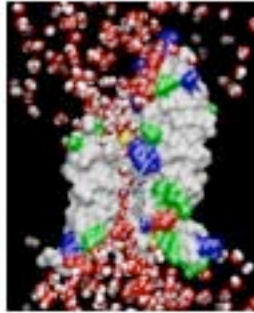


## Sequence and Structure Alignment - Illustrated for the Water Channel Aquaporin

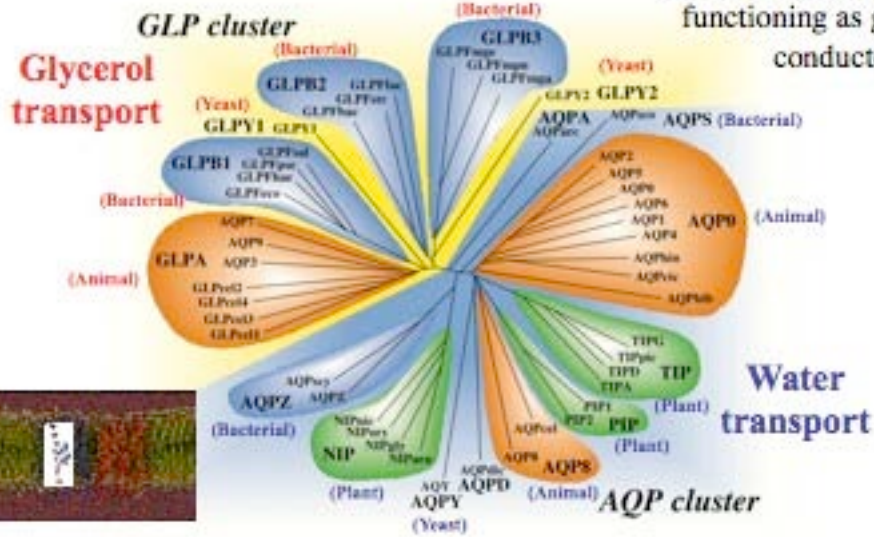


### Physical Bioinformatics - A Case Study

Sequence and structure information are the bedrock on which an understanding of cellular functions and the underlying physical mechanisms can be built. This lecture illustrates how the two sources of information are combined to investigate by means of the program VMD function and mechanism of the aquaporin family of membrane channels that transport water and certain small solutes across cell walls. Introducing first the key architectural features of a single aquaporin, structures and sequences of four aquaporins are aligned and common features recognized. The shared and distinct features are examined closely and used as guideposts leading quickly to key questions regarding the mechanism underlying aquaporin's efficient conduction and selection.

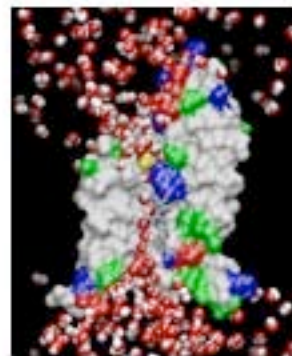
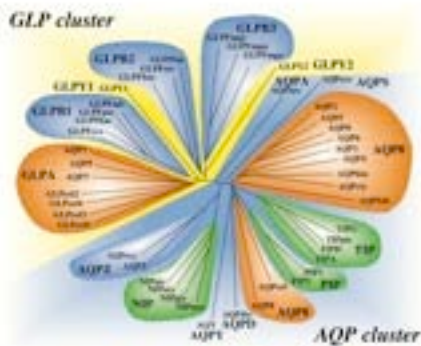
# The Aquaporin Superfamily

aquaporins are also  
functioning as gas  
conductors



Heymann and Engel *News Physiol. Sci.* 14, 187 (1999)

