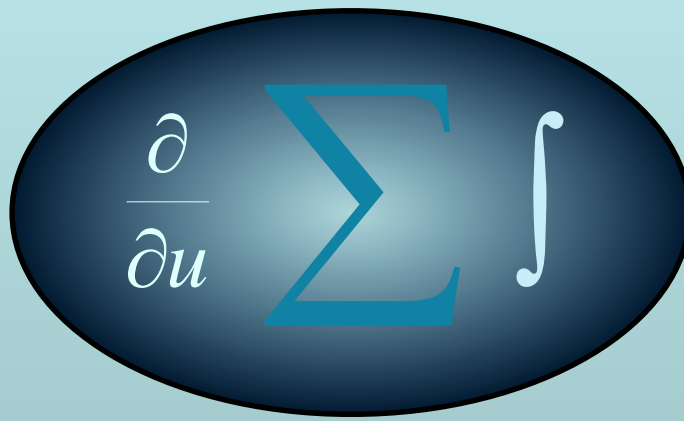


Thanks
to
PSU Math
particularly
Xiantao Li,
Chun Liu
Jinchao Xu, and
Yuxi Zheng

**It is a privilege (and joy) to visit many times and
work together.**



**Mathematics
describes only a tiny part of life,**

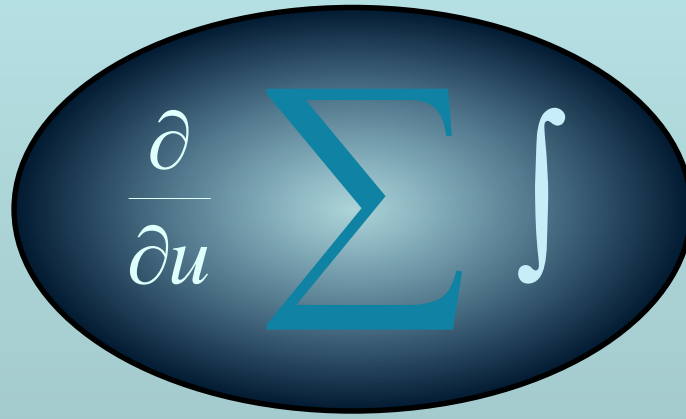
But

Mathematics* Creates

our

Standard of Living

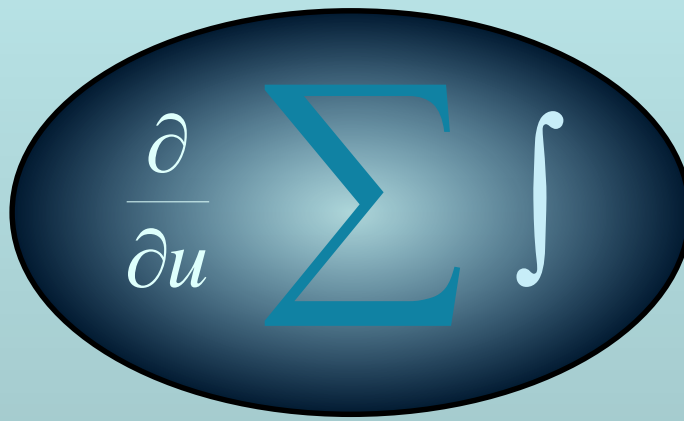
**e.g., Electricity, Computers, Fluid Dynamics, Optics, Structural Mechanics,*



**Mathematics* Creates
our
Standard of Living**

**Mathematics replaces
Trial and Error
with Computation**

**e.g., Electricity, Computers, Fluid Dynamics,
Optics, Structural Mechanics,*



***Chemistry and Biology
occur in Salt Solutions
with
Multiple Components, Interactions, and Dissipation***

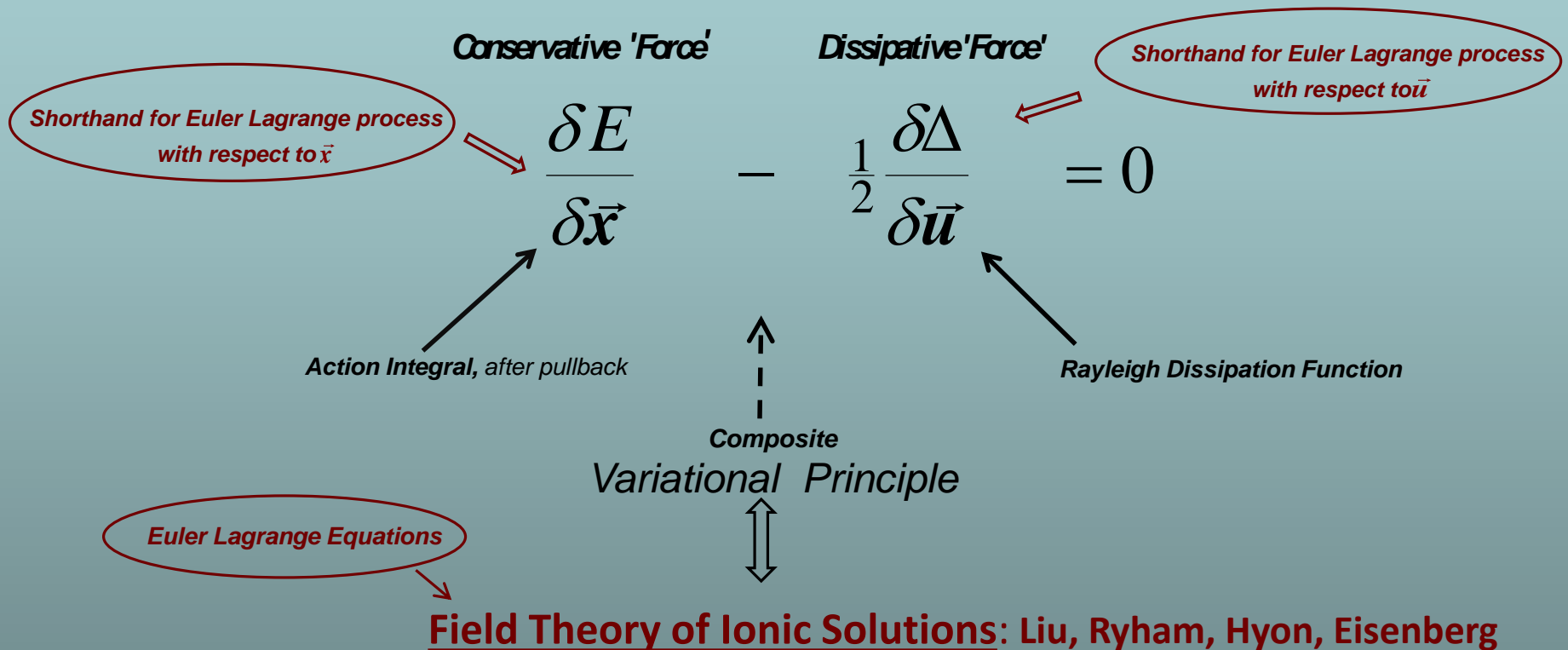
***Mathematics is now available
Energetic Variational Approach***

Energetic Variational Approach

EnVarA

Chun Liu, Rolf Ryham, and Yunkyong Hyon

Mathematicians and Modelers: two different 'partial' variations written in one framework, using a 'pullback' of the action integral



Allows boundary conditions and flow
Deals Consistently with Interactions of Components

All of Biology occurs in Salt Solutions

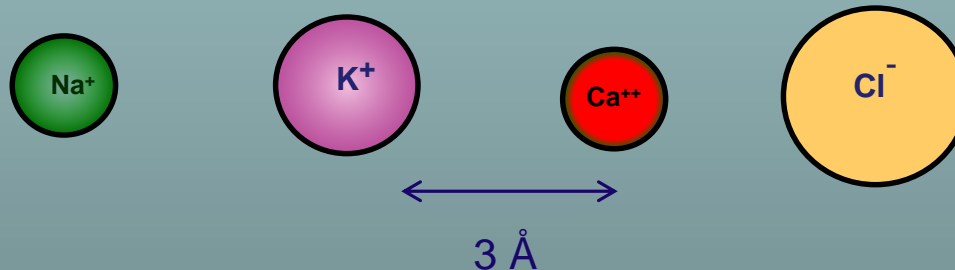
of definite composition and concentration
and that matters!

Salt Water is the Liquid of Life

Pure H_2O is toxic to cells and molecules!

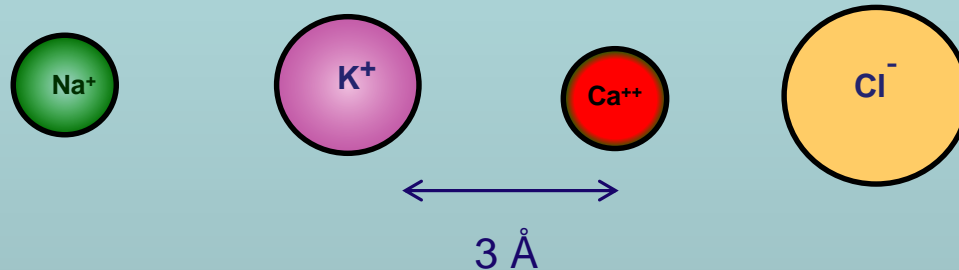
Main Ions are Hard Spheres, close enough

Sodium Na^+ Potassium K^+ Calcium Ca^{2+} Chloride Cl^-



Main Biological Ions are Hard Spheres, close enough

Sodium Na^+ Potassium K^+ Calcium Ca^{2+} Chloride Cl^-



General Theory of Hard Spheres is now available

Thanks to Chun Liu, more than anyone else

Took a long time, because dissipation, multiple fields, and multiple ion types had to be included

VARIATIONAL APPROACH IS NEEDED

Dissipation Principle

Conservative Energy dissipates into Friction

Conservative

$$\begin{aligned}
 & \frac{d}{dt} \int \left\{ k_B T \sum_{i=n,p} c_i \log c_i + \frac{1}{2} \left(\rho_0 + \sum_{i=n,p} z_i e c_i \right) \phi + \sum_{i,j=n,p} \frac{c_i}{2} \int \tilde{\Psi}_{i,j} c_j d\vec{y} \right\} d\vec{x} \\
 &= - \int \left\{ \underbrace{\sum_{i=n,p} \frac{D_i c_i}{k_B T} \left| k_B T \frac{\nabla c_i}{c_i} + z_i e \nabla \phi - \sum_{j=n,p} \nabla \int \tilde{\Psi}_{i,j} c_j d\vec{y} \right|^2}_{\text{Dissipative}} \right\} d\vec{x}
 \end{aligned}$$

Annotations in the diagram:

- $\frac{d}{dt}$: time
- $k_B T$: Thermal Energy
- c_i : Number Density
- ρ_0 : Permanent Charge of protein
- z_i : valence proton charge
- $\tilde{\Psi}_{i,j}$: Hard Sphere Terms

c_i number density; $k_B T$ thermal energy; D_i diffusion coefficient; n negative; p positive; z_i valence; ϵ dielectric constant

Note that $\epsilon \frac{|\nabla \phi|^2}{2} = \frac{1}{2} \left(\rho_0 + \sum_{i=n,p} z_i e c_i \right) \phi$ with suitable boundary conditions

Energetic Variational Approach

EnVarA

Conservative 'Force'

Dissipative 'Force'

$$\frac{\delta E}{\delta \vec{x}} - \frac{1}{2} \frac{\delta \Delta}{\delta \vec{u}} = 0$$

is defined by the Euler Lagrange Process,

as I understand the pure math from Craig Evans

which gives

Equations like PNP

BUT

I leave it to you (all)

to argue/discuss with Craig

about the purity of the process

when two variations are involved

PNP (Poisson Nernst Planck) for Spheres

Non-equilibrium variational field theory *EnVarA*

Nernst Planck Diffusion Equation

for **number density** c_n of negative n ions; positive ions are analogous

Diffusion Coefficient

$$\frac{\partial c_n}{\partial t} = \nabla \cdot \left[D_n \left\{ \nabla c_n + \frac{c_n}{k_B T} \left(z_n e \nabla \phi - \int \frac{12 \varepsilon_{n,n} (a_n + a_n)^{12} (\vec{x} - \vec{y})}{|\vec{x} - \vec{y}|^{14}} c_n(\vec{y}) d\vec{y} - \int \frac{6 \varepsilon_{n,p} (a_n + a_p)^{12} (\vec{x} - \vec{y})}{|\vec{x} - \vec{y}|^{14}} c_p(\vec{y}) d\vec{y} \right) \right\} \right],$$

