

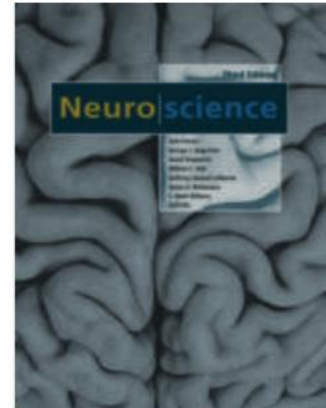
# Voltage Gated Ion Channels

The Machines  
That Make It  
Possible...

Topics I	Topics II
Introduction & Electrochemical Gradients	Synaptic Transmission
Passive Membrane Properties	Electrophysiology Techniques
Action Potentials	Basic Circuits (Spinal Cord)
Voltage-Gated Ion Channels	Sensory Systems Overview
Ligand-Gated Ion Channels	Synaptic Plasticity

# Study Material

- NEUROSCIENCE Third Edition
  - Dale Purves
- Chapter 4 pages 69-85



**THE COVER**  
Dorsal view of the human brain.  
(Courtesy of S. Mark Williams.)

NEUROSCIENCE: Third Edition  
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# Aims for this Lecture

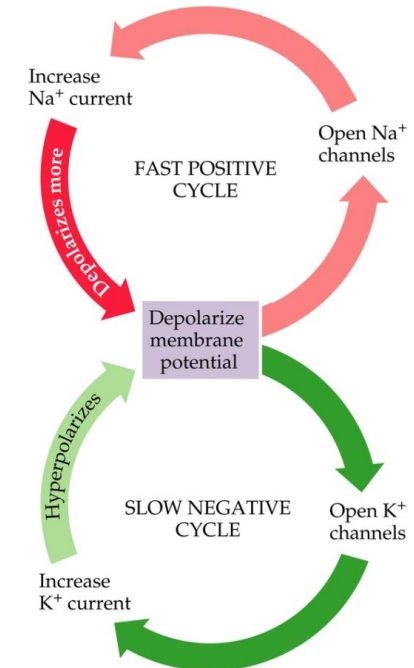
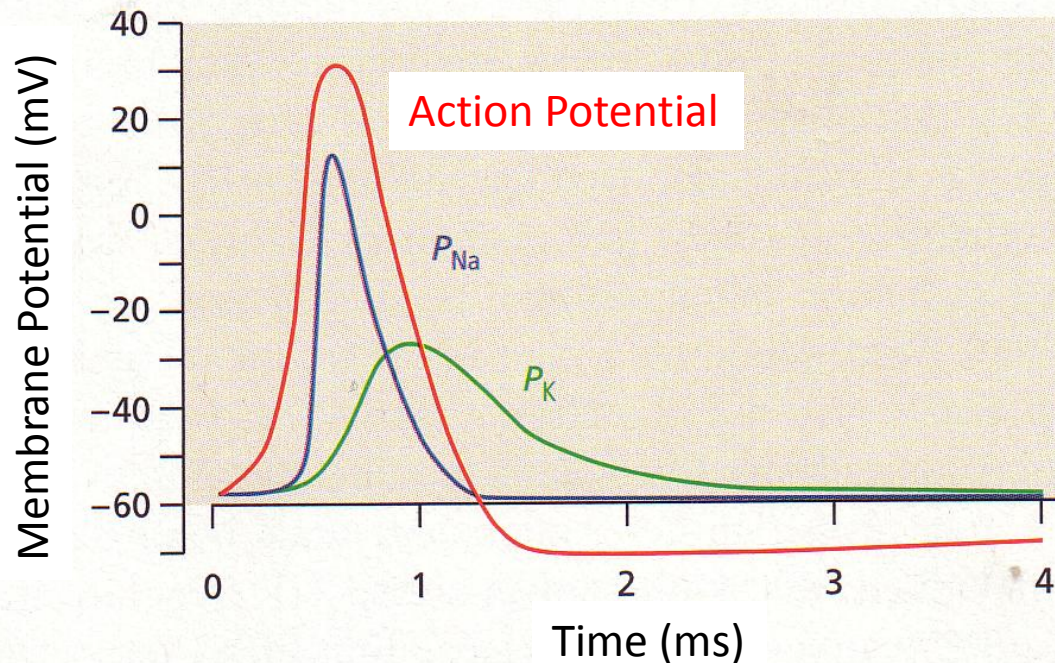
- Get an idea of the molecular structure of voltage-gated ion channels.
- Understand mechanisms for ion selectivity.
- Understand channel opening and closing and its stochastic nature.
- Understand activation and inactivation curves.

# Recapitulation L4

- Action potentials are fast stereotypical signals produced by neurons when depolarized above a threshold.
- Na conductances open and let Na flow into the neuron. The neuron further depolarizes. The Na conductances rapidly inactivate.
- K conductances open with a delay and cause an outflow of K ions and a repolarization of the neuron.

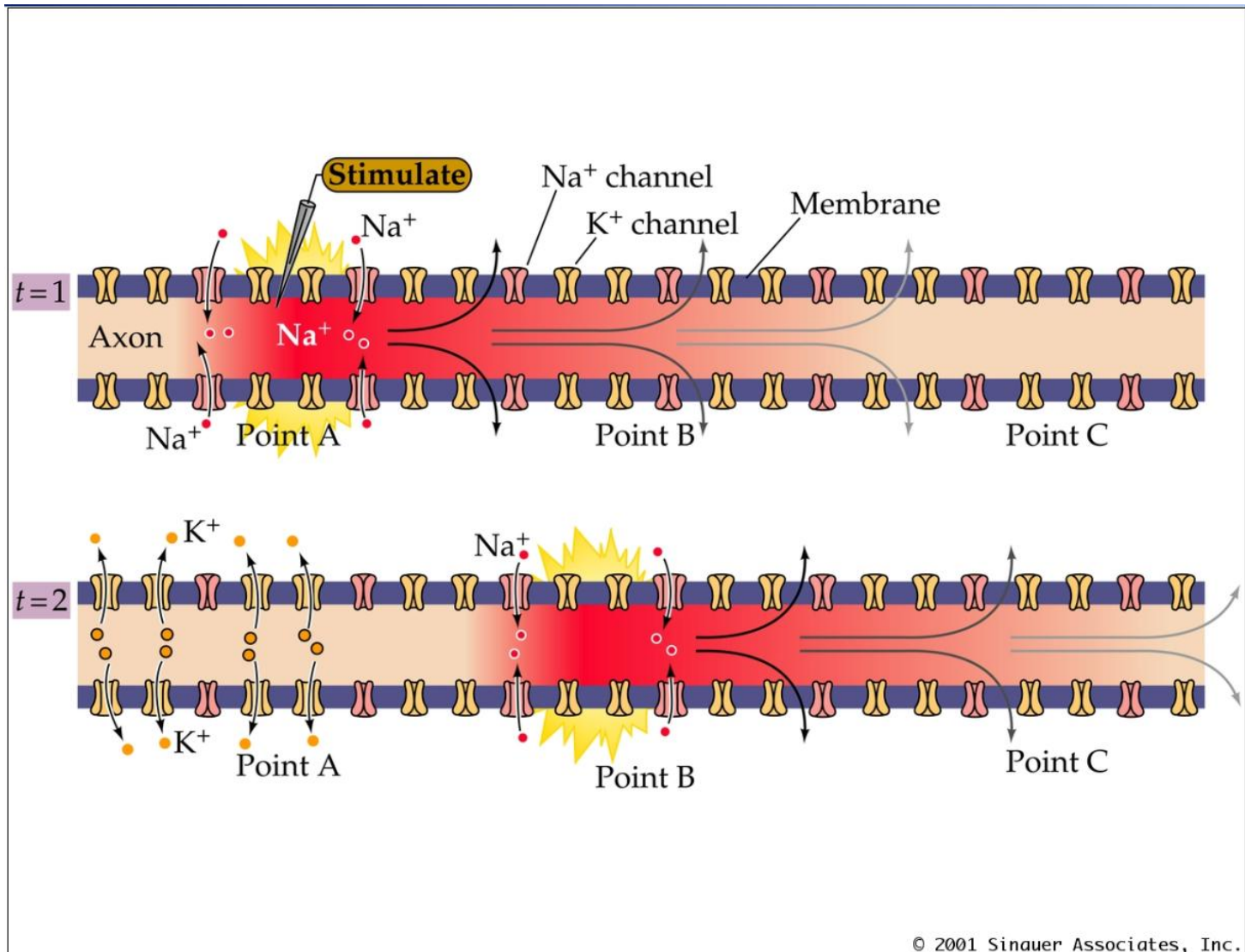
# Recapitulation L4

$$E_i = \frac{RT}{F} \ln \left( \frac{P_K [K^+]_a + P_{Na} [Na^+]_a}{P_K [K^+]_i + P_{Na} [Na^+]_i} \right)$$