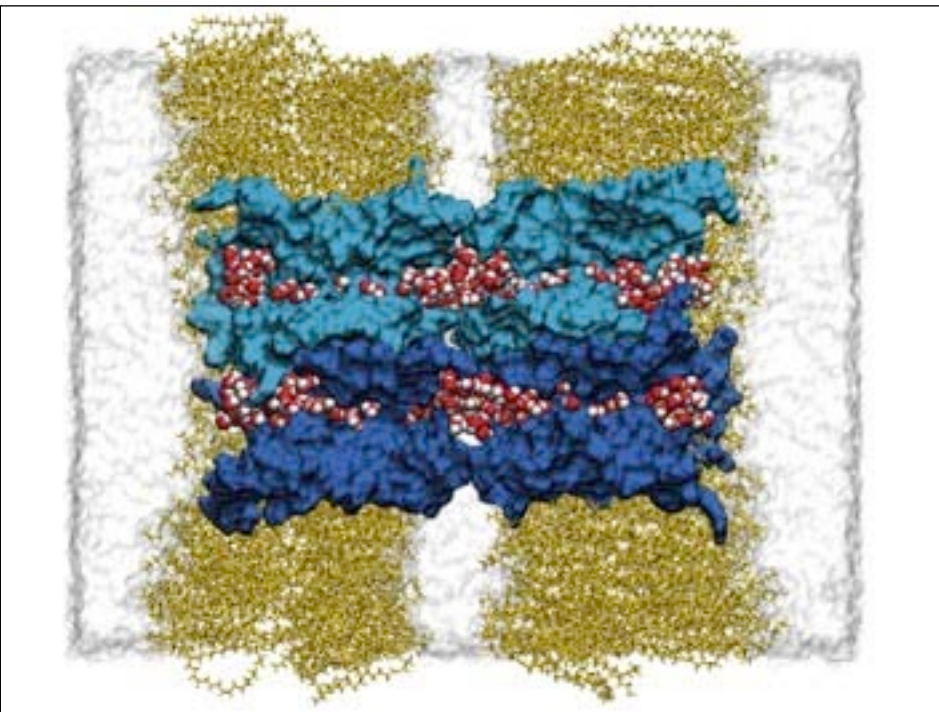
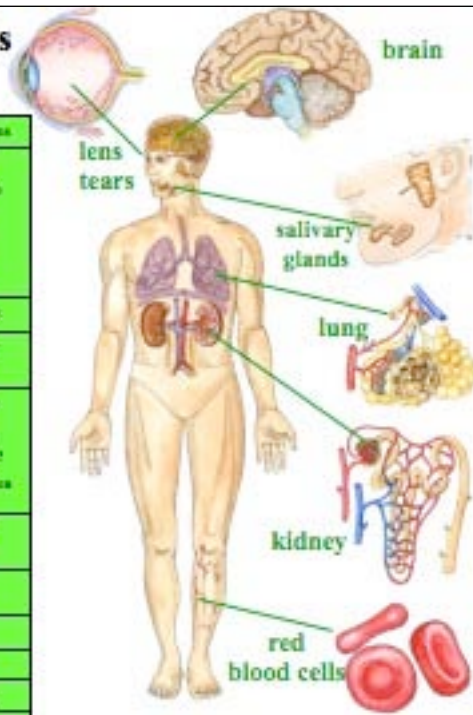
## Aquaporin Function and Human Aquaporins



AQP0 HUMAN	---LNTLHPAVVGGATVVEIFLQAVVLCIFATYDE-ERNRQLGSEVALAVGPELALGHLFQDQVGGAGN	183
AQP1 HUMAN	---RNE LADSVNQQQLGIEIIFLQVLCVLAATSR-ERRDGGHAPLAIQLFVALGHLAIDVYGGGI	191
AQP2 HUMAN	---VVALGNSVTAAGAVVVELFLLQVLCIFATYDE-ERNRQFVALVIGPELALGHLAIDVYGGGI	183
AQP3 HUMAN	KEFATTFPGLHINQVDFQPIGALIVCVLAIVGYNQVPPGLKAPVQLVYLVIGVHMDPNSGQAV	214
AQP4 HUMAN	---VTVHNSLQAGQLLVLLIIFQLVLCIFATYDE-ERTDVTGSLALGPELVALGHLAIDVYGGAGN	212
AQP5 HUMAN	---VVALGNSVTAAGAVVVELIIFQLVLCIFATYDE-ERTDVTGSLALGPELVALGHLAIDVYGGAGN	184
AQP6 HUMAN	---EYVFNHNVYVQAVAVVELLQVLCVFAITDE-NQTS-DEPDAHIGIIPALGHLAIDVYGGAGN	195
AQP7 HUMAN	KEFATTFPGLHINQVDFQPIGALIVCVLAIVGYNQVPPGLKAPVQLVYLVIGVHMDPNSGQAV	225
AQP8 HUMAN	---AAPVYVQDQVADALVAVIILQLLAVCVLAIVGYNQVPPGLKAPVQLVYLVIGVHMDPNSGQAV	209
AQP9 HUMAN	KEFATTFPGLHINQVDFQPIGALIVCVLAIVGYNQVPPGLKAPVQLVYLVIGVHMDPNSGQAV	215
GLPF ECOLI	KEFATTFPGLHINQVDFQPIGALIVCVLAIVGYNQVPPGLKAPVQLVYLVIGVHMDPNSGQAV	202
rule	...180.....190.....200.....210.....220.....230.....240....	