Thermal Energy

Coupling Parameters

Ion Radii

Number Densities

Poisson Equation

Dielectric Coefficient

$$\nabla \cdot (\varepsilon \nabla \phi) = - \left(\rho_0 + \sum_{i=1}^N z_i e c_i \right) \quad i = n \text{ or } p$$

Permanent Charge of Protein

valence proton charge

Semiconductor *PNP* Equations

For Point Charges

Poisson's Equation

$$-\frac{\epsilon_0}{A(x)} \frac{d}{dx} \left(\epsilon(x) A(x) \frac{d\phi}{dx} \right) = eP(x) + e \sum_i z_i \rho_i(x)$$

Dielectric Coefficient ϵ_0
 Cross sectional Area $A(x)$
 Permanent Charge of Protein $P(x)$
 Valence Proton charge z_i
 Number Densities $\rho_i(x)$

Drift-diffusion & Continuity Equation

$$\frac{dJ_i}{dx} = 0 \quad -J_i = D_i(x) A(x) \rho_i(x) \frac{d\mu_i}{dx}$$

Flux J_i
 Diffusion Coefficient $D_i(x)$
 Number Densities $\rho_i(x)$
 Chemical Potential $\mu_i(x)$

Chemical Potential $\mu_i(x)$

$$\mu_i(x) = z_i e \phi(x) + kT \ln \left(\frac{\rho_i(x)}{\rho^*} \right) + \underbrace{\mu_i^{\text{ex}}(x)}_{\text{Finite Size Special Chemistry}}$$

valence proton charge z_i
 Thermal Energy kT
 Finite Size Special Chemistry $\mu_i^{\text{ex}}(x)$

All we have to do is

Solve it/them!

with boundary conditions

All we have to do is

Solve it/them!

Boundary conditions:

STRUCTURES of Ion Channels

**STRUCTURES of semiconductor
devices and integrated circuits**

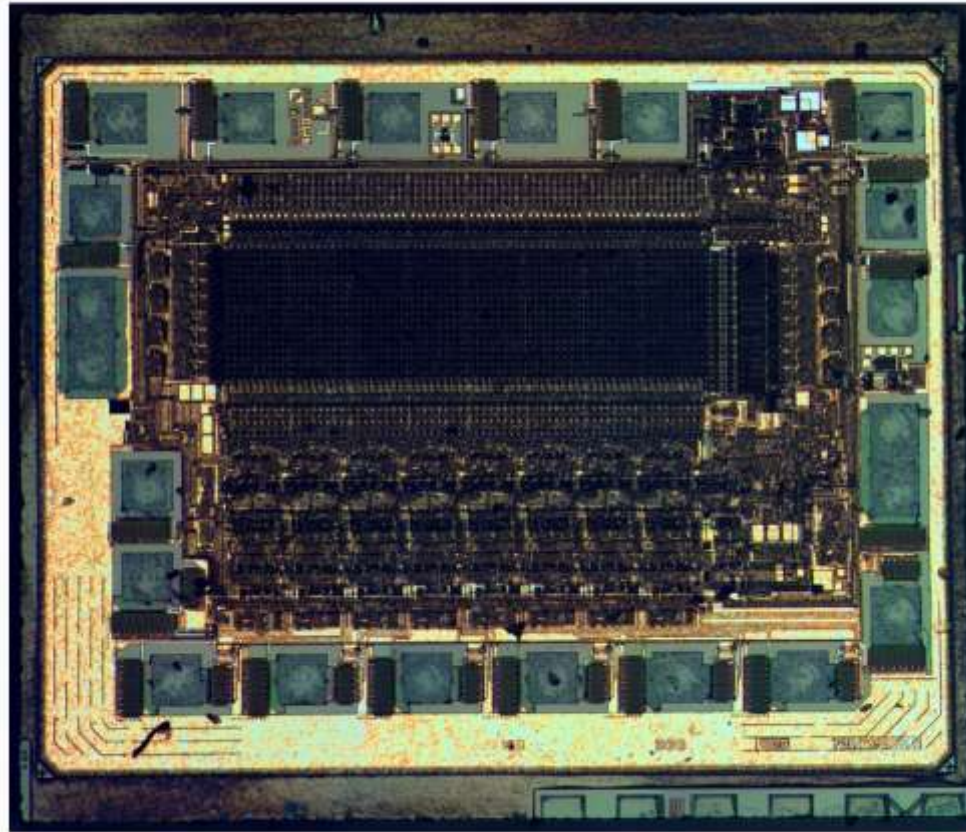
Integrated Circuit

Technology as of ~2005

AMD Palce16V8h : Weekend die-shot

GAL (Generic array logic) microchips are FPGA and CPLD grandfathers.
AMD Palce16V8h is an 32x64 array of AND elements.

Die size - 2434x2079µm, 1µm technology.



**Too
small
to see!**

Semiconductor Devices

PNP equations describe many robust input output relations

Amplifier

Limiter

Switch

Multiplier

Logarithmic convertor

Exponential convertor

These are SOLUTIONS of PNP for different boundary conditions
with ONE SET of CONSTITUTIVE PARAMETERS

PNP of POINTS IS TRANSFERRABLE

Analytical - Numerical Analysis

should be attempted using techniques of

Weishi Liu University of Kansas

Tai-Chia Lin National Taiwan University & Chun Liu PSU

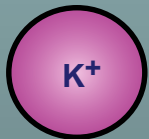
Ion Channels are Devices

Ion Channels are the Main Controllers of Biological Function

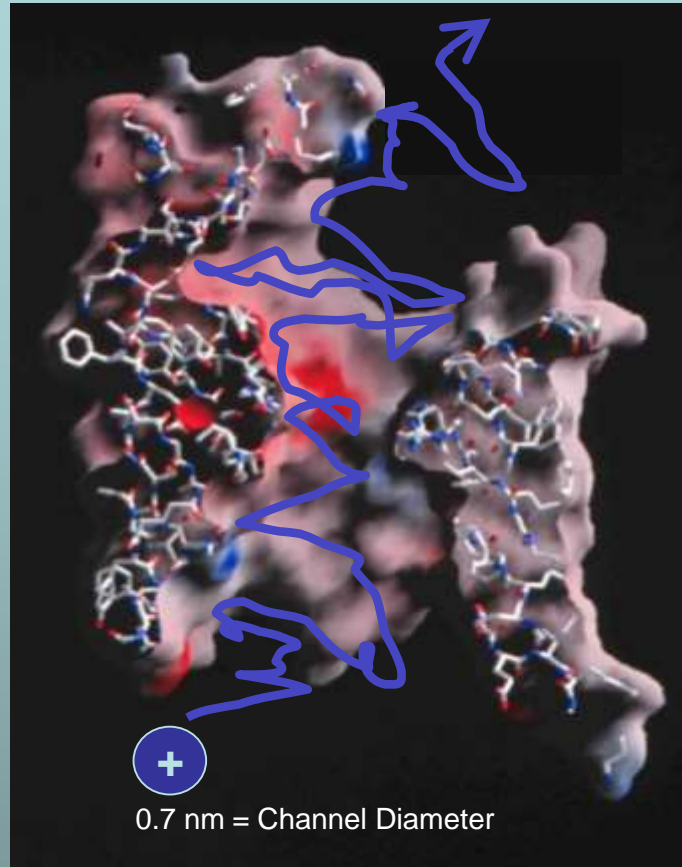
Biological Diodes

nonlinear current/voltage
curves

Different Ions
Different Diameters
carry
Different Signals



3 Å



~30 Å

Figure of ompF porin by Raimund Dutzler

Ions in Water*

are the
Liquid of Life

*Pure H₂O is toxic to cells & proteins

Chemical Bonds are lines
Surface is Electrical Potential
Red is negative (acid)
Blue is positive (basic)

General Theme

**Mathematics of Molecular Biology
Provides Great Opportunity**

Biology Provides the Data

Engineering Provides the Approach

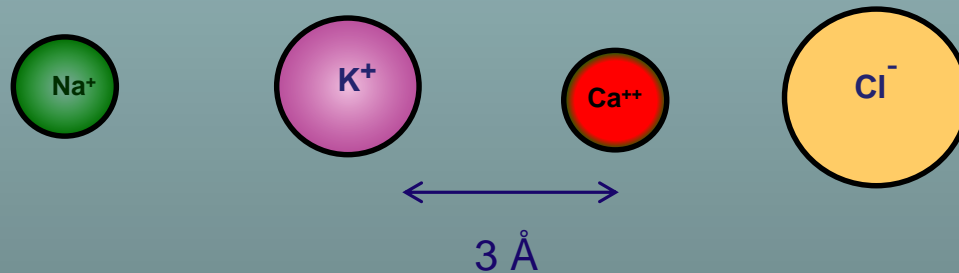
Mathematics Provides the Tools

particularly variational methods that allow
'everything' to interact with 'everything' else

Chemistry and Biology
Need a
TRANSFERRABLE Theory of
Salt Water
the liquid of life

Main Ions are Hard Spheres, close enough

Sodium Na^+ Potassium K^+ Calcium Ca^{2+} Chloride Cl^-



Classical text
Robinson and Stokes
still in print after sixty years

Not otherwise noted for its emotional content
gives a glimpse of these feelings when it says

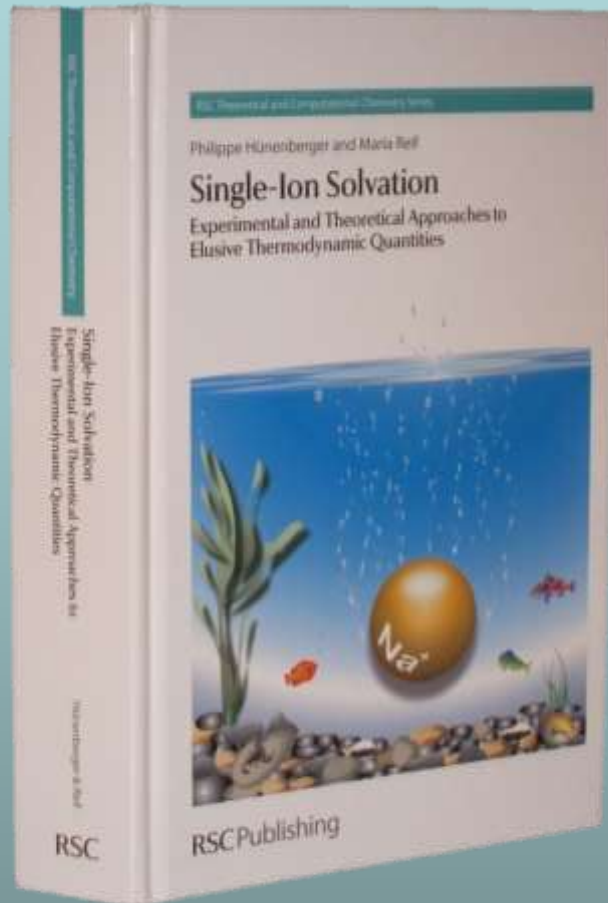
**“In regard to concentrated solutions,
many workers adopt a
counsel of despair,**

confining their interest to concentrations below about 0.02 M, ... ”

Biology occurs in concentrations > 0.2 M

p. 302 *Electrolyte Solutions* (1959) Butterworths , also Dover (2002)

Ionic Solutions are Complex Fluids



After 664 pages and 2604 references, properties of

SINGLE Ions
are
Elusive

because

Every Ion
Interacts
with
Everything

Hünenberger & Reif (2011) Single-Ion Solvation
Experimental and Theoretical Approaches to
Elusive Thermodynamic Quantities
in infinitely dilute solutions!

