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Lab 3&4 Introduction:

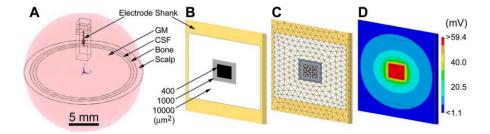
In this report, we delve into the realm of electrostatic models, focusing on their pivotal role in deep brain stimulation (DBS) research. By drawing upon historical experiment data, such as resistivity, conductivity, and neuronal structure, we aim to elucidate the intricate relationship between externally applied voltages and the internal currents necessary for neuronal activation. This exploration is further enriched by our construction of deep brain simulation models within the COMSOL framework, alongside the simulation of electrode data both in the presence and absence of noise. A significant portion of our analysis is dedicated to understanding the K matrix, which represents the voltage distribution along a neuron's axis, and its correlation with external voltages generated by internal currents. Through our comprehensive study, we strive to pinpoint the minimal internal current required for neuronal activation via external stimulation and to map the effective distance range for such activation, thereby offering novel insights into the optimization of DBS strategies.

Methods:

Lab-3:

Moffit & McIntyre (2005), assumed that the tissue surrounding the electrode behaves as a purely resistive and linear medium for electric currents, following Ohm's law. The models are static and do not account for the dynamic changes in tissue properties or electrode-tissue interfaces over time. Despite using high-impedance measurement equipment, the model assumes the electrode-electrolyte interfacial impedance's effect on the recording or stimulation is negligible.

To model the electrical properties of the head, concentric spheres are used, which simplifies the geometric and compositional complexities of the head and brain with current source and sinks in the brain and skull as the ground.



The rat head was modeled as 4 concentric spheres representing the brain, cerebrospinal fluid (CSF), skull, and scalp with radii of 8000, 8500, 9000, and 10,000 mm, respectively. The figures were 10x for the human brain. As we can observe the aim is to create an electric mesh.

Lab-4:

Warman and Durand (1992), made certain assumptions for predicting axonal excitation due to electric fields simplifying biological reality to make computations feasible. Key simplifications include treating myelin as a perfect insulator, assuming a homogeneous isotropic extracellular medium, and using linear cable models for subthreshold behavior. The paper also leverages the response of a passive cable to predict excitation. The axon is modeled as an electrical network with distinct nodes and internodal spacing, with myelin assumed to be a perfect insulator. Axonal excitation is determined by the extracellular potentials, which are themselves assumed not to be significantly affected by the fiber response. The use of linear cable models for subthreshold behavior and the approximation of membrane activation based on induced polarization exceeding a certain transmembrane voltage.

We calculated the effect the internal currents of a neuron from all its compartments, at a certain distance from the electrode, will have on the external potential at the electrode. When obtaining the signal all the neurons in the vicinity of the electrode will have contributions to the resultant signal. We can decide the firing rate for each signal from individual neurons and combine. After adding noise to the resultant recording, the simulation of the neural signal looks as good as the original. It's better to discard the signal from neurons which are in 25 um vicinity from the neuron, as it's a dead zone.

We calculated the external voltage from internal currents and K matrix from COMSOL, by interpolating the voltages at the coordinates of our present neuron.

We calculated the internal currents of the neuron from the stimulus current and electrode distance from the neuron by using the predefined equations from Warman & Durand (1992), it loops over all the neuron compartments multiplying the conductivity and the second difference of voltage at that compartment.

In the neural recording model, various elements were altered to examine their effects on recordings. These modifications included adding multiple neurons to simulate a more complex neural network, adjusting cell sizes to reflect differences between large and small neurons, incorporating noise to mimic real-world recording conditions, and utilizing a K matrix for modeling variable conductivity environments. Each change aimed to enhance the model's realism and investigate how such factors influence neural signal recordings, providing insights into the complexities of electrophysiological recordings and deep brain stimulation.

Results:

Lab-3:

The decreasing trend can be observed in figure 1, in the voltage magnitude as the distance from the point source increases. The voltage drops sharply close to the current source, reflecting the strong electric field near the point of injection. The rate of decrease in voltage should become less steep as the distance increases, indicating the diffusive spread of the electric field through the head model's conductive media. In the figure 2, we can observe the model as a mesh of electric field.

