



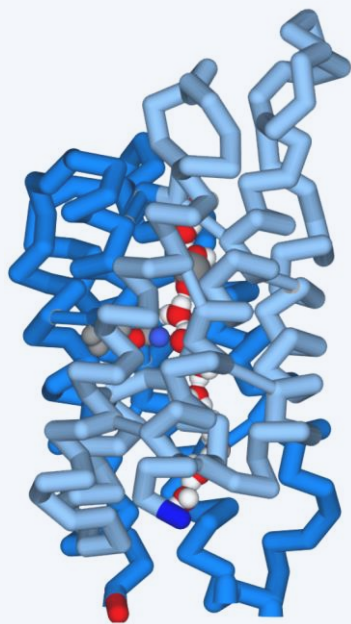
Introducing Mighty Models...

*...from the water channel
to action potentials*

Tim Herman
3D Molecular Designs

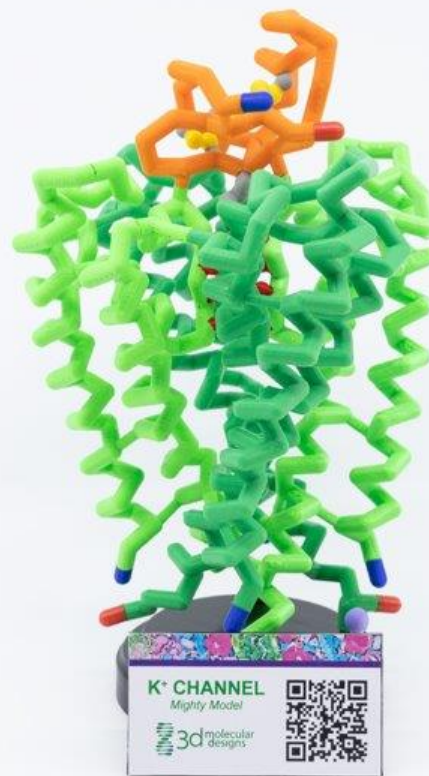


Aquaporin Mighty Model



Hg atom

Potassium Channel Mighty Model



Scorpion toxin

Sodium Channel Mighty Model



Small molecule inhibitor

The Discovery of AQP1

(...in the early 1990's...)

“The field was essentially stuck, but following the well-known scientific approach known as *sheer blind luck*, we stumbled upon the protein that is the answer to the question ...

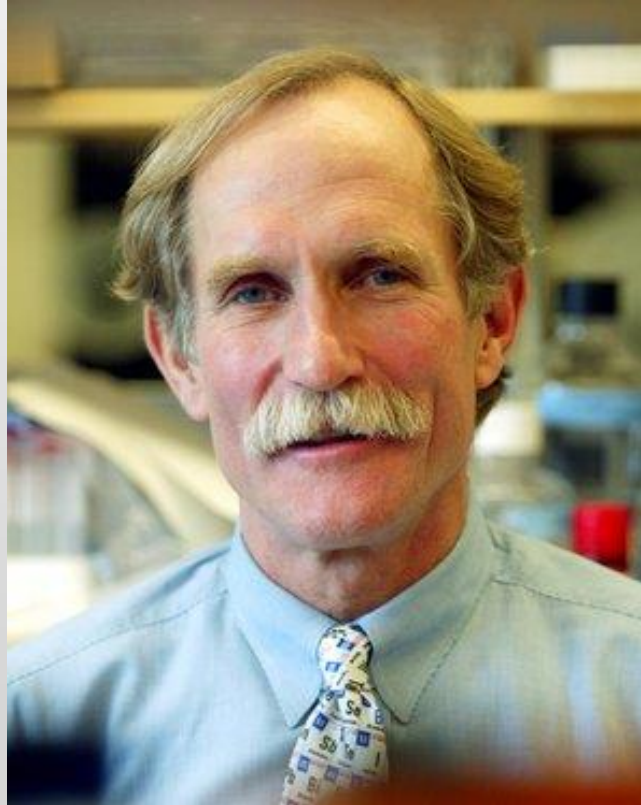
Do water channels exist?”

From Peter Agre's Nobel Lecture



2003 Nobel Prize in Chemistry

The Nobel Prize in Chemistry 2003 was awarded "for discoveries concerning channels in cell membranes" jointly with one half to Peter Agre "for the discovery of water channels" and with one half to Roderick MacKinnon "for structural and mechanistic studies of ion channels."



Peter Agre



Roderick MacKinnon

Appearance of Water Channels in *Xenopus* Oocytes Expressing Red Cell CHIP28 Protein

**Gregory M. Preston, Tiziana Piazza Carroll,
William B. Guggino, Peter Agre***

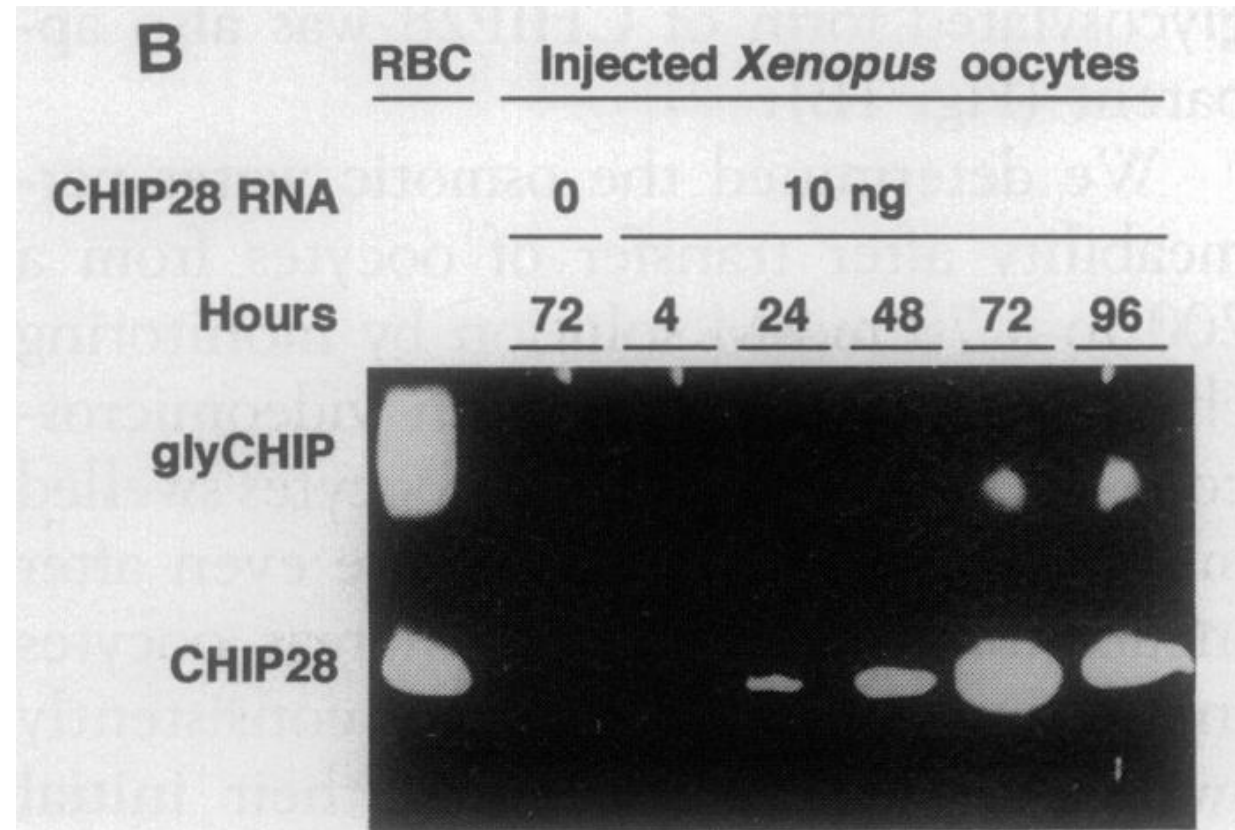
Water rapidly crosses the plasma membrane of red blood cells (RBCs) and renal tubules through specialized channels. Although selective for water, the molecular structure of these channels is unknown. The CHIP28 protein is an abundant integral membrane protein in mammalian RBCs and renal proximal tubules and belongs to a family of membrane proteins with unknown functions. Oocytes from *Xenopus laevis* microinjected with in vitro-transcribed CHIP28 RNA exhibited increased osmotic water permeability; this was reversibly inhibited by mercuric chloride, a known inhibitor of water channels. Therefore it is likely that CHIP28 is a functional unit of membrane water channels.

