

AMATH 422 HW 5 Submission

I Dwell time distributions: theory.

Consider an ion channel with 4 open states and 2 closed states. Give a mathematical argument, similar to that in class, that derives the typical functional form of the dwell time distribution for the channel being in any one of the open states. Note: I am looking for a derivation here, not just a statement of the answer or a "mantra" or result from class.

Four open states (S_1, S_2, S_3, S_4) Two closed states (S_5, S_6)

$$\Omega = \{Open, Closed\} = \{S_1, S_2, S_3, S_4, S_5, S_6\}$$

Indistinguishable between S and the other open states

We want to find the mathematical argument for the dwell time: $P(\text{Dwell time in } (C_1, C_2, C_3, C_4) \text{ for } k \text{ steps})$

We are going to assume time is homogenous and time does not affect the markov chain probabilities. We are going to assume the gate starts in the open condition $X_0 \in \{Open\}$

Want to find the probability Open for K steps $P(T_o \geq k)$

We know that the probability of going to other open steps is $A p_0 = p_1$ Only interested in open states

$$\begin{bmatrix} a_{11} & a_{12} & a_{13} & a_{14} & a_{15} & a_{16} \\ a_{21} & a_{22} & a_{23} & a_{24} & a_{25} & a_{26} \\ a_{31} & a_{32} & a_{33} & a_{34} & a_{35} & a_{36} \\ a_{41} & a_{42} & a_{43} & a_{44} & a_{45} & a_{46} \\ a_{51} & a_{52} & a_{53} & a_{54} & a_{55} & a_{56} \\ a_{61} & a_{62} & a_{63} & a_{64} & a_{65} & a_{66} \end{bmatrix} \begin{bmatrix} p_1(0) \\ p_2(0) \\ p_3(0) \\ p_4(0) \\ 0 \\ 0 \end{bmatrix} = \begin{bmatrix} p_1(1) \\ p_2(1) \\ p_3(1) \\ p_4(1) \\ p_5(1) \\ p_6(1) \end{bmatrix}$$

We can remove the two right columns due to the zeros in $p(0)$ we can remove the bottom two row because those would release us from the dwell time. Simplifying becomes

$$\begin{bmatrix} a_{11} & a_{12} & a_{13} & a_{14} \\ a_{21} & a_{22} & a_{23} & a_{24} \\ a_{31} & a_{32} & a_{33} & a_{34} \\ a_{41} & a_{42} & a_{43} & a_{44} \end{bmatrix} \begin{bmatrix} p_1(0) \\ p_2(0) \\ p_3(0) \\ p_4(0) \end{bmatrix} = \begin{bmatrix} p_1(1) \\ p_2(1) \\ p_3(1) \\ p_4(1) \end{bmatrix}$$

This holds for k and $k+1$ steps because

$$P_{1,2,3,4}^c(k) = P(X_k = S_i | X_t \in \{S_1, S_2, S_3, S_4\} 0 \leq t \leq k-1)$$

$$\begin{bmatrix} a_{11} & a_{12} & a_{13} & a_{14} \\ a_{21} & a_{22} & a_{23} & a_{24} \\ a_{31} & a_{32} & a_{33} & a_{34} \\ a_{41} & a_{42} & a_{43} & a_{44} \end{bmatrix} \begin{bmatrix} p_1(k) \\ p_2(k) \\ p_3(k) \\ p_4(k) \end{bmatrix} = \begin{bmatrix} p_1(k+1) \\ p_2(k+1) \\ p_3(k+1) \\ p_4(k+1) \end{bmatrix}$$

Linear map of matrix A can be expressed as. A is 4x4 so there should be 4 terms in the final math expression

$$A = \sum_{i=1}^n C_i \lambda_i \mathbf{v}_i$$

Where λ_i is the corresponding eigenvalue \mathbf{v}_i is the corresponding eigenvector and C_i is the corresponding arbitrary constant

Thus at time step k the probability the channel is in the open state

$$P_{1,2,3,4}^c(k) = C_1 \lambda_1^k \mathbf{v}^1 + C_2 \lambda_2^k \mathbf{v}^2 + C_3 \lambda_3^k \mathbf{v}^3 + C_4 \lambda_4^k \mathbf{v}^4$$

