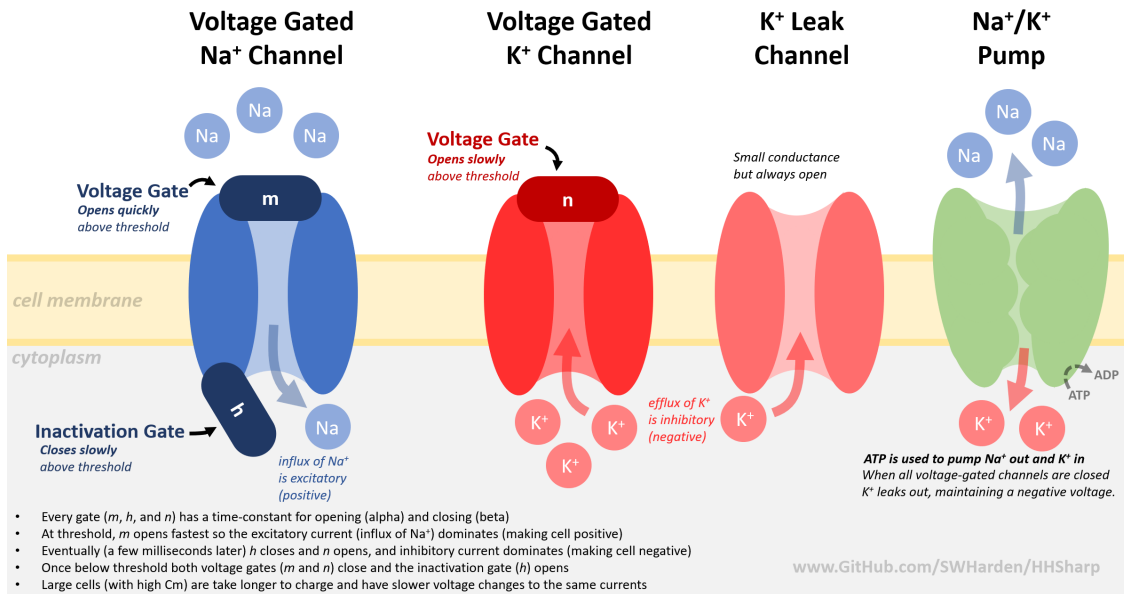


February 11, 2024

1 PSY 452 - Advanced Neurophysiological Psychology

1.1 Hodgkin-Huxley Neuromorphic Model Demonstration for Presentation by Dan Jang

1.1.1 The Hodgkin-Huxley Model [1], Illustrated [2]



1.1.2 The Full Hodgkin-Huxley Model [1] Equations (in LaTeX [3])

"The Hodgkin-Huxley equations using modern conventions are given by:

$$\frac{dV}{dt} = [I_{inj} - \bar{g}_{Na} m^3 h (V - V_{Na}) - \bar{g}_K n^4 (V - V_K) - g_L (V - V_L)] / C \quad (1)$$

$$\frac{dn}{dt} = \alpha_n(V)(1 - n) - \beta_n(V)n \quad (2)$$

$$\frac{dm}{dt} = \alpha_m(V)(1 - m) - \beta_m(V)m \quad (3)$$

$$\frac{dh}{dt} = \alpha_h(V)(1 - h) - \beta_h(V)h \quad (4)$$

where:

$$\alpha_n(V) = \frac{0.01(V + 55)}{1 - \exp[-(V + 55)/10]} \quad (5)$$

$$\beta_n(V) = 0.125 \exp[-(V + 65)/80] \quad (6)$$

$$\alpha_m(V) = \frac{0.1(V + 40)}{1 - \exp[-(V + 40)/10]} \quad (7)$$

$$\beta_m(V) = 4 \exp[-(V + 65)/18] \quad (8)$$

$$\alpha_h(V) = 0.07 \exp[-(V + 65)/20] \quad (9)$$

$$\beta_h(V) = \frac{1}{1 + \exp[-(V + 35)/10]} \quad (10)$$

The values of the constants are:

- $C = 1\mu F/cm^2$;
- $\bar{g}_{Na} = 120mS/cm^2$;
- $V_{Na} = 50mV$;
- $\bar{g}_K = 36mS/cm^2$;
- $V_K = -77mV$;
- $g_L = 0.3mS/cm^2$;
- $V_L = -54mV$;

In these equations, voltages are in mV , current densities in $\mu A/cm^2$, capacitance in $\mu F/cm^2$, and time in ms .” [3]

1.1.3 Quick & Concise Variate & Verbose Variables List Made Easier-to-Understand [1]

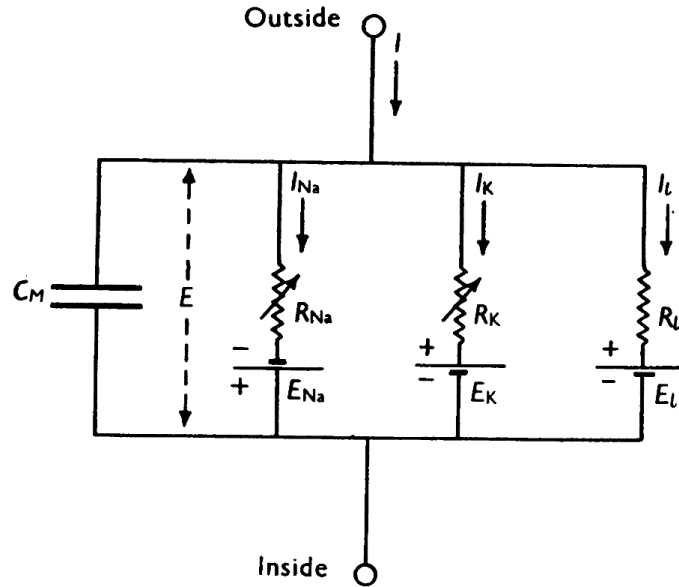


Fig. 1. Electrical circuit representing membrane. $R_{Na}=1/g_{Na}$; $R_K=1/g_K$; $R_L=1/\tilde{g}_l$. R_{Na} and R_K vary with time and membrane potential; the other components are constant.

1. **V_m**: Membrane Potential - Describes the electrochemical gradient across the neuron membrane. This variable is what the Hodgkin-Huxley model primarily focuses on to predict, etc!
2. **C_m**: Membrane Capacitance - Represents the ability of the neuron membrane to store charge as a constant variable!
3. **I**: External Current - The current, in our case, represents Squidward's impulse as output, applied to Rob, the Motor Neuron, to start positioning his clarinet!
4. **g_{Na}, g_K, g_L**: Maximum Conductances for the Sodium (Na⁺) Cation, Potassium (K⁺) Cation, and Leak (L {e.g., Chlorine Cation (Cl⁻) or Calcium Anion (Ca²⁺)}) Channels!
5. **E_{Na}, E_K, E_L**: Equilibrium Potentials for Sodium (Na⁺) Cation, Potassium (K⁺) Cation, and Leak (L {e.g., Chlorine Cation (Cl⁻) or Calcium Anion (Ca²⁺)}) Channels! These values represent the membrane potential, when reached, results / indicates no net flux in-between the membrane (intracellular space {e.g., axoplasm} or extracellular space)!
6. **m, h**: Sodium (Na⁺) channel activation and inactivation probability variables, they are also called & part of the gate variables!
7. **n**: Potassium (K⁺) channel activation probability variable, also one of the gate variables!
8. **alpha/beta + __ + n/m/h**: These represent rate constants. Alpha for opening & Beta for closing. Followed by either n, m, or h, representing the gate variables as described above!
9. **I_{Na}, I_K, I_L**: Currents that are following through the Sodium (Na⁺) Cation, Potassium (K⁺) Cation, and Leak (L {e.g., Chlorine Cation (Cl⁻) or Calcium Anion (Ca²⁺)}) Ionic Channels!

1.1.4 Part 0: Initialization

```
[1]: ##### PSY 452 (Winter 2024) - Advanced Neurophysiological Psychology: Part 0,
      ↪ Initialization
      ### By Dan Jang

      ## I.) Installing necessary packages
      # !pip install numpy
      # !pip install matplotlib.pyplot
      # !pip install scipy

      # ## II.) Importing essential packages + pyHodgkin-Huxley package [1]

      # import sys
      # import os

      # currpath = os.path.join(os.getcwd(), 'libraries', 'pyhh')

      # if currpath not in sys.path:
      #     sys.path.append(currpath)

      ## II.) Importing essential packages + pyHodgkin-Huxley package [1]

      import numpy as np
      import matplotlib.pyplot as plot
      import scipy.integrate
      from scipy.integrate import odeint

      import pyhh as hodhux

      ## III.) Create full Hodgkin-Huxley model-based action potential stimulation

      ## III.A) Create Squidward's Rob Motor Neuron :~)
      class Squidward:
          def __init__(self):
              self.robert, self.gNa, self.gK, self.gL, self.Cm, self.ENa, self.EK,
              ↪ self.EL, self.I, self.Vm, self.m, self.h, self.n, self.t, self.pulse, self.
              ↪ pulsex, self.pulsey, self.movement, self.playtime = None, 120.0, 36.0, 0.3,
              ↪ 1.0, 50.0, -77.0, -54.387, 10.0, -65.0, 0.052, 0.596, 0.317, 50, 7000,
              ↪ 13000, None, 0.01, np.zeros(20000)
              self.timestep = 0.01
              self.robert = Robert()

              # Essentially, a funny named new-custom constructor
              def tentacles(self, robert, gNa, gK, gL, Cm, ENa, EK, EL, I, Vm, m, h, n,
              ↪ t, pulse, pulsex, pulsey, movement, timestep):
```

```

    if robert is None:
        if self.robert is None:
            self.robert = Robert()

    else:
        try:
            self.robert.activation(robert, gNa, gK, gL, Cm, ENa, EK, EL, I,
↳Vm, m, h, n, t, pulse, pulsex, pulsey)
        except:
            print("[E1<Squid-Init>] bruh u might have missed a neuronal_
↳parameter for rob lol")

    self.playtime = np.zeros(t) #self.robert.t
    self.playtime[pulsex:pulsey] = pulse

    #if clarinet is None:
    #    if self.clarinet is None:
    if timestep is None:
        timestep = 0.01
        self.timestep = timestep
    else:
        self.timestep = timestep
    if movement is None:
        self.movement = hodhux.Simulation(self.robert.model)
        self.clarinet(self.timestep, self.movement)
    else:
        self.clarinet(self.timestep, movement)
        #self.clarinet = clarinet(t, timestep, )

    #self.playtime = self.robert.t

def clarinet(self, timestep, movement):
    # if playtime is None:
    #     playtime = np.zeros(20000)

    if timestep is None:
        timestep = 0.01

    if movement is None:
        self.movement = hodhux.Simulation(self.robert.model)
        self.movement.Run(stimulusWaveform=self.playtime,
↳stepSizeMs=timestep)
        self.movement = movement
        #print("[E4<Squid-Clarinet>] bruh u gotta pass a movement thingy so_
↳u can actually graph the results u silly head lol")

    else:

```

```

#     try:
#         movement.Run(stimulusWaveform=self.playtime, stepSizeMs=timestep)
#         self.movement = movement
#     except:
#         print("[E3<Squid-Clarinet>] bruh squidward's robert motor
↳neuron model might be misconfigured lol")

def sheetmusic(self, figsizex, figsizey, color1, color2, color3, color4,
↳color5, color6, color7, color8, title, file):
    #try:
    if figsizex:
        if figsizey:
            plot.figure(figsize=(figsizex, figsizey))
        else:
            plot.figure(figsize=(10, 8))
    # except:
    #     print("[E4<Squid-SheetMusic>] bruh u might have missed the figure
↳sizes for the sheet music lol")

    if title is None:
        title = "PSY 452 (Winter 2024) - Hodgkin-Huxley Neuromorphic
↳Computational Graphical Model"

    else:
        title = "PSY 452 (Winter 2024) - Hodgkin-Huxley Neuromorphic
↳Computational Graphical Model: " + title

    if file is None:
        file = "PSY452-Hodgkin-Huxley-Model-Waveform-DanJ.png"

    if self.movement is None:
        print("[E4<Squid-SheetMusic>] bruh squidward apparently hasnt moved
↳his clarinet or if he did, it wasn't recorded u silly goose lol")

    if self.robert is None:
        print("[E5<Squid-SheetMusic>] bruh somehow at this stage, Robert
↳the Motor Neuron still has not been created yet, you are one silly goose dan
↳lmao")

    else:
        movement = self.movement # lazy but saves typing self. each time
↳lmao

        # Potential (mV) Color
        if color1 is None:
            color1 = 'r'

```

```

# Input Stimulation ( $\mu\text{A}/\text{cm}^2$ ) Color
if color2 is None:
    color2 = 'b'

# Sodium Activation Gating Variable (m) Color
if color3 is None:
    color3 = 'purple'

# Sodium Inactivation Gating Variable (h) Color
if color4 is None:
    color4 = 'orange'

# Potassium Activation Gating Variable (n) Color
if color5 is None:
    color5 = 'yellow' # BANANA

# Sodium Current ( $I_{\text{Na}}$ ) ( $\mu\text{A}/\text{cm}^2$ ) Color
if color6 is None:
    color6 = 'purple'

# Potassium Current ( $I_{\text{K}}$ ) ( $\mu\text{A}/\text{cm}^2$ ) Color
if color7 is None:
    color7 = 'yellow'

# Leak Current ( $I_{\text{L}}$ ) ( $\mu\text{A}/\text{cm}^2$ ) Color
if color8 is None:
    color8 = 'turquoise'

## Credits to [2] for parts from README.md Example Code

# Potential (mV) Plot #1
potentialplt = plot.subplot(411)
potentialplt.set_title(title)
potentialplt.set_ylabel("Rob's Membrane Potential (mV)")
potentialplt.plot(movement.times, movement.Vm - 70, color=color1)

# Input Stimulation ( $\mu\text{A}/\text{cm}^2$ ) Plot #2
stimulationplt = plot.subplot(412)
stimulationplt.set_ylabel("Squidward Stimulation ( $\mu\text{A}/\text{cm}^2$ )")
playtime = self.playtime
## Sanity Debug Print Statement
#print(movement.times, playtime, color2)
stimulationplt.plot(movement.times, playtime, color=color2)

# Gating Variable Activation (m/h/n) Plot #3
activationplt = plot.subplot(413, sharex=potentialplt)
activationplt.set_ylabel("Rob's Channel Activation (fractional)")

```

```

        activationplt.plot(movement.times, movement.StateM, label='m (Na+  

↪Channel Open Prob.)', color=color3)
        activationplt.plot(movement.times, movement.StateH, label='h (Na+  

↪Channel Close Prob.)', color=color4)
        activationplt.plot(movement.times, movement.StateN, label='n (K+  

↪Channel Open Prob.)', color=color5)
        activationplt.legend()

        # Ionic Channel Current Plot (INa/IK/Ileak) Plot #4
        currentplt = plot.subplot(414, sharex=potentialplt)
        currentplt.set_xlabel("Clarinet Action Potential Time (in_  

↪milliseconds)")
        currentplt.set_ylabel("Rob Membrane Ionic Current (μA/cm²)")
        currentplt.plot(movement.times, movement.INa, label='INa (Sodium_  

↪Current)', color=color6)
        currentplt.plot(movement.times, movement.IK, label='IK (Potassium_  

↪Current)', color=color7)
        currentplt.plot(movement.times, movement.IKleak, label='IKleak_  

↪(KLeakage Current)', color=color8)
        currentplt.legend()

        plot.tight_layout()
        plot.savefig(file)
        plot.show()

class Robert:
    def __init__(self):
        self.model = hodhux.HHModel()
        self.t = None

    def activation(self, model, gNa, gK, gL, Cm, ENa, EK, EL, I, Vm, m, h,  

↪n, t, pulse, pulsex, pulsey):

        if model is None:
            if self.model is None:
                robert = hodhux.HHModel()

                # Squidward's Rob Motor Neuron Parameter, gNa: Avg. Na+  

↪(Sodium cation) conductance per unit area (mS/cm²)
                robert.gNa = 120

                # Squidward's Rob Motor Neuron Parameter, gK: Avg. K+  

↪(Potassium cation) conductance per unit area (mS/cm²)
                robert.gK = 36

```



```

# Squidward's Rob Motor Neuron Parameter, gL: Avg. Leak (e.
↪g., Chlorine anions / Calcium {2+} cations) conductance per unit area (mS/
↪cm2)

robert.gKleak = 0.3

# Squidward's Rob Motor Neuron Parameter, Cm: Membrane
↪capacitance per unit area (uF/cm2)

robert.Cm = 1.0

# Squidward's Rob Motor Neuron Parameter, ENa: Sodium
↪cation Nernst reversal potential (mV)

robert.ENa = 115

# Squidward's Rob Motor Neuron Parameter, EK: Potassium
↪cation Nernst reversal potential (mV)

robert.EK = -12

# Squidward's Rob Motor Neuron Parameter, EL: Leak Nernst
↪reversal potential (mV)

robert.EKleak = 10.6

# Squidward's Rob Motor Neuron Parameter, I: External
↪current (uA/cm2)

#robert.I = 10.0

# Squidward's Rob Motor Neuron Parameter, Vm: Membrane
↪potential (mV)

#robert.Vm = 0

# Squidward's Rob Motor Neuron Parameter, m: Sodium cation
↪activation gating variable

#robert.m = 0

# Squidward's Rob Motor Neuron Parameter, h: Sodium cation
↪inactivation gating variable

#robert.h = 0

# Squidward's Rob Motor Neuron Parameter, n: Potassium
↪cation activation gating variable

#robert.n = 0

# Stimulus Waveform thingy
self.t = np.zeros(20000)

# Stimulus Waveform <+ Square Pulse
self.t[7000:13000] = 50

```

```

        self.model = robert
try:

    robert = self.model

    if gNa is not None:
        robert.gNa = gNa

    if gK is not None:
        robert.gK = gK

    if gL is not None:
        robert.gKleak = gL

    if Cm is not None:
        robert.Cm = Cm

    if ENa is not None:
        robert.ENa = ENa

    if EK is not None:
        robert.EK = EK

    if EL is not None:
        robert.EKleak = EL

    # if I is not None:
    #     robert.I = I

    # if Vm is not None:
    #     robert.Vm = Vm

    # if m is not None:
    #     robert.m = m

    # if h is not None:
    #     robert.h = h

    # if n is not None:
    #     robert.n = n

    if t is not None:
        # Stimulus Waveform thingy
        self.t = np.zeros(t)

    if pulse is not None:

```

```

        if pulsex is not None:
            if pulsey is not None:
                self.t[pulsex:pulsey] = pulse

        self.model = robert
    except:
        print("[E2<SubSquid-Rob-Init>] bruh u might have missed a_
↳neuronal parameter for rob within squidward initialization thingy lol")

    def model(self):
        return self.model

### Finally, the grand test for class Squidward (main wrapper), with a subclass_
↳Robert (le Motor Neuron model encompassing wrapper)

# Reference - Squidward(self, robert, gNa, gK, gL, Cm, ENa, EK, EL, I, Vm, m,
↳h, n, t, pulse, pulsex, pulsey, movement, timestep)
squidward = Squidward()

# Reference -
# Squidward's Rob Motor Neuron Parameter, gNa: Avg. Na+ (Sodium cation)_
↳conductance per unit area (mS/cm^2)
# robert.gNa = 120.0
# # Squidward's Rob Motor Neuron Parameter, gK: Avg. K+ (Potassium cation)_
↳conductance per unit area (mS/cm^2)
# robert.gK = 36.0
# # Squidward's Rob Motor Neuron Parameter, gL: Avg. Leak (e.g., Chlorine_
↳anions / Calcium {2+} cations) conductance per unit area (mS/cm^2)
# robert.gKleak = 0.3
# # Squidward's Rob Motor Neuron Parameter, Cm: Membrane capacitance per unit_
↳area (uF/cm^2)
# robert.Cm = 1.0
# # Squidward's Rob Motor Neuron Parameter, ENa: Sodium cation Nernst reversal_
↳potential (mV)
# robert.ENa = 115.0
# # Squidward's Rob Motor Neuron Parameter, EK: Potassium cation Nernst_
↳reversal potential (mV)
# robert.EK = -12.0
# # Squidward's Rob Motor Neuron Parameter, EL: Leak Nernst reversal potential_
↳(mV)
# robert.EKleak = 10.6
# # Squidward's Rob Motor Neuron Parameter, I: External current (uA/cm^2)
# robert.I = 10.0
# # Squidward's Rob Motor Neuron Parameter, Vm: Membrane potential (mV)
# robert.Vm = -65.0

```

```

# # Squidward's Rob Motor Neuron Parameter, m: Sodium cation activation gating
    ↪variable
# robert.m = 0.052
# # Squidward's Rob Motor Neuron Parameter, h: Sodium cation inactivation
    ↪gating variable
# robert.h = 0.596
# # Squidward's Rob Motor Neuron Parameter, n: Potassium cation activation
    ↪gating variable
# robert.n = 0.317
# tentacles(self, robert, gNa, gK, gL, Cm, ENa, EK, EL, I, Vm, m, h, n, t,
    ↪pulse, pulsex, pulsey, movement, timestep):

squidward.tentacles(None, 120.0, 36.0, 0.3, 1.0, 115, -12, 10.6, 0, 0, 0, 0, 0,
    ↪10000, 50, 1000, 9000, None, 0.01)

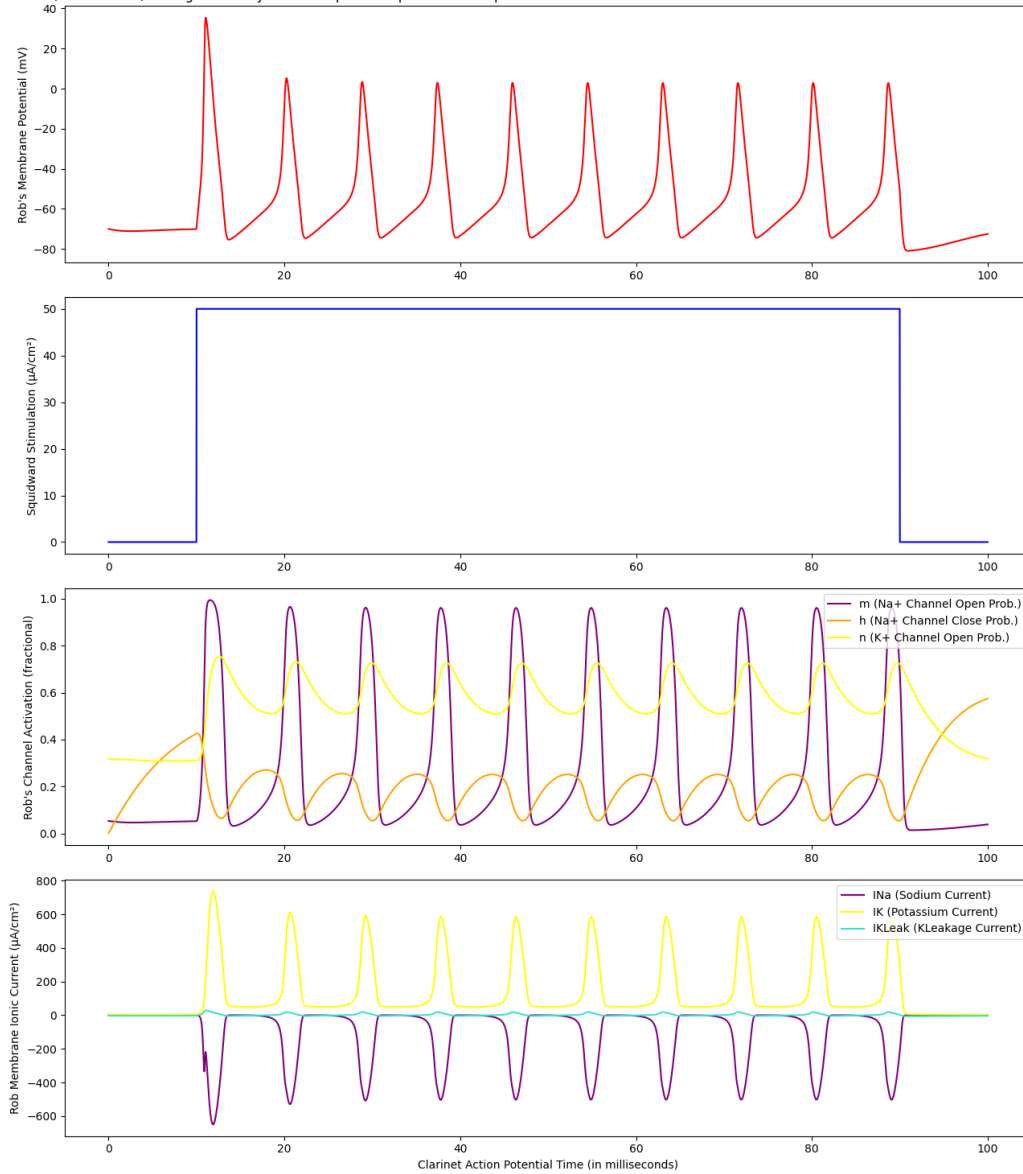
# Reference - sheetmusic(self, figsizex, figsizey, color1, color2, color3,
    ↪color4, color5, color6, color7, color8, title)
#         movement = self.movement # lazy but saves typing self. each time
    ↪lmao
# # Potential (mV) Color
#     color1 = 'r'
# # Input Stimulation ( $\mu\text{A}/\text{cm}^2$ ) Color
#     color2 = 'b'
# # Sodium Activation Gating Variable (m) Color
#     color3 = 'purple'
# # Sodium Inactivation Gating Variable (h) Color
#     color4 = 'orange'
# # Potassium Activation Gating Variable (n) Color
#     color5 = 'yellow' # BANANA
# # Sodium Current (INa) ( $\mu\text{A}/\text{cm}^2$ ) Color
#     color6 = 'purple'
# # Potassium Current (IK) ( $\mu\text{A}/\text{cm}^2$ ) Color
#     color7 = 'yellow'
# # Leak Current (IL) ( $\mu\text{A}/\text{cm}^2$ ) Color
#     color8 = 'turquoise'

squidward.sheetmusic(13, 15, 'r', 'b', 'purple', 'orange', 'yellow', 'purple',
    ↪'yellow', 'turquoise', "Full Clarinet Orchestral Positionization Action
    ↪Potential Stimulation Waveform",
    ↪"PSY452-FullSquidwardClarinetActionPotentialStimulationWave-DanJ.png")

```

Simulating 10000 time points...
Simulation complete!

PSY 452 (Winter 2024) - Hodgkin-Huxley Neuromorphic Computational Graphical Model: Full Clarinet Orchestral Positionization Action Potential Stimulation Waveform



1.1.5 Part 1: *Resting Membrane Potential (RMP)*

```
[2]: ##### PSY 452 (Winter 2024) - Advanced Neurophysiological Psychology: Part 1,
      ↪ Initialization
      ### By Dan Jang
      ### Credits to Scott W. Harden for pyhh package [1]

      class Squidward:
          def __init__(self):
```

```

        self.robert, self.gNa, self.gK, self.gL, self.Cm, self.ENa, self.EK,
↪self.EL, self.I, self.Vm, self.m, self.h, self.n, self.t, self.pulse, self.
↪pulsex, self.pulsey, self.movement, self.playtime = None, 120.0, 36.0, 0.3,
↪1.0, 50.0, -77.0, -54.387, 10.0, -65.0, 0.052, 0.596, 0.317, 50, 7000,
↪13000, None, 0.01, np.zeros(20000)
        self.timestep = 0.01
        self.robert = Robert()

    # Essentially, a funny named new-custom constructor
    def tentacles(self, robert, gNa, gK, gL, Cm, ENa, EK, EL, I, Vm, m, h, n,
↪t, pulse, pulsex, pulsey, movement, timestep):

        if robert is None:
            if self.robert is None:
                self.robert = Robert()

        else:
            try:
                self.robert.activation(robert, gNa, gK, gL, Cm, ENa, EK, EL, I,
↪Vm, m, h, n, t, pulse, pulsex, pulsey)
            except:
                print("[E1<Squid-Init>] bruh u might have missed a neuronal
↪parameter for rob lol")

        self.playtime = np.zeros(t) #self.robert.t
        self.playtime[pulsex:pulsey] = pulse

        #if clarinet is None:
        #if self.clarinet is None:
        if timestep is None:
            timestep = 0.01
            self.timestep = timestep
        else:
            self.timestep = timestep
        if movement is None:
            self.movement = hodhux.Simulation(self.robert.model)
            self.clarinet(self.timestep, self.movement)
        else:
            self.clarinet(self.timestep, movement)
            #self.clarinet = clarinet(t, timestep, )

        #self.playtime = self.robert.t

    def clarinet(self, timestep, movement):
        # if playtime is None:
        #     playtime = np.zeros(20000)