Science 256, 365-387 (1992)

Appearance of Water Channels in *Xenopus* Oocytes Expressing Red Cell CHIP28 Protein

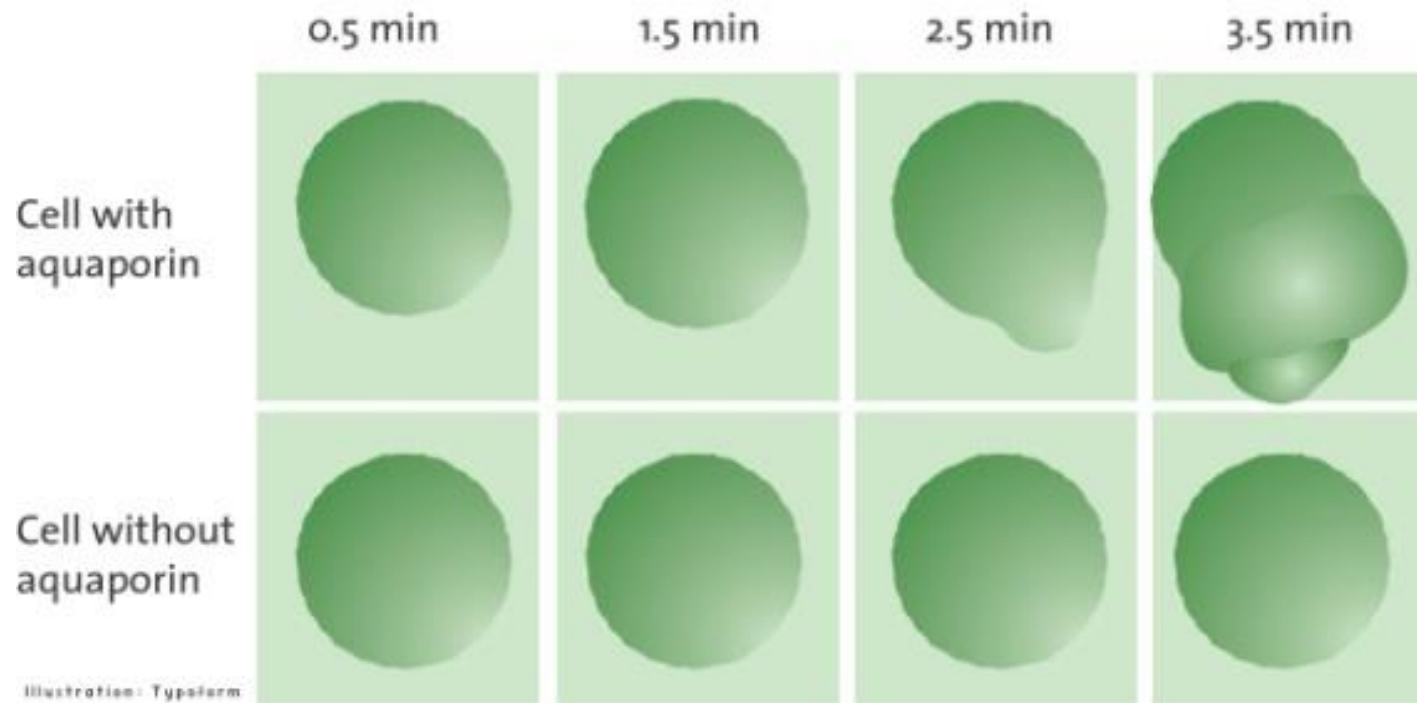
Gregory M. Preston, Tiziana Piazza Carroll,
William B. Guggino, **Peter Agre***

