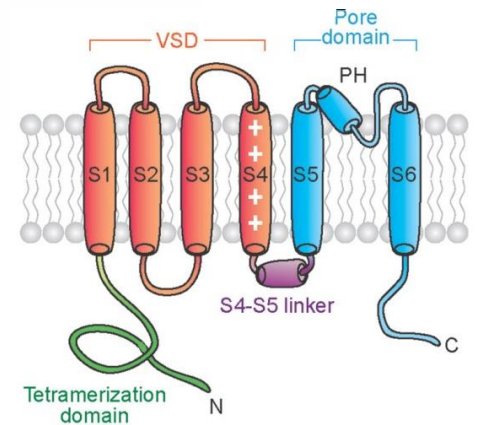
# Pore Region - Gate



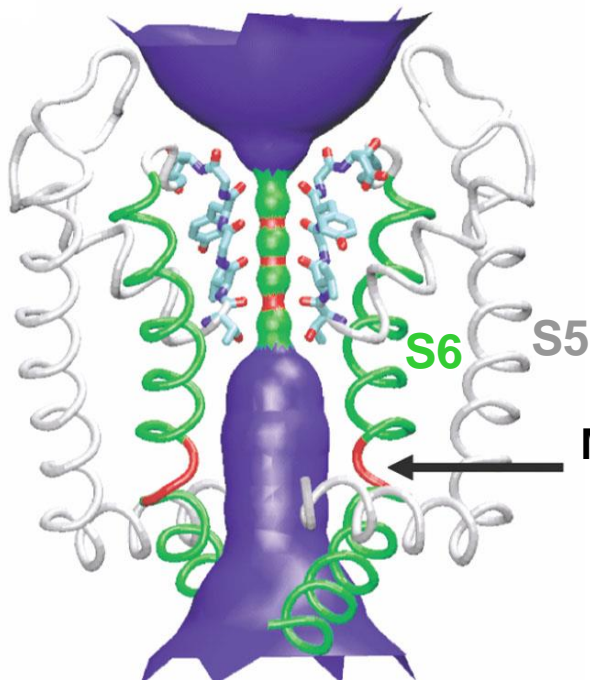
**Gate**

**“open” S6**

**“closed” S6**

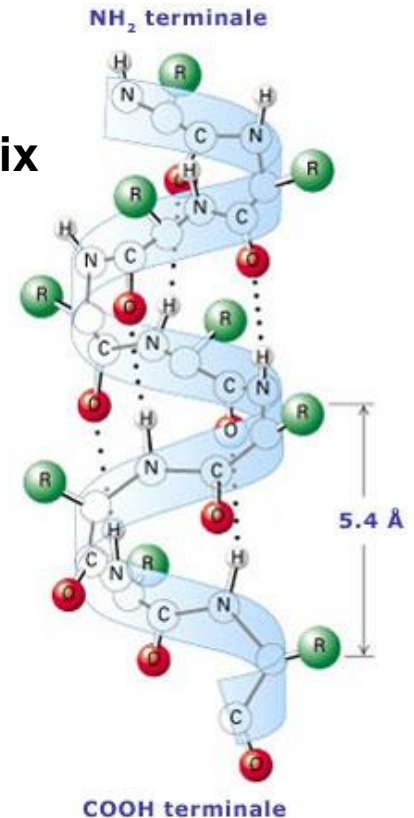


# Pore Region - Gate

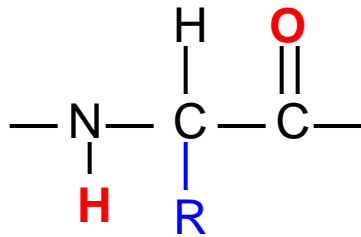


Molecular "hinge" of the "gate",  
the Proline-X-Proline motif

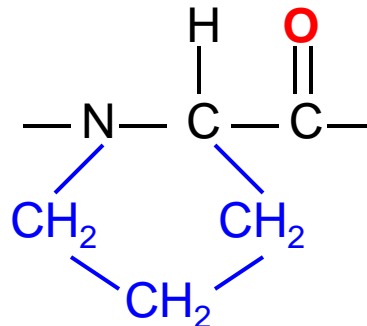
$\alpha$ -helix



Regular amino acid

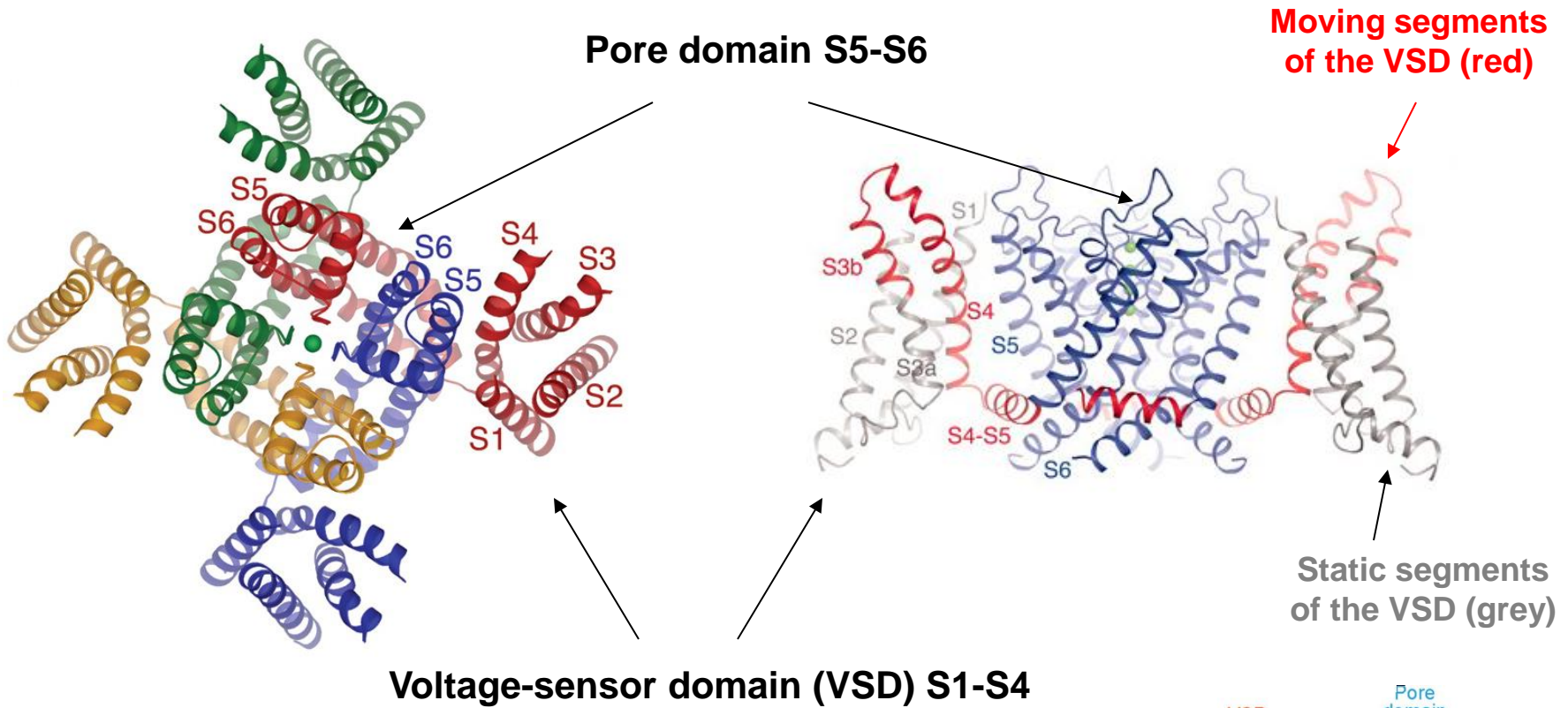


Proline

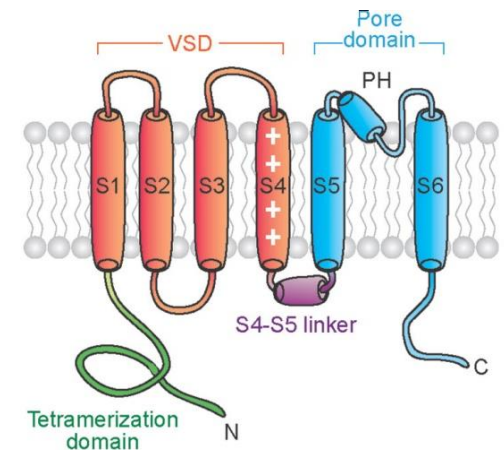


In the  $\alpha$ -helix, N-H group donates a hydrogen bond to the backbone C=O group of the amino acid 4 residues earlier

