CS 178: Machine Learning & Data Mining

Homework 2: Due Friday, 2 February 2024 (11:59 PM)

Instructions

This homework (and many subsequent ones) will involve data analysis and reporting on methods and results using Python code. You will submit a **single PDF file** that contains everything to Gradescope. This includes any text you wish to include to describe your results, the complete code snippets of how you attempted each problem, any figures that were generated, and scans of any work on paper that you wish to include. It is important that you include enough detail that we know how you solved the problem, since otherwise we will be unable to grade it.

Your homeworks will be given to you as Jupyter notebooks containing the problem descriptions and some template code that will help you get started. You are encouraged to modify these starter Jupyter notebooks to complete your assignment and to write your report. You may add additional cells (containing either code or text) as needed. This will help you not only ensure that all of the code for the solutions is included, but also will provide an easy way to export your results to a PDF file (for example, doing *print preview* and *printing to pdf*). Before submitting, ensure that your ubmission is complete, all text and code is legible (i.e. not cut off), and the pdf includes page breaks (i.e. not one very long page). We recommend liberal use of Markdown cells to create headers for each problem and subproblem, explaining your implementation/answers, and including any mathematical equations. For parts of the homework you do on paper, scan it in such that it is legible (there are a number of free Android/iOS scanning apps, if you do not have access to a scanner), and include it as an image in the Jupyter notebook.

If you have any questions/concerns about using Jupyter notebooks, ask us on Piazza. There you can also find additional instructions on how to convert to a .pdf.

Summary of Assignment: 100 total points

- Problem 1: Nearest Centroids on MNIST Dataset (25 points)
 - Problem 1.1: Visualizing MNIST (5 points)
 - Problem 1.2: Implementing Nearest Centroids (10 points)
 - Problem 1.3: Evaluating Nearest Centroids (10 points)
- Problem 2: kNN on Penguins Dataset (20 points)
 - Problem 2.1: Plot decision boundary for various k (10 points)
 - Problem 2.2: Plot tr/te error for various k (10 points)
- Problem 3: Logistic Regression (50 points)
 - Problem 3.1: Implement Forward Pass of Logistic Regression and fit to Penguins Dataset (10 points)
 - Problem 3.2: Explore Yelp dataset (10 points)
 - Problem 3.3: Fit a logistic regression classifier on Yelp dataset. (10 points)

- Problem 3.4: Find the top k most negative/most positive features and corresponding coefficients. (10 points)
- Problem 3.5: Tune reguarlization parameter on Yelp dataset (10 points)
- Statement of Collaboration (5 points)

Before we get started, let's import some libraries that you will make use of in this assignment. Make sure that you run the code cell below in order to import these libraries.

Important: In the code block below, we set seed=1234. This is to ensure your code has reproducible results and is important for grading. Do not change this. If you are not using the provided Jupyter notebook, make sure to also set the random seed as below.

```
In [44]:
         # JUST RUN - DO NOT EDIT THIS CODE BLOCK
         import numpy as np
         import pandas as pd
         import math
          import matplotlib.pyplot as plt
         from sklearn.datasets import fetch_openml
         from sklearn.neighbors import KNeighborsClassifier, NearestCentroid
         from sklearn.metrics import accuracy_score, confusion_matrix, ConfusionMatrixDisplay
         from sklearn.model_selection import train_test_split
         from sklearn.inspection import DecisionBoundaryDisplay
         from sklearn.preprocessing import LabelEncoder
         from sklearn.linear_model import LogisticRegression
         from sklearn.feature_extraction.text import CountVectorizer
         # Fix the random seed for reproducibility
         # !! Important !! : do not change this
         seed = 1234
         np.random.seed(seed)
```

Problem 1: Nearest Centroid on MNIST Dataset

In this problem, you will implement a nearest centroid classifier and train it on the MNIST dataset. The MNIST dataset is an image dataset consisting of 70,000 hand-written digits (from 0 to 9), each of which is a 28x28 grayscale image. For each image, we also have a label, corresponding to which digit is written. Run the following block of code to load the MNIST dataset:

In the following block, we split the MNIST dataset into training and testing sets -- 75% of the data is used for training, and 25% is used for testing. The function train_test_split is provided by scikit-learn, and will automatically shuffle our data for us if we use the flag shuffle=True.

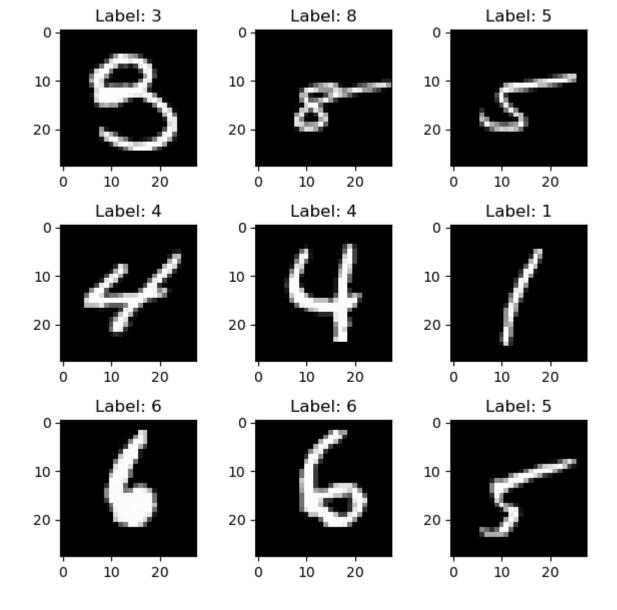
NOTE: For this homework, do not alter the flag random_state=seed, as this is necessary for obtaining reproducible results.

Problem 1.1 (5 points): MNIST Visualization

Let's begin by visualizing a few of the images in the MNIST dataset.

- Plot the first 9 images in mnist_X_tr in a 3x3 grid.
- Include a title for each subplot indicating the label of the image.

```
# Some default settings for our plots
In [48]:
         plt.rcParams['image.interpolation'] = 'nearest'
         plt.rcParams['image.cmap'] = 'gray'
         # Create a figure with 3 rows and 3 columns
         figure, axes = plt.subplots(3, 3, figsize=(6, 6))
         ### YOUR CODE STARTS HERE ###
         # Plot the first 9 images in our dataset.
         # Include a title on each subplot to indicate the corresponding label.
         k=0
         for i in range(0,3):
             for j in range(0,3):
                 image = mnist_X_tr[k,:]
                  imgreshape = image.reshape((28,28))
                 axes[i][j].imshow(imgreshape)
                 axes[i][j].set_title(f"Label: {mnist_y_tr[k]}")
                 k+=1
         ### YOUR CODE ENDS HERE ###
         plt.tight_layout()
```



Problem 1.2 (20 points): Implementing a Nearest Centroid Classifier

In the code given below, we define the class NearestCentroidClassifier which has an unfinished implementation of a nearest centroid classifier. For this problem, you will complete this implementation.