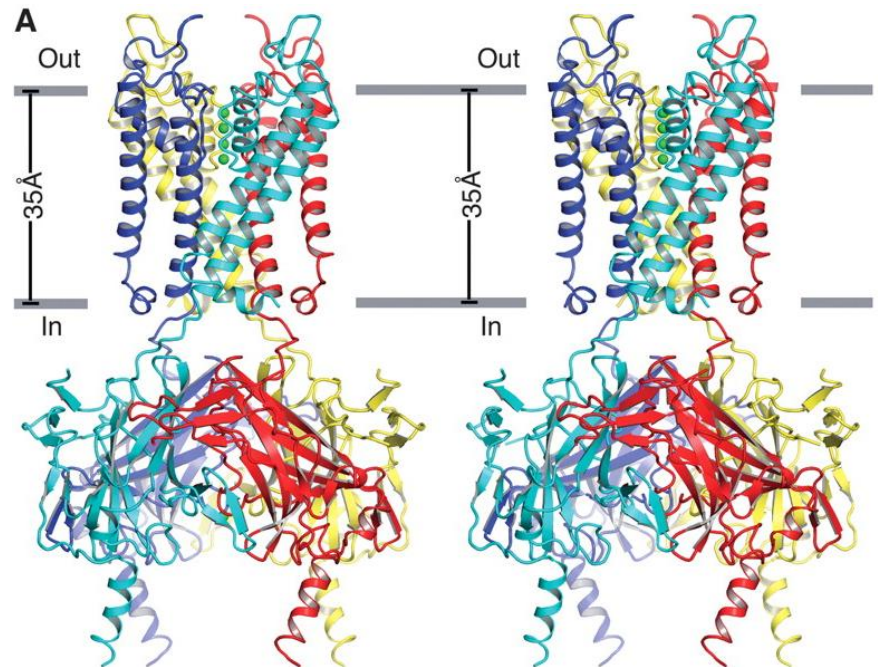
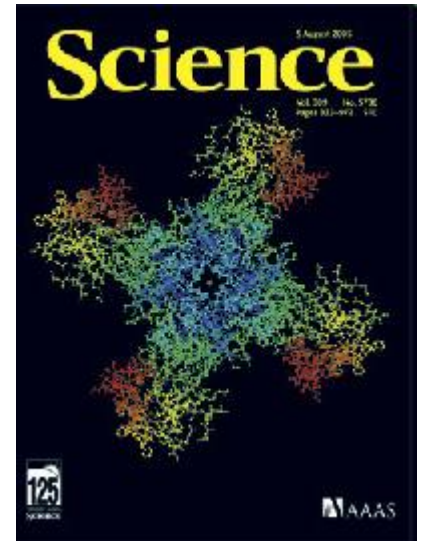
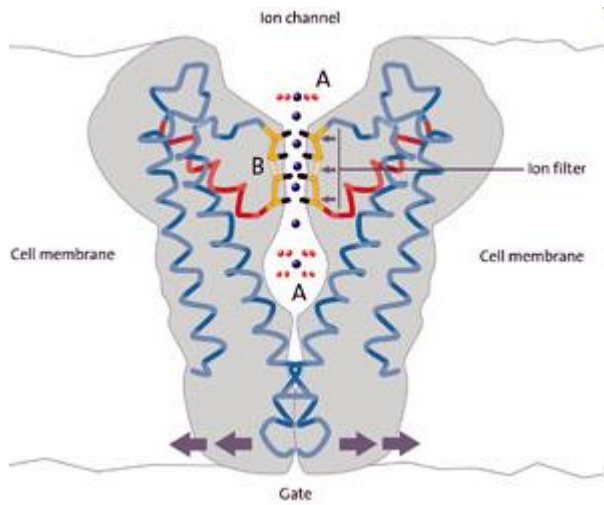
- Positive feedback of voltage dependent Na-conductance
- Negative feedback of voltage dependent K-conductance
- Explosive depolarization and subsequent repolarization..

# Recapitulation L4



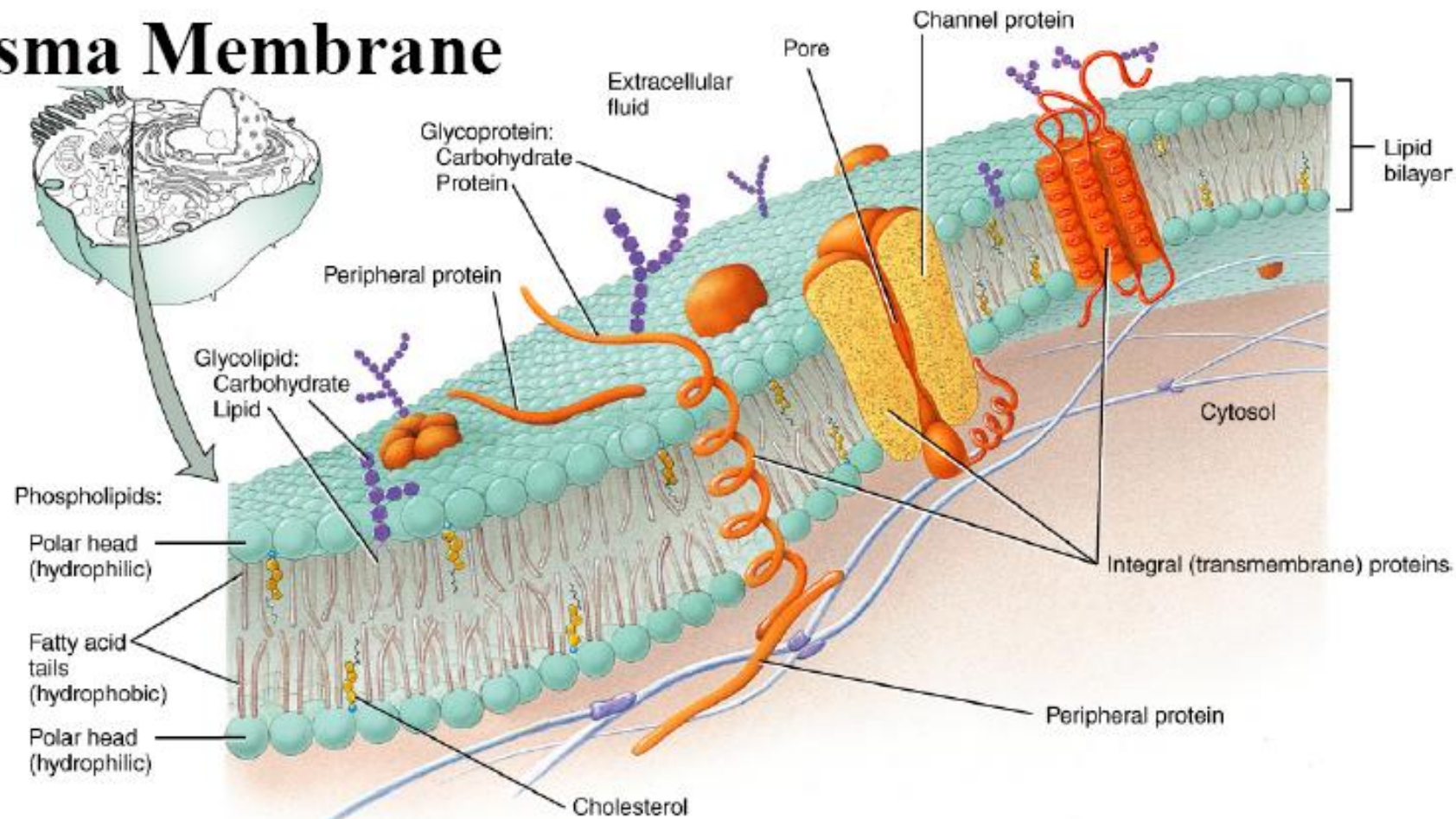


# Ion Channels



# The Cell Membrane

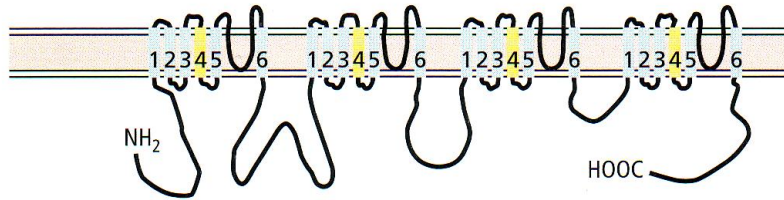
## Plasma Membrane



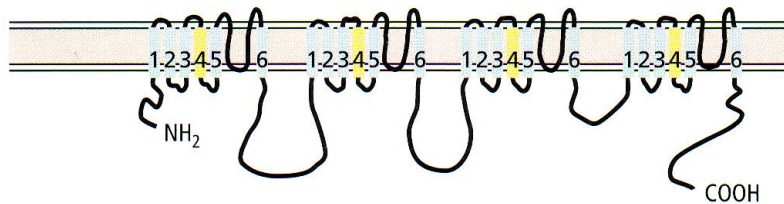


# Ion Channels - Structures

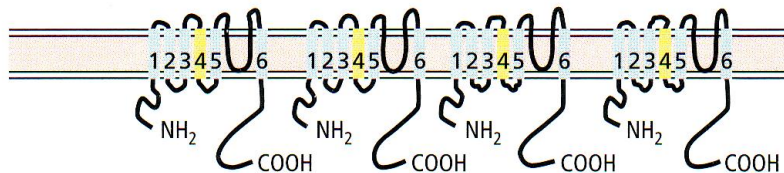
Sodium Channel



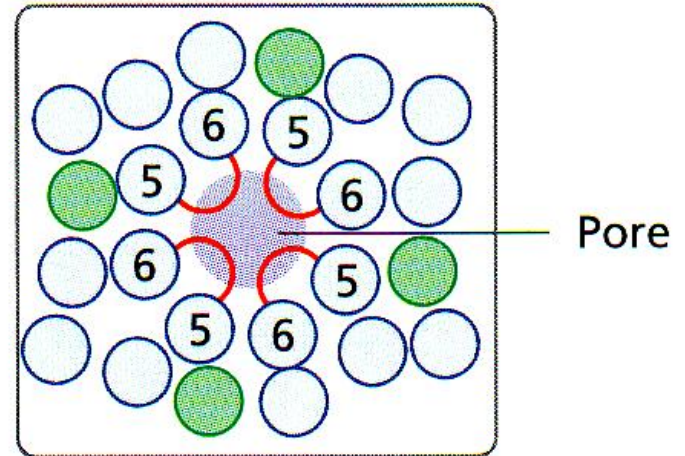
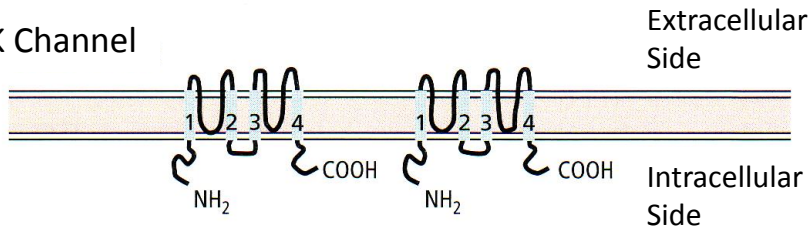
Calcium Channel