It is difficult to even define

in a unique way

Properties of One Ion

in infinitely dilute solution

when

Everything

Interacts

with

Everything

**Tremendous Opportunity
for Mathematics:**

Numerics and Variational Approach

Note
Emotion
IN
TITLE



664 pages, 2604 references

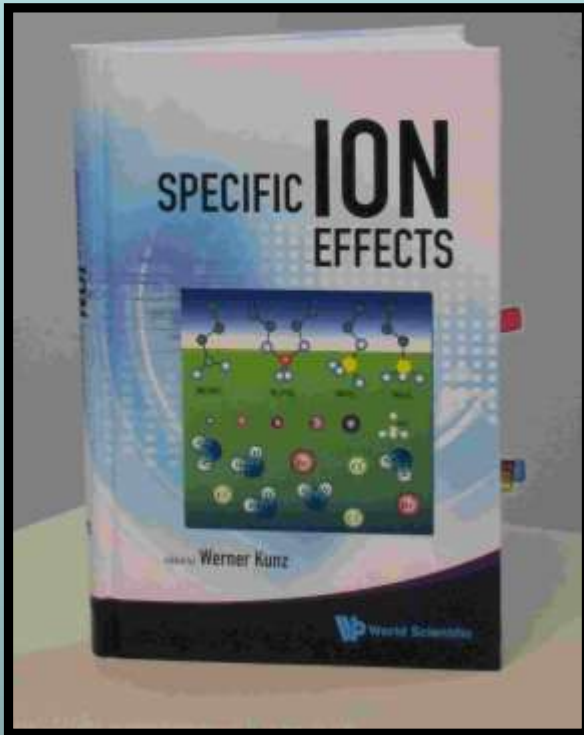
Deals with infinitely dilute solutions

Hünenberger, P. and M. Reif, (2011) *Single-Ion Solvation. Experimental and Theoretical Approaches to*

Elusive Thermodynamic Quantities*

Royal Society of Chemistry.

***Emphasis Bob E.**



Kunz, W. "Specific Ion Effects"
World Scientific Singapore, 2009; p 11.



Werner Kunz

“It is still a fact that over the last decades,
**it was easier to fly to the
moon
than to describe the
free energy
of even the simplest salt
solutions**

beyond a concentration of 0.1M or so.”

It is not surprising that
Inconsistent Treatments
of ionic solutions
have been

Unsuccessful

despite more than a century of work by fine scientists
and mathematicians

Good Data



Good Data

Compilations of Specific Ion Effect

1. **>139,175 Data Points** *on-line*
IVC-SEP Tech Univ of Denmark
http://www.cere.dtu.dk/Expertise/Data_Bank.aspx
2. Kontogeorgis, G. and G. Folas, 2009:
Models for Electrolyte Systems. Thermodynamic
John Wiley & Sons, Ltd. 461-523.
3. Zemaitis, J.F., Jr., D.M. Clark, M. Rafal, and N.C. Scrivner, 1986,
Handbook of Aqueous Electrolyte Thermodynamics.
American Institute of Chemical Engineers
4. Pytkowicz, R.M., 1979,
Activity Coefficients in Electrolyte Solutions. Vol. 1.
Boca Raton FL USA: CRC. 288.

**“Sometimes it is necessary to put a veil on
the past, for the sake of the future”**

Henry Clay

p. 375 of Henry Clay, the Essential American

David Heidler, Jeanne Heidler Random House

**Mathematics can remove
the veil!**



Mathematics
is needed to replace
Trial and Error
Experiments and Simulations

with
Computations
and
Consistent Theories

Mathematics of Chemistry

must deal

Naturally

with

Interactions

‘Law of Mass Action’ assumes nothing interacts

So this is a great opportunity for new mathematics and applications!



***Chemists have reliable quantitative data
but almost all their theory ignores interactions***

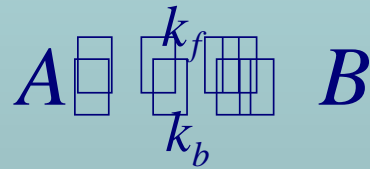
**Enormous opportunities
for
mathematics!**

***But you have to know which chemistry
That is where I can help, I hope!***

Chemistry
is about
Chemicals
not signals

Law of Mass Action

is what how chemists describe chemicals



$$J_{AB} = -\frac{d}{dt}[A] = k_f [A]; \quad J_{BA} = -\frac{d}{dt}[B] = k_b [B]$$

k is constant

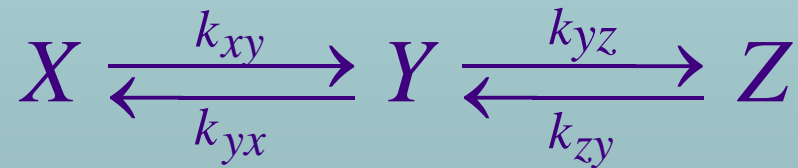
[A] means the activity or approximately the concentration of species A,
i.e., the number density of A

Law of Mass Action

is about

Conservation of Mass and Matter

It is not about conservation of charge



$$J_{xy}^{net} = J_{xy} - J_{yx}$$

$$= k_{xy} [X] - k_{yx} [Y]$$

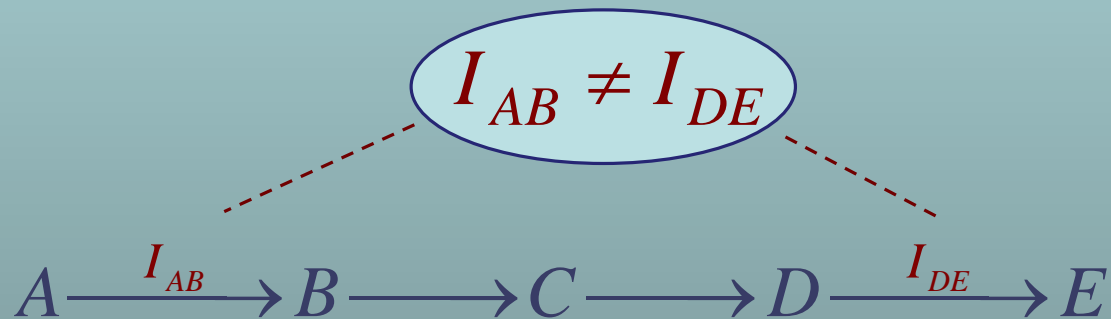
$$I_{xy}^{net} = z_x F k_{xy} [X] - z_y F k_{yx} [Y]$$

[X] means the concentration, really activity of species X, i.e., concentration is the number density

F is Faraday constant 96,500 coulombs/mole = 6.023×10^{23} particles

Units of current I_{xy} are (cou/sec)/liter

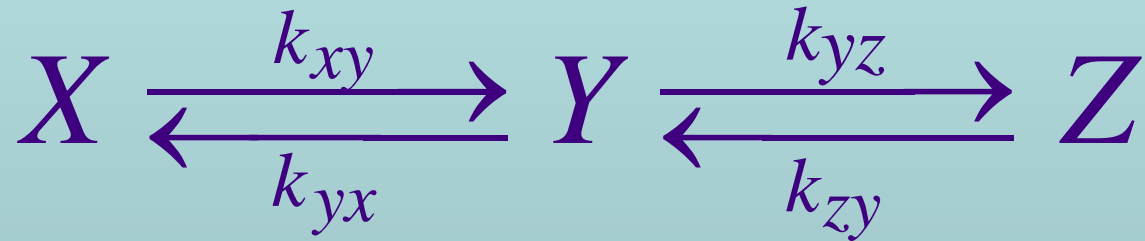
‘Current-in’
does not equal
‘Current-out’
in
Law of Mass Action



but **Kirchoff Current Law (i.e., Maxwell Eqns)**
requires

$$I_{AB} = I_{DE}$$

More specifically



$$I_{XY} - I_{YZ} = z_x F k_{xy} [X] - z_y F k_{yx} [Y] \\ - z_y F k_{yz} [Y] + z_z F k_{zy} [Z]$$

$$I_{XY} \neq I_{YZ}$$

Significance of Error

Asymmetry Determines Size of the Error

Special Case A*: Set all **charges equal to one**, along with **concentrations equal to one**,

$$\frac{\hat{I}_{XY} - \hat{I}_{YZ}}{F \cdot 1 \frac{\text{mole}}{\text{liter}}} = k_{xy} - k_{yx} - k_{yz} + k_{zy};$$

$$\text{Concentrations} = 1 \frac{\text{mole}}{\text{liter}}; z_X = z_Y = z_Z = 1$$

Special Case B: Alternatively, set all rate constants and all concentrations equal to one,

$$\frac{\hat{I}_{XY} - \hat{I}_{YZ}}{F \cdot 1 \frac{\text{mole}}{\text{liter}} \frac{1}{\text{sec}}} = z_X - z_Y - z_Y + z_Z;$$

$$\text{Concentrations} = 1 \frac{\text{mole}}{\text{liter}}; \text{rate constants} = 1 \frac{1}{\text{sec}}$$

Discontinuities in Current have Large Effects

$$I_{XY} - I_{YZ}$$

has large effects
exceeding breakdown voltages in
microseconds

Chemistry
is about
Isolated Substances

Engineering
is about
Signals
interactions with outside world,
not substances

Maxwell's Equations

Kirchoff's Current Law

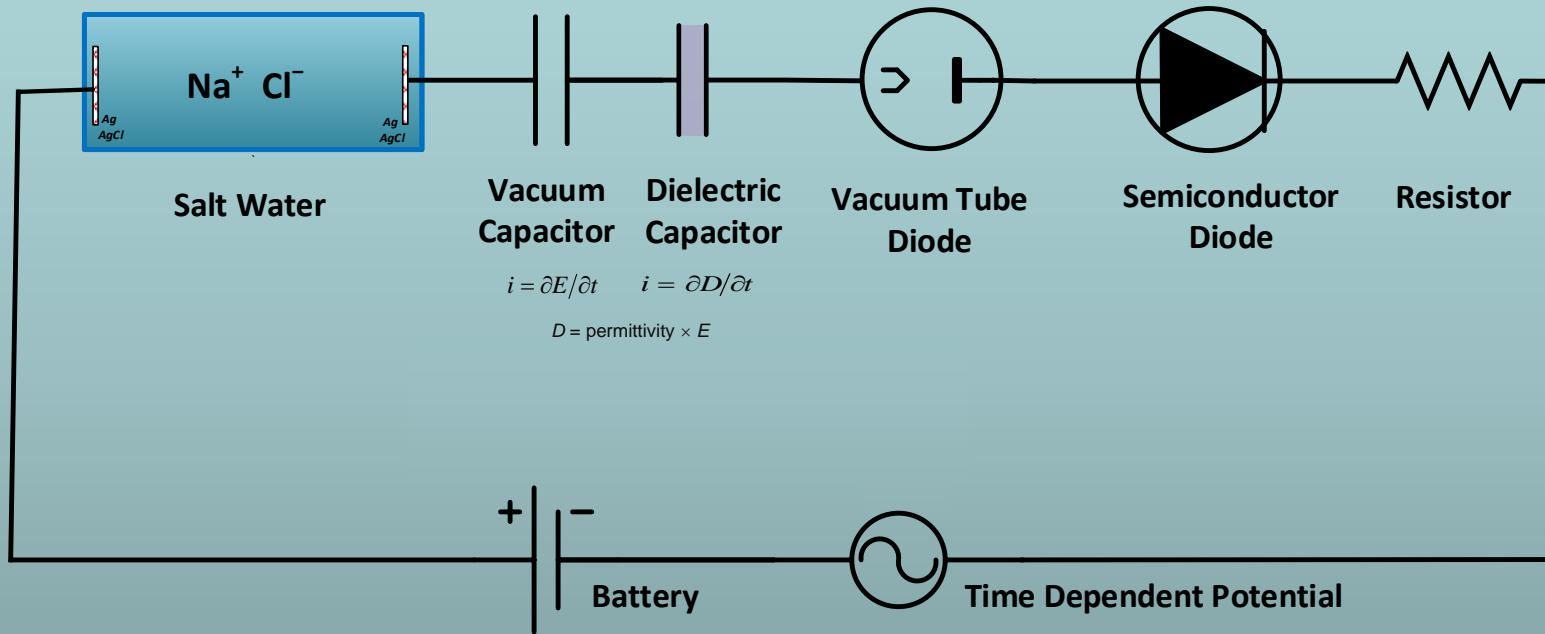
compute

Signals

from Conservation of Charge
and
Continuity of Current,
including displacement current

'Charge' is an Abstraction

with different physics in different systems



but **Continuity of Current is Exact**

No matter what carries the current!