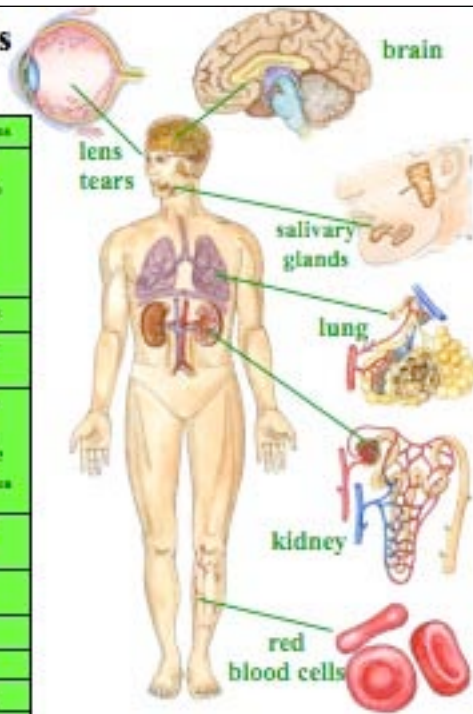
# Water and Glycerol Channels in the Human Body

Aquaporin-0	Eye: lens fiber cells	Fluid balance of the lens
Aquaporin-1	Red blood cells Kidney: proximal tubules Eye: ciliary epithelium Brain: choroid plexus Lung: alveolar epithelial cells	Osmotic protection Concentration of urine Aqueous humor Production of CSF Alveolar hydration
Aquaporin-2	Kidney: collecting ducts	ADH hormone activity
Aquaporin-3	Kidney: collecting ducts Trachea: epithelial cells	Reabsorption of water Secretion of water
Aquaporin-4	Kidney: collecting ducts Brain: astrocyte cells Brain: hypothalamus Lung: bronchial epithelium	Reabsorption of water CSF fluid balance Osmoregulation function? Bronchial fluid secretion
Aquaporin-5	Salivary glands Lacrimal glands	Production of saliva Production of tears
Aquaporin-6	Kidney	Very low water permeability!
Aquaporin-7	Testis and sperm	
Aquaporin-8	Testis, pancreas, liver	
Additional	Leakage channels are suspected to exist	



## Water and Glycerol Channels in the Human Body

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Aquaporin-8	Testis, pancreas, liver	
Additional members	are suspected to exist	



## Functionally Important Features of Aquaporins

- Water, gas, and glycerol transport
- Exclusion of ions and protons
- Tetrameric arrangement in membrane



Aquaporins of known structure:

GlpF - E. coli glycerol channel (aquaglyceroporin)

- Fu, et al., Science (2000)