Potassium Channel

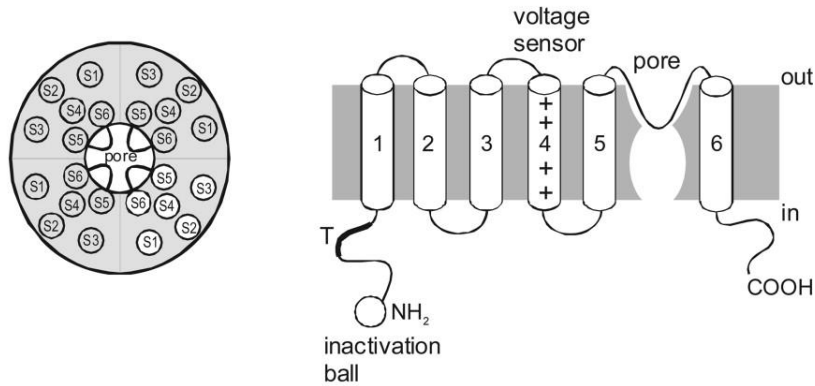


Leak K Channel

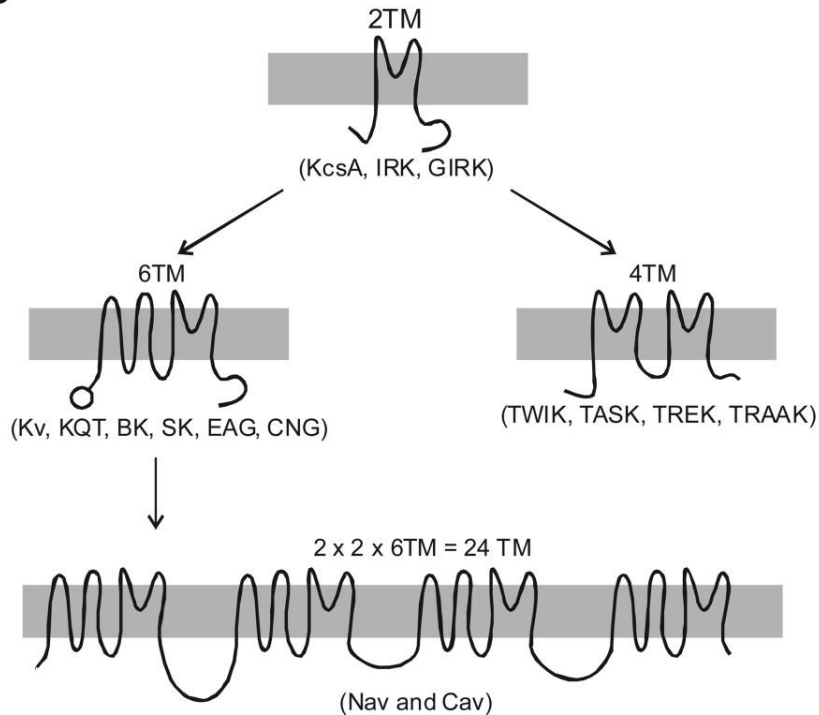


# Ion Channel Evolution

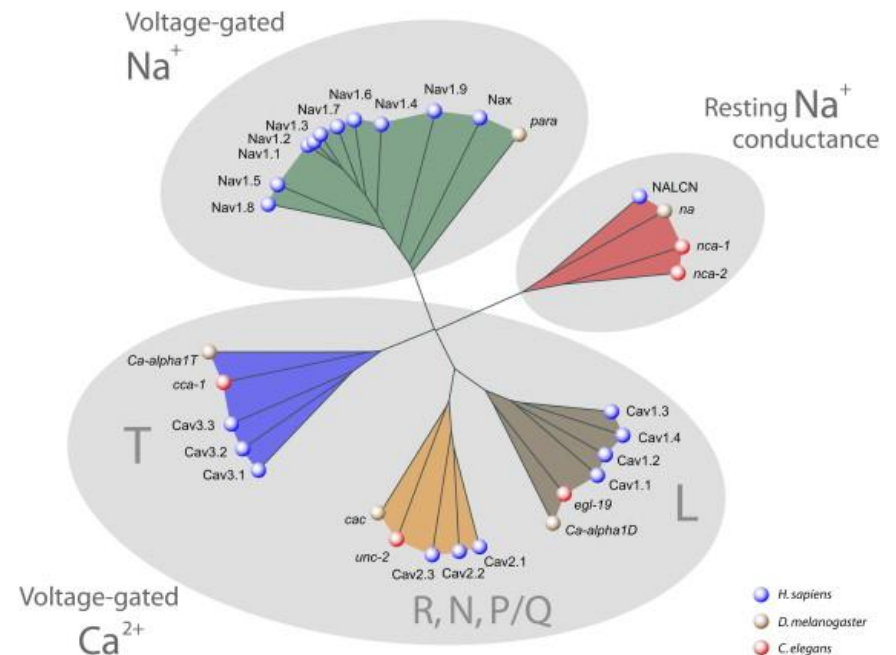
A



B



## Four-Domain Channels



# Potassium Channels

- The most diverse and most ancient class of ion channels.
- All sequenced genomes contain at least one, most contain many genes for potassium channels.
- Voltage-gated potassium channels are members of this class of channels

# K Channels

TABLE 1  
*K<sub>v</sub> channel families*

Gene names shown are those assigned by the IUPHAR (Catterall et al., 2002) and HGNC (<http://www.gene.ucl.ac.uk>) in addition to some other commonly used names.

IUPHAR	HGNC	Other
<i>K<sub>v</sub>1.1</i>	<i>KCNA1</i>	<i>Shaker</i> -related family
<i>K<sub>v</sub>1.2</i>	<i>KCNA2</i>	
<i>K<sub>v</sub>1.3</i>	<i>KCNA3</i>	
<i>K<sub>v</sub>1.4</i>	<i>KCNA4</i>	
<i>K<sub>v</sub>1.5</i>	<i>KCNA5</i>	
<i>K<sub>v</sub>1.6</i>	<i>KCNA6</i>	<i>Shab</i> -related family
<i>K<sub>v</sub>1.7</i>	<i>KCNA7</i>	
<i>K<sub>v</sub>1.8</i>	<i>KCNA10</i>	
<i>K<sub>v</sub>2.1</i>	<i>KCNB1</i>	
<i>K<sub>v</sub>2.2</i>	<i>KCNB2</i>	
<i>K<sub>v</sub>3.1</i>	<i>KCNC1</i>	<i>Shaw</i> -related family
<i>K<sub>v</sub>3.2</i>	<i>KCNC2</i>	
<i>K<sub>v</sub>3.3</i>	<i>KCNC3</i>	
<i>K<sub>v</sub>3.4</i>	<i>KCNC4</i>	
<i>K<sub>v</sub>4.1</i>	<i>KCND1</i>	<i>Shal</i> -related family
<i>K<sub>v</sub>4.2</i>	<i>KCND2</i>	
<i>K<sub>v</sub>4.3</i>	<i>KCND3</i>	
<i>K<sub>v</sub>5.1</i>	<i>KCNF1</i>	Modifier
<i>K<sub>v</sub>6.1</i>	<i>KCNG1</i>	Modifiers
<i>K<sub>v</sub>6.2</i>	<i>KCNG2</i>	
<i>K<sub>v</sub>6.3</i>	<i>KCNG3</i>	
<i>K<sub>v</sub>6.4</i>	<i>KCNG4</i>	
<i>K<sub>v</sub>7.1</i>	<i>KCNQ1</i>	<i>KVLQT</i>
<i>K<sub>v</sub>7.2</i>	<i>KCNQ2</i>	
<i>K<sub>v</sub>7.3</i>	<i>KCNQ3</i>	<i>EQT2</i>
<i>K<sub>v</sub>7.4</i>	<i>KCNQ4</i>	
<i>K<sub>v</sub>7.5</i>	<i>KCNQ5</i>	
<i>K<sub>v</sub>8.1</i>	<i>KCNV1</i>	Modifiers
<i>K<sub>v</sub>8.2</i>	<i>KCNV2</i>	
<i>K<sub>v</sub>9.1</i>	<i>KCNS1</i>	Modifiers
<i>K<sub>v</sub>9.2</i>	<i>KCNS2</i>	
<i>K<sub>v</sub>9.3</i>	<i>KCNS3</i>	
<i>K<sub>v</sub>10.1</i>	<i>KCNH1</i>	<i>eag1</i>
<i>K<sub>v</sub>10.2</i>	<i>KCNH5</i>	<i>eag2</i>
<i>K<sub>v</sub>11.1</i>	<i>KCNH2</i>	<i>erg1</i>
<i>K<sub>v</sub>11.2</i>	<i>KCNH6</i>	<i>erg2</i>
<i>K<sub>v</sub>11.3</i>	<i>KCNH7</i>	<i>erg3</i>
<i>K<sub>v</sub>12.1</i>	<i>KCNH8</i>	<i>elk1, elk3</i>
<i>K<sub>v</sub>12.2</i>	<i>KCNH3</i>	<i>elk2</i>
<i>K<sub>v</sub>12.3</i>	<i>KCNH4</i>	<i>elk1</i>

TABLE 1  
*K<sub>Ca</sub> channels*

IUPHAR names of the members of the K<sub>Ca</sub> group of potassium channels are shown, together with their HGNC designations and other commonly used names.

IUPHAR	HGNC	Other
<i>K<sub>Ca</sub>1.1</i>	<i>KCNMA1</i>	Slo, Slo1, BK
<i>K<sub>Ca</sub>2.1</i>	<i>KCNN1</i>	SK <sub>Ca</sub> 1
<i>K<sub>Ca</sub>2.2</i>	<i>KCNN2</i>	SK <sub>Ca</sub> 2
<i>K<sub>Ca</sub>2.3</i>	<i>KCNN3</i>	SK <sub>Ca</sub> 3
<i>K<sub>Ca</sub>3.1</i>	<i>KCNN4</i>	IK <sub>Ca</sub> 1
<i>K<sub>Ca</sub>4.1</i>	<i>KCNT1</i>	Slack, Slo2.2
<i>K<sub>Ca</sub>4.2</i>	<i>KCNT2</i>	Slick, Slo2.1
<i>K<sub>Ca</sub>5.1</i>	<i>KCNU1</i>	Slo3

BK, big-conductance K<sup>+</sup> channel; SK, small-conductance K<sup>+</sup> channel; IK, intermediate-conductance K<sup>+</sup> channel.

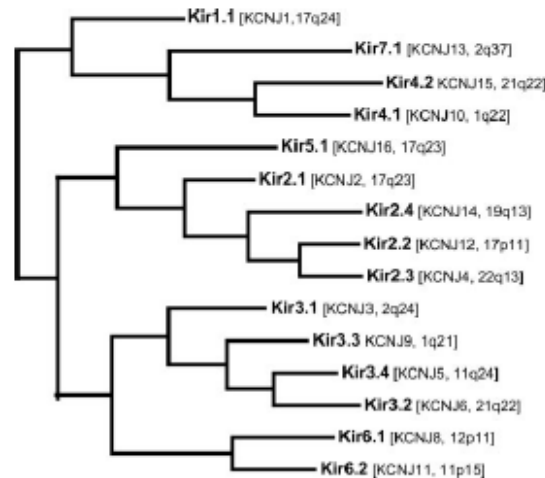


FIG. 1. Phylogenetic tree of K<sub>v</sub> channels. Amino acid sequence alignments and phylogenetic analysis for the 15 known members of the human K<sub>v</sub> family were generated as described in the legend for Fig. 1 of "LIII. Nomenclature and Molecular Relationships of Voltage-Gated Potassium Channels". No new channels have been added to this topology since it appeared in the earlier edition of this compendium. International Union of Pharmacology and HUGO Gene Nomenclature Committee names of the genes are shown together with their chromosomal localization.

