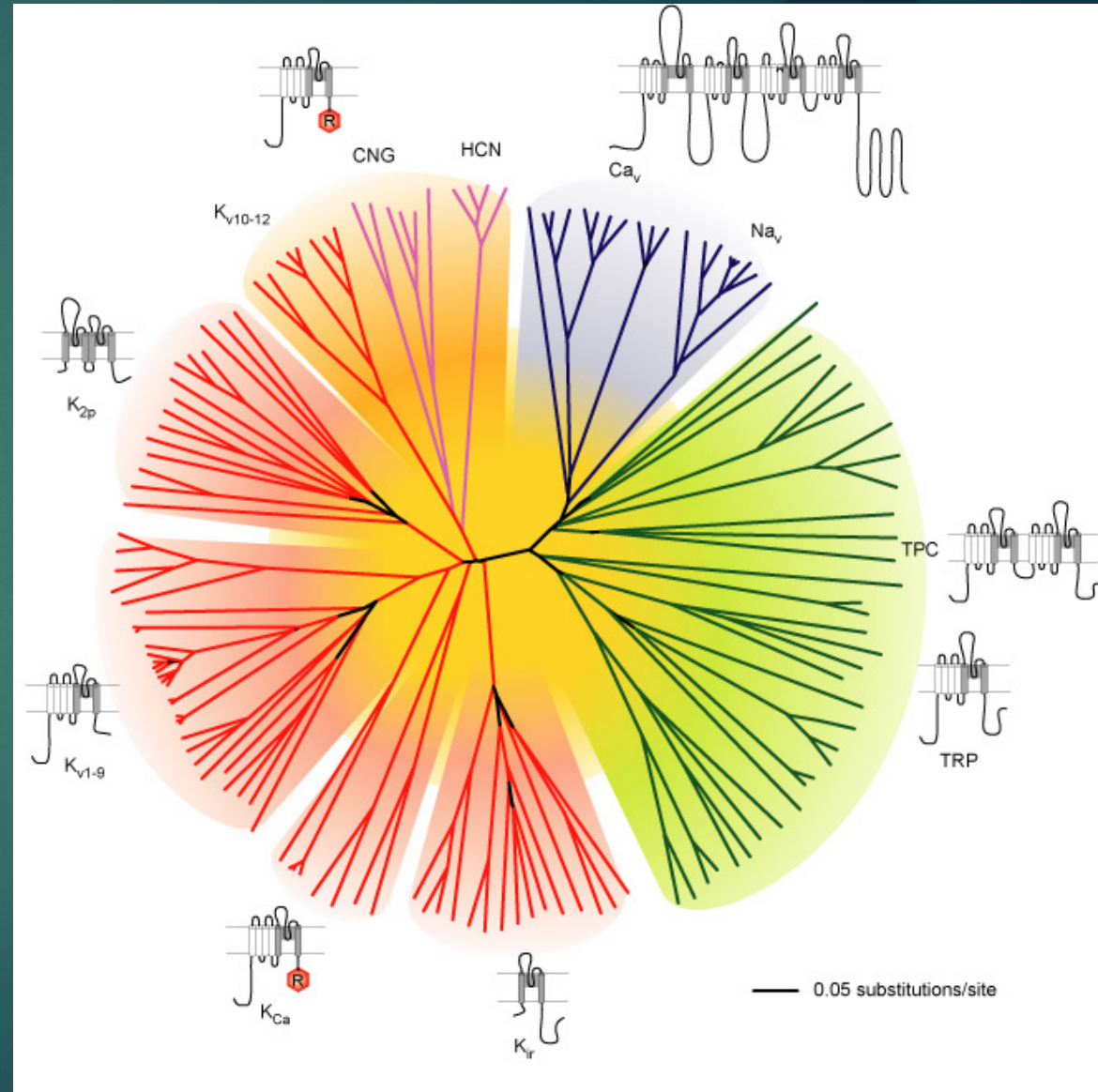


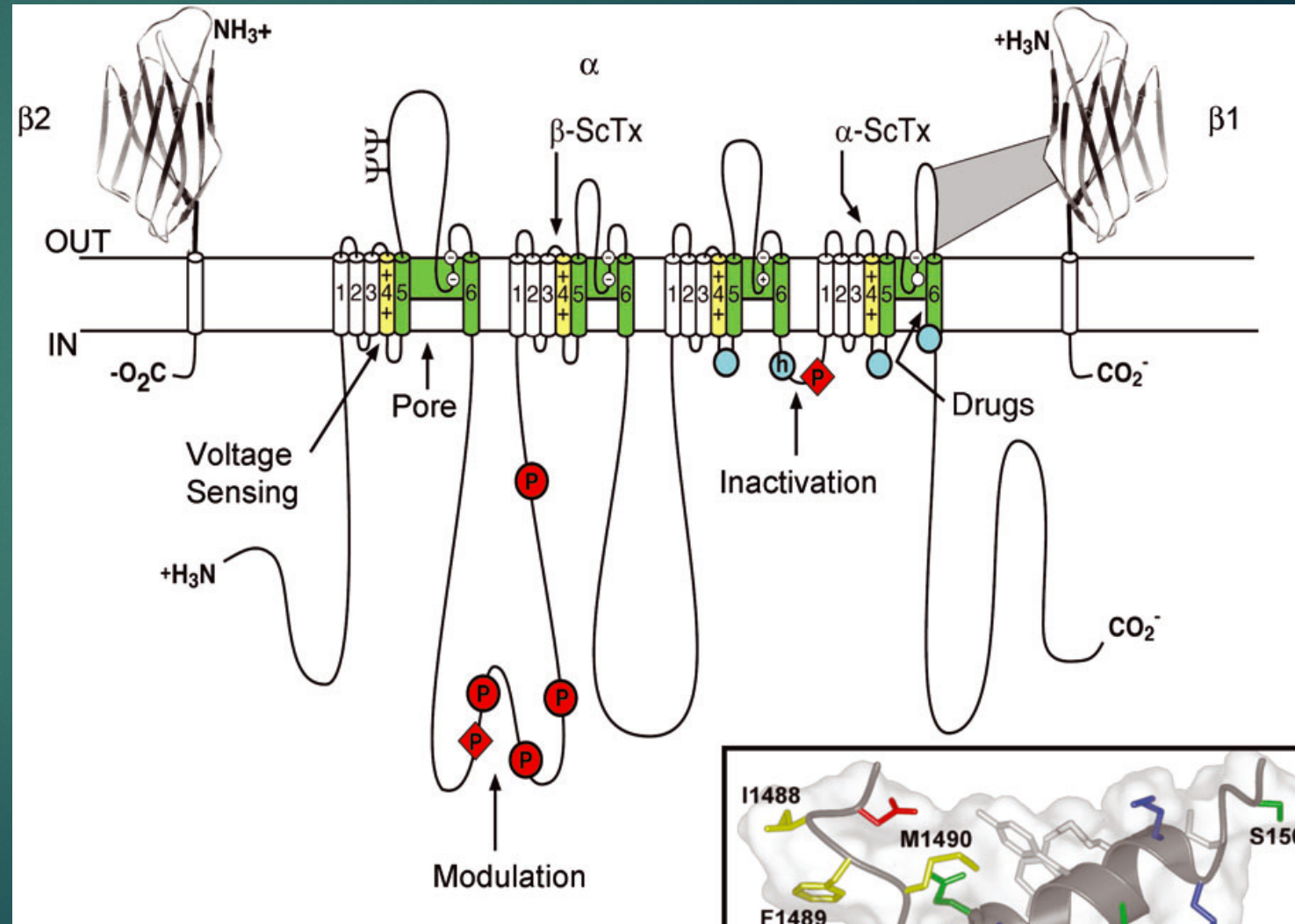
Ion Channels

STRUCTURE AND FUNCTION



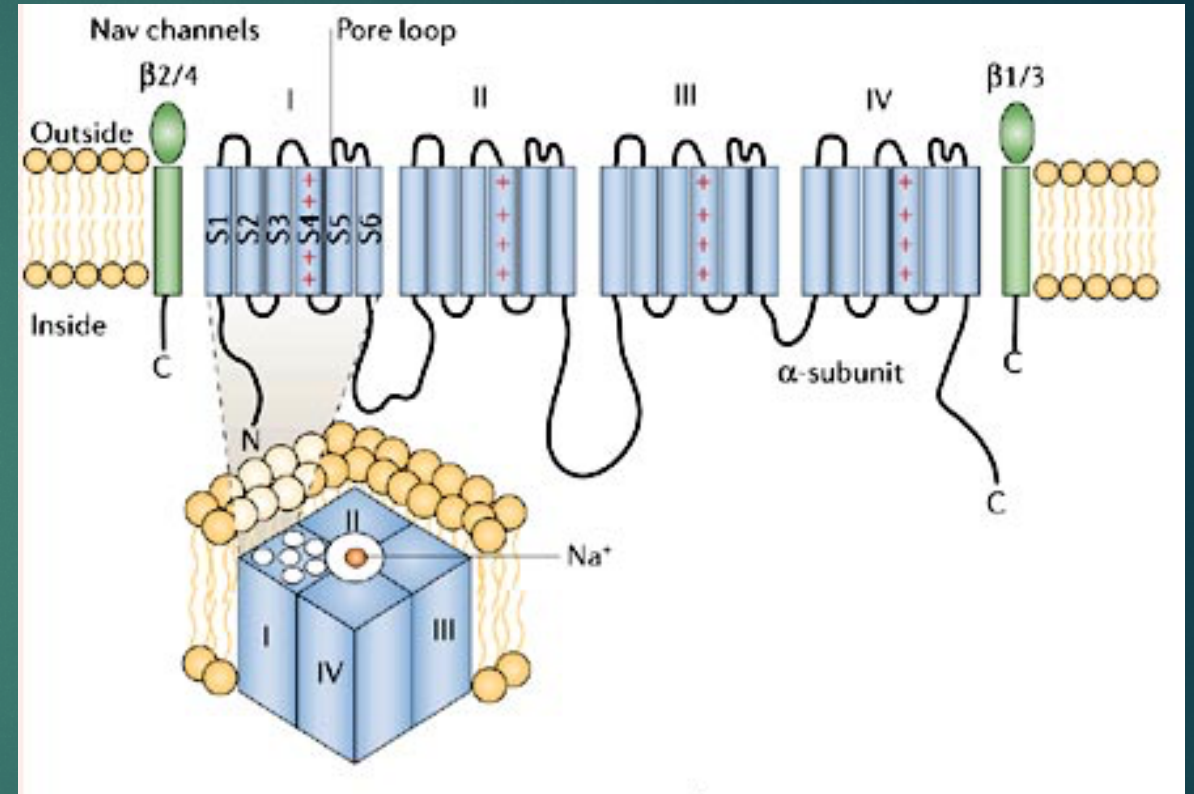
Voltage gated Na⁺ channels

Sodium channel α subunits are composed of approximately 2000 amino acid residues organized in four homologous domains, which each contains six transmembrane segments. Later biochemical analyses and cDNA cloning showed that sodium channel β subunits are composed of an N-terminal extracellular immunoglobulin-like fold, a single transmembrane segment, and a short intracellular segment.



Voltage gated Na⁺ channels

These subunits are thought to form heterodimeric and heterotrimeric complexes composed of a single α subunit and one or two β subunits in excitable cell membranes, and co-expression of β subunits modulates the kinetics and voltage dependence of sodium channel activation and inactivation.



Voltage gated Na⁺ channels

Sodium channel **α subunits are encoded by 10 genes**, which are expressed in different excitable tissues.

- NaV1.1, 1.2, 1.3 and 1.6 are the primary sodium channels in the central nervous system.

- NaV1.7, 1.8 and 1.9 are the primary sodium channels in the peripheral nervous system.

- NaV1.4 is the primary sodium channel in skeletal muscle, whereas

- NaV1.5 is primary in heart.

Most of these sodium channels also have significant levels of expression outside of their primary tissues.