II Simulating Markov chains and neural spiking

```
In [1]: import matplotlib.pyplot as plt    # That gives plotting, and the next line makes plots appear inline
%matplotlib inline
import numpy as np    # That gives numerical arrays and tools for manipulating them
import scipy.optimize as opt
import scipy.linalg as la

from scipy.optimize import curve_fit
```

```
In [2]: # Initialize random number generator
rng = np.random.default_rng()
#Define Transition Matrices
A = np.array([[0.98, 0.10, 0],
              [0.02, 0.70, 0.05],
              [0, 0.20, 0.95]]) #Inward Channel

B = np.array([[0.90, 0.10, 0],
              [0.10, 0.60, 0.10],
              [0, 0.30, 0.90]]) #BOutward Channel

print(A)
print(B)
```

```
[[0.98 0.1  0.  ]
 [0.02 0.7  0.05]
 [0.   0.2  0.95]]
[[0.9 0.1 0.  ]
 [0.1 0.6 0.1]
 [0.  0.3 0.9]]
```

```
In [3]: def voltage_gate(Tmax, A, count, Direction):
    #matrix A
    states=np.zeros(Tmax,dtype=int)
    states[0]=0
    voltage=np.zeros(Tmax,dtype=int)
    voltage[0]=0
    plt.figure()
    for i in range(count):
        for t in np.arange(Tmax - 1):
            r = rng.uniform(0, 1) # draw random variable (uniformly distributed in 0,1)
            if states[t] == 0:
                if r < A[0, 0]: #transition 0 to 0
                    states[t + 1] = 0
                    voltage[t + 1] += 0
                else: #transition 0 to 1
                    states[t + 1] = 1
            elif states[t] == 1:
```

```

        if r < A[1, 1]: #transition 1 to 1.
            states[t + 1] = 1
        elif A[1, 1] < r < A[1, 1] + A[0, 1]: #transition 1 to 0.
            states[t + 1] = 0
            #transition 1 to 1 < r < transition 1 to 1 + transition 1 to 0
        else: #transition 1 to 2
            states[t + 1] = 2
            if Direction == "Inward":
                voltage[t + 1] += 1
            else:
                voltage[t + 1] -= 1

    elif states[t] == 2:
        if r < A[2, 2]: #transition 2 to 2
            states[t + 1] = 2
            if Direction == "Inward":
                voltage[t + 1] += 1
            else:
                voltage[t + 1] -= 1
        else: #transition 2 to 1
            states[t + 1] = 1

    #Plot of states
    plt.plot(states)
    plt.xlabel('state')
    plt.ylabel('timestep')
    plt.yticks([0, 1, 2]) # Limit y-axis to only show values 0, 1, 2
    #print(A)

    return states, voltage