# Voltage Sensor and Gating

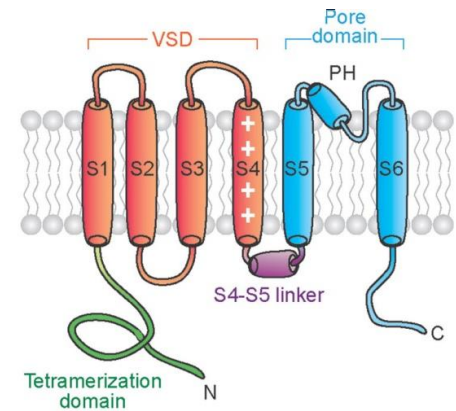
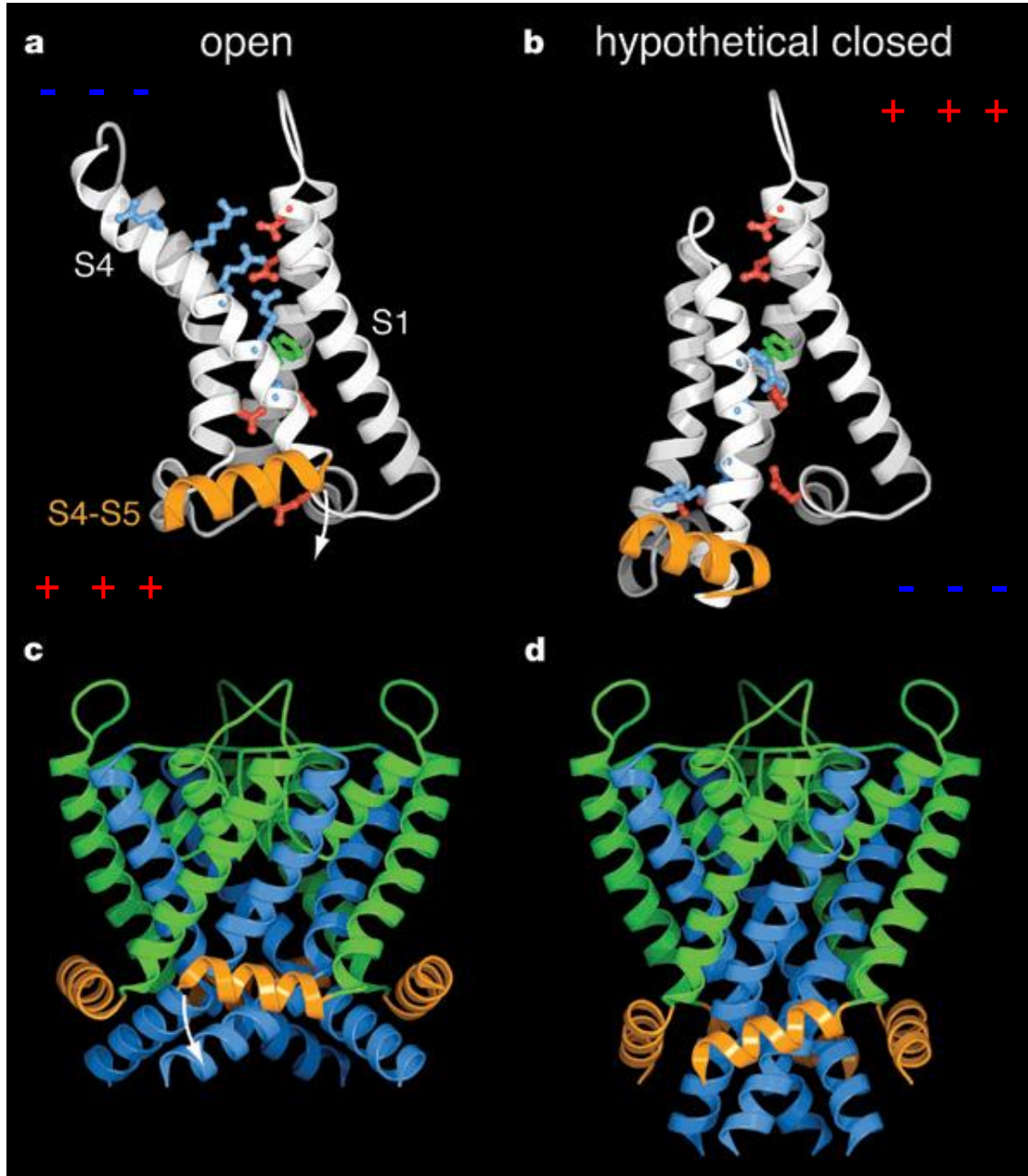


- Pore domain and VSD do not only have separate functions, but are also separated in space.
- Pore and VSD domains have only two contact points: S4-S5 linker and the top of the S1 transmembrane helix.





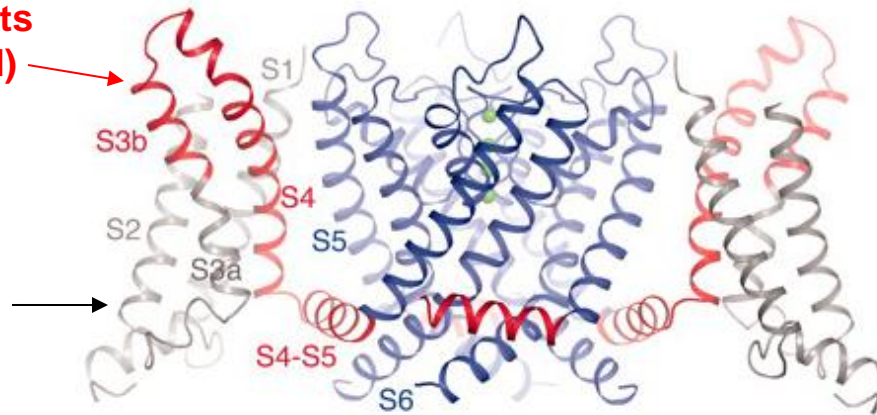
# Voltage Sensor and Gating



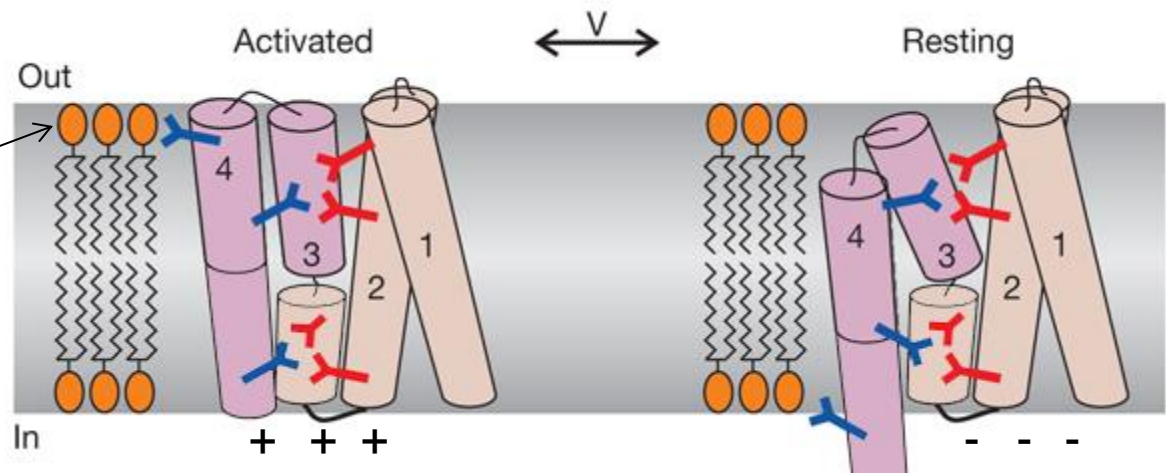
# Stabilization of the voltage sensor in the lipid environment

Moving segments of the VSD (red)

Static segments of the VSD (grey)