- Implement the method fit, which takes in an array of features X and an array of labels y and trains our classifier.
- Test your implementation of fit by training a NearestCentroidClassifier on the MNIST training set, and using the provided method plot_centroids to visualize the centroids. If your implementation is correct, the centroids should resemble the corresponding class label in the plot.
- Implement the method predict, which takes in an (array of) feature vectors X and predicts their class labels.
- Print the predicted labels (using your predict function) and the true labels for the first ten images in the MNIST testing set.

```
In [49]: class NearestCentroidClassifier:
    def __init__(self):
        # A list containing the centroids; to be filled in with the fit method.
        self.centroids = []

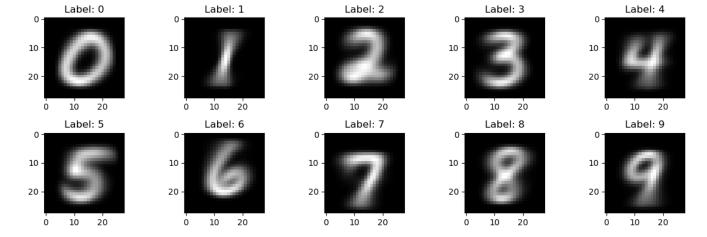
def plot_centroids(self):
    # Some default settings for our plots
```

```
plt.rcParams['image.interpolation'] = 'nearest'
        plt.rcParams['image.cmap'] = 'gray'
        # Create a figure with 2 rows and 5 columns
        figure, axes = plt.subplots(2, 5, figsize=(12, 4))
        # Plot the centroids
        for i in range(10):
            axes[i//5, i%5].imshow(self.centroids[i].reshape(28, 28))
            axes[i//5, i%5].set_title(f'Label: {i}')
        plt.tight_layout()
        plt.show()
    def fit(self, X, y):
        """ Fits the nearest centroid classifier with training features X and training labels
        X: array of training features; shape (n, d), where n is the number of datapoints,
        and d is the number of features.
        y: array training labels; shape (n, ), where n is the number of datapoints.
        ### YOUR CODE STARTS HERE ###
        # Hint: you should append to self.centroids with the corresponding centroids.
        \#np.where(my array == 3)[0]
        # Calculate the column means for the selected rows
        #column_means = np.mean(data[selected_rows], axis=0)
        for i in range(0,10):
            index = np.where(y == i)[0] #find Label i
            mean = np.mean(X[index], axis=0) # Calculate the column means for the selected re
            self.centroids.append(mean)
        ### YOUR CODE ENDS HERE
    def predict(self, X):
        """ Makes predictions with the nearest centroid classifier on the features in X.
        X: array of features; shape (n, d), where n is the number of datapoints,
        and d is the number of features.
        Returns:
        y_pred: a numpy array of predicted labels; shape (n, ), where n is the number of data
        ### YOUR CODE STARTS HERE ###
        centroids = np.array(self.centroids)
        y pred = []
        for i in range(X.shape[0]):
            #find each mean of the feature compare the centroid of 0-9
            preds = np.mean((centroids-X[i])**2, axis=1).flatten()
            #find the index of smallest mean(which is also label)
            pred = np.argmin(preds)
            y_pred.append(pred)
        ### YOUR CODE ENDS HERE ###
        return y_pred
# JUST RUN - DO NOT EDIT THIS CODE BLOCK
nc classifier = NearestCentroidClassifier()
```

```
In [51]: # JUST RUN - DO NOT EDIT THIS CODE BLOCK
    nc_classifier.plot_centroids()
```

nc_classifier.fit(mnist_X_tr, mnist_y_tr)

In [50]:



```
In [52]: # Print the predicted and true labels for the first ten images in the MNIST testing set
    ### YOUR CODE STARTS HERE ###
    y_pred = nc_classifier.predict(mnist_X_te)
    print(f"predicted label: {y_pred[:10]}")
    print(f"true label: {list(mnist_y_te[:10])}")
    ### YOUR CODE ENDS HERE ###

predicted label: [6, 6, 7, 7, 4, 7, 3, 4, 3, 5]
```

Problem 1.3 (10 points): Evaluating the Nearest Centroids Classifier

true label: [6, 6, 7, 7, 4, 7, 3, 4, 3, 5]

Now that you've implemented the nearest centroid classifier, it is time to evaluate its performance.

- Write a function compute_accuracy that computes the accuracy of a model's predictions. That is, your function should take in an array of true labels y and an array of predicted labels y_pred, and return the accuracy of the predictions. You may use numpy to do this, but do not use sklearn or any other machine learning libraries.
- Write a function that computes the confusion matrix of a model's predictions. That is, your function should take in an array of true labels y and an array of predicted labels y_pred, and return the confusion matrix as a numpy array. You may use numpy to do this, but do not use sklearn or any other machine learning libraries.
- Verify that your implementations of NearestCentroidClassifier, compute_accuracy, and compute_confusion_matrix are correct. To help you do this, you are given the functions eval_sklearn_implementation and eval_my_implementation. The function eval_sklearn_implementation will use the relevant sklearn implementations to compute the accuracy and confusion matrix of a nearest centroid classifier. The function eval_my_implementation will do the same, but for your implementations. If your code is correct, the outputs of the two functions should be the same.

```
In [53]: def compute_accuracy(y, y_pred):
    """ Computes the accuracy of an array of predictions.

y: true labels; shape (n, ), where n is the number of datapoints.
y_pred: predicted labels; shape (n, ), where n is the number of datapoints.

Returns:
    accuracy: the accuracy of y_pred compared to y; scalar expressed as a decimal (e.g. 0.5)
    """

### YOUR CODE STARTS HERE ###
# Create a new array without the specified value
#new_array = my_array[my_array != value_to_remove]
```

```
y_difference = y-y_pred
error = y_difference[y_difference != 0] #remove 0 from the array=>error
accuracy = 1.0-(np.size(error)/np.size(y)) #1-error%
### YOUR CODE ENDS HERE ###
return accuracy
```

```
In [54]:
         def compute_confusion_matrix(y, y_pred):
             """ Computes the confusion matrix of an array of predictions.
             y: true labels; shape (n, ), where n is the number of datapoints.
             y_pred: predicted labels; shape (n, ), where n is the number of datapoints.
             confusion_matrix: a numpy array corresponding to the confusion matrix from y and y_pred;
             where C is the number of unique classes.
             ### YOUR CODE STARTS HERE ###
             #Given a list of pairs (true_class = y, predicted_class = y_pred)
             #Initialize the confusion matrix entries to be all zeros
             conf_matrix = np.zeros((10, 10), dtype=int)
             yny_pred = np.column_stack((y,y_pred))
             #Go through the list, and for each pair add 1 to the corresponding row/column cell in the
             #I used np.unique to count each pairs=>faster than using loop
             unique_values, counts = np.unique(yny_pred, axis=0, return_counts=True)
             for value, count in zip(unique_values, counts):
                 conf_matrix[value[0]][value[1]] = count
             ### YOUR CODE ENDS HERE ###
             return conf_matrix
```

