

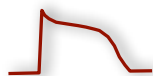
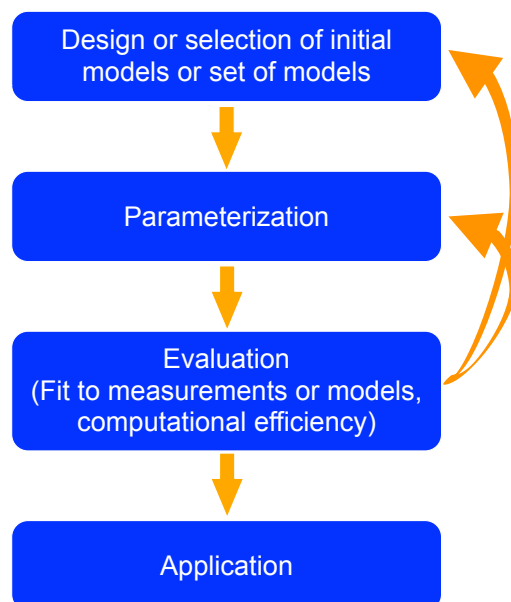
Whole Cell Simulation



Whole Cell Simulation

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Motivation and Approach




From Sachse Lectures


Whole Cell Simulation

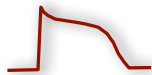
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Types of Cellular Electrophysiology Models

- **Molecular Models**
 - Structure and dynamics
 - Molecular interactions, drug binding, ion movement
- **Markov Models**
 - Current through a single channel and population of channels as well as gating currents
 - Based on states and transitions


Lecture Week 6 (Frank Sachse)
- **Hodgkin-Huxley Models**
 - Current through a population of channels and single channels as well as gating currents
 - Based on gating variables and rate coefficients

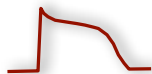
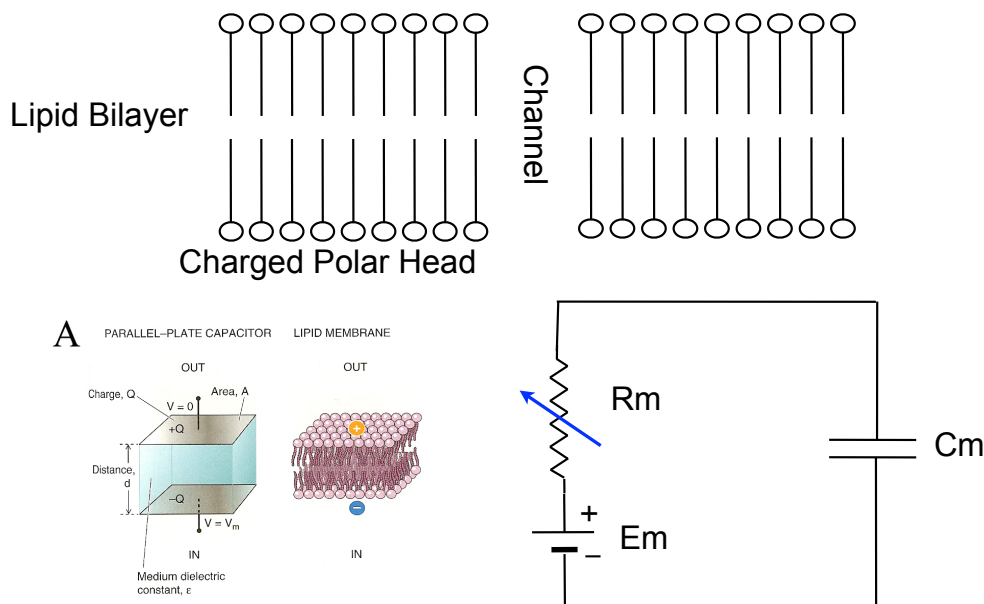

This week!



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Membrane Equivalent Circuit

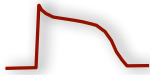


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Modeling Excitable Membranes

- Main concepts
 - Driving Force
 - V-Veq for each ion independently
 - Model Formalism (Hodgkin-Huxley)
 - continuous vs. discrete/stochastic
 - describes whole cell not single channels
 - Changes required for cardiac cells
 - additional currents, different coefficients
 - Numerical solutions
 - solve set of ODEs at discrete time steps
- Background
 - Week #7 lectures/slides (Frank Sachse)