Negatively charged polar heads of phospholipids

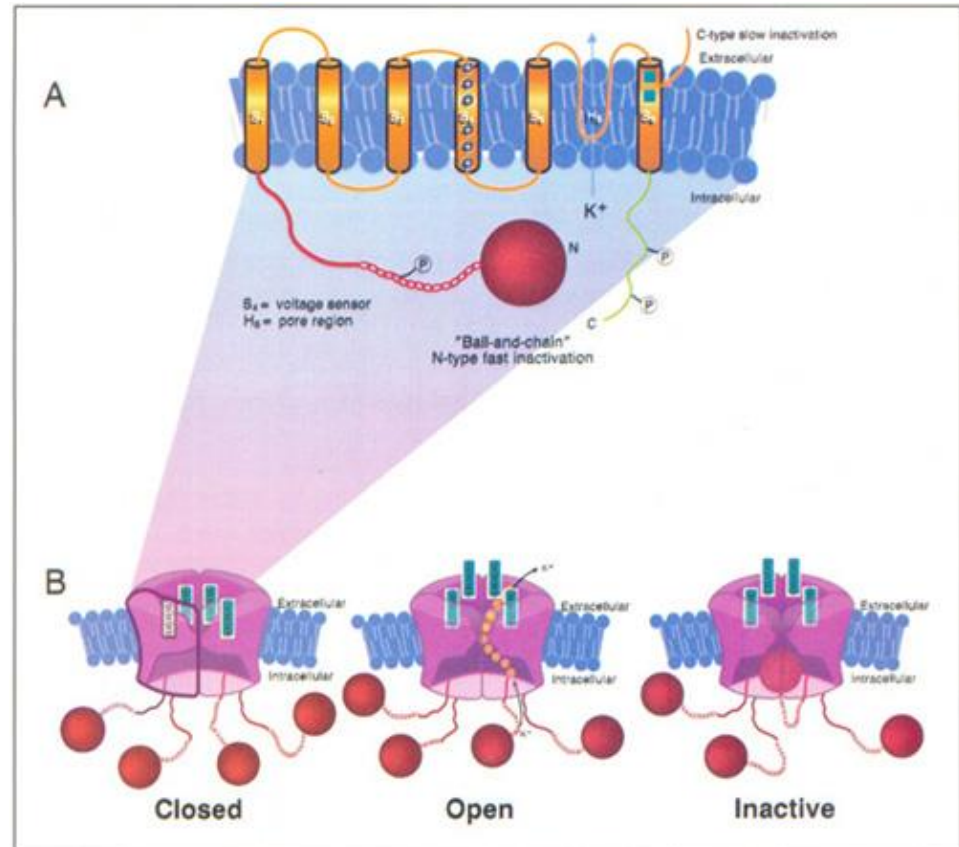


Positive charges of the S4 domain are stabilized by forming ion pairs with negative charges of the static segments of VSD and phospholipids.

# Inactivation of $K_v$ channels

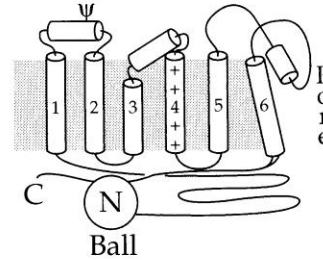
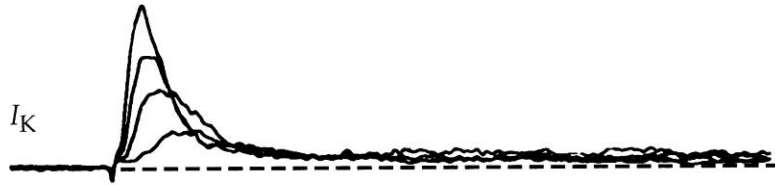
What is the difference between channel inactivation and closure?

- N-type or ball-and-chain inactivation: an N-terminal ball plugs the pore from the cytoplasmic side.
- C-type inactivation results from a localized constriction in the outer mouth of the channel pore.



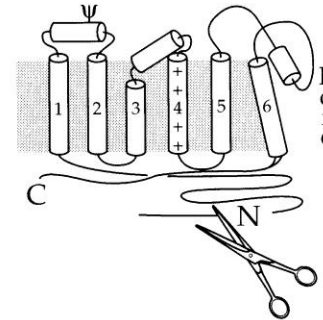
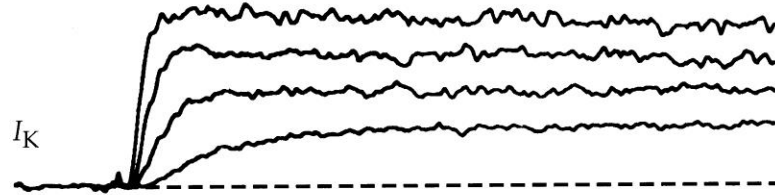
# N-type inactivation: ball and chain

(A) WILD-TYPE ShB



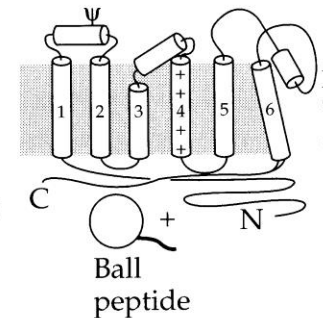
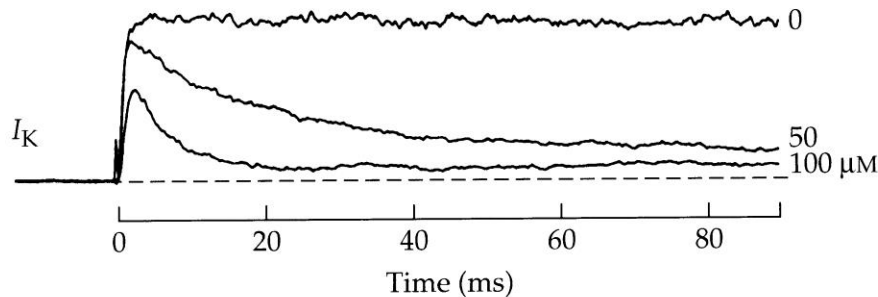
'ball' at  
N-terminus