In the figure 3, we can observe a peak at (0,0,0) as there's a current source of 5mA. We can also observe two dips because of current sinks at +3 mm and -3mm in y direction, the superior point with 1.25 mA current sink

and the inferior point a 3.75 mA current sink. In figure 4, we can observe the presence of two current sinks in the bright regions. In figure 5, we can observe the points of current source and sinks.

In the figure 6, we can observe the second spatial derivative of the voltage, it portrays how the curvature of the voltage distribution changes as you move away from the active DBS contact, perpendicular to the probe.

Lab-4:

Figure 7 & 8 consists of external voltage plots at the electrode at distances of 50, 100, 200, 300 um from the neuron, with internal currents as Big currents and small currents respectively.

Figure 9 consists of a visualization of multiple neurons passing through the cube making sure that their axon hillocks are inside the cube. With the electrode at the origin of the cube (0,0,0).

Figure 10 consists of the combined signals from all neuronal voltage contributions with and without the dropped neurons from the dead zone, which is 25um from the electrode."Dead zone" refers to an area immediately surrounding the electrode where neuronal signals are not effectively recorded or stimulated. This can be due to tissue damage from electrode insertion, changes in tissue conductivity, or the physical separation between neurons and electrode surfaces. The dead zone can significantly affect the efficacy of neural interfaces by limiting the spatial resolution of recordings and reducing the precision of targeted stimulation. Recognizing and accounting for the dead zone is crucial in designing and interpreting neural interface experiments.

Figure 11 consists of the combined signals from all neuronal voltage contributions with added noise, which makes the simulated signal more realistic to the actual electrode recordings.

Figure 12 consists of the modeled voltage waveform from the interpolated voltage matrices from COMSOLS and treating them as K matrix and along with internal currents, finding the external voltage at the electrode.

Minimum current for action potential in an axon that is 20 um in diameter and 5mm in length is 40pA. (NEURON MODEL)

Minimum external current from from 1mm distance to activate the axon is >1A

I(internal)= ga*second difference of voltage at the particular node.

I(n) = ga*[Ve(n-1)-2Ve(n)+Ve(n+1)]

Increasing model complexity in neural recording and deep brain stimulation simulations can significantly enhance the accuracy and realism of the predictions. More complex models can account for the heterogeneous and anisotropic nature of neural tissue, interactions between multiple neurons, and the impact of various noise sources on signal fidelity. This leads to better understanding and prediction of how neural interfaces interact with biological tissue, allowing for the optimization of electrode designs and stimulation protocols. However, increased complexity also demands more computational resources and sophisticated modeling techniques, potentially making simulations more challenging to execute and interpret.

Discussions:

Lab-3:

The curvature of the second spatial derivative of voltage provides insight into how sharply or smoothly the voltage changes in space, which is crucial for understanding the spatial specificity of DBS effects: High Magnitude of Derivative: Sharp changes in voltage suggest that DBS effects are highly localized around the active contact. High peaks in the second derivative indicate regions with rapid changes in the electric field, potentially corresponding to areas of strong neural activation or inhibition. Low Magnitude of Derivative: Smoother curves or smaller magnitudes suggest a more gradual spread of the DBS effect, indicating that the stimulation might affect a broader area with less specificity.

The electric field generated by DBS electrodes and to understand its spatial distribution within the brain allows for the optimization of electrode placement and stimulation parameters, ensuring maximum efficacy while minimizing side effects. This approach could lead to the development of patient-specific DBS therapies, where treatment parameters are tailored based on the individual's unique anatomical and physiological characteristics, thereby enhancing therapeutic outcomes.

Lab-4:

From Warman and Durand (1992), the equation to find the internal current essentially calculates the second spatial derivative of the external voltage along the axon, which corresponds to the activating function. This function plays a critical role in determining the sites of action potential initiation and how the external electric field influences the neuron's internal current. We can also determine the stimulus current needed from a certain distance, to generate the action potential in the cell.

Looking forward, the models developed in this lab can serve as a foundational tool for advancing neural interface technologies. By accurately simulating how neurons respond to external stimuli, these models can be used to design more effective neural prosthetics, optimize deep brain stimulation (DBS) parameters for therapeutic applications, and enhance the resolution of brain-machine interfaces (BMIs). Furthermore, the ability to model complex conductivity environments and account for noise in neural recordings can lead to the development of more sensitive and selective recording devices, enabling the detection of subtler neural signals amidst the noise. These advancements could significantly improve the quality of life for individuals with neurological disorders, offer new insights into brain function, and pave the way for novel neurotechnological applications.