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Hodgkin Huxley Formalism

$$i_m = i_1 + i_2 + i_3 \dots + I_c$$

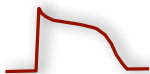
$$i_m = i_K + i_{Na} + i_L + I_c$$

$$i_K(V, t) = (V_m - E_K)g_K(V, t)$$

$$i_{Na}(V, t) = (V_m - E_{Na})g_{Na}(V, t)$$

$$i_L = (V_m - E_L)g_L$$

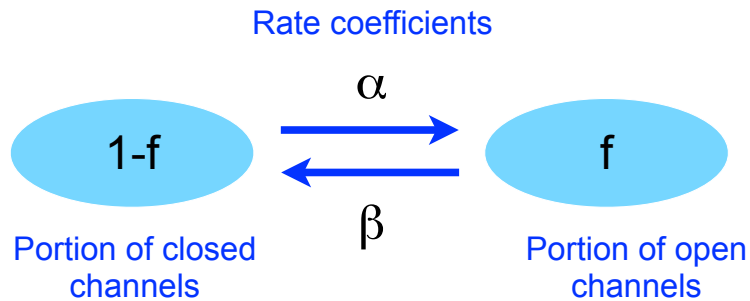
$$I_C = -C \frac{dV}{dt}$$



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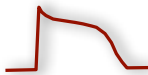
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Basic HH Concept



$$\frac{df}{dt} = \alpha_f(1 - f) - \beta_f f$$

f = Gating variable



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Potassium Current

$$i_K(V, t) = (V_m - E_K)g_K(V, t)$$

We can write an expression for the conductivity of this ion as

$$g_K = \bar{g}_K n^4(V, t)$$

And then assume a simple first order kinetic behavior of the gating variable n

$$\frac{dn}{dt} = \alpha_n(V)(1 - n) - \beta_n(V)n$$

Which has a solution for the case of constant V (voltage clamp) of the form

$$n(t) = n_\infty(v_o) - (n_\infty(v_o) - n_0)e^{-t/\tau_n(v_o)}$$



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Potassium Current II

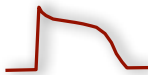
With the voltage dependent variables defined as follows:

$$\alpha_n(V) = \frac{n_\infty}{\tau_n} \quad \beta_n(V) = \frac{(1 - n_\infty)}{\tau_n}$$

and

$$n_0 = \frac{\alpha_{n0}}{\alpha_{n0} + \beta_{n0}}$$

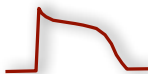
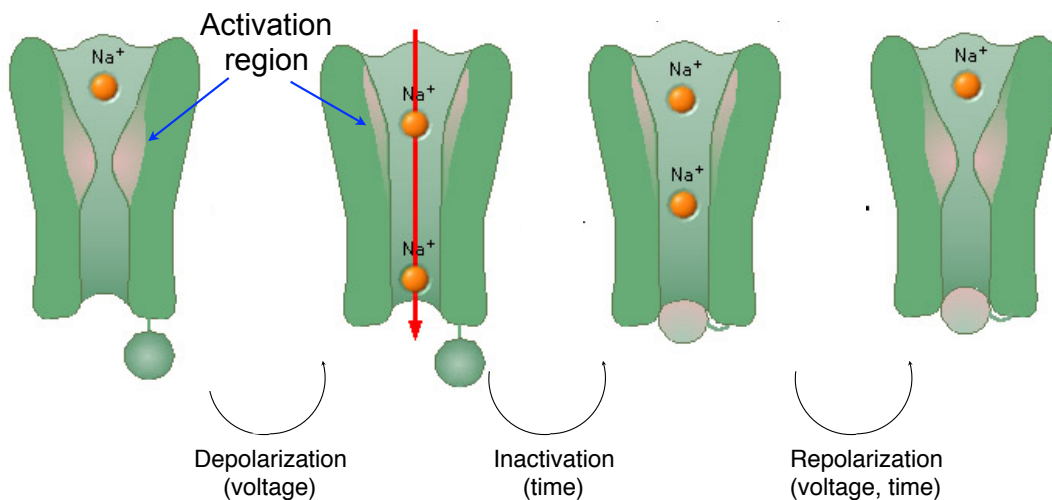
the value of n with the membrane at rest



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Sodium Channel Behavior



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Sodium Current Activation I

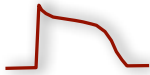
$$i_{Na}(V, t) = (V_m - E_{Na})g_{Na}(V, t)$$

which we write as the product of two first order kinetic terms

$$g_{Na} = \bar{g}_{Na} m^3(V, t) h(V, t)$$

Taking the first of these, we can write

$$\frac{dm}{dt} = \alpha_m(V)(1 - m) - \beta_m(V)m = \frac{1}{\tau_m}(m_\infty(V) - m)$$



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Sodium Current Activation II

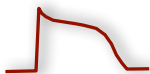
With constant voltage (voltage clamp)

$$m = m_\infty(v_o) - (m_\infty(v_o) - m_0)e^{-t/\tau_m(v_o)}$$

with the constants

$$\tau_m = \frac{1}{\alpha_m(V) + \beta_m(V)}$$

$$m_\infty = \frac{\alpha_m(V)}{\alpha_m(V) + \beta_m(V)}$$



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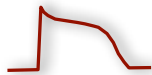
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Sodium Current (Inactivation)

$$g_{Na} = \bar{g}_{Na} m^3(V, t) h(V, t)$$

for which we write a similar ordinary differential equation

$$\frac{dh}{dt} = \alpha_h(V)(1 - h) - \beta_h(V)h = \frac{1}{\tau_h}(h_\infty(V) - h)$$