(B) DELETION MUTANT  $\Delta 6-46$



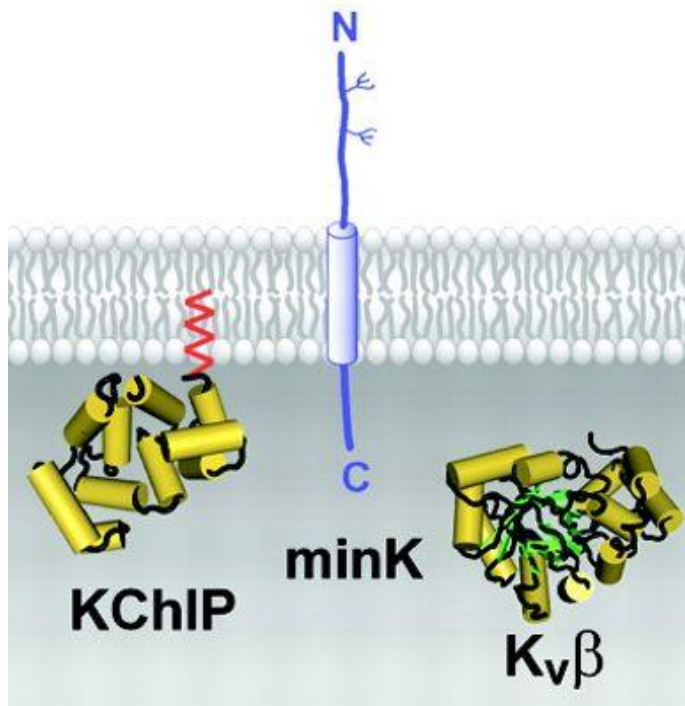
N-terminal  
truncated  
channel

(C) MUTANT + ShB PEPTIDE



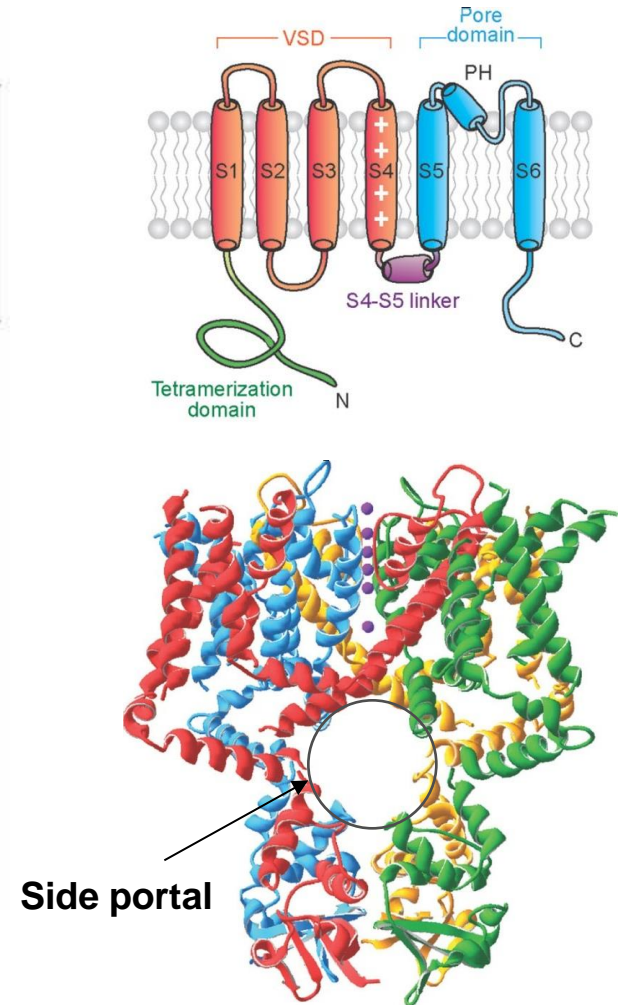
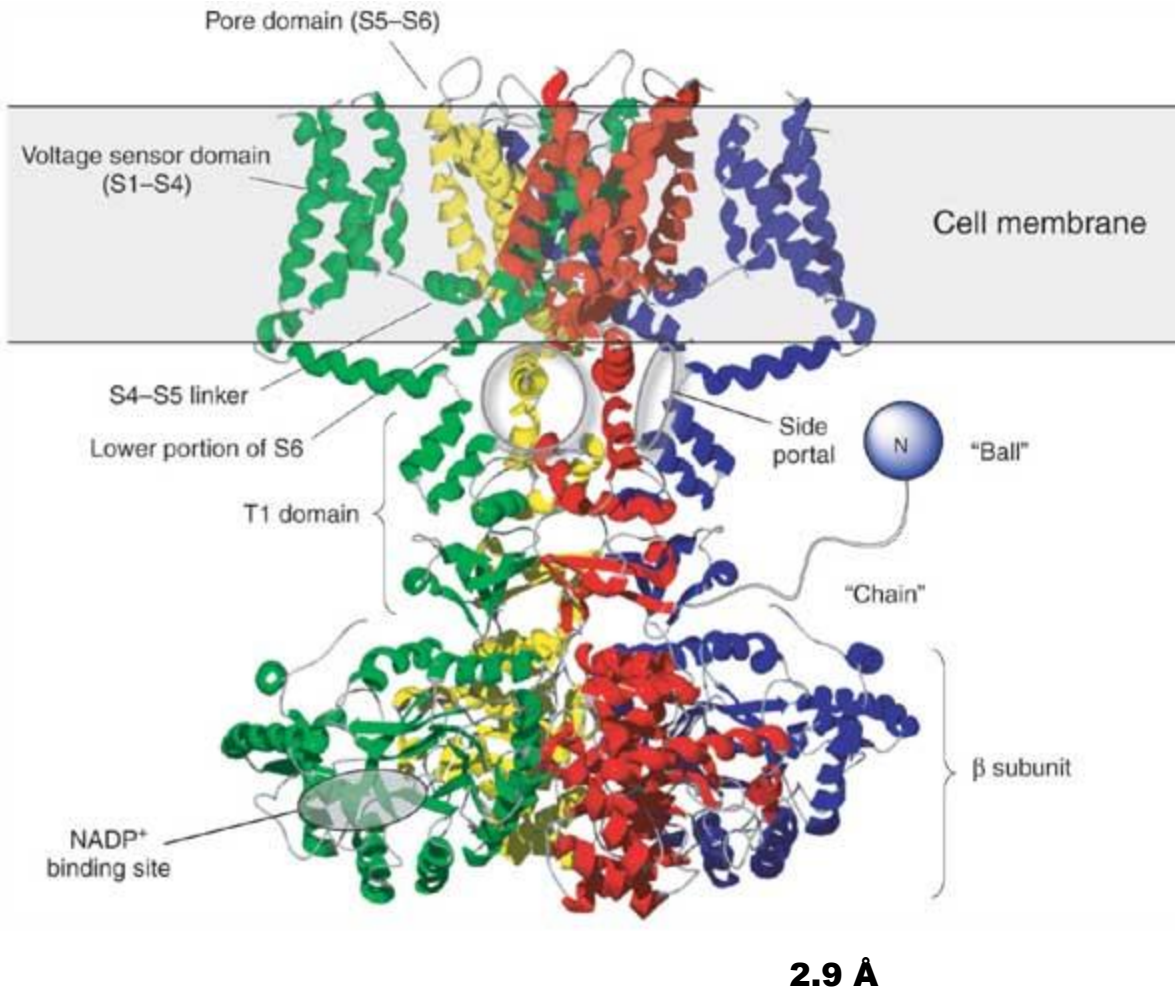
N-terminal  
truncated  
channel  
+  
Free N-terminal  
peptide

# Auxiliary subunits of K<sub>v</sub> channels



- K<sub>v</sub>1 channels are often associated with an intracellular subunit (K<sub>v</sub>β 1-3). The N terminus of K<sub>v</sub>β subunits serves as an N-type inactivation gate for K<sub>v</sub>1 α subunits. The β subunits bind to the N terminus of the α subunits.
- K<sub>v</sub>4 channels interact with the K channel interacting proteins KChIP1-4. The KChIPs enhance expression of K<sub>v</sub>4 channels and modify their functional properties.
- The K<sub>v</sub>3, K<sub>v</sub>4, K<sub>v</sub>7, K<sub>v</sub>10, and K<sub>v</sub>11 channels associate with the minK subunits (there 5 of them). The minK subunits are important regulators of K<sub>v</sub> channel function. In particular, they significantly slow down C-type inactivation.

# Crystal structure of the $K_v1.2/K_v\beta$ complex



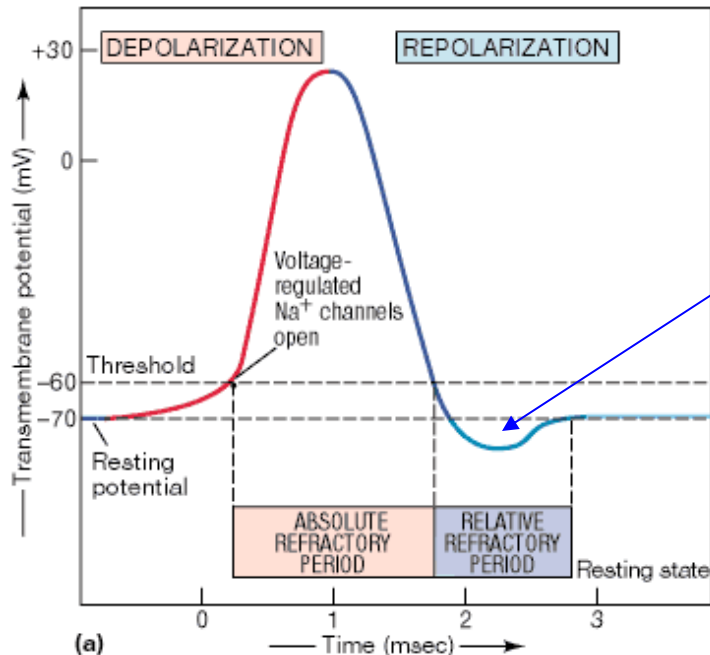
# $K_{Ca}$ , $Ca^{2+}$ -activated $K^+$ channels

Single-channel patch-clamp recordings identified two types of  $K_{Ca}$  channels: small conductance (SK) and high-conductance (BK)

Gating: opened by elevation of intracellular  $Ca^{2+}$  with  $K_d \sim 0.5 \mu M$  (SK), opened by elevation of intracellular  $Ca^{2+}$  & depolarization (BK)

Location: Plasma membrane of neurons, muscle cells and some non-excitabile cells

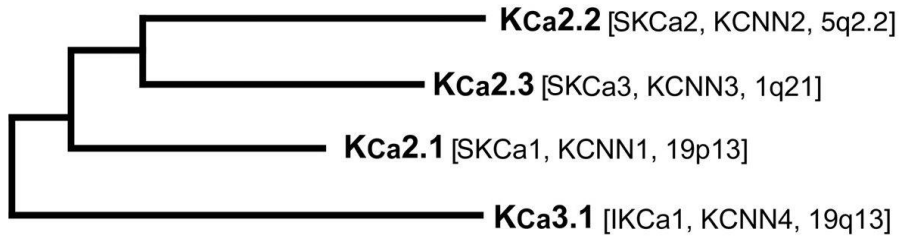
Function: **negative-feedback system for  $Ca^{2+}$  entry in many cell types**, slow afterhyperpolarization (up to 1 second, SK channels), fast afterhyperpolarization (several milliseconds, BK channels) and presynaptic regulation of neurotransmitter release (BK channels)