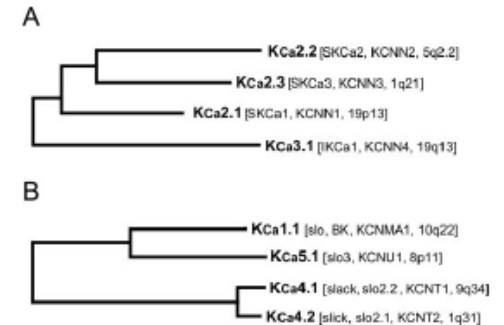
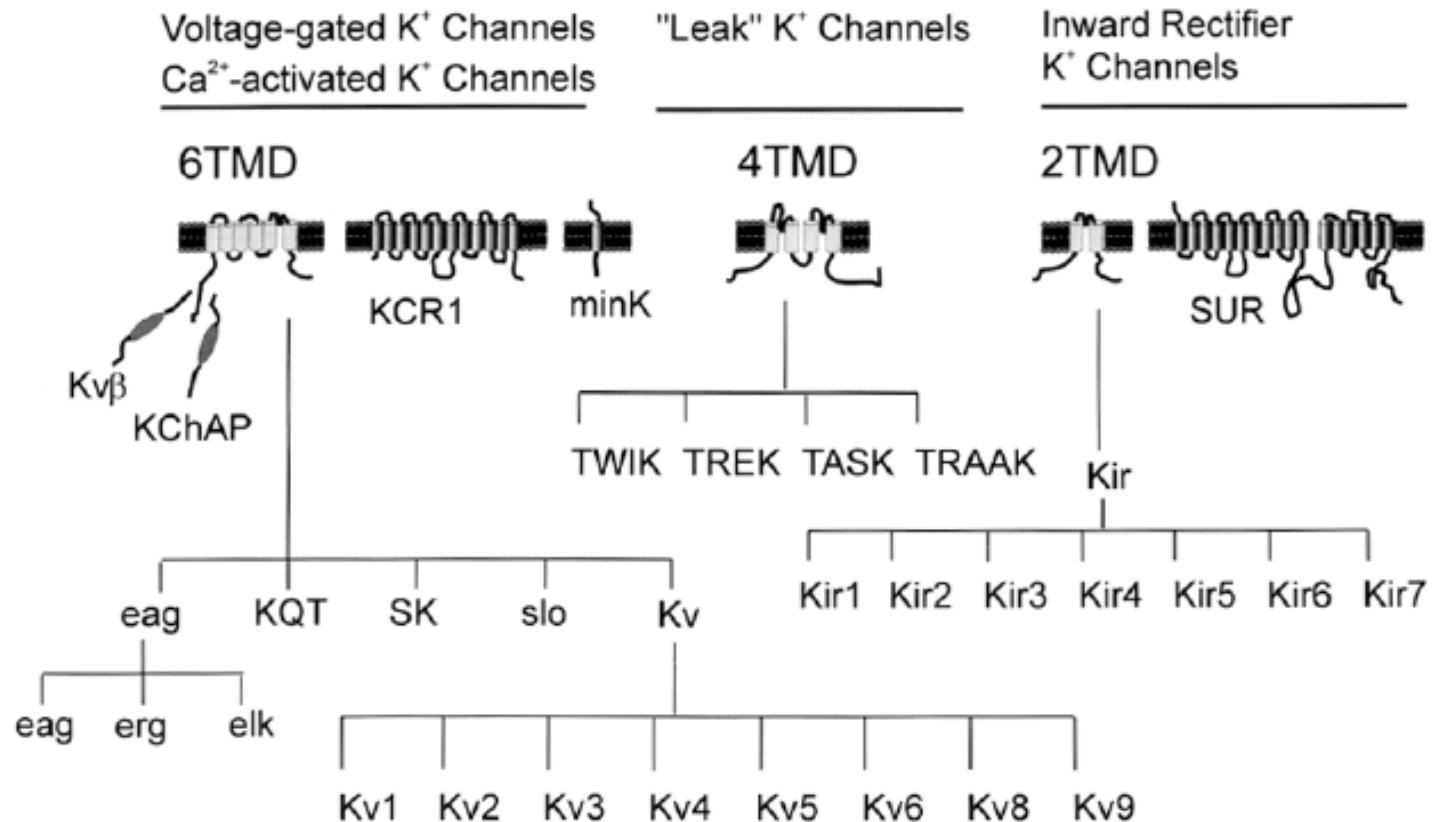


FIG. 1. Phylogenetic tree for K<sub>Ca</sub> channels. A, K<sub>Ca</sub>2/3 group. B, K<sub>Ca</sub>1/4/5 group. Amino acid sequence alignments and phylogenetic analysis for these two groups of four human K<sub>Ca</sub> channels were generated as described in the legend for Fig. 1 of "International Union of Pharmacology LIII. Nomenclature and Molecular Relationships of Voltage-Gated Potassium Channels". No new channels have been added to these topologies since they appeared in the earlier edition of this compendium. IUPHAR and HGNC names of the genes are shown together with other commonly used names and their chromosomal localization.

Pharmacol Rev. 2005 Dec;57(4):463-72.  
Pharmacol Rev. 2005 Dec;57(4):473-508.  
Pharmacol Rev. 2005 Dec;57(4):509-26.

Found in all fully sequenced genomes (Archaea, Eubacteria, Eucaryotes).  
70 genes in vertebrates code for potassium channels.  
Other channels probably evolved from potassium channels.

# Potassium Channel Classes

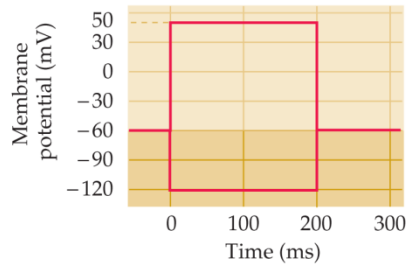


## Molecular Diversity of K<sup>+</sup> Channels

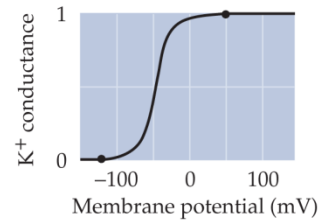
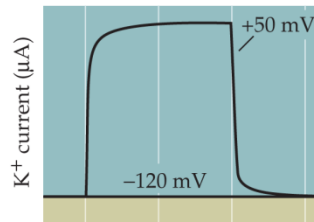
ANNALS NEW YORK ACADEMY OF SCIENCES



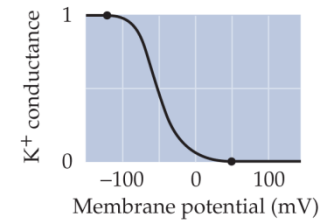
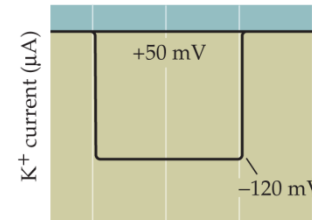
# Different K Channels



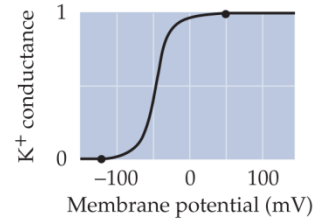
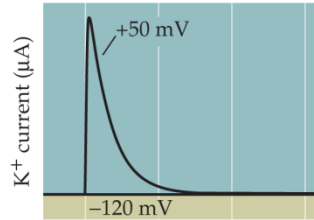
(A)  $K_{V2.1}$



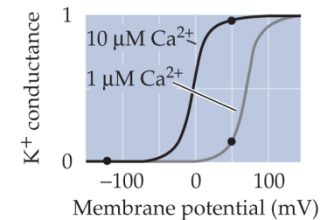
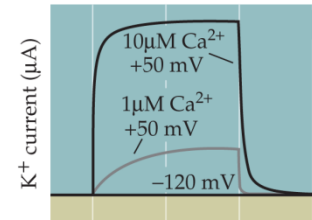
(D) Inward rectifier



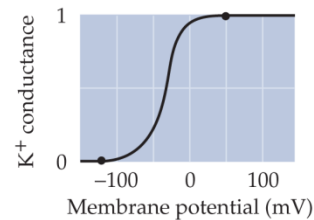
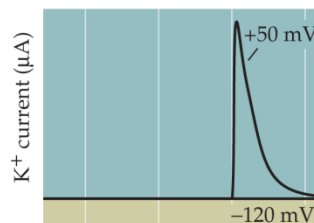
(B)  $K_{V4.1}$



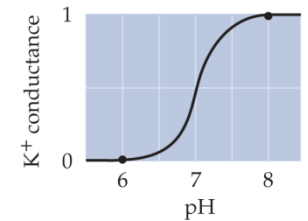
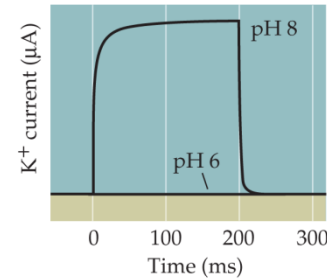
(E)  $Ca^{2+}$ -activated



(C) HERG



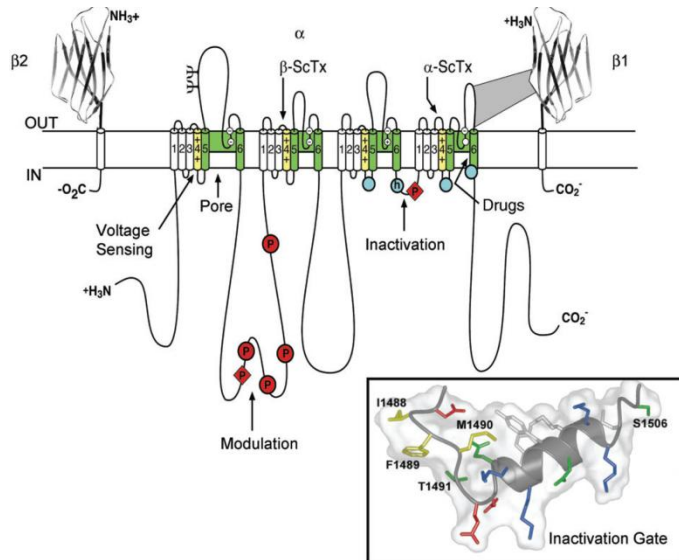
(F) 2-pore



# Voltage-Gated Sodium Channels

- Less diverse – most prominent in excitable tissue that can produce action potentials.
- Four six-transmembrane domain units strung together.
- Prominent inactivation.

