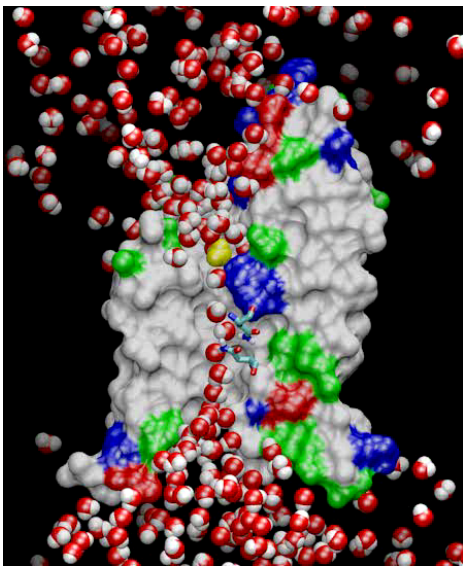
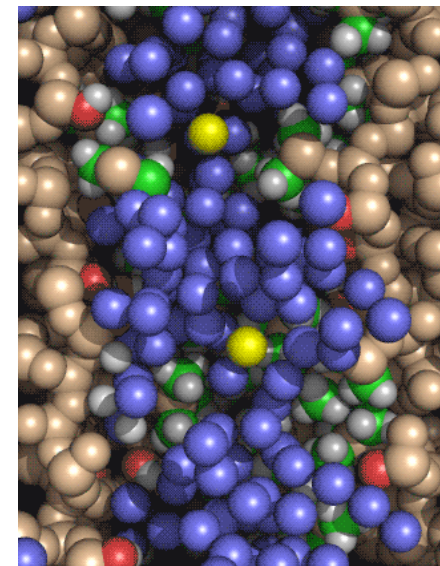


Biomimetic Membranes for Water Purification



protein



synthetic

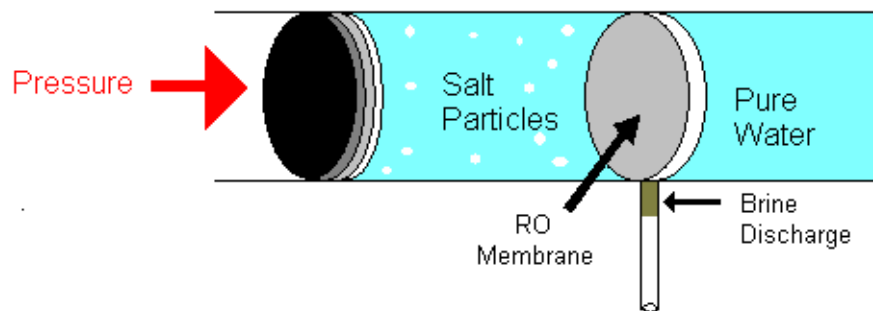
Susan Rempe
Sandia National Laboratories
Albuquerque, NM

Sandia Mountain, Albuquerque, NM



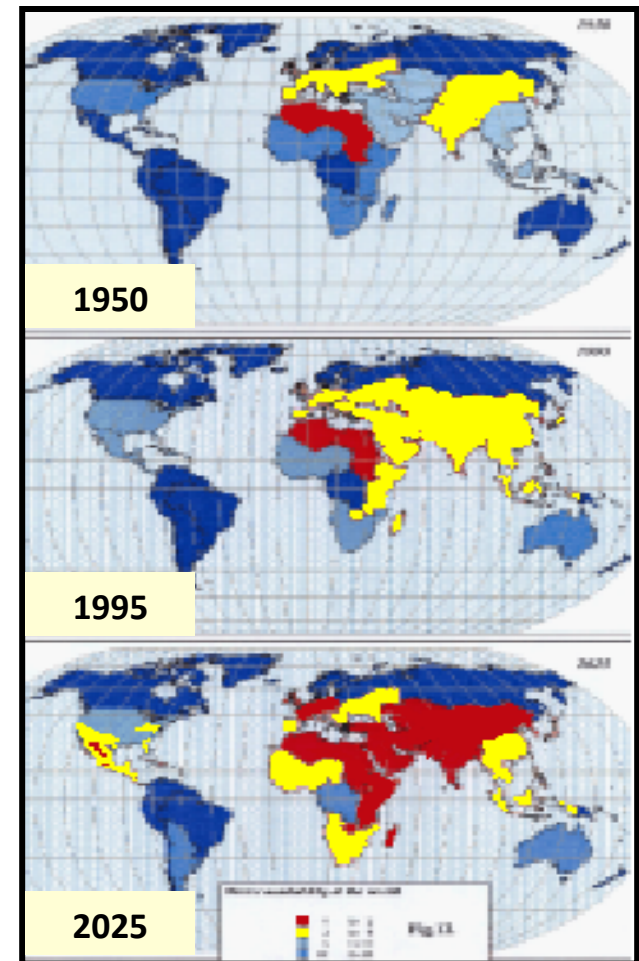
Clean Water: A Global Crisis

- Half of world's population lacks clean water:
 - public health crises
 - Water / energy link
 - RO membranes: expensive & overly sterile H₂O
 - R. Electrodialysis: high resistance to ion permeation
- Membrane needs new design for efficiency:
 - fast (**barrier-less**) water (or ion) transport
 - select ion exclusion (mineral water)



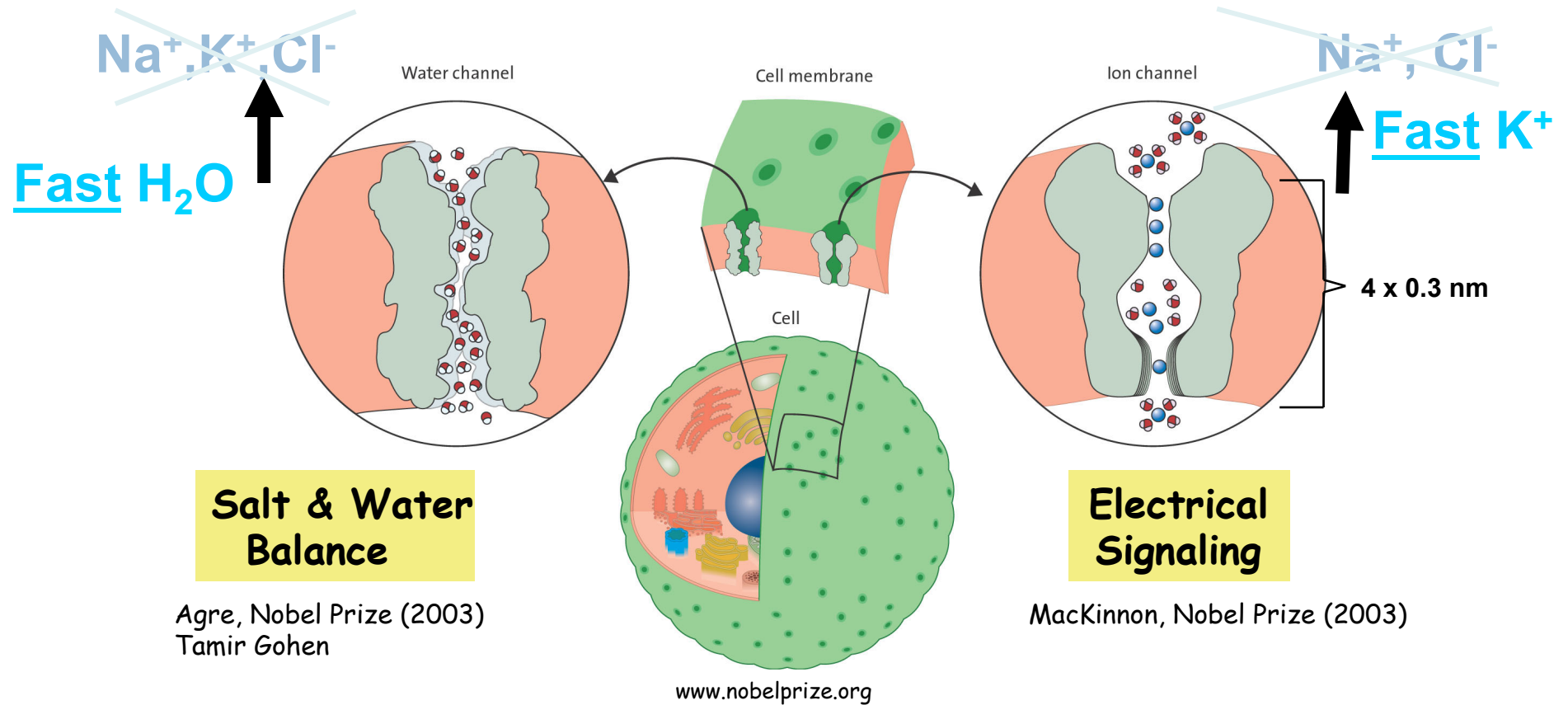
“Water promises to be to the 21st century what oil was to the 20th century: the precious commodity that determines the wealth of nations.”

Fortune Magazine, May 15, 2000



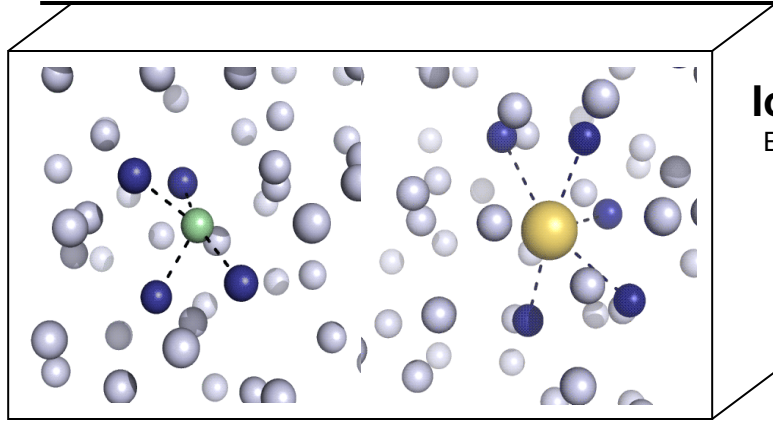
shortage

Biological Channels Inspire: Catalyze fast, selective permeation



- **Structurally Similar / Diverse Functions:**
 - General: narrow + thin + uniform/robust
 - Specific: polar

Natural protein channels: Not just simple holes!



Ions in water
Biophys Chem 2006

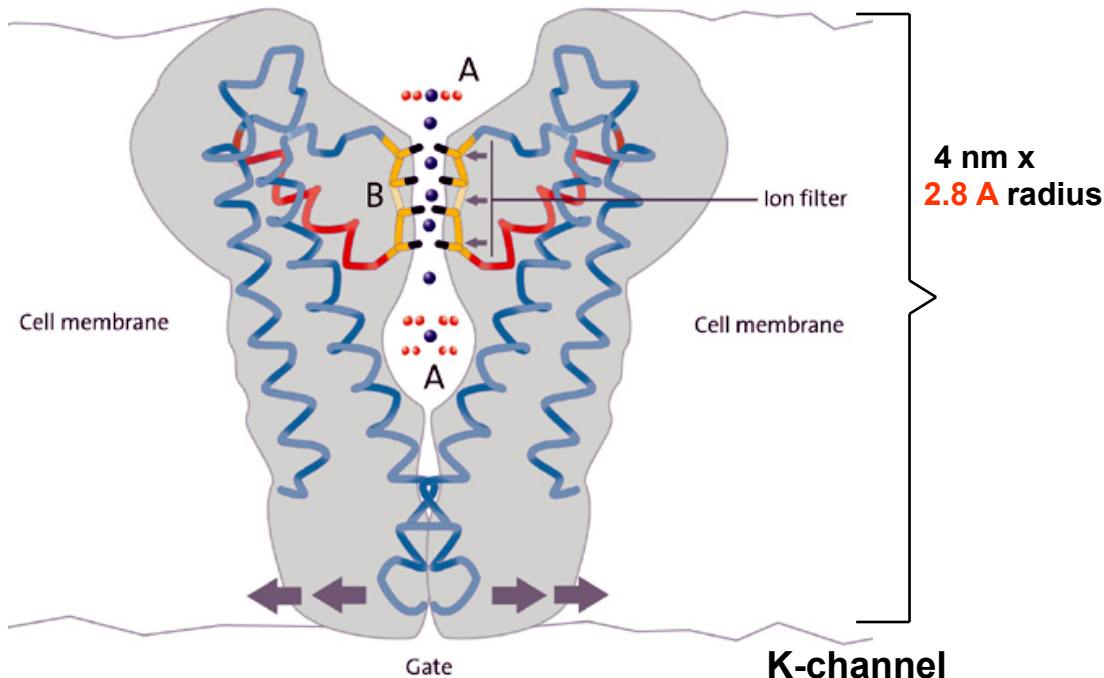
- Ions 'happy' in liquid water

How do channels work?

- K^+ equa-energy?
- Na^+ barrier?
- same structure – varied function?