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Sodium Current (Inactivation) II

If we hold $V=v_0$, a constant (voltage clamp) then

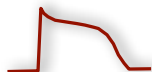
$$h = h_\infty(v_0) - (h_\infty(v_0) - h_0)e^{-t/\tau_h(v_0)}$$

with

$$h_\infty = \frac{\alpha_h(V)}{\alpha_h(V) + \beta_h(V)}$$

and

$$\tau_h = \frac{1}{\alpha_h(V) + \beta_h(V)}$$



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Summary of Fitted Rate Constants

$$\alpha_m = 0.1 \frac{25 - v}{e^{\frac{25-v}{10}} - 1}$$

$$\alpha_n = 0.01 \frac{10 - v}{e^{\frac{10-v}{10}} - 1}$$

$$\beta_m = 4e^{-v/18}$$

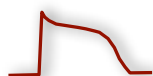
$$\beta_n = 0.125e^{-v/80}$$

$$\alpha_h = 0.07e^{-v/20}$$

$$\beta_h = \frac{1}{e^{\frac{30-v}{10}} + 1}$$

$$v = V - V_{eq}$$

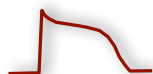
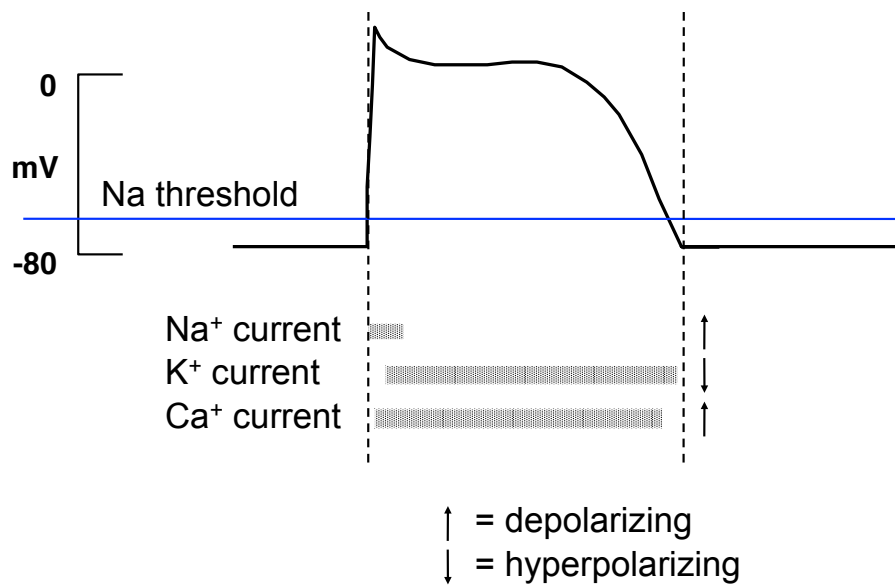
$$\bar{g}_{Na} = 120; \bar{g}_K = 36; g_L = 0.3$$



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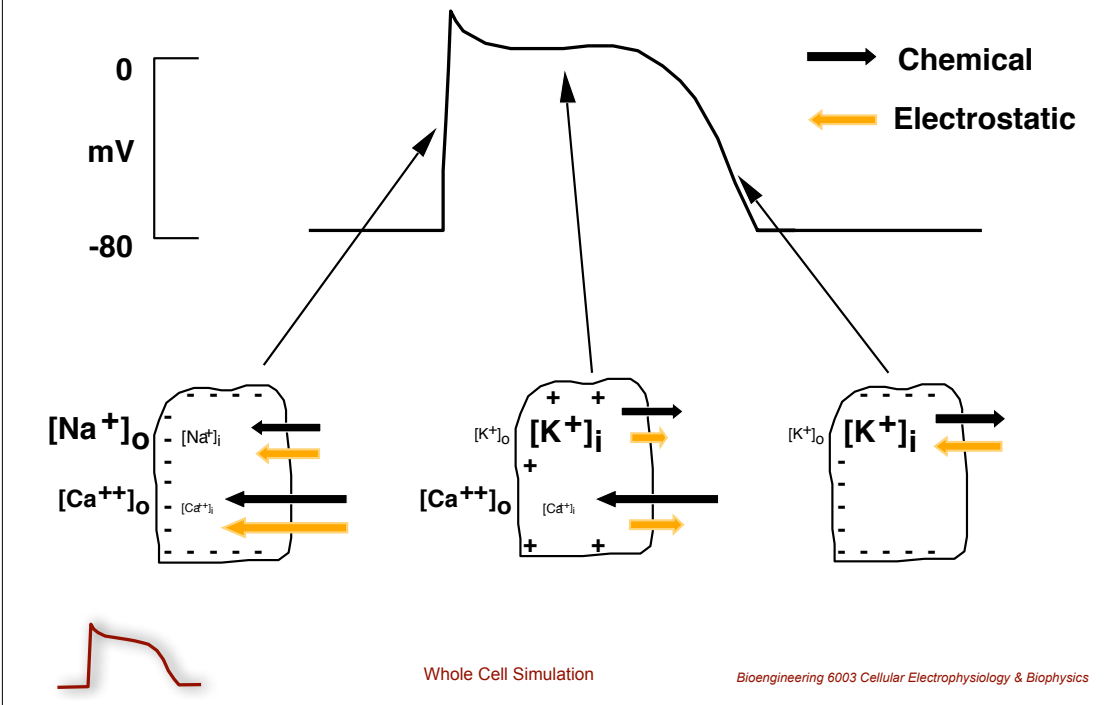
Cardiac Action Potential