```

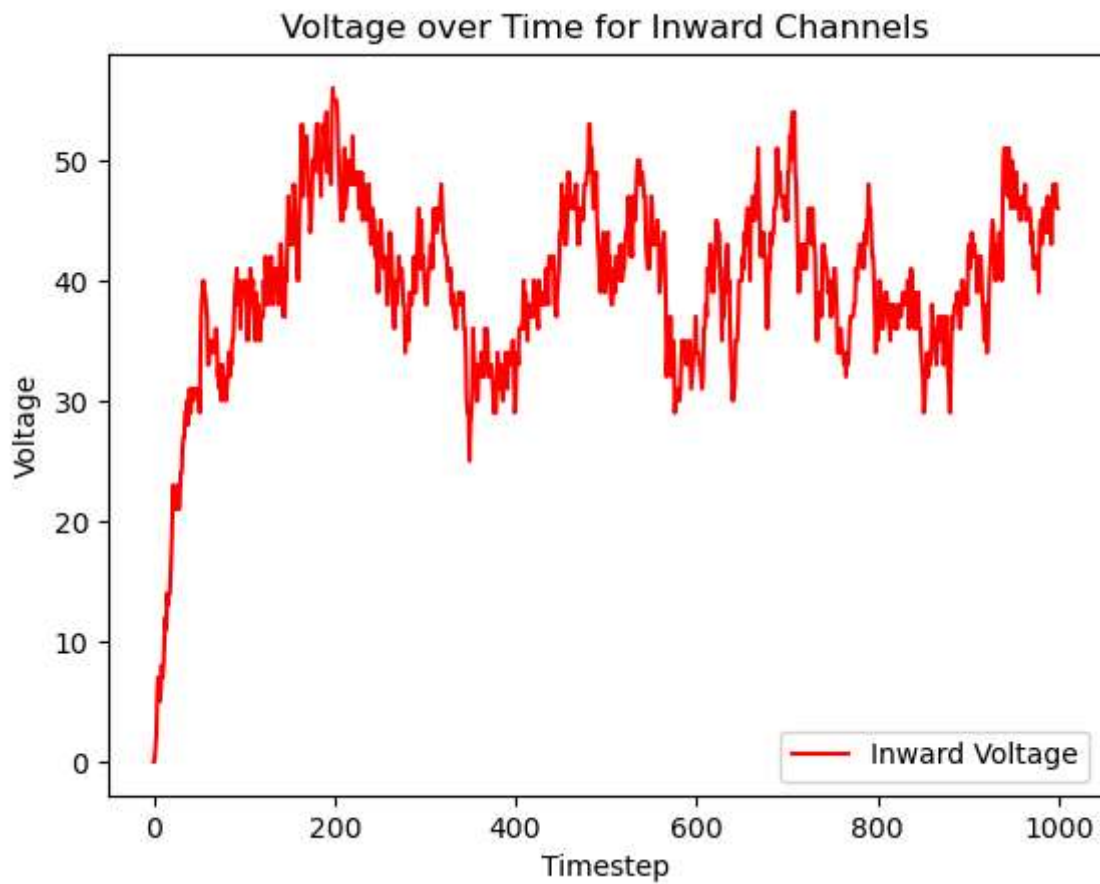
Assume there are a total of $n_{inward} = 100$ inward channels, each evolving independently under a realization of the Markov kinetics above. If n_{inward} of these channels are in the open configuration at timestep t , then the total inward current is $+n_{inward}$

```

In [4]: states_inward, voltage_inward = voltage_gate(Tmax=1000,A=A, count=100,Direction="Inward")
plt.figure()
plt.plot(voltage_inward, label="Inward Voltage", color="red")
plt.xlabel("Timestep")
plt.ylabel("Voltage")
plt.legend()
plt.title("Voltage over Time for Inward Channels")
plt.show()

