Mathematical Modeling of Action Potential with Transmission Equations and Hodgkin-Huxley Model

BENG 221 Problem Solving Report

Introduction

Action potential, a process during which the electrical membrane potential rapidly rises and falls with a distinctive pattern, is almost a universal process in all organisms. Action potential exists in neurons, muscle cells, and other types of endocrine cells. In neurons, propagation of action potential enables the communication between neurons, leading to cognitive functions. In other types of cells, action potential triggers cascades of intracellular processes. For instance, in muscle cells, the propagation of action potential triggers the release of calcium and further results in muscle contraction. Action potential is generated by voltage-gated ion channels that respond to membrane potentials. When the incoming membrane potential is above a certain threshold, these ion channels open. The electrochemical gradient then drives sodium and potassium ions across membrane. Rushing in of the sodium ions is responsible for the depolarization phase (increase in membrane potential), while outward movement of potassium ions, which lags behind the movement of sodium ions, results in repolarization and hyperpolarization phases. The sodium/potassium ion transporters then actively transport these ions against their gradients to restore the original electrochemical gradients.

Propagation of action potential is such an essential process in all organisms as they are responsible for cellular processes in multiple organs. The impairment of conduction of action potential can lead to diseases such as multiple sclerosis or even sudden death. Here we try to use transmission equations and Hodgkin-Huxley model to mathematically model the propagation of action potentials as a function of time and distance. As the model depends on a wide array of parameters, such as the input voltage, membrane capacitance, resistance, etc., with such a model established, one can simply modify the parameters, such as these affected by a certain disease, and examine the effects on propagation of action potential. Also, knowing the spatial and temporal distribution of action potential of a specific can let us reversely model the parameters and shed light on the physiological origins of certain diseases.

Set-up

An unmyelinated axon with radius a would be modeled. Current is allowed to leak back and forth across a cylindrical membrane, every point in

the membrane, to the interstitial fluid through capacitive and ion-transport mechanisms. Modeling of membrane as a capacitor is reasonable because it is thin so that the accumulation of charged particles on one side will pull the oppositely charged particles to the other side of the membrane. Interstitial fluid is treated as a shunt, so it does not have resistance. The differential equations that will be derived are also



Figure 1 Excitatory postsynaptic potential and an action potential.

adapted to muscle fibers as muscle fibers are not myelinated. In our model, a voltage v is implemented at x=0 as an initial -15 mV sawtooth impulse returning to V=0 after 3 ms (more

detailed initial and boundary conditions would be described in the next section). A sawtooth impulse is chosen to mimic the shape of an excitatory postsynaptic potential (EPSP), depicted in figure 1. In addition, as current passes through the axon, it generates self-inductance. In summary, the axon model can be described as the circuit diagram, which is also the circuit for telegrapher's equations, in figure 2.



Figure 2 Circuit diagram for axon model and telegrapher's equations.

The differential equations that are derived from the circuit are described as following:

$$-\frac{\partial i_a}{\partial x} = i + \pi a^2 C_a \frac{\partial v}{\partial t} \tag{1}$$

$$-\frac{\partial v}{\partial x} = ri_a + \frac{L}{\pi a^2} \frac{\partial i_a}{\partial t}$$
(2)

In which v is potential difference across membrane, i is membrane current per unit length, I is membrane current density, i_a is axon current per unit length, r is resistance per unit length of axon material, R is specific resistance of axon, L is axon specific self-inductance, and C_a is axon self-capacitance per unit area per unit length.

Solution

Figure 3 is the circuit diagram of Hodgkin-Huxley model. The lipid bilayer is represented as a capacitance (*Cm*). Voltage-gated and leak ion channels are represented by nonlinear (g_n) and linear (g_L) conductances, respectively. The electrochemical gradients driving the flow of ions are represented by batteries (E), and ion pumps and exchangers are represented by current sources (I_p).



Figure 3 Circuit diagram for Hodgkin-Huxley model

(Reference: http://en.wikipedia.org/wiki/File:Hodgkin-Huxley.jpg)

The time derivative of the potential across the membrane is proportional to the sum of the currents in the circuit. This is represented as follows,

$$\frac{\mathrm{dVm}}{\mathrm{dt}} = -\frac{1}{\mathrm{Cm}} \left(\sum_{i} \mathrm{I}i \right)$$

The Hodgkin-Huxley expression for I can be separated into four parallel components, the capacitive current (I_c) , ion currents of potassium and sodium $(I_k \text{ and } I_{Na})$, and a smaller current (I_l) made up of chloride and other ions.

$$\mathbf{I} = \mathbf{I}_{c} + \mathbf{I}_{k} + \mathbf{I}_{Na} + \mathbf{I}_{l}$$