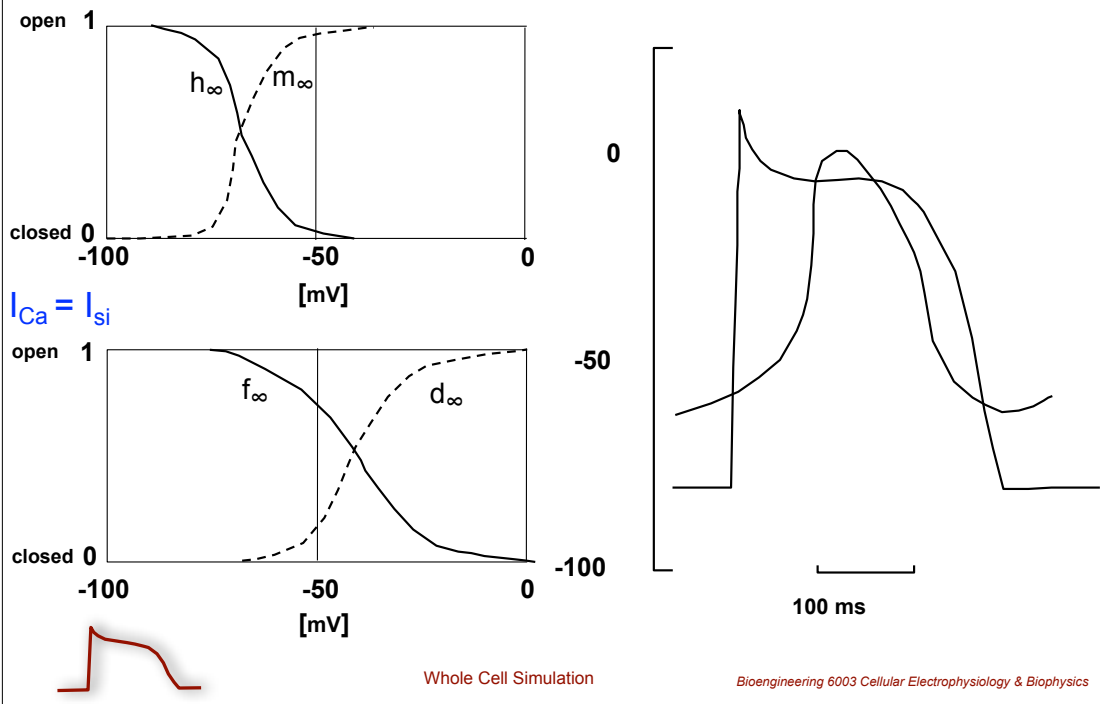
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Cardiac Cell Currents

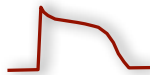


I_{Na} HH Gating Parameters in Cardiac Cells



Elements of Cell Simulation

- Currents
 - Sodium
 - Potassium (20+ types)
 - Calcium (L type and T type)
 - Leak/background currents
- Pumps
 - Sodium/potassium
 - Calcium
 - Proton
- Exchangers
 - Calcium/Sodium
- Buffers
 - Sarcoplasmic Reticulum (SR)
 - Mitochondria

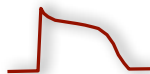


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Challenges of Cardiac Cell Modeling

- Range of ionic channels
 - Across cells and within single cells
- Range of cell types
 - Discrete (e.g., SA nodal, AV nodal, ventricular)
 - Heterogeneity within the heart (e.g., endocardial, epicardial, and midmyocardial)
- Range of species
- Results:
 - Models are more qualitative than quantitative
 - Each requires balance of detail and tweaking to match experiments

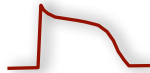


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Cardiac Membrane Models (a sampling)