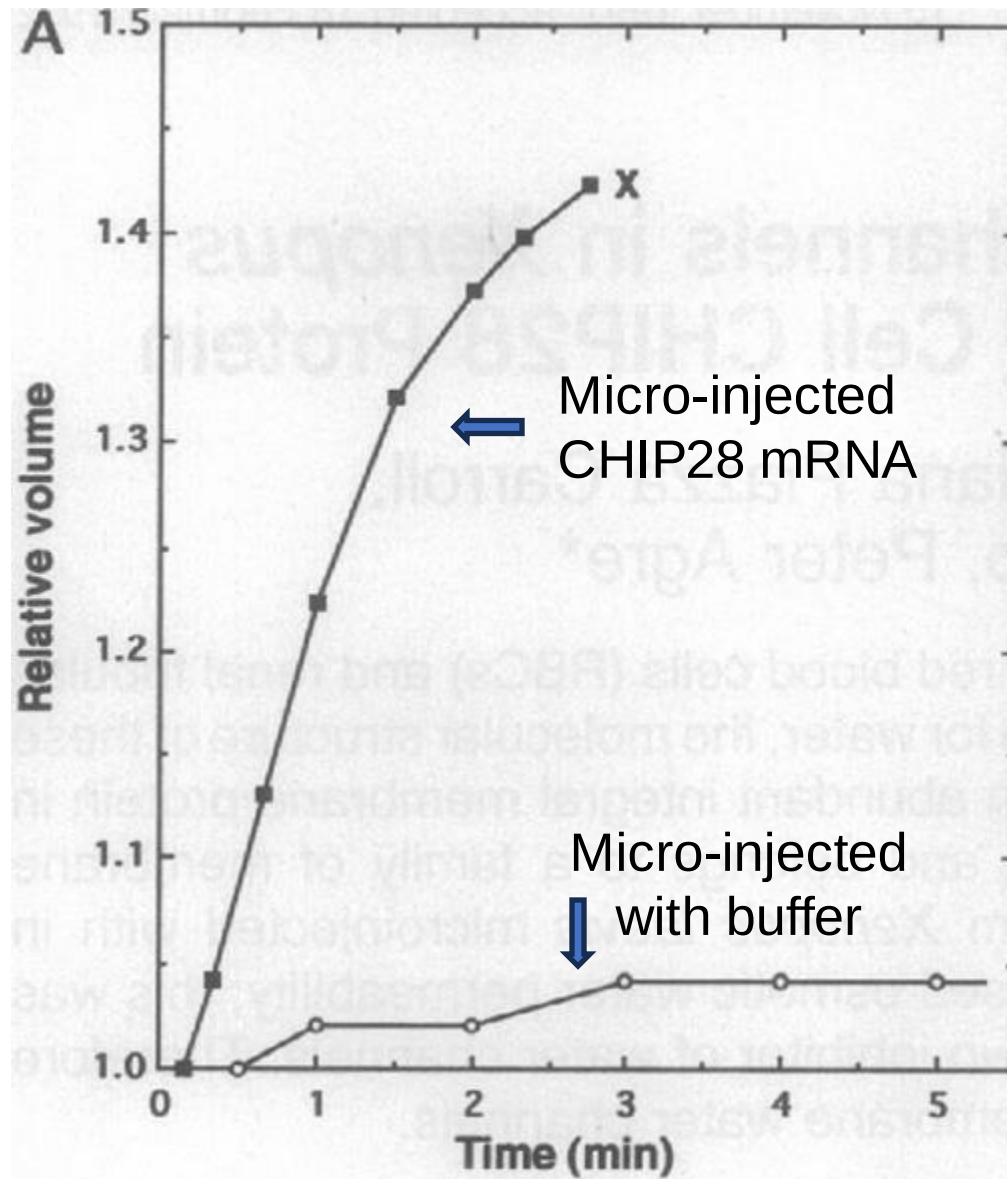
SCIENCE * VOL. 256 * 17 APRIL 1992



The Famous *Exploding Frog Egg Experiment*

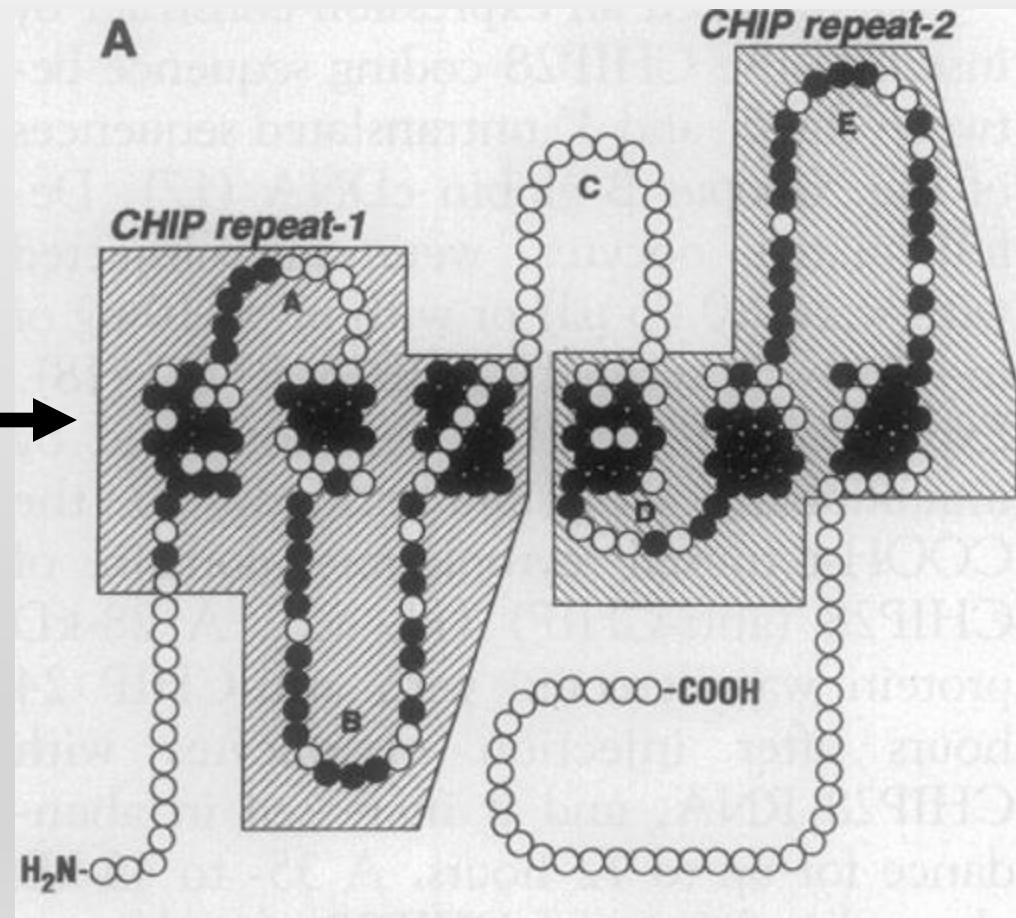
Aquaporin mRNA micro-injected into *Xenopus* eggs





1992 “Model” of Aquaporin

- Based on its Amino Acid Sequence
- 6 Transmembrane Helices
- 2-Fold Symmetry



Structural determinants of water permeation through aquaporin-1

Kazuyoshi Murata^{*†}, Kaoru Mitsuoka^{†‡}, Teruhisa Hirai^{†‡}, Thomas Walz[§], Peter Agre^{||}, J. Bernard Heymann[¶], Andreas Engel[¶] & Yoshinori Fujiyoshi[‡]

^{*} National Institute for Physiological Sciences, Okazaki 444-8585, Japan

[‡] Department of Biophysics, Faculty of Science, Kyoto University, Kitashirakawa, Sakyo-ku, Kyoto, 606-8502, Japan

