hw1

September 11, 2017

1 CSCI 5992 - HW1 - Kelly/Milne

1.1 Part 1

1.1.1 Solution

```
1a) w1 = -2.04427331 \text{ } w2 = 3.99683416 \text{ } b = -0.92427055
   1b) Levenberg-Marquardt as per implementation below
In [1]: #%%
        import os
        import numpy as np
        import scipy.io
        import scipy.optimize as optimization
        #from IPython.core.debugger import set_trace
        # Data
        filedir = os.path.dirname(os.path.realpath('__file__'))
        datapath = os.path.join(filedir, 'assign1_data.mat')
        data = scipy.io.loadmat(datapath)
        # fitter wants shape (k,M) (k number of predictors)
        # data['x'] has shape (M,k) so we fix that
        x1 = data['x'][:,0]
        x2 = data['x'][:,1]
        X = np.array([x1,x2])
        # argg - shape (100,1) - gives obstuse error in the fitter
        # we fix that too (flatten)
        y = np.array(data['y']).flatten()
        # Initial guess
        w0 = np.array([1, 1, 1])
        # Objective function
        # y = w1 * x1 + w2 * x2 + b.
        def func(X, b, w1, w2):
          # unpack independent vars
          rval = w1*X[0] + w2*X[1] + b
```

```
return rval

result = optimization.curve_fit(func, X, y, w0)
print(result[0])

[-0.92427055 -2.04427331 3.99683416]
```

1.2 Part 2

1.2.1 Solution

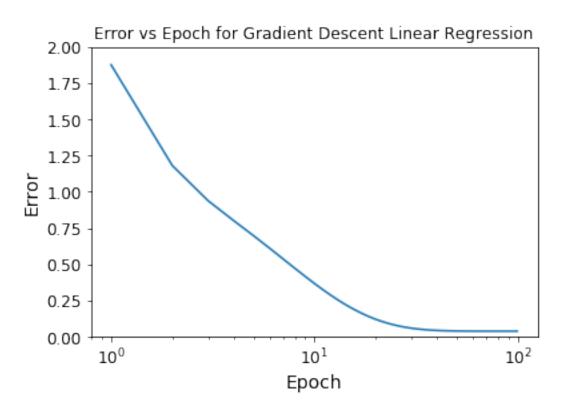
```
2a) w1 = -2.0427036 w2 = 3.99155586 b = -0.92233149
```

2b) Batch processed. A step size of 0.05 diverged so used 0.01. Approx N=100 epochs to recover the results from the LM method. Termination was based on plot below of error vs epoch, which amounts to a % change in the error function of \sim 10 $^{\circ}$ 4 between epochs.

```
In [2]: import os
        import numpy as np
        import scipy.io
        import scipy.optimize as optimization
        #from IPython.core.debugger import set_trace
        EPS = 0.01
        N EPOCH = 100
        err = []
        def dw(w,d,X):
            # using numpy einstein summation to vectorize the computation
            # vector of coefficients (d-w.x) for each pattern (alpha)
            C = d - np.einsum('i,ij->j', w, X) # shape is (100,)
            e = 1.0/len(C)*np.sum([c**2 for c in C])
            err.append(e)
            \# C_i * X_j i \text{ or } C * transpose(X)
            dw = EPS*np.einsum('i,ji->j', C, X) # shape is (3,)
            return dw
        def main():
            # data
            filedir = os.path.dirname(os.path.realpath('__file__'))
            datapath = os.path.join(filedir, 'assign1_data.mat')
            data = scipy.io.loadmat(datapath)
            x1 = data['x'][:,0]
            x2 = data['x'][:,1]
            # inputs - with bias input tied high, shape is (3,100)
            X = np.array([np.full(len(x1),1),x1,x2])
            # output
            d = np.array(data['y']).flatten()
            # starting vector of weights
            w = [1, 1, 1]
```

Out[3]: <matplotlib.text.Text at 0x115214278>

ax = plt.subplot(111)
plt.semilogx(err)
ax.set_ylim([0, 2])
plt.ylabel('Error')
plt.xlabel('Epoch')



plt.title('Error vs Epoch for Gradient Descent Linear Regression')