- The 10th sodium channel protein is not voltage-gated and is involved in salt sensing.

Table 1. Mammalian sodium channel α subunits

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

Voltage gated Na⁺ channels

β subunits have been identified by genomic analyses and cDNA cloning to give a small family of four Navβ subunits in total.

β1 and β3 are associated non-covalently with α subunits and resemble each other most closely in amino acid sequence, whereas β2 and β4 form disulfide bonds with α subunits and also resemble each other closely.

The structure of Navβ subunits resembles the family of cell adhesion molecules, and increasing evidence supports their role in localization and immobilization of sodium channels in specific locations in excitable cells.

Voltage gated Ca²⁺ channels

Ca²⁺ entering the cell through voltage-gated Ca²⁺ channels serves as the second messenger of electrical signaling, initiating many different cellular events:

- In cardiac and smooth muscle cells, activation of Ca²⁺ channels initiates contraction directly by increasing cytosolic Ca²⁺ concentration and indirectly by activating calcium-dependent calcium release by ryanodine-sensitive Ca²⁺ release channels in the sarcoplasmic reticulum.
- In skeletal muscle cells, voltage-gated Ca²⁺ channels in the transverse tubule membranes interact directly with ryanodine-sensitive Ca²⁺ release channels in the sarcoplasmic reticulum and activate them to initiate rapid contraction.