**Currents are
NOT
just the sum of
Currents of Particles**

Current is Abstract

it includes

Displacement Current

and

Current of quasi-particles

for example

Currents are NOT just the sum of fluxes of Charged particles

Current is Abstract

it includes

Displacement Current

$$i_{\text{displacement}} = C(\partial V / \partial t) = i_{\Sigma} = \text{sum of fluxes of charges}$$

$\partial V / \partial t$ adjusts itself so generalized current has no discontinuities

Generalized Current $i_{\text{displacement}} + i_{\Sigma}$ is continuous

C is capacitance “to ground”, i. e. to infinity
Capacitance is charge divided by “self-energy”

Electricity is Different

it has a life of its own,
beyond mass

Mass accumulates

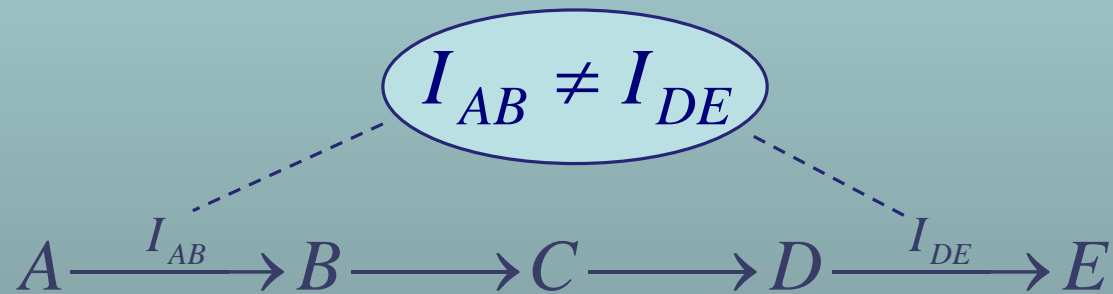
but voltage always changes so

GENERALIZED CURRENT

$$\dot{i}_{displacement} + \dot{i}_{\Sigma}$$

NEVER ACCUMULATES

‘Current-in’
does not equal
‘Current-out’
in
Law of Mass Action



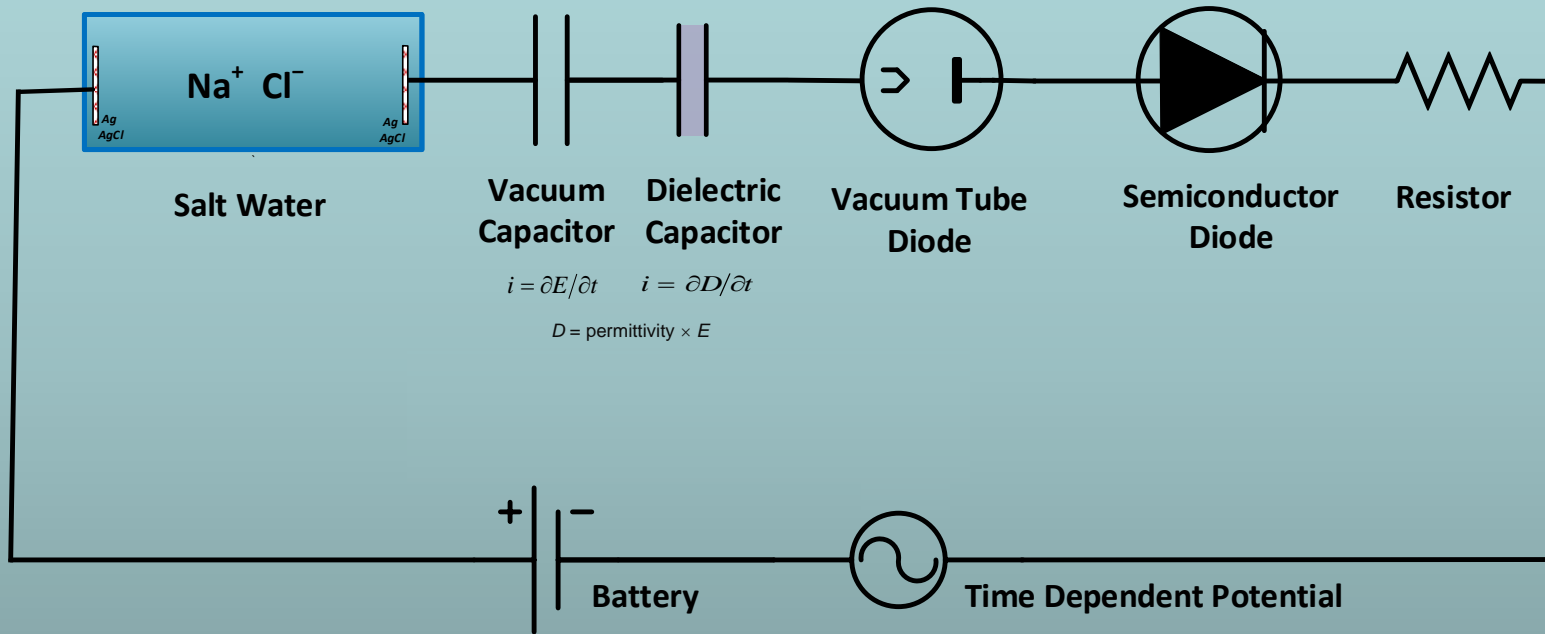
but **Kirchoff Current Law** (*i.e., Maxwell Eqns*)

requires

$$I_{AB} = I_{DE}$$

'Charge' is an Abstraction

with different physics in different systems



but **Continuity of Current is Exact**

No matter what carries the current!

Transferrable Models are not Possible

Rate constants chosen at one boundary charge or one potential
cannot work for different charges or potentials.

Currents in Rate Models

are

Independent of Charge and Potential

but

in the real world

Currents depend on Charge and Potential

Correlation between Currents

0.999 999 999 999 999 999

because

Conservation of Charge is exact

Kirchoff Continuity of Current Law

**Classical Chemical Reactions Assume
INDEPENDENT
uncorrelated
Rate Constants**

**Transferrable Models are not Possible
with this assumption**

Reconciling
Mass Action
and
Maxwell-Kirchoff

will no doubt be a

Long Journey

**“Journey
of a thousand miles
starts
with a single step”**

**in the right direction,
I beg to add to this Chinese saying**

Let's do Channels!
They are easier than bulk solution!!!

Biology is Easier than Physics
because reduced models exist!

Biology is Easier than Physics

Reduced Models Exist*

for important biological functions
or the

**Animal would not survive
to reproduce**

*Evolution provides the existence theorems and uniqueness conditions
so hard to find in theory of inverse problems.

*(Some biological systems – the human shoulder – are not robust,
probably because they are incompletely evolved,
i.e. they are in a local minimum 'in fitness landscape'.*

I do not know how to analyze these.

I can only describe them in the classical biological tradition.)

General Theme

**Mathematics of Molecular Biology
Provides Great Opportunity**

Biology Provides the Data

Engineering Provides the Approach

Mathematics Provides the Tools

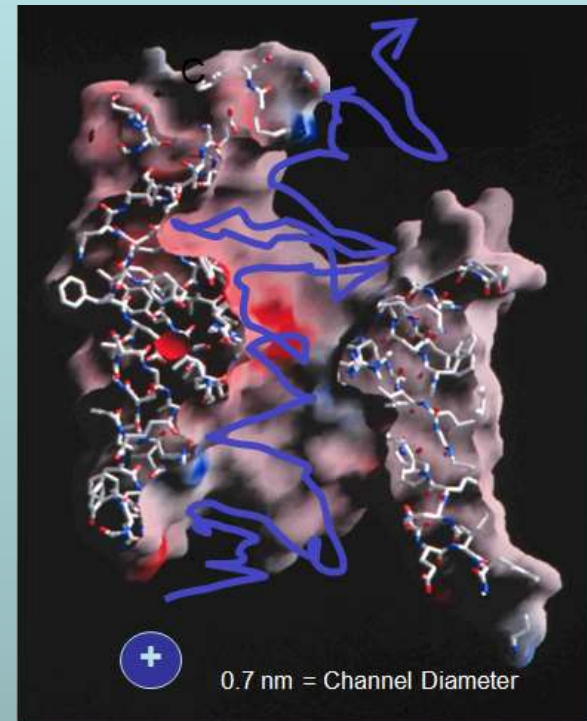
particularly variational methods that allow
'everything' to interact with 'everything' else



**Mathematics* Creates
our
Standard of Living**

Mathematics replaces Trial and Error with Computation

**e.g., Electricity, Computers, Fluid Dynamics,
Optics, Structural Mechanics,*



**Thousands of Molecular Biologists
Study Ion Channels Everyday,
One protein molecule at a time
using amplifiers like the
*AxoPatch***



Ion Channels are the Valves of Cells

Ion Channels are the Main Controllers of Biological Function

Selectivity

Different Ions
carry
Different Signals

Chemical Bonds are lines
Surface is Electrical Potential
Red is negative (acid)
Blue is positive (basic)



~30 Å

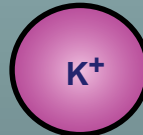
Figure of ompF porin by Raimund Dutzler

Ions in Water*

are the
Liquid of Life

*Pure H₂O is toxic to cells & proteins

Hard Spheres



3 Å

3 Å

ION CHANNELS – Biological Role

Ion channels coordinate contraction of cardiac muscle making the heart a pump

Ion channels coordinate contraction in skeletal muscle

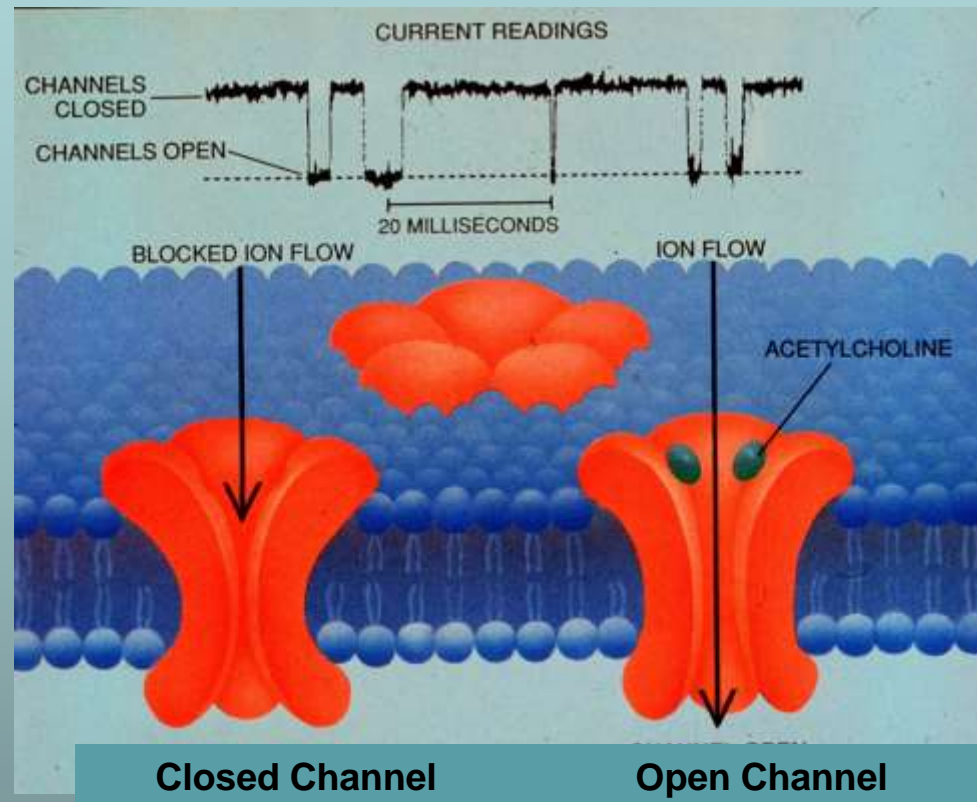
Ion channels control all electrical activity and produce nerve signals

Ion channels are involved in secretion and absorption in all cells: kidney, intestine, liver, adrenal glands, etc.