1.3 Part 3

1.3.1 Solution

```
3a) b = -16. w1 = -50.43661604 w2 = 81.86640442
   3b) See plot in 'Visualization of Perceptron Results'
In [4]: import os as os
        import numpy as np
        import scipy.io
        import scipy.optimize as optimization
        from IPython.core.debugger import set_trace
        N_EPOCH = 30
        n mis = []
        def dwp(w,d,X):
             # using numpy einstein summation to vectorize the computation
             \# calculating w.x.d -- w.x.d < 0 \Rightarrow incorrect classification
             C = d*np.einsum('i,ij->j', w, X) # shape is (100,)
            XT = X.transpose()
             # use enumerate for the equivalent to each_with_index (Ruby)
            xd = [ d[i]*XT[i] for i,c in enumerate(C) if c < 0]</pre>
             # for N vs EPOCH plot
            n_mis.append(len(xd))
             # sum xd element-wise
             \# dw is x.t summed over misclassified teachers
             dw = np.einsum('ij->j',xd) if(len(xd) > 0) else 0
             return dw
        def main():
             \# d.a.t.a.
            filedir = os.path.dirname(os.path.realpath('__file__'))
             datapath = os.path.join(filedir, 'assign1_data.mat')
             data = scipy.io.loadmat(datapath)
            x1 = data['x'][:,0]
            x2 = data['x'][:,1]
             # inputs - with bias input tied high, shape is (3,100)
            X = np.array([np.full(len(x1),1),x1,x2])
             # output
             d = np.array(data['z']).flatten()
             # use domain -1,1 for teachers so we can use x.w.d<0 for the classification test
             d = [z \text{ if } z == 1 \text{ else } -1 \text{ for } z \text{ in } d]
             # starting vector of weights
            w = [1, 1, 1]
             \# sweep through the data N_EPOCH times
```

2 Visualization of perceptron results

Read the data into a panda dataframe for visualization

```
In [5]: # Common imports
        import numpy as np
        import os
        # to make this notebook's output stable across runs
        np.random.seed(42)
        # To plot pretty figures
        %matplotlib inline
        import matplotlib
        import matplotlib.pyplot as plt
        plt.rcParams['axes.labelsize'] = 14
        plt.rcParams['xtick.labelsize'] = 12
        plt.rcParams['ytick.labelsize'] = 12
        # Where to save the figures
        PROJECT_ROOT_DIR = "."
        import pandas as pd
        def load_data():
            csv_path = os.path.join(".", "assign1_data.txt")
            return pd.read_csv(csv_path, sep="\s+")
        def load_data():
            csv_path = os.path.join(".", "assign1_data.txt")
            return pd.read_csv(csv_path, sep="\s+")
        data = load_data()
        data.head()
        c0 = data[(data.z==0)]
        c1 = data[(data.z==1)]
```

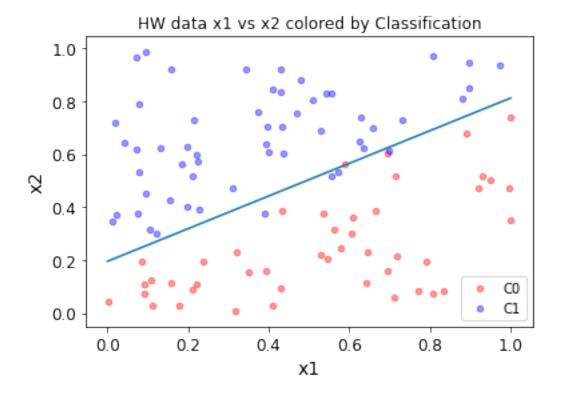
2.0.1 HW data in parameter space with classification indicated by color