```

```

    if timestep is None:
        timestep = 0.01

    if movement is None:
        self.movement = hodhux.Simulation(self.robert.model)
        self.movement.Run(stimulusWaveform=self.playtime,
↪stepSizeMs=timestep)
        self.movement = movement
        #print("[E4<Squid-Clarinet>] bruh u gotta pass a movement thingy so
↪u can actually graph the results u silly head lol")

    else:
        # try:
            movement.Run(stimulusWaveform=self.playtime, stepSizeMs=timestep)
            self.movement = movement
        # except:
            # print("[E3<Squid-Clarinet>] bruh squidward's robert motor
↪neuron model might be misconfigured lol")

    def sheetmusic(self, figsizex, figsizey, color1, color2, color3, color4,
↪color5, color6, color7, color8, title, file):
        #try:
            if figsizex:
                if figsizey:
                    plot.figure(figsize=(figsizex, figsizey))
                else:
                    plot.figure(figsize=(10, 8))
            # except:
                # print("[E4<Squid-SheetMusic>] bruh u might have missed the figure
↪sizes for the sheet music lol")

        if title is None:
            title = "PSY 452 (Winter 2024) - Hodgkin-Huxley Neuromorphic
↪Graphical Model"

        else:
            title = "PSY 452 (Winter 2024) - Hodgkin-Huxley Neuromorphic
↪Graphical Model: " + title

        if self.movement is None:
            print("[E4<Squid-SheetMusic>] bruh squidward apparently hasnt moved
↪his clarinet or if he did, it wasn't recorded u silly goose lol")

        if self.robert is None:

```

```

        print("[E5<Squid-SheetMusic>] bruh somehow at this stage, Robert_
↳the Motor Neuron still has not been created yet, you are one silly goose dan_
↳lmao")

    else:
        movement = self.movement # lazy but saves typing self. each time_
↳lmao

        # Potential (mV) Color
        if color1 is None:
            color1 = 'r'

        # Input Stimulation ( $\mu A/cm^2$ ) Color
        if color2 is None:
            color2 = 'b'

        # Sodium Activation Gating Variable (m) Color
        if color3 is None:
            color3 = 'purple'

        # Sodium Inactivation Gating Variable (h) Color
        if color4 is None:
            color4 = 'orange'

        # Potassium Activation Gating Variable (n) Color
        if color5 is None:
            color5 = 'yellow' # BANANA

        # Sodium Current ( $I_{Na}$ ) ( $\mu A/cm^2$ ) Color
        if color6 is None:
            color6 = 'purple'

        # Potassium Current ( $I_K$ ) ( $\mu A/cm^2$ ) Color
        if color7 is None:
            color7 = 'yellow'

        # Leak Current ( $I_L$ ) ( $\mu A/cm^2$ ) Color
        if color8 is None:
            color8 = 'turquoise'

    ## Credits to [2] for parts from README.md Example Code

    # Potential (mV) Plot #1
    potentialplt = plot.subplot(411)
    potentialplt.set_title(title)
    potentialplt.set_ylabel("Rob's Membrane Potential (mV)")
    potentialplt.plot(movement.times, movement.Vm - 70, color=color1)

```



```

# Input Stimulation ( $\mu\text{A}/\text{cm}^2$ ) Plot #2
stimulationplt = plot.subplot(412)
stimulationplt.set_ylabel("Squidward Stimulation ( $\mu\text{A}/\text{cm}^2$ )")
playtime = self.playtime
## Sanity Debug Print Statement
#print(movement.times, playtime, color2)
stimulationplt.plot(movement.times, playtime, color=color2)

# Gating Variable Activation (m/h/n) Plot #3
activationplt = plot.subplot(413, sharex=potentialplt)
activationplt.set_ylabel("Rob's Channel Activation (fractional)")
activationplt.plot(movement.times, movement.StateM, label='m (Na+
↪Channel Open Prob.)', color=color3)
activationplt.plot(movement.times, movement.StateH, label='h (Na+
↪Channel Close Prob.)', color=color4)
activationplt.plot(movement.times, movement.StateN, label='n (K+
↪Channel Open Prob.)', color=color5)
activationplt.legend()

# Ionic Channel Current Plot (INa/IK/Ileak) Plot #4
currentplt = plot.subplot(414, sharex=potentialplt)
currentplt.set_xlabel("Clarinet Action Potential Time (in
↪milliseconds)")
currentplt.set_ylabel("Rob Membrane Ionic Current ( $\mu\text{A}/\text{cm}^2$ )")
currentplt.plot(movement.times, movement.INa, label='INa (Sodium
↪Current)', color=color6)
currentplt.plot(movement.times, movement.IK, label='IK (Potassium
↪Current)', color=color7)
currentplt.plot(movement.times, movement.IKleak, label='IKLeak
↪(KLeakage Current)', color=color8)
currentplt.legend()

plot.tight_layout()
plot.savefig(file)
plot.show()

class Robert:
    def __init__(self):
        self.model = hodhux.HHModel()
        self.t = None

    def activation(self, model, gNa, gK, gL, Cm, ENa, EK, EL, I, Vm, m, h,
↪n, t, pulse, pulsex, pulsey):

        if model is None:
            if self.model is None:

```

```

robert = hodhux.HHModel()

# Squidward's Rob Motor Neuron Parameter, gNa: Avg. Na+
→(Sodium cation) conductance per unit area (mS/cm2)
robert.gNa = 120

# Squidward's Rob Motor Neuron Parameter, gK: Avg. K+
→(Potassium cation) conductance per unit area (mS/cm2)
robert.gK = 36

# Squidward's Rob Motor Neuron Parameter, gL: Avg. Leak (e.
→g., Chlorine anions / Calcium {2+} cations) conductance per unit area (mS/
→cm2)
robert.gKleak = 0.3

# Squidward's Rob Motor Neuron Parameter, Cm: Membrane
→capacitance per unit area (uF/cm2)
robert.Cm = 1.0

# Squidward's Rob Motor Neuron Parameter, ENa: Sodium
→cation Nernst reversal potential (mV)
robert.ENa = 115

# Squidward's Rob Motor Neuron Parameter, EK: Potassium
→cation Nernst reversal potential (mV)
robert.EK = -12

# Squidward's Rob Motor Neuron Parameter, EL: Leak Nernst
→reversal potential (mV)
robert.EKleak = 10.6

# Squidward's Rob Motor Neuron Parameter, I: External
→current (uA/cm2)
#robert.I = 10.0

# Squidward's Rob Motor Neuron Parameter, Vm: Membrane
→potential (mV)
#robert.Vm = 0

# Squidward's Rob Motor Neuron Parameter, m: Sodium cation
→activation gating variable
#robert.m = 0

# Squidward's Rob Motor Neuron Parameter, h: Sodium cation
→inactivation gating variable
#robert.h = 0

```

```

        # Squidward's Rob Motor Neuron Parameter, n: Potassium
        cation activation gating variable
        #robert.n = 0

        # Stimulus Waveform thingy
        self.t = np.zeros(20000)

        # Stimulus Waveform <+ Square Pulse
        self.t[7000:13000] = 50

        self.model = robert
    try:

        robert = self.model

        if gNa is not None:
            robert.gNa = gNa

        if gK is not None:
            robert.gK = gK

        if gL is not None:
            robert.gKleak = gL

        if Cm is not None:
            robert.Cm = Cm

        if ENa is not None:
            robert.ENa = ENa

        if EK is not None:
            robert.EK = EK

        if EL is not None:
            robert.EKleak = EL

        # if I is not None:
        #     robert.I = I

        # if Vm is not None:
        #     robert.Vm = Vm

        # if m is not None:
        #     robert.m = m

        # if h is not None:

```

```

        #      robert.h = h

        # if n is not None:
        #      robert.n = n

        if t is not None:
            # Stimulus Waveform thingy
            self.t = np.zeros(t)

        if pulse is not None:
            if pulsex is not None:
                if pulsey is not None:
                    self.t[pulsex:pulsey] = pulse

        self.model = robert
    except:
        print("[E2<SubSquid-Rob-Init>] bruh u might have missed a
↳neuronal parameter for rob within squidward initialization thingy lol")

    def model(self):
        return self.model

### Finally, the grand test for class Squidward (main wrapper), with a subclass
↳Robert (le Motor Neuron model encompassing wrapper)

# Reference - Squidward(self, robert, gNa, gK, gL, Cm, ENa, EK, EL, I, Vm, m,
↳h, n, t, pulse, pulsex, pulsey, movement, timestep)
squidward = Squidward()

# tentacles(self, robert, gNa, gK, gL, Cm, ENa, EK, EL, I, Vm, m, h, n, t,
↳pulse, pulsex, pulsey, movement, timestep):

squidward.tentacles(None, 120.0, 36.0, 0.3, 1.0, 115, -12, 10.6, 0, 0, 0, 0, 0,
↳500, 0, 0, 0, None, 0.01)

# Reference - sheetmusic(self, figsizex, figsizey, color1, color2, color3,
↳color4, color5, color6, color7, color8, title)
#      movement = self.movement # lazy but saves typing self. each time
↳lmao

# # Potential (mV) Color
#      color1 = 'r'
# # Input Stimulation ( $\mu\text{A}/\text{cm}^2$ ) Color
#      color2 = 'b'
# # Sodium Activation Gating Variable (m) Color
#      color3 = 'purple'
# # Sodium Inactivation Gating Variable (h) Color