Voltage gated Ca²⁺ channels

- In endocrine cells, voltage-gated Ca²⁺ channels mediate Ca²⁺ entry that initiates secretion of hormones.
- In neurons, voltage-gated Ca²⁺ channels initiate synaptic transmission.
- In many different cell types, Ca²⁺ entering the cytosol via voltage-gated Ca²⁺ channels regulates enzyme activity, gene expression, and other biochemical processes

Voltage gated Ca²⁺ channels

Table 1. Subunit composition and function of Ca²⁺ channel types

Ca ²⁺ current type	α1 Subunits	Specific blocker	Principal physiological functions	Inherited diseases
L	Ca _v 1.1	DHPs	Excitation-contraction coupling in skeletal muscle, regulation of transcription	Hypokalemic periodic paralysis
	Ca _v 1.2	DHPs	Excitation-contraction coupling in cardiac and smooth muscle, endocrine secretion, neuronal Ca ²⁺ transients in cell bodies and dendrites, regulation of enzyme activity, regulation of transcription	Timothy syndrome: cardiac arrhythmia with developmental abnormalities and autism spectrum disorders
	Ca _v 1.3	DHPs	Endocrine secretion, cardiac pacemaking, neuronal Ca ²⁺ transients in cell bodies and dendrites, auditory transduction	
	Ca _v 1.4	DHPs	Visual transduction	Stationary night blindness
N	Ca _v 2.1	ω-CTx-GVIA	Neurotransmitter release, Dendritic Ca ²⁺ transients	
P/Q	Ca _v 2.2	ω-Agatoxin	Neurotransmitter release, Dendritic Ca ²⁺ transients	Familial hemiplegic migraine, cerebellar ataxia
R	Ca _v 2.3	SNX-482	Neurotransmitter release, Dendritic Ca ²⁺ transients	
T	Ca _v 3.1	None	Pacemaking and repetitive firing	Absence seizures
	Ca _v 3.2		Pacemaking and repetitive firing	
	Ca _v 3.3			

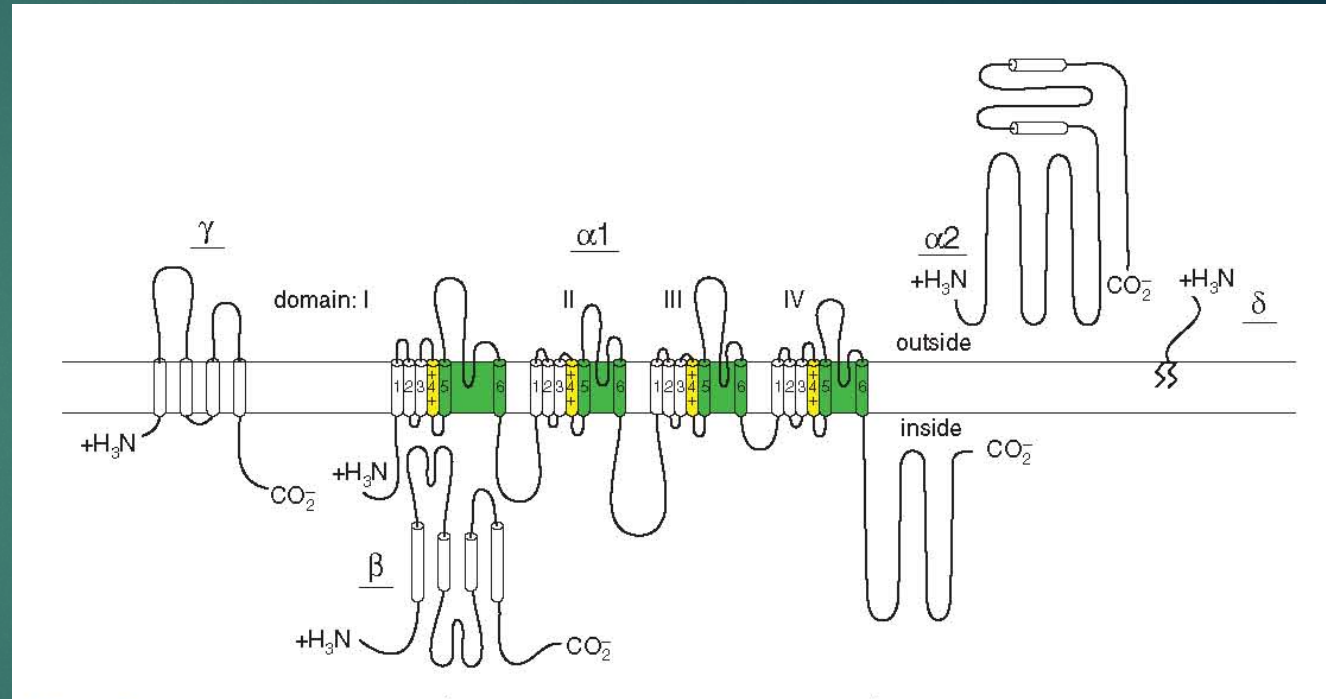
Abbreviations: DHP, dihydropyridine; ω-CTx-GVIA, ω-conotoxin GVIA from the cone snail *Conus geographus*; SNX-482, a synthetic version of a peptide toxin from the tarantula *Hysterocrates gigas*.

Voltage gated Ca²⁺ channels

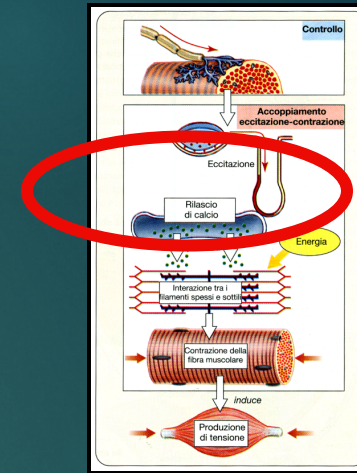
Ca²⁺ channels purified from skeletal muscle transverse tubules are complexes of $\alpha 1$, $\alpha 2$, β , γ , and δ subunits.

The principal transmembrane $\alpha 1$ subunit of 190 kDa in association with a disulfide-linked $\alpha 2\delta$ dimer of 170 kDa, an intracellular phosphorylated β subunit of 55 kDa, and a transmembrane γ subunit of 33 kDa.