Smaller Na^+
rejected

Larger K^+
permeate fast



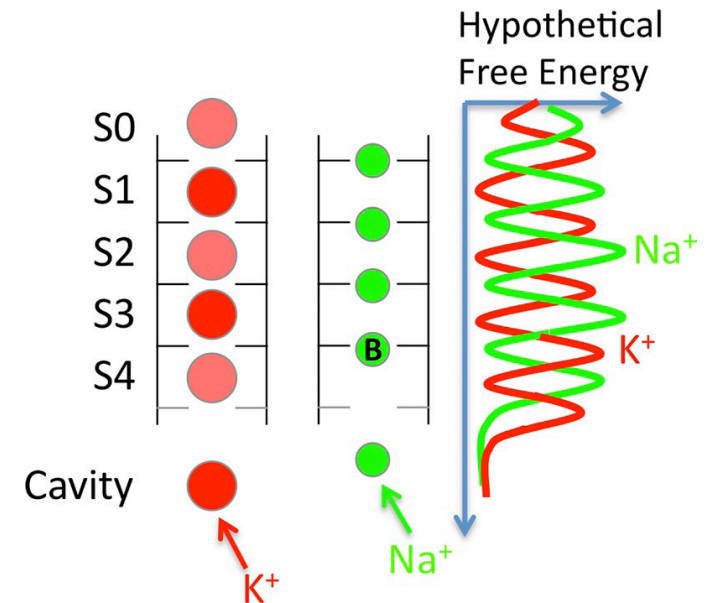
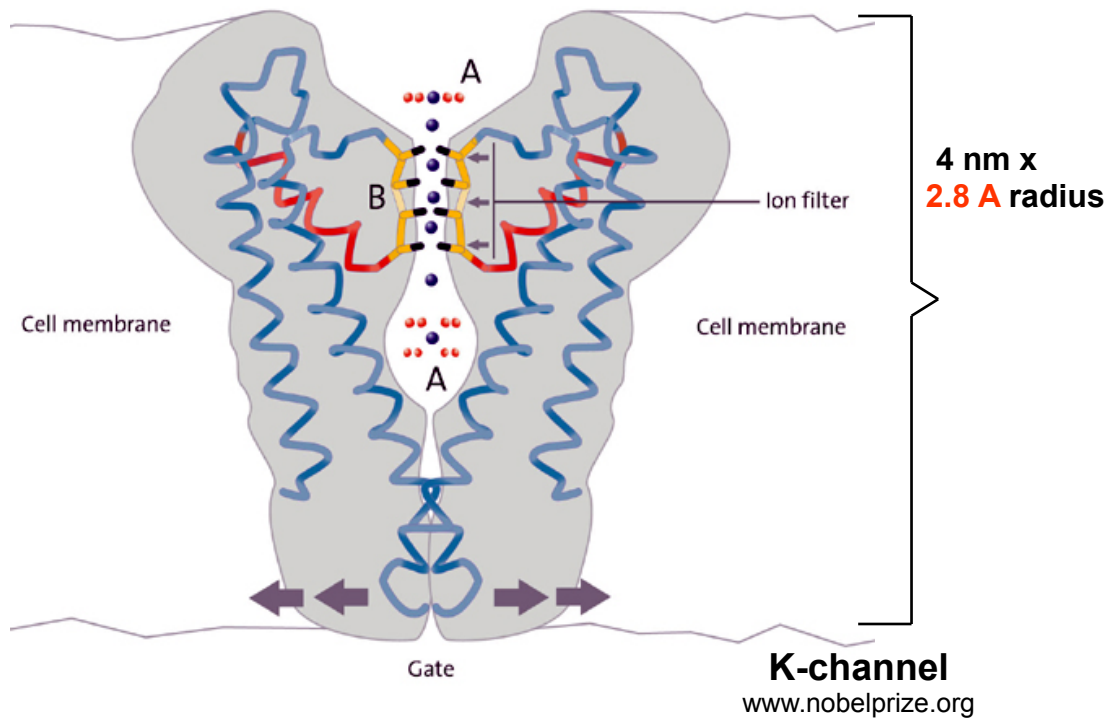
K-channel
www.nobelprize.org

Natural protein channels: Not just simple holes!

- Ions 'happy' in liquid water

How do channels work?

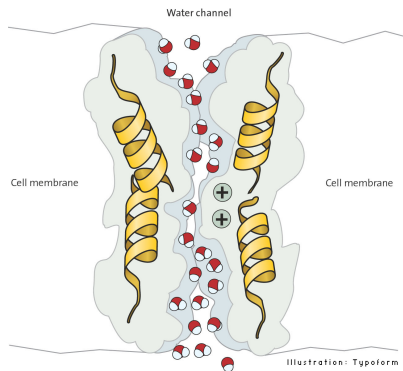
- K^+ equa-energy?
- Na^+ barrier?
- same structure – varied function?



Efficient Membranes: Biomimetic

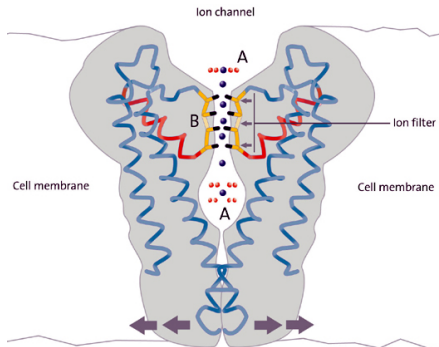
Understand, design, engineer nano-channels for desalination

Water channels



- transport H₂O fast

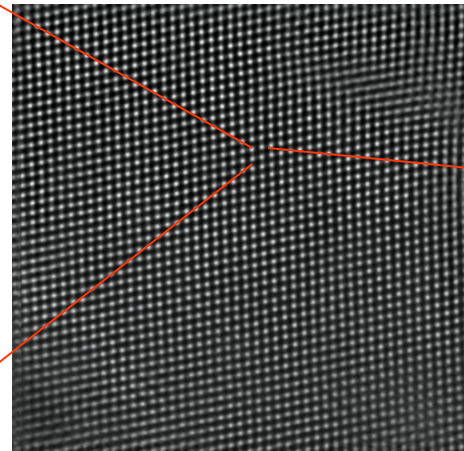
Ion channels



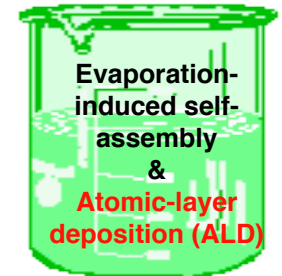
- select minerals fast

- Bio-inspired design

- Theory to reveal mechanism



- Inorganic membrane synthesis



Brinker Lab

J Membr Sci (2007)

JACS (2007)

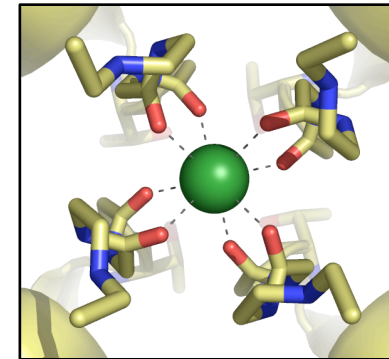
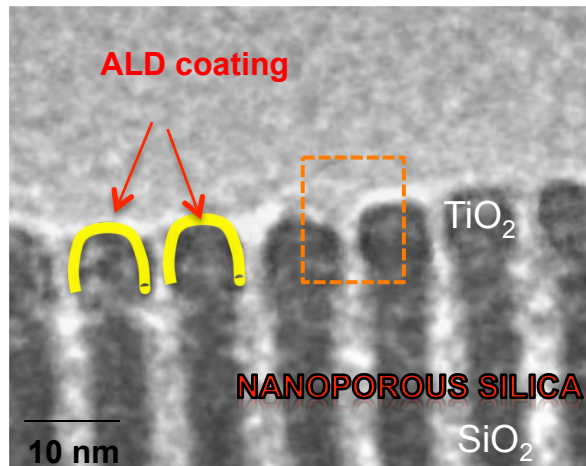
Solution: Harness molecular biomechanisms.
Gain 100x in water flux + minerals.

Models to study ion discrimination

- Link pore structure -- solvation -- & water/ion mobility

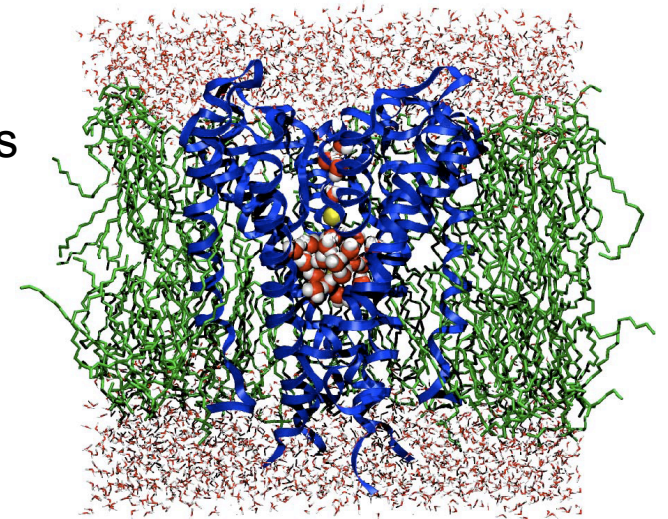
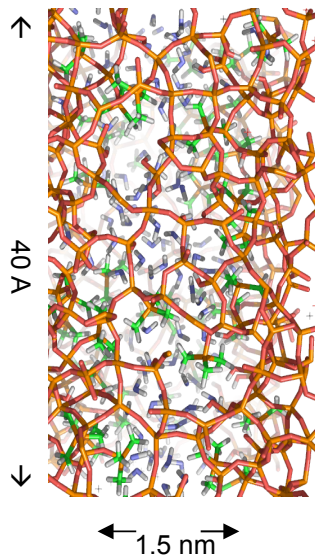
(ΔG)

(D)



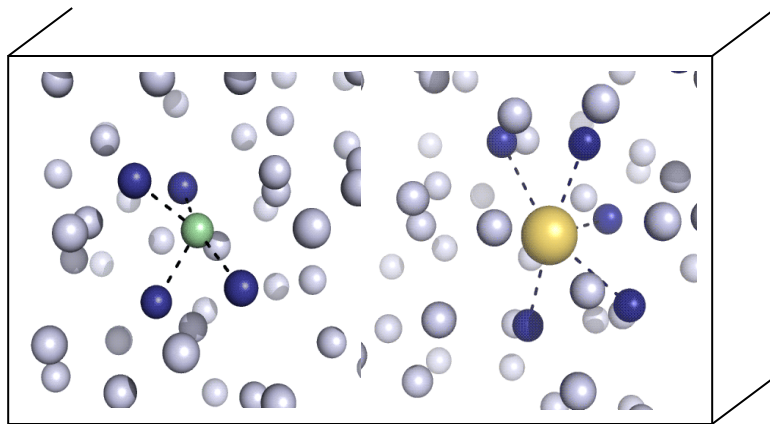
Synthetic ← → Biological

Whole Channels



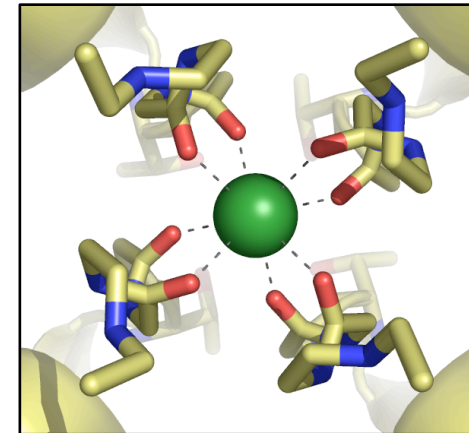
Modeling to study ion discrimination

- Link pore structure -- solvation -- & water/ion mobility
(ΔG) (D)



Dynamic structures

$$F = ma$$



Static structures

$$F = 0$$

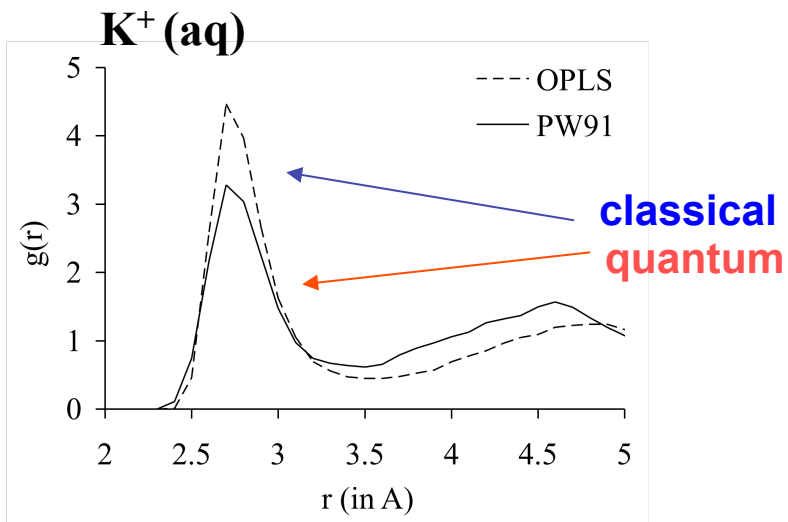
Forces?

capture discriminating interactions

- **Quantum (*ab initio*) interactions:**
 - expensive; describe multi-body interactions
 - **Classical atomic interactions:**
 - inexpensive; parameterized
-

• Example 1

Classical ions bind more ligands??



Whitfield, Varma, Rempe, Roux et al. *JCTC* (2007)
Leung & Rempe, *PCCP* (2006) Ab initio rigid water
Klein & co, *JCP* (2013) Dispersion over-structures

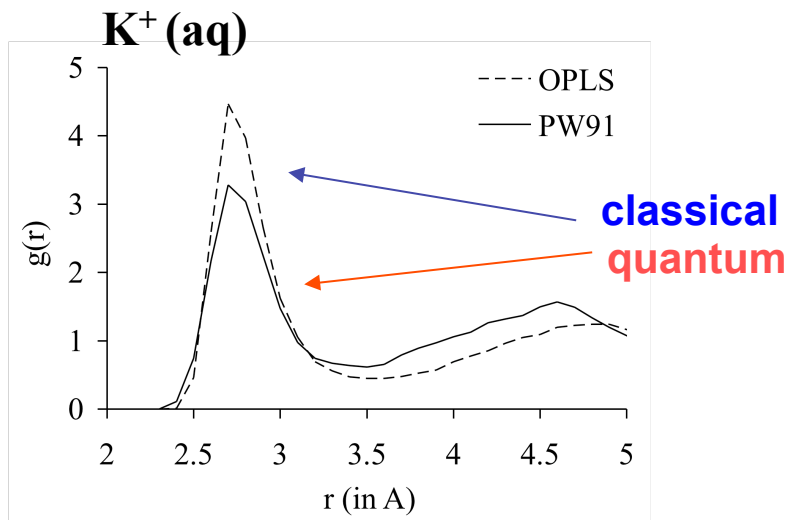
Forces?

capture discriminating interactions

- **Quantum (*ab initio*) interactions:**
 - expensive; describe multi-body interactions
- **Classical atomic interactions:**
 - inexpensive; parameterized

• Example 1

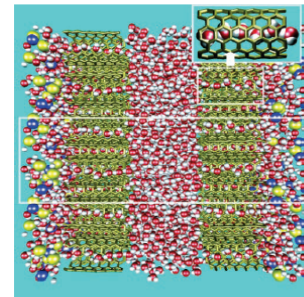
Classical ions bind more ligands??



Whitfield, Varma, Rempe, Roux et al. *JCTC* (2007)
 Leung & Rempe, *PCCP* (2006) Ab initio rigid water
 Klein & co, *JCP* (2013) Dispersion over-structures

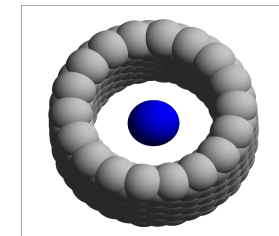
• Example 2

Classical nanotubes
Exclude ions??



Hummer & co *PNAS* (2003)

Quantum tubes
Admit ions!!