```

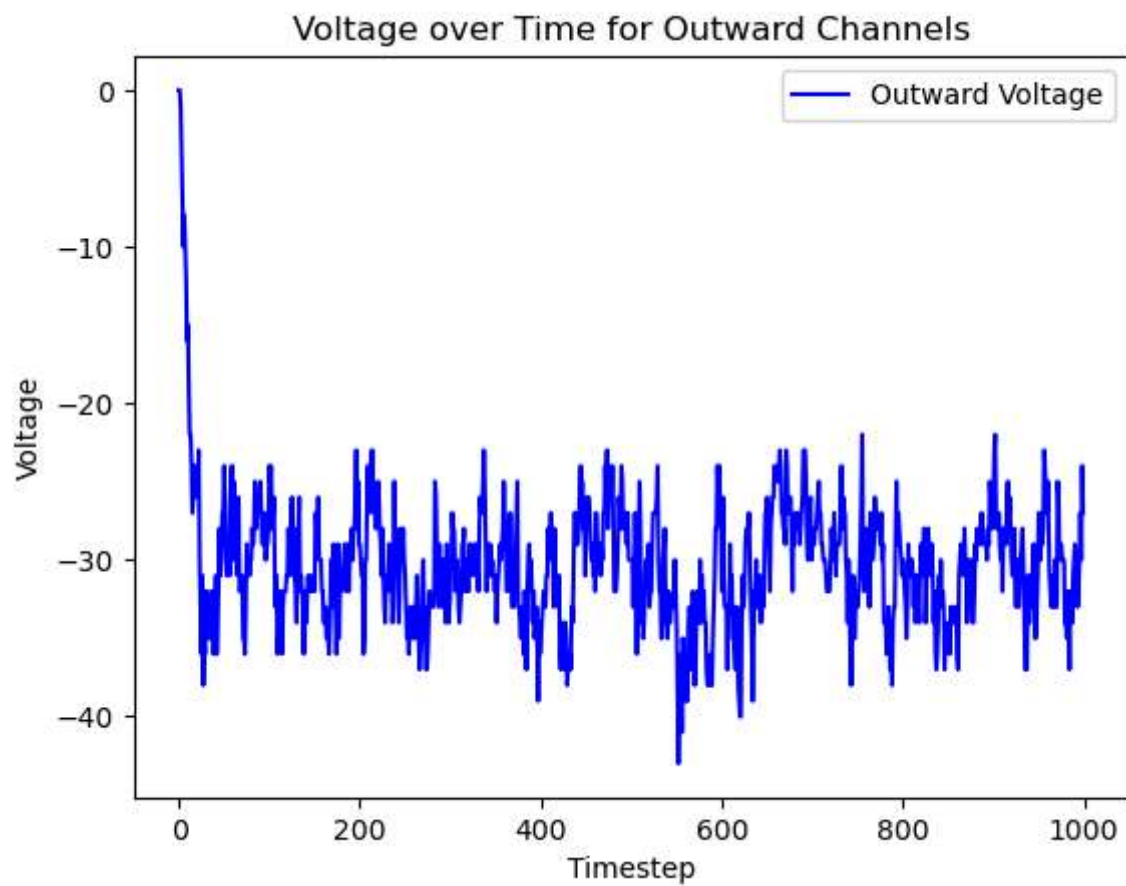
<Figure size 640x480 with 0 Axes>



Assume there are a total of $N_{outward} = 50$ outward channels, each evolving independently under a realization of the Markov kinetics above. If $n_{outward}$ of these channels are in the open configuration at timestep t , then the total outward current is $-n_{outward}$ units.

```
In [5]: states_outward, voltage_outward = voltage_gate(Tmax=1000,A=B, count=50,Direction="Outward")
plt.figure()
plt.plot(voltage_outward, label="Outward Voltage", color="blue")
plt.xlabel("Timestep")
plt.ylabel("Voltage")
plt.legend()
plt.title("Voltage over Time for Outward Channels")
plt.show()
```