# Na Channels



**Table 1. Mammalian sodium channel  $\alpha$  subunits**

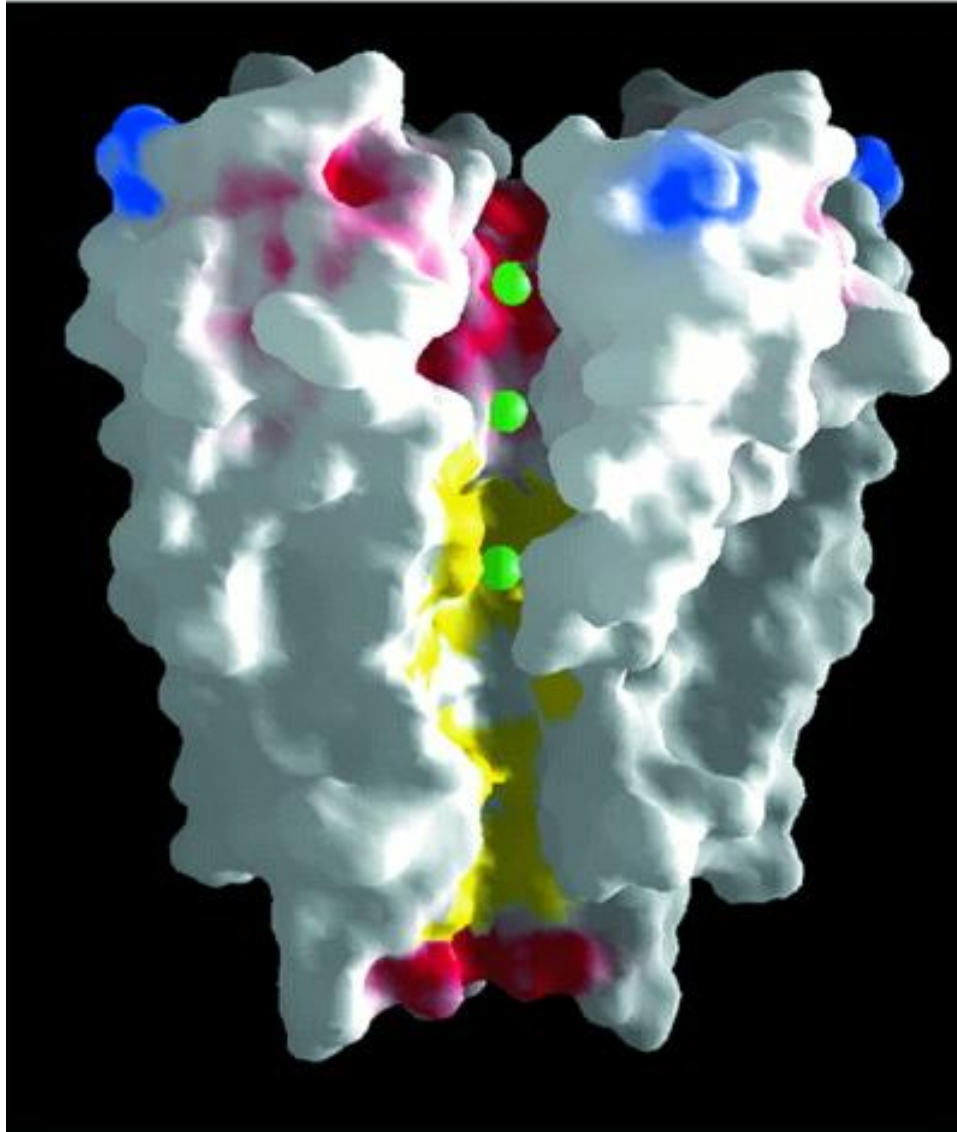
Type	Gene symbol	Chromosomal location	Primary tissues
Na <sub>v</sub> 1.1	SCN1A	Mouse 2 Human 2q24	CNS neurons
Na <sub>v</sub> 1.2	SCN2A	Mouse 2 Human 2q23–24	CNS neurons
Na <sub>v</sub> 1.3	SCN3A	Mouse 2 Human 2q24	CNS neurons
Na <sub>v</sub> 1.4	SCN4A	Mouse 11 Human 17q23–25	SkM
Na <sub>v</sub> 1.5	SCN5A	Mouse 9 Human 3p21	Uninnervated SkM, heart
Na <sub>v</sub> 1.6	SCN8A	Mouse 15 Human 12q13	CNS neurons
Na <sub>v</sub> 1.7	SCN9A	Mouse 2 Human 2q24	PNS neurons
Na <sub>v</sub> 1.8	SCN10A	Mouse 9 Human 3p22–24	DRG neurons
Na <sub>v</sub> 1.9	SCN11A	Mouse 9 Human 3p21–24	DRG neurons
Na <sub>x</sub>	SCN7A SCN6A	Mouse 2 Human 2q21–23	uterus, astrocytes, hypothalamus

TOPICAL REVIEW

## Voltage-gated sodium channels at 60: structure, function and pathophysiology

William A. Catterall

# Ion Selectivity



Blue - positive charges  
Red - negative charges  
Yellow - hydrophobic

Rough structure of a potassium channel (fourth subunit removed). Negatively charged mouth repels negatively charged ions.

Gruppen	1	2	3
Perioden	$s^1$	$s^2$	$d^1$

# Pore Structure

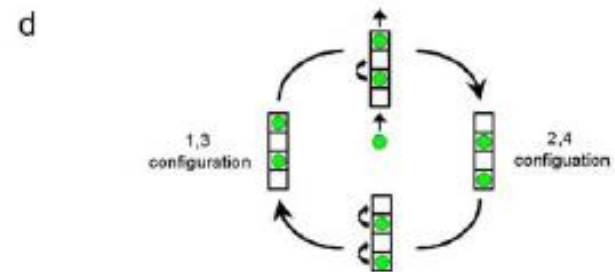
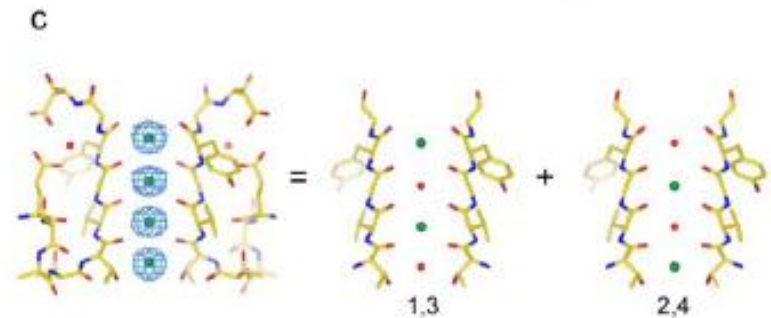
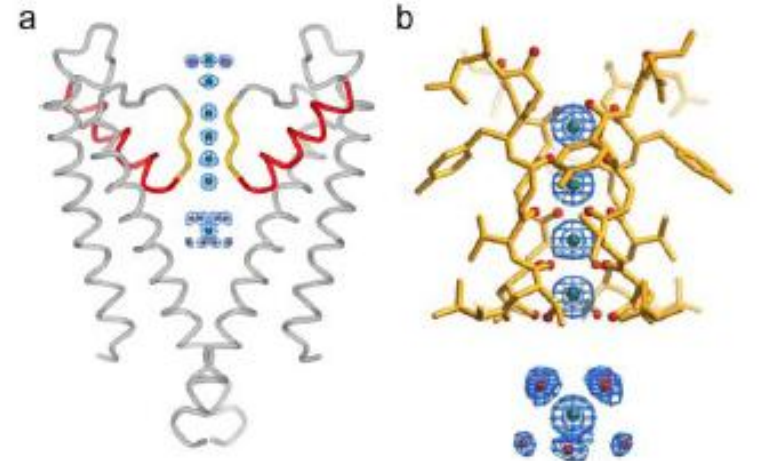
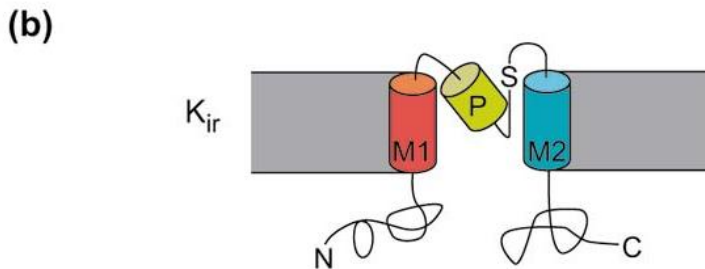
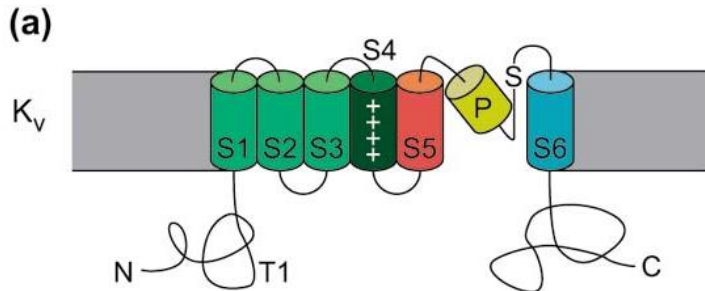
Minireview