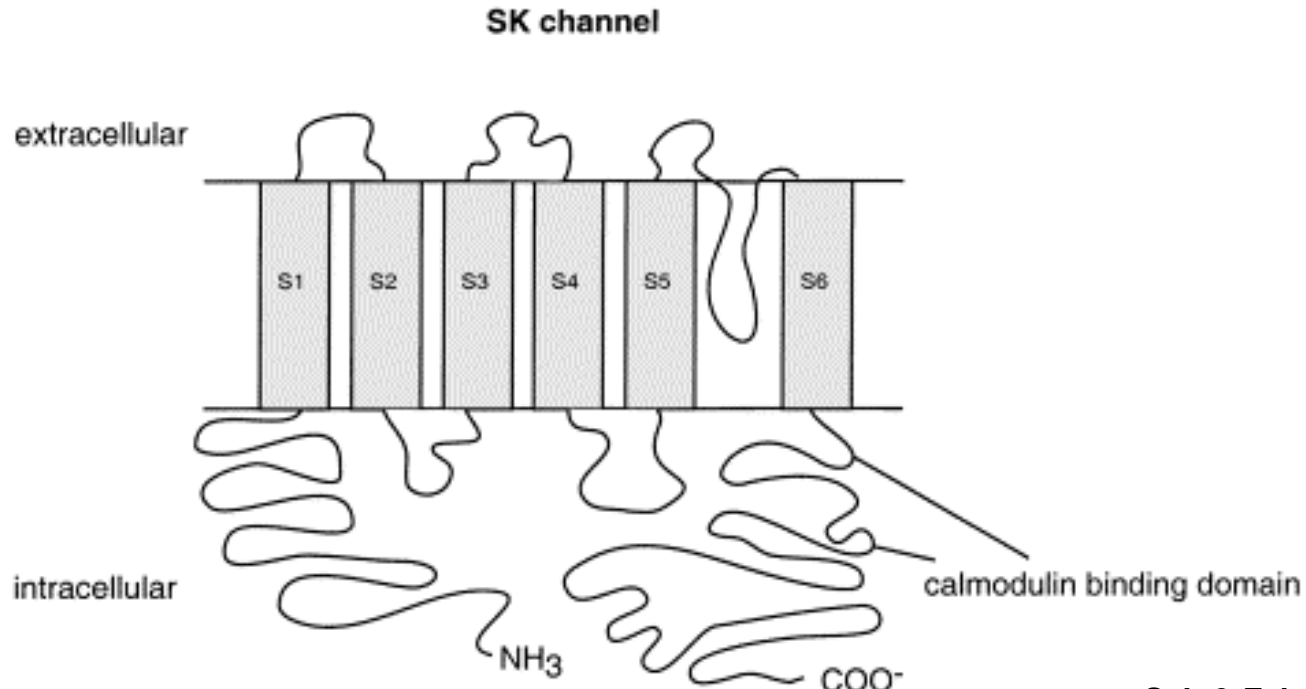
Afterhyperpolarization can be long, up to 1 second

# K<sub>Ca</sub>, Ca<sup>2+</sup>-activated K<sup>+</sup> channels

SK



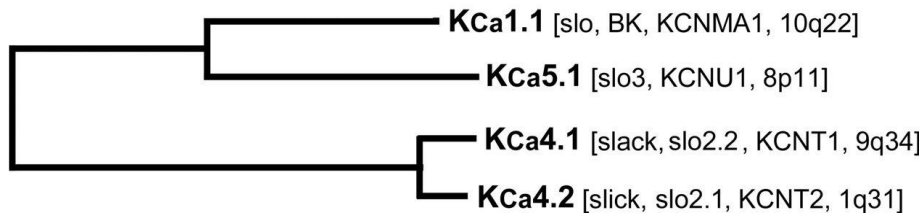
- SK  $\alpha$  subunits make homo-tetrameric and likely hetero-tetrameric complexes.
- There is no charge in S4 domain.
- Calmodulin is constitutively bound to C-terminal domain of the channel.





# K<sub>Ca</sub>, Ca<sup>2+</sup>-activated K<sup>+</sup> channels

## BK



- Interestingly, K<sub>Ca</sub>4.1, K<sub>Ca</sub>4.2 and K<sub>Ca</sub>5.1 are insensitive to intracellular Ca<sup>2+</sup>.

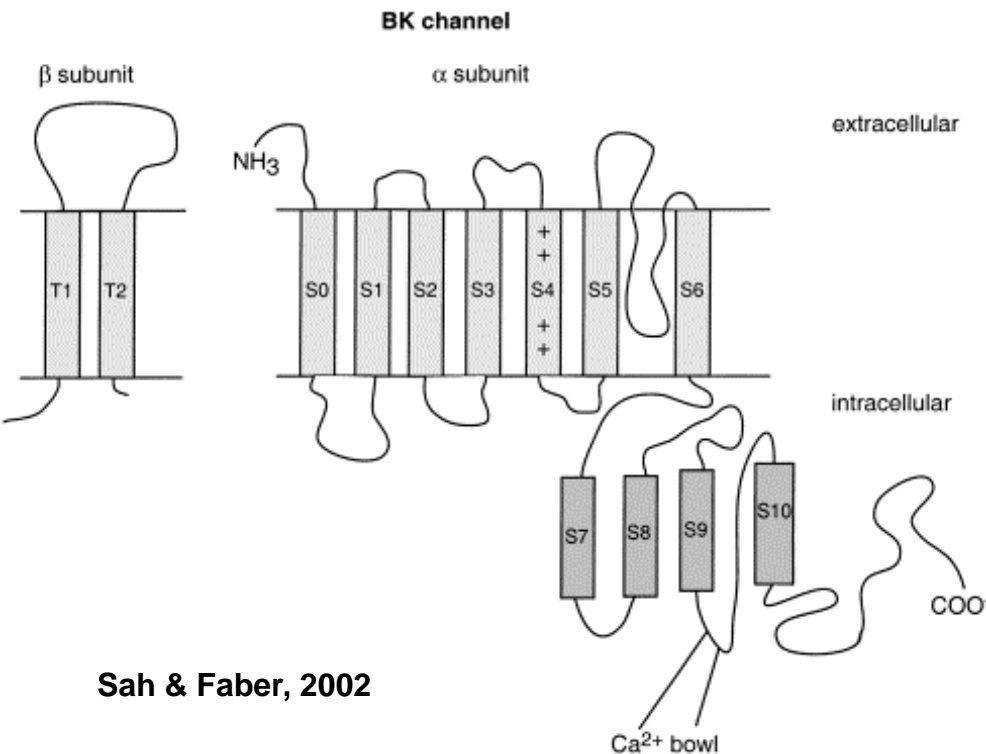
- K<sub>Ca</sub>4.1, K<sub>Ca</sub>4.2 are gated by depolarization and synergistically by elevation of intracellular Na<sup>+</sup> and Cl<sup>-</sup>.

- K<sub>Ca</sub>5.1 is gated by voltage and elevation of intracellular pH. This channel is sperm-specific and controls sperm membrane potential.

- BK α subunits make homo-tetrameric and likely hetero-tetrameric complexes.

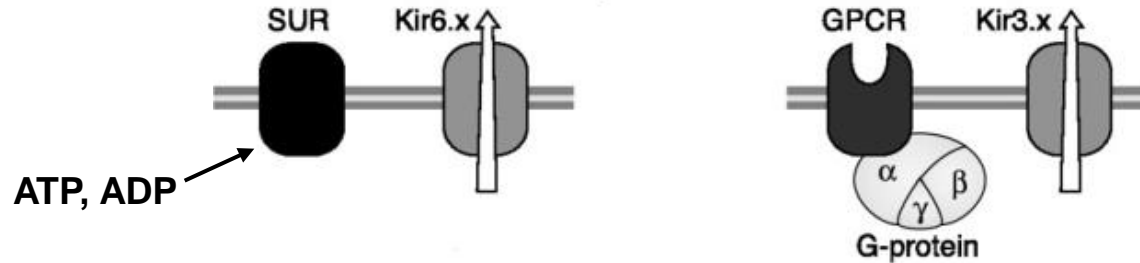
- BK channels interact with four β subunits (β1-4) that can confer a higher Ca<sup>2+</sup> sensitivity and faster inactivation on the channel.

