## CS441: Applied ML - HW 1

### Parts 1-2: MNIST

Include all the code for generating MNIST results below

```
In [48]: # initialization code
         import numpy as np
          from keras.datasets import mnist
         %matplotlib inline
          from matplotlib import pyplot as plt
          from scipy import stats
         def load mnist():
            Loads, reshapes, and normalizes the data
            (x train, y train), (x test, y test) = mnist.load data() # loads MNIST dat
            x train = x train[::-1]
            y train = y train[::-1]
           x train = np.reshape(x train, (len(x train), 28*28)) # reformat to 784-d
           x \text{ test} = \text{np.reshape}(x \text{ test}, (\text{len}(x \text{ test}), 28*28))
            maxval = x train.max()
           x train = x train/maxval # normalize values to range from 0 to 1
            x_{test} = x_{test/maxval}
            return (x_train, y_train), (x_test, y_test)
          def display mnist(x, subplot rows=1, subplot cols=1):
            Displays one or more examples in a row or a grid
            if subplot rows>1 or subplot cols>1:
              fig, ax = plt.subplots(subplot rows, subplot cols, figsize=(15,15))
              for i in np.arange(len(x)):
                ax[i].imshow(np.reshape(x[i], (28,28)), cmap='gray')
                ax[i].axis('off')
            else:
                plt.imshow(np.reshape(x, (28,28)), cmap='gray')
                plt.axis('off')
            plt.show()
```

In [49]: # example of using MNIST load and display functions
 (x\_train, y\_train), (x\_test, y\_test) = load\_mnist()
 display\_mnist(x\_train[:10],1,10)
 print('Total size: train={}, test ={}'.format(len(x\_train), len(x\_test)))



#### 1. Retrieval, Clustering, and NN Classification

**Retrieval**: Implement the function get\_nearest using Euclidean (L2) distance. Check that get\_nearest( $x_{test}[0]$ ,  $x_{train}$ ) returns i=6156. Report the index of the closest example in x train to x test[1].

```
In [50]: # Retrieval
         def get nearest(X query, X):
           minimum distance = np.Inf
           index = -1
           for i in range(len(X)):
             dist = np.linalg.norm(X[i] - X query)
             if dist < minimum distance:</pre>
               minimum distance = dist
               index = i
           return index
           ''' Return the index of the sample in X that is closest to X query accordi
               to L2 distance '''
           # TO DO
         j = get nearest(x test[0], x train)
         print(j)
         j = get nearest(x test[1], x train)
         print(j)
```

6156 31117

**K-means**: Using your get\_nearest function, write a function kmeans that iteratively assigns each data point to the nearest cluster center. Apply it to only the first 1000 examples,  $x_{\text{train}}[:1000]$ . Try this with K = 10 and K = 30, and display the cluster centers after each iteration. Include the displays from after the 1st and 10th iteration for K=30 in your report. See the note in the assignment if your cluster centers do not seem to be changing.

```
for j in range(len(X)): #
       # index = get nearest(X[j], centroids);
       # print(index)
       cluster index.append(get nearest(X[j], centroids))
     new centroid = []
     for j in range(K):
       cluster = []
       for l in range(len(X)):
        if j == cluster index[l] :
          cluster.append(X[l])
       new centroid.append(np.mean(cluster, axis=0))
     # are equal = sorted(list1) == sorted(list2)
     # if(np.array equal(np.array(centroids), np.array(new centroid))):
     # break
     centroids = new centroid
     # print(centroids)
     if (i == 0 \text{ or } i == 9):
       print("For K = " + str(K) + " after iteration " + str(i+1))
       display mnist(centroids[:K], 1, K)
       # plot clusters(X, centroids, cluster index, K, iteration=i+1)
   final cluster index = cluster index
   # return centroids, final cluster index
   Starting with the first K samples in X as cluster centers, iteratively ass
   point to the nearest cluster and compute the mean of each cluster.
   Input: X[i] is the ith sample, K is the number of clusters, niter is the r
   Output: K cluster centers
   # TO DO -- implement kmeans and add code to display cluster centers at eac
 K = 30
 print("for K = " + str(K))
 kmeans(x train[:1000], K)
 K = 10
 print("for K = " + str(K))
 kmeans(x train[:1000], K)
for K = 30
For K = 30 after iteration 1
865384592987062706631376429022
For K = 30 after iteration 10
86938 # 57 2 9 8 7 0 6 2 1 0 6 6 3 1 3 7 6 4 4 9 0 2 2
for K = 10
For K = 10 after iteration 1
8653815729
```



