Worked with: Joe Thurakal

```
In [1]: import numpy as np
from matplotlib import pyplot as plt
%matplotlib inline
```

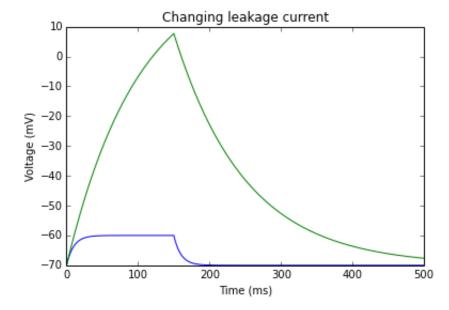
1. Membrane Model

(a)

```
In [2]: t = 500.
    N = 5000
    dt = t/N
    v_r = -70.
    c_m = 100
    i_in = lambda tt: 100. if tt<150. else 0.
    v_t = np.zeros(shape=(2,N))
    v_t[:,0] = v_r
    time = np.linspace(0,t, N)</pre>
```

```
In [5]: plt.plot(time, v_t[0], time, v_t[1])
   plt.title('Changing leakage current')
   plt.ylabel('Voltage (mV)')
   plt.xlabel('Time (ms)')
```

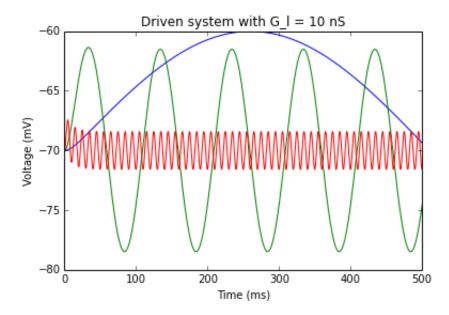
Out[5]: <matplotlib.text.Text at 0x10c671790>



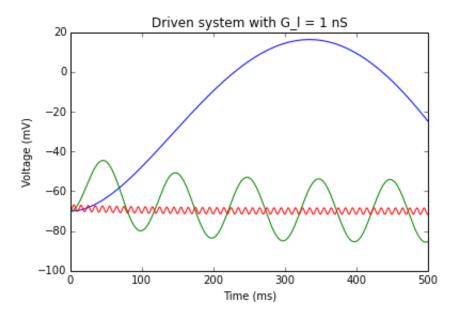
(b)

```
In [6]: t = 500.
    N = 5000
    dt = t/N
    i_in2 = lambda tt, f: 100.*np.sin(2.*np.pi*f*tt/1000.)
    v_t2 = np.zeros(shape=(2,3,N))
    v_t2[...,0] = v_r
    time = np.linspace(0,t, N)
    fs = [1,10,100]
```

Out[7]: <matplotlib.text.Text at 0x10c660f50>

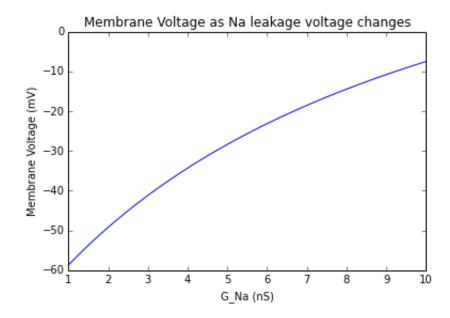


Out[8]: <matplotlib.text.Text at 0x10c6b9890>

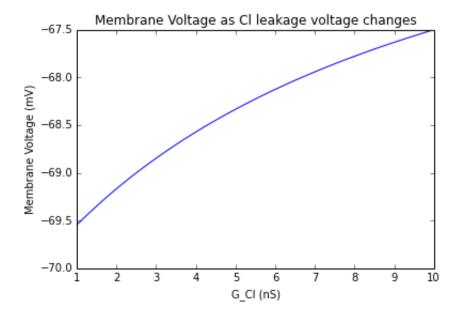


2. Shunting Inhibition

Out[9]: <matplotlib.text.Text at 0x10d118890>

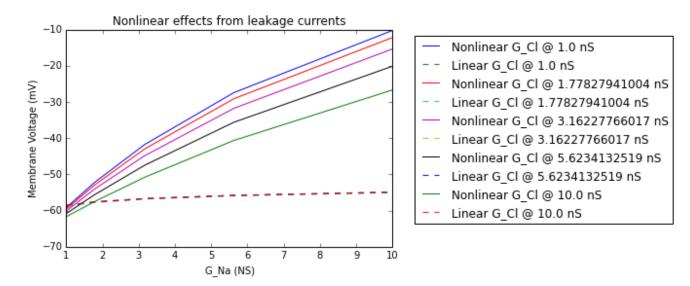


Out[10]: <matplotlib.text.Text at 0x10d37f190>



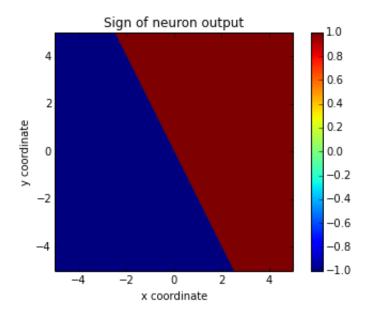
```
In [11]:
         N=5
         def V all(G Na, G Cl):
             return (-70.*10.-65.*G_Cl+55.*G_Na)/(10.+G_Cl+G_Na)
         g na = np.logspace(0,1,N)
         g_cl = np.logspace(0,1,N)
         v = np.zeros(N)
         v2 = np.zeros(N)
         for jj in xrange(N):
             for ii in xrange(N):
                 v[ii] = V all(g na[ii],g cl[jj])
                 v2[ii] = v_na[ii]
             plt.plot(g_na, v, label='Nonlinear G_Cl @ '+str(g_cl[jj])+' nS')
             plt.plot(g_na,v2+v_cl[jj]-v_cl[0],'--',
                       label='Linear G_Cl @ '+str(g_cl[jj])+' nS')
         plt.legend(bbox to anchor=(1.05,1),loc=2)
         plt.xlabel('G_Na (NS)')
         plt.ylabel('Membrane Voltage (mV)')
         plt.title('Nonlinear effects from leakage currents')
```

Out[11]: <matplotlib.text.Text at 0x10d4ae950>



3. Linear Neuron

Out[15]: <matplotlib.text.Text at 0x11563ca10>

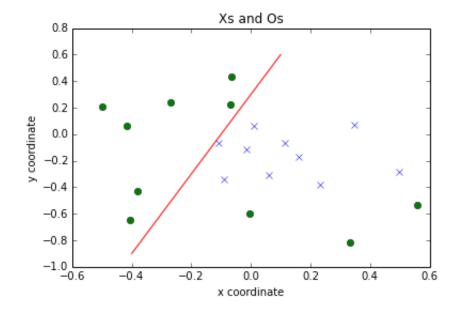


4. Pattern Discrimination

```
In [13]: from scipy.io import loadmat
  data = loadmat('data.mat')
  X = data['X'].T
  O = data['O'].T
```

```
In [14]: plt.plot(X[0],X[1],'x',O[0],O[1],'o')
    plt.title('Xs and Os')
    x = np.linspace(-.4, .1,100)
    y = 3.*x+.3
    plt.plot(x,y)
    plt.xlabel('x coordinate')
    plt.ylabel('y coordinate')
```

Out[14]: <matplotlib.text.Text at 0x115534f50>



A hand-tuned kernel that could be used would be the distance from the center of the x's.

More generally, one could take combinations of (x^ny^m) .