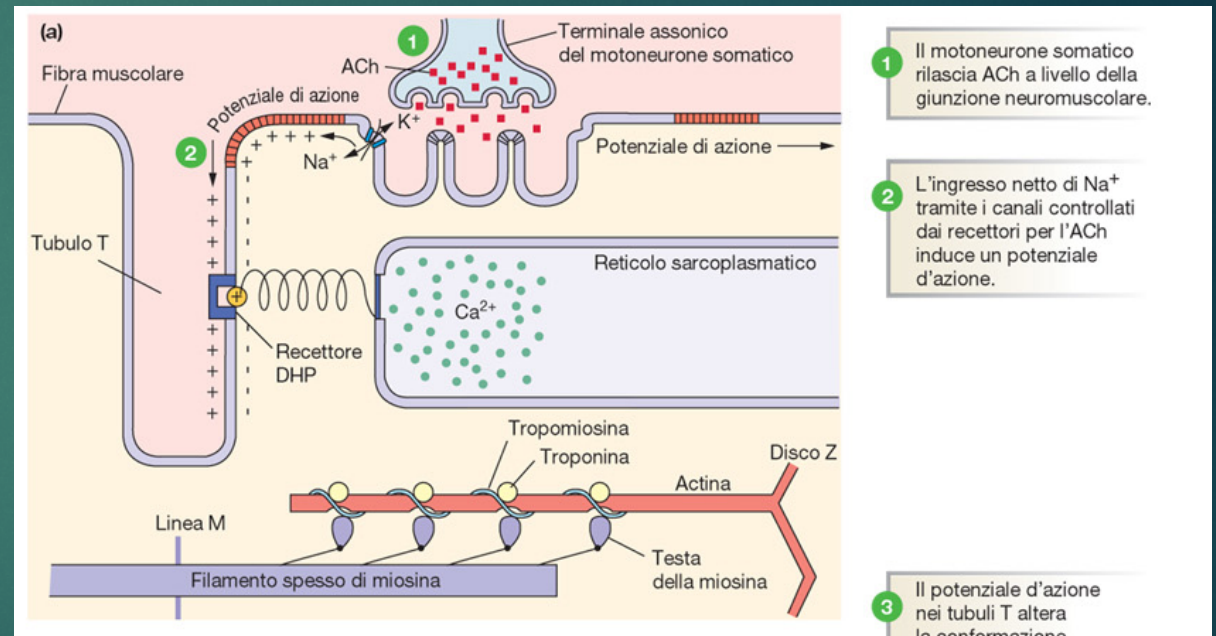
The $\alpha 1$ subunit is a protein of about 2000 amino acid residues in length with an amino acid sequence and predicted transmembrane structure like the previously characterized, pore-forming α subunit of voltage-gated sodium channels



Excitation-contraction coupling in skeletal muscle

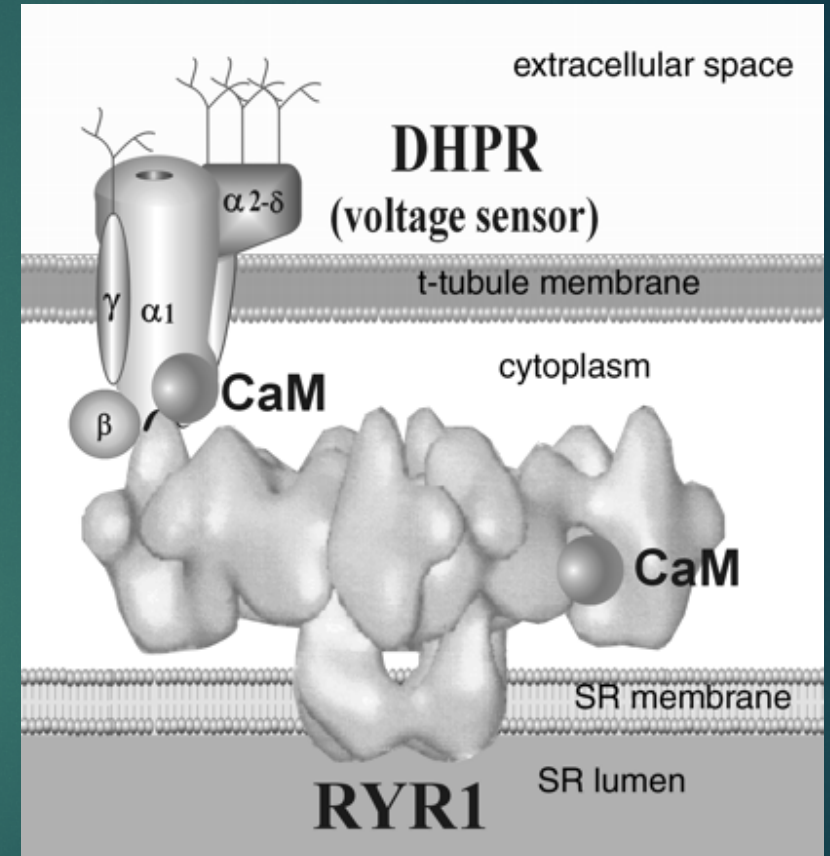


CaV1.1 channels in the transverse tubules are thought to interact directly with the ryanodine-sensitive Ca²⁺ release channels (RyR1) of the sarcoplasmic reticulum, as observed in high-resolution electron microscopy, and the voltage-driven conformational changes in their voltage-sensing domains are thought to directly induce activation of RyR1.



In skeletal muscle, entry of external Ca²⁺ is not required for initiation of contraction

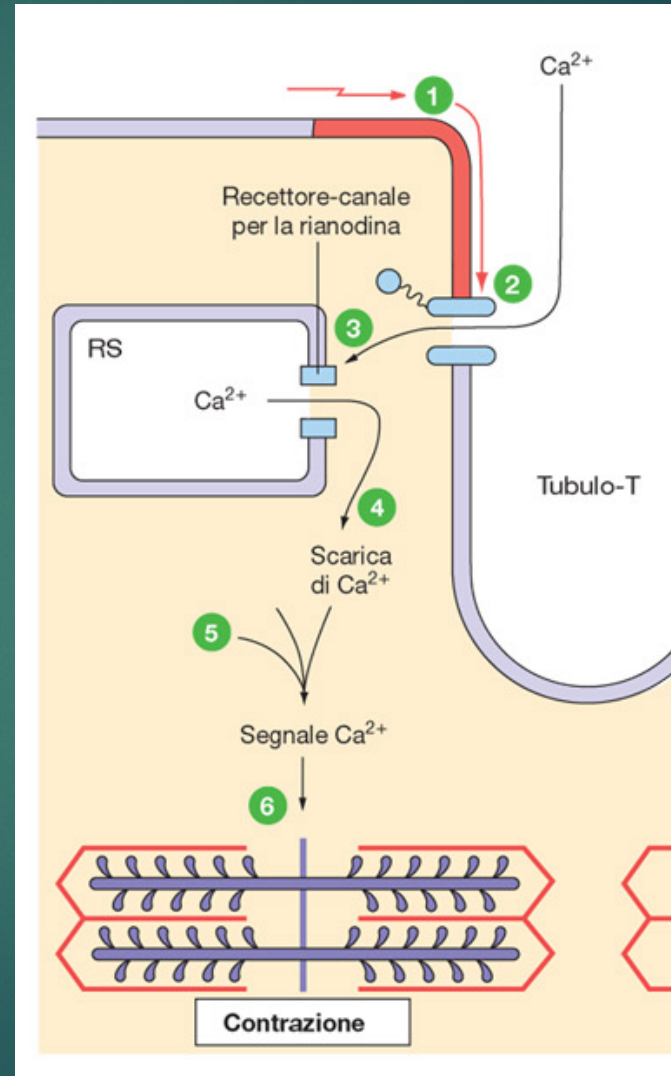
This is because the cytoplasmic domain of these channels is physically coupled to ryanodine receptor (RyR1) Ca^{2+} release channels on internal membranes. Even though Cav1.1 proteins can act as *bona fide* Ca^{2+} channels, they also function as voltage sensors that directly produce conformational changes in the ryanodine receptor/Cav1.1 complex, resulting in the release of Ca^{2+} from internal stores.