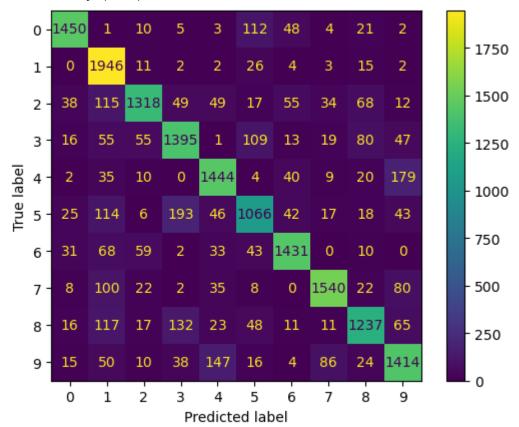
Run the two code blocks below to compare your implementation to the implementation in sklearn. Make sure you read and understand this code.

```
In [55]: # JUST RUN - DO NOT EDIT THIS CODE BLOCK
        ### Results with the sklearn implementation ###
         def eval_sklearn_implementation(X_tr, y_tr, X_te, y_te):
            # Nearest centroid classifier implemented in sklearn
            sklearn_nearest_centroid = NearestCentroid()
            # Fit on training dataset
            sklearn_nearest_centroid.fit(X_tr, y_tr)
            # Make predictions on training and testing data
            sklearn_y_pred_tr = sklearn_nearest_centroid.predict(X_tr)
            sklearn_y_pred_te = sklearn_nearest_centroid.predict(X_te)
            # Evaluate accuracies using the sklearn function accuracy score
            sklearn_acc_tr = accuracy_score(y_tr, sklearn_y_pred_tr)
            sklearn_acc_te = accuracy_score(y_te, sklearn_y_pred_te)
            print(f'Sklearn Results:')
            print(f'--- Accuracy (train): {sklearn_acc_tr}')
            print(f'--- Accuracy (test): {sklearn_acc_te}')
            # Evaluate confusion matrix using the sklearn function confusion matrix
            sklearn_cm = confusion_matrix(y_te, sklearn_y_pred_te)
            sklearn_disp = ConfusionMatrixDisplay(confusion_matrix = sklearn_cm)
            sklearn_disp.plot();
```

```
# Call the function
eval_sklearn_implementation(mnist_X_tr, mnist_y_tr, mnist_X_te, mnist_y_te)
```

Sklearn Results:

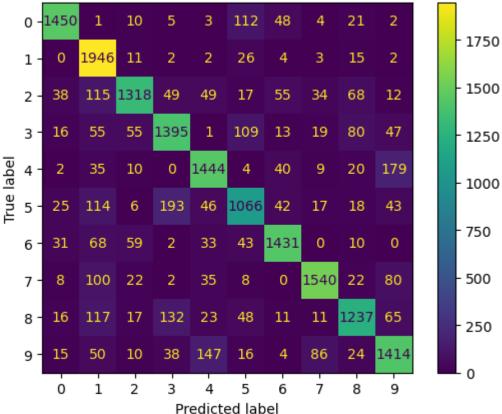
```
--- Accuracy (train): 0.8092571428571429
--- Accuracy (test): 0.8137714285714286
```



```
In [56]:
        # JUST RUN - DO NOT EDIT THIS CODE BLOCK
         ### Results with your implementation ###
         def eval_my_implementation(X_tr, y_tr, X_te, y_te):
            # Now test your implementation of NearestCentroidClassifier
            nearest_centroid = NearestCentroidClassifier()
            # Fit on training dataset
            nearest_centroid.fit(X_tr, y_tr)
            # Make predictions on training and testing data
            y_pred_tr = nearest_centroid.predict(X_tr)
            y_pred_te = nearest_centroid.predict(X_te)
            # Evaluate accuracies using your function compute_accuracy
            acc_tr = compute_accuracy(y_tr, y_pred_tr)
            acc_te = compute_accuracy(y_te, y_pred_te)
            print(f'Your Results:')
            print(f'--- Accuracy (train): {acc_tr}')
            print(f'--- Accuracy (test): {acc_te}')
            # Evaluate confusion matrix using your function compute_confusion_matrix
            cm = compute_confusion_matrix(y_te, y_pred_te)
            disp = ConfusionMatrixDisplay(confusion_matrix = cm)
            disp.plot();
```

```
# Call the function
eval_my_implementation(mnist_X_tr, mnist_y_tr, mnist_X_te, mnist_y_te)
```

```
Your Results:
--- Accuracy (train): 0.8092571428571429
--- Accuracy (test): 0.8137714285714286
```



Problem 2: kNN on Penguins Dataset

For the final problem of this homework, you will explore the k-nearest-neighbors algorithm using the Penguins dataset.