```

```

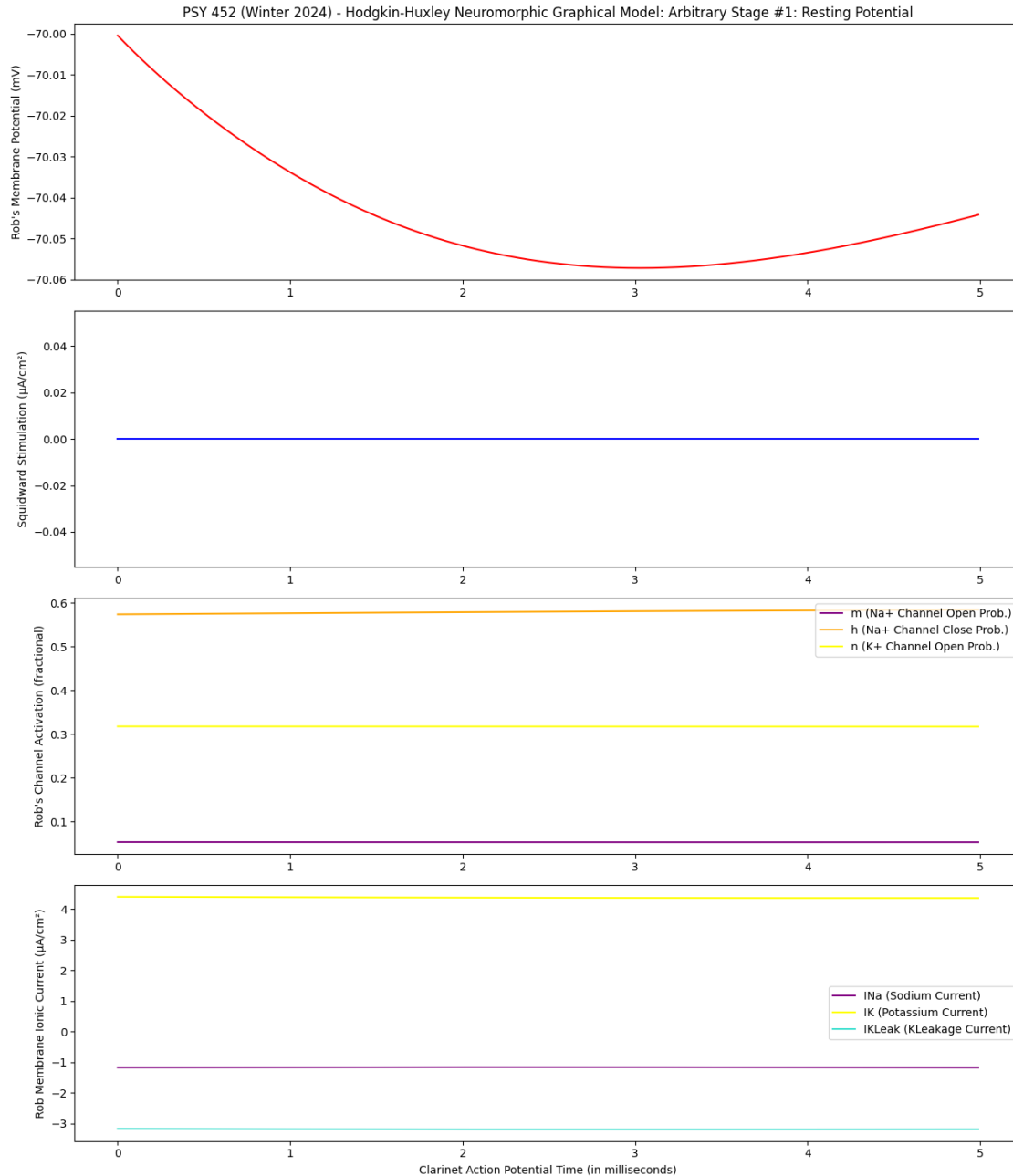
#     color4 = 'orange'
# # Potassium Activation Gating Variable (n) Color
#     color5 = 'yellow' # BANANA
# # Sodium Current (INa) ( $\mu\text{A}/\text{cm}^2$ ) Color
#     color6 = 'purple'
# # Potassium Current (IK) ( $\mu\text{A}/\text{cm}^2$ ) Color
#     color7 = 'yellow'
# # Leak Current (IL) ( $\mu\text{A}/\text{cm}^2$ ) Color
#     color8 = 'turquoise'

squidward.sheetmusic(13, 15, 'r', 'b', 'purple', 'orange', 'yellow', 'purple',
    ↪ 'yellow', 'turquoise', "Arbitrary Stage #1: Resting Potential",
    ↪ "PSY452-Phase1-SquidwardClarinetActionPotentialStimulationWaveform-DanJ.png")

```

Simulating 500 time points...

Simulation complete!



1.1.6 Part 2: *Depolarization / Beginning of Action Potential*

```
[3]: ##### PSY 452 (Winter 2024) - Advanced Neurophysiological Psychology: Part 2,
      ↳ Depolarization / Beginning of Action Potential
      ### By Dan Jang
      ### Credits to Scott W. Harden for pyhh package [1]

      class Squidward:
```

```

def __init__(self):
    self.robert, self.gNa, self.gK, self.gL, self.Cm, self.ENa, self.EK,
    ↪self.EL, self.I, self.Vm, self.m, self.h, self.n, self.t, self.pulse, self.
    ↪pulsex, self.pulsey, self.movement, self.playtime = None, 120.0, 36.0, 0.3,
    ↪1.0, 50.0, -77.0, -54.387, 10.0, -65.0, 0.052, 0.596, 0.317, 50, 7000,
    ↪13000, None, 0.01, np.zeros(20000)
    self.timestep = 0.01
    self.robert = Robert()

    # Essentially, a funny named new-custom constructor
    def tentacles(self, robert, gNa, gK, gL, Cm, ENa, EK, EL, I, Vm, m, h, n,
    ↪t, pulse, pulsex, pulsey, movement, timestep):

        if robert is None:
            if self.robert is None:
                self.robert = Robert()

        else:
            try:
                self.robert.activation(robert, gNa, gK, gL, Cm, ENa, EK, EL, I,
    ↪Vm, m, h, n, t, pulse, pulsex, pulsey)
            except:
                print("[El<Squid-Init>] bruh u might have missed a neuronal
    ↪parameter for rob lol")

        self.playtime = np.zeros(t) #self.robert.t
        self.playtime[pulsex:pulsey] = pulse

        #if clarinet is None:
        #if self.clarinet is None:
        if timestep is None:
            timestep = 0.01
            self.timestep = timestep
        else:
            self.timestep = timestep
        if movement is None:
            self.movement = hodhux.Simulation(self.robert.model)
            self.clarinet(self.timestep, self.movement)
        else:
            self.clarinet(self.timestep, movement)
            #self.clarinet = clarinet(t, timestep, )

        #self.playtime = self.robert.t

    def clarinet(self, timestep, movement):
        # if playtime is None:
        #     playtime = np.zeros(20000)

```

```

    if timestep is None:
        timestep = 0.01

    if movement is None:
        self.movement = hodhux.Simulation(self.robert.model)
        self.movement.Run(stimulusWaveform=self.playtime,
↳stepSizeMs=timestep)
        self.movement = movement
        #print("[E4<Squid-Clarinet>] bruh u gotta pass a movement thingy so
↳u can actually graph the results u silly head lol")

    else:
        # try:
            movement.Run(stimulusWaveform=self.playtime, stepSizeMs=timestep)
            self.movement = movement
        # except:
            print("[E3<Squid-Clarinet>] bruh squidward's robert motor
↳neuron model might be misconfigured lol")

    def sheetmusic(self, figsizex, figsizey, color1, color2, color3, color4,
↳color5, color6, color7, color8, title, file):
        #try:
            if figsizex:
                if figsizey:
                    plot.figure(figsize=(figsizex, figsizey))
                else:
                    plot.figure(figsize=(10, 8))
            # except:
            # print("[E4<Squid-SheetMusic>] bruh u might have missed the figure
↳sizes for the sheet music lol")

        if title is None:
            title = "PSY 452 (Winter 2024) - Hodgkin-Huxley Neuromorphic
↳Graphical Model"

        else:
            title = "PSY 452 (Winter 2024) - Hodgkin-Huxley Neuromorphic
↳Graphical Model: " + title

        if self.movement is None:
            print("[E4<Squid-SheetMusic>] bruh squidward apparently hasnt moved
↳his clarinet or if he did, it wasn't recorded u silly goose lol")

        if self.robert is None:

```



```

        print("[E5<Squid-SheetMusic>] bruh somehow at this stage, Robert_
↳the Motor Neuron still has not been created yet, you are one silly goose dan_
↳lmao")

    else:
        movement = self.movement # lazy but saves typing self. each time_
↳lmao

        # Potential (mV) Color
        if color1 is None:
            color1 = 'r'

        # Input Stimulation ( $\mu A/cm^2$ ) Color
        if color2 is None:
            color2 = 'b'

        # Sodium Activation Gating Variable (m) Color
        if color3 is None:
            color3 = 'purple'

        # Sodium Inactivation Gating Variable (h) Color
        if color4 is None:
            color4 = 'orange'

        # Potassium Activation Gating Variable (n) Color
        if color5 is None:
            color5 = 'yellow' # BANANA

        # Sodium Current ( $I_{Na}$ ) ( $\mu A/cm^2$ ) Color
        if color6 is None:
            color6 = 'purple'

        # Potassium Current ( $I_K$ ) ( $\mu A/cm^2$ ) Color
        if color7 is None:
            color7 = 'yellow'

        # Leak Current ( $I_L$ ) ( $\mu A/cm^2$ ) Color
        if color8 is None:
            color8 = 'turquoise'

    ## Credits to [2] for parts from README.md Example Code

    # Potential (mV) Plot #1
    potentialplt = plot.subplot(411)
    potentialplt.set_title(title)
    potentialplt.set_ylabel("Rob's Membrane Potential (mV)")
    potentialplt.plot(movement.times, movement.Vm - 70, color=color1)

```

```

# Input Stimulation ( $\mu\text{A}/\text{cm}^2$ ) Plot #2
stimulationplt = plot.subplot(412)
stimulationplt.set_ylabel("Squidward Stimulation ( $\mu\text{A}/\text{cm}^2$ )")
playtime = self.playtime
## Sanity Debug Print Statement
#print(movement.times, playtime, color2)
stimulationplt.plot(movement.times, playtime, color=color2)

# Gating Variable Activation (m/h/n) Plot #3
activationplt = plot.subplot(413, sharex=potentialplt)
activationplt.set_ylabel("Rob's Channel Activation (fractional)")
activationplt.plot(movement.times, movement.StateM, label='m (Na+
↪Channel Open Prob.)', color=color3)
activationplt.plot(movement.times, movement.StateH, label='h (Na+
↪Channel Close Prob.)', color=color4)
activationplt.plot(movement.times, movement.StateN, label='n (K+
↪Channel Open Prob.)', color=color5)
activationplt.legend()

# Ionic Channel Current Plot (INa/IK/IKleak) Plot #4
currentplt = plot.subplot(414, sharex=potentialplt)
currentplt.set_xlabel("Clarinet Action Potential Time (in
↪milliseconds)")
currentplt.set_ylabel("Rob Membrane Ionic Current ( $\mu\text{A}/\text{cm}^2$ )")
currentplt.plot(movement.times, movement.INa, label='INa (Sodium
↪Current)', color=color6)
currentplt.plot(movement.times, movement.IK, label='IK (Potassium
↪Current)', color=color7)
currentplt.plot(movement.times, movement.IKleak, label='IKLeak
↪(KLeakage Current)', color=color8)
currentplt.legend()

plot.tight_layout()
plot.savefig(file)
plot.show()

class Robert:
    def __init__(self):
        self.model = hodhux.HHModel()
        self.t = None

    def activation(self, model, gNa, gK, gL, Cm, ENa, EK, EL, I, Vm, m, h,
↪n, t, pulse, pulsex, pulsey):

        if model is None:
            if self.model is None:

```

```

robert = hodhux.HHModel()

# Squidward's Rob Motor Neuron Parameter, gNa: Avg. Na+
→ (Sodium cation) conductance per unit area (mS/cm2)
robert.gNa = 120

# Squidward's Rob Motor Neuron Parameter, gK: Avg. K+
→ (Potassium cation) conductance per unit area (mS/cm2)
robert.gK = 36

# Squidward's Rob Motor Neuron Parameter, gL: Avg. Leak (e.
→ g., Chlorine anions / Calcium {2+} cations) conductance per unit area (mS/
→ cm2)
robert.gKleak = 0.3

# Squidward's Rob Motor Neuron Parameter, Cm: Membrane
→ capacitance per unit area (uF/cm2)
robert.Cm = 1.0

# Squidward's Rob Motor Neuron Parameter, ENa: Sodium
→ cation Nernst reversal potential (mV)
robert.ENa = 115

# Squidward's Rob Motor Neuron Parameter, EK: Potassium
→ cation Nernst reversal potential (mV)
robert.EK = -12

# Squidward's Rob Motor Neuron Parameter, EL: Leak Nernst
→ reversal potential (mV)
robert.EKleak = 10.6

# Squidward's Rob Motor Neuron Parameter, I: External
→ current (uA/cm2)
#robert.I = 10.0

# Squidward's Rob Motor Neuron Parameter, Vm: Membrane
→ potential (mV)
#robert.Vm = 0

# Squidward's Rob Motor Neuron Parameter, m: Sodium cation
→ activation gating variable
#robert.m = 0

# Squidward's Rob Motor Neuron Parameter, h: Sodium cation
→ inactivation gating variable
#robert.h = 0

```

```

        # Squidward's Rob Motor Neuron Parameter, n: Potassium
        cation activation gating variable
        #robert.n = 0

        # Stimulus Waveform thingy
        self.t = np.zeros(20000)

        # Stimulus Waveform <+ Square Pulse
        self.t[7000:13000] = 50

        self.model = robert
    try:

        robert = self.model

        if gNa is not None:
            robert.gNa = gNa

        if gK is not None:
            robert.gK = gK

        if gL is not None:
            robert.gKleak = gL

        if Cm is not None:
            robert.Cm = Cm

        if ENa is not None:
            robert.ENa = ENa

        if EK is not None:
            robert.EK = EK

        if EL is not None:
            robert.EKleak = EL

        # if I is not None:
        #     robert.I = I

        # if Vm is not None:
        #     robert.Vm = Vm

        # if m is not None:
        #     robert.m = m

        # if h is not None:

```

```

        #      robert.h = h

        # if n is not None:
        #      robert.n = n

        if t is not None:
            # Stimulus Waveform thingy
            self.t = np.zeros(t)

        if pulse is not None:
            if pulsex is not None:
                if pulsey is not None:
                    self.t[pulsex:pulsey] = pulse

        self.model = robert
    except:
        print("[E2<SubSquid-Rob-Init>] bruh u might have missed a
↳neuronal parameter for rob within squidward initialization thingy lol")

    def model(self):
        return self.model

### Finally, the grand test for class Squidward (main wrapper), with a subclass
↳Robert (le Motor Neuron model encompassing wrapper)

# Reference - Squidward(self, robert, gNa, gK, gL, Cm, ENa, EK, EL, I, Vm, m,
↳h, n, t, pulse, pulsex, pulsey, movement, timestep)
squidward = Squidward()

# tentacles(self, robert, gNa, gK, gL, Cm, ENa, EK, EL, I, Vm, m, h, n, t,
↳pulse, pulsex, pulsey, movement, timestep):

squidward.tentacles(None, 120.0, 36.0, 0.3, 1.0, 115, -12, 10.6, 0, 0, 0, 0, 0,
↳500, 500, 200, 1000, None, 0.01)

# Reference - sheetmusic(self, figsizex, figsizey, color1, color2, color3,
↳color4, color5, color6, color7, color8, title)
#      movement = self.movement # lazy but saves typing self. each time
↳lmao

# # Potential (mV) Color
#      color1 = 'r'
# # Input Stimulation ( $\mu\text{A}/\text{cm}^2$ ) Color
#      color2 = 'b'
# # Sodium Activation Gating Variable (m) Color
#      color3 = 'purple'
# # Sodium Inactivation Gating Variable (h) Color

```

```

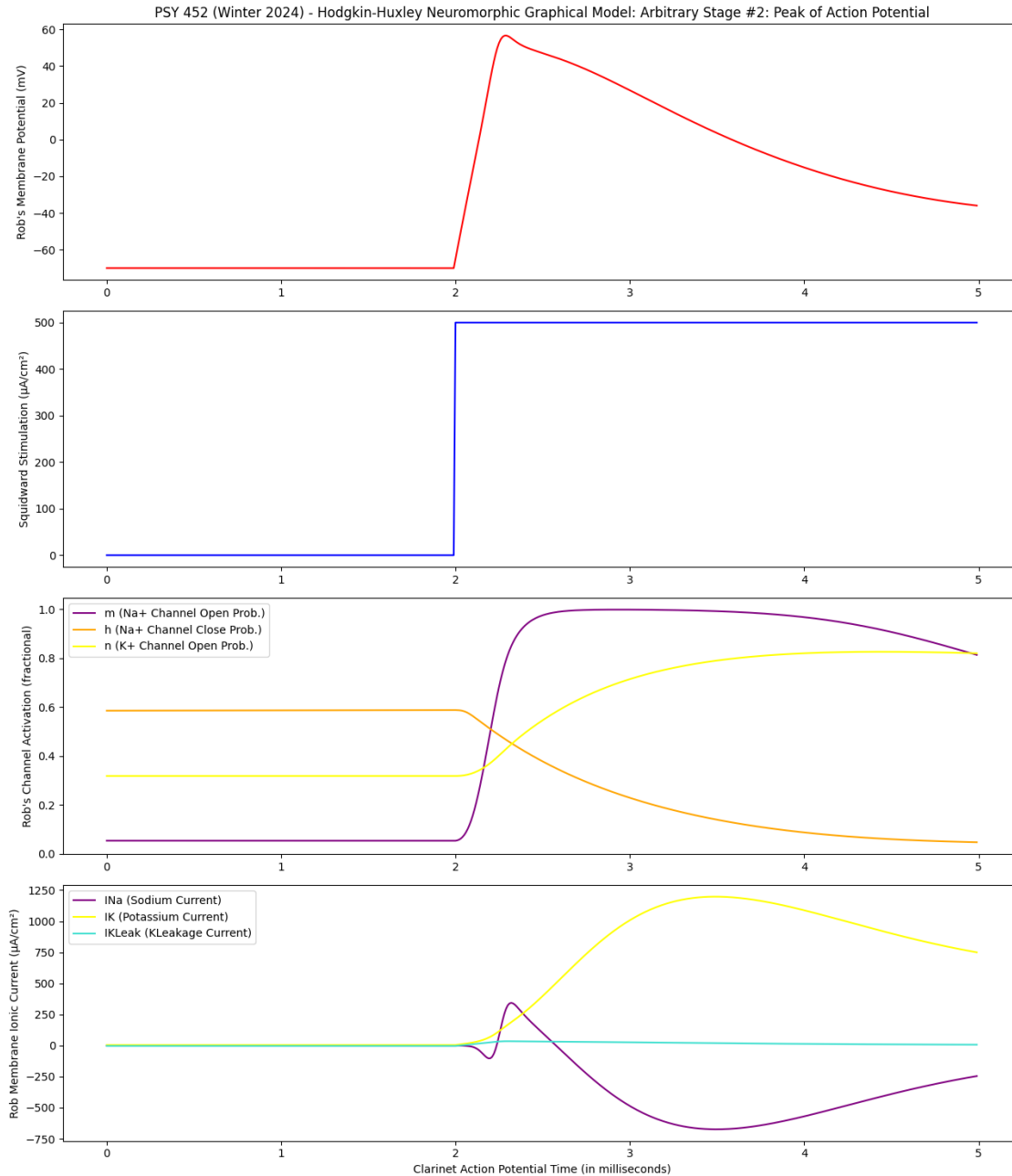
#     color4 = 'orange'
# # Potassium Activation Gating Variable (n) Color
#     color5 = 'yellow' # BANANA
# # Sodium Current (INa) ( $\mu\text{A}/\text{cm}^2$ ) Color
#     color6 = 'purple'
# # Potassium Current (IK) ( $\mu\text{A}/\text{cm}^2$ ) Color
#     color7 = 'yellow'
# # Leak Current (IL) ( $\mu\text{A}/\text{cm}^2$ ) Color
#     color8 = 'turquoise'

squidward.sheetmusic(13, 15, 'r', 'b', 'purple', 'orange', 'yellow', 'purple',
    ↪ 'yellow', 'turquoise', "Arbitrary Stage #2: Peak of Action Potential",
    ↪ "PSY452-Phase2-SquidwardClarinetActionPotentialStimulationWaveform-DanJ.png")