Appendix:

Figures & Tables:

Lab 3:

Figure 1

Electric Potential vs. Distance from Stimulus

1400

1200

1200

0 0.01 0.02 0.03 0.04 0.05 0.06 0.07 0.08 0.09

Distance from stimulus (m)

Figure 2

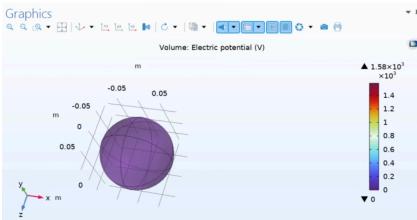


Figure 3

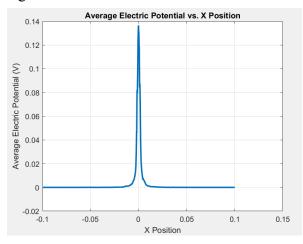


Figure 4

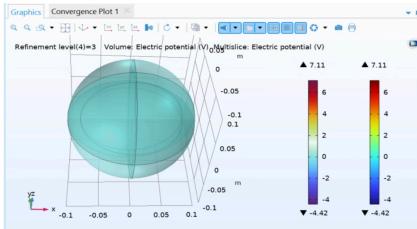


Figure 5

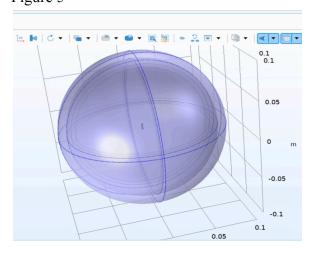
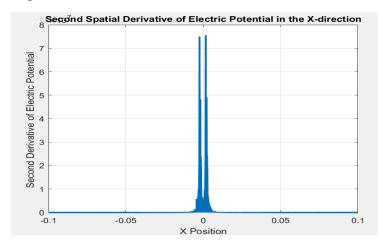
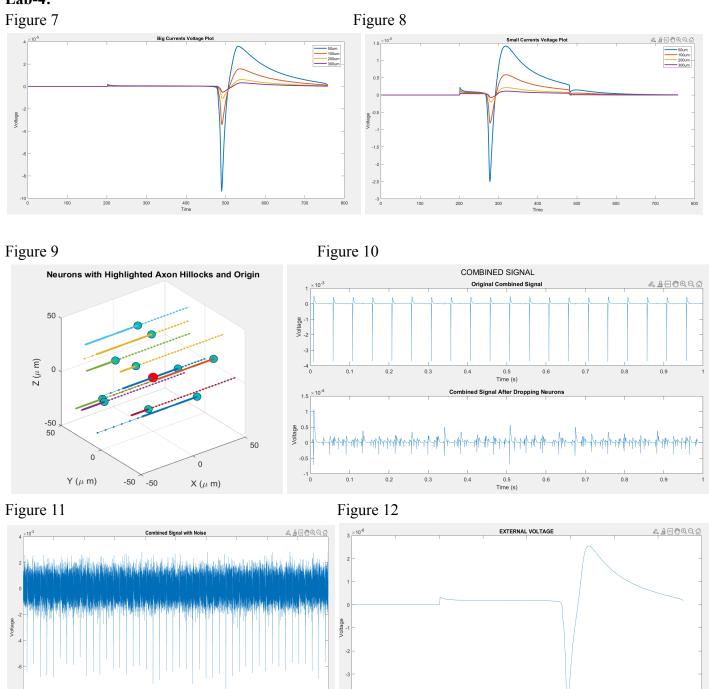


Figure 6



Lab-4:



```
Code:
```

Lab 3:

<u>Part 2:</u>

x grid(i) if exists

```
numHeaderLines = 9; % Adjust this number based on the actual number of header lines in your
file.
% Read the file into a table, skipping the header lines
data = readtable('multislice.txt', 'HeaderLines', numHeaderLines);
x pos=data.Var1;
y pos=data.Var2;
z pos=data.Var3;
electric potential = data.Var4;
% Calculate the distance from the stimulus for each data point
distances = sqrt(x pos.^2 + y pos.^2 + z pos.^2);
% Create a plot of electric potential vs. distance
figure;
plot(distances, electric potential, 'o');
xlabel('Distance from stimulus (m)');
ylabel('Electric potential (V)');
title('Electric Potential vs. Distance from Stimulus');
Part3:
% Specify the number of header lines in the file
numHeaderLines = 8;
% Read the file into a table, skipping the header lines
data = readtable('brain part3.txt', 'HeaderLines', numHeaderLines);
% Extract positions and electric potential
x pos = data.Var1;
y pos = data. Var2;
z pos = data.Var3;
electric potential = data. Var4;
% Calculate the distance from the stimulus for each data point
distances = sqrt(x_pos.^2 + y pos.^2 + z pos.^2);
% Plot electric potential vs. distance
figure;
plot(electric potential, distances, 'o', 'MarkerSize', 3);
ylabel('Distance from stimulus (units)');
xlabel('Electric potential (V)');
title('Electric Potential vs. Distance from Stimulus');
% Define a new grid for x positions
x \text{ grid} = \text{linspace}(\min(x \text{ pos}), \max(x \text{ pos}), 30000);
% Initialize the array for the calculated potentials
custom avg V = zeros(size(x grid));
for i = 1:length(x grid)
   % Find indices and distances for points on the negative and positive sides
   neg side indices = find(x pos < x grid(i));</pre>
   pos side indices = find(x pos >= x grid(i)); % Use >= to include the point exactly at
```

```
% Initialize averages for negative and positive sides
   avg neg = NaN; % Use NaN to handle cases where we don't have enough data points
   avg pos = NaN;
   % Calculate for negative side if there are enough points
   if length(neg side indices) >= 4
       neg distances = abs(x pos(neg side indices) - x grid(i));
       [~, neg indices] = mink(neg distances, 4);
       direct avg neg = mean(electric potential(neg side indices(neg indices(1:2))));
       weights neg = 1 ./ neg distances(neg indices(3:4));
       normalized weights neg = weights neg / sum(weights neg);
           weighted avg neg = sum(electric potential(neg side indices(neg indices(3:4))) .*
normalized weights neg);
       avg neg = mean([direct avg neg, weighted avg neg]);
   end
   % Calculate for positive side if there are enough points
   if length(pos side indices) >= 4
       pos distances = abs(x pos(pos side indices) - x grid(i));
       [~, pos indices] = mink(pos distances, 4);
       direct avg pos = mean(electric potential(pos side indices(pos indices(1:2))));
       weights pos = 1 ./ pos distances(pos indices(3:4));
       normalized weights pos = weights pos / sum(weights pos);
           weighted avg pos = sum(electric potential(pos side indices(pos indices(3:4))) .*
normalized weights pos);
       avg pos = mean([direct avg pos, weighted avg pos]);
   end
   % Combine the averages from both sides, ignoring NaN values
   custom avg V(i) = mean([avg neg, avg pos], 'omitnan');
end
% Plot the custom average electric potential vs. x grid
windowSize=400;
smoothV1 = smoothdata(custom avg V, 'movmean', windowSize);
plot(x grid, smoothV1, 'LineWidth', 2);
xlabel('X Position');
ylabel('Average Electric Potential (V)');
title('Average Electric Potential vs. X Position');
% Calculate the second derivative using finite differences on the custom averages
h = x \operatorname{grid}(2) - x \operatorname{grid}(1); % Assuming uniform spacing in x \operatorname{grid}
second derivative custom = diff(custom avg V, 2) / h^2;
% Adjust x grid to match the size of the second derivative custom array after diff operation
x grid adjusted for custom = x grid(2:end-1);
% Plot the second derivative of electric potential based on the custom averages
windowSize=600;
smoothV = smoothdata(second derivative custom, 'movmean', windowSize);
smoothV=abs(smoothV);
```

```
figure;
plot(x_grid_adjusted_for_custom, smoothV, '-', 'LineWidth', 2);
xlabel('X Position');
ylabel('Second Derivative of Electric Potential ');
title('Second Spatial Derivative of Electric Potential in the X-direction');
grid on;
```