- S0 transmembrane domain of BK channels is important for interaction with the β subunit.



# $K_{ir}$ , inwardly rectifying $K^+$ channels

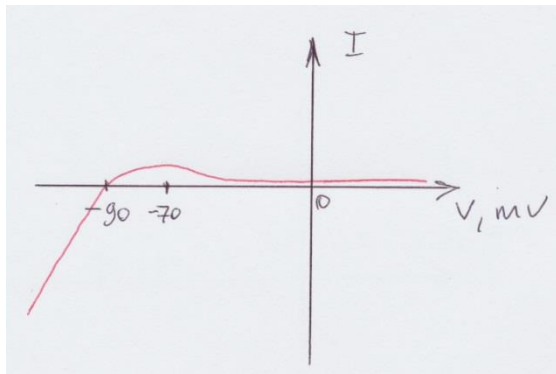
Gating: constitutively active ( $K_{ir}1.1$ ,  $K_{ir}4$  subfamily,  $K_{ir}5.1$ ), opened at voltages negative to  $E_K$  and closed at voltages positive to  $E_K$  ( $K_{ir}2$  subfamily,  $K_{ir}7.1$ ), opened by  $G_{\beta\gamma}$  subunits ( $K_{ir}3$  subfamily), opened by ADP ( $K_{ir}6$  subfamily), closed by intracellular ATP ( $K_{ir}6.2$ ).



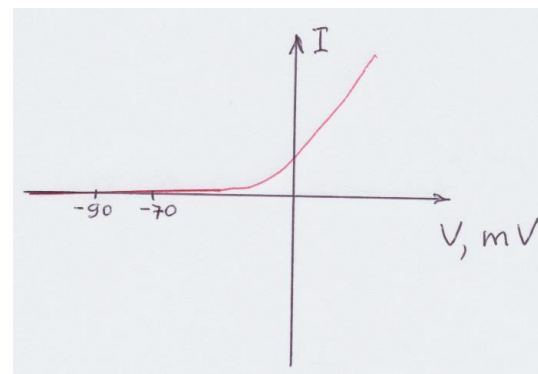
Location: plasma membrane of excitable and non-excitable cells

Function: as a generalization,  $K_{ir}$  help maintain the resting membrane potential and control excitability of neurons and muscle cells; transepithelial  $K^+$  transport ( $K_{ir}1.1$ ,  $K_{ir}7.1$ ), G protein coupled (serotonin, metabotropic glutamate, etc.) receptor-dependent membrane hyperpolarization, regulation of insulin secretion in pancreatic  $\beta$ -cells ( $K_{ir}6.2$ ), **oxygen and glucose sensor in brain ( $K_{ir}6.2$ )**, cytoprotection during cardiac and brain ischemia ( $K_{ir}6.2$ ).

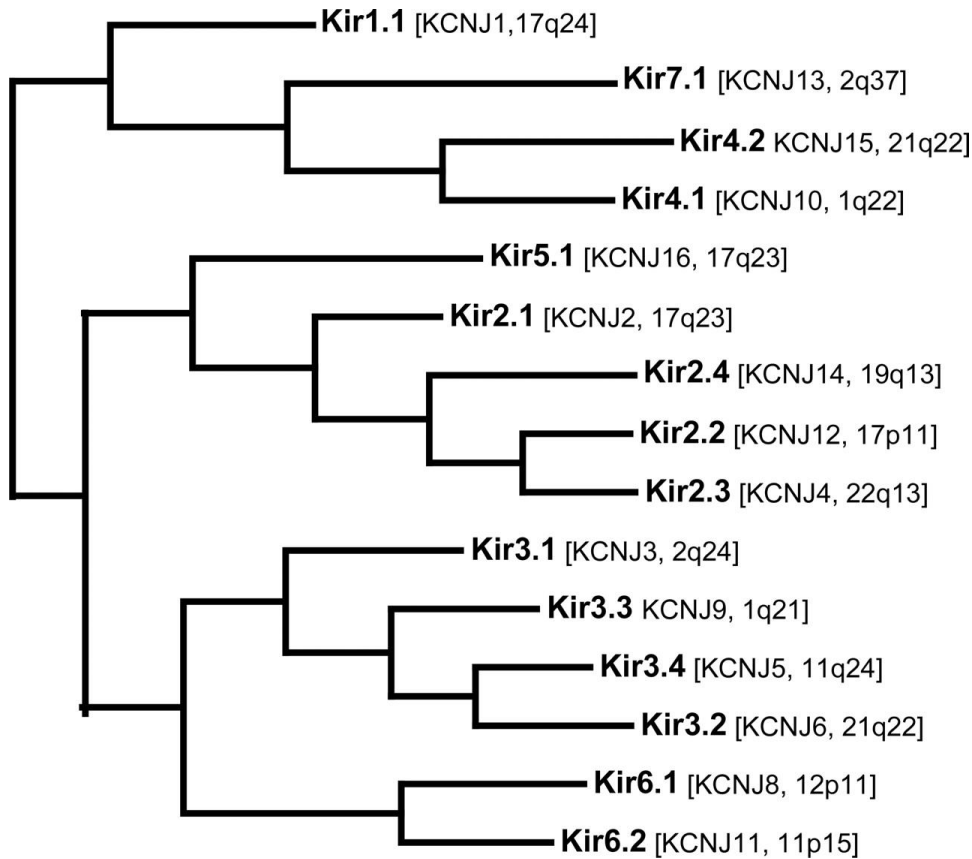
Inwardly rectifying  $K^+$  channel



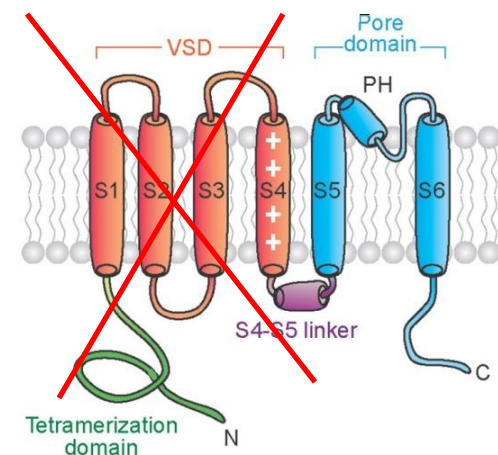
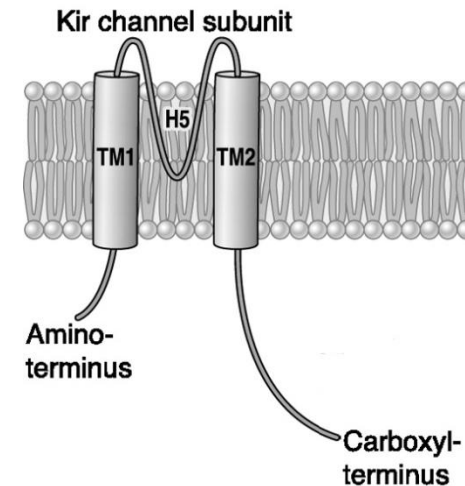
Outwardly rectifying  $K^+$  channel