Excitation-contraction coupling in cardiac muscle

In contrast to skeletal muscle, entry of Ca^{2+} is required for excitation-contraction coupling in cardiac myocytes, and Ca^{2+} entry via CaV1.2 channels triggers activation of the RyR2 and initiates Ca^{2+} -induced Ca^{2+} -release, activation of actomyosin, and contraction

Release of Ca^{2+} from the sarcoplasmic reticulum via RyR2 greatly amplifies the cellular Ca^{2+} transient and is required for effective initiation of contraction. All three steps in the cascade of Ca^{2+} transport processes— Ca^{2+} entry via CaV1.2 channels, Ca^{2+} release via RyR , and Ca^{2+} uptake into the sarcoplasmic reticulum by SERCA Ca^{2+} pumps—are tightly regulated by second messenger signaling networks



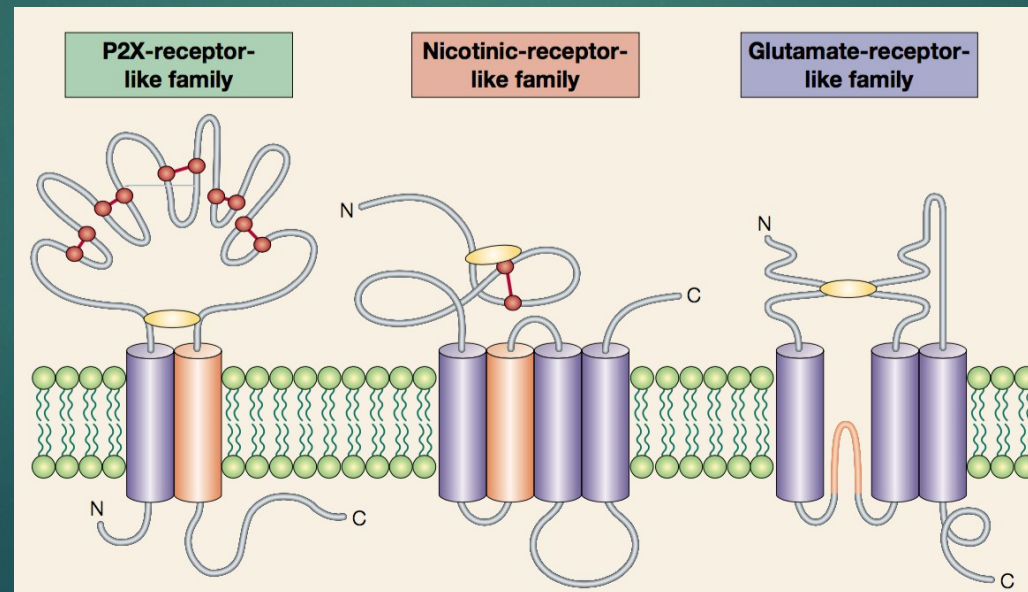
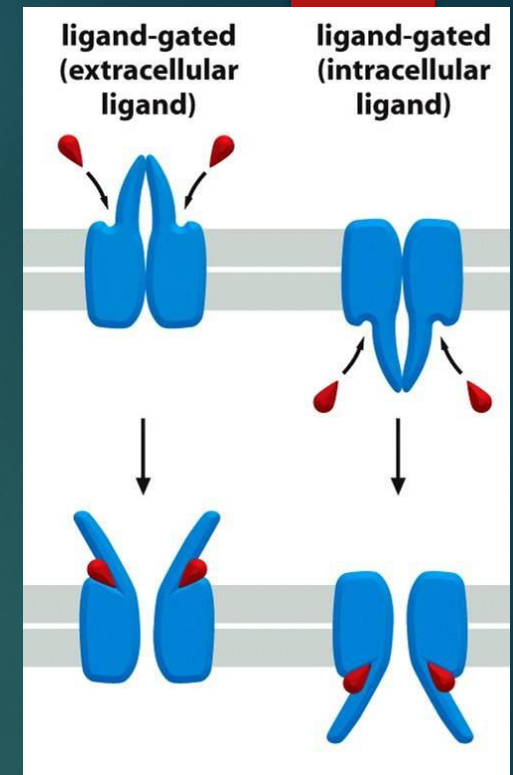
- 1 Un potenziale d'azione invade la membrana cellulare provenendo da una cellula adiacente.
- 2 I canali voltaggio-dipendenti per il Ca^{2+} si aprono. Il Ca^{2+} entra nella cellula.
- 3 L'ingresso di Ca^{2+} innesca il rilascio di altro Ca^{2+} dal reticolo sarcoplasmatico attraverso i recettori-canali della rianodina (RyR).
- 4 Il rilascio localizzato di calcio provoca la «scarica» di Ca^{2+} .
- 5 Le scariche di Ca^{2+} si sommano per produrre un segnale di Ca^{2+} .
- 6 Gli ioni calcio si legano alla troponina e inizia la contrazione.

Ligand gated ion channels

This is a highly heterogenous family of channels that includes several families

- the **extracellular ligand-activated channels** which includes channels such as **GABA and glycine receptor channels**, most of which are regulated by ligands that are "neurotransmitters". These channels are often named according to the ligand they bind to. Other examples are: **nicotinic receptors ; P2X receptors**

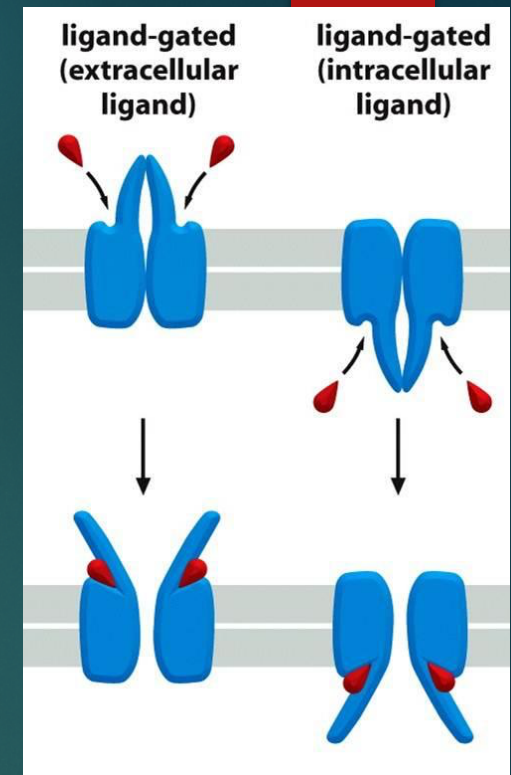
We will discuss different examples in the course