```

Simulating 500 time points...

Simulation complete!



1.1.7 Part 3: Re- / Hyperpolarization to Reach Resting Membrane Potential (RMP)

[4]: ##### PSY 452 (Winter 2024) - Advanced Neurophysiological Psychology: Part 3,
 ↳ Re- / Hyperpolarization to Reach Resting Membrane Potential (RMP)

By Dan Jang

Credits to Scott W. Harden for pyhh package [1]

```
class Squidward:
```

```

def __init__(self):
    self.robert, self.gNa, self.gK, self.gL, self.Cm, self.ENa, self.EK,
    ↪self.EL, self.I, self.Vm, self.m, self.h, self.n, self.t, self.pulse, self.
    ↪pulsex, self.pulsey, self.movement, self.playtime = None, 120.0, 36.0, 0.3,
    ↪1.0, 50.0, -77.0, -54.387, 10.0, -65.0, 0.052, 0.596, 0.317, 50, 7000,
    ↪13000, None, 0.01, np.zeros(20000)
    self.timestep = 0.01
    self.robert = Robert()

    # Essentially, a funny named new-custom constructor
    def tentacles(self, robert, gNa, gK, gL, Cm, ENa, EK, EL, I, Vm, m, h, n,
    ↪t, pulse, pulsex, pulsey, movement, timestep):

        if robert is None:
            if self.robert is None:
                self.robert = Robert()

        else:
            try:
                self.robert.activation(robert, gNa, gK, gL, Cm, ENa, EK, EL, I,
    ↪Vm, m, h, n, t, pulse, pulsex, pulsey)
            except:
                print("[El<Squid-Init>] bruh u might have missed a neuronal
    ↪parameter for rob lol")

        self.playtime = np.zeros(t) #self.robert.t
        self.playtime[pulsex:pulsey] = pulse

        #if clarinet is None:
        #if self.clarinet is None:
        if timestep is None:
            timestep = 0.01
            self.timestep = timestep
        else:
            self.timestep = timestep
        if movement is None:
            self.movement = hodhux.Simulation(self.robert.model)
            self.clarinet(self.timestep, self.movement)
        else:
            self.clarinet(self.timestep, movement)
            #self.clarinet = clarinet(t, timestep, )

        #self.playtime = self.robert.t

    def clarinet(self, timestep, movement):
        # if playtime is None:
        #     playtime = np.zeros(20000)

```



```

        if timestep is None:
            timestep = 0.01

        if movement is None:
            self.movement = hodhux.Simulation(self.robert.model)
            self.movement.Run(stimulusWaveform=self.playtime,
↪stepSizeMs=timestep)
            self.movement = movement
            #print("[E4<Squid-Clarinet>] bruh u gotta pass a movement thingy so
↪u can actually graph the results u silly head lol")

        else:
            # try:
            movement.Run(stimulusWaveform=self.playtime, stepSizeMs=timestep)
            self.movement = movement
            # except:
            #     print("[E3<Squid-Clarinet>] bruh squidward's robert motor
↪neuron model might be misconfigured lol")

    def sheetmusic(self, figsizex, figsizey, color1, color2, color3, color4,
↪color5, color6, color7, color8, title, file):
        #try:
        if figsizex:
            if figsizey:
                plot.figure(figsize=(figsizex, figsizey))
            else:
                plot.figure(figsize=(10, 8))
        # except:
        #     print("[E4<Squid-SheetMusic>] bruh u might have missed the figure
↪sizes for the sheet music lol")

        if title is None:
            title = "PSY 452 (Winter 2024) - Hodgkin-Huxley Neuromorphic
↪Graphical Model"

        else:
            title = "PSY 452 (Winter 2024) - Hodgkin-Huxley Neuromorphic
↪Graphical Model: " + title

        if self.movement is None:
            print("[E4<Squid-SheetMusic>] bruh squidward apparently hasnt moved
↪his clarinet or if he did, it wasn't recorded u silly goose lol")

        if self.robert is None:

```

```

        print("[E5<Squid-SheetMusic>] bruh somehow at this stage, Robert_
↳the Motor Neuron still has not been created yet, you are one silly goose dan_
↳lmao")

    else:
        movement = self.movement # lazy but saves typing self. each time_
↳lmao

        # Potential (mV) Color
        if color1 is None:
            color1 = 'r'

        # Input Stimulation ( $\mu A/cm^2$ ) Color
        if color2 is None:
            color2 = 'b'

        # Sodium Activation Gating Variable (m) Color
        if color3 is None:
            color3 = 'purple'

        # Sodium Inactivation Gating Variable (h) Color
        if color4 is None:
            color4 = 'orange'

        # Potassium Activation Gating Variable (n) Color
        if color5 is None:
            color5 = 'yellow' # BANANA

        # Sodium Current ( $I_{Na}$ ) ( $\mu A/cm^2$ ) Color
        if color6 is None:
            color6 = 'purple'

        # Potassium Current ( $I_K$ ) ( $\mu A/cm^2$ ) Color
        if color7 is None:
            color7 = 'yellow'

        # Leak Current ( $I_L$ ) ( $\mu A/cm^2$ ) Color
        if color8 is None:
            color8 = 'turquoise'

        ## Credits to [2] for parts from README.md Example Code

        # Potential (mV) Plot #1
        potentialplt = plot.subplot(411)
        potentialplt.set_title(title)
        potentialplt.set_ylabel("Rob's Membrane Potential (mV)")
        potentialplt.plot(movement.times, movement.Vm - 70, color=color1)

```

```

# Input Stimulation ( $\mu\text{A}/\text{cm}^2$ ) Plot #2
stimulationplt = plot.subplot(412)
stimulationplt.set_ylabel("Squidward Stimulation ( $\mu\text{A}/\text{cm}^2$ )")
playtime = self.playtime
## Sanity Debug Print Statement
#print(movement.times, playtime, color2)
stimulationplt.plot(movement.times, playtime, color=color2)

# Gating Variable Activation (m/h/n) Plot #3
activationplt = plot.subplot(413, sharex=potentialplt)
activationplt.set_ylabel("Rob's Channel Activation (fractional)")
activationplt.plot(movement.times, movement.StateM, label='m (Na+
↪Channel Open Prob.)', color=color3)
activationplt.plot(movement.times, movement.StateH, label='h (Na+
↪Channel Close Prob.)', color=color4)
activationplt.plot(movement.times, movement.StateN, label='n (K+
↪Channel Open Prob.)', color=color5)
activationplt.legend()

# Ionic Channel Current Plot (INa/IK/IKleak) Plot #4
currentplt = plot.subplot(414, sharex=potentialplt)
currentplt.set_xlabel("Clarinet Action Potential Time (in
↪milliseconds)")
currentplt.set_ylabel("Rob Membrane Ionic Current ( $\mu\text{A}/\text{cm}^2$ )")
currentplt.plot(movement.times, movement.INa, label='INa (Sodium
↪Current)', color=color6)
currentplt.plot(movement.times, movement.IK, label='IK (Potassium
↪Current)', color=color7)
currentplt.plot(movement.times, movement.IKleak, label='IKLeak
↪(KLeakage Current)', color=color8)
currentplt.legend()

plot.tight_layout()
plot.savefig(file)
plot.show()

class Robert:
    def __init__(self):
        self.model = hodhux.HHModel()
        self.t = None

    def activation(self, model, gNa, gK, gL, Cm, ENa, EK, EL, I, Vm, m, h,
↪n, t, pulse, pulsex, pulsey):

        if model is None:
            if self.model is None:

```

```

robert = hodhux.HHModel()

# Squidward's Rob Motor Neuron Parameter, gNa: Avg. Na+
→ (Sodium cation) conductance per unit area (mS/cm2)
robert.gNa = 120

# Squidward's Rob Motor Neuron Parameter, gK: Avg. K+
→ (Potassium cation) conductance per unit area (mS/cm2)
robert.gK = 36

# Squidward's Rob Motor Neuron Parameter, gL: Avg. Leak (e.
→ g., Chlorine anions / Calcium {2+} cations) conductance per unit area (mS/
→ cm2)
robert.gKleak = 0.3

# Squidward's Rob Motor Neuron Parameter, Cm: Membrane
→ capacitance per unit area (uF/cm2)
robert.Cm = 1.0

# Squidward's Rob Motor Neuron Parameter, ENa: Sodium
→ cation Nernst reversal potential (mV)
robert.ENa = 115

# Squidward's Rob Motor Neuron Parameter, EK: Potassium
→ cation Nernst reversal potential (mV)
robert.EK = -12

# Squidward's Rob Motor Neuron Parameter, EL: Leak Nernst
→ reversal potential (mV)
robert.EKleak = 10.6

# Squidward's Rob Motor Neuron Parameter, I: External
→ current (uA/cm2)
#robert.I = 10.0

# Squidward's Rob Motor Neuron Parameter, Vm: Membrane
→ potential (mV)
#robert.Vm = 0

# Squidward's Rob Motor Neuron Parameter, m: Sodium cation
→ activation gating variable
#robert.m = 0

# Squidward's Rob Motor Neuron Parameter, h: Sodium cation
→ inactivation gating variable
#robert.h = 0

```

```

        # Squidward's Rob Motor Neuron Parameter, n: Potassium
        cation activation gating variable
        #robert.n = 0

        # Stimulus Waveform thingy
        self.t = np.zeros(20000)

        # Stimulus Waveform <+ Square Pulse
        self.t[7000:13000] = 50

        self.model = robert
try:

    robert = self.model

    if gNa is not None:
        robert.gNa = gNa

    if gK is not None:
        robert.gK = gK

    if gL is not None:
        robert.gKleak = gL

    if Cm is not None:
        robert.Cm = Cm

    if ENa is not None:
        robert.ENa = ENa

    if EK is not None:
        robert.EK = EK

    if EL is not None:
        robert.EKleak = EL

    # if I is not None:
    #     robert.I = I

    # if Vm is not None:
    #     robert.Vm = Vm

    # if m is not None:
    #     robert.m = m

    # if h is not None:

```

```

        #      robert.h = h

        # if n is not None:
        #      robert.n = n

        if t is not None:
            # Stimulus Waveform thingy
            self.t = np.zeros(t)

        if pulse is not None:
            if pulsex is not None:
                if pulsey is not None:
                    self.t[pulsex:pulsey] = pulse

        self.model = robert
    except:
        print("[E2<SubSquid-Rob-Init>] bruh u might have missed a
↳neuronal parameter for rob within squidward initialization thingy lol")

    def model(self):
        return self.model

### Finally, the grand test for class Squidward (main wrapper), with a subclass
↳Robert (le Motor Neuron model encompassing wrapper)

# Reference - Squidward(self, robert, gNa, gK, gL, Cm, ENa, EK, EL, I, Vm, m,
↳h, n, t, pulse, pulsex, pulsey, movement, timestep)
squidward = Squidward()

# tentacles(self, robert, gNa, gK, gL, Cm, ENa, EK, EL, I, Vm, m, h, n, t,
↳pulse, pulsex, pulsey, movement, timestep):

squidward.tentacles(None, 120.0, 36.0, 0.3, 1.0, 115, -12, 10.6, 0, 0, 0, 0, 0,
↳500, 500, 0, 500, None, 0.01)

# Reference - sheetmusic(self, figsizex, figsizey, color1, color2, color3,
↳color4, color5, color6, color7, color8, title)
#      movement = self.movement # lazy but saves typing self. each time
↳lmao

# # Potential (mV) Color
#      color1 = 'r'
# # Input Stimulation ( $\mu\text{A}/\text{cm}^2$ ) Color
#      color2 = 'b'
# # Sodium Activation Gating Variable (m) Color
#      color3 = 'purple'
# # Sodium Inactivation Gating Variable (h) Color

```

```

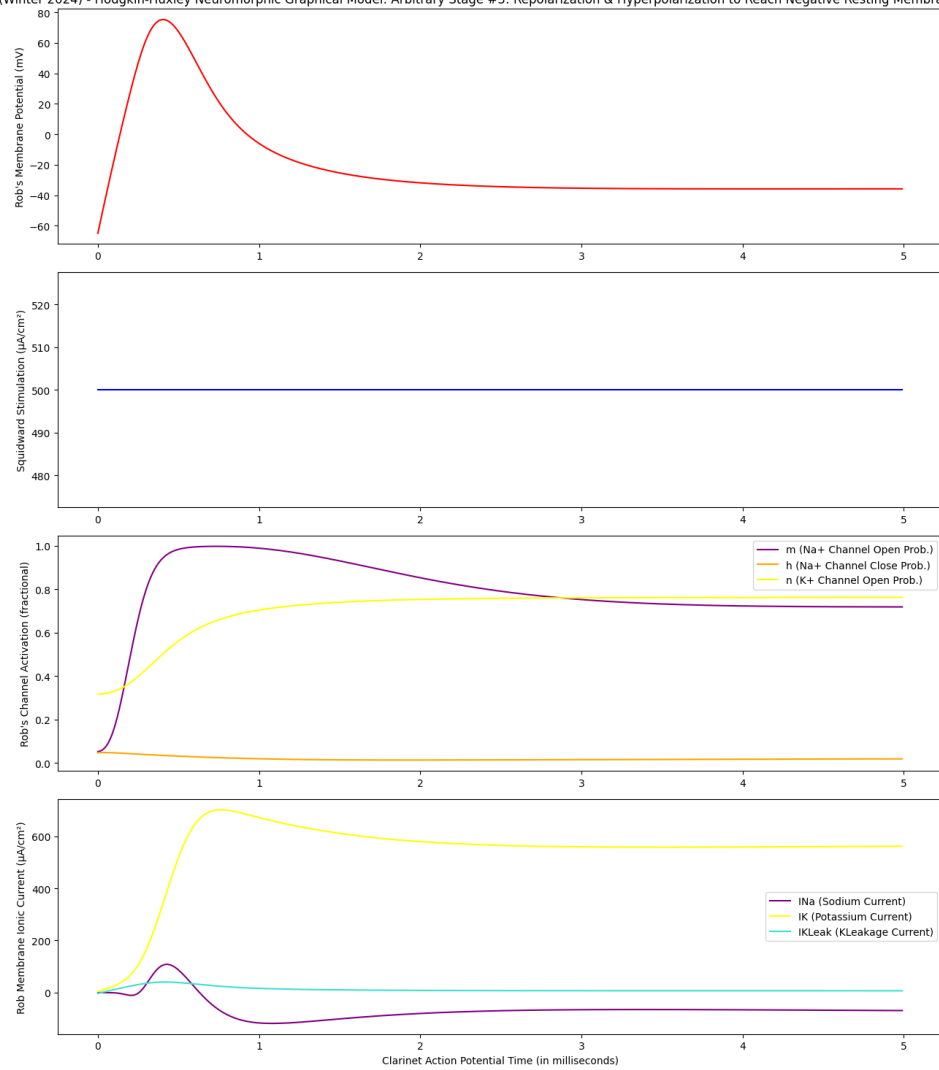
#     color4 = 'orange'
# # Potassium Activation Gating Variable (n) Color
#     color5 = 'yellow' # BANANA
# # Sodium Current (INa) ( $\mu\text{A}/\text{cm}^2$ ) Color
#     color6 = 'purple'
# # Potassium Current (IK) ( $\mu\text{A}/\text{cm}^2$ ) Color
#     color7 = 'yellow'
# # Leak Current (IL) ( $\mu\text{A}/\text{cm}^2$ ) Color
#     color8 = 'turquoise'

squidward.sheetmusic(13, 15, 'r', 'b', 'purple', 'orange', 'yellow', 'purple',
    ↪ 'yellow', 'turquoise', "Arbitrary Stage #3: Repolarization &
    ↪ Hyperpolarization to Reach Negative Resting Membrane Potential (RMP)",
    ↪ "PSY452-Phase3-SquidwardClarinetActionPotentialStimulationWaveform-DanJ.png")

```

Simulating 500 time points...

Simulation complete!



[4]

1.2 References

- [1] S. W. Harden, “swharden/pyHH.” Feb. 11, 2024. [Online]. Available: <https://github.com/swharden/pyHH>
- [2] A. L. Hodgkin and A. F. Huxley, “A quantitative description of membrane current and its application to conduction and excitation in nerve,” *J Physiol*, vol. 117, no. 4, pp. 500–544, Aug. 1952. [Online]. Available: <https://www.ncbi.nlm.nih.gov/pmc/articles/PMC1392413/>
- [3] J. Żygierewicz, “Exercise 1: The Hodgkin Huxley model.” [Online]. Available: https://www.fuw.edu.pl/~jarekz/MODELOWANIE/Hh/HH_exercises.tex
- [4] C. J. Schwiening, “A brief historical perspective: Hodgkin and Huxley,” *J Physiol*, vol. 590, no. Pt 11, pp. 2571–2575, Jun. 2012, doi: 10.1113/jphysiol.2012.230458.