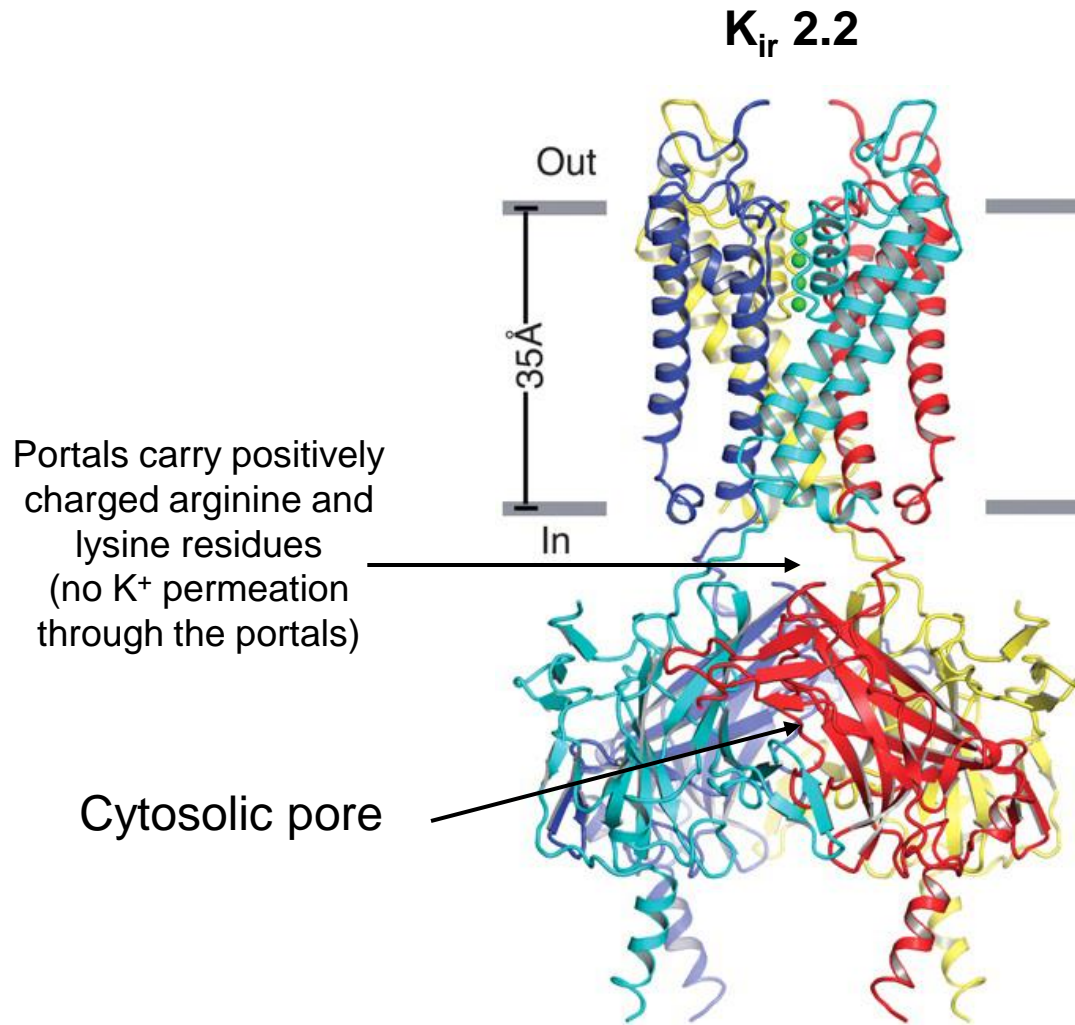
# K<sub>ir</sub>, inwardly rectifying K<sup>+</sup> channels



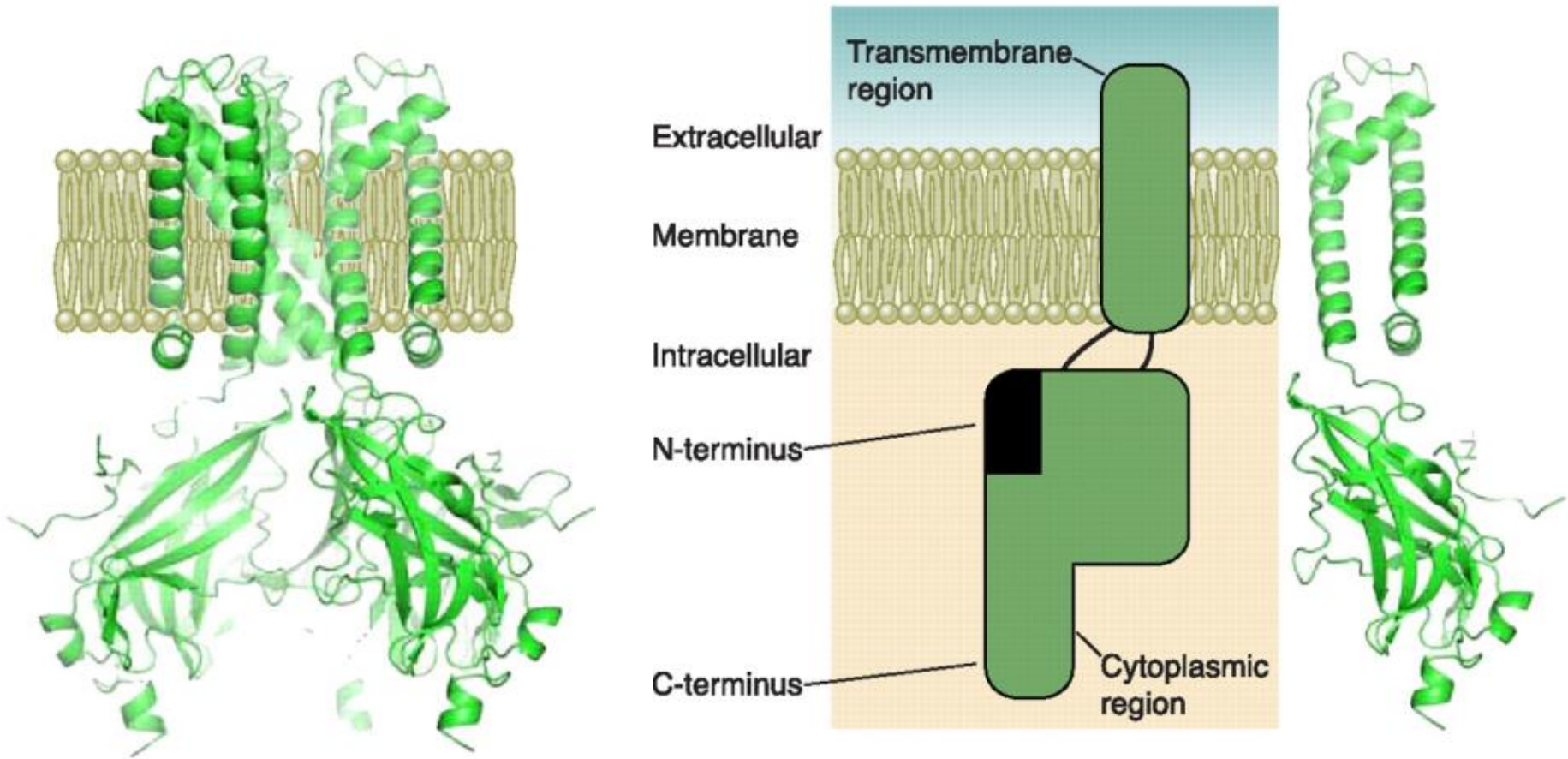
- K<sub>ir</sub> subunits form homo-tetrameric and hetero-tetrameric channels



# Structure of inwardly rectifying K<sup>+</sup> channel



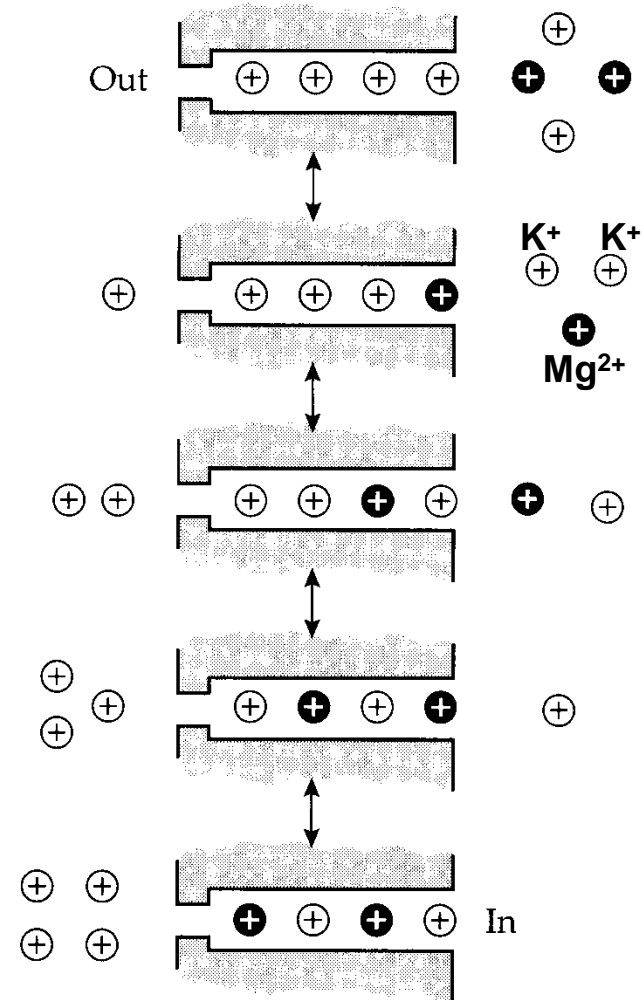
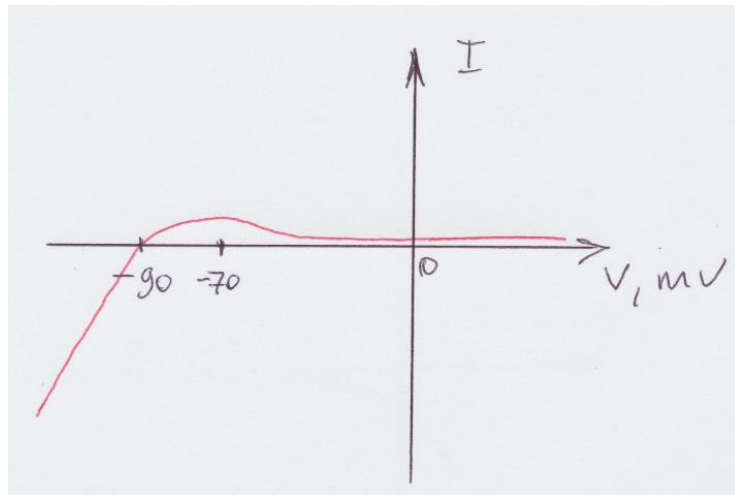
# Structure of inwardly rectifying K<sup>+</sup> channel