The parameters used in the Hodgkin-Huxley equation are as follows, the specific resistances corresponding to the component ion currents can be denoted by R_k , R_{Na} , and R_l .

$$g_k = \frac{1}{R_k}$$
, $g_{Na} = \frac{1}{R_{Na}}$, $g_l = \frac{1}{R_l}$

From here on the $\frac{\partial 0}{\partial t}$ will be written as ()_t. n, *m*, and *h* are quantities of empirical convenience. They can also be thought of as the probabilities of given ion in a specific location.

$$n_{t} = \alpha_{n}(1 - n) - \beta_{n}n$$

$$m_{t} = \alpha_{m}(1 - m) - \beta_{m}m$$

$$h_{t} = \alpha_{h}(1 - h) - \beta_{h}h$$
(3)

Where

$$\alpha_n(v) = \frac{0.01(v+10)}{\left\{-1 + exp\left[\frac{v+10}{10}\right]\right\}}$$
$$\beta_n(v) = 0.125 exp\left(\frac{v}{80}\right)$$
$$\alpha_m(v) = \frac{0.1(v+25)}{\left\{-1 + exp\left[\frac{v+25}{10}\right]\right\}}$$

$$\beta_m(v) = 4 \exp\left(\frac{v}{18}\right)$$
$$\alpha_h(v) = 0.07 \exp\left(\frac{v}{20}\right)$$
$$\beta_h(v) = \left(1 + \exp\frac{v + 30}{10}\right)^{-1}$$

Also, let the Vk, VNa, and Vt denote the equilibrium potential of the corresponding ions and CM the membrane capacitance per unit area.

The full Hodgkin-Huxley excitation equation is

$$I = C_M v_t + \bar{g}_K n^4 (v - v_K) + \bar{g}_{N_a} m^3 h (v - v_{N_a}) + \bar{g}_l (v - v_l)$$
(4)

Now we apply $\left(\gamma + \frac{\partial (L/\pi a^2)}{\partial t}\right)$ to equation (1), $-\left(r + \frac{L}{\pi a^2} \frac{\partial}{\partial t}\right) \frac{\partial i_a}{\partial x} = \left(r + \frac{L}{\pi a^2} \frac{\partial}{\partial t}\right) (i + \pi a^2 C_a) \frac{\partial v}{\partial t},$ Or $-\frac{\partial}{\partial x} \left(ri_a + \frac{L}{\pi a^2} \frac{\partial i_a}{\partial t}\right) = ri + \frac{L}{\pi a^2} \frac{\partial}{\partial t} + \pi a^2 r C_a \frac{\partial v}{\partial t} + L C_a \frac{\partial^2 v}{\partial t^2}.$

While using the left term of equation (2), we can get

$$\frac{\partial^2 v}{\partial x^2} - LC_a \frac{\partial^2 v}{\partial t^2} = (\pi a^2) rC_a \frac{\partial v}{\partial t} + ri + \frac{L}{\pi a^2} \frac{\partial i}{\partial t}$$

Since I = $\frac{i}{2\pi a}$ and $r = \frac{R}{\pi a^2}$,

The transmission equation can be written as

$$\frac{\partial^2 v}{\partial x^2} - LC_a \frac{\partial^2 v}{\partial t^2} = RC_a \frac{\partial v}{\partial t} + \frac{2}{a}RI + \frac{2}{a}L\frac{\partial I}{\partial t}$$
(5)

Replacing the current by the value we got in equation (4), the following equation for membrane voltage can be obtained, which combines both the processes of transmission down the axoplasm and excitation across the membrane.

$$\begin{split} v_{xx} &- \frac{2}{a} L \left(\frac{a}{2} C_a + C_M \right) v_{tt} \\ &= \frac{2}{a} R \left(\frac{a}{2} C_a + C_M \right) v_t + \frac{2}{a} L \left(\bar{g}_K m^4 + \bar{g}_{N_a} m^3 h + \bar{g}_l \right) v_t \\ &+ \bar{g}_K \left(\frac{2}{a} R n^4 + 4 \frac{2L}{a} n^3 n_t \right) (v - v_K) \\ &+ \bar{g}_{N_a} \left[\frac{2}{a} R m^3 h + \frac{2}{a} L (3 m^2 h m_t + m^3 h_t) \right] (v - v_{N_a}) + \bar{g}_l \frac{2}{a} R (v - v_l). \end{split}$$

The *C* can be replaced by C_M without significant loss since the $(\frac{a}{2}C_a)$ is pretty small compared to C_M .