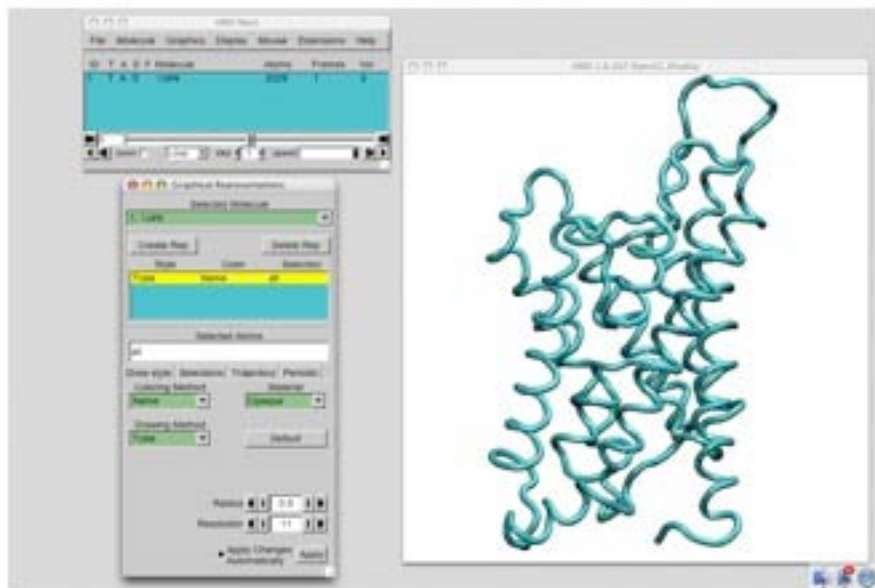
AQP1 - Mammalian aquaporin-1 (pure water channel) - Sui et al, Nature (2001)

AQP1 - Bovine - Murata et al, Nature (2000)

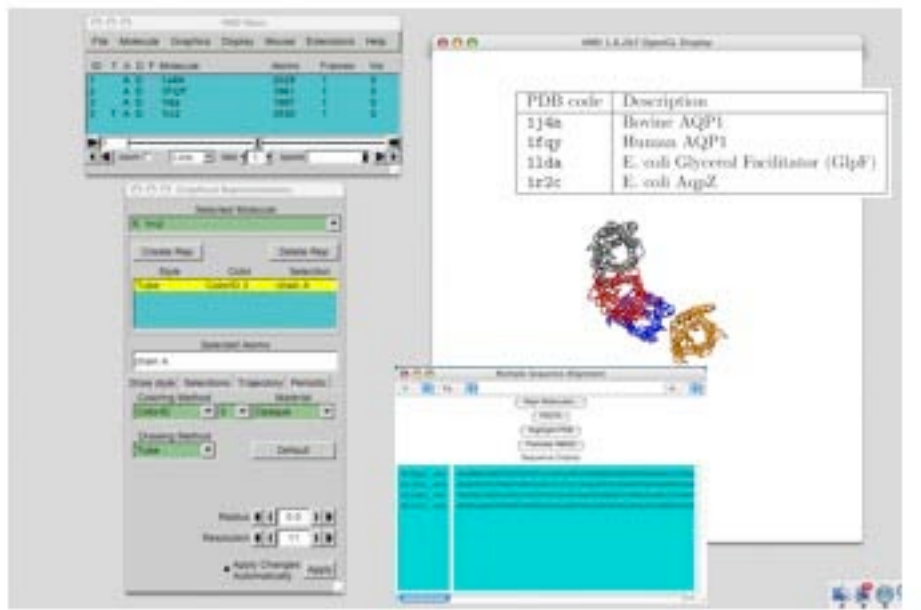
AQPZ - E. coli water channel - Savage et al, PLOS Biol (2003)