Ligand gated ion channels

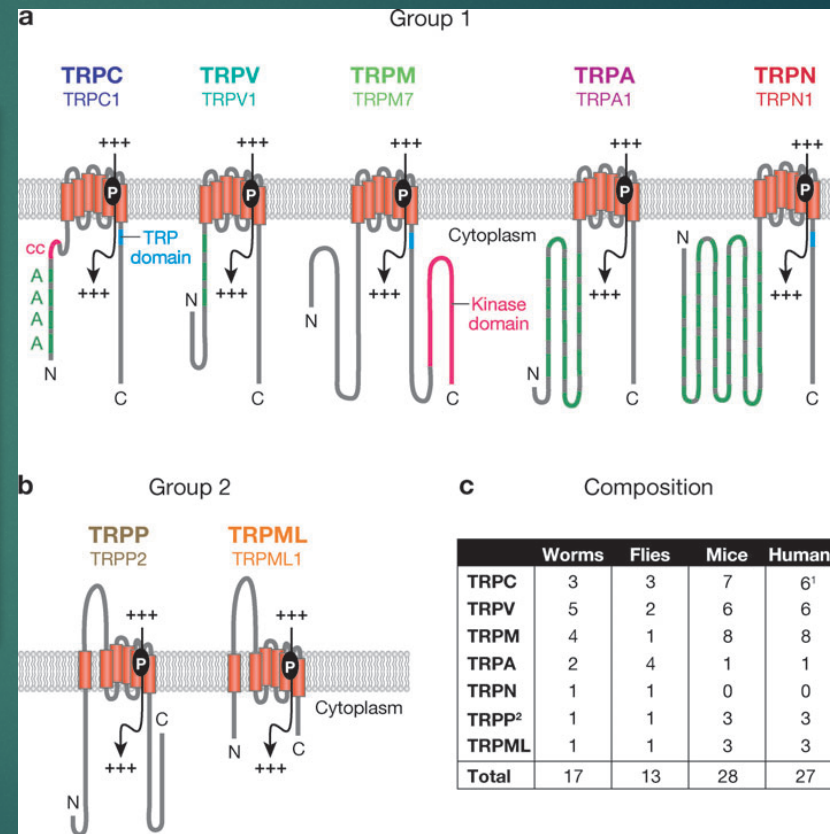
This is a highly heterogeneous family of channels that includes several families

- **Intracellular ligand-gated ion channels.** These include **CFTR** and some other ABC family members as well as ion channels involved in sense perception; **TRP channels; CNGC**; **These are often activated indirectly by GCPRs.** Other common intracellular ligands which activate these kinds of channels include calcium ions, **ATP, cyclic AMP and GMP as well as phosphatidylinositol (PI).** There are additional systems of nomenclature which have joined the second and third groups into the "chemically activated" or just simply "ligand gated" ion channels.



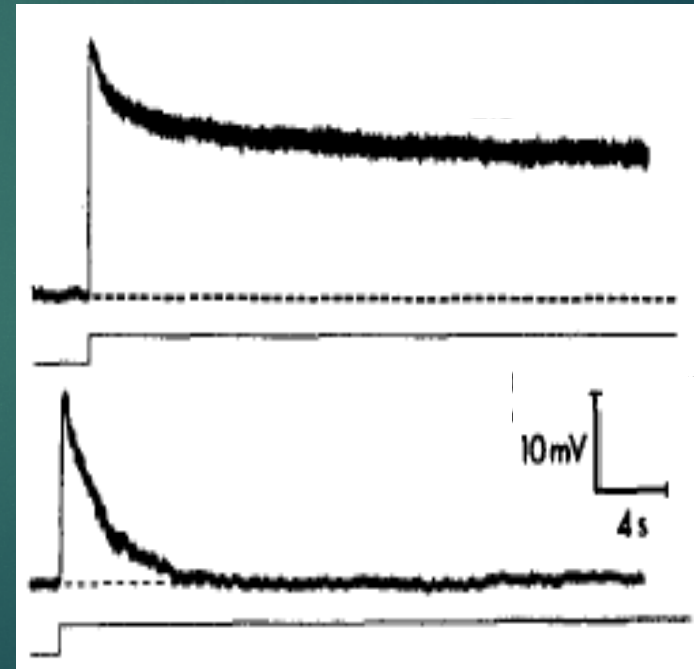
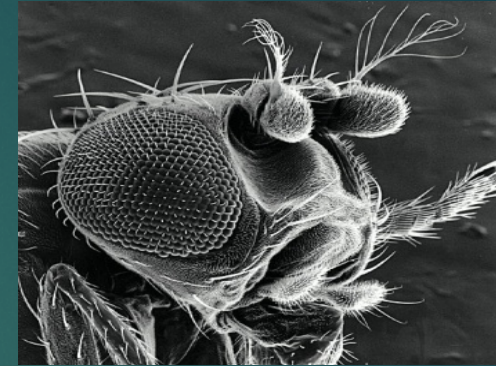
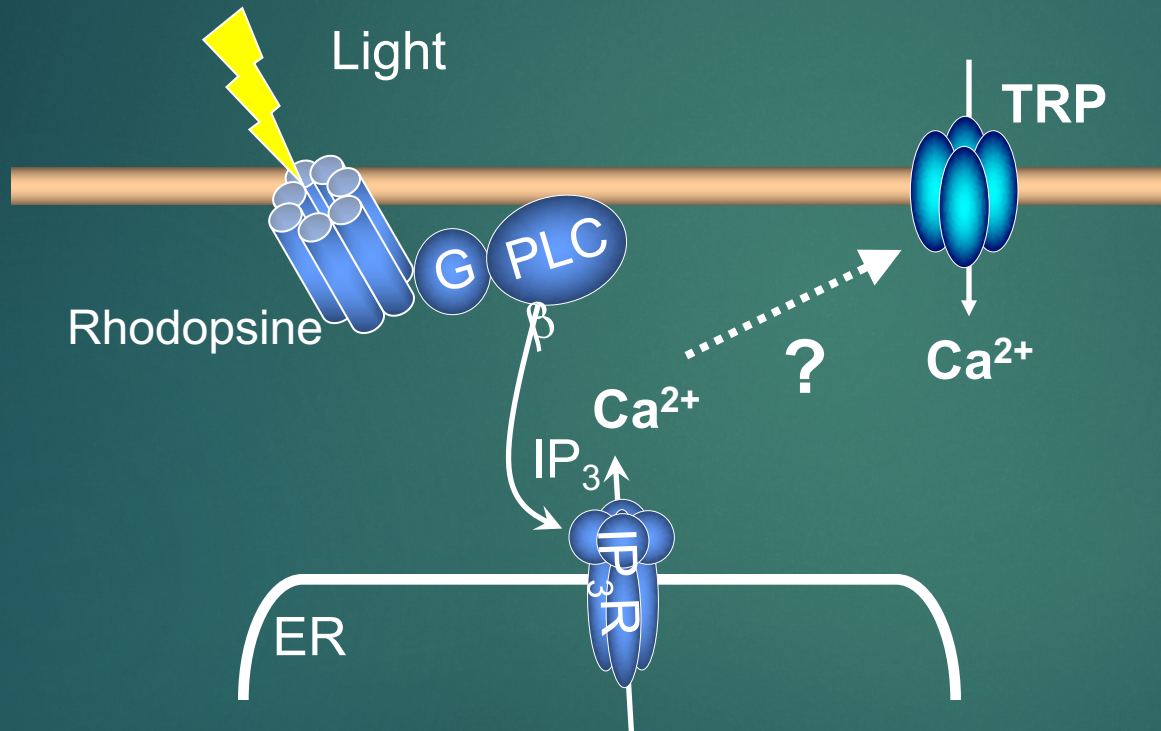
TRP family of channels