Important: Be sure to download the `penguins.csv` file from Canvas and place it in the same directory of this notebook.

```
In [57]: # JUST RUN - DO NOT EDIT THIS CODE BLOCK
    # Load the Penguins dataset
    peng_data = pd.read_csv("./penguins.csv") # Note that this is in the form of a Pandas datafr
    peng_data.head()
    #print(np.array(peng_data['bill_length_mm']))
    #np.unique(peng_data['species'])=>'Adelie', 'Chinstrap', 'Gentoo'
```

Out[57]:		species	island	bill_length_mm	bill_depth_mm	flipper_length_mm	body_mass_g	sex
	0	Adelie	Torgersen	39.1	18.7	181	3750	MALE
	1	Adelie	Torgersen	39.5	17.4	186	3800	FEMALE
	2	Adelie	Torgersen	40.3	18.0	195	3250	FEMALE
	3	Adelie	Torgersen	36.7	19.3	193	3450	FEMALE
	4	Adelie	Torgersen	39.3	20.6	190	3650	MALE

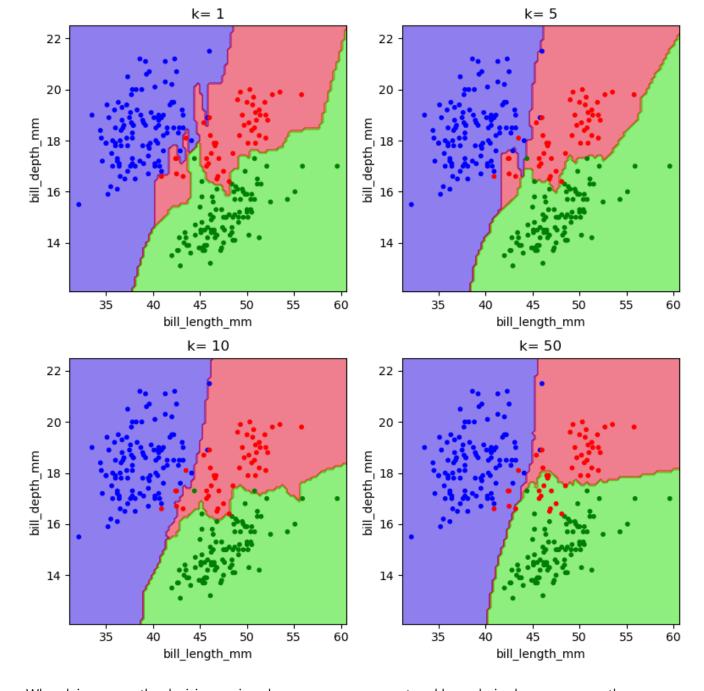
Problem 2.1: Decision Boundaries (15 points)

- Using the code given in Problem 1 and Problem 2, to create a train/test split of the Penguins dataset only containing the features "bill_length_mm" and "bill_depth_mm", as well as the class label "species". Use 75% of the data for training, and 25% of the data for testing. Set shuffle=True and be sure to use random_state=seed. Save the train/test split of the input features as peng_X_tr and peng_X_te respectively. For the train/test split of the class labels, store these under variable names peng_y_tr and peng_y_te respectively.
- For the values of k=[1, 5, 10, 50], fit a kNN classifier on this new training set with each value for k and plot the resulting decision boundary. Your plot should have 4 subplots total, clearly labeled and arranged in 2 rows and 2 columns. Each plot should showcase the decision boundary as well as display the individual training points with the colors of the point corresponding to the class.
- Write a short description of what you see happen as you increase the value of k.

Note that you are intended to use the sklearn implementation of kNN for this problem and 2.2.

```
# Create a 75%/25% train/test split using only the "bill_length_mm" and "bill_depth_mm" featu
In [58]:
         # along with the class labels "species".
         ### YOUR CODE STARTS HERE ###
         blm = np.array(peng_data['bill_length_mm']).reshape(-1, 1)
         bdm = np.array(peng_data['bill_depth_mm']).reshape(-1, 1)
         peng_X= np.hstack((blm, bdm))
         peng_y = np.array(peng_data['species'])
         peng_X_tr, peng_X_te, peng_y_tr, peng_y_te = train_test_split(
             peng_X, peng_y, test_size=0.25, random_state=seed, shuffle=True
         ### YOUR CODE ENDS HERE ###
         print(peng_X_tr.shape, peng_X_te.shape, peng_y_tr.shape, peng_y_te.shape)
         (256, 2) (86, 2) (256,) (86,)
In [59]: # Some keyword arguments for making nice looking plots.
         # Feel free to change grid_resolution to a higher number -- this results in better looking pl
         # but may result in your code running more slowly.
         plot_kwargs = {'cmap': 'viridis',
                         'response_method': 'predict',
                         'plot_method': 'pcolormesh',
                         'shading': 'auto',
                         'alpha': 0.5,
                         'grid_resolution': 100}
         # Create a figure with 2 rows and 2 columns
         figure, axes = plt.subplots(2, 2, figsize=(8, 8))
         ### YOUR CODE STARTS HERE ###
```

```
# Feel free to use the LabelEncoder to convert the class label strings into integer values fo
label_encoder = LabelEncoder()
peng_y_tr_encoded = label_encoder.fit_transform(peng_y_tr)
#grid_resolution
h = 0.2
ks=[1, 5, 10, 50]
k=0
colors = []
for c in range(peng_y_tr.shape[0]):
   if peng_y_tr[c] =='Adelie':
       colors.append("blue")
   elif peng_y_tr[c] == 'Chinstrap':
        colors.append("red")
   else:
        colors.append("green")
for i in range(peng_X_tr.shape[1]):
   for j in range(peng_X_tr.shape[1]):
        neigh = KNeighborsClassifier(n_neighbors=ks[k])
        classifier = neigh.fit(peng_X_tr, peng_y_tr_encoded)
        disp = DecisionBoundaryDisplay.from_estimator(classifier, peng_X_tr, ax = axes[i,j],
        axes[i,j].scatter(peng_X_tr[:,0], peng_X_tr[:,1], c=colors, s=10)
        axes[i,j].set_xlabel('bill_length_mm')
        axes[i,j].set_ylabel('bill_depth_mm')
        axes[i,j].set_title(f"k= {ks[k]}")
       k+=1
plt.tight_layout()
### YOUR CODE ENDS HERE ###
```



When k increases, the decision regions become more compact and boundaries becomes smoother.

Problem 2.2: Error Rates vs k (15 points)

Now, we will vary the value of k and see what effect this has on our predictions.

- Again using only the "bill_length_mm" and "bill_depth_mm" features of the Penguins dataset, compute the error rate on both the training and testing data as a function of k. Do this for all values of k = [1, 2, 5, 10, 50, 100, 110]. You may use your own implementation of the accuracy, or the scikit-learn function sklearn.metrics.accuracy_score.
- Plot the resulting error rate functions using a semi-log plot (i.e. the x-axis is on a logarithmic scale), with the training error in red and the validation error in green.
- What value of k would you recommend, and why?