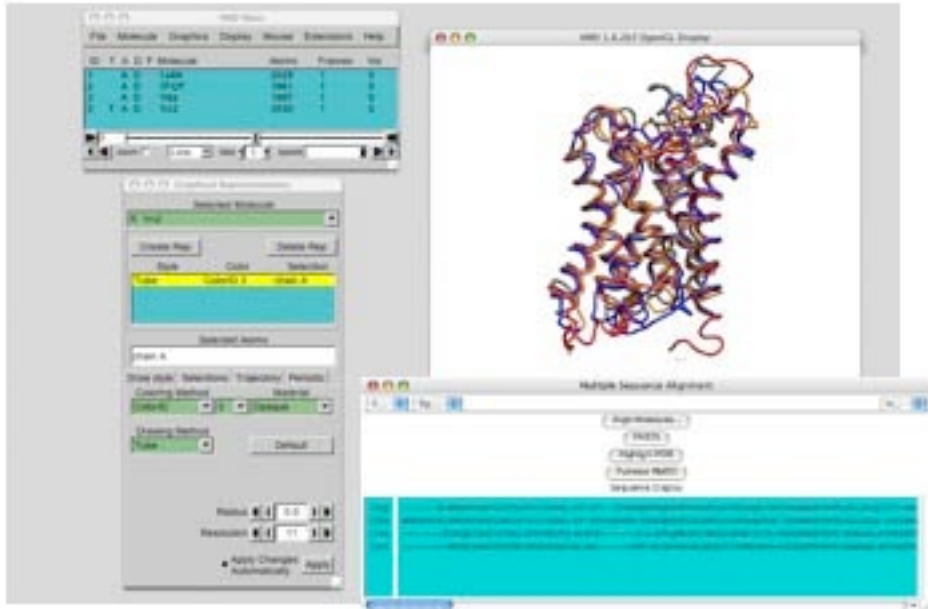
## Load Aquaporin 1J4N into VMD



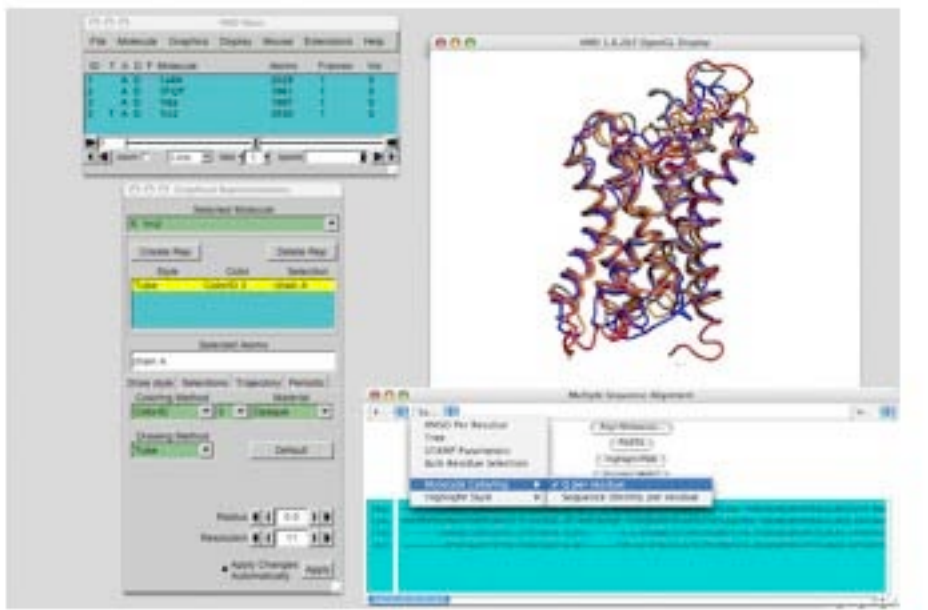
## Load Aquaporins 1j4n, 1fqy, 1lda, 1rc2 into VMD



# Aligning Structures and Sequences



# Comparing Structures by Similarity - Q Value



## Comparing Structures by Similarity - Q Value

The screenshot displays the PyMOL interface. On the left, the 'Command Line' window shows a list of molecules:

ID	Name	Residues	Parent	Vis
1	1L4K	252	1	✓
2	1L4P	252	1	✓
3	1L4Q	252	1	✓
4	1L4R	252	1	✓

The main window shows a ribbon representation of a protein structure. The 'Multiple Sequence Alignment' window at the bottom displays a sequence alignment with a 'Q-Value' column highlighted in blue, indicating the similarity score for each alignment.