- Nobel(1962) Purkinje cells (first cardiac cell model)
- McAllister, Noble, & Tsein (1975) Purkinje muscle
- Beeler-Reuter (1977) mammalian ventricle (first of its kind)
- Ebihara & Johnson (1981) B&R + updated sodium channel
- Luo-Rudy Phase I (1991) guinea pig ventricular muscle
- Luo-Rudy Phase II (1994) guinea pig ventricular muscle
- Demir, Clark Murphey, & Gilles (1994) rabbit SA Node
- Priebe & Beukelmann (1998) human ventricle
- Courtemanche & Ramirez (1998) human atrium
- Nobel, Varghese, Kohl, & Noble (1998) guinea pig ventricle
- Winslow, Rice, et al. (1999) canine ventricle + EC coupling
- Sachse, Seemann, Chaisaowong & Wieß (2003) human ventricle
- Ten Tusscher, Noble, Noble, Panfilov (2003) human ventricle



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Nobel (1962): Purkinje

Noble, D. *The Initiation of the Heartbeat*, 1975

Goal: show that oscillation possible in a membrane model

$$i_m = i_{Na} + i_{Na,b} + i_{K1} + i_{K2} + i_{An}$$

and a background current with constant conductivity

$$i_{Na,b} = g_{Na,b}(V_m - E_{Na})$$

as well an anion current with constant conductivity

$$i_{An} = g_{An}(V_m - E_{An})$$



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Nobel (1962): Sodium Currents

There are two different inward (Na^+) currents,

1) Voltage dependent that switches on and then off very quickly (with increasing V_m)

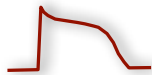
$$i_{Na} = g_{Na}(V_m - E_{Na})$$

$$g_{Na} = g_{\bar{Na}} m^3 h$$

2) Time dependent, acts like a classic HH K^+ current but with long time constant, i.e., 100 times longer than in nerve. “Delayed rectifier” because it is slow and primarily outward.

$$i_{K2} = g_{K2}(V_m - E_k)$$

$$g_{K2} = g_{\bar{K}2} n^4$$



Whole Cell Simulation

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Nobel (1962): Potassium Currents

There are two different outward (K^+) currents,

1) Voltage dependent that switches on and then off very quickly (with increasing V_m)

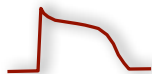
$$i_{K1} = g_{K1}(V_m - E_K)$$

$$g_{K1} = 1.2 e^{-\frac{V_m}{50}} + 0.015 e^{\frac{V_m+90}{60}}$$

2) Time dependent, acts like a classic HH K^+ current but with long time constant, i.e., 100 times longer than in nerve. “Delayed rectifier” because it is slow and primarily outward.

$$i_{K2} = g_{K2}(V_m - E_k)$$

$$g_{K2} = g_{\bar{K}2} n^4$$



Whole Cell Simulation

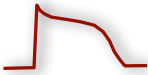
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Nobel (1962): Results

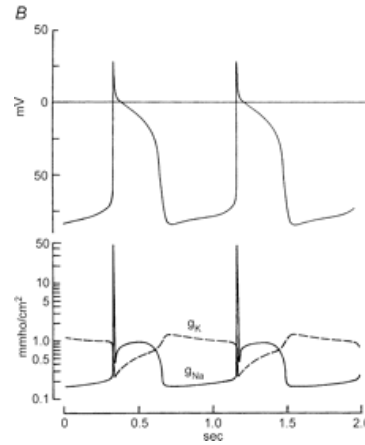
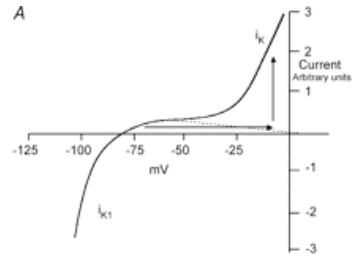
"So my research day started at 1:30 a.m.; a quick coffee, and then two hours at the *Mercury* computer. Then on to the slaughterhouse at 5 a.m. to pick up the sheep hearts with which the day's experiments would be done. Those experiments sometimes lasted until the time came to return to programming *Mercury*. I think that experience completely wrecked my circadian rhythms, but let's return to that kind of rhythm later in this chapter."

Denis Noble. *The Music of Life: Biology beyond the Genome*. (Oxford University Press, USA, 2006). Page 61.

- Modeled pacemaker activity without explicit oscillator
- Physiologically incorrect:
 - Developed before voltage clamp of cardiac cells
 - Prolonged AP produced by Na current rather than Ca current, which was missing!



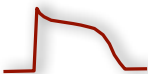
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McAllister, Noble, and Tsien (1975)

- Extension of Noble's 1962
- Included Ca^{2+} , 9 gating parameters
- Inward currents
 - I_{Na} : typical HH current
 - I_{SI} : similar to I_{Na} but with slower kinetics
 - $I_{\text{Na,b}}$: background sodium current
- Outward currents
 - I_{K1} : background current
 - I_{K2} : pacemaker current, activation variable
 - I_{X1} : plateau current, activation variable
 - I_{X2} : plateau current, activation variable
 - I_{Cl} : time and voltage dependent Cl current