Ion channels are involved in thousands of diseases and many drugs act on channels

Ion channels are proteins with genes (blueprints) manipulated by molecular genetics

Ion channels have structures shown by x-ray crystallography in favorable cases



**Reduced models exist
because
they are the adaptation
created by evolution
to perform a biological function
like selectivity**

Reduced Models
and its parameters
are found by
Inverse Methods
of Reverse Engineering

**I have presented evidence for a satisfactory
reduced model of Calcium Channels in many
previous talks so I did not want to bore you with
again.**

Just send an email!!

**bob.eisenberg@gmail.com
or
beisenbe@rush.edu**

The End

Any Questions?

Evidence

Best Evidence is from the
RyR Receptor

Dirk Gillespie

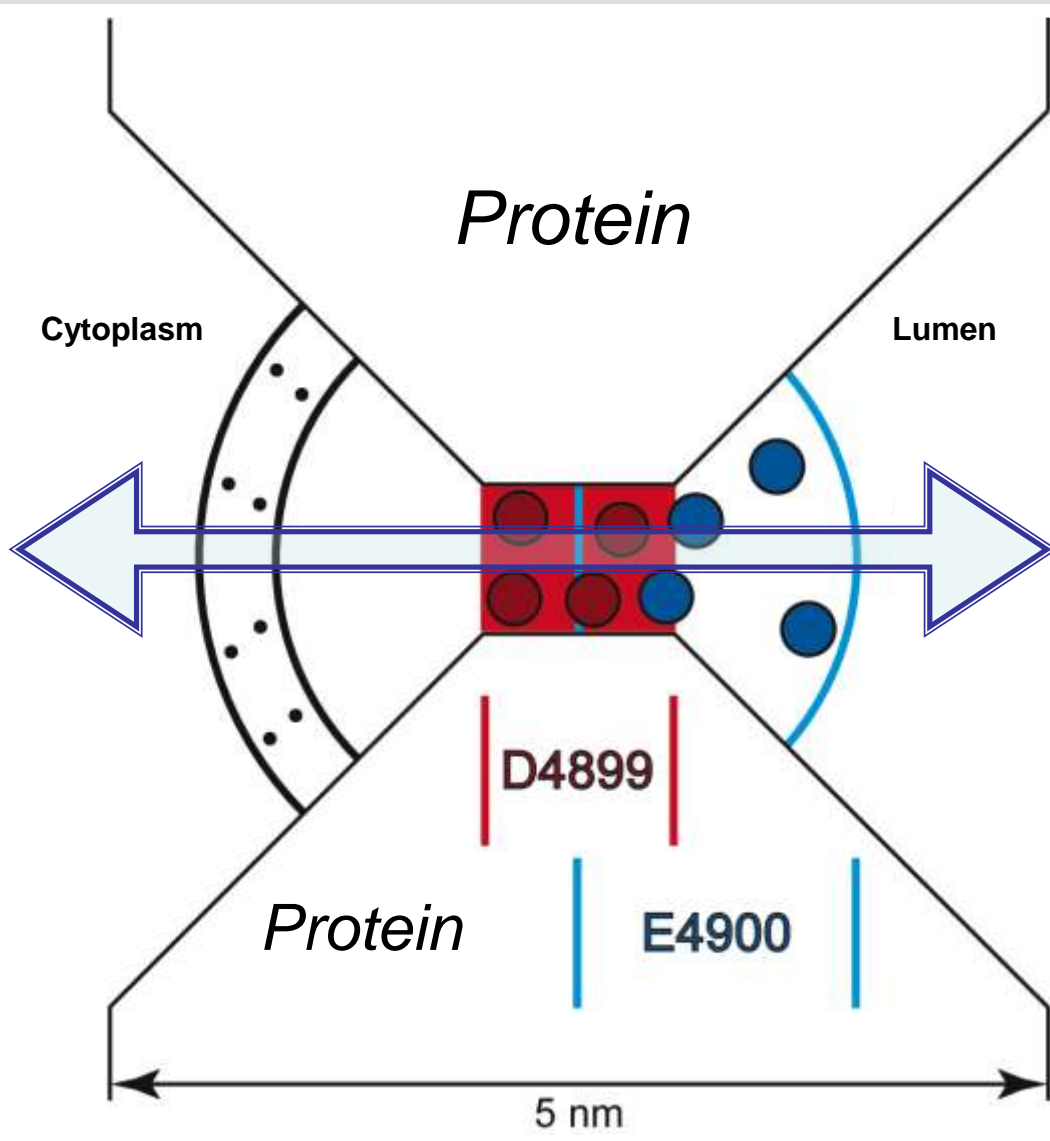


Gerhard Meissner, Le Xu, et al,
not Bob Eisenberg

- **More than 120 combinations of solutions & mutants**
- **7 mutants with significant effects fit successfully**

1. Gillespie, D., Energetics of divalent selectivity in a calcium channel: the ryanodine receptor case study. *Biophys J*, 2008. 94(4): p. 1169-1184.
2. Gillespie, D. and D. Boda, Anomalous Mole Fraction Effect in Calcium Channels: A Measure of Preferential Selectivity. *Biophys. J.*, 2008. 95(6): p. 2658-2672.
3. Gillespie, D. and M. Fill, Intracellular Calcium Release Channels Mediate Their Own Countercurrent: Ryanodine Receptor. *Biophys. J.*, 2008. 95(8): p. 3706-3714.
4. Gillespie, D., W. Nonner, and R.S. Eisenberg, Coupling Poisson-Nernst-Planck and Density Functional Theory to Calculate Ion Flux. *Journal of Physics (Condensed Matter)*, 2002. 14: p. 12129-12145.
5. Gillespie, D., W. Nonner, and R.S. Eisenberg, Density functional theory of charged, hard-sphere fluids. *Physical Review E*, 2003. 68: p. 0313503.
6. Gillespie, D., Valisko, and Boda, Density functional theory of electrical double layer: the RFD functional. *Journal of Physics: Condensed Matter*, 2005. 17: p. 6609-6626.
7. Gillespie, D., J. Giri, and M. Fill, Reinterpreting the Anomalous Mole Fraction Effect. The ryanodine receptor case study. *Biophysical Journal*, 2009. 97: p. pp. 2212 - 2221
8. Gillespie, D., L. Xu, Y. Wang, and G. Meissner, (De)constructing the Ryanodine Receptor: modeling ion permeation and selectivity of the calcium release channel. *Journal of Physical Chemistry*, 2005. 109: p. 15598-15610.
9. Gillespie, D., D. Boda, Y. He, P. Apel, and Z.S. Siwy, Synthetic Nanopores as a Test Case for Ion Channel Theories: The Anomalous Mole Fraction Effect without Single Filing. *Biophys. J.*, 2008. 95(2): p. 609-619.
10. Malasics, A., D. Boda, M. Valisko, D. Henderson, and D. Gillespie, Simulations of calcium channel block by trivalent cations: Gd(3+) competes with permeant ions for the selectivity filter. *Biochim Biophys Acta*, 2010. 1798(11): p. 2013-2021.
11. Roth, R. and D. Gillespie, Physics of Size Selectivity. *Physical Review Letters*, 2005. 95: p. 247801.
12. Valisko, M., D. Boda, and D. Gillespie, Selective Adsorption of Ions with Different Diameter and Valence at Highly Charged Interfaces. *Journal of Physical Chemistry C*, 2007. 111: p. 15575-15585.
13. Wang, Y., L. Xu, D. Pasek, D. Gillespie, and G. Meissner, Probing the Role of Negatively Charged Amino Acid Residues in Ion Permeation of Skeletal Muscle Ryanodine Receptor. *Biophysical Journal*, 2005. 89: p. 256-265.
14. Xu, L., Y. Wang, D. Gillespie, and G. Meissner, Two Rings of Negative Charges in the Cytosolic Vestibule of T Ryanodine Receptor Modulate Ion Fluxes. *Biophysical Journal*, 2006. 90: p. 443-453.

The Geometry



Selectivity Filter

- is 10 Å long and 8 Å in diameter
- confines four **D4899** negative amino acids
- Four **E4900** positive amino acids are on luminal side, overlapping D4899
- **Cytosolic distributed charge**

Ryanodine Receptor Pore

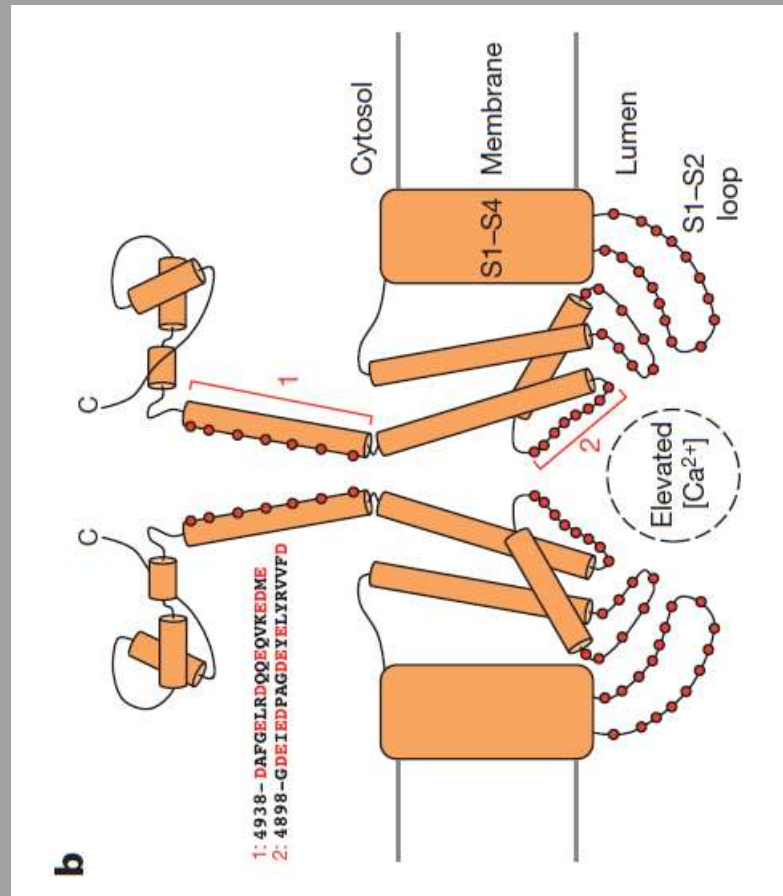
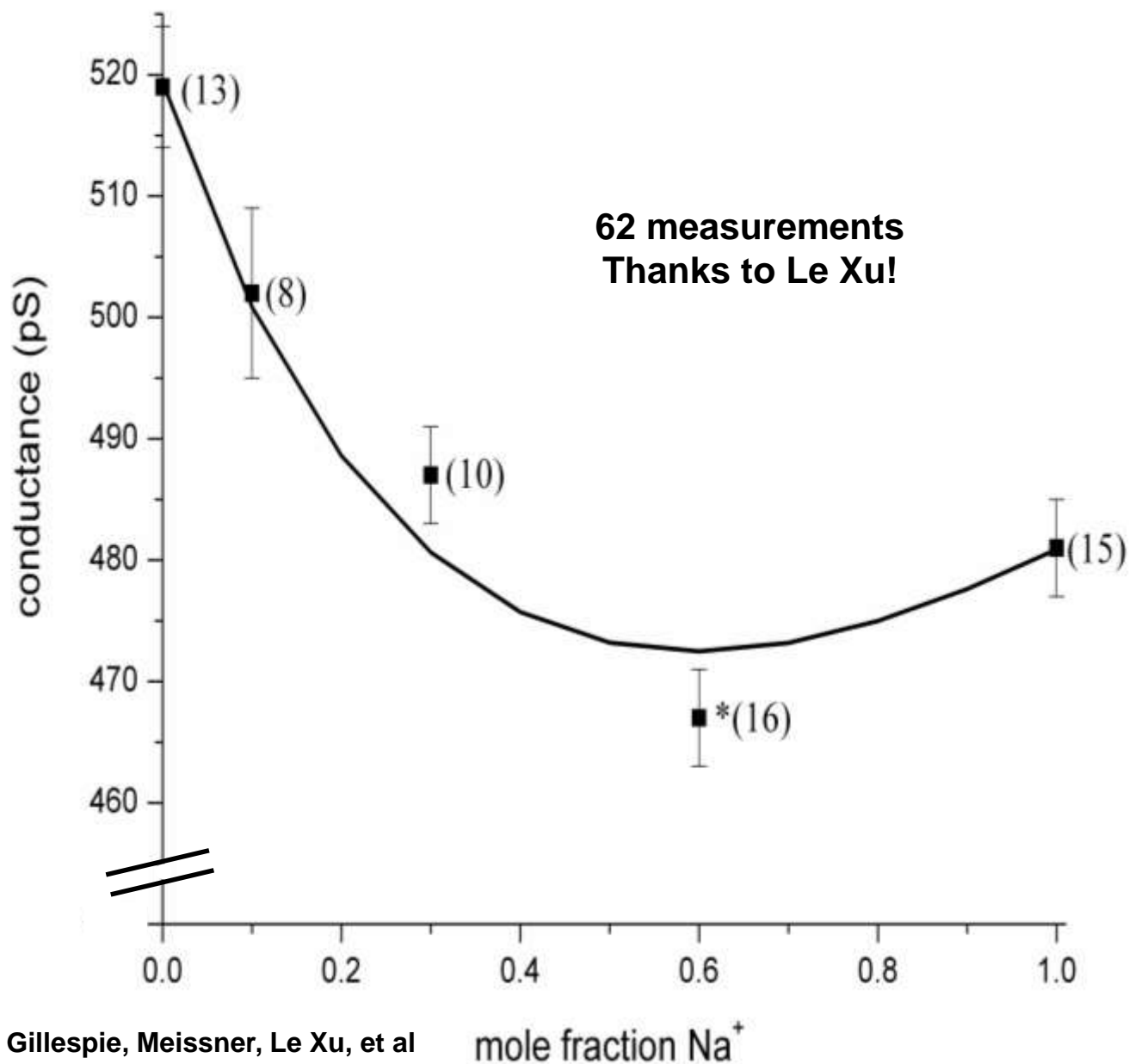