Potassium channels

Roderick MacKinnon\*

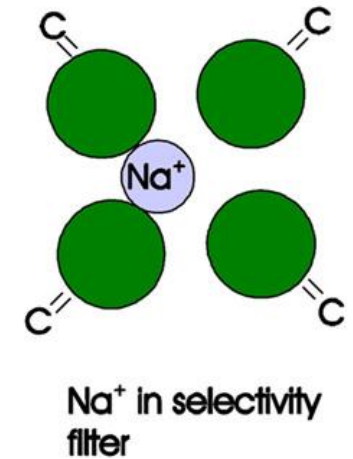
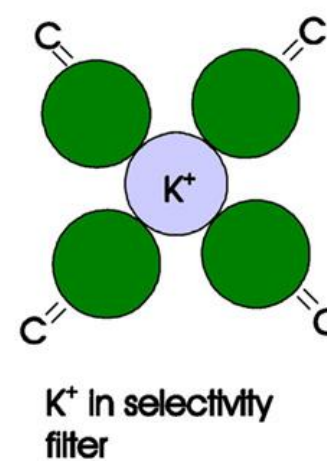
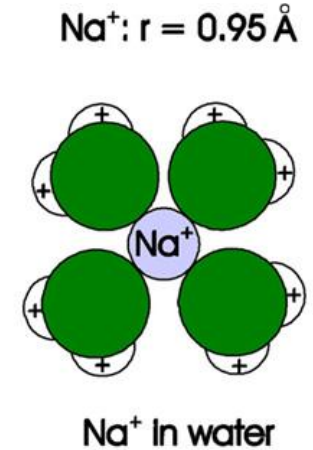
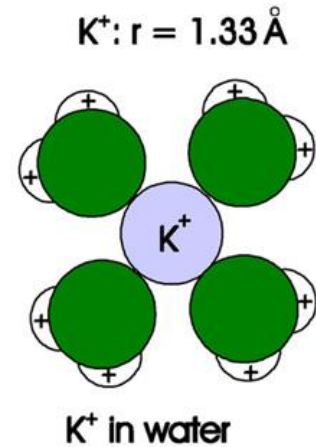
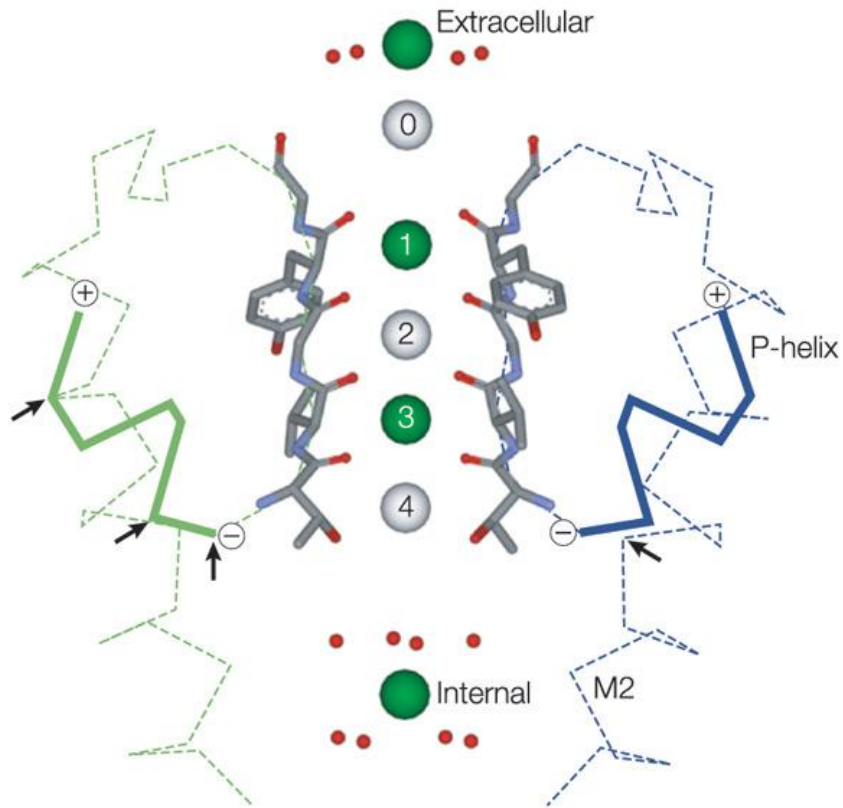
FEBS Letters 555 (2003) 62–6

1	1,00 1H		
2	6,94 3Li	9,01 4Be	
3	22,99 11Na	24,31 12Mg	←
4	39,10 19K	40,08 20Ca	44,96 21Sc
5	85,47 37Rb	87,62 38Sr	88,91 39Y
6	132,91 55Cs	137,33 56Ba	138,91 57La*
7	223,02 87Fr	226,83 88Ra	227,03 89Ac#

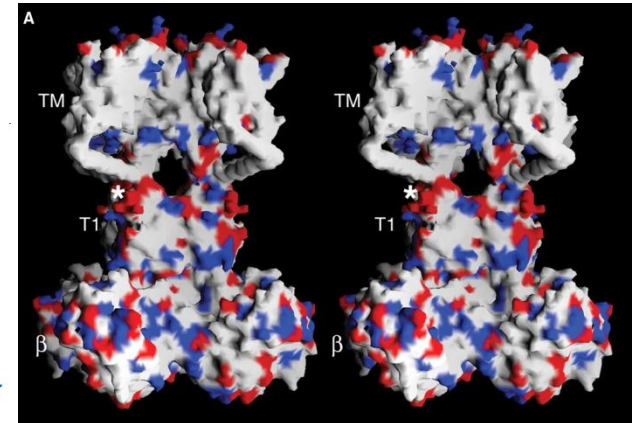
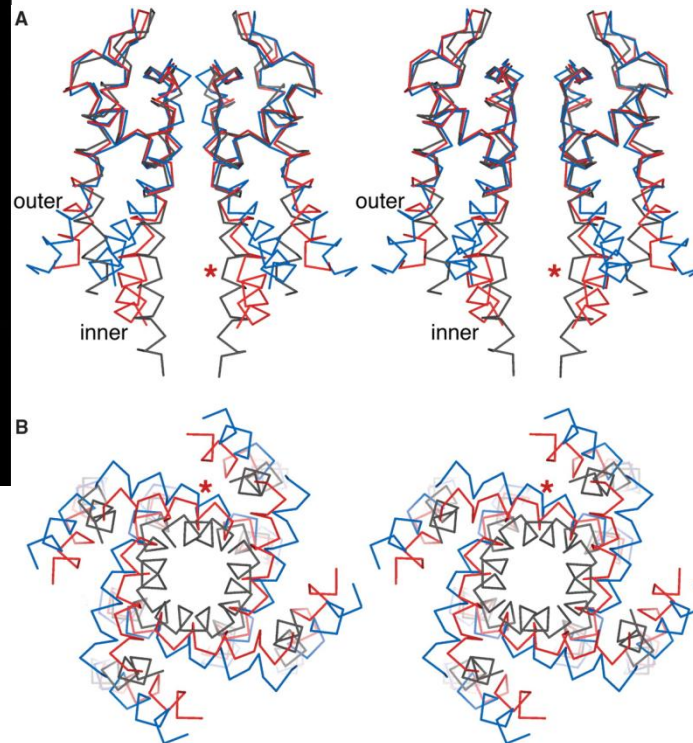
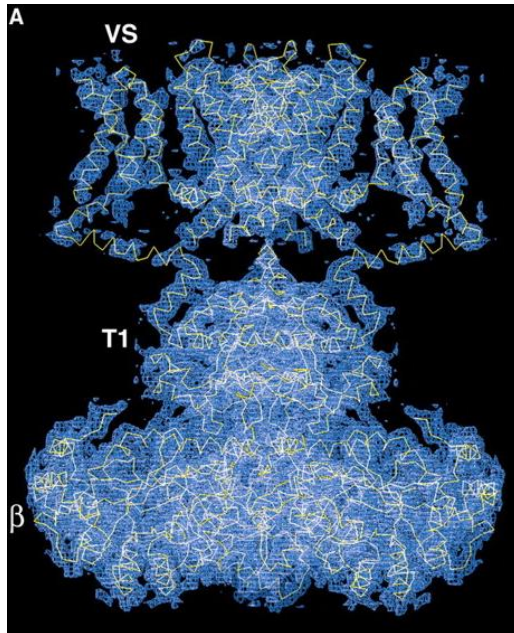




# Selectivity Filter



# Mammalian K-Channel



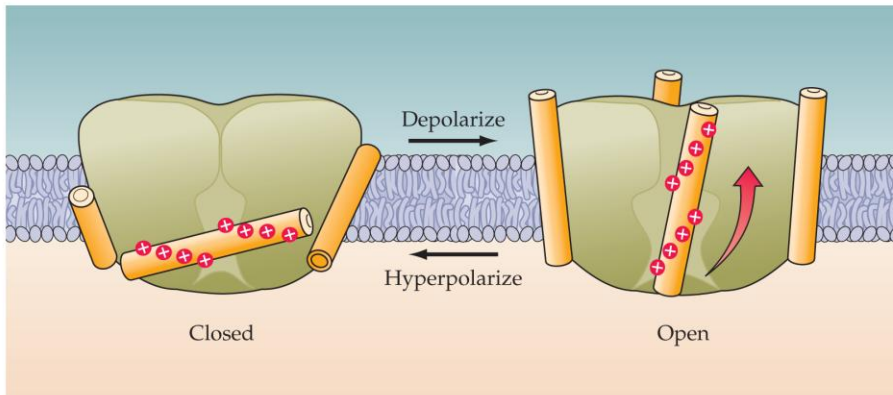
Science 5 August 2005:  
Vol. 309, no. 5736, pp. 897 - 903

**Crystal Structure of a Mammalian Voltage-Dependent *Shaker* Family K<sup>+</sup> Channel**  
Stephen B. Long, Ernest B. Campbell, Roderick MacKinnon\*

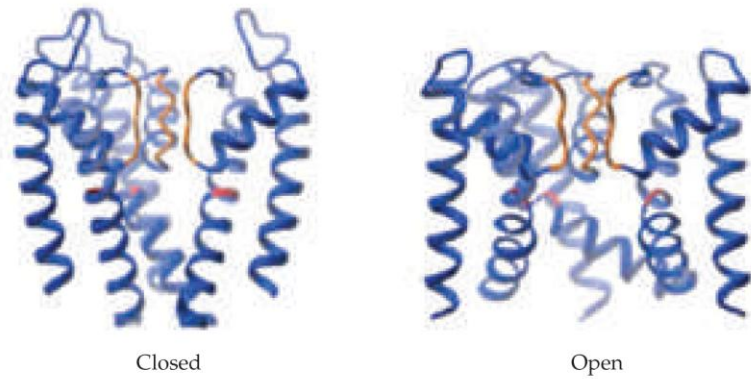
3D if watched cross-eyed

# K Channel Gating

(A)



(B)