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Beeler-Reuter (1977)

$$i_m = i_{Na} + i_{Na,b} + i_{K1} + i_{X1} + i_S$$

$$i_{Na} = g_{\bar{Na}} m^3 h j (V_m - E_{Na})$$

Modified Na current
(h,j inactivation)

$$i_{Na,b} = g_{Na,b} (V_m - E_{Na})$$

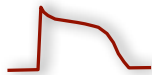
Background Na current
(time dependent)

$$i_{K1} = f(t, V_M)$$

Outward K current,
contributes to plateau

$$i_{X1} = x1 f(t, V_M)$$

Mostly K outward current,
slow to activate,
contributes to
repolarization



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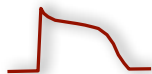
Beeler-Reuter (1977)

$$i_S = g_s df(V_m - E_s)$$

Inward, slow, largely Ca
current, d activation, f
inactivation.

$$\frac{\partial [Ca^{2+}]_i}{\partial t} = -10^{-7} i_s + 0.07(10^{-7} - [Ca^{2+}]_i)$$

Ca concentration changes with
 i_s current!

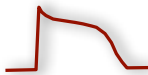
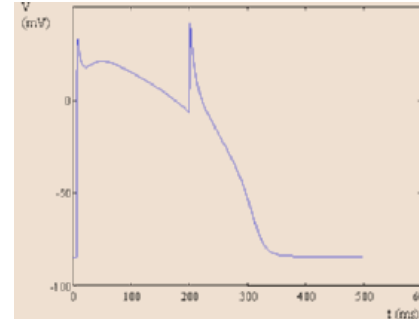
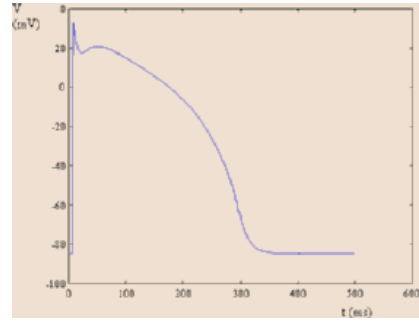


Whole Cell Simulation

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Beeler-Reuter (1977)

- Online version:
thevirtualheart.org/java/BR.html
- Simulated rate dependence of AP duration
- Included refractory period
- Supported premature stimulus (see example)
- Implicit Ca current, similar to Na, activation and inactivation variables



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Luo-Rudy I (1991)