Fig 3 The RyR1 conduction pathway

from Zalk et al, Nature, 2014, 10.1038/nature13950

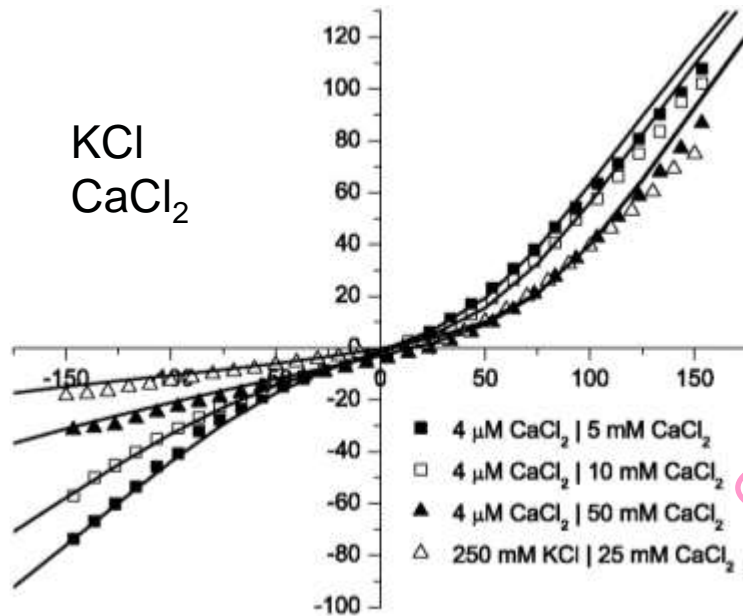
“b, Scheme ... of all the negatively charged residues in the ionic pathway (red dots) and the [other] negatively charged residues”

The model predicted an AMFE for Na⁺/Cs⁺ mixtures before it had been measured

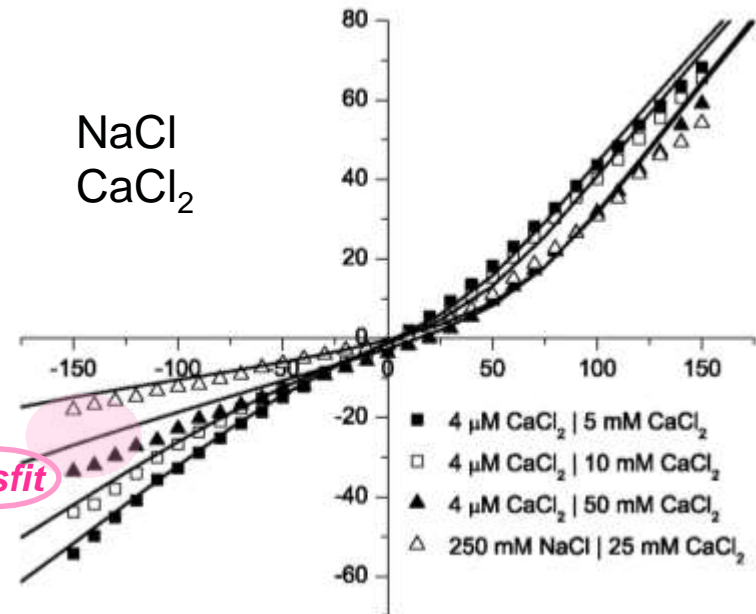


Divalents

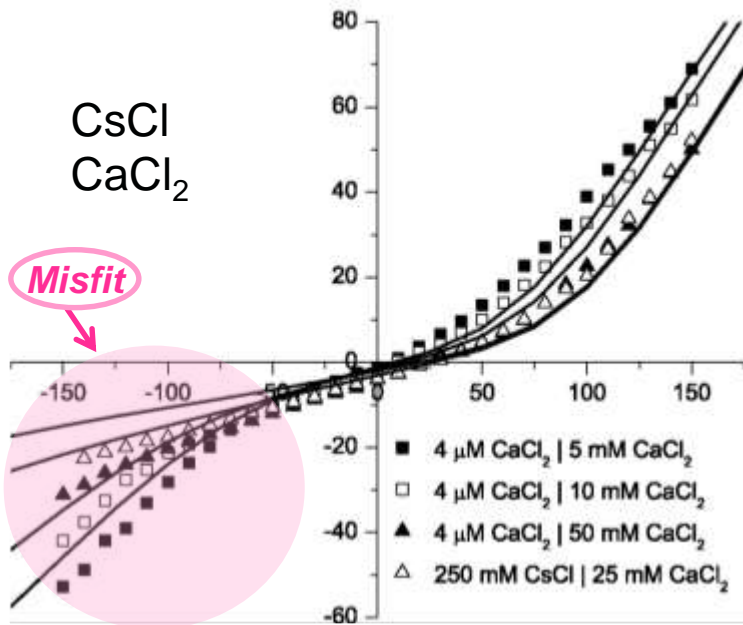
KCl
CaCl₂



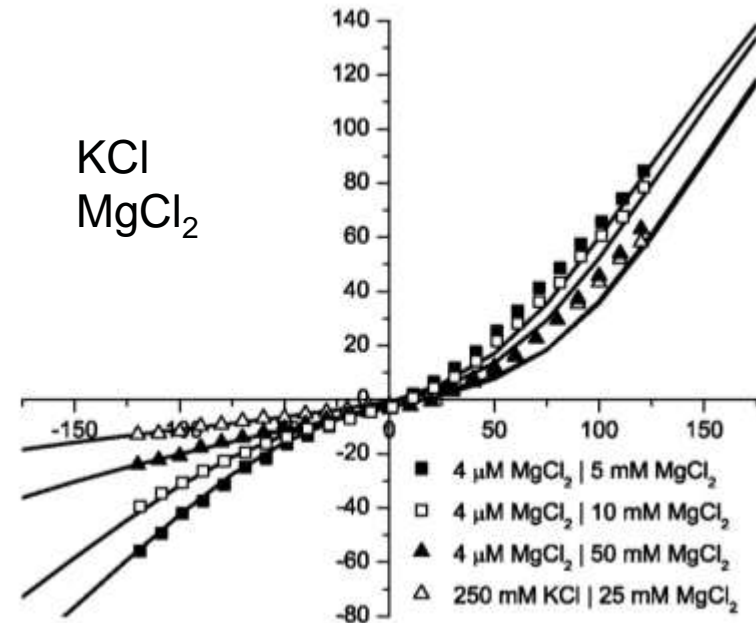
NaCl
CaCl₂

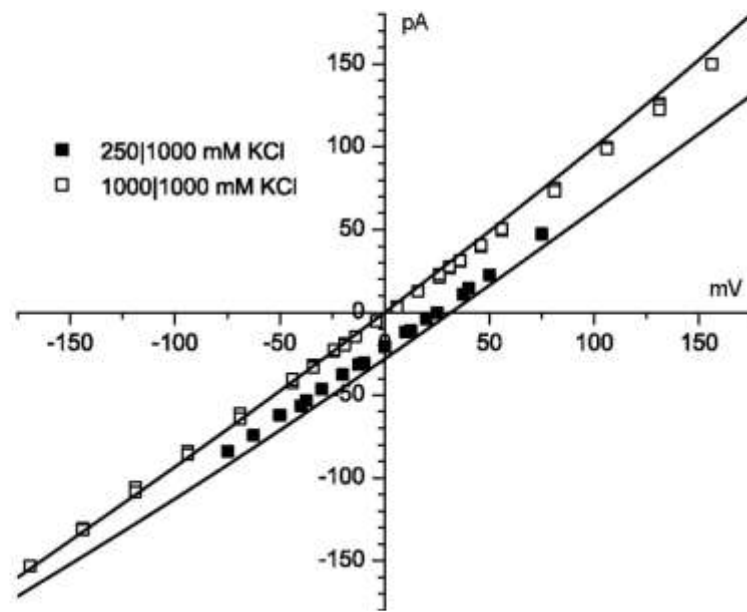
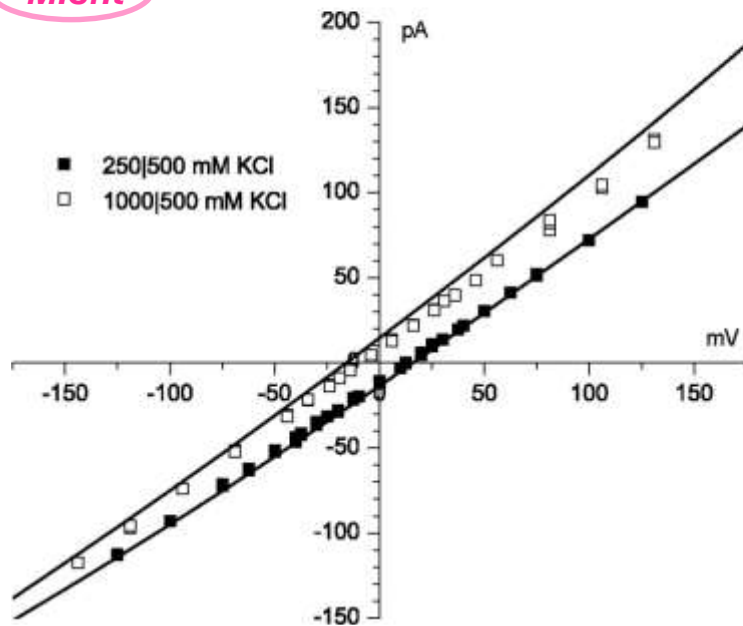
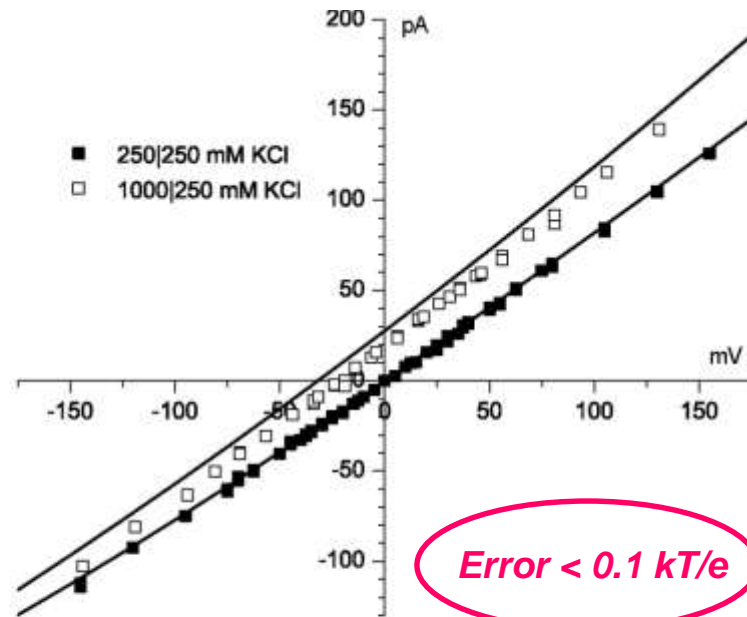
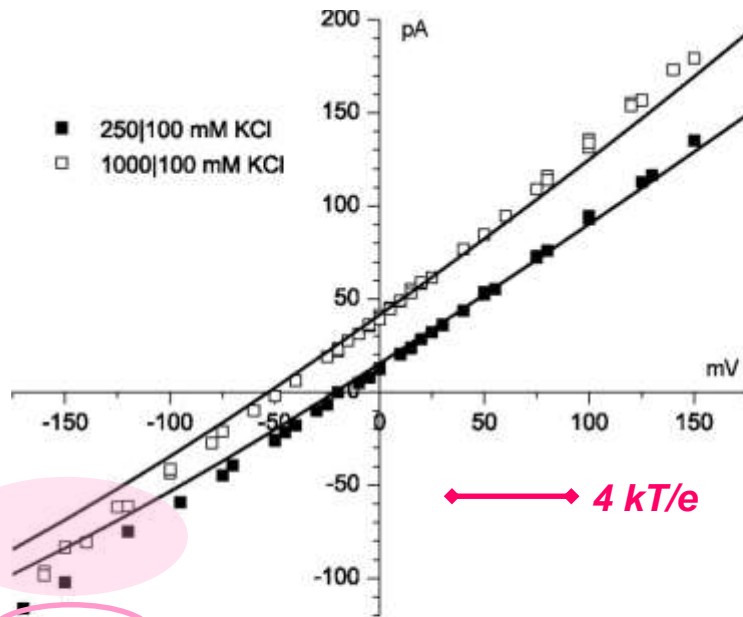