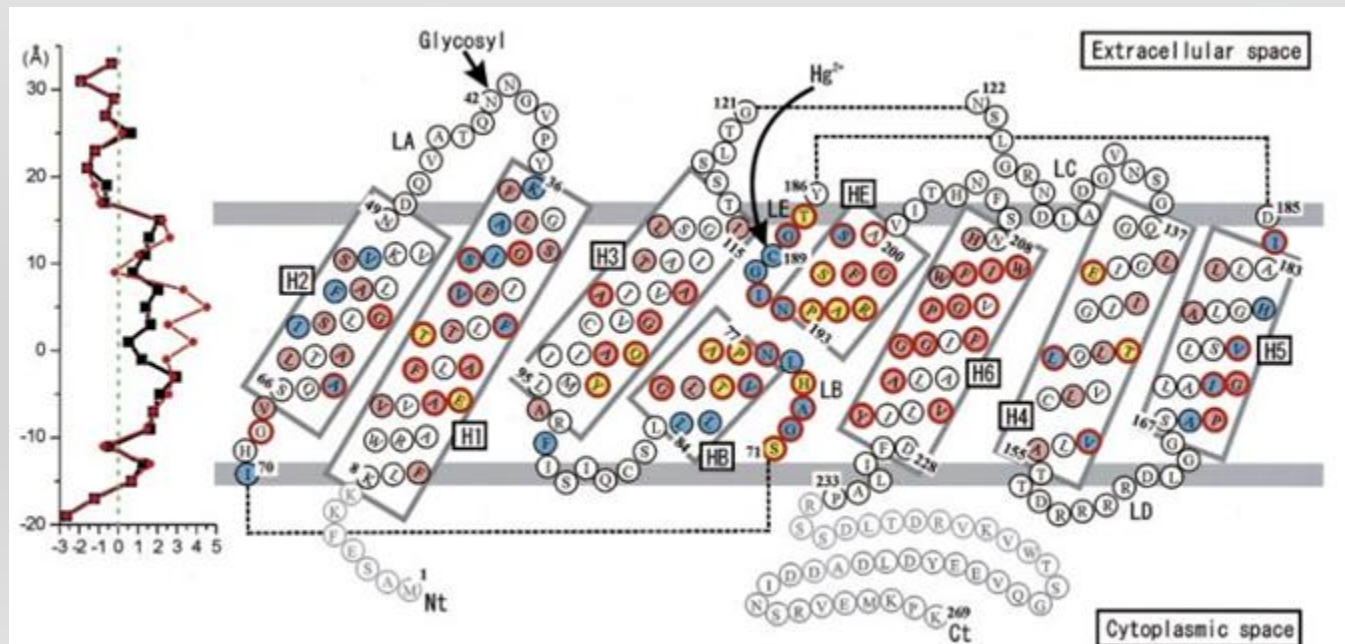
[§] Department of Cell Biology, Harvard Medical School, 240 Longwood Avenue, Boston, Massachusetts, 02115-5730, USA

^{||} Departments of Biological Chemistry and Medicine, Johns Hopkins University School of Medicine, Baltimore, Maryland 21205-2185, USA

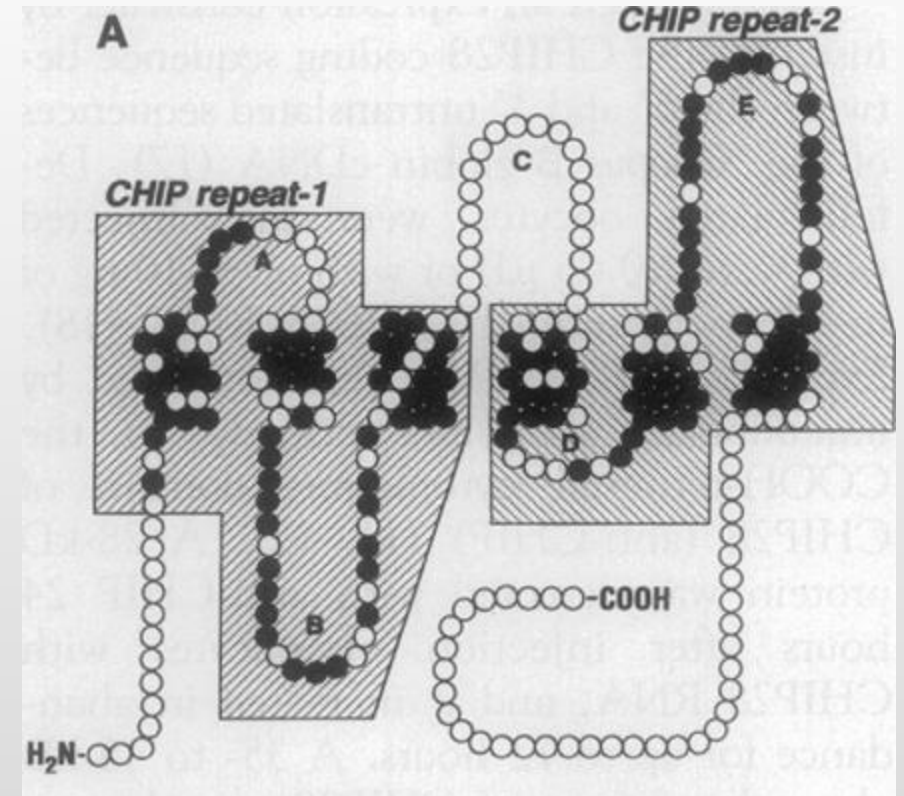
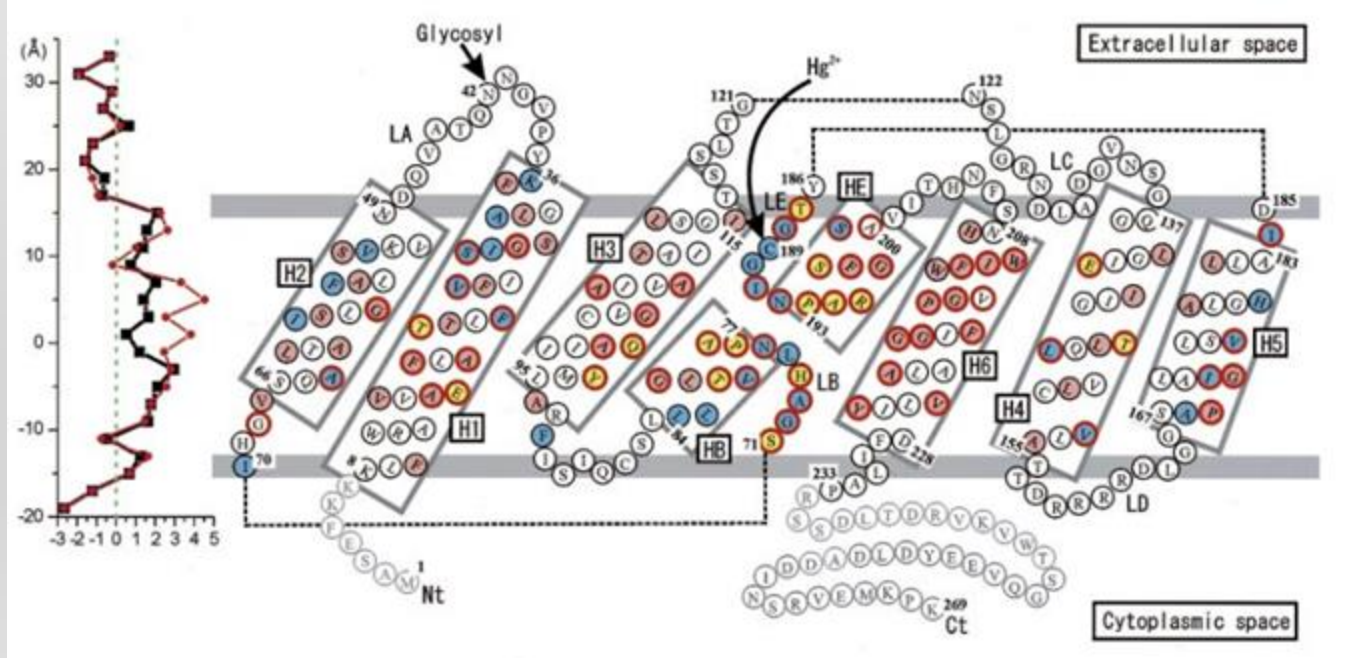
[¶] M. E. Müller-Institute for Microscopy at the Biozentrum, University of Basel, Basel CH-4056, Switzerland