**1-NN**: Now, use your get\_nearest function to perform 1-nearest neighbor. For each test sample, find the index of the closest sample in the training data to predict its label. To check your method, calculate the error for the first 100 test samples using only the first 1,000 training samples; the error should be 19%. Report the percent error for the first 100 test samples using the first 10,000 training samples.

```
In [55]: # 1-NN
         # TO DO
         def nearest neaighbour classifier(x train, y train, x test):
          y test = []
           for i in np.arange(len(x test)):
             y test.append(y train[get nearest(x test[i], x train)])
           return np.array(y test)
         trained y test = nearest neaighbour classifier(x train[:1000], y train[:1000
         wrong values = 0
         for i in range(len(trained y test)):
           if(trained y test[i] != y test[i]):
             wrong values += 1
         sample = 100
         error = (wrong values/ sample) * 100
         print(error)
        19.0
```

```
In [56]: trained_y_test = nearest_neaighbour_classifier(x_train[:10000], y_train[:100]
    wrong_values = 0
    for i in range(len(trained_y_test)):
        if(trained_y_test[i] != y_test[i]):
            wrong_values += 1
    sample = 100
    error = (wrong_values/ sample) * 100
    print(error)
```

#### 7.0000000000000001

#### 2. Make it fast

```
In [57]: # install libraries you need for part 2
!apt install libomp-dev
!pip install faiss-cpu
import faiss
import time
```

```
Reading package lists... Done
Building dependency tree... Done
Reading state information... Done
libomp-dev is already the newest version (1:14.0-55~exp2).
0 upgraded, 0 newly installed, 0 to remove and 49 not upgraded.
Requirement already satisfied: faiss-cpu in /usr/local/lib/python3.10/dist-p ackages (1.8.0.post1)
Requirement already satisfied: numpy<2.0,>=1.0 in /usr/local/lib/python3.10/dist-packages (from faiss-cpu) (1.26.4)
Requirement already satisfied: packaging in /usr/local/lib/python3.10/dist-p ackages (from faiss-cpu) (24.1)
```

**Retrieval**: Exact search can be performed using the code below.

```
index = faiss.IndexFlatL2(X.shape[1]) # set for exact search
index.add(x_train) # add the data
dist, idx = index.search(x_test[:2],1) # returns index and sq
err for each sample
```

Check that idx matches your retrieved indices from Part 1.

```
In [58]: # retrieval
   index = faiss.IndexFlatL2(x_train.shape[1]) # set for exact search
   index.add(x_train) # add the data
   dist, idx = index.search(x_test[:2],1) # returns index and sq err for each s
   print(dist)
   print(idx)
# TO DO (check that you're using FAISS correctly)

[[ 7.0398464]
   [20.79831 ]]
   [[ 6156]
   [31117]]
```

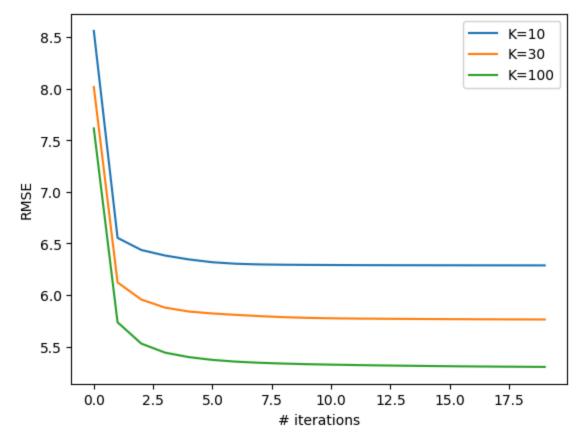
**K-means**: Complete fast\_kmeans using FAISS for the retrieval instead of your get\_nearest function. In each iteration, create a new index, add the cluster centers, and find the nearest center to all samples. Once it's working, disable any print or display functions inside the kmeans loop.