Part 1:

```
%lest keep origin as the first value of XYZ.
%we need to obtain a point 50um in perpendicular direction to that..y axis.
origin=XYZ(200,:);
current origin=currents(200,:);
sigma=3.33*10^-7;%um 0.3333S/m
distance1=50; %um
radius= 20;%um
POI = [0.5, 50, 0];
%calculate Vext for that for origin
Vext 50 origin=current origin/(4*3.14*sigma*distance1);
%Calculate Vext for every point along the neuron
%voltages from everypoint is added
%sqrt(XYZ-POI)
distances = sqrt((XYZ(:,1) - POI(:,1)).^2 + (XYZ(:,2) - POI(:,2)).^2 + (XYZ(:,3) - POI(:,3).^2));
Vext 50 ALL=(currents./(4*3.14*sigma.*distances));
Vext 50 ALL=sum(Vext 50 ALL,1);
%plot(Vext 50 ALL(500,:))
%Now we need to repeat this process for points 100um, 200um, 300um from the axon hilloc: Vext
for all
%%100um
POI 100=[0.5,100,0];
%calculate Vext for that for origin
Vext 100 origin=current origin/(4*3.14*sigma*100);
%Calculate Vext for every point along the neuron
%sqrt(XYZ-POI)
distances100=sqrt((XYZ(:,1)-POI 100(:,1)).^2+(XYZ(:,2)-POI 100(:,2)).^2+(XYZ(:,3)-POI 100(:,
3).^2));
Vext 100 ALL=sum(currents./(4*3.14*sigma.*distances100),1);
Vext 100 ALL=sum(Vext 100 ALL,1);
%%200um
POI 200=[0.5,200,0];
%calculate Vext for that for origin
Vext 200 origin=current origin/(4*3.14*sigma*200);
%Calculate Vext for every point along the neuron
%sqrt(XYZ-POI)
distances200=sqrt((XYZ(:,1)-POI 200(:,1)).^2+(XYZ(:,2)-POI 200(:,2)).^2+(XYZ(:,3)-POI 200(:,
Vext 200 ALL=sum(currents./(4*3.14*sigma.*distances200),1);
Vext 200 ALL=sum(Vext 200 ALL,1);
%%300um
POI 300 = [0.5, 300, 0];
%calculate Vext for that for origin
Vext 300 origin=current origin/(4*3.14*sigma*300);
%Calculate Vext for every point along the neuron
%sqrt(XYZ-POI)
 \texttt{distances} \texttt{300} = \texttt{sqrt} ((\texttt{XYZ}(:,1) - \texttt{POI}\_\texttt{300}(:,1)) . ^2 + (\texttt{XYZ}(:,2) - \texttt{POI}\_\texttt{300}(:,2)) . ^2 + (\texttt{XYZ}(:,3) - \texttt{POI}\_\texttt{300}(:,3)) . ^2 + (\texttt{XYZ}(:,3) - \texttt{YZ}(:,3) - \texttt{YZ}(:,3)) . ^2 + (\texttt{YZ}(:,3) - \texttt{YZ}(:,3) - \texttt{YZ}(
3).^2);
```

```
Vext 300 ALL=sum(currents./(4*3.14*sigma.*distances300),1);
Vext 300 ALL=sum(Vext 300 ALL,1);
%distance at which the AP won't be detectable above a 10uV peakk to peak noise floor?
V noise floor = 10 ;%uV
% Function to compare voltage against noise floor
is detectable = @(voltage, V noise floor) voltage > V noise floor;
% Check detectability at each distance
detectable_at_50_um = is_detectable(Vext_50_ALL, V_noise_floor);
detectable at 100 um = is detectable (Vext 100 ALL, V noise floor);
detectable at 200 um = is detectable (Vext 200 ALL, V noise floor);
detectable at 300 um = is detectable (Vext 300 ALL, V noise floor);
%%Plotting together
figure;
plot(Vext 50 ALL, 'LineWidth', 2);
hold on;
plot(Vext 100 ALL, 'LineWidth', 2);
hold on;
plot(Vext 200 ALL, 'LineWidth', 2);
hold on;
plot(Vext 300 ALL, 'LineWidth', 2);
title('Small Currents Voltage Plot ')
legend('50um','100um','200um','300um')
xlabel('Time');
ylabel('Voltage')
hold off;
Part 2:
%ques1
%%distance of origin of cube from neuron 1 comp
origin=[0,0,0];
neuron1=[XYZ(:,1),XYZ(:,2),XYZ(:,3)];
num neurons=10;
hilloc pos=neuron1(200,:);
neurons = zeros(size(neuron1, 1), 3, num neurons);
neurons(:,:,1) = neuron1;
relative_positions = neuron1 - hilloc_pos;
% Cube boundaries for the axon hillocks
cube min = -50;
cube max = 50;
hillock positions = zeros(num neurons, 3); % 10 neurons, 3 coordinates (x, y, z)
hillock positions(1, :) = hilloc pos;
%%randon hilloc pos and other coordinates
for i = 2:num neurons
   % Random position for the axon hillock within the cube
   random hillock position = cube min + (cube max - cube min) .* rand(1, 3);