[†] These authors contributed equally to this work.

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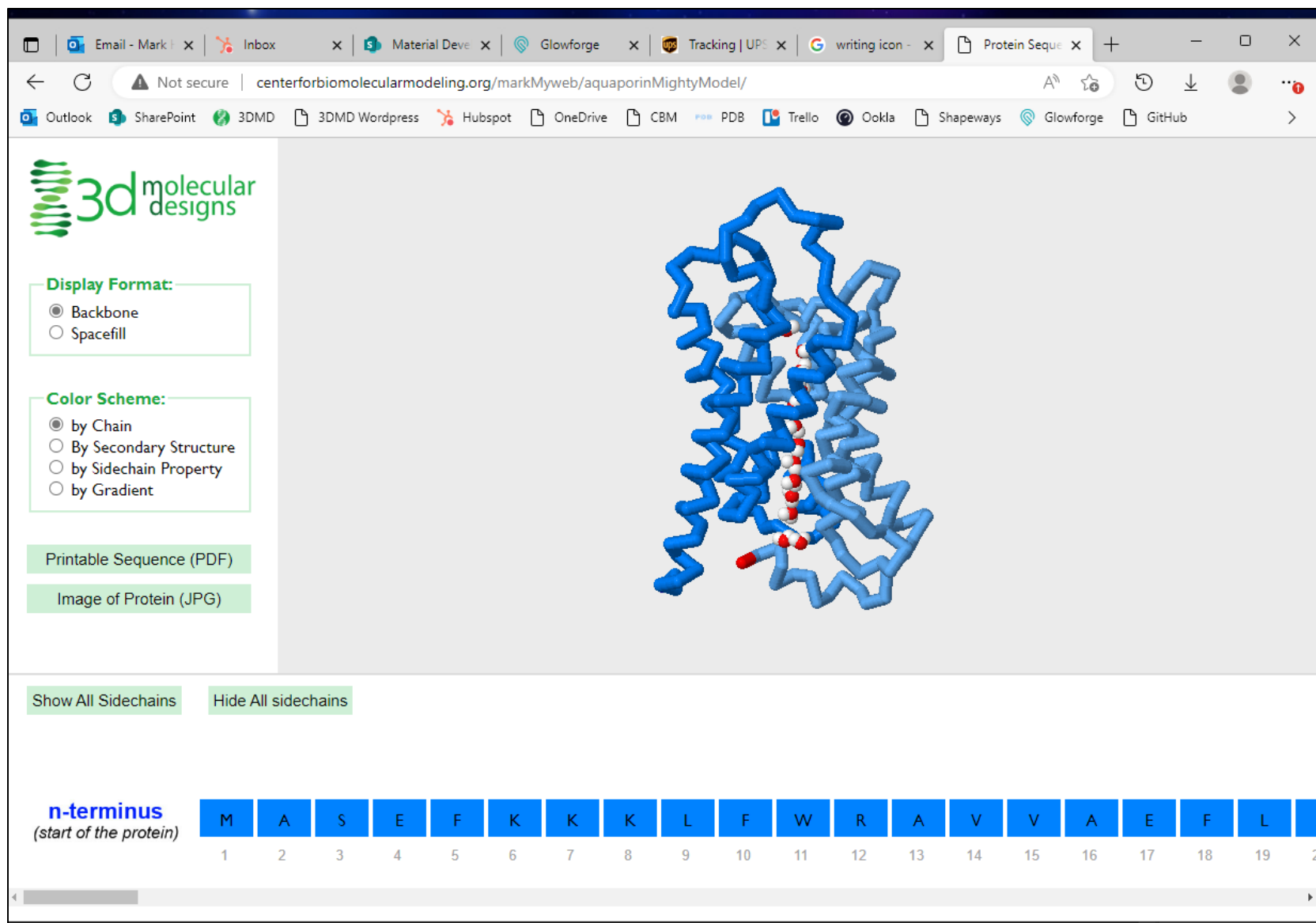


Agre “model”, 1992 →



← Murata model, ...
based on x-ray
diffraction, 2000

Protein Exploratorium



The screenshot shows a web browser window with the URL centerforbiomolecularmodeling.org/markMyweb/aquaporinMightyModel/. The page features the 3d molecular designs logo and a 3D ribbon model of a protein structure (Aquaporin) in blue, with a red and white molecular model in the center. On the left, there are controls for 'Display Format' (Backbone selected, Spacefill unselected) and 'Color Scheme' (by Chain selected, By Secondary Structure, by Sidechain Property, and by Gradient unselected). Below these are buttons for 'Printable Sequence (PDF)' and 'Image of Protein (JPG)'. At the bottom, there are buttons for 'Show All Sidechains' and 'Hide All sidechains', and a sequence viewer showing the n-terminus (start of the protein) with amino acids: M, A, S, E, F, K, K, K, L, F, W, R, A, V, V, A, E, F, L.