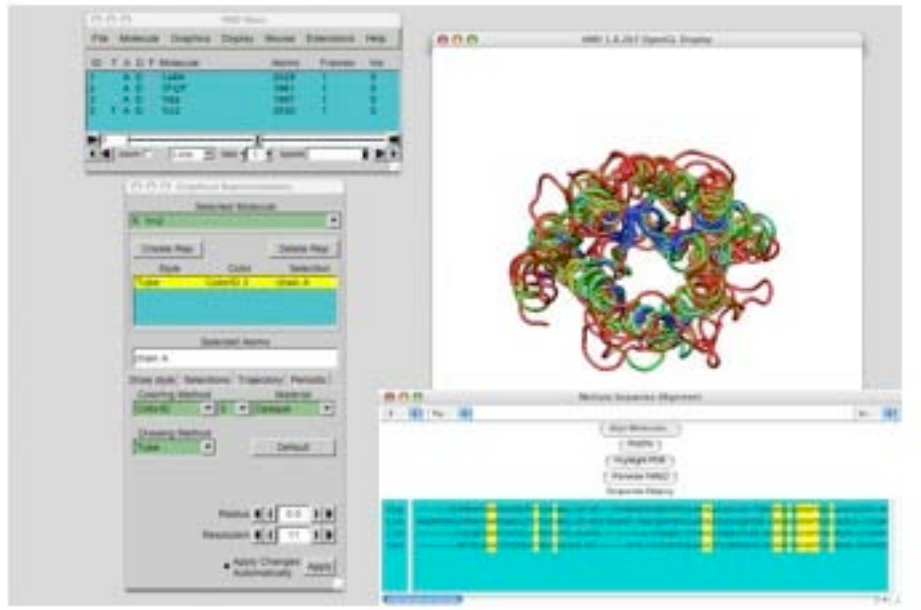
## Exhibiting Sequence Identity - Side View

The screenshot displays the PyMOL interface. On the left, the 'Command Line' window shows a list of molecules:

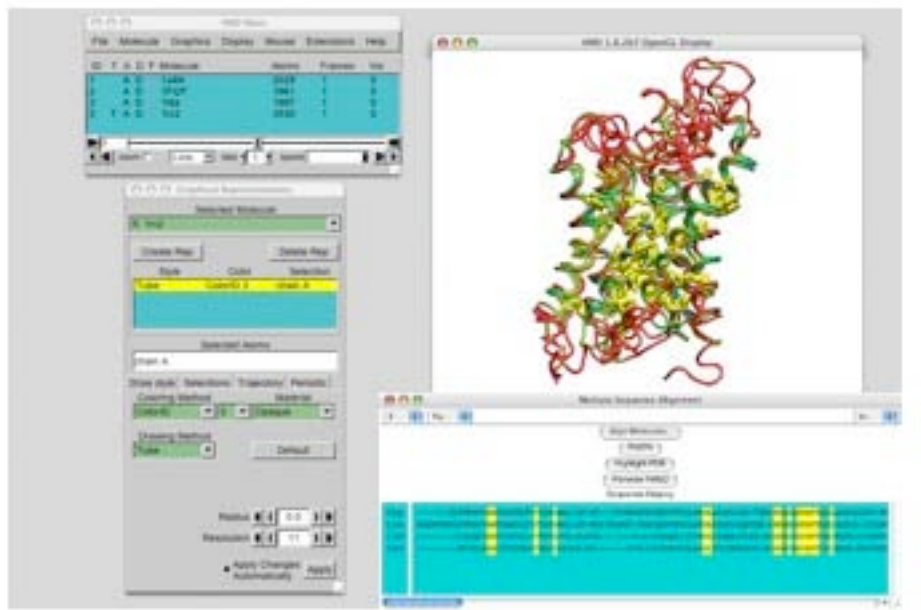
ID	Name	Residues	Parent	Vis
1	1L4K	252	1	✓
2	1L4P	252	1	✓
3	1L4Q	252	1	✓
4	1L4R	252	1	✓

The main window shows a ribbon representation of a protein structure in a side view. The 'Multiple Sequence Alignment' window at the bottom displays a sequence alignment with yellow vertical bars highlighting regions of high sequence identity across the different structures.