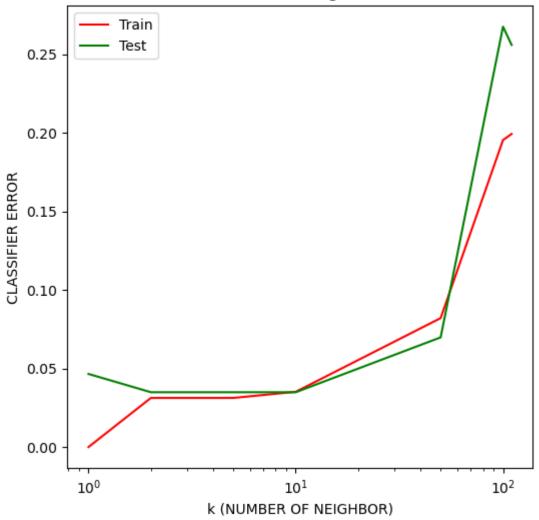
```
In [60]: # Create a figure with only one subplot
figure, axes = plt.subplots(1, figsize=(6, 6))
### YOUR CODE STARTS HERE ###
```

```
k = [1, 2, 5, 10, 50, 100, 110]
test_error=[]
train_error=[]
for i in k:
   neigh = KNeighborsClassifier(n_neighbors=i)
   neigh.fit(peng_X_tr, peng_y_tr)
   peng_y_tepred = neigh.predict(peng_X_te)
   teaccuracy = accuracy_score(peng_y_te, peng_y_tepred)
   test_error.append(1-teaccuracy)
   peng_y_trpred = neigh.predict(peng_X_tr)
   traccuracy = accuracy_score(peng_y_tr, peng_y_trpred)
   train_error.append(1-traccuracy)
print(f'test error: {test error}')
print(f'train error: {train_error}')
axes.semilogx(k,train_error,label='Train', color="red")
axes.semilogx(k,test_error,label='Test', color="green")
axes.set_xlabel('k (NUMBER OF NEIGHBOR)')
axes.set_ylabel('CLASSIFIER ERROR')
axes.set_title('Error Rate on Penguins Data')
axes.legend()
### YOUR CODE ENDS HERE ###
```

test error: [0.046511627906976716, 0.03488372093023251, 0.03488372093023251, 0.03488372093023251, 0.06976744186046513, 0.2674418604651163, 0.2558139534883721] train error: [0.0, 0.03125, 0.03125, 0.03515625, 0.08203125, 0.1953125, 0.19921875] <matplotlib.legend.Legend at 0x1d211217710>

Out[60]:

Error Rate on Penguins Data



k=2 or 5, since both train(0.03488372093023251) and test errors(0.03125) are relatively low (They have the same value).

Problem 3: Logistic Regression

Important: We will need to install three more libraries for this assignment, do not skip this. Also, be sure to download the 'yelp_reviews.json' file from Canvas and place it in the same directory of this notebook

We need to run conda install -c conda-forge scipy simplejson nltk in order to install scipy, nltk, and simplejson. This can also be done directly inside this notebook by running !conda install ... inside a code cell.

We need scipy to work with certain mathematical optimization techniques, nltk to work with natural language data, and simplejson to open json files.

In this problem, you will first implement a basic version of logistic regression on the Penguins dataset, and then you will work with the sklearn implementation of logistic regression on a more realistic dataset.

```
# JUST RUN - DO NOT EDIT THIS CODE BLOCK
In [61]:
         from scipy.optimize import minimize
         import simplejson as json
         import nltk
         nltk.download('punkt')
         nltk.download('averaged_perceptron_tagger')
         nltk.download('universal_tagset')
         from nltk import word_tokenize
         [nltk_data] Downloading package punkt to
                         C:\Users\allye\AppData\Roaming\nltk_data...
         [nltk_data]
         [nltk_data]
                       Package punkt is already up-to-date!
          [nltk_data] Downloading package averaged_perceptron_tagger to
         [nltk_data]
                         C:\Users\allye\AppData\Roaming\nltk_data...
         [nltk_data]
                       Package averaged_perceptron_tagger is already up-to-
         [nltk_data]
         [nltk_data] Downloading package universal_tagset to
                         C:\Users\allye\AppData\Roaming\nltk_data...
         [nltk_data]
```

Problem 3.1 (10 points): Implement Forward Pass of Logistic Regression and fit to Penguins Dataset

Package universal_tagset is already up-to-date!

[nltk_data]

Recall that logistic regression models the probability of a datapoint with features x_1, x_2 having class y=1 via:

$$p(y=1|x_1,x_2)=\sigma(heta_0+ heta_1x_1+ heta_2x_2)$$

where $\sigma(z)=\frac{1}{1+e^{-z}}$. Likewise, it models the probability of class y=0 via $p(y=0|x_1,x_2)=1-\sigma(\theta_0+\theta_1x_1+\theta_2x_2)$. We typically learn parameter values $\theta_0,\theta_1,\theta_2$ by minimizing the cross-entropy of a given dataset:

$$\mathcal{L}(heta_0, heta_1, heta_2; ec{y}, ec{x}_1, ec{x}_2) = rac{1}{n} \sum_{i=1}^n -y_i \log p(y=1|x_{i1}, x_{i2}) - (1-y_i) \log p(y=0|x_{i1}, x_{i2})$$

- Implement the logistic_prob function that takes in parameter values theta_0, theta_1, and theta_2, as well as two vectors of input data x_1 and x_2 of length n, and produces the n corresponding probability values of $y=1|x_1,x_2$ as an <code>np.array</code>.
- Implement the logistic_loss function that takes in parameter values theta_0, theta_1, and theta_2, as well as three data vectors of input features x_1 and x_2 and the class values y of length n, and produces the cross-entropy value as defined above. Note this should call the logistic_prob function.
- Implement the plot_boundary_with_intercept function that takes in parameter values theta_0, theta_1, and theta_2, as well as three data vectors of input features x_1 and x_2 and the class values y of length n, and plots all datapoints x_1 on the x-axis, x_2 on the y-axis, with colors determined by class values y, and the decision boundary determined via parameter values.
- From the Penguins dataset, let x_1 be the "bill_length_mm" feature, let x_2 be the "bill_depth_mm" feature, and let y be a 1 when a penguin is of the "Adelie" species and a 0 when not. Use scipy 's minimize function to find the optimal parameter values $\theta_0, \theta_1, \theta_2$ that minimize the negative cross-entropy over this dataset. Plot the resulting decision boundary using plot_boundary_with_intercept (note this can be done exactly since the model is simple).

```
def sigmoid(z):
In [62]:
             if isinstance(z, np.ndarray):
                 return 1 / (1 + np.exp(-z))
             else:
                 return 1 / (1 + math.exp(-z))
         def logistic_prob(theta_0, theta_1, theta_2, x_1, x_2):
             '''Return probability of y=1 given x_1 and x_2 for specific parameter values theta_0,
             theta_1, and theta_2.'''
             ### YOUR CODE STARTS HERE ###
             z = theta_0 + theta_1*x_1 + theta_2 * x_2
             prob_y1 = sigmoid(z)
             ### YOUR CODE ENDS HERE ###
             return prob_y1
         def logistic_loss(theta_0, theta_1, theta_2, x_1, x_2, y):
In [63]:
             '''Compute the cross-entropy for logistic regression model parameterized by
             theta_0, theta_1, and theta_2 over dataset y \mid x_1, x_2.''
             ### YOUR CODE STARTS HERE ###
             prob_y1 = logistic_prob(theta_0, theta_1, theta_2, x_1, x_2)
             cross_entropy = np.mean(-y*np.log(prob_y1)-(1-y)*np.log(1-prob_y1))
             ### YOUR CODE ENDS HERE ###
             return cross_entropy
In [64]:
         def plot_boundary_w_intercept(theta_0, theta_1, theta_2, x_1, x_2, y):
             theta_0: the bias of linear model
             theta_1: the first parameter of linear model
             theta_2: the second parameter of linear model
                      [# feature vectors], feature vectors
             x_1:
             x_2:
                        [# feature vectors], feature vectors
                      [# feature vectors], labels
             y:
```