Display Format:

- ☒ Backbone
- ☐ Spacefill

Color Scheme:

- ☒ by Chain
- ☐ By Secondary Structure
- ☐ by Sidechain Property
- ☐ by Gradient

Printable Sequence (PDF)

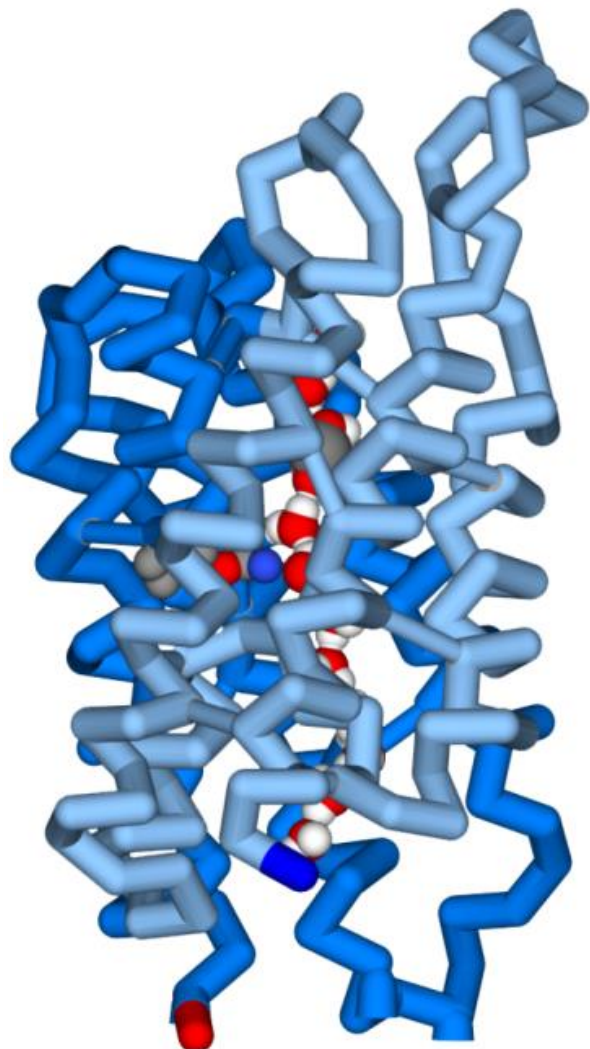
Image of Protein (JPG)

Show All Sidechains Hide All sidechains

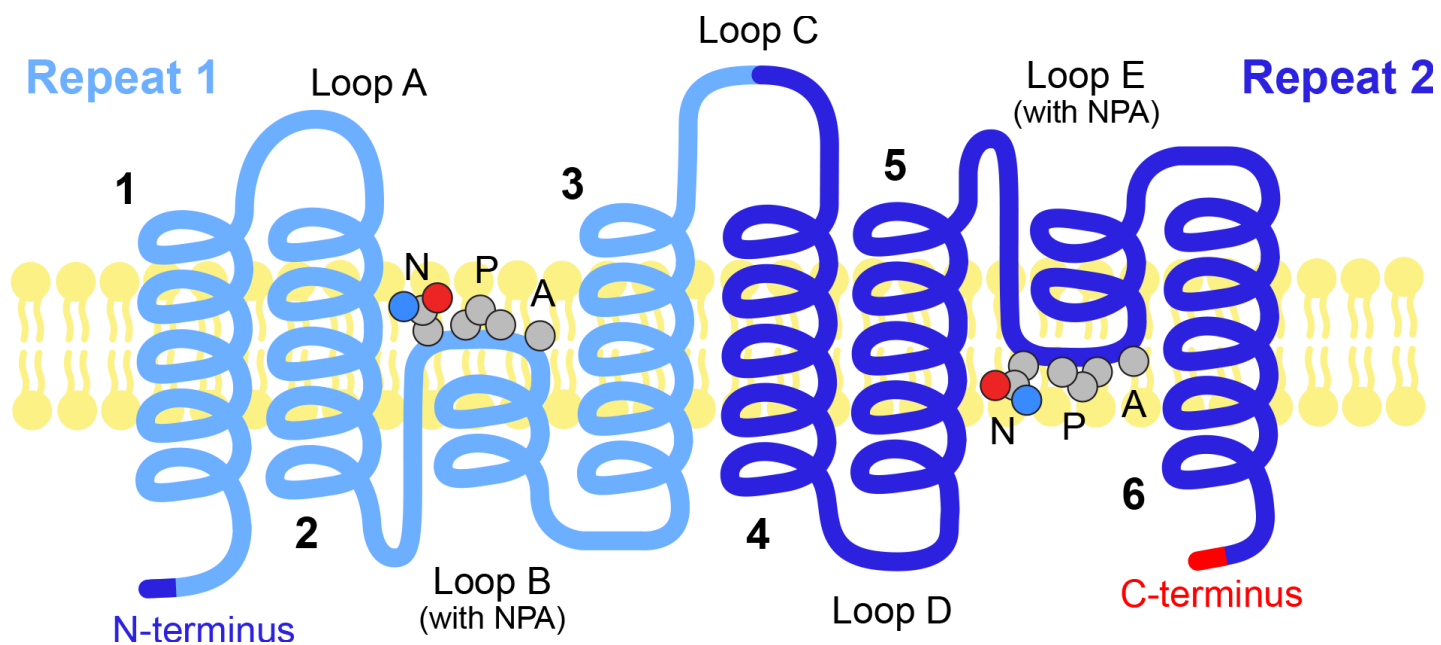
n-terminus
(start of the protein)

M	A	S	E	F	K	K	K	L	F	W	R	A	V	V	A	E	F	L
1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19

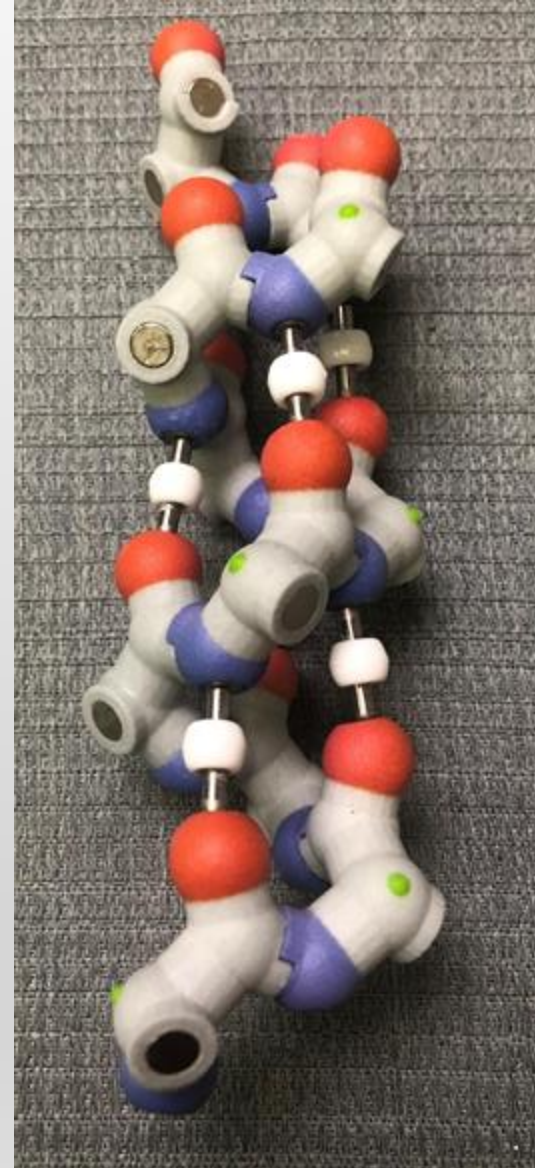




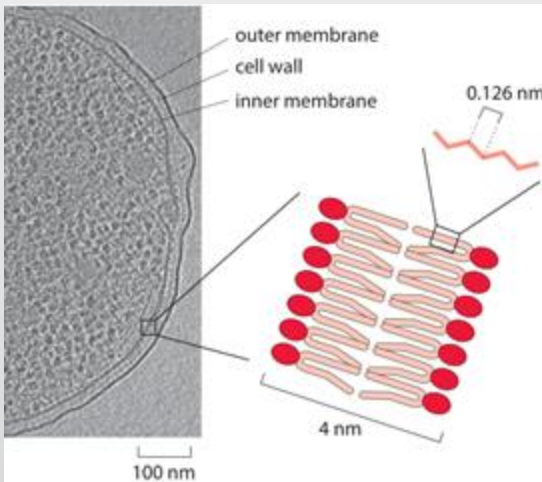
What is Aquaporin?