$Na^+ : -210$ kJ/mol

Leung, Marsman *JCP* (2007)

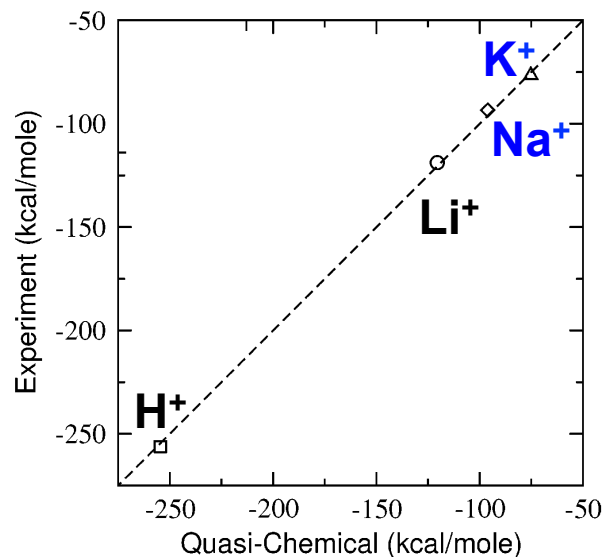
“When in doubt,
solve the Schroedinger equation”

- courtesy P. Jeff Hay, LANL

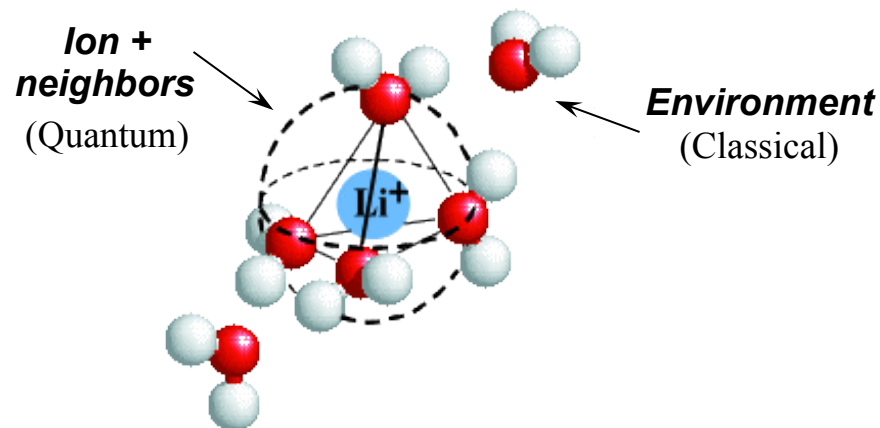
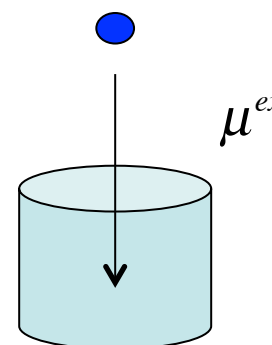
... locally!

Apply QM locally via **Q**uasi-**C**hemical **T**heory:

- Theory well-tested for ion hydration:



JACS (2004)

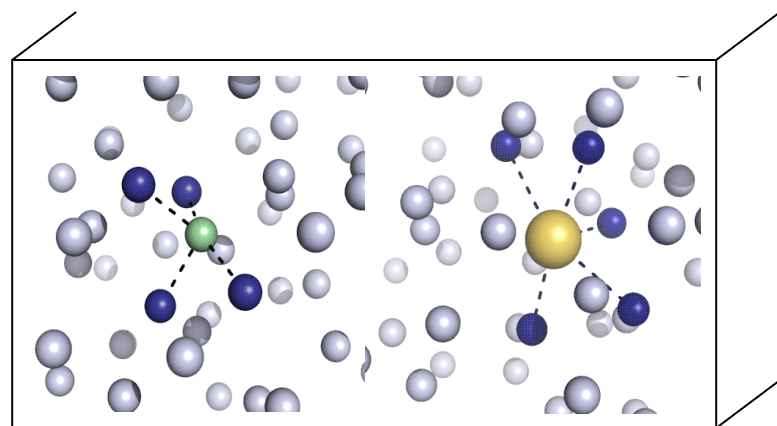


- Adaptable to ion solvation in binding sites??

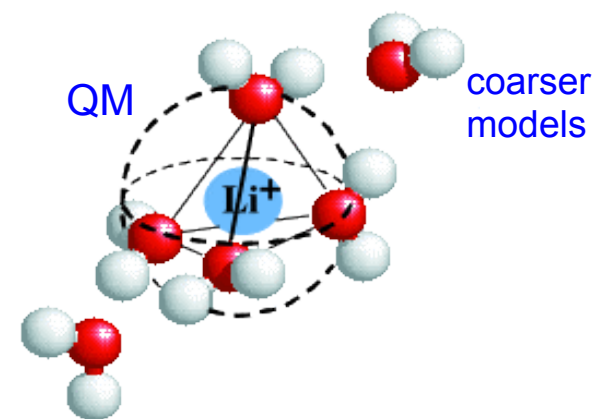
State-of-the-art all-QM for Na⁺(aq) agrees with QCT:
Leung, Rempe, von Lilienfeld *JCP* (2009)

QCT (Pratt & co):
Pratt & Rempe 'Red Book' (1999),
Chem Phys Lett (2010), *Ann Rev Comp Chem* (2012)

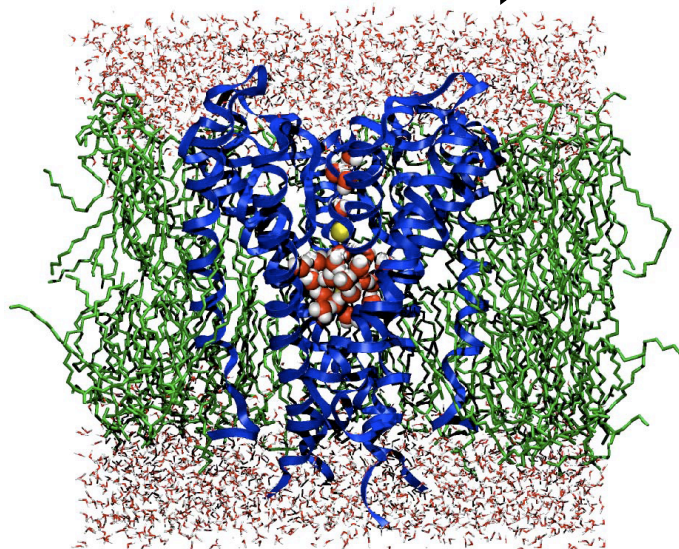
Same theory for ion solvation in channels:



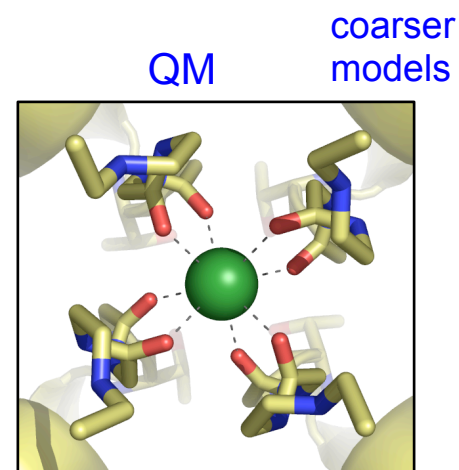
Ions in water



$\Delta\Delta G$



Ions in channels



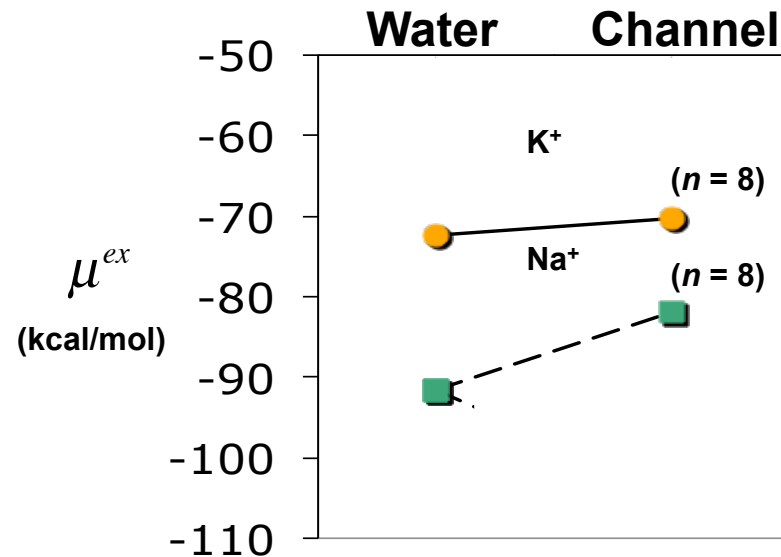
Ion Selectivity = free energy differences

- Ion solvation: local + environment

$$\mu_{\text{Ba}^{2+}}^{(\text{ex})} = \underbrace{-kT \ln K_n^{(0)} \rho_{\text{H}_2\text{O}}^n}_{\text{chemistry + ligand density}} + \underbrace{kT \ln p_{\text{Ba}^{2+}}(n)}_{\text{environment}} + \mu_{\text{Ba}(\text{H}_2\text{O})_n^{2+}}^{(\text{ex})} - n\mu_{\text{H}_2\text{O}}^{(\text{ex})}$$

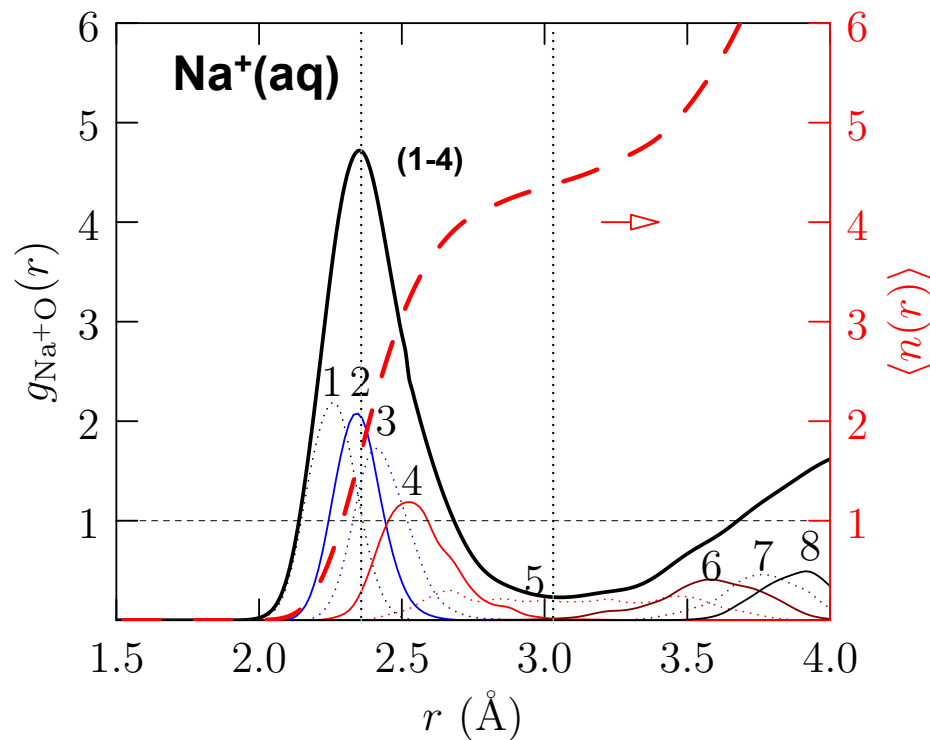
absolute value

- Selectivity: subtle balance between hydration vs solvation by channel



Ab initio structures tested against experiment

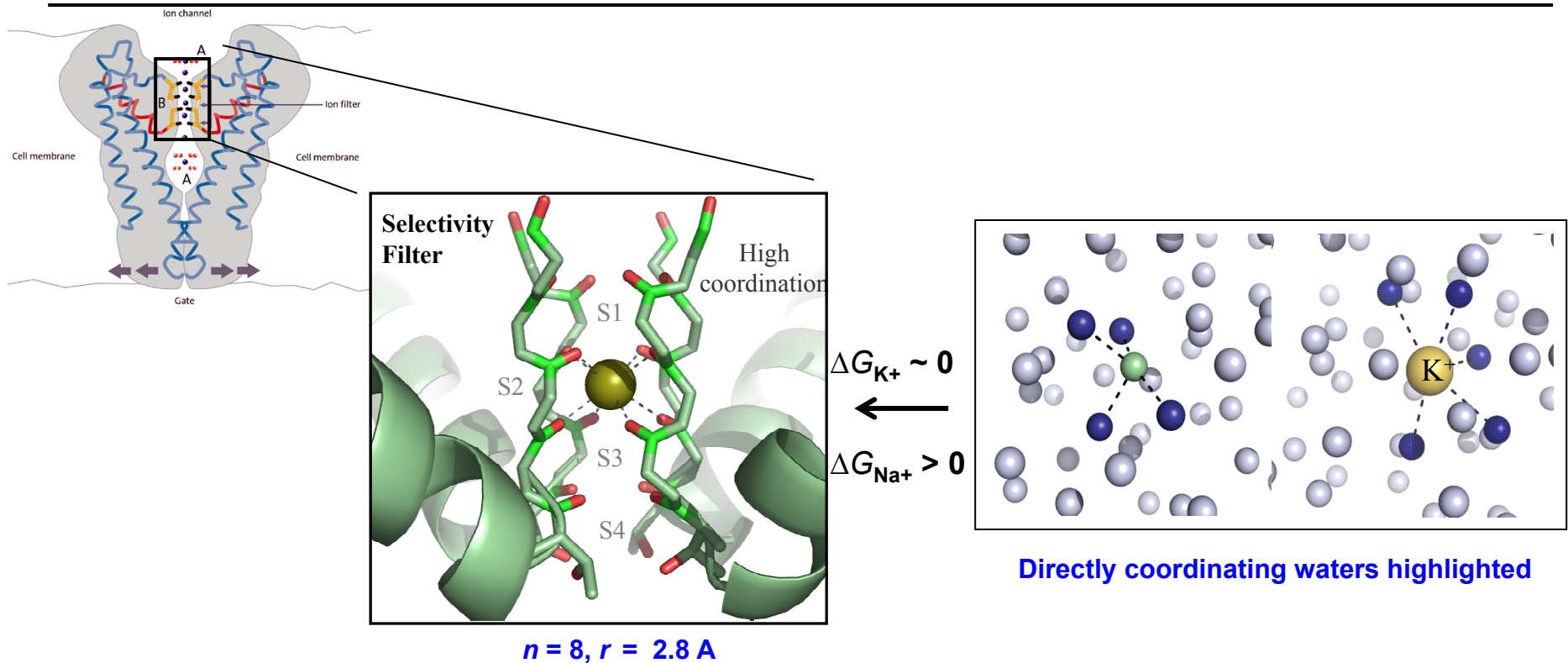
- Peaks & wells match scattering experiments:



- **Additional info obtained by AIMD:**
 - individual water distributions
 - $n(r)$

K⁺/Na⁺ Ion Discrimination Problem:

Prominent Views



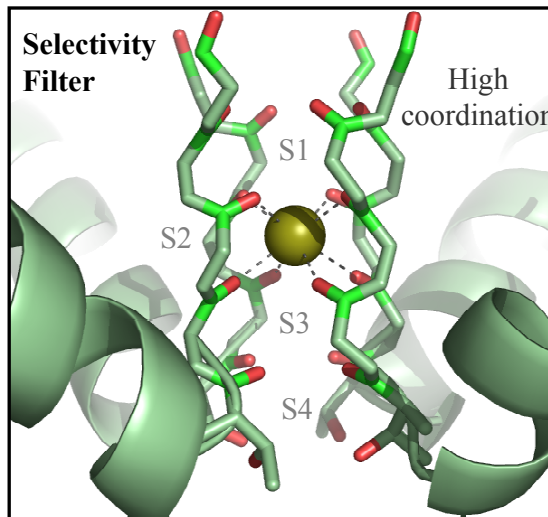
1. K⁺ equi-energy? Mimic K⁺ ion hydration structures
2. Na⁺ rejection? Specific cavity size fits K⁺^{1,2} vs Liquid flexibility & specific chemistry^{3,4}
3. Same filter structure vs varied selectivity? X

¹Bezanilla & Armstrong *JGP* (1972), ²Zhou *et al.* *Nature* (2001), ³Eisenman & Horn *JMB* (1983) ⁴Noskov *et al.* *Nature* (2004)
 Perspectives on Ion Selectivity *JGP* (2011)

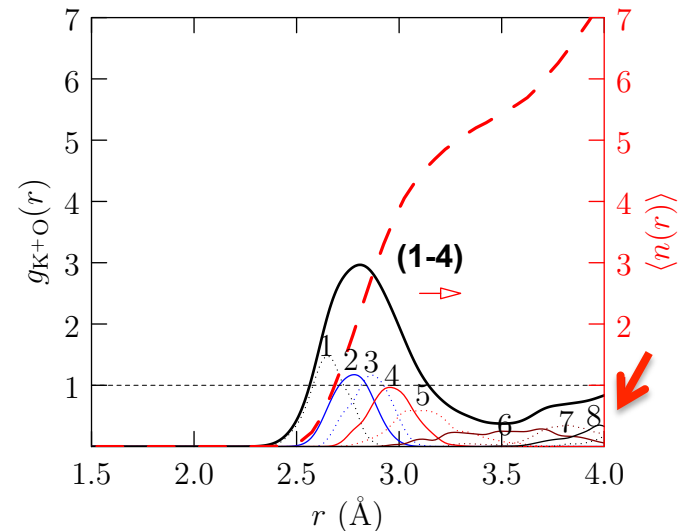
K⁺/Na⁺ Ion Discrimination Problem:

1. Ion hydration not mimicked!

C=O density around K⁺
n = 8 at r = 2.8 Å



Water density around K⁺
n = 4 at r_{max} = 2.8 Å



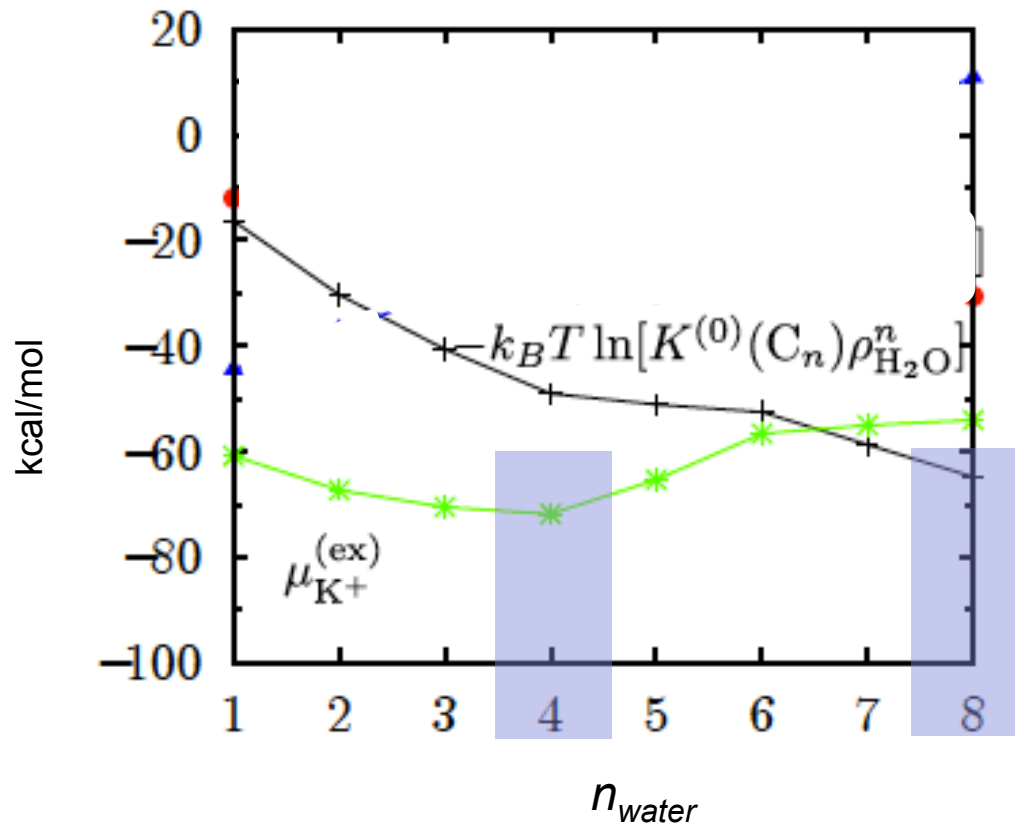
Ann Rep Comp Chem (2012)

1. K⁺ equa-energy? **Mimic K⁺ ion hydration structures**
2. Na⁺ rejection? Specific cavity size fits K⁺^{1,2} vs Liquid flexibility & specific chemistry^{3,4}
3. Same filter structure vs varied selectivity? X

¹Bezanilla & Armstrong *JGP* (1972), ²Zhou *et al.* *Nature* (2001), ³Eisenman & Horn *JMB* (1983) ⁴Noskov *et al* *Nature* (2004)
Perspectives on Ion Selectivity *JGP* (2011)

How can K^+ be stable with $n = 8$ in channels?

- K^+ hydration free energy vs coordinating waters

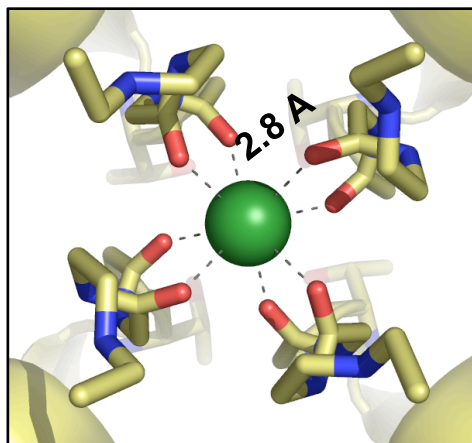


- Solvation (chem + density)
 - $n = 8$
- Solvation (aq)
 - $n = 4$
 - ligand desolvation penalty $\mu_{H_2O}^{(ex)}$

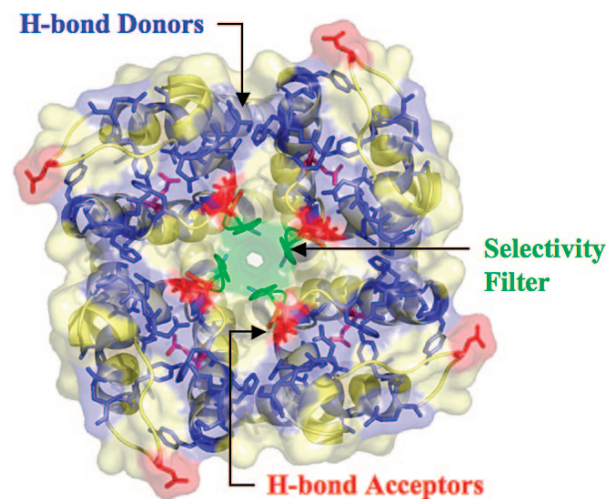
Does high ligand availability + poor ligand solvation environment account for high n in proteins?

Highly Selective K Channels:

High ligand availability + poor ligand solvation environ $\rightarrow n = 8$



K⁺ binding site



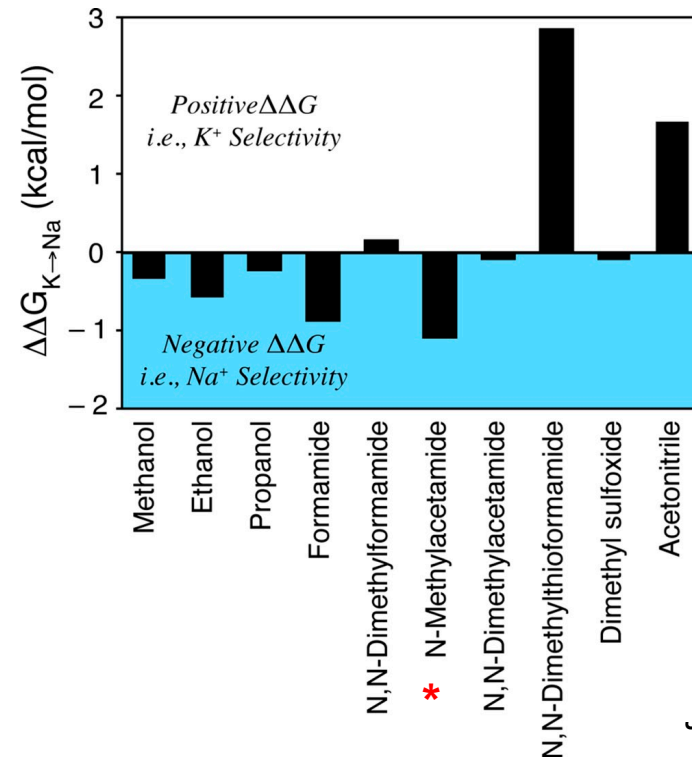
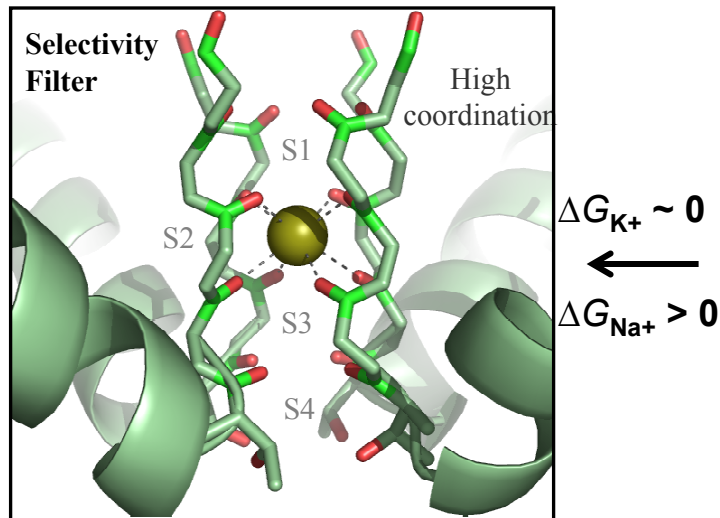
Highly selective K channels:
-- high CO density
-- few CO solvators

- To solvate K⁺ without barrier or well, binding site + surrounds must 'mimic' $\mu^{ex}(aq)$

\rightarrow not necessary to mimic K⁺(aq) local structure

K⁺/Na⁺ Ion Discrimination Problem:

2a. C=O does not preferentially solvate K⁺!

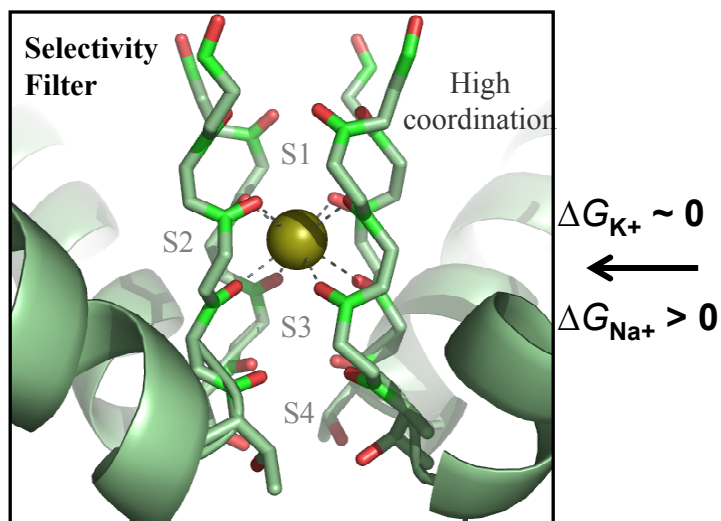


1. K⁺ equa-energy? Mimic K⁺ ion hydration structures
2. Na⁺ rejection? Specific cavity size fits K⁺^{1,2} vs Liquid flexibility & specific chemistry^{3,4}
3. Same filter structure vs varied selectivity? X

¹Bezanilla & Armstrong *JGP* (1972), ²Zhou *et al.* *Nature* (2001), ³Eisenman & Horn *JMB* (1983) ⁴Noskov *et al.* *Nature* (2004)

⁵Perspectives on Ion Selectivity *JGP* (2011)

K⁺/Na⁺ Ion Discrimination Problem: Prevailing Views



$n = 8, r = 2.7 - 3.1 \text{ \AA}$

Ion-water distances (A) (r_{max} in AIMD)

- K⁺ 2.8
- Rb⁺ 3.0
- Ba²⁺ 2.8
- Sr²⁺ 2.7

- Na⁺ 2.4

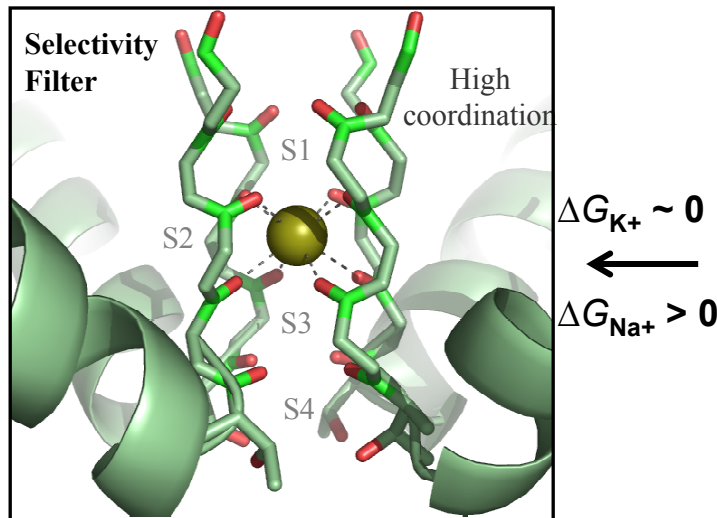
1. K⁺ equa-energy? Mimic K⁺ ion hydration structures
2. Na⁺ rejection? Specific cavity size fits K⁺^{1,2} vs Liquid flexibility & specific chemistry^{3,4}
3. Same filter structure vs varied selectivity? X

¹Bezanilla & Armstrong *JGP* (1972), ²Zhou *et al.* *Nature* (2001), ³Eisenman & Horn *JMB* (1983) ⁴Noskov *et al.* *Nature* (2004)

⁵Perspectives on Ion Selectivity *JGP* (2011)

K⁺/Na⁺ Ion Discrimination Problem:

More complexity than just cavity size



$n = 8, r = 2.7 - 3.1 \text{ \AA}$

B-factors: **rmsf = +/- 0.8 \AA**
Static vs dynamic disorder?