Make a figure with 1 subplot
fig, axes = plt.subplots()

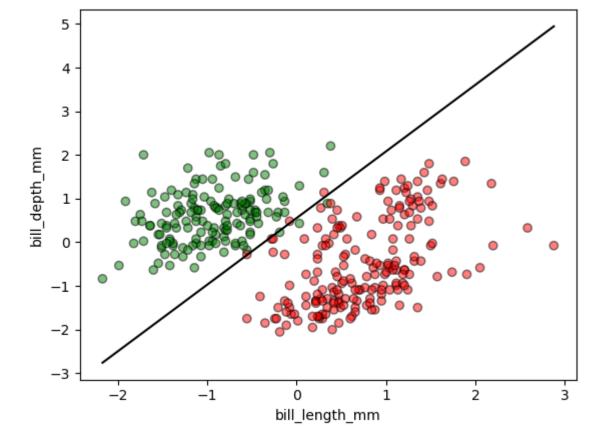
```
if y[c] ==0:
                                               colors.append("red")
                                               colors.append("green")
                              axes.scatter(x_1, x_2, alpha=0.5,c=colors, edgecolors="k")
                              axes.set xlabel('bill length mm')
                              axes.set ylabel('bill depth mm')
                              x_{boundary} = np.linspace(min(x_1), max(x_1), 100)
                             y_boundary = (-theta_0 - theta_1 * x_boundary) / theta_2
                              axes.plot(x_boundary, y_boundary, label='Decision Boundary', color='black')
                              ### YOUR CODE ENDS HERE ###
                     \# Set `x_1` and `x_2` to be the training values for the "bill_length_mm" and "bill_depth_mm"
In [65]:
                     # Set `y` to be an array with values of 1 if the training observation is of species "Adelie"
                     # \dot{y} should be of same length as \dot{x}_1 and \dot{x}_2.
                     ### YOUR CODE STARTS HERE ###
                     x_1 = np.array(peng_data['bill_length_mm'])
                     x_2 = np.array(peng_data['bill_depth_mm'])
                     species = np.array(peng_data['species'])
                     y=np.zeros(len(x_1))
                     for i in range(len(x 1)):
                             if(species[i]=="Adelie"):
                                      y[i]=1
                     ### YOUR CODE ENDS HERE ###
                     print(x_1.shape, x_2.shape, y.shape)
                     (342,) (342,) (342,)
In [66]: # JUST RUN - DO NOT EDIT THIS CODE BLOCK
                     # Standardize inputs
                     std_x_1 = (x_1-x_1.mean()) / x_1.std()
                     std_x_2 = (x_2-x_2.mean()) / x_2.std()
                     loss_function = lambda thetas: logistic_loss(thetas[0], thetas[1], thetas[2], x_1=std_x_1, x
                     bound = 20 # Limit search of theta values to have this magnitude or less for stability reaso
                     initial_parameter_value = np.array([0.0, 0.0, 0.0])
                     r = minimize(loss_function, initial_parameter_value, bounds=[(-bound, bound) for _ in range(3
                     print("theta_0={}, theta_1={}, theta_2={}".format(*r.x))
                     plot_boundary_w_intercept(theta_0=r.x[0], theta_1=r.x[1], theta_2=r.x[2], x_1=std_x_1, x_2=std_x_1, x_2=std_x_1, x_2=std_x_1, x_2=std_x_1, x_3=std_x_1, x_4=std_x_1, x_4=std_x
                    theta_0=-4.361620222112985, theta_1=-12.051533943600239, theta_2=7.886921802207681
```

YOUR CODE STARTS HERE

for c in range(len(y)):

colors = []

prob_y1 = logistic_prob(theta_0, theta_1, theta_2, x_1, x_2)



Problem 3.2 (10 points): Explore Yelp dataset

The Yelp dataset can be used in a sentiment analysis task, where we are given a review and we predict whether it's positive or negative. We will pre-process the dataset for you with bag-of-words approach. You will create a bag of words (BOW) representation from text documents, using the Vectorizer function in scikit-learn.

A bag of words representation of text is a way to turn text into a fixed size vectorized output, this is done by deciding on a fixed-size vocabulary of words that are often used and then assigning each word a number, then, every sentence in a dataset is turned into a vector \mathbf{x} , where \mathbf{x}_i denotes whether the ith word in our list is used in the sentence. For example, if our vocabulary is [he, she, his, her, loves, likes, dogs, cats, cute, nice], the sentence "he loves cat and she loves dog" can be vectorized as [1, 1, 0, 0, 2, 0, 1, 1, 0, 0]. This representation allows us to compare sentences in the same way we have been doing to tabular data and images.

The inputs are

- a filename (you will use "yelp_reviews.json") containing the reviews in JSON format
- the min_pos and max_neg parameters (use the default values.)