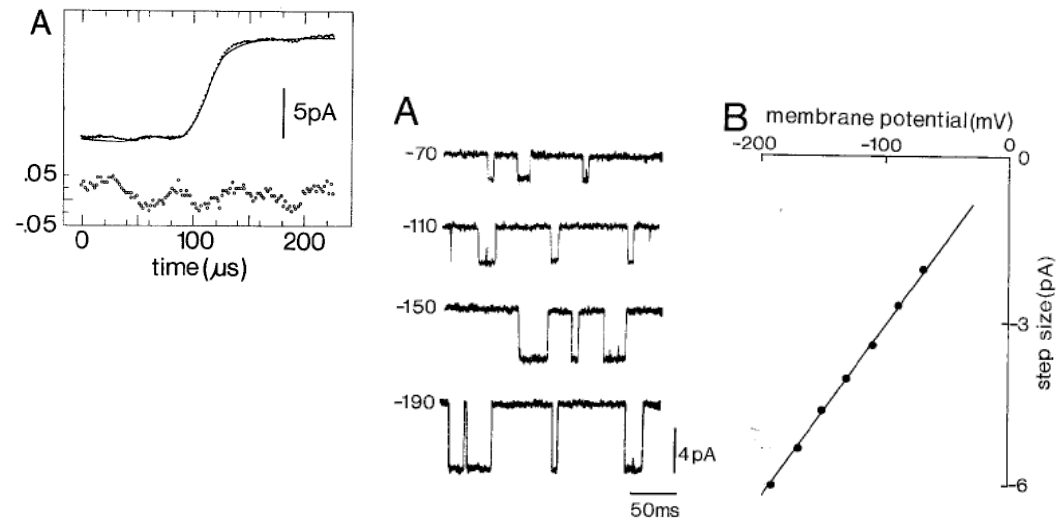
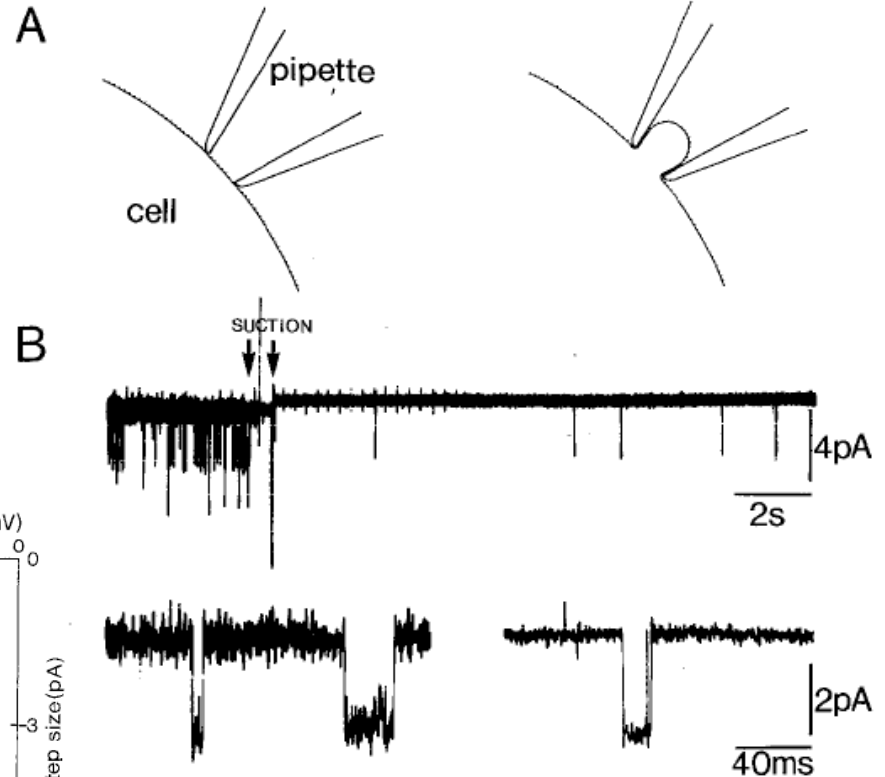
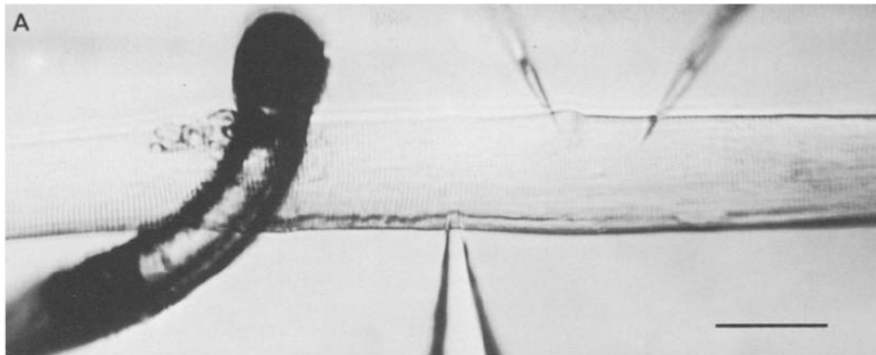
# Measuring Individual Channels

Pflügers Arch (1981) 391:85–100

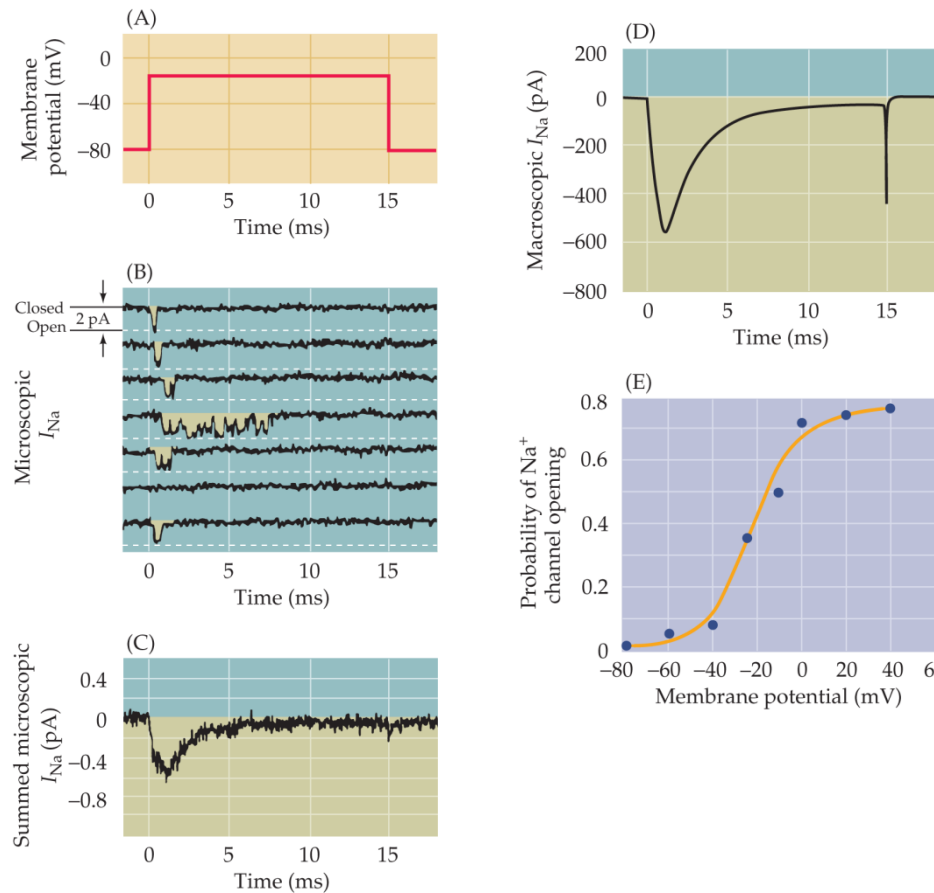
## Improved Patch-Clamp Techniques for High-Resolution Current Recording from Cells and Cell-Free Membrane Patches

O. P. Hamill, A. Marty, E. Neher, B. Sakmann, and F. J. Sigworth

Max-Planck-Institut für biophysikalische Chemie, Postfach 968, Am Fassberg, D-3400 Göttingen, Federal Republic of Germany



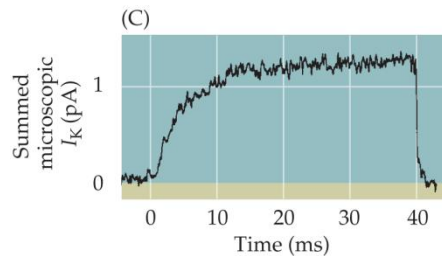
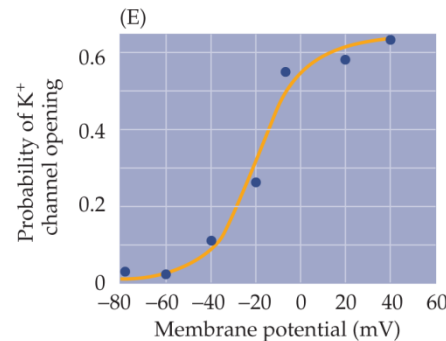
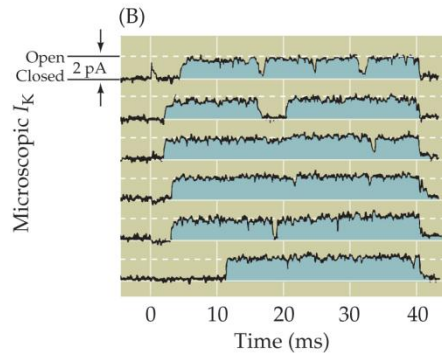
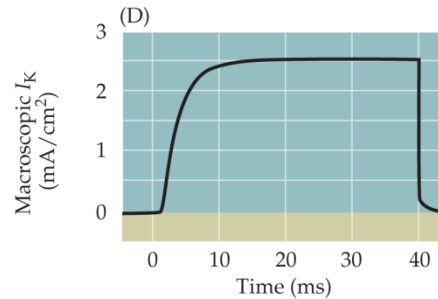
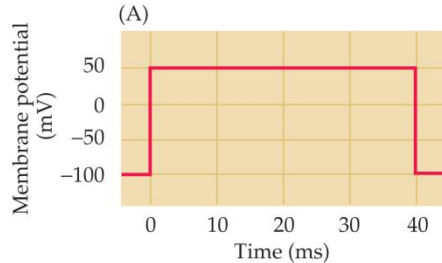
# Na Channel Behavior



Individual channels behave stochastically – only the probability of their opening can be predicted. Because many channels are typically involved the macroscopic current can still be predicted with high accuracy