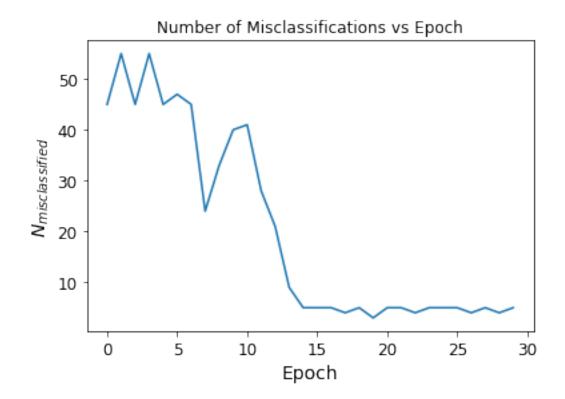
The plotted line is the surface of separation given by x2 = -1/w2(w1x1 + w0) where w is the weight vector calculated using perceptron algorithm. As is indicated in the figure the data is not linearly separable so the perceptron algorithm does not converge, instead it jitters around near the minimum.

```
In [6]: X = np.linspace(0, 1, 128, endpoint=True)
    W = w_perceptron
    L = -(1/W[2])*(W[1]*X + W[0])

ax = c0.plot.scatter(x="x1", y="x2", color='Red', alpha=0.4, label='C0')
    c1.plot.scatter( x="x1", y="x2", color='Blue', alpha=0.4, label='C1', ax=ax)
    ax.plot(X,L)
    plt.title('HW data x1 vs x2 colored by Classification')
```

Out[6]: <matplotlib.text.Text at 0x1169c1a58>





3 Part 4

Plot of performance vs number of training samples below.

```
In [8]: import os as os
        import numpy as np
        import scipy.io
        import scipy.optimize as optimization
        N_EPOCH = 20
        n_mis = []
        def dwp(w,d,X):
            # using numpy einstein summation to vectorize the computation
            \# calculating w.x.d -- w.x.d < 0 => incorrect classification
            C = d*np.einsum('i,ij->j', w, X) # shape is (100,)
            XT = np.einsum('ij->ji', X)
            # use enumerate for the equivalent to each_with_index (Ruby)
            xd = [ d[i]*XT[i] for i,c in enumerate(C) if c < 0]</pre>
            # sum xd element-wise
            # dw is x.t summed over misclassified teachers
            dw = np.einsum('ij->j',xd) if(len(xd) > 0) else 0 # shape is (3,)
```

```
return dw
        def train(n, X, d):
            # use n_sample for training
            X = X[:,0:n]
            d = d[0:n]
             # starting vector of weights
            w = [1, 1, 1]
             # sweep through the data N_EPOCH times
            for _ in range(N_EPOCH):
                 w = w + dwp(w,d,X)
            return w
        def performance(w,X,d):
            X = X[:,75:]
            d = d[75:]
            C = d*np.einsum('i,ij->j', w, X) # shape is (100,)
            XT = np.einsum('ij->ji', X)
             # d.x.t for misclassified
            xd = [ d[i]*XT[i] for i,c in enumerate(C) if c < 0]</pre>
            return len(xd)/25.0
        # invoke main
        def main():
             # data
            filedir = os.path.dirname(os.path.realpath('__file__'))
            datapath = os.path.join(filedir, 'assign1_data.mat')
            data = scipy.io.loadmat(datapath)
            x1 = data['x'][:,0]
            x2 = data['x'][:,1]
            # inputs - with bias input tied high, shape is (3,100)
            X = np.array([np.full(len(x1),1),x1,x2])
             # output
            d = np.array(data['z']).flatten()
             # use domain -1,1 for teachers so we can use x.w.d<0 for the classification test
            d = [z \text{ if } z == 1 \text{ else } -1 \text{ for } z \text{ in } d]
            samples = [5, 10, 25, 50, 75]
            perf = []
            for n in samples:
                w = train(n, X, d)
                 p = performance(w, X, d)
                 perf.append(1-p)
            return samples, perf
        samples, perf = main()
In [9]: plt.ylabel('% Correct')
```

```
plt.xlabel(r'$N_{training}$')
plt.title(r'Percent Correctly Classified vs Number Training Samples')
plt.plot(samples, perf)
```

Out[9]: [<matplotlib.lines.Line2D at 0x116b02c50>]