The outputs are

- X: Feature Matrix in compressed sparse row format.
 - **X** is in shape of [# reviews, # terms in vocabulary]
 - **\mathbf{X}_{ij}** indicates the number of the jth term in the vocabulary existing in the ith review.
- y: Review label vector
- **text**: Raw reviews
- vectorizerBOW.vocabulary: A mapping of terms to feature indices, somes terms are excluded

• please read the scikit-learn tutorial on text feature extraction before you start this problem: https://scikit-learn.org/stable/modules/feature extraction.html#text-feature-extraction

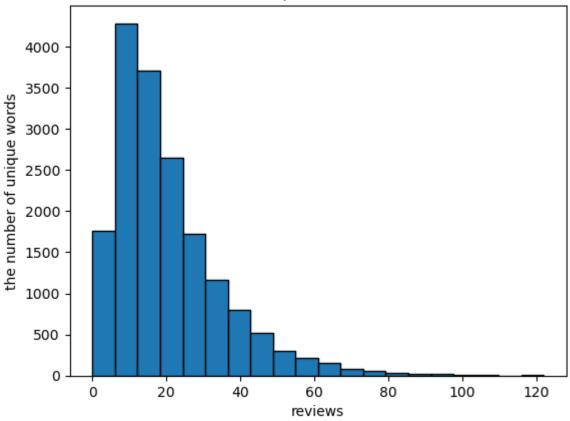
```
# JUST RUN - DO NOT EDIT THIS CODE BLOCK
In [67]:
         def create_bow_from_reviews(filename, min_pos=4, max_neg=2):
             print('Loading the file:', filename)
             with open(filename, 'r') as jfile:
                 data = json.load(jfile)
             print('Total number of reviews extracted =', len(data))
             text = []
             y = []
             print('Extracting tokens from each review.....(can be slow for a large number of reviews)
                                # can substitute data[0:9] here if you want to test this function on
             for d in data:
                 review = d['text'] # keep only the text and label
                 stars = int(d['stars'])
                 if stars >= min_pos: # represent scores > min_pos as "1"
                     score = 1
                 elif stars <= max_neg: # represent scores < max_neg as "0"</pre>
                 else: # do not consider reviews with scores above max_neg and below min_pos (these re
                     continue
                 text.append(review)
                 y.append(score)
             # create an instance of a CountVectorizer, using
             # (1) the standard 'english' stopword set
             # (2) only keeping terms in the vocabulary that occur in at least 1% of documents
             # (3) allowing only unigrams in the vocabulary (use "ngram_range=(1, 1)" to do this)
             vectorizer = CountVectorizer(stop_words='english',min_df=0.02, ngram_range=(1, 1))
             # create a sparse BOW array from 'text' using vectorizer
             X = vectorizer.fit_transform(text)
             print('Data shape: ', X.shape)
             # you can uncomment this next line if you want to see the full list of tokens in the voca
             #print('Vocabulary: ', vectorizer.get feature names())
             return X, y, vectorizer, text
In [68]: # JUST RUN - DO NOT EDIT THIS CODE BLOCK
         X, y, vectorizer_BOW, text = create_bow_from_reviews('yelp_reviews.json')
         Loading the file: yelp_reviews.json
         Total number of reviews extracted = 20000
         Extracting tokens from each review.....(can be slow for a large number of reviews).....
         Data shape: (17501, 404)
         Implement the function review_word_count_histogram , which generates a histogram to show how
         many unique words (from the vocabulary) are in each review. That is, for each review, count the number
```

```
In [69]: def review_word_count_histgram(X):
    ### YOUR CODE STARTS HERE ###
    unique_words = X.toarray()
    num_unique_words = [row[row != 0].tolist() for row in unique_words]
```

of unique words in the review, and plot a histogram illustrating these counts.

```
num_unique_words = [len(row) for row in num_unique_words]
#print(num_unique_words)
fig, axes = plt.subplots()
axes.hist(num_unique_words, bins=20 ,edgecolor='black')
axes.set_title('number of unique words in the review')
axes.set_xlabel("reviews")
axes.set_ylabel("the number of unique words")
### YOUR CODE ENDS HERE ###
review_word_count_histgram(X)
```

number of unique words in the review



Problem 3.3 (10 points): Fit a logistic regression classifier on Yelp dataset.

Separate an X,Y dataset (X=features, Y=labels) into training and test subsets

- Build a logistic classifier (from sklearn, not our previous implementation from problem 3.1) on the training subset. Please use 'I1' penality type, 'liblinear' solver and enable fit_intercept
- Evaluate performance on the test subset

NOTE: before starting this problem please read the scikit-learn documentation on logistic classifiers: https://scikit-learn.org/stable/modules/linear_model.html and logistic regression https://scikit-learn.org/stable/modules/generated/sklearn.linear_model.LogisticRegression.html#sklearn.linear_model.Log

```
In [70]: def logistic_classification(X, y):
    X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.3, random_state=see
# set the state of the random number generator so that we get the same results across rule.
```

```
print('Number of training examples: ', X_train.shape[0])
print('Number of testing examples: ', X_test.shape[0])
print('Vocabulary size: ', X_train.shape[1])

### YOUR CODE STARTS HERE ###
# Specify the Logistic classifier model
# Please use 'l1' penality type, 'liblinear' solver and enable fit_intercept
classifier = LogisticRegression(penalty='l1', solver='liblinear', fit_intercept=True)
classifier.fit(X_train, y_train)

# Compute and print accuracy on the test data
test_predictions = classifier.predict(X_test)
test_accuracy = accuracy_score(y_test, test_predictions)

### YOUR CODE ENDS HERE ###

print('\nTesting accuracy:', format( 100*test_accuracy , '.2f') )
return classifier
```

```
In [71]: # JUST RUN - DO NOT EDIT THIS CODE BLOCK
    logistic_classifier = logistic_classification(X, y)

Number of training examples: 12250
    Number of testing examples: 5251
    Vocabulary size: 404
```

Problem 3.4 (10 points): Find the top k most negative/most positive features

In this problem, you will complete the function <code>most_significant_terms</code> to

- print out and return the most significant positive and negative weights
- print our the associated terms

Testing accuracy: 89.51

most_significant_terms takes as input

- a scikit-learn trained logistic regression classifier (e.g., trained in Problem 2.3)
- a scikit-learn vectorizer object that produced the BOW features for the classifier

and prints out

- the terms in the vocabulary tokens with the **top10** largest positive weights
- the terms in the vocabulary with the top10 smallest (i.e., largest in terms of absolute value) negative weights