- Cation channels
- Non voltage-dependent
- Diversity in activation mechanisms
- Implication in diverse physiological functions



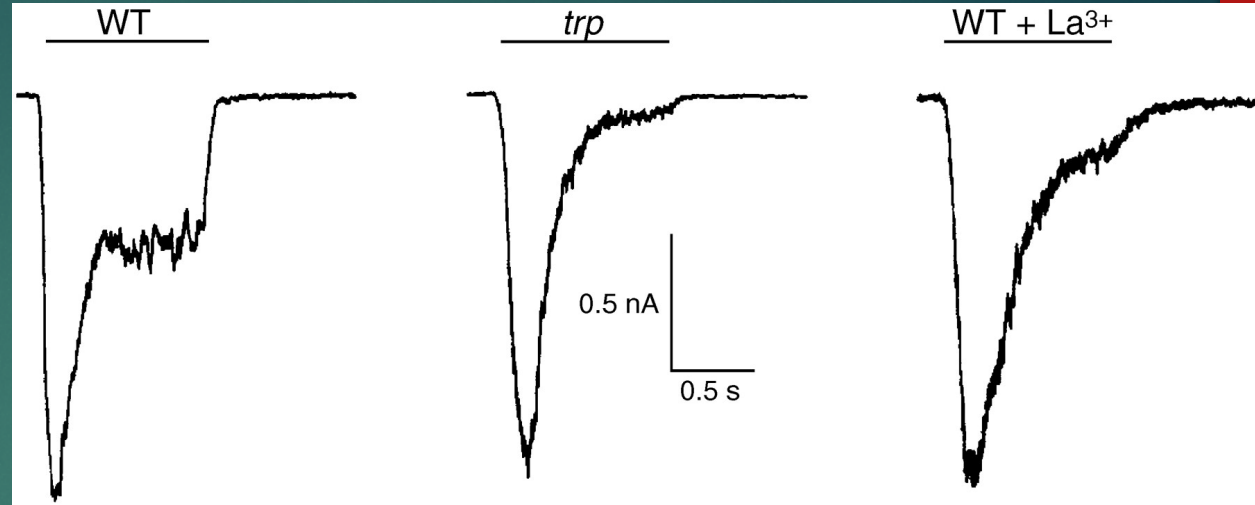
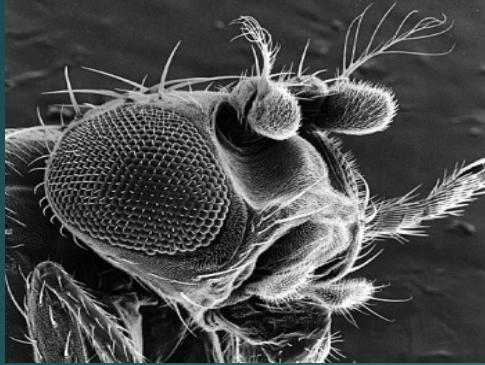
First TRP channel identification

Drosophila melanogaster :