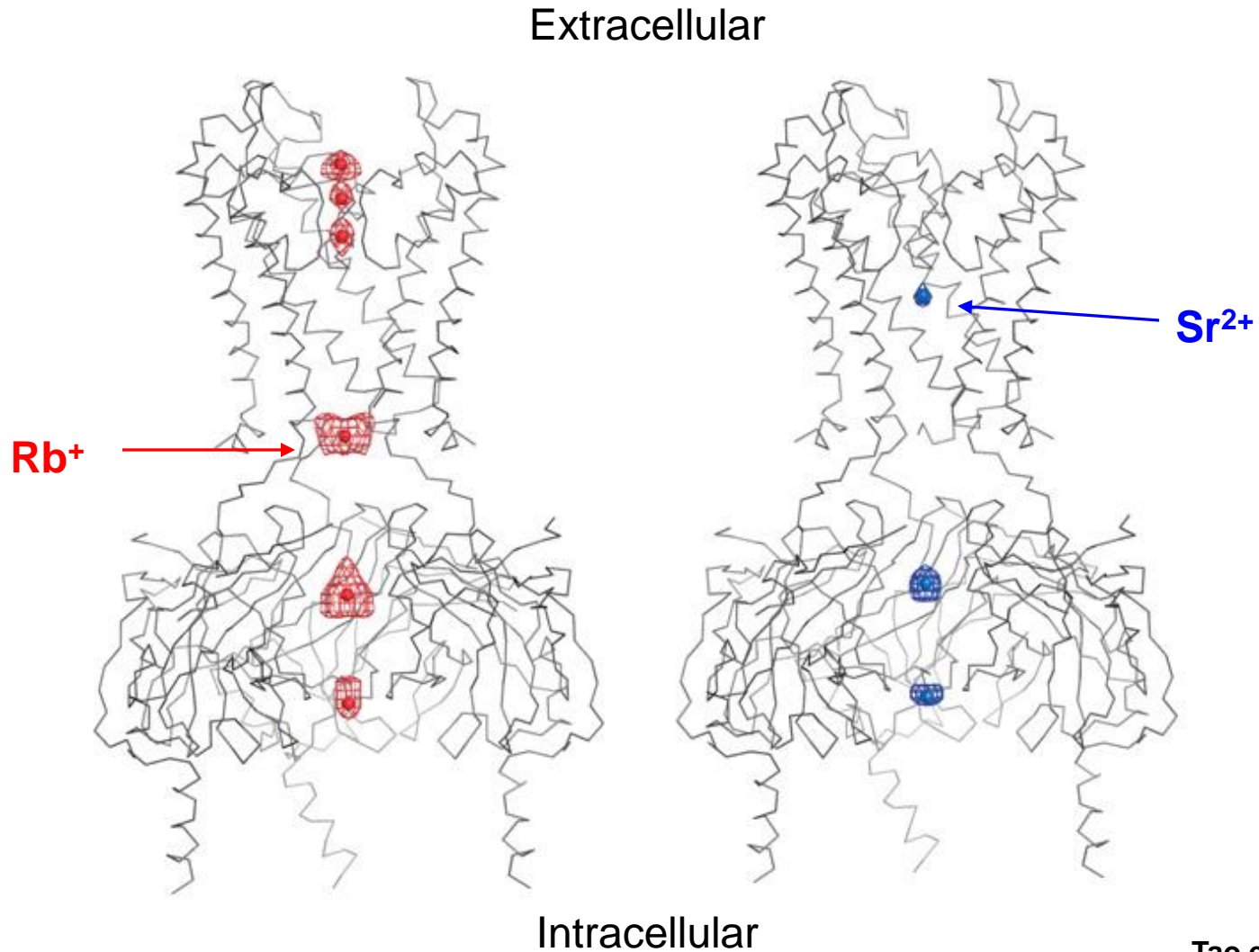
# Inward rectification of $K_{ir}$ channels

## 16.12 Multi-Ion Block in a Long Pore

Hypothetical occupancy states of a pore with four occupied ion binding sites. The pore is permeable to the white cation, but the selectivity filter is too narrow to pass the black ion. Current from right to left will draw the black ion from the cytoplasm into the pore, blocking further flow. Current from left to right draws permeant ions from the outside through the selectivity filter and clears the pore.



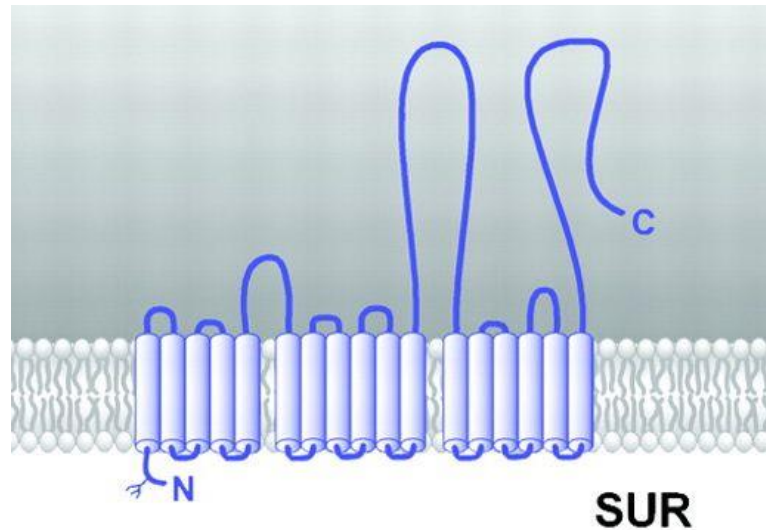
# Structural basis of inward rectification of $K_{ir}$ channels



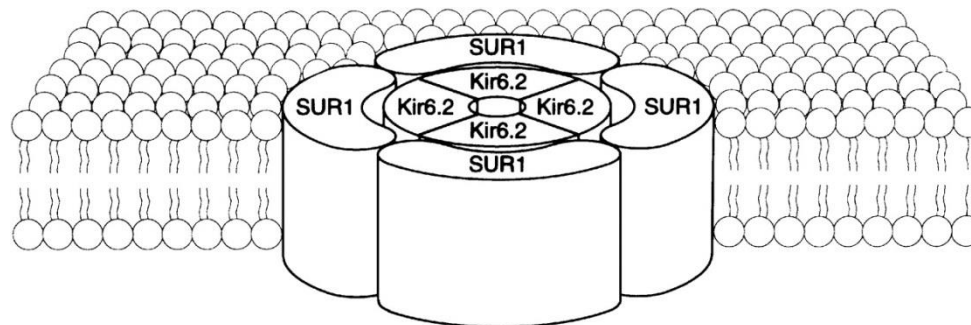
# $K_{ir}$ , inwardly rectifying $K^+$ channels

Sulfonylurea receptors (SUR1 and SUR2) confer ATP/ADP sensitivity on the  $K_{ir6}$  ion channels.

Auxiliary subunit of  $K_{ir6}$  subfamily



Modified from Yu et al, 2005



Seino, 1999