$$C = \frac{a}{2}C_a + C_M$$

Let

$$\theta = \left(\frac{a}{2LC}\right)^{\frac{1}{2}}, \tau = \theta t$$

and thus ()_t = ()_{τ} · τ_t = θ · ()_{τ}

The final partial differential equation is as follow,

$$\begin{aligned} v_{xx} - v_{\tau\tau} &= \frac{2}{a} R C \theta v_{\tau} + \frac{1}{\theta C} \left(\bar{g}_{K} n^{4} + \bar{g}_{N_{a}} m^{3} h + \bar{g}_{1} \right) v_{\tau} + \bar{g}_{K} \left(\frac{2}{a} R n^{4} + \frac{4}{\theta C} n^{3} n_{\tau} \right) (v - v_{K}) \\ &+ \bar{g}_{N_{a}} \left[\frac{2}{a} R m^{3} h + \frac{1}{\theta C} (3m^{2} h m_{\tau} + n^{3} h_{\tau}) \right] (v - v_{N_{a}}) + \bar{g}_{l} \frac{2}{a} R (v - v_{l}). \end{aligned}$$

Numerical Simulation

The analytical equation derived above is not one that can be easily solved. To simplify, the equations above can be rewritten as a system of first order equations and simulated using a finite difference method. This is accomplished by defining variables ψ and φ where $\psi = v_{\tau}$ and $\varphi = v_{x}$. Thus the final differential equation we obtained can be rewritten as,

$$\varphi_{x} - \psi_{\tau} = \frac{2}{a}RC\theta\psi + \frac{1}{\theta C}\left(\bar{g}_{K}n^{4} + \bar{g}_{N_{a}}m^{3}h + \bar{g}_{1}\right)\psi + \bar{g}_{K}\left(\frac{2}{a}Rn^{4} + \frac{4}{\theta C}n^{3}n_{\tau}\right)(v - v_{K}) + \bar{g}_{N_{a}}\left[\frac{2}{a}Rm^{3}h + \frac{1}{\theta C}\left(3m^{2}hm_{\tau} + n^{3}h_{\tau}\right)\right]\left(v - v_{N_{a}}\right) + \bar{g}_{l}\frac{2}{a}R(v - v_{l}) = F(\psi, v, n, m, k)$$

Using a finite difference approach, the above equation and the hodgkin huxley parameters can be written as,

$$\begin{split} n(i,j+1) &= n(i,j) + dt * N(n(i,j), v(i,j)) \\ m(i,j+1) &= m(i,j) + dt * M(m(i,j), v(i,j)) \\ h(i,j+1) &= h(i,j) + dt * H(h(i,j), v(i,j)) \\ v(i,j+1) &= v(i,j) + dt * \psi(i,j) \\ \varphi(i+1,j) &= \varphi(i,j) + s(\psi(i+1,j) - \psi(i,j)) \\ \psi(i,j+1) &= \psi(i,j) + s(\varphi(i+1,j) - \varphi(i,j)) - dt * F(\psi(i,j), v(i,j), n(i,j), m(i,j), h(i,j)) \end{split}$$

By iterating the variables i and j, the full space and time dependent profile of voltage can be obtained.





The first graph shows the action potential over time across different x points which is distance away from the input wave point. This shows how the action potential is initiated by the square wave of -15mV and propagates along the axon (x axis) with the same peak. This makes sense because action potential is all or nothing event unlike graded potentials. The undershoot caused by high transient potassium permeability is also apparent in the graph.





The second graph shows how the action potential is not achieved when an input wave of much lower amplitude is introduced. This is important because it verifies the property of action potential that high enough difference in membrane potential has to be introduced to trigger an action potential. The input signal just dies out over time.

Conclusion

Our project involved using Hodgkin Huxley's current law to derive a wave equation and simulating the membrane potential in x and t domain using Eulers finite difference method. From the analytical solution, the behavior of an action potential was understood as a wave equation.

From the numerical simulation with high enough input wave to induce an action potential, the well-known property of an action potential was verified: An action potential is all or nothing event that is only triggered above the certain threshold. Understanding the propagation of action potentials is very important because it is responsible for not only communication between the cells but also many cellular processes in multiple organs. The solutions match the current knowledge of action potential propagation in time and space domain. And with modifications, they could be used to model the propagation of action potential when there is a change in certain parameter or in intensity of an input wave.

Future Work

Our model could be improved by assuming myelinated axon in which the majority of the axon is myelinated which, in circuit, means high membrane resistance and low capacitance. This would result in faster propagation of action potential across the axon. Using the myelination, multiple sclerosis could be modeled and its behavior could be compared to normal myelinated axon as well. Instead of a large change like myelination, simple modifications could be introduced to the model when, for example, certain ion channels are damaged or ion permeability across a membrane are altered, resulting in parameter changes in the model. The numerical approximation method could be improved as well. In this study, Eulers finite element difference method was used but other numerical approximation could be used such as Gram–Schmidt process.

