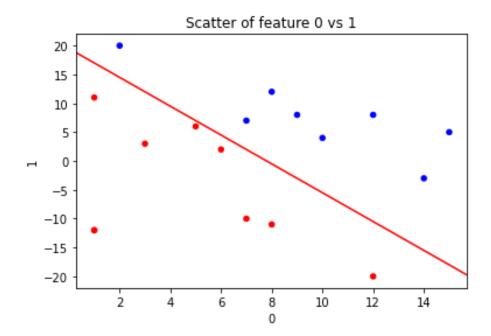
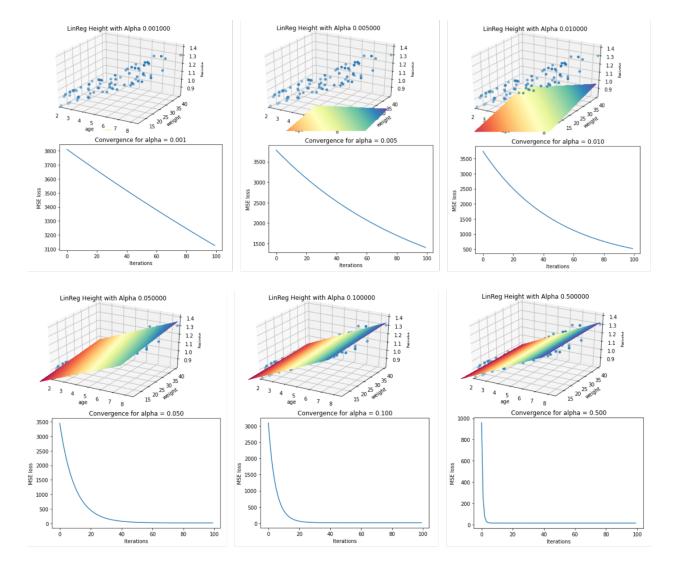
Due: 11:59 pm (New York, EDT); Tuesday, 29 November 2022

1 **Perceptron** decision boundary:



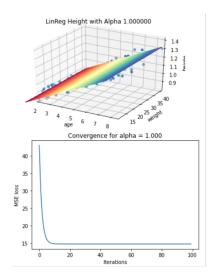
2 **Linear Regression** results (custom choice: alpha = 0.6 for 30 iterations)

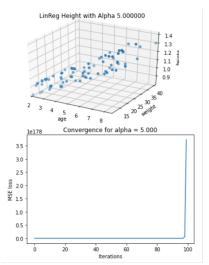
The objective in choosing alpha, the learning rate for the Linear Regression (LR) algorithm, is to select a value that converges on a loss minimum in a reasonable time. If alpha is too high, the algorithm may skip the minima and fail to converge, while if alpha is too low, the algorithm will converge too slowly. The plots below show the Mean Squared Error (MSE) loss against the number of iterations, using the nine provided alpha values (0.001, 0.005, 0.01, 0.05, 0.1, 0.5, 1, 5, and 10) as well as the custom value that was selected. As instructed, for all of the provided alphas, the algorithm is set to run for 100 iterations. Based on the illustrations, it can be seen that the MSE converges faster as alpha increases, with the fastest convergence occurring for an alpha of 0.5. Although the model still converges at the higher alpha value of 1 (albeit slightly more slowly than for 0.5), the model does not converge at all when alpha becomes too large (5 or 10). When alpha is 0.5 or 1, it can be observed that the model converges in fewer than 30 iterations. The choice for a tenth rate and number of iterations is alpha 0.6 with 30 iterations. From the last set of plots below, it can be seen that this choice converges quickly to a minimum MSE loss in a relatively small number of iterations.