   hillock positions(i, :) = random hillock position;
   neurons(:,:,i) = relative positions + random hillock position;
end
% Plotting each neuron
```

```
figure; hold on; % Create a figure and hold it for multiple plots
% Plotting each neuron and highlighting axon hillocks
for i = 1:10 % Assuming 10 neurons
   neuron positions = neurons(:,:,i); % Extract the ith neuron's positions
   plot3(neuron positions(:,1), neuron positions(:,2), neuron positions(:,3), '.-'); % Plot
neuron compartments
   % Highlighting the axon hillock for this neuron
     scatter3(hillock positions(i,1), hillock positions(i,2), hillock positions(i,3), 100,
'filled', 'MarkerEdgeColor', 'k', 'MarkerFaceColor', [0 .75 .75]);
end
scatter3(0, 0, 0, 150, 'MarkerFaceColor', 'r');
xlabel('X (\mu m)');
ylabel('Y (\mu m)');
zlabel('Z (\mu m)');
title('Neurons with Highlighted Axon Hillocks and Origin');
axis equal; grid on; % Equal aspect ratio and grid on
xlim([-50 50]); ylim([-50 50]); zlim([-50 50]); %cube boundaries
view(3); % 3D view
hold off;
%%distances from origin of cube:
distances = zeros(10, 531);
neurons keep = false(num neurons, 1); % Initialize as logical array
for i = 1:10
   % Extracting the positions for the i-th neuron (531x3)
  neuron positions = neurons(:,:,i);
   % Calculating the distances for all compartments of the i-th neuron
   distances i = sqrt(sum((origin-neuron positions).^2, 2));
   distances(i,:) = distances i';
   if distances i > 25
      neurons keep(i) = true;
   end
end
%%%ques2
%RANDOM SPIKE TIMES
simulation duration = 1; % Duration in ms
dt=2.5*10^{-5}; %0.025ms=>s
spike duration ms = 2*10^-4; %0.18ms>s;
refractory period = 0.001; % 1 ms => s, example refractory period
neuron spike times = cell(num neurons, 1); % Cell array to store spike times for each neuron
for i = 1:num neurons
   % Randomly pick the number of spikes for the i-th neuron
   num spikes = randi([2, 50]);
   % Calculating the total duration that spikes occupy plus refractory periods
      total duration needed = (num spikes * spike duration ms) + ((num spikes - 1) *
refractory period);
   % Calculating the available time for intervals between spikes
   available time = simulation duration - total duration needed;
   % Adjust num spikes if not enough time
```

```
while available time < 0 && num spikes > 2
      num spikes = num spikes - 1;
          total duration needed = (num spikes * spike duration ms) + ((num spikes - 1) *
refractory period);
      available time = simulation duration - total duration needed;
   end
   % Calculating evenly distributed intervals between spikes
   interval between spikes = available time / (num spikes - 1) + refractory period;
   % Generating spike times starting with a random offset to stagger spikes between neurons
   random offset = rand() * refractory period; % Stagger start within one refractory period
   spike times = cumsum([random offset, repmat(spike duration ms + interval between spikes,
1, num spikes - 1)]);
  neuron spike times{i} = spike times;
end
%making sure none of the spike trails are overlapping
figure; hold on;
scatter(neuron spike times{1}, ones(size(neuron spike times{1})), 'filled', 'DisplayName',
'Neuron 1', 'SizeData', 10);
scatter(neuron spike times{2},
                               ones(size(neuron spike times{2})), 'filled', 'DisplayName',
'Neuron 2', 'SizeData', 10);
scatter(neuron spike times{3},
                               ones(size(neuron spike times{3})), 'filled', 'DisplayName',
'Neuron 3', 'SizeData', 10);
scatter(neuron_spike_times{4}, ones(size(neuron spike times{4})), 'filled', 'DisplayName',
'Neuron 4', 'SizeData', 10);