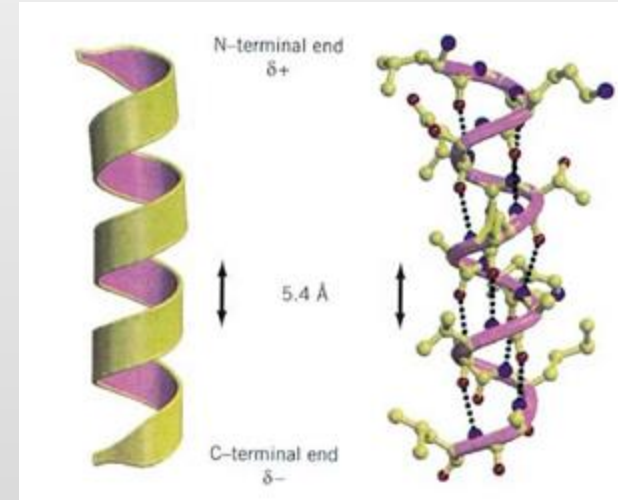
Why are transmembrane proteins often composed of bundles of alpha helices?



How many turns of the helix are required to span a membrane?



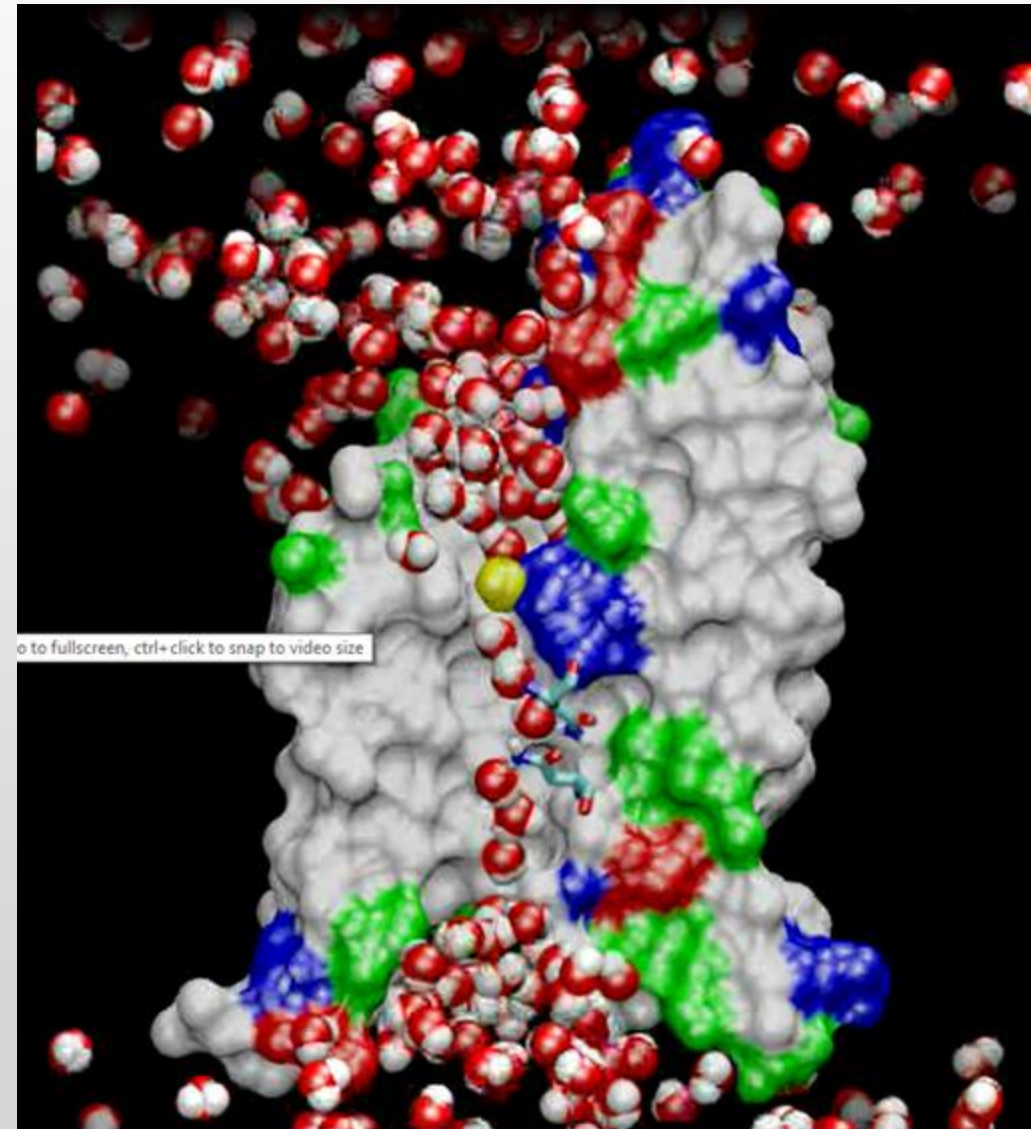
A lipid bilayer is ~ 3nm thick



An alpha helix makes on complete turn in 0.54 nm

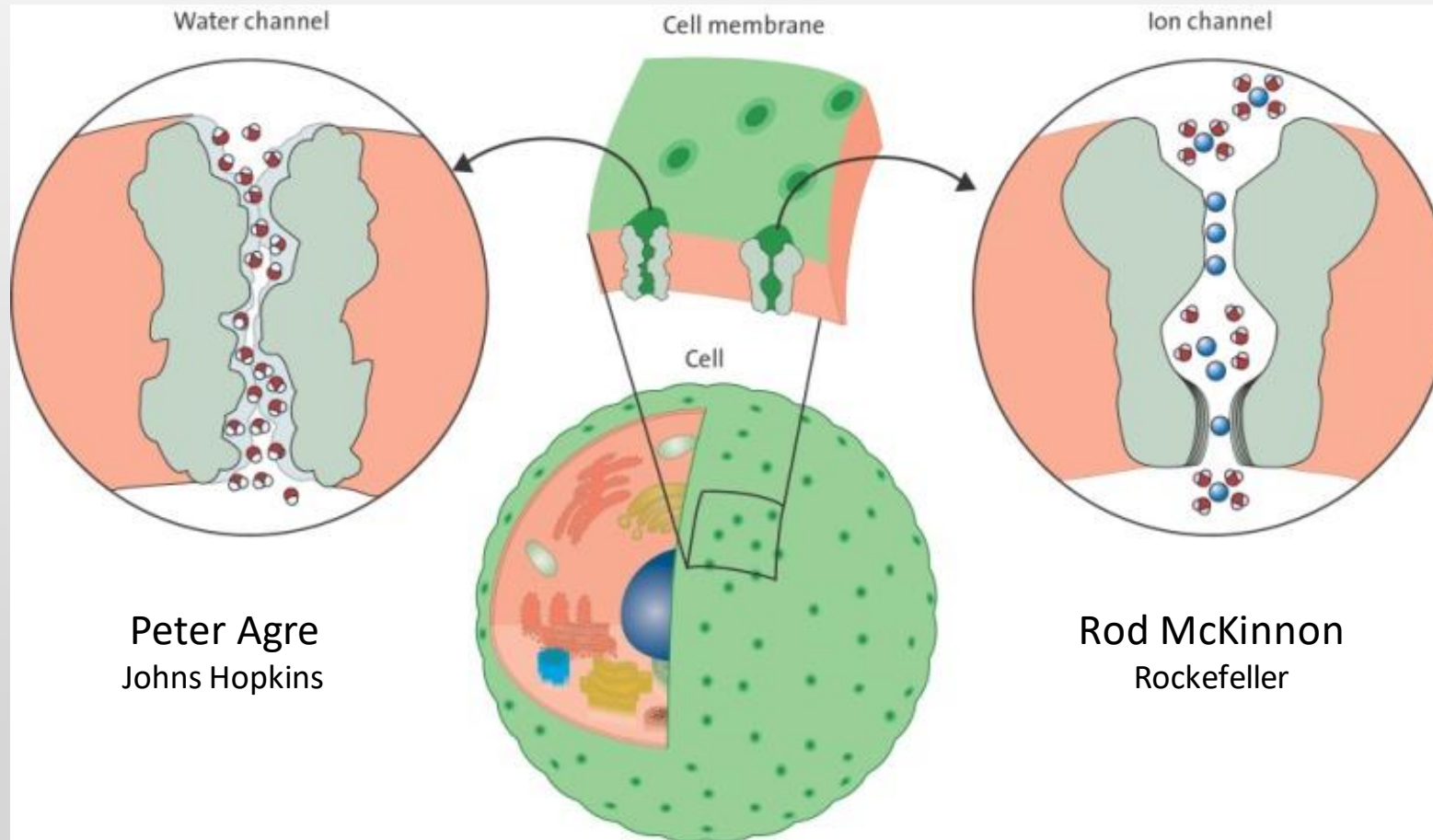
$3\text{nm} / .54 \text{ nm/turn} = \sim 6 \text{ turns}$ of an alpha helix to span a membrane.
 (22 amino acids)

A molecular dynamics simulation from the Klaus Shulten group at UIUC



<https://www.ks.uiuc.edu/Research/aquaporins/>