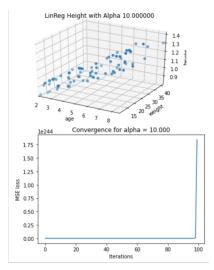


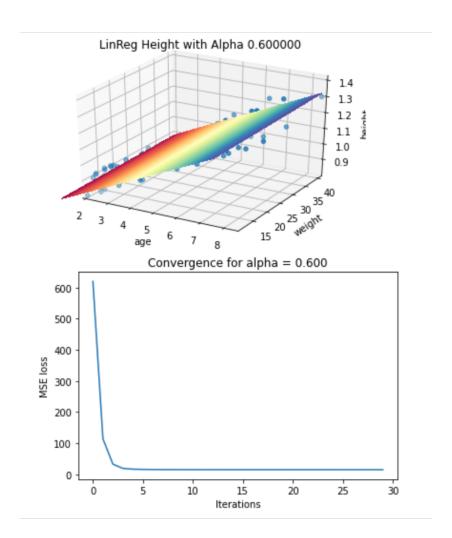
COMS 4701 Artificial Intelligence: Homework 04 Programming – SVW2112

Due: 11:59 pm (New York, EDT); Tuesday, 29 November 2022



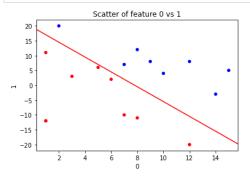






```
In [ ]: # plot db.py
        import matplotlib.pyplot as plt
        import numpy as np
        import pandas as pd
         import sys
         from matplotlib import cm
         import matplotlib.lines as mlines
        from mpl_toolkits.mplot3d import Axes3D
        Author: Kelsey D'Souza
         This file contains two functions for visualizing 2-feature labeled datasets.
         Its purpose is to give you ideas on how to vizualize your data and use pandas
         and matplotlib, feel free to snippets of any code in here or import the file
         into your programs.
         This file *does not* need to be included in your submission unless imported.
         visualize scatter
                Assumes binary labels and creates a line between the data using the given
                 feature and bias weights.
                Note: weights should be input as [w1, w2, bias]
         visualize 3d
                Plots data points in 3D space using feat1 x feat2 on the x-y base, and
                label as the data point height along the z-axis.
                It then creates a 3D surface plot of the continuous label model using
                the given linear regressor weights.
        def visualize_scatter(df, feat1=0, feat2=1, labels=2, weights=[-1, -1, 1],
                               title=''):
                Scatter plot feat1 vs feat2.
                Assumes +/- binary labels.
                Plots first and second columns by default.
                Args:
                  - df: dataframe with feat1, feat2, and labels
                   - featl: column name of first feature
                   - feat2: column name of second feature
                   - labels: column name of labels
                  - weights: [w1, w2, b]
            # Draw color-coded scatter plot
            colors = pd.Series(['r' if label > 0 else 'b' for label in df[labels]])
            ax = df.plot(x=feat1, y=feat2, kind='scatter', c=colors)
            # Get scatter plot boundaries to define line boundaries
            xmin, xmax = ax.get_xlim()
            # Compute and draw line. ax + by + c = 0 \Rightarrow y = -a/b*x - c/b
            a = weights[0]
            b = weights[1]
            c = weights[2]
            def y(x):
                return (-a/b)*x - c/b
            line_start = (xmin, xmax)
            line_end = (y(xmin), y(xmax))
            line = mlines.Line2D(line_start, line_end, color='red')
            ax.add_line(line)
            if title == '':
                title = 'Scatter of feature %s vs %s' %(str(feat1), str(feat2))
            ax.set_title(title)
            plt.show()
        def visualize_3d(df, lin_reg_weights=[1,1,1], feat1=0, feat2=1, labels=2,
                         xlim=(-1, 1), ylim=(-1, 1), zlim=(0, 3),
alpha=0., xlabel='age', ylabel='weight', zlabel='height',
title=''):
            3D surface plot.
            Main args:
              - df: dataframe with feat1, feat2, and labels
              - featl: int/string column name of first feature
              - feat2: int/string column name of second feature
              - labels: int/string column name of labels
              - lin_reg_weights: [b_0, b_1 , b_2] list of float weights in order
            Optional args:
              - x,y,zlim: axes boundaries. Default to -1 to 1 normalized feature values.
               - alpha: step size of this model, for title only
              - x,y,z labels: for display only
              - title: title of plot
```

```
In [ ]: # Perceptron Learning Algorithm (pla.py)
         import pandas as pd
         import numpy as np
         import sys
         def perceptron(X, y, output_file):
             num_samples = X.shape[0]
num_features = X.shape[1]
             output = []
             # Initialize weights and bias to zero
             w = np.zeros((num_features, 1))
             # Iterate until the model converges
             model_not_converged = True
             while model_not_converged:
                 # Assume convergence to start
model_not_converged = False
                  # Loop through each sample