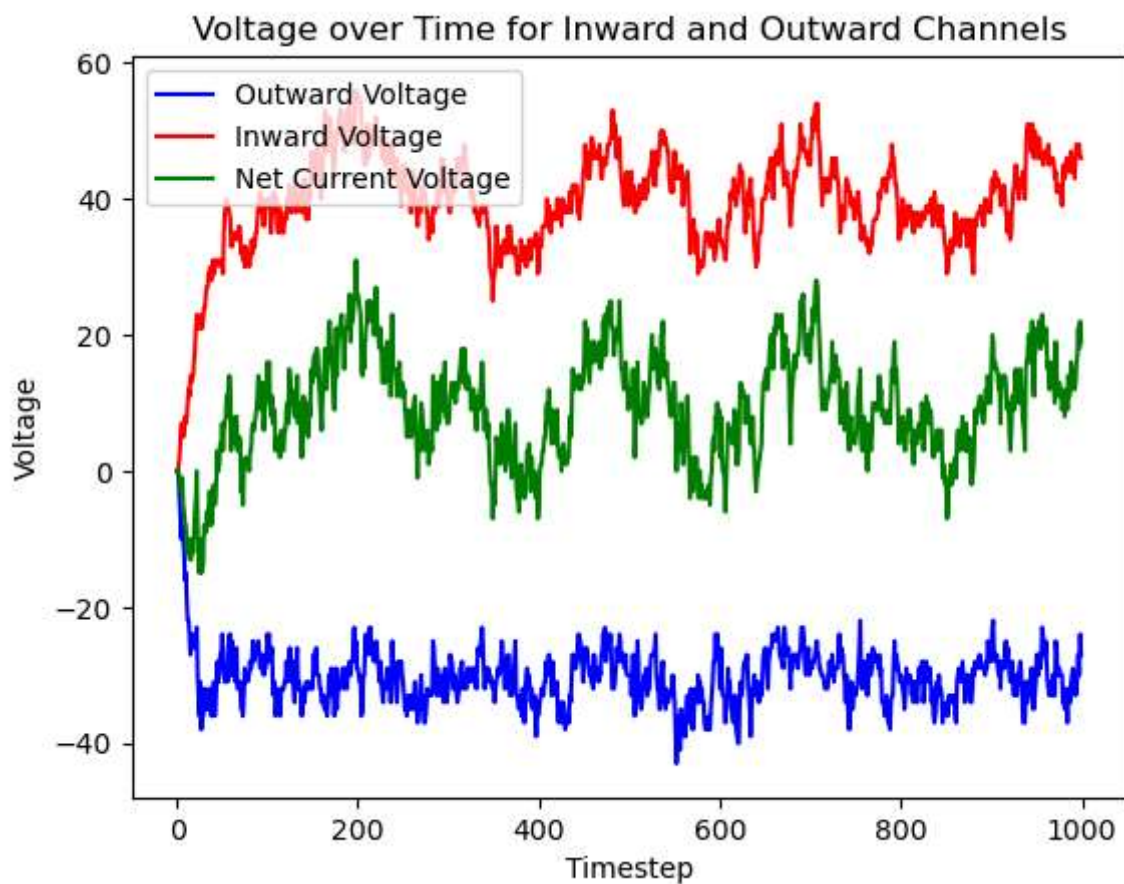
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Thus, the net current into the cell at timestep t is $n_{inward} - n_{outward}$: the number of open inward channels minus the number of open outward channels.

```
In [6]: net_current = voltage_inward + voltage_outward

#Plot of Net Current
plt.figure()
plt.plot(voltage_outward, label="Outward Voltage", color="blue")
plt.plot(voltage_inward, label="Inward Voltage", color="red")
plt.plot(net_current, label="Net Current Voltage", color="green")
plt.xlabel("Timestep")
plt.ylabel("Voltage")
plt.legend()
plt.title("Voltage over Time for Inward and Outward Channels")
plt.show()
```