## Exhibiting Sequence Identity - Top View

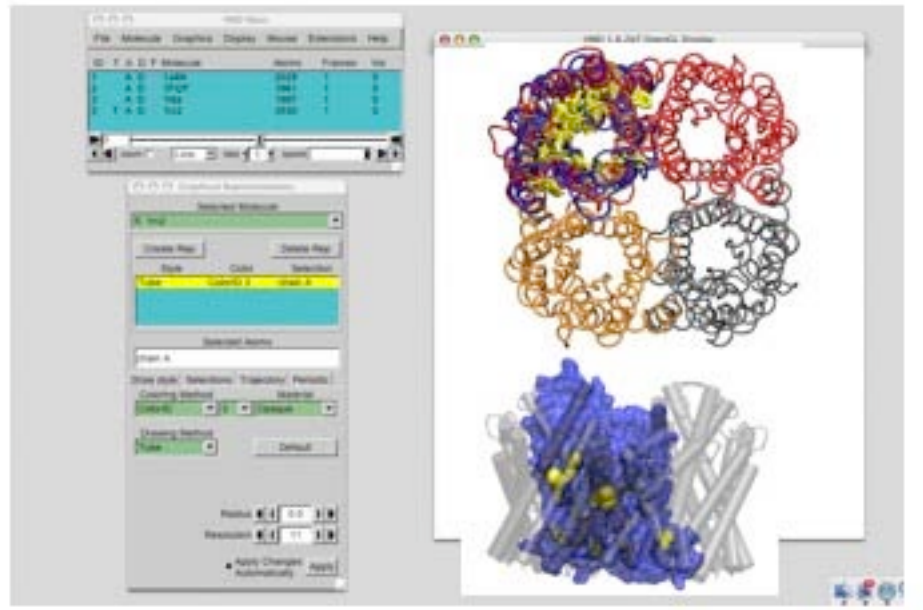


## Showing Conserved Residues - Monomer

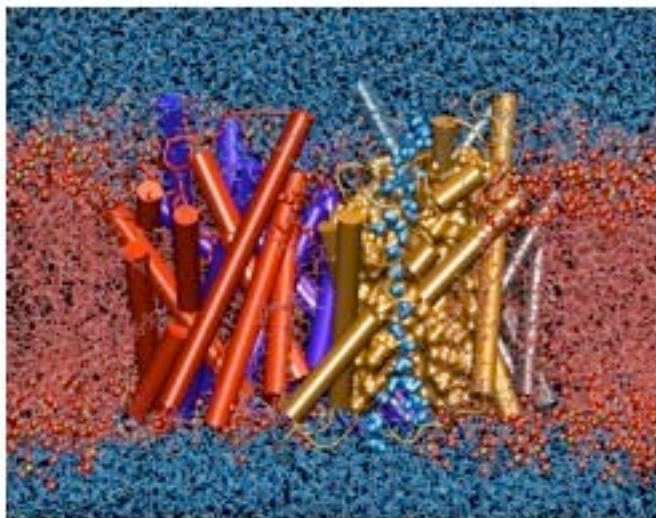




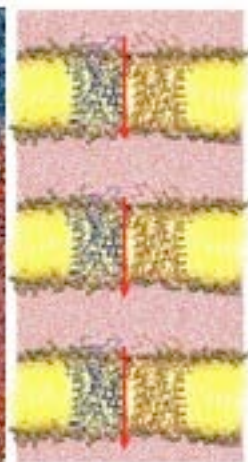
## Showing Conserved Residues - Tetramer