                  for i in range(num_samples):
                      # Get the prediction
                      y_pred = b + np.dot(X[[i], :], w)
                      y_pred = 1 if y_pred > 0 else -1
                      # Check if the true and predicted labels have different signs
                      if y[i] * y_pred <= 0:
    # Yes, update weights and bias, and flag model as not converged</pre>
                          w = w + y[i] * np.transpose(X[[i], :])
b = b + y[i]
                          model_not_converged = True
                  # Record weights
                  out_line = []
                  for w_value in w:
                      out_line.append(str(int(w_value[0])))
                  out_line.append(str(int(b)))
                  output.append(out_line)
              # Write output CSV file
             pd.DataFrame(output).to_csv(output_file, header=False, index=False)
             # Return combined weights vector
             weights = np.zeros((num_features + 1, 1))
             weights[:-1] = w
weights[-1] = b
             return weights
         # Read data
         df = pd.read_csv('data1.csv', header=None)
         M = df.to_numpy()
         X = M[:, [0, 1]]
         y = M[:, 2]
         # Output file
output_file = str('output1.csv')
         weights = perceptron(X, y, output_file)
         visualize_scatter(df, weights = weights)
```



```
In [ ]: # Linear Regression Model (lr.py)
         import pandas as pd
         import numpy as np
         import sys
         def GradientDescent(alpha, X, y, means, stds, df, output, iterations = 100):
             n = len(X)
             beta = np.zeros(3)
             loss_list = np.zeros(iterations)
             # Iterate: update beta values and compute loss
             for i in range(iterations):
                 beta = beta - alpha*(1.0/n)*np.transpose(X).dot(X.dot(beta)-y)
                 loss_list[i] = (1.0/2*n) * np.sum(np.square(X.dot(beta)-y))
             # De-normalize weights
             weights = np.zeros(3)
             weights[1] = beta[1] / stds[0]
weights[2] = beta[2] / stds[1]
             weights[0] = beta[0] - (weights[1] * means[0]) - (weights[2] * means[1])
             # Plot model
             xlim = (min(df[0]), max(df[0]))
             ylim = (min(df[1]), max(df[1]))
             zlim = (min(df[2]), max(df[2]))
             visualize_3d(df, lin_reg_weights = weights, alpha = alpha,
                          xlim = xlim, ylim = ylim, zlim = zlim)
             # Plot loss versus iterations
             plt.plot(loss list)
             plt.title(f'Convergence for alpha = {alpha:.3f}')
            plt.xlabel('Iterations')
            plt.ylabel('MSE loss')
            plt.show()
             # File content
            output.append([alpha, iterations, f"{weights[0]:0.8f}", f"{weights[1]:0.8f}", f"{weights[2]:0.8f}"])\\
             # Done
            return weights
         # Alpha values
         alphas = [0.001, 0.005, 0.01, 0.05, 0.1, 0.5, 1, 5, 10]
         Custom alpha and iterations
         When alpha is too high it can skip the minima for convergence, but if alpha is too low then it will converge too slowly. We want to find a balance of the alpha and the number of iterations in order to
         converge on the minima with the lowest error rates.
         my_alpha = 0.6
         my_iterations = 30
         # Import data and get output file
         df = pd.read_csv('data2.csv', header=None)
        output_file = str('output2.csv')
         # Convert to numpy
        data = df.to_numpy()
         X = data[:, [0, 1]]
        y = data[:, 2]
         # Normalize features
         # Track the means and std devs to de-normalize later when plotting
         means = X.mean(axis=0)
         stds = X.std(axis=0)
         for i in range(2):
          X[:, i] = (X[:, i] - means[i]) / stds[i]
         # Add a column of ones at the start
         X = np.hstack([np.ones((X.shape[0], 1)), X])
         # Process the pre-set alpha values
         output = []
         for alpha in alphas:
             GradientDescent(alpha, X, y, means, stds, df, output)
         # Process the custom alpha value and iterations count
        GradientDescent(my_alpha, X, y, means, stds, df, output, my_iterations)
         # Write output CSV file
        pd.DataFrame(output).to_csv(output_file, header=False, index=False)
```