Ion-water distances (Å) (r_{\max} in AIMD)

• K ⁺	2.8	permeates
• Rb ⁺	3.0	permeates
• Ba ²⁺	2.8	blocks*
• Sr ²⁺	2.7	excluded
• Na ⁺	2.4	excluded

*PNAS, 2013

1. K⁺ equa-energy? Mimic K⁺ ion hydration structures
2. Na⁺ rejection? **Specific cavity size fits K⁺**^{1,2} vs Liquid flexibility & specific chemistry^{3,4}
3. Same filter structure vs varied selectivity? X

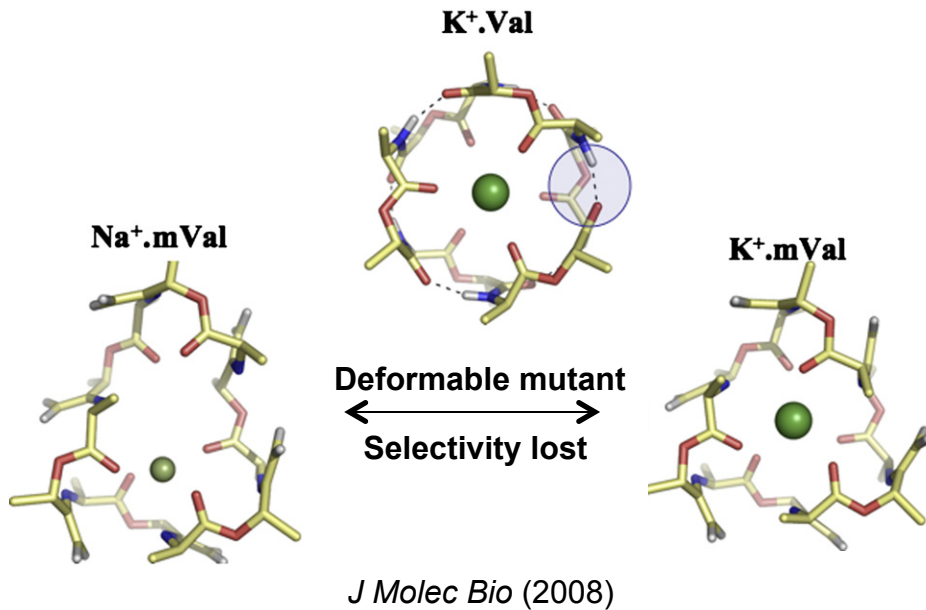
¹Bezanilla & Armstrong *JGP* (1972), ²Zhou *et al.* *Nature* (2001), ³Eisenman & Horn *JMB* (1983) ⁴Noskov *et al.* *Nature* (2004)

⁵Perspectives on Ion Selectivity *JGP* (2011)

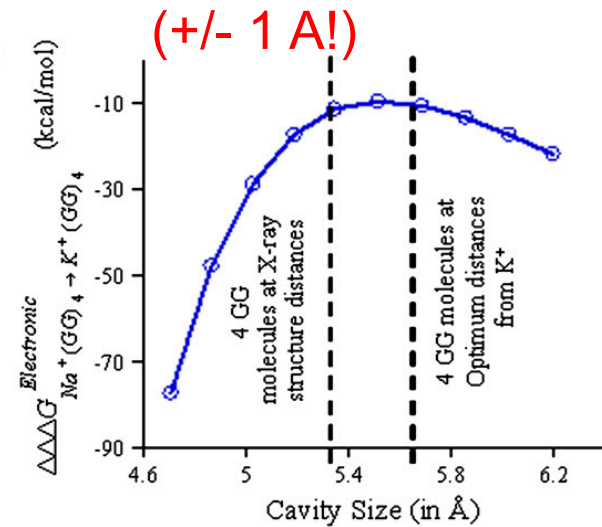
K⁺/Na⁺ Ion Discrimination Problem:

2b. Specific cavity size works - but not required!

- **Valinomycin ($n = 6$):**
Selective by specific cavity size



- **K channel ($n = 8$):**
Selective with varied cavity size



Biophys J (2007)

1. K⁺ equa-energy? Mimic K⁺ ion hydration structures
2. Na⁺ rejection? Specific cavity size fits K⁺^{1,2} vs Liquid flexibility & specific chemistry^{3,4}
3. Same filter structure vs varied selectivity? X

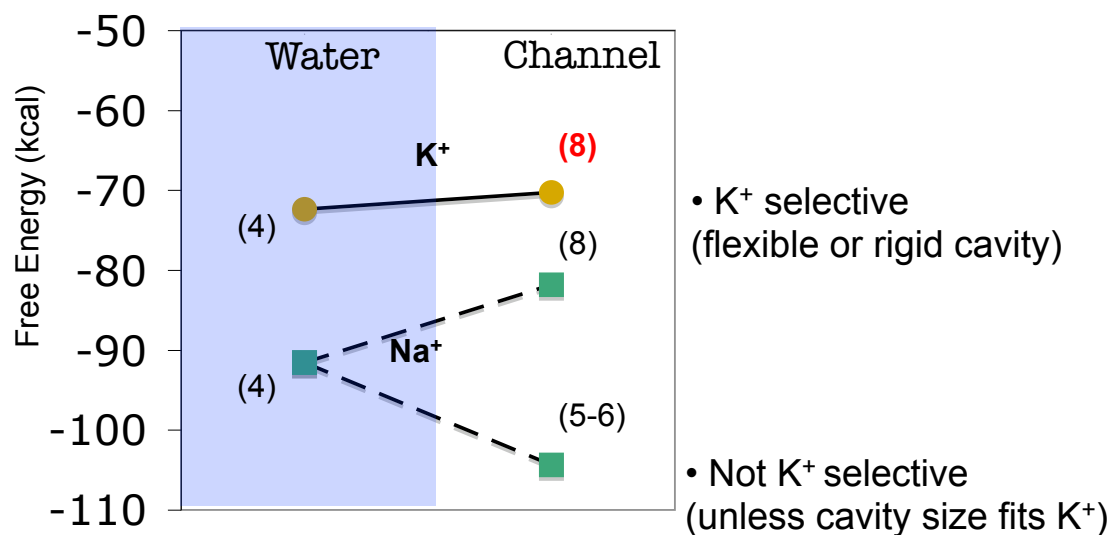
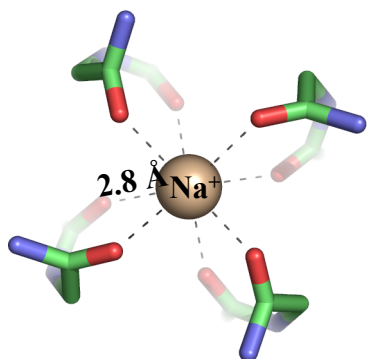
¹Bezanilla & Armstrong *JGP* (1972), ²Zhou *et al.* *Nature* (2001), ³Eisenman & Horn *JMB* (1983) ⁴Noskov *et al* *Nature* (2004)

⁵Perspectives on Ion Selectivity *JGP* (2011)

K⁺/Na⁺ Ion Discrimination Problem:

1-3. Environment → maintain crowded (> 6) ligands → K⁺/Na⁺ discrimination

K⁺ Channels



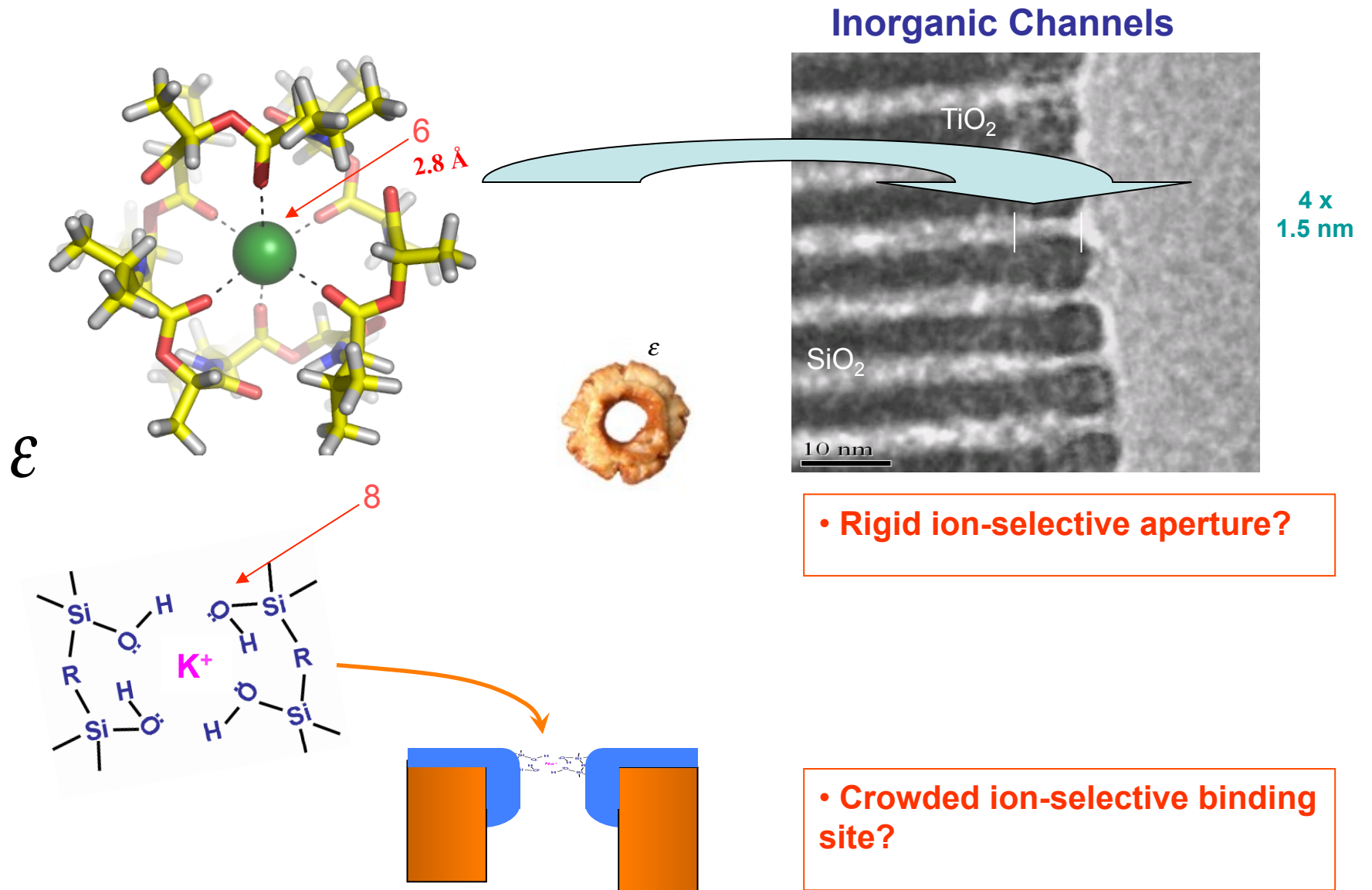
1. K⁺ equa-energy? Over-coordinate K⁺ with CO to match $\mu^{ex}(aq)$
2. Na⁺ rejection? Environment maintains specific, high n or specific cavity size
3. Same filter structure vs varied selectivity? Environment lets n (or cavity size) adjust

Biophys J (2007); Bostick & Brooks *PNAS* (2007)
J Mol Bio (2008); *JACS* (2008);
Ann Rep Comput Chem (2012)

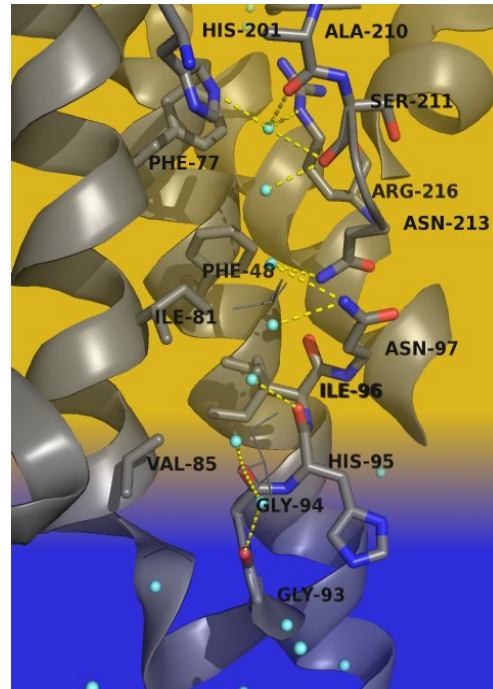
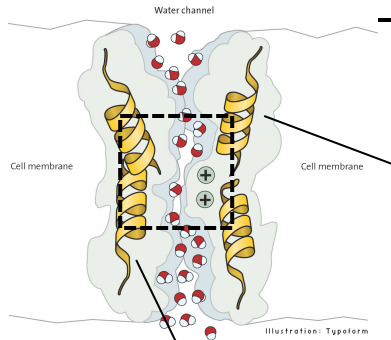
New hypothesis consistent with experiments:

Jiang & co Nature (2011)

Translate design to synthetic membranes?