scatter(neuron spike times{5},
                               ones(size(neuron spike times{5})), 'filled', 'DisplayName',
'Neuron 5', 'SizeData', 10);
scatter(neuron spike times{6}, ones(size(neuron spike times{6})), 'filled', 'DisplayName',
'Neuron 6', 'SizeData', 10);
scatter(neuron spike times{7}, ones(size(neuron spike times{7})), 'filled', 'DisplayName',
'Neuron 7', 'SizeData', 10);
scatter(neuron spike times{8}, ones(size(neuron spike times{8})), 'filled', 'DisplayName',
'Neuron 8', 'SizeData', 10);
scatter(neuron spike times{9}, ones(size(neuron spike times{9})), 'filled', 'DisplayName',
'Neuron 9', 'SizeData', 10);
scatter(neuron spike times{10}, ones(size(neuron spike times{10})), 'filled', 'DisplayName',
'Neuron 10', 'SizeData', 10);
hold off;
xlabel('Time (s)');
ylabel('Spike');
title(sprintf('Spiking Pattern for 10 Neurons %d'));
xlim([0 1]);
%%dropped spikes times
neurons droppped=neurons;
neurons drop = find(~neurons keep);
neurons droppped(:,:,neurons drop) = [];
neuron dropped times=neuron spike times;
neuron dropped times(neurons drop) = [];
%%%ques3
```

```
sigma=3.33*10^-7;%um
                         0.3333S/m
Voltage neuron1=sum((currents./(4*3.14*sigma.*distances(1,:)')),1);
Voltage neuron2=sum((currents./(4*3.14*sigma.*distances(2,:)')),1);
Voltage neuron3=sum((currents./(4*3.14*sigma.*distances(3,:)')),1);
Voltage neuron4=sum((currents./(4*3.14*sigma.*distances(4,:)')),1);
Voltage neuron5=sum((currents./(4*3.14*sigma.*distances(5,:)')),1);
Voltage neuron6=sum((currents./(4*3.14*sigma.*distances(6,:)')),1);
Voltage neuron7=sum((currents./(4*3.14*sigma.*distances(7,:)')),1);
Voltage neuron8=sum((currents./(4*3.14*sigma.*distances(8,:)')),1);
Voltage neuron9=sum((currents./(4*3.14*sigma.*distances(9,:)')),1);
Voltage neuron10=sum((currents./(4*3.14*sigma.*distances(10,:)')),1);
voltage contribution=[Voltage neuron1; Voltage neuron2; Voltage neuron3; Voltage neuron4; Voltage
e neuron5; Voltage neuron6; Voltage neuron7; Voltage neuron8; Voltage neuron9; Voltage neuron10];
voltage contribution dropped=[Voltage neuron2; Voltage neuron4; Voltage neuron6; Voltage neuron
7; Voltage neuron8; Voltage neuron9; Voltage neuron10];
%%combined signal
simulation duration = 1; % Simulation duration in seconds
dt = 2.5e-5;
% Generate the time vector
time vector = 0:dt:simulation duration-dt;
combined signal = zeros(1, length(time vector));
for i = 1:10 % For each of the 10 neurons
   voltage trace = voltage contribution(i, :);
   for spike time = neuron spike times{i}
       spike index = round((spike time / dt)) + 1;
       % Determining the range of indices in combined signal where the voltage trace will be
added
       start idx = spike index;
       end idx = min(spike index + length(voltage trace) - 1, length(combined signal));
           % Adjust the length of the voltage trace if it extends beyond the end of
combined signal
       trace length = end idx - start idx + 1;
               combined signal(start idx:end idx) = combined signal(start idx:end idx)
voltage trace(1:trace length);
   end
end
%%combined signal with dropped
combined signal dropped = zeros(1, length(time vector));
for i = 1:7 % For each of the 10 neurons
   voltage trace = voltage contribution dropped(i, :);
   for spike time = neuron dropped times{i}
       spike index = round((spike time / dt)) + 1;
       % Determining the range of indices in combined signal where the voltage trace will be
added
       start idx = spike index;
                        end idx
                                       min(spike index
                                                              length(voltage trace)