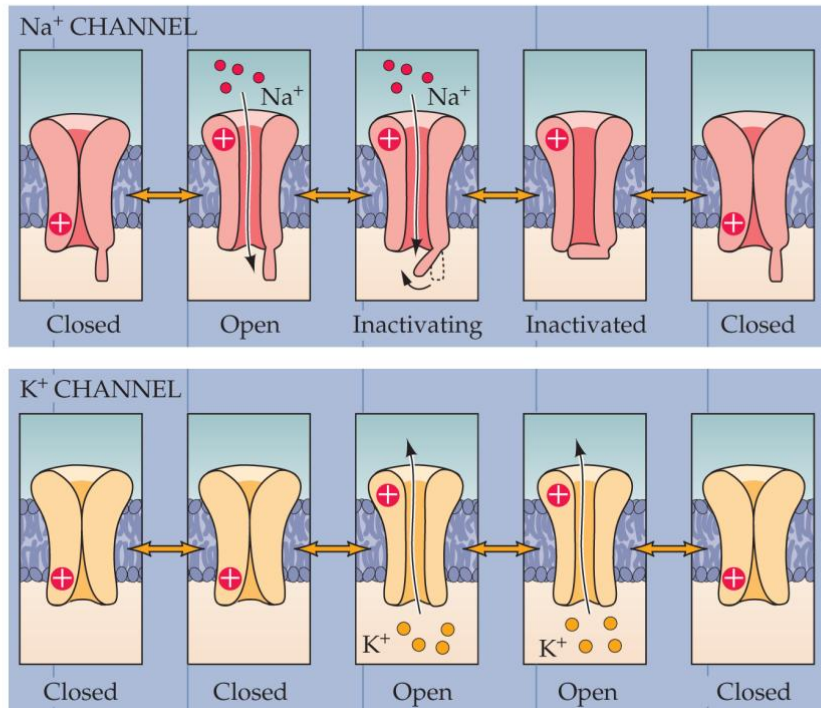
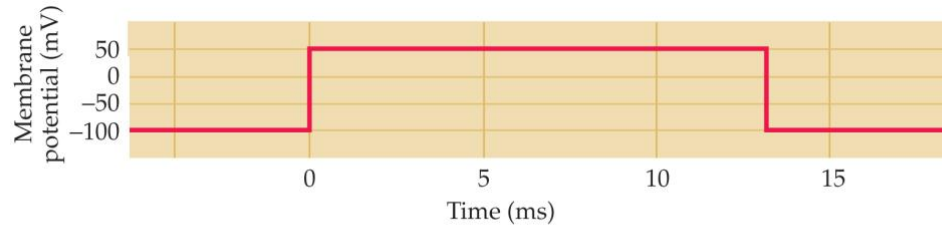


# K Channel Behavior



Individual channels behave stochastically – only the probability of their opening can be predicted. Because many channels are typically involved the macroscopic current can still be predicted with high accuracy

# AP and Channel States



# Activation and Inactivation Curves

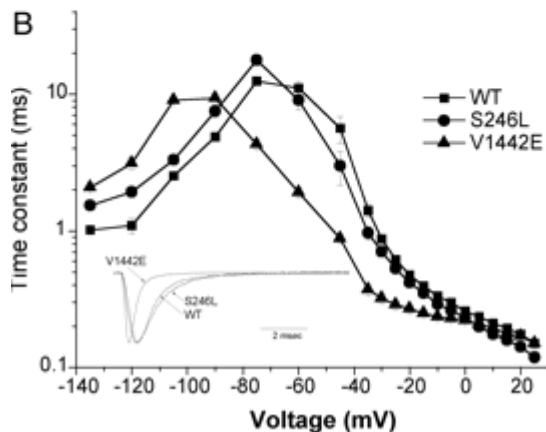
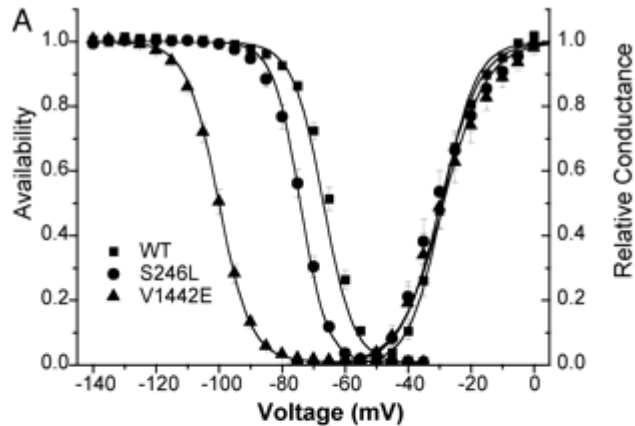


FIGURE 9 Absence of shift in steady-state activation curve with test-pulse height. Steady-state inactivation curves were obtained by giving a 100-ms prepulse to a variable potential, returning to the holding potential for 300  $\mu$ s, and stepping to the test-pulse potential. The peak currents during the test pulses are plotted as a function of the prepulse potential, normalized to the value for a step from  $-84$  mV, the holding potential. The data points for  $-94$  and  $-104$  mV for the  $-34$ -mV test pulse show that there was virtually no inactivation at  $-84$  mV; peak current was about 5% increased by the 100-ms prepulse to  $-104$  mV, and much of this may have been due to removal of slow inactivation. Peak conductance at  $-34$  mV was 0.19 of that at  $+51$  mV.  $\frac{1}{2}$  Na,  $\frac{1}{2}$  K, 1 mM 4-AP, 3.3–3.5°C.

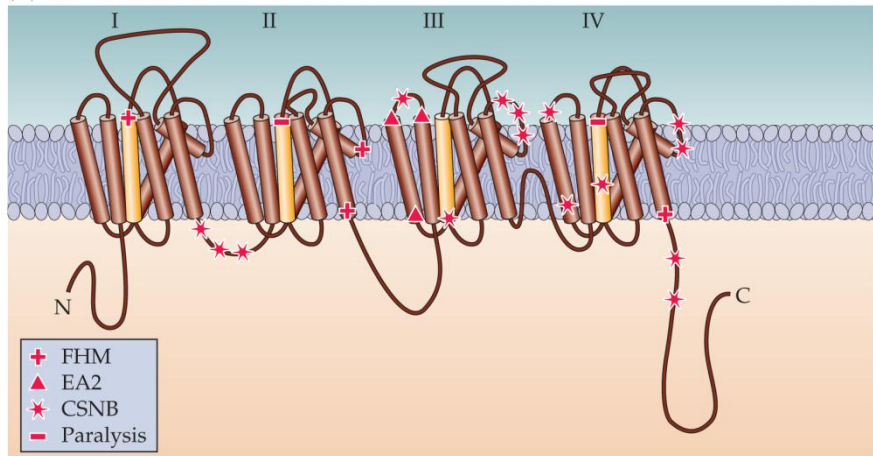
## SODIUM CHANNEL INACTIVATION IN THE CRAYFISH GIANT AXON MUST CHANNELS OPEN BEFORE INACTIVATING?

Myasthenic syndrome caused by  
mutation of the *SCN4A* sodium  
channel

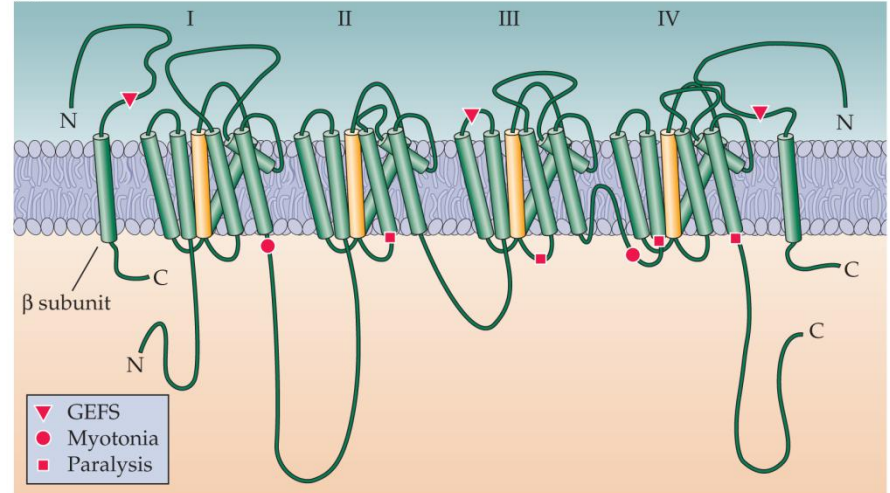
*PNAS June 10, 2003 vol. 100*

# Channelopathy

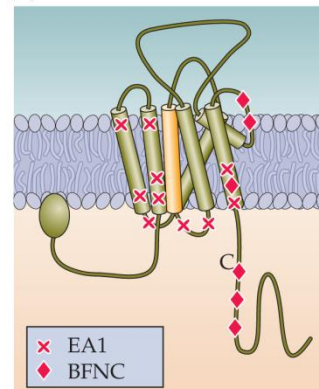
(A)  $\text{Ca}^{2+}$  CHANNEL



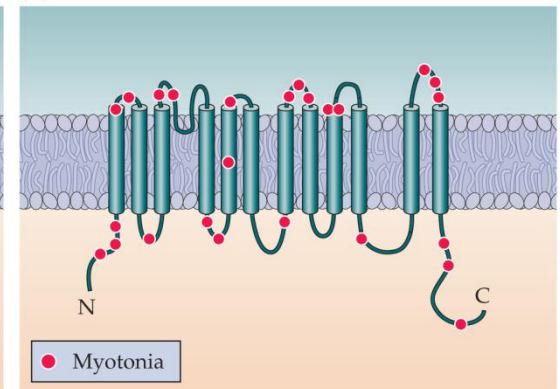
(B)  $\text{Na}^{+}$  CHANNEL



(C)  $\text{K}^{+}$  CHANNEL



(D)  $\text{Cl}^{-}$  CHANNEL



Familial Hemiplegic Migraine  
 Episodic Ataxia Type 2  
 Congenital Stationary Night Blindness

Generalized Epilepsy with Febrile Seizures  
 Benign Familial Neonatal Convulsion

# What Do You Want To Know....

- ... about a channel?
- How is it opened (activation curve)?
- How is it closed (de-activation, inactivation curve)?
- Which ion type(s) go through?
- What are its kinetics?
- What is its single channel conductance?

*These parameters may vary depending on subunit assembly, phosphorylation, ion-, cAMP-, ATP-binding, .....*