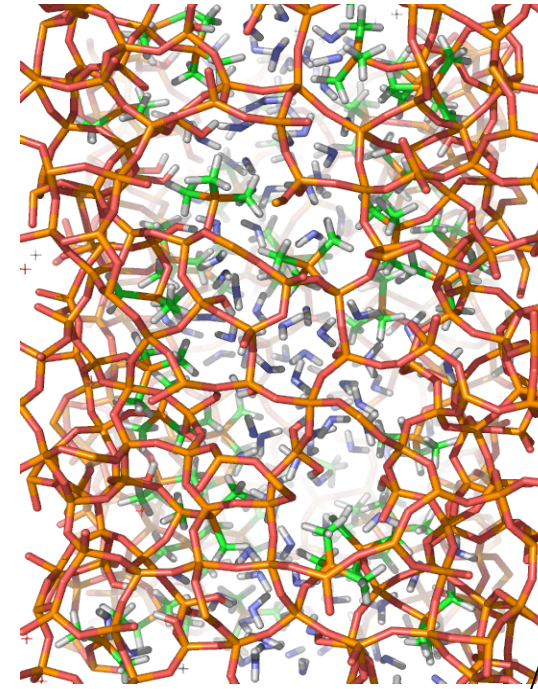


Water Discrimination: Fast water transport & ion rejection?



**Aq water channel
(crystal structure)**

(www.nobelprize.org)



**Model inorganic pore
(simulation results)**

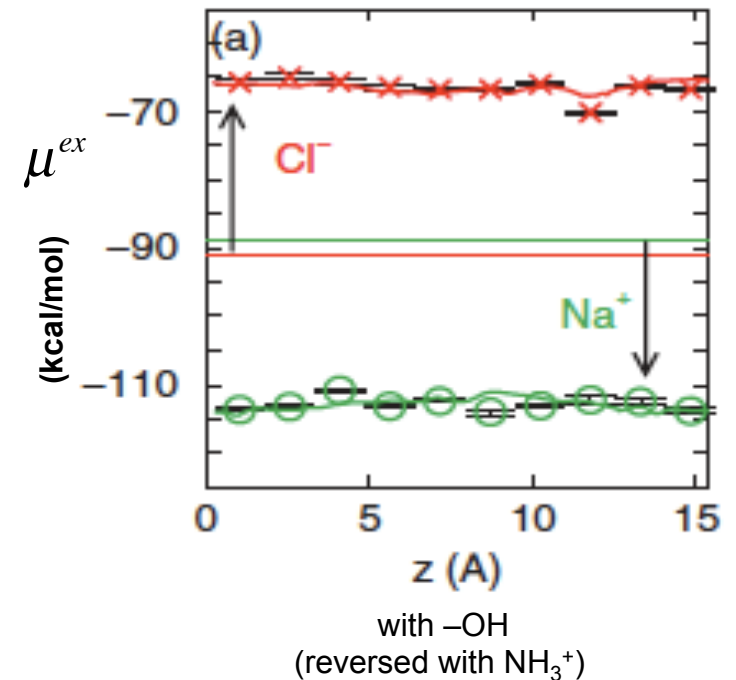
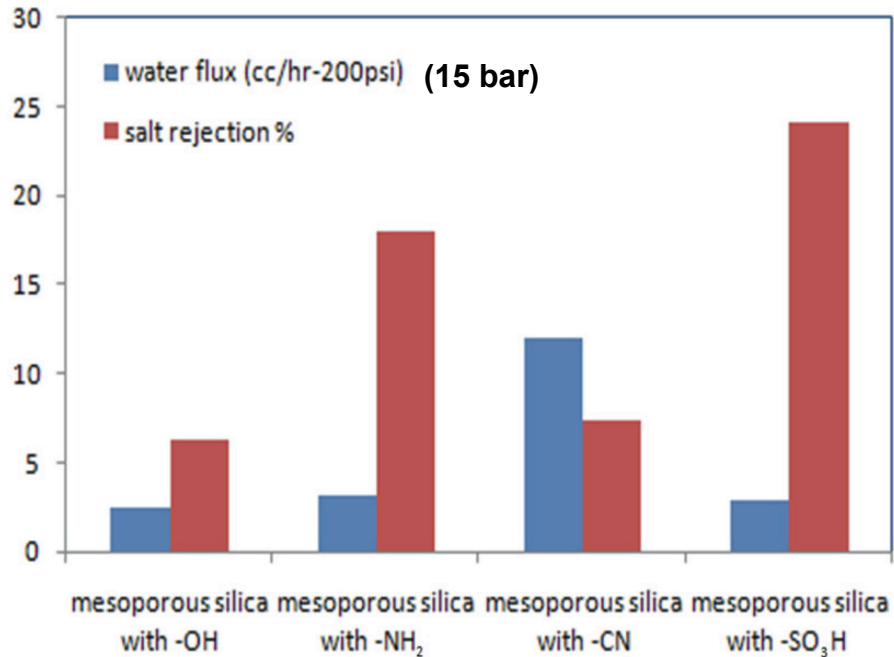
- thin, straight, narrow passage
- repulsive hydrophobic walls
- staircase of stabilizing dipoles
(C=O, N-H)
- no symmetric dipole clusters

Leung, Rempe, Lorenz *Phys Rev Lett* 2006
Cygan, .. Leung, Rempe *Mat Res Bull* (2008)
Leung & Rempe *J Comp Theor Nanosci* (2009)

Water channels with wide polar pores

Fast water transport, poor ion rejection

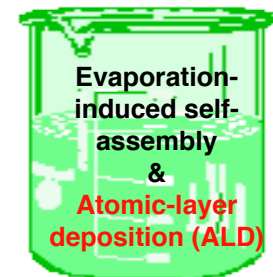
- Atomic layer deposition modifies pore size & chemistry



Over 1 nm polar pores:

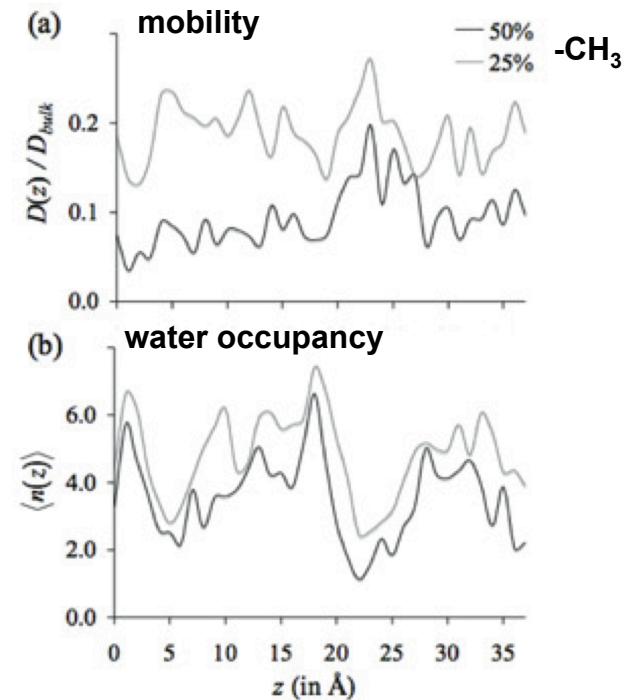
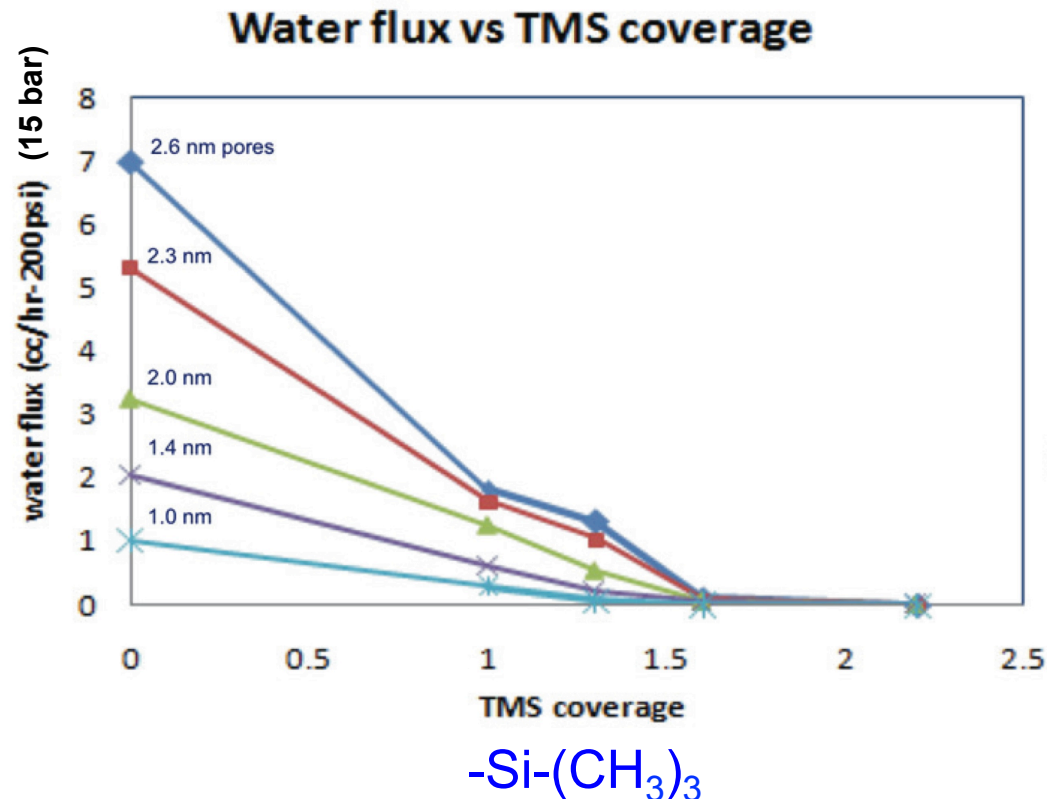
- waters stabilized → high flux, little salt rejection
- predict alternating chemistry → full salt rejection

Leung, Rempe, Lorenz PRL (2006)



Water Channels with non-polar pores

Slow water transport, great ion rejection

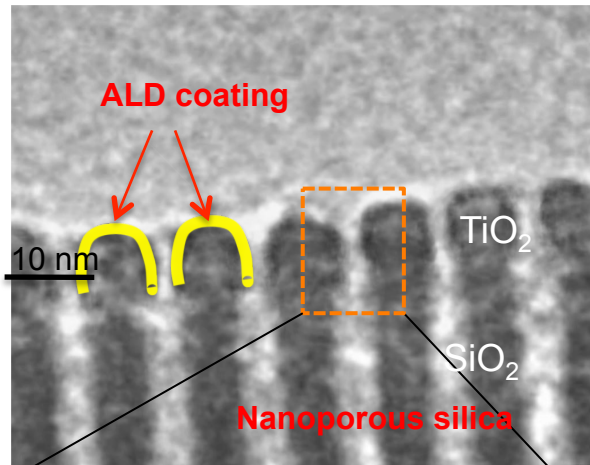


Over 1 nm hydrophobic pores:

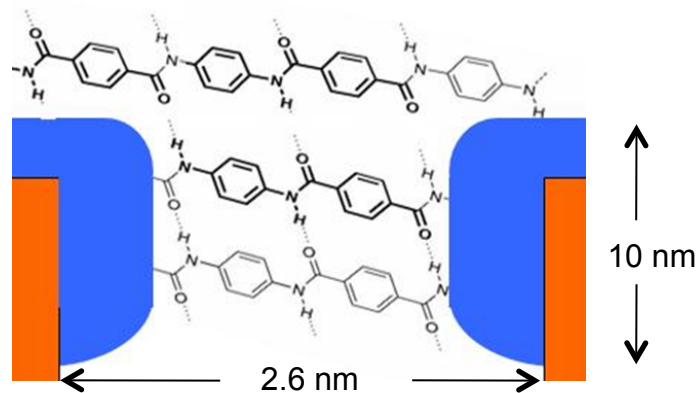
- ions rejected
- waters destabilized & immobilized

Molecular-layer deposition of polymer:

Translate key water channel designs?



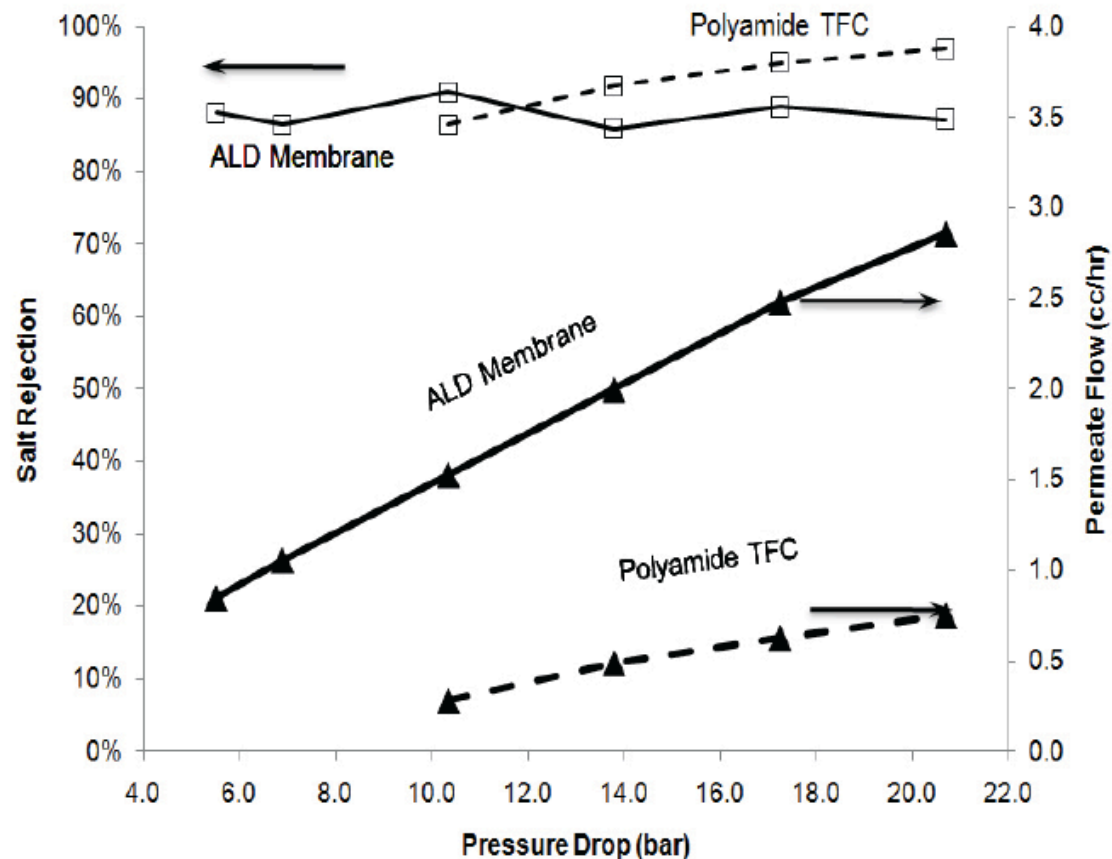
- C=O, N-H dipoles stabilize water
- hydrophobic rings prevent sticking
- no symmetric dipole clusters for ions
- narrow passageway excludes hydrated ions