In our model, the cell will produce an action potential (spike) in a given timestep if this net current is greater than a threshold value T . Assume that the channels have settled into equilibrium (i.e., that a time has passed that is large enough since a simulation was initialized). Plot the probability that the cell will produce a spike in a given timestep, as a function of the spiking threshold T .

There are at least two ways of doing this: (1) by computing the equilibrium state probabilities, and simulating many coin tossings, or (2) by computing the equilibrium state probabilities, and using the form of the binomial distribution.

```
In [7]: # Define threshold range and Tmax
thresholds = np.arange(-20, 50, 1)
Tmax = 1000

# Define spike probability calculation function
def spike_probability(voltage, thresholds):
    spike_probs = []
    for T in thresholds:
        spikes = np.sum(voltage >= T) # Count timesteps where voltage >= T
        spike_prob = spikes / len(voltage) # Fraction of timesteps with spikes
        spike_probs.append(spike_prob)
    return spike_probs

# Create subplots with a 2x3 grid
fig, axes = plt.subplots(2, 3, figsize=(18, 10)) # Adjust figsize as needed
counts = [10, 100, 1000]

for i, count in enumerate(counts):
    # Generate voltage data
    states_inward, voltage_inward = voltage_gate(Tmax=Tmax, A=A, count=count, Direction="Inward")
    states_outward, voltage_outward = voltage_gate(Tmax=Tmax, A=B, count=count // 2, Direction="Outward")

    # Calculate net current and spike probabilities
```

```

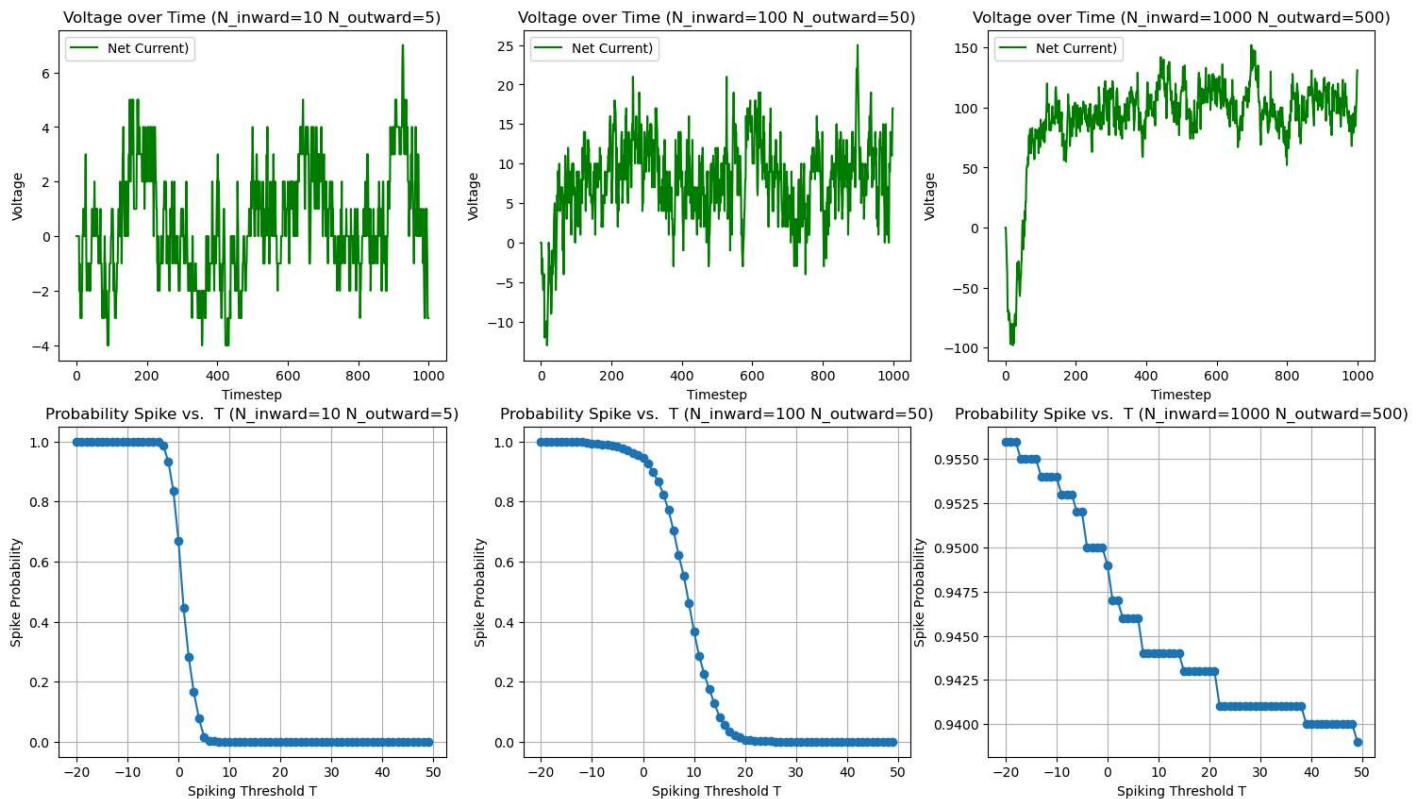
net_current = voltage_inward + voltage_outward
spike_probs = spike_probability(net_current, thresholds)

# Plot net current voltage over time
ax1 = axes[0, i] # First row
ax1.plot(net_current, label=f"Net Current)", color="green")
ax1.set_xlabel("Timestep")
ax1.set_ylabel("Voltage")
ax1.legend()
ax1.set_title(f"Voltage over Time (N_inward={count} N_outward={count//2})")

# Plot spike probability vs. threshold
ax2 = axes[1, i] # Second row
ax2.plot(thresholds, spike_probs, marker='o', linestyle='-')
ax2.set_xlabel("Spiking Threshold T")
ax2.set_ylabel("Spike Probability")
ax2.set_title(f"Probability Spike vs. T (N_inward={count} N_outward={count//2})")
ax2.grid(True)

# Adjust layout and display
plt.tight_layout()
plt.show()

```



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(2) by computing the equilibrium state probabilities, and using the form of the binomial distribution.

Write a few sentences reporting on any qualitative changes that you observe.

In [8]: # At Longer time the average net current is more settled on one value. At shorter Time max there
 # The more outward and inward channels there are the larger the magnitude of the net current.

When there are double the amount of inward channels compared to outward channels there is a mor
It appears the I_{net} current is
The Longer the timestep the more smooth the function of the thresholds is.
The threshold probability shape is relatively constant
There is no major difference between the amounts on the size of the net current.
The net membrane can be either positive or negative