The 2003 Nobel Prize in Chemistry



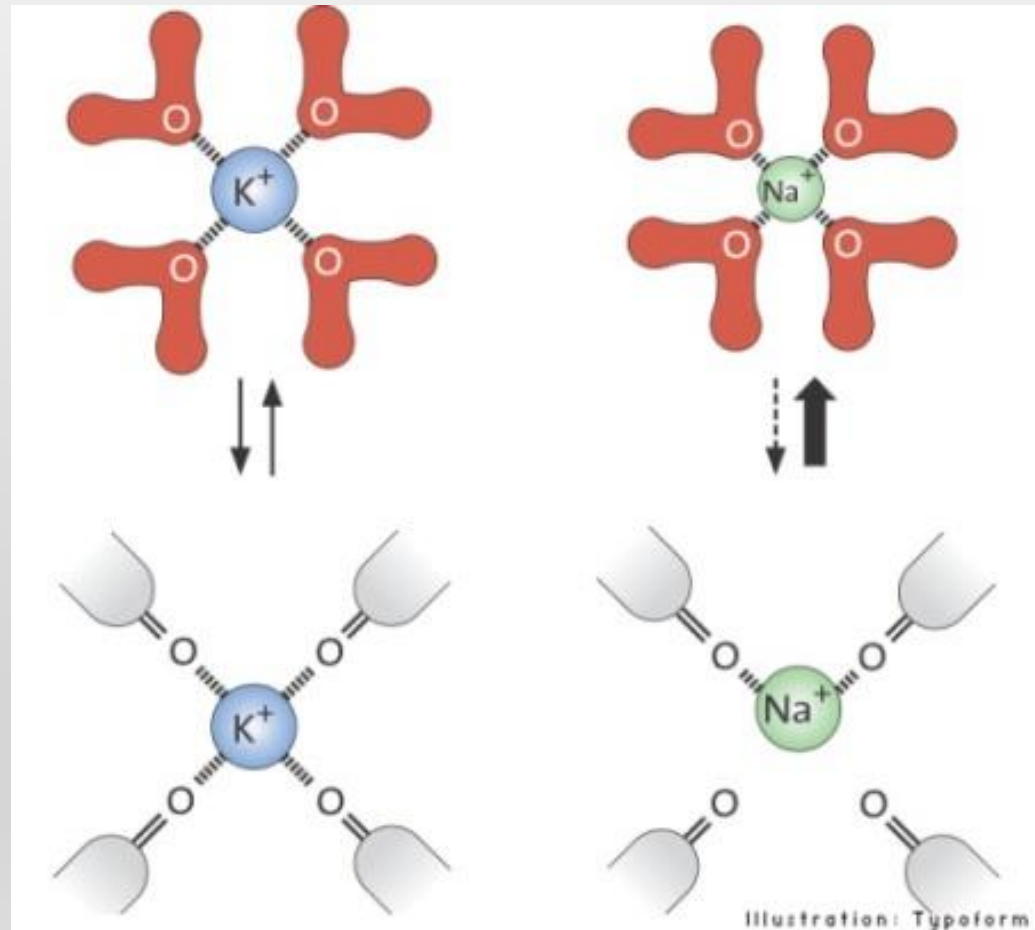
The ion channels.... for K^+ and Na^+

These two ion channels are responsible for the action potentials that drive our neurons...

- *They are very fast...*
- *They are very specific...*
- *They are regulated by voltage sensor domains...*



Why K^+ ,... but not Na^+



...and now,... yet another Nobel Prize-winning Channel Protein

2021 Nobel Prize in Physiology
or Medicine to **David Julius**

The Pain Receptor



TRPV1



A 2015 SMART Team Project



Peter Agre with Aquaporin Model