Molecular layer deposition near surface via plasma → fundamentally different membrane architecture:

- TMOS (tri-methoxy silane)
- APS (amino-propyl silane)

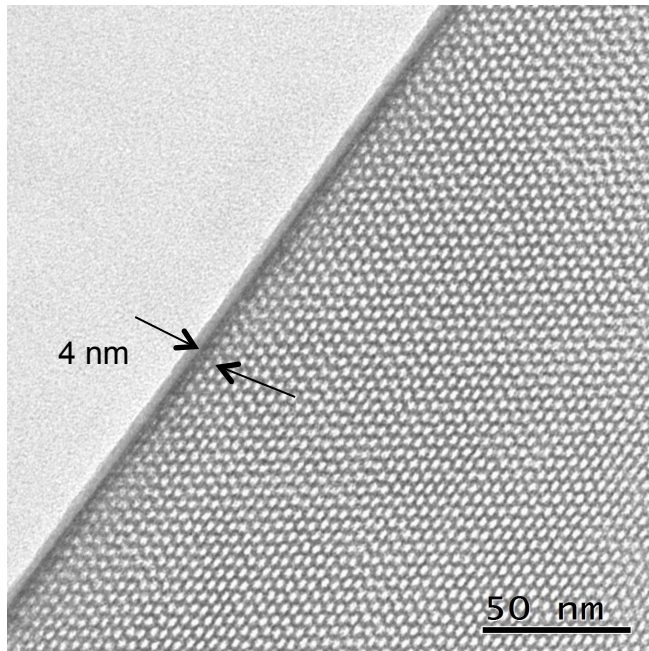
ALD Biomimetic Membranes = behave differently than (TFC) membranes



- ALD: Rejection high even at low P
- Different mechanism?

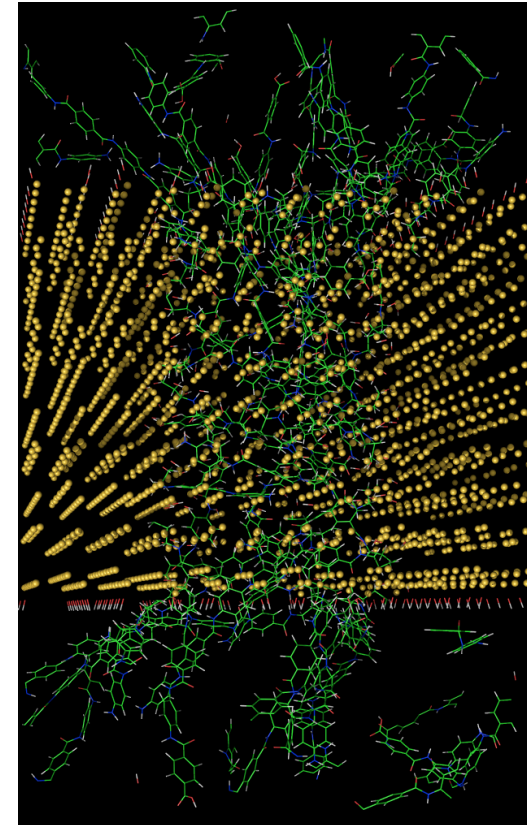
- 0.1 M NaCl, osmotic $\Pi = 2.1$ bar, $1/\kappa = 1.2$ -nm

What structural elements can be improved?



TEM image of thin membrane coating

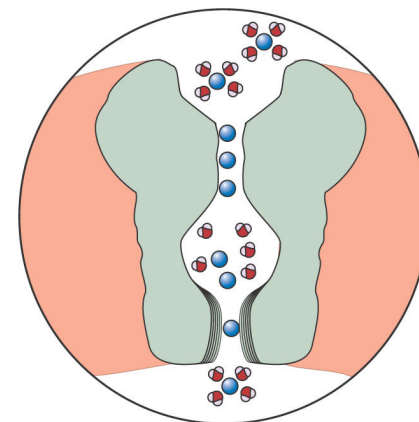
- thin
 - uniform
 - robust
 - Coordination chemistry?
- room for improvement



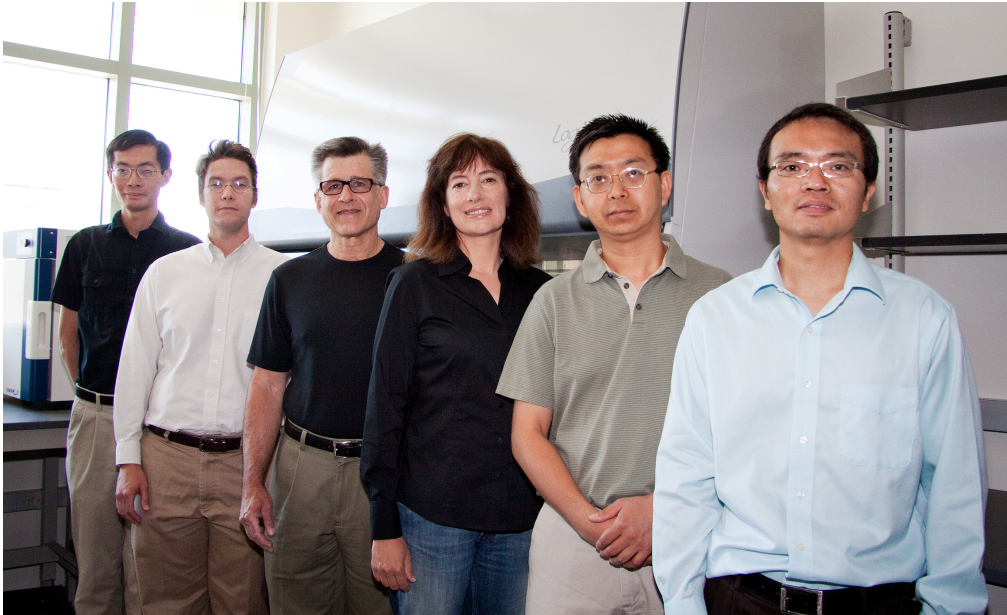
Simulated polymer ALD membrane

Summary: Membranes Designed to Mimic Protein Channels

- K^+/Na^+ discrimination & tunability (QM locally with new theory):
 - surroundings \rightarrow maintain crowded (>6) binding site fit to K^+
 - surroundings \rightarrow maintain rigid pore size fit to K^+
- Specific water selectivity & reject ions (fabricated ALD):
 - narrow
 - polar/non-polar mix
 - no ion binding sites
- General biomimetic features (fabricated EISA/PA-ALD):
 - thin & straight
 - uniformly active
 - robust structure



Team



Acknowledgements

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1. DOE: Sandia's Water Desalination Program
2. DOE: Sandia's LDRD program (BST, ERN)
3. NIH: Nanomedicine Center

- **Compute time**

1. Sandia Computing: Red Sky
2. National Centers: NCSA, NERSC

- **Theory Collaborators**

- David Rogers (postdoc → U S Florida)
- Dian Jiao (postdoc → UT)
- Sameer Varma (postdoc → U S Florida)
- Dubravko Sabo (postdoc → NYU)
- Chris Lorenz (postdoc → King's College)
- Mangesh Chaudhari (postdoc)

- Kevin Leung (SNL)
- Peter Feibelman (SNL)
- Lawrence Pratt (Tulane)

- **Experimental Collaborators**

- Jeff Brinker (SNL, UNM)
- Ying-Bing Jiang (UNM postdoc → UNM)
- Zhu Chen (UNM postdoc → Ford)
- Shaorong Yang (UNM postdoc → CNM)
- Seema Singh (UNM postdoc → SNL)
- Caroline Rempe (UNM student → U Tenn)
- Todd Alam (SNL)
- Tom Mayer (project manager SNL → UNM)

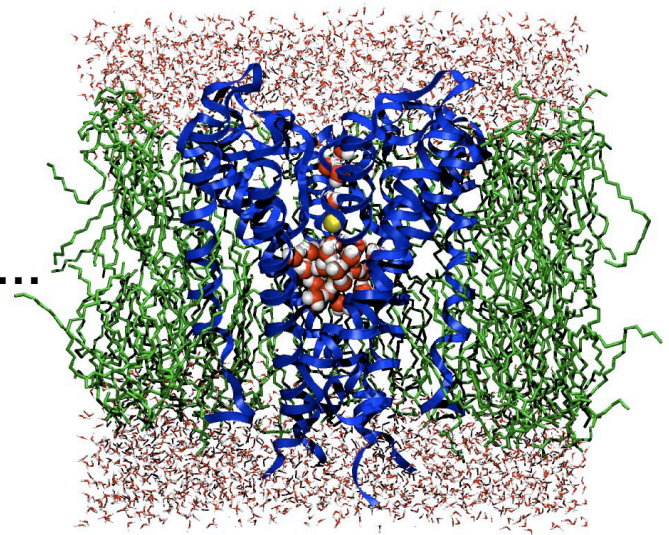


Prior Simulation Studies of Fast Separations

- *J of General Physiology: Perspectives on Ion Selectivity (2011)*

Noskov, Roux, Allen, Asthagiri, Pratt, Rempe, ...

Corry, Bostick, Brooks, Weaver, Green Sansom, Domene, Carloni, Aqvist, ...



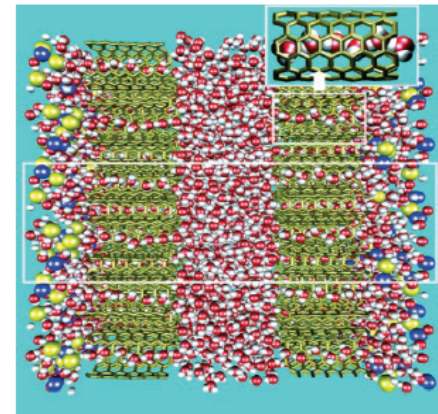
Rogers & Rempe, *J Stat Phys* (2011)

- **Aquaporins**

- De Groot, Grubmuller
- Tajkhorshid, Schulten
- Warshel

- **Other Pores**

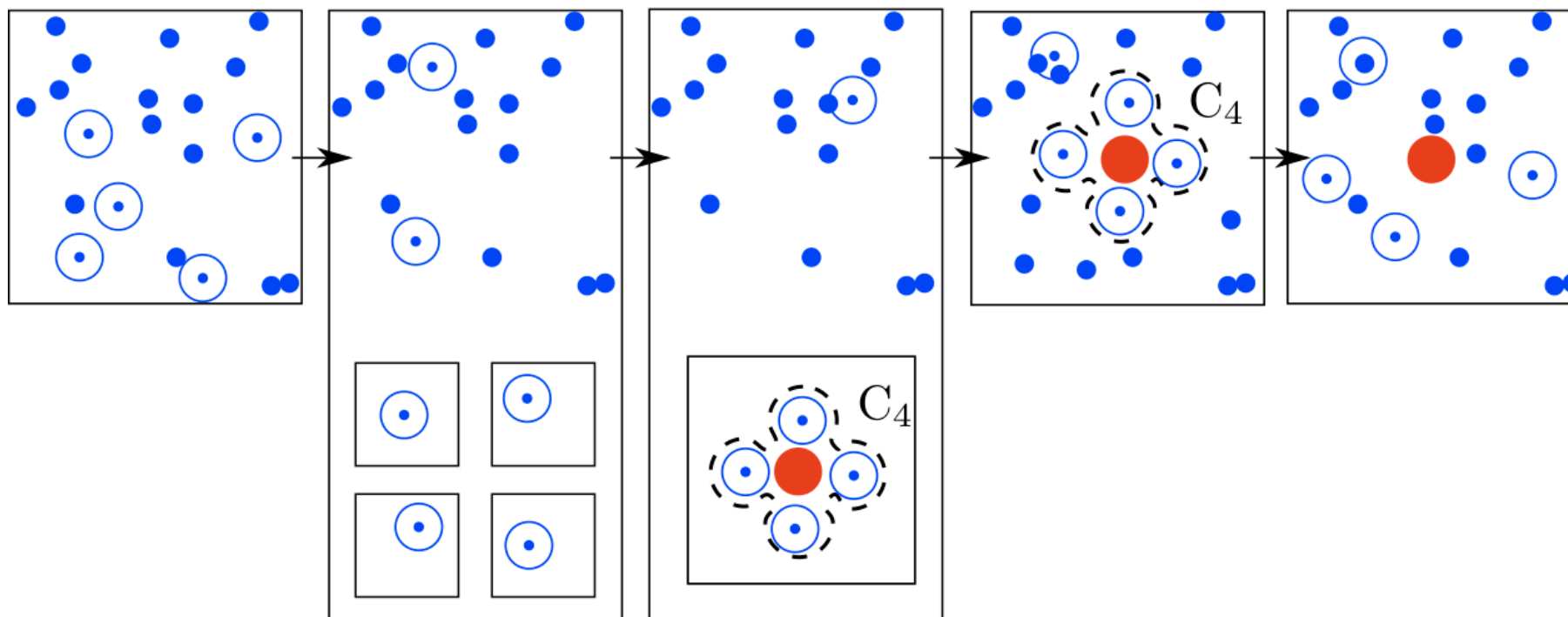
- Hummer, Patel, Leung, Aluru
- Corry, Roux
- Lynden-Bell



Hummer & co *PNAS* (2003)