CsCl
CaCl₂



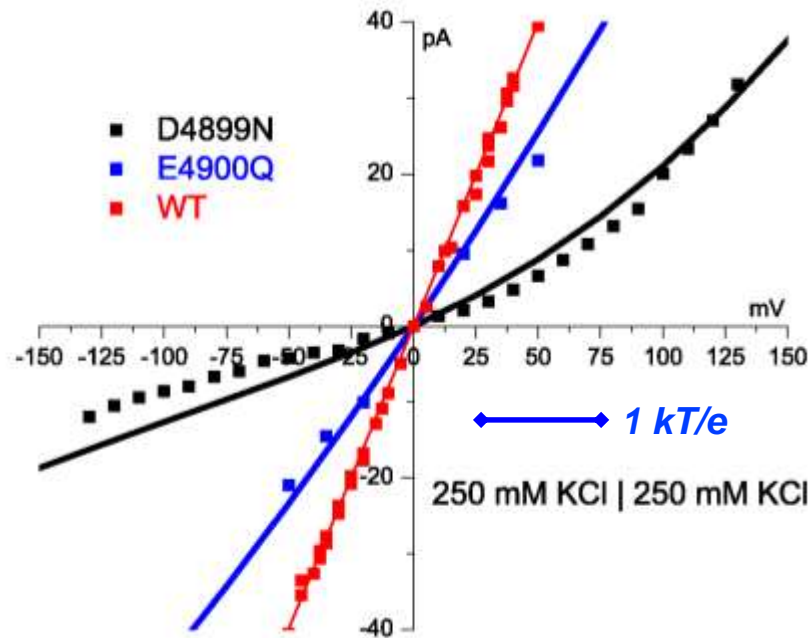
KCl
MgCl₂





Theory fits Mutation with Zero Charge

Theory Fits Mutant in K

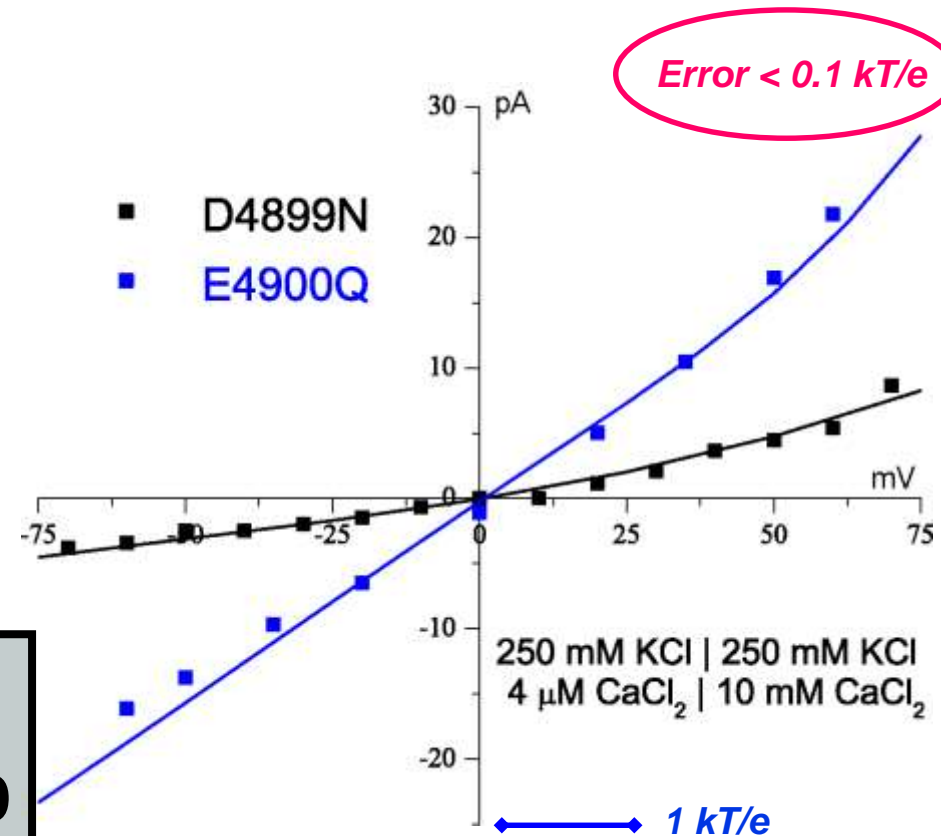


Protein charge density
wild type* $13 \text{ M} \Rightarrow 0 \text{ M}$ in D4899

Water is 55 M

*some wild type curves not shown, 'off the graph'

Theory Fits Mutant in K + Ca



Gillespie *et al*

J Phys Chem 109 15598 (2005)

Calcium Channel

has been examined in ~35 papers, e.g.,



- Nonner, W., D. P. Chen, and B. Eisenberg. 1998. Anomalous Mole Fraction Effect, Electrostatics, and Binding in Ionic Channels. *Biophysical Journal* 74:2327-2334.
- Nonner, W., L. Catacuzzeno, and B. Eisenberg. 2000. Binding and Selectivity in L-type Ca Channels: a Mean Spherical Approximation. *Biophysical Journal* 79:1976-1992.
- Nonner, W., D. Gillespie, D. Henderson, and B. Eisenberg. 2001. Ion accumulation in a biological calcium channel: effects of solvent and confining pressure. *J Physical Chemistry B* 105:6427-6436.
- Boda, D., W. Nonner, D. Henderson, B. Eisenberg, and D. Gillespie. 2008. Volume exclusion in calcium selective channels. *Biophys. J.:biophysj*.107.122796.
- Boda, D., M. Valisko, B. Eisenberg, W. Nonner, D. Henderson, and D. Gillespie. 2006. Effect of Protein Dielectric Coefficient on the Ionic Selectivity of a Calcium Channel. *Journal of Chemical Physics* 125:034901.
- Boda, D., T. Varga, D. Henderson, D. Busath, W. Nonner, D. Gillespie, and B. Eisenberg. 2004. Monte Carlo simulation study of a system with a dielectric boundary: application to calcium channel selectivity. *Molecular Simulation* 30:89-96.
- Boda, D., M. Valisko, B. Eisenberg, W. Nonner, D. Henderson, and D. Gillespie. 2007. The combined effect of pore radius and protein dielectric coefficient on the selectivity of a calcium channel. *Physical Review Letters* 98:168102.

Most of the papers are available at

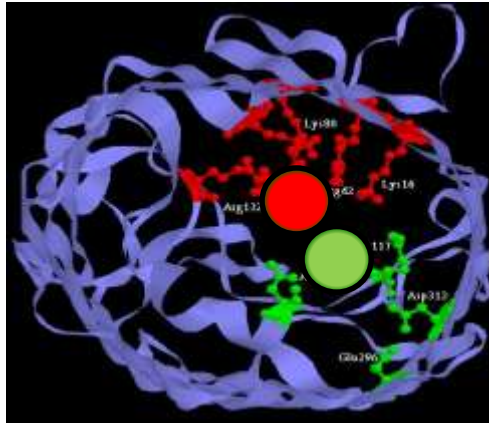
ftp://ftp.rush.edu/users/molebio/Bob_Eisenberg/Reprints

<http://www.phys.rush.edu/RSEisenberg/physioeis.html>

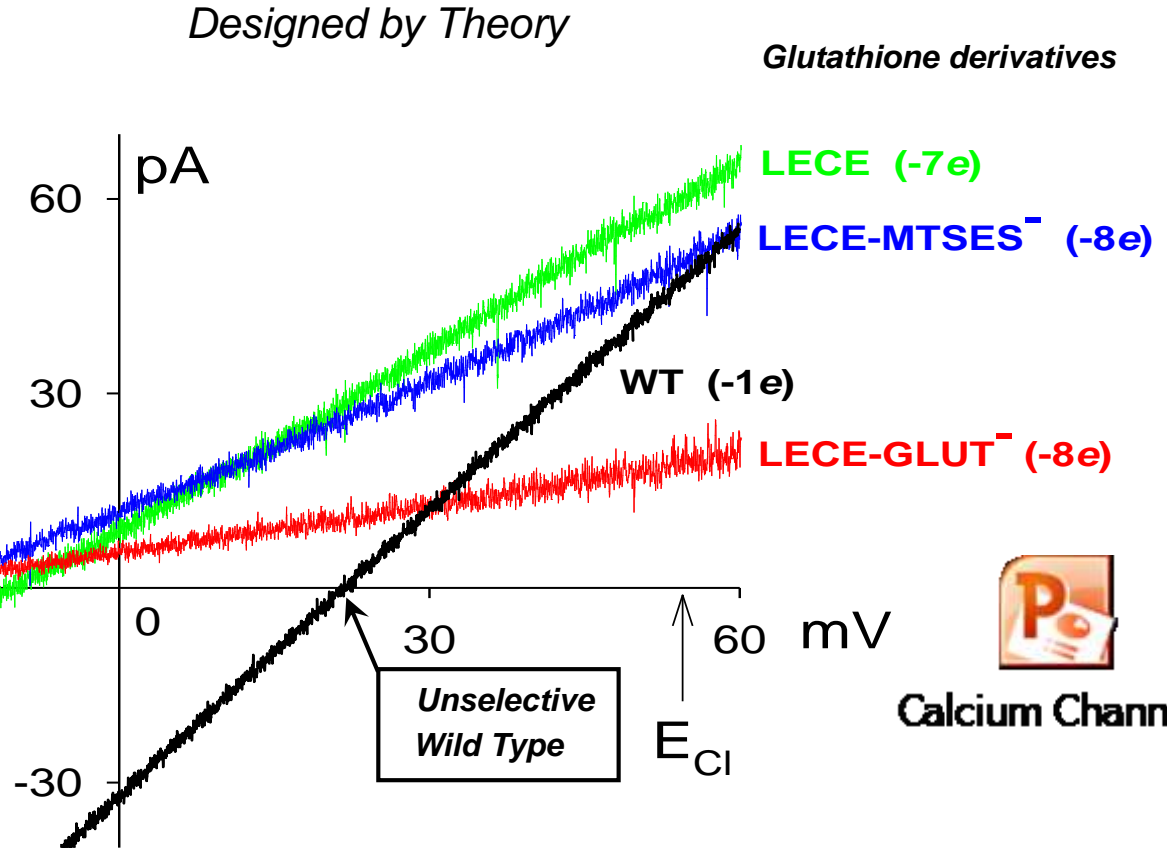
Experiments have built

Atomic Scale

Two Synthetic Calcium Channels



Calcium selective



Calcium Channels

As density of permanent charge increases, channel becomes calcium selective

$E_{\text{rev}} \rightarrow E_{\text{Ca}}$ in 0.1M || 1.0 M CaCl₂

Macro Scale

built by Henk Miedema, Wim Meijberg of BioMade Corp., Groningen, Netherlands

69

Miedema et al, *Biophys J* 87: 3137–3147 (2004)