Hint: Check the attributes section of the vectorizer documentation (https://scikit-learn.org/stable/modules/generated/sklearn.feature_extraction.text.CountVectorizer.html) to find where the mapping between words and indices are located.

```
In [72]: def most_significant_terms(classifier, vectorizer, K):
    # cycle through the positive weights, in the order of largest weight first and print out
    # K lines where each line contains
    # (a) the term corresponding to the weight (a string)
    # (b) the weight value itself (a scalar printed to 3 decimal places)

### YOUR CODE STARTS HERE ###
```

```
# extract coefficients from trained model and sort based on coefficient
coefficients = classifier.coef [0]
coefficients = coefficients[1:] #reomve theta 0
features = np.arange(1, len(coefficients)+1)
coefficients_dict = dict(zip(features, coefficients))
# sort from large to small
sort_coef = list(sorted(coefficients_dict.items(), key=lambda x: x[1], reverse=True))
names = vectorizer.get_feature_names_out()
# extract topK positive terms
topK_pos_index = [item[0] for item in sort_coef[:K]][::-1]
topK_pos_terms= [names[i] for i in topK_pos_index]
# extract corresponding weights
topK_pos_weights = [item[1] for item in sort_coef[:K]][::-1]
# extract topK negative terms #[::-1] reverse
topK_neg_index = [item[0] for item in sort_coef[-K:]][::-1]
topK_neg_terms = [names[i] for i in topK_neg_index]
# extract corresponding weights
topK_neg_weights = [item[1] for item in sort_coef[-K:]][::-1]
### YOUR CODE ENDS HERE ###
print('topK_pos_weights', topK_pos_weights)
print('topK_pos_terms', topK_pos_terms)
print('topK_neg_weights', topK_neg_weights)
print('topK_neg_terms',topK_neg_terms)
return (topK_pos_weights, topK_neg_weights, topK_pos_terms, topK_neg_terms)
```

```
In [73]: # JUST RUN - DO NOT EDIT THIS CODE BLOCK
  output = most_significant_terms(logistic_classifier, vectorizer_BOW, K=10)
```

```
topK_pos_weights [1.5574633076300772, 1.6342102917019183, 1.6730570344539843, 1.726028425698 2228, 1.7268897127051854, 1.760880422467794, 1.986403486830437, 2.032124775696237, 2.3056578 018656473, 2.3116872481142483] topK_pos_terms ['easy', 'enjoyed', 'perfect', 'thank', 'wonderful', 'awesome', 'amazing', 'excellent', 'delicious', 'unique'] topK_neg_weights [-3.3223280059894273, -1.5438213919708186, -1.2990807123655226, -1.26948635 27040765, -1.1738281149135814, -1.109455321411859, -1.0753783837776383, -0.902560056495337, -0.899239179494491, -0.8571106959195339] topK_neg_terms ['worst', 'okay', 'disappointed', 'tasted', 'ok', 'guess', 'won', 'maybe', 'slow', 'money']
```

Problem 3.5 (10 points): Tune reguarlization parameter on Yelp dataset

In this problem, you will complete the function accuracy_vs_reg

- plot curve of train/test accuracy vs regularization strength.
- choose a final value for the regularization parameter and explain why.

```
In [74]: def accuracy_vs_reg(X, y):

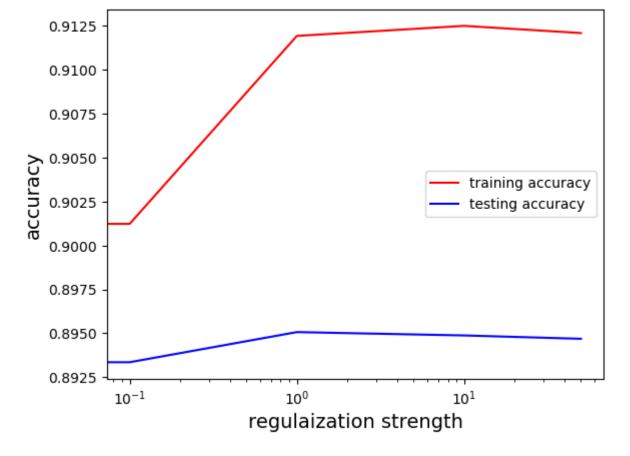
    X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.3, random_state=see
    # set the state of the random number generator so that we get the same results across ru

    coefs_ = []
    train_accuracies = []
    test_accuracies = []
    cs = [0, 0.1, 1, 10, 50]

### YOUR CODE STARTS HERE ###
```

```
for c in cs:
    # Specify the logistic classifier modle
    if c == 0:
        classifier = LogisticRegression(penalty=None, fit_intercept=True)
   else:
        # Refer https://scikit-learn.org/stable/modules/generated/sklearn.linear_model.Ld
        # to see which argument can adjust the regularization regularization,
        # Additionally, please use 'l1' penalty type, 'liblinear' solver and enable fit_i
        classifier = LogisticRegression(penalty='l1', solver='liblinear', fit_intercept=1
   classifier.fit(X_train, y_train)
   train_predictions = classifier.predict(X_train)
   train_accuracy = accuracy_score(y_train, train_predictions)
   test_predictions = classifier.predict(X_test)
   test_accuracy = accuracy_score(y_test, test_predictions)
    # log model coefficients
    coefs_.append(classifier.coef_[0])
   # log model training and test accuracies
   train_accuracies.append(train_accuracy)
   test_accuracies.append(test_accuracy)
### YOUR CODE ENDS HERE ###
fig, axes = plt.subplots()
axes.semilogx(cs, train_accuracies, color='red', label='training accuracy')
axes.semilogx(cs, test_accuracies, color='blue', label='testing accuracy')
axes.set_xlabel('regulaization strength', fontsize=14)
axes.set_ylabel('accuracy', fontsize=14)
axes.legend()
return train_accuracies, test_accuracies, coefs_
```

```
In [75]: # JUST RUN - DO NOT EDIT THIS CODE BLOCK
train_accuracies, test_accuracies, coefs = accuracy_vs_reg(X, y)
```



Choose a final value for the regularization parameter and justify your choice. You should choose a single value from the list cs given in the code above.

1 is the best value since both training and testing accuracy are relatively high.

Statement of Collaboration (5 points)

It is **mandatory** to include a Statement of Collaboration in each submission, with respect to the guidelines below. Include the names of everyone involved in the discussions (especially in-person ones), and what was discussed. If you did not collaborate you can simply write N/A.

All students are required to follow the academic honesty guidelines posted on the course website. For programming assignments, in particular, I encourage the students to organize (perhaps using Piazza) to discuss the task descriptions, requirements, bugs in my code, and the relevant technical content before they start working on it. However, you should not discuss the specific solutions, and, as a guiding principle, you are not allowed to take anything written or drawn away from these discussions (i.e. no photographs of the blackboard, written notes, referring to Piazza, etc.). Especially after you have started working on the assignment, try to restrict the discussion to Piazza as much as possible, so that there is no doubt as to the extent of your collaboration.

N/A