Treat Ion Binding & Solvation via QM Efficiently

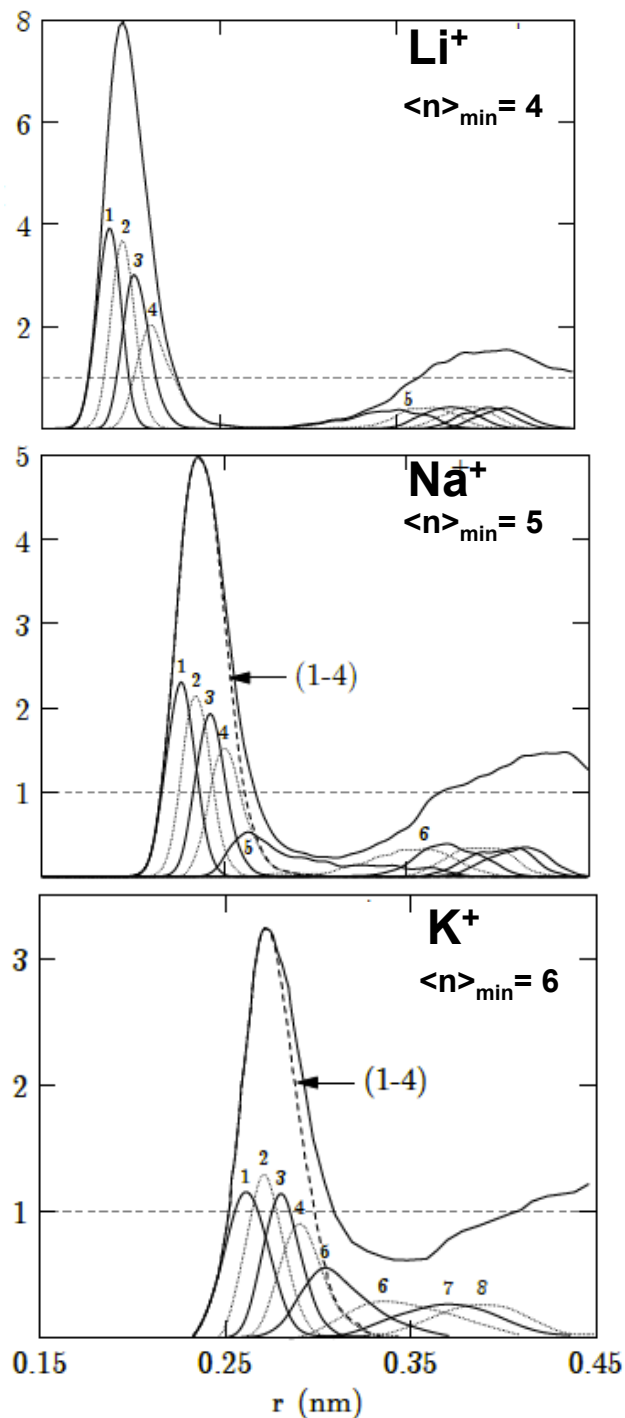
$$\beta\mu_X^{(\text{ex})} = -n\beta\mu_L^{(\text{ex})} - \ln[K^{(0)}(C_n)\rho_L^n] + \beta\mu_{X\cdot L_n}^{(\text{ex})} + \ln p_X(C_n)$$



Quasi-chemical theory (QCT) advantages:

- study binding sites with QM
- study environment ('architectural constraints')

Lessons: Ions (aq) (full QM)



- Ions differ:
 - Structure, DG
- Common structural theme:
 - $n = 4$ directly coordinating
 - Non 'split-shell' convenient
- Higher n in proteins?
 - $n = 4$ within 2.8 Å (water)
 - $n = 8$ within 2.8 Å (channel)

Agreement with exp
*Neilson
*Soper

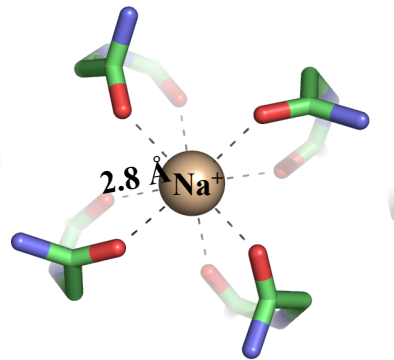
How do channels discriminate
between ions?

Ann Rep Comp Chem (2012)

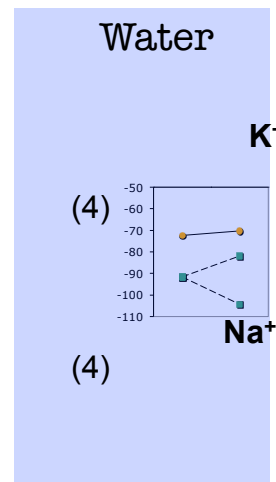
Varma & Rempe *Biophys Chem* (2006)

Environment maintains a specific binding site coordination that fits K^+

K^+ Channels



Free Energy (kcal)



Channel

(8)

(8)

- K^+ selective (flexible or rigid cavity)

(5-6)

- Not K^+ selective (unless cavity size fits K^+)

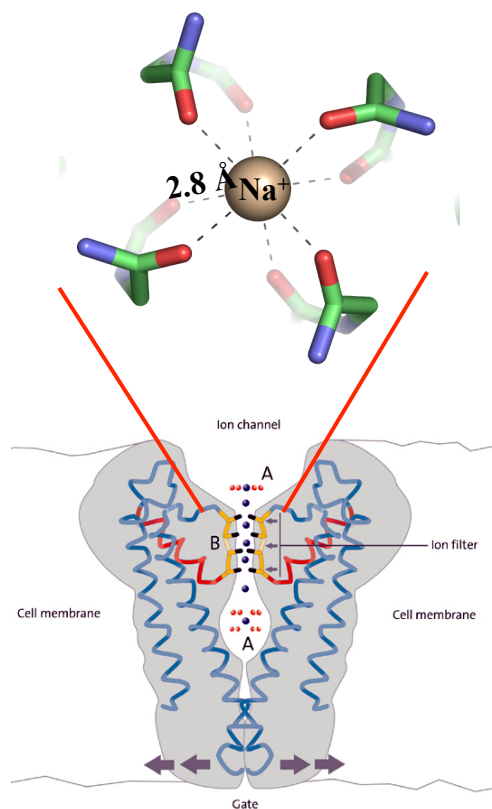
Environment \rightarrow maintain crowded (> 6) ligands \rightarrow K^+/Na^+ discrimination:
***Same chemistry \rightarrow environment tunes binding site composition \rightarrow varied function**

Biophys J (2007); ²Bostick & Brooks *PNAS* (2007)
J Mol Bio (2008); *JACS* (2008); *Ann Rep Comput Chem* (2012)

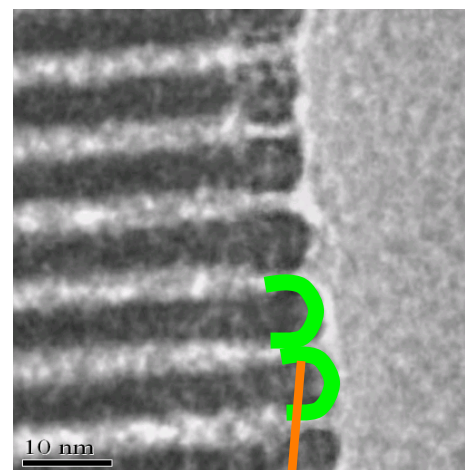
Jiang & co *Nature* (2011) Environment of multiple binding sites

Translate design for mineral water

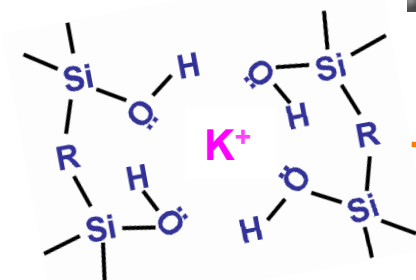
Natural Channels¹



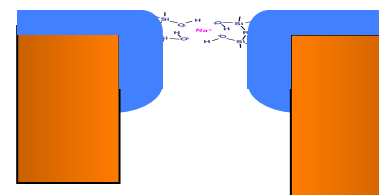
Inorganic Channels



Bio mechanisms
→
Engineered Mimicry



•³ Plasma-limited ALD +
molecular imprinting?



4 nm x 1.5 nm

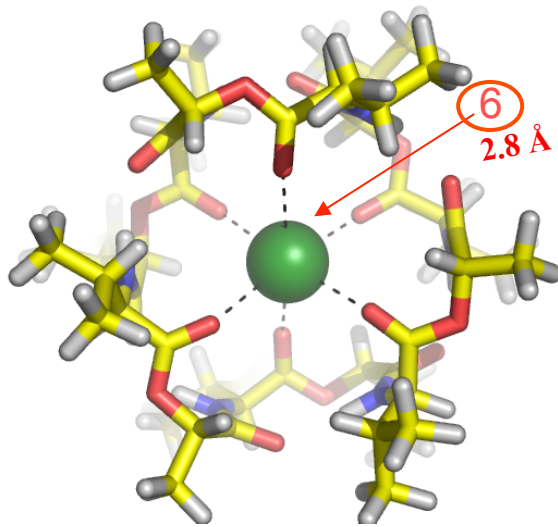
Nanoscale design parameters to target¹:

“The caress of the surroundings”² to maintain crowded ligands in fluctuating cavity

¹*Biophys J* (2007); ²*Jordan Biophys J* (2007); ³Jiang, Brinker, et al. *JACS* (2006,2007)

Translate design for mineral water

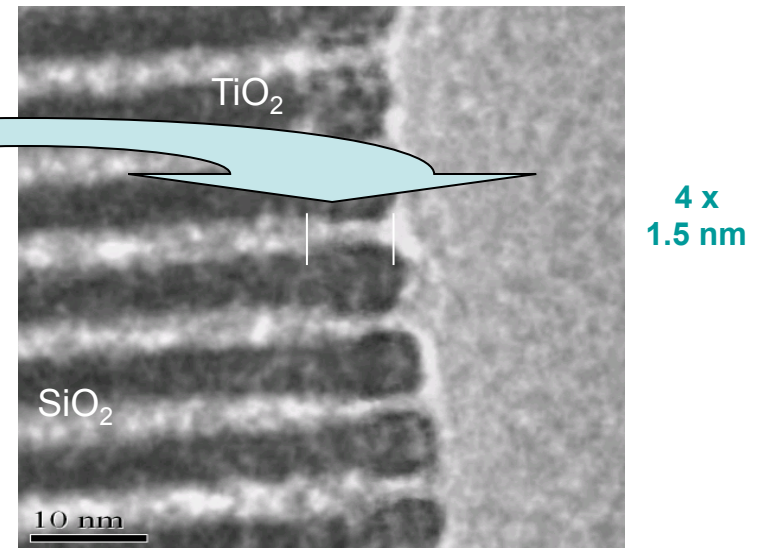
Natural K-selective Molecule



- **Specific cavity** fits K^+ , not Na^+
- **H-bonds** & ring maintain cavity size
- **Environment** may disrupt H-bonds, selectivity

Valinomycin, JMB (2008)

Inorganic Channels



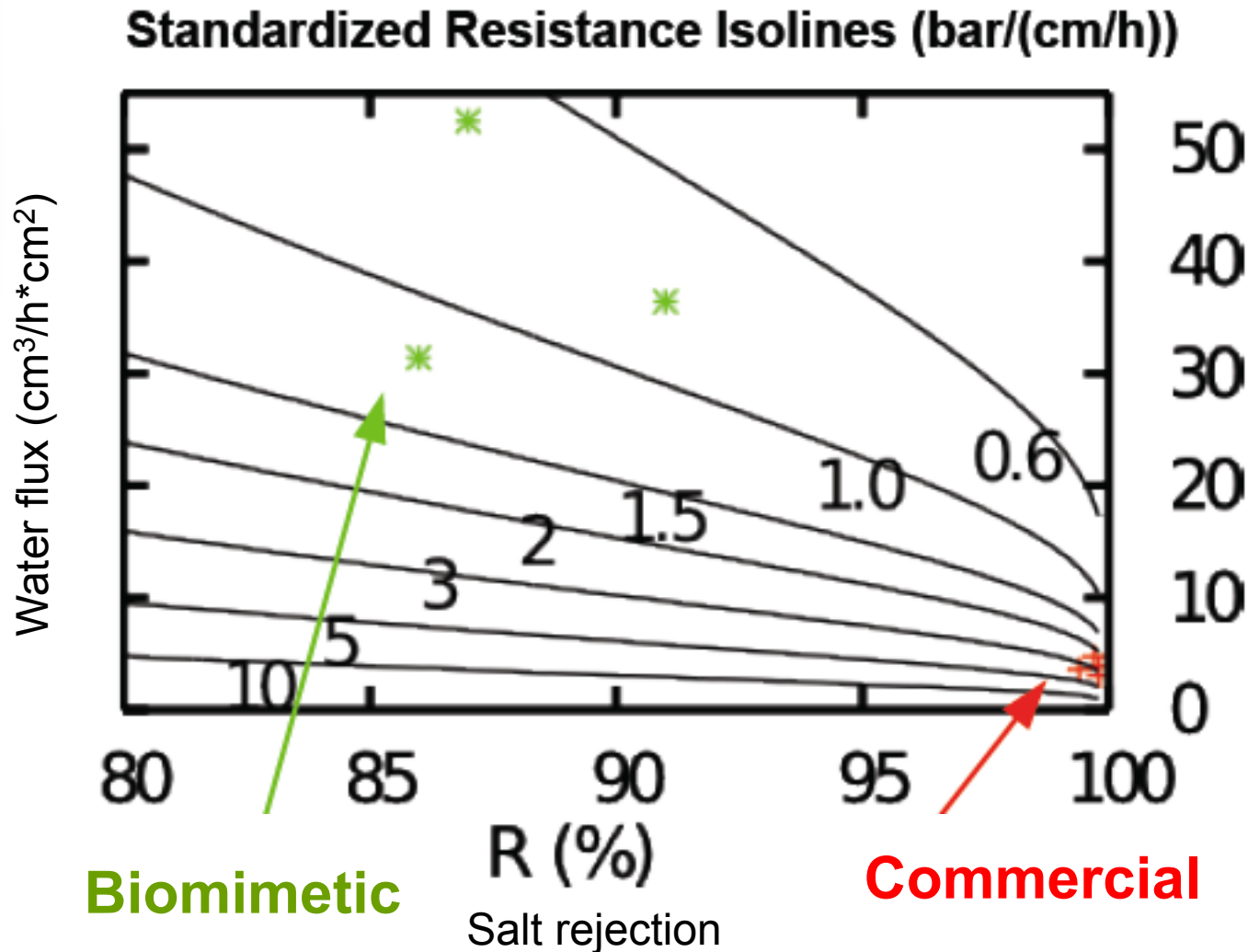
- **Rigid ion-selective aperture?**



• Two ways to discriminate K^+ over Na^+ :

- Environment → **overcoordinate** to fit K^+ → pore size can fluctuate
- Environment → **fixed cavity** fits K^+ → any number of ligands that solvate K^+

ALD Biomimetic membrane: 5-fold less resistance than Commercial



Gold standard:
FilmTec