$$i_m = i_{Na} + i_{si} + i_K + i_{K1} + i_{Kp} + i_b$$

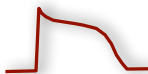
i_{si} Slow inward current, Ca

i_K Time dependent K current

i_{K1} Time independent K current

i_{Kp} Plateau K current

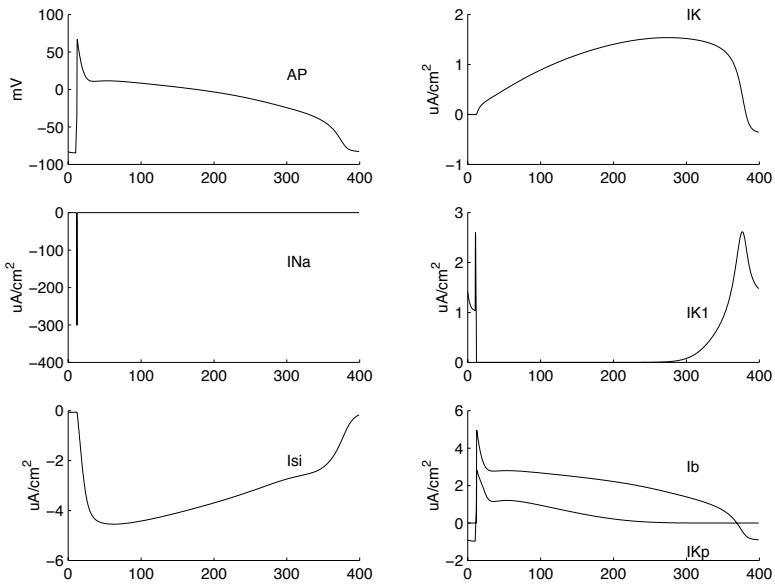
i_b Background K current



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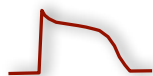
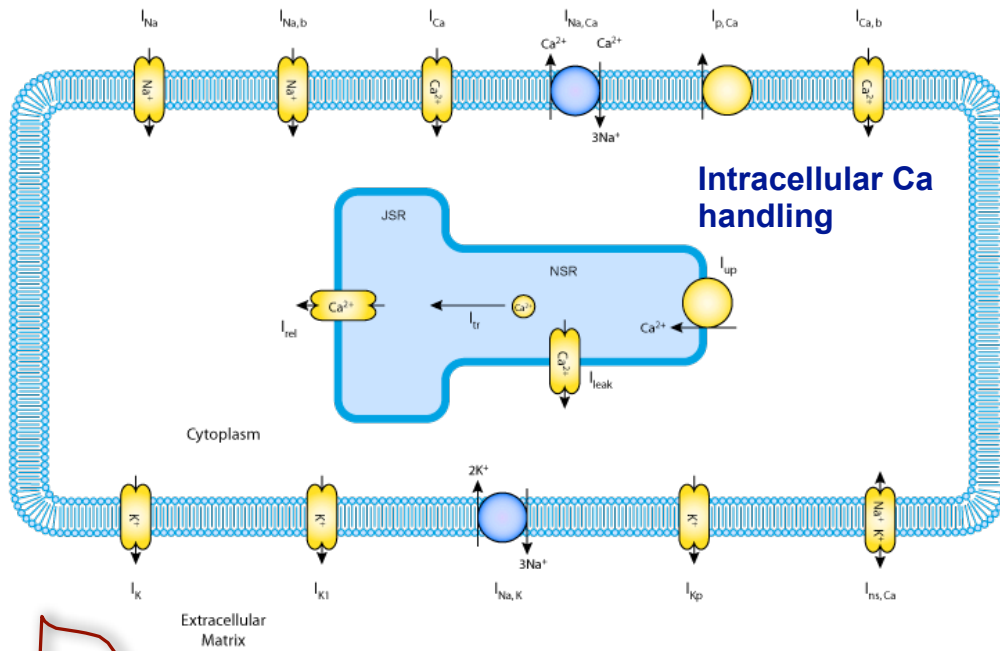
Luo-Rudy I (1991)



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Luo-Rudy II (1994)



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Multiscale Simulation

Perspectives on: Molecular dynamics and computational methods

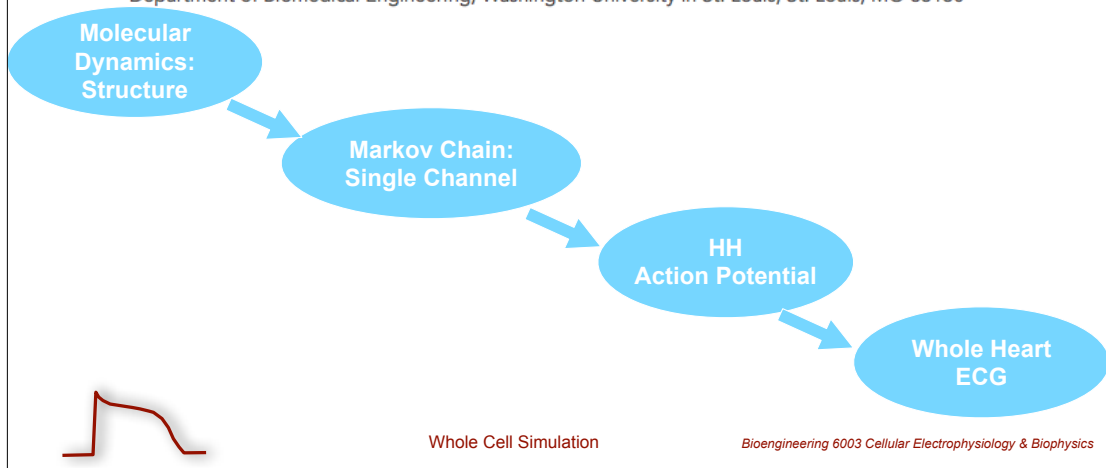
Multi-scale electrophysiology modeling: from atom to organ

Jonathan R. Silva¹ and Yoram Rudy²

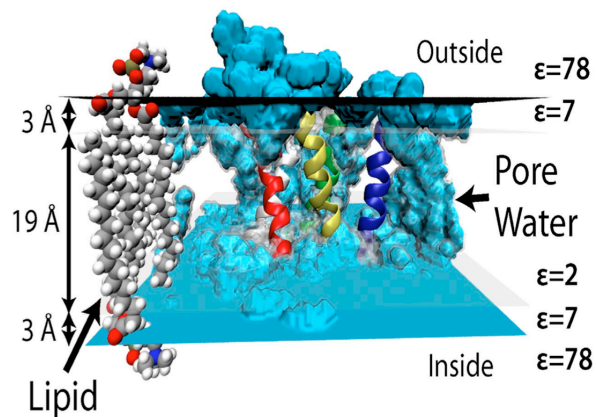
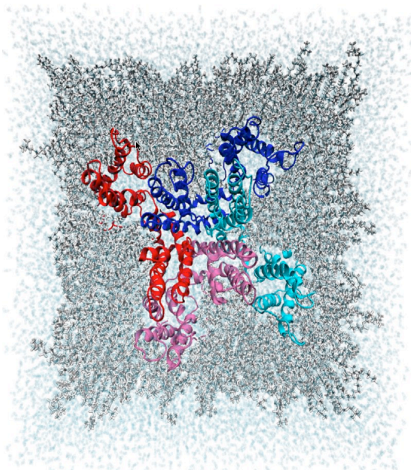
J Gen Physiol. 135(6): 575-581