References

H. M. Lieberstein, On the Hodgkin-Huxley partial differential equation, Mathematical Biosciences, 1967; 1:45-69

Stephen Waxman, "Ion channels and Neuronal Dysfunction in Multiple Sclerosis", *Arch Neurol.* 2002; 59:1377-1380

Christof Koch, "Methods in Neuronal Modeling: From synapses to networks", 1988; ISBN 0-262-61071-X

MATLAB SIMULATION CODE

```
project.m
clc;
close all;
clear all;
dt=0.0001;
dx=1;
s=dt/dx;
t=0:dt:10;
x=0:dx:10;
alpha n0=0.01*(0+10)/(-1+exp((0+10)/10));
beta n0=0.125 \exp(0/80);
alpha m0=0.1*(0+25)/(-1+\exp((0+25)/10));
beta m0=4 \exp(0/18);
alpha h0=0.07 \exp(0/20);
beta h0=(1+exp((0+30)/10))^{-1};
n=zeros(length(x),length(t));
m=zeros(length(x),length(t));
h=zeros(length(x),length(t));
v=zeros(length(x),length(t));
psi=zeros(length(x),length(t));
phi=zeros(length(x),length(t));
n(:,1)=alpha n0/(alpha n0+beta n0);
m(:,1)=alpha m0/(alpha m0+beta m0);
h(:,1)=alpha h0/(alpha h0+beta h0);
%-15mV impulse
for i=1:1:30000
    v(1,i) = -15;
end
%below threshold
%for i=1:1:30000
% v(1,i)=−5;
%end
for i=1:1:length(x)-1
    phi(i,1)=1.5;
    v(i+1,1)=v(i,1)+dx*phi(i,1);
end
for j=1:1:length(t)-1
    %indicator for completion (in percent)
    j/(10/dt)*100
    for i=1:1:length(x)-1
        n(i,j+1)=n(i,j) + dt*N(n(i,j),v(i,j));
        m(i,j+1) = m(i,j) + dt M(m(i,j), v(i,j));
        h(i,j+1)=h(i,j) + dt + H(h(i,j),v(i,j));
        v(i,j+1)=v(i,j) + dt*psi(i,j);
```

```
phi(i+1,j)=phi(i,j) + s*(psi(i+1,j)-psi(i,j));
    psi(i,j+1)=psi(i,j) + s*(phi(i+1,j)-phi(i,j)) -
dt*F(psi(i,j),v(i,j),n(i,j),m(i,j),h(i,j));
    end
end
figure(1)
plot(t,v(1,:))
figure(2)
plot(t,v(1,:),t,v(2,:),t,v(3,:),t,v(4,:),t,v(5,:),t,v(6,:),t,v(7,:),t,v(8,:),
t,v(9,:),t,v(10,:))
```

```
legend('1','2','3','4','5','6','7','8','9','10')
```

F.m

```
function f=F(psi,v,n,m,h)
R2 a=2974.8991;
theta=1.23138148;
C=0.001;
g k=0.036;
g n=0.12;
g l=0.0003;
v k=12;
v n=-115;
v l=-10.5989;
N f=N(n,v);
M f=M(m,v);
H f=H(h,v);
f=R2 a*C*theta*psi + (1/(theta*C))*(g k*n^4 + g n*m^3*h + g l)*psi +
g k*(R2 a*n^4 +(4/(theta*C))*n^3*N f)*(v-v k) + g n*(R2 a*m<sup>3</sup>*h +
(1/(theta*C))*(3*m^2*h*M f + m^3*H f))*(v-v n) + g l*R2 a*(v-v l);
```

```
end
```

H.m

```
function [H_prime] = H(h,v)
theta=1.23138148;
```

```
alpha_h=0.07*exp(v/20);
beta_h=(1+exp((v+30)/10))^-1;
H_prime=(1/theta)*(alpha_h*(1-h)-beta_h*h);
end
```

M.m

function [M_prime] = M(m,v)
theta=1.23138148;

 $alpha_m=0.1*(v+25)/(-1+exp((v+25)/10));$

```
beta_m=4*exp(v/18);
M_prime=(1/theta)*(alpha_m*(1-m)-beta_m*m);
end
```

N.m

```
function [N_prime] = N(n,v)
theta=1.23138148;
```

```
alpha_n=0.01*(v+10)/(-1+exp((v+10)/10));
beta_n=0.125*exp(v/80);
N_prime=(1/theta)*(alpha_n*(1-n)-beta_n*n);
end
```