Selectivity Filter

Crowded with Charge

L type Ca Channel

Selectivity
Filter

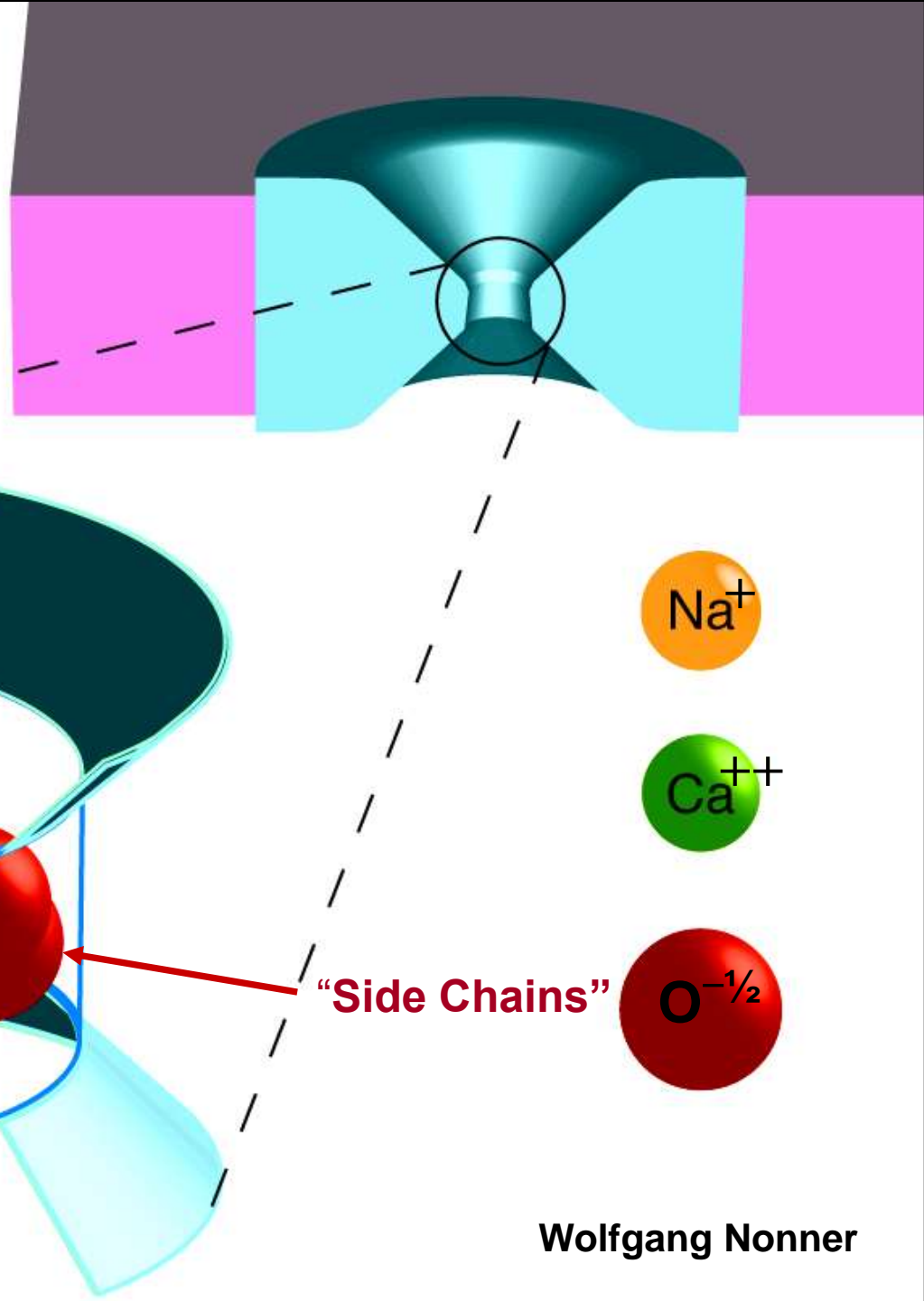
“Side Chains”

Na^+

Ca^{++}

$\text{O}^{-1/2}$

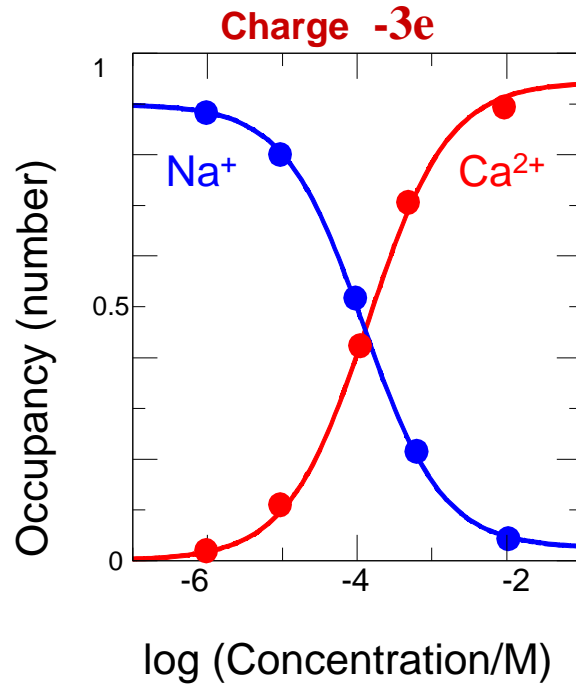
Wolfgang Nonner



Ca Channel



E
E
E
A



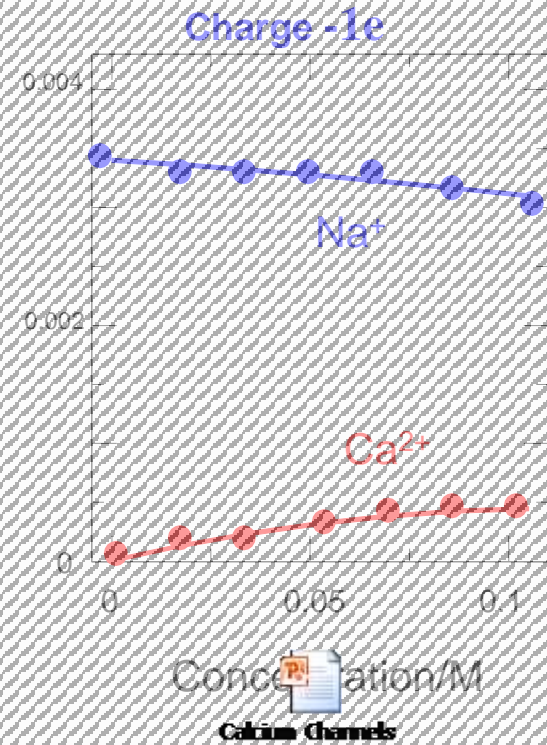
EEEE has full biological selectivity
in similar simulations

Mutation

Same Parameters

Na Channel

D
E
K
A



Boda, et al

Selectivity

comes from

Electrostatic Interaction

and

Steric Competition for Space



Repulsion

Location and Strength of Binding Sites
Depend on Ionic Concentration and
Temperature, etc

Rate Constants are Variables

Sodium Channel

Voltage controlled channel responsible for signaling in nerve and coordination of muscle contraction

Challenge

from leading biophysicists

Walter Stühmer and Stefan Heinemann

Göttingen

Leipzig

Max Planck Institutes

**Can THEORY explain the MUTATION
Calcium Channel into Sodium Channel?**

DEEA  **DEKA**

*Calcium
Channel*

*Sodium
Channel*

Ca Channel

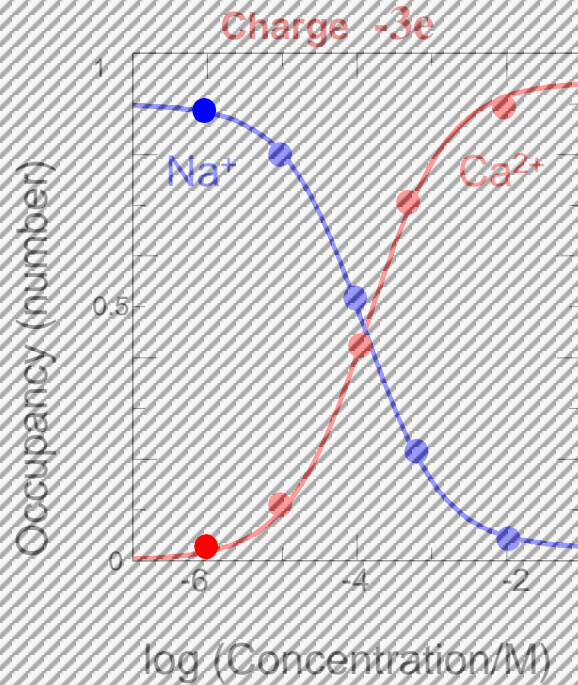
E
E
E
A

Mutation

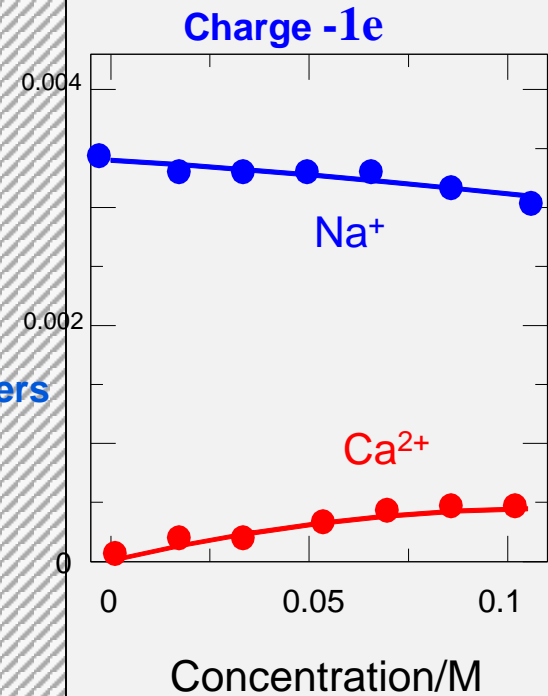
Same Parameters

Na Channel

D
E
K
A



EEEE has full biological selectivity
in similar simulations



Boda, et al

Nothing was changed
from the
EEEE Ca channel
except the amino acids

**Calculated DEKA Na Channel
Selects**

Ca²⁺ vs. Na⁺ and also K⁺ vs. Na⁺



Metropolis Monte Carlo

Simulates Location of Ions

both the mean and the variance

Details:

- 1) Start with Configuration A , with computed energy E_A
- 2) Move an ion to location B , with computed energy E_B
- 3) If spheres overlap, $E_B \rightarrow \infty$ and configuration is rejected
- 4) If spheres do not overlap, $E_B \rightarrow 0$ and configuration is accepted
- 5) If $E_B < E_A$: accept new configuration.
- 6) If $E_B > E_A$: accept new configuration with probability $\exp[-(E_A - E_B)/k_B T]$

Key idea

MMC chooses configurations with a Boltzmann probability and weights them evenly
instead of
choosing them from uniform distribution and then weighting them with $\exp(-E/k_B T)$