¹Department of Pediatrics, University of Chicago, Chicago, IL 60637

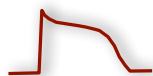
²Department of Biomedical Engineering, Washington University in St. Louis, St. Louis, MO 63130



Multiscale: Molecular Dynamics



- Purpose
 - Channel shape, conformational changes, interaction with environment
- Challenges
 - Parameter choices, computational, validation

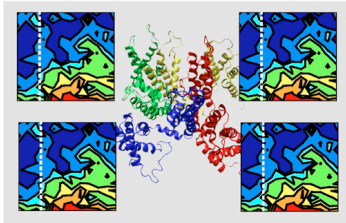


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Multiscale: Markov Models

A

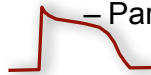


Transition takes place when all 4 voltage sensors in permissive state

$$\rho \uparrow \sigma$$

Flickery Blocked State

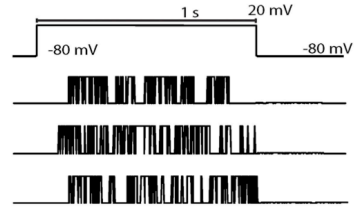
- Purpose
 - Capture conformational states, voltage and time dependence, single channel behavior (normal and after mutations)
- Challenges
 - Parameter choices, computational, validation



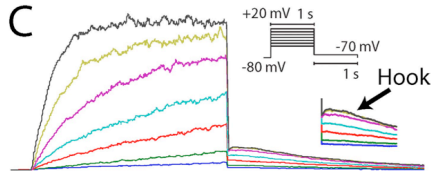
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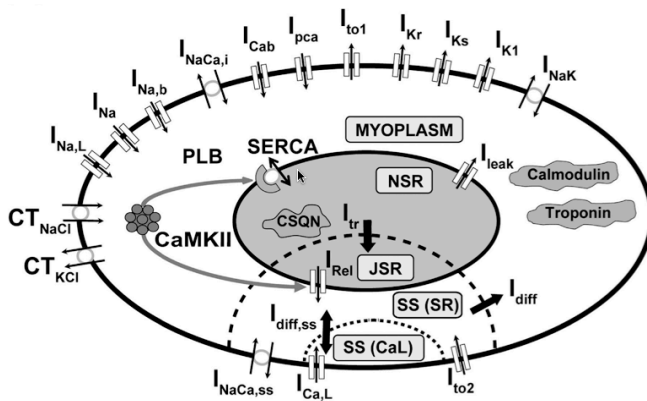
B



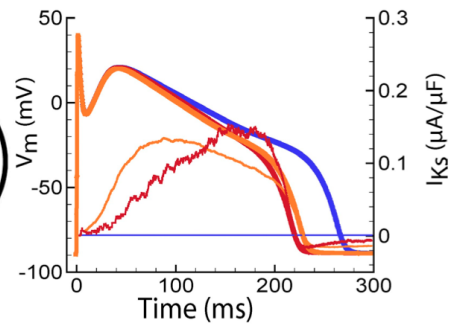
C



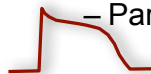
Multiscale: Action Potentials



AP and Currents



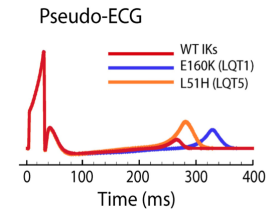
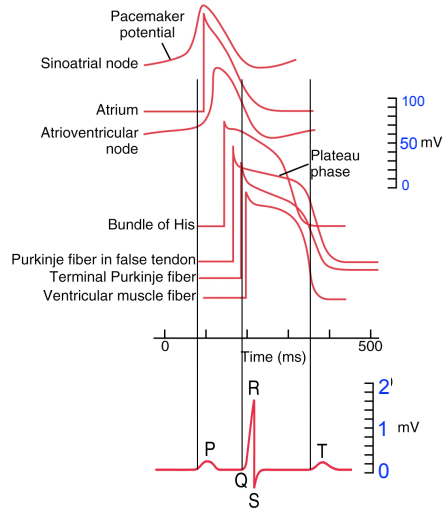
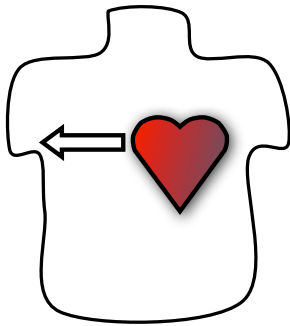
- Purpose
 - Whole cell behavior, EP and ECC, ionic currents, effect of environment, multiple beat behavior
- Challenges
 - Parameter choices, computational, geometry, diffusion



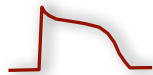
Whole Cell Simulation

Bioengineering 6003 Cellular Electrophysiology & Biophysics

Multiscale: ECG



- Purpose
 - Whole heart behavior, link to clinic, integration
- Challenges
 - Parameter choices, geometry, computational



Whole Cell Simulation

Bioengineering 6003 Cellular Electrophysiology & Biophysics