## Water Transport in Aquaporins



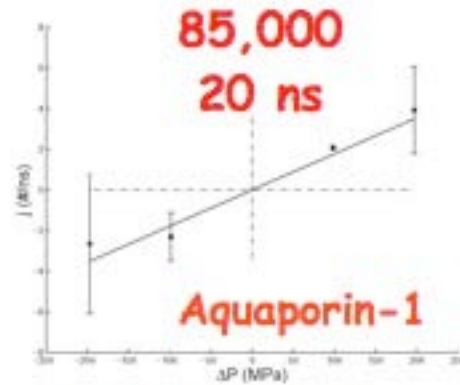
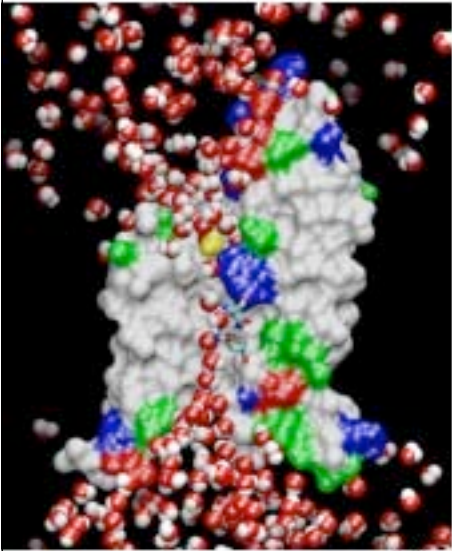
100,000 atoms



**Simulation:**

*Apply constant force  
on bulk water  
molecules*

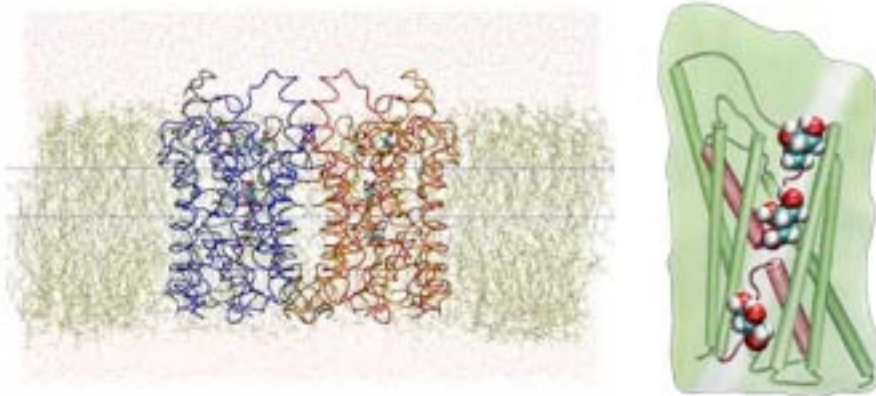
## Osmotic permeability of water channels



$p_f$ :  $7.0 \pm 0.9 \times 10^{-14} \text{ cm}^3/\text{s}$   
Exp:  $5.4 - 11.7 \times 10^{-14} \text{ cm}^3/\text{s}$

F. Zhu, E. Tajkhorshid, K. Schulten, *Biophys. J.* 86: 50-57 (2004)  
F. Zhu, E. Tajkhorshid, K. Schulten, *Phys. Rev. Lett.* 93: 224501 (2004)

## Dynamics of Protein, Lipid, Water System

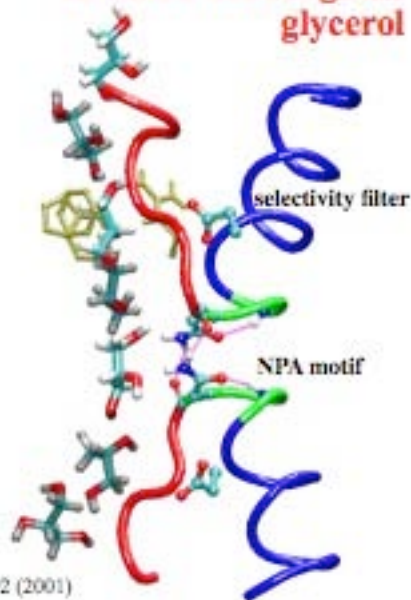


M. Jensen, E. Tajkhorshid, K. Schulten, *Structure* 9, 1083-1092 (2001)

# Glycerol Conduction



Inverted helices guide glycerol



M. Jensen, E. Tajkhorshid, K. Schulten, *Structure* 9, 1083-1092 (2001)

University of Illinois at Urbana-Champaign  
 NIH Resource for Macromolecular Modeling and Bioinformatics  
 Beckman Institute

## Aquaporins

Case study, see at  
<http://www.ks.uiuc.edu/Training/CaseStudies/>

**Aquaporins in Human Body**

More are expected to exist.  
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Dejost Dhalluin

Zem Longhui-Schulten