Minke B, Biophys Struct Mech., 1977

Identification du 1er Canal TRP

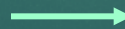


Light

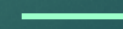
Light

Light

trp
Transient Receptor Potential



Cationic channel

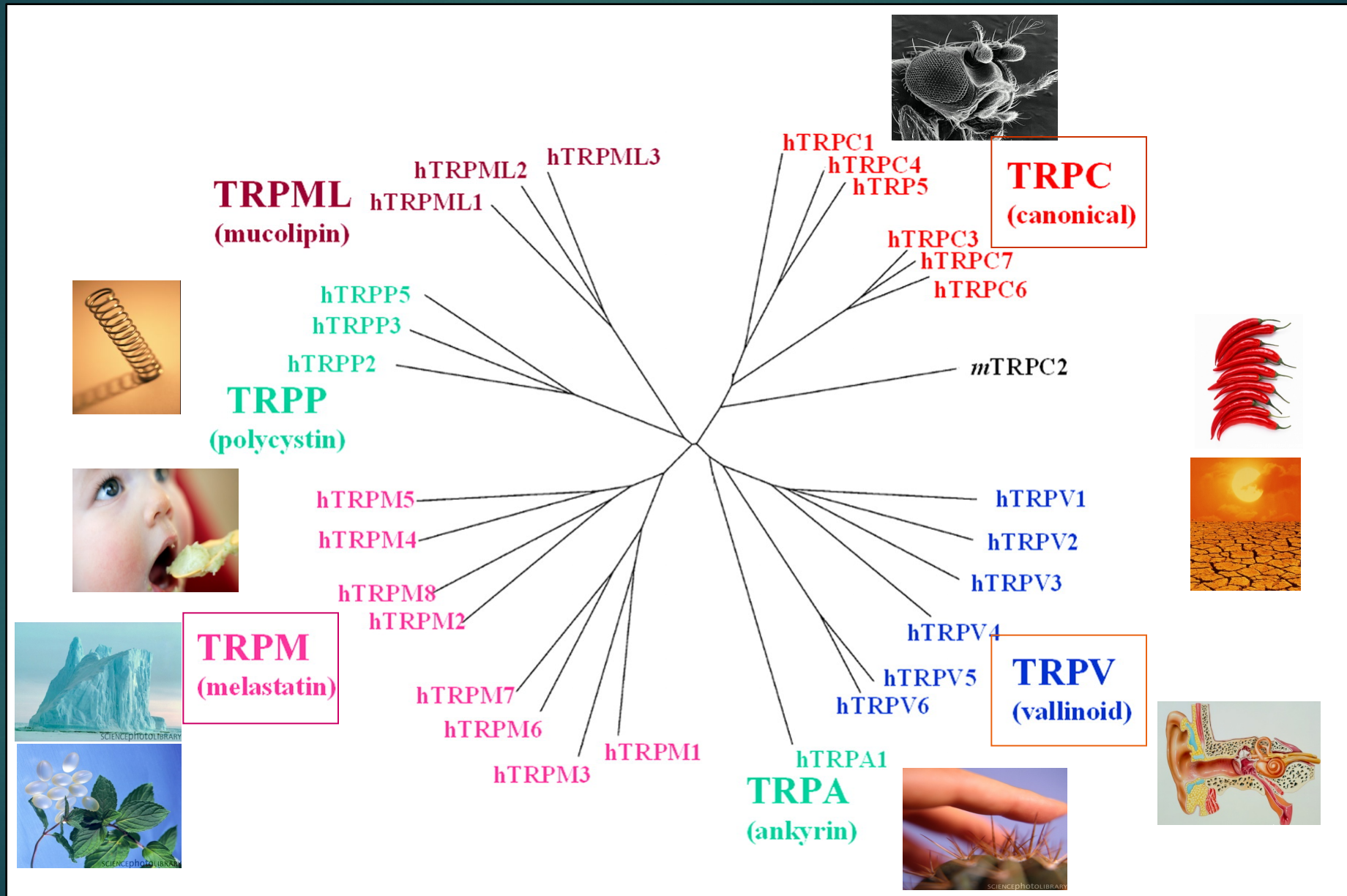


Conserved from
c.Elegans to human

TRP family composition in worms, flies, mice and humans

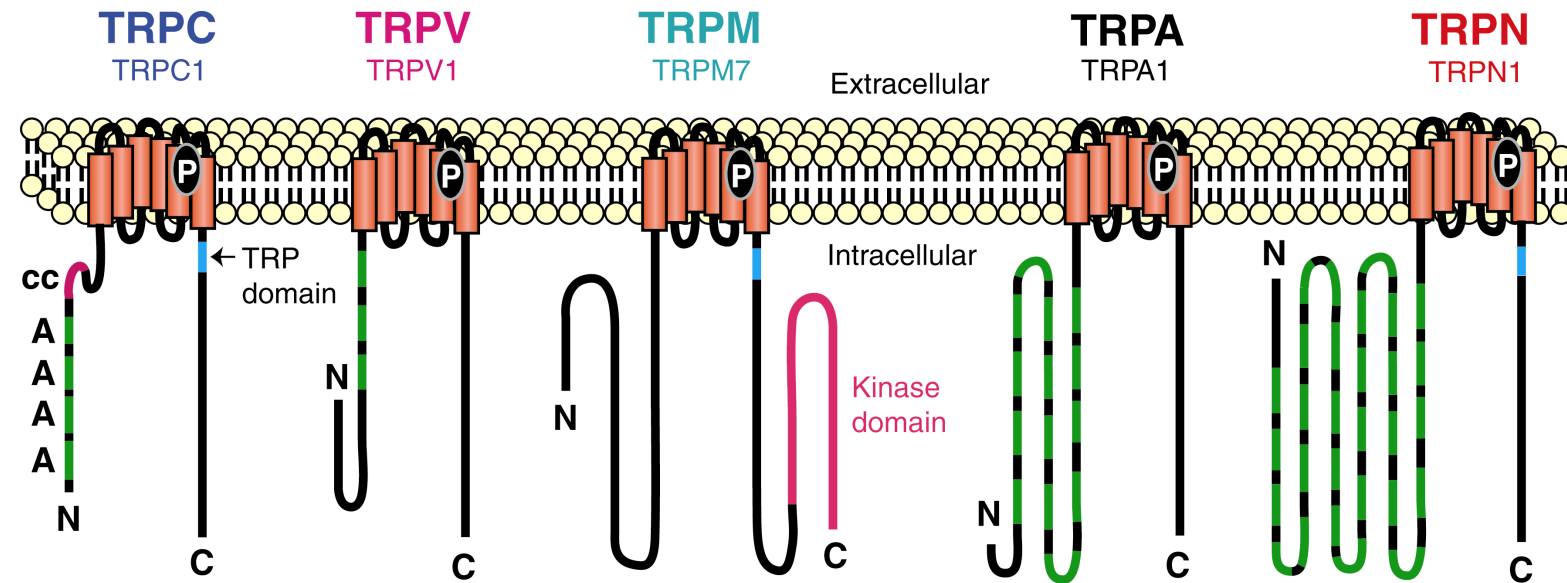
Subfamily	Worms	Flies	Mice	Humans
TRPC	3	3	7	6 ¹
TRPV	5	2	6	6
TRPM	4	1	8	8
TRPA	2	4	1	1
TRPN	1	1	0	0
TRPP ²	1	1 ³	3	3
TRPML	1	1	3	3
Total	17	13	28	27

The TRP channels and 5 senses

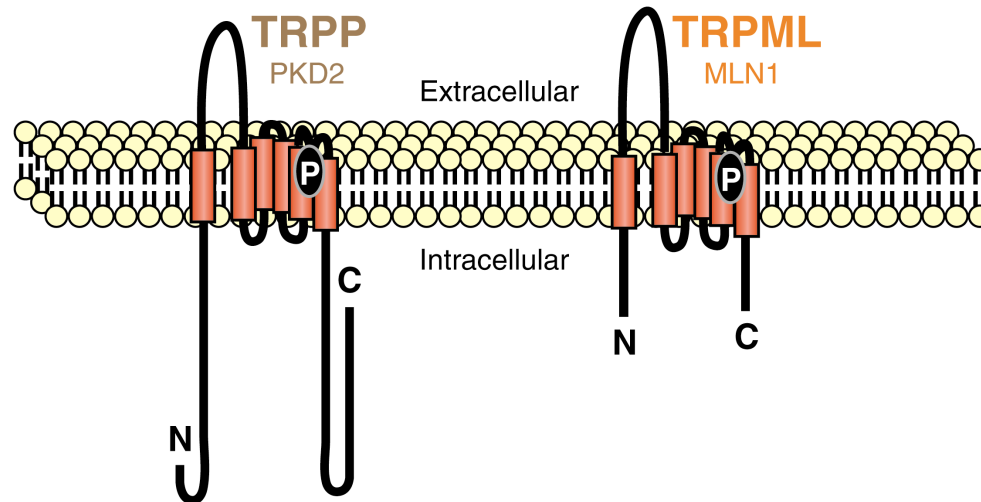


Different topologies of TRP channels

Group 1 TRPs



Group 2 TRPs



The quaternary structure of TRP channels allows homo- or heteromeric configurations