                                                                                           1,
length(combined signal dropped));
           % Adjust the length of the voltage trace if it extends beyond the end of
combined signal
```

```
trace length = end idx - start idx + 1;
                                            combined signal dropped(start idx:end idx)
combined signal dropped(start idx:end idx) + voltage trace(1:trace length);
   end
end
%%%%specifically focusing on the reduction of larger spikes
%due to the exclusion of neurons within 25um of the electrode
subplot(2, 1, 1);
plot(time vector, combined signal);
title('Original Combined Signal');
xlabel('Time (s)');
ylabel('Voltage');
subplot(2, 1, 2);
plot(time vector, combined signal dropped);
title ('Combined Signal After Dropping Neurons');
xlabel('Time (s)');
ylabel('Voltage');
sgtitle('COMBINED SIGNAL');
%%ques4
%%adding noise to the combined signal
numSamples = round(simulation duration / dt);
recordedNoise = randn(1, numSamples);
rangeVext = max(combined signal) - min(combined signal);
scaledNoise = recordedNoise * (0.1 * rangeVext) / std(recordedNoise);
combined signal noise = combined signal + scaledNoise;
% Plot the combined signal with noise
plot(time vector, combined signal noise);
xlabel('Time (s)');
ylabel('Voltage');
title('Combined Signal with Noise');
Part 3
origin=[0,0,0];
neuron1=[XYZ(:,1),XYZ(:,2),XYZ(:,3)];
hilloc pos=neuron1(200,:);
nwpos=neuron1-[0,50,0];
%%Kmatrix with 1A current
numHeaderLines = 8; % Adjust this number based on the actual number of header lines in your
file.
% Read the file into a table, skipping the header lines
data = readtable('brain part3.txt', 'HeaderLines', numHeaderLines);
%%units: volatge from comsol is in V, xyz is in m
%%units from current big: voltage is in
                                         , xyz is in um
x pos=(data.Var1)*10^6;%um
y pos=(data.Var2)*10^6;%um
z pos=(data.Var3)*10^6;%um
electric potential = data. Var4;
%%we need to interpolate the x,y,z coordonates to match our neuron's
```

```
%%coordinate system to use V
%%we need V for our neuron's coordinates
% Interpolate K matrix data to neuron positions
% V interpolated = griddata(x pos, y pos, z pos, electric potential, neuron1(:,1),
neuron1(:,2), neuron1(:,3), 'cubic');
% V interpolated(isnan(V interpolated)) = 0; % Setting NaNs to 0
F = scatteredInterpolant(x pos, y pos, z pos, electric potential, 'nearest', 'none');
% Use the interpolant to find values at neuron positions
K = F(nwpos(:,1), nwpos(:,2), nwpos(:,3));
K(isnan(K)) = 0; % Setting NaNs to 0
Vext=currents.*K;
%%total voltage at the electrode
Vext elec=sum(Vext,1);
plot(Vext elec);
xlabel('Time');
ylabel('Voltage');
title('EXTERNAL VOLTAGE');
Part 4
axonLength = 5; % mm
numCompartments = 200; % Number of compartments
current = 1000; %mA
sigma = 0.00033; % S/mm,
radius=10; %um
x coordinates = linspace(-axonLength / 2, axonLength / 2, numCompartments);
y coordinates=zeros(1, numCompartments);
z coordinates=zeros(1, numCompartments);
axon pos=[x coordinates;y coordinates;z coordinates]';
axon hilloc=axon pos(100,:);
%%electrode position is 1mm away from ceneter
electrode=axon hilloc+[0,1,0];
distances=sqrt((electrode(1)-axon pos(:,1).^2)+(electrode(2)-axon pos(:,2).^2)+
(electrode(3)-axon pos(:,3).^2));
qa=3*10^-5; %S
Vext=current/4*3.14*sigma*distances; %mV
Iint = zeros(size(Vext));
for n = 2:length(Vext) - 1
   Iint(n) = ga * (Vext(n-1) - 2*Vext(n) + Vext(n+1));
% Handling boundary conditions for n = 1 and n = 200
Iint(1) = ga * (-2*Vext(1) + Vext(2));
Iint(end) = qa * (Vext(end-1) - 2*Vext(end));
Iint=(Iint);
plot(Iint)
%%ques1: 40pA to cross nthe threshold
%%ques4: mA>pA >>> 10^15pA
%>>1A
```