Record the root mean squared error (RMSE) at the start of each iteration, and plot the RMSE for each iteration for K=10, K=30, and K=100 when clustering the full training set with 20 iterations.

```
centroids = X[:K].copy()
  rmse list = []
 # for i in range(K):
    index = np.random.randint(len(X));
 # centroids.append(X[index])
 final cluster index = []
 for i in range(niter):
    index = faiss.IndexFlatL2(X.shape[1])
    index.add(np.array(centroids))
    temp dist, cluster index = index.search(X,1)
    # print(cluster index)
    # for j in range(len(X)):
    # cluster index.append(get nearest(X[j], centroids))
    # temp rmse k = []
    new centroid = []
    for j in range(K):
      cluster = []
     # cluster dist = []
     for l in range(len(X)):
        if j == cluster index[l] :
          cluster.append(X[l])
          # cluster dist.append(temp dist[l])
      new centroid.append(np.mean(cluster, axis=0))
     # temp rmse = 0
     # for dist in cluster dist:
     # temp rmse += dist*dist
     # temp rmse = np.sqrt(temp rmse/len(cluster dist))
     # temp rmse k.append(temp rmse)
    t = 0
    # for dist in temp dist:
   # t += dist*dist
    rmse list.append(np.sqrt(np.mean(temp dist)))
    # if(centroids == new centroid):
    # break
   centroids = new centroid
   # print(centroids)
   # if (i == 0 \text{ or } i == 9):
     \# print("For K = " + str(K) + " after iteration " + str(i+1))
      # display_mnist(centroids[:K], 1, K)
 final cluster index = cluster index
 return centroids, rmse list
 # TO DO (you can base this on part 1, but use FAISS for search)
 # if you include display code, you need to re-organize the plotting code b
K = 10
centers, rmse = kmeans fast(x train, K, niter=20)
plt.plot(np.arange(len(rmse)), rmse, label='K=10')
```

```
K=30
centers, rmse = kmeans_fast(x_train, K, niter=20)
plt.plot(np.arange(len(rmse)), rmse, label='K=30')

K=100
centers, rmse = kmeans_fast(x_train, K, niter=20)
plt.plot(np.arange(len(rmse)), rmse, label='K=100')
plt.legend(), plt.ylabel('RMSE'), plt.xlabel('# iterations')
plt.show()
```



**1-NN**: Use FAISS to evaluate 1-NN on the full training and test sets. Try this with both the exact search and LSH approximate search. The only difference is how you set up the index.

For LSH, use:

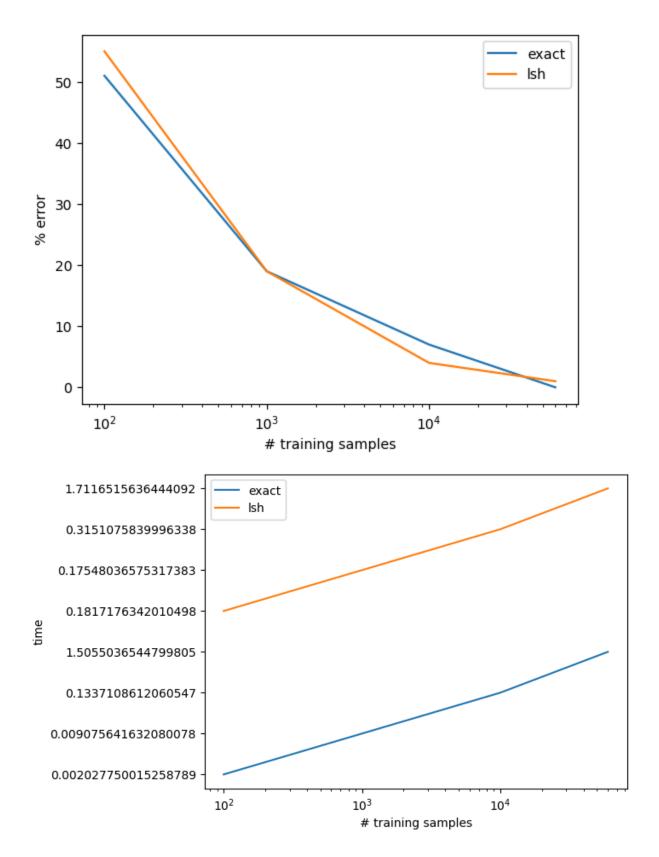
```
dim = X.shape[1]
index = faiss.IndexLSH(dim, dim)
```

Evaluate 1-NN using each search method, while varying the number of training samples: s in [100,1000, 10000, 60000]. In each case, use  $x_{train}$ [:s] as the train set. Plot number of samples vs. percent error on a semilogx plot for both exact and LSH (on the same plot). Also, record timings using time.time() and plot samples vs. time on a semilogx plot.

```
In [68]: # 1-NN
         def nearest neaighbour classifier Faiss(x train, y train, x test):
           index = faiss.IndexFlatL2(x train.shape[1]) # set for exact search
           index.add(x train)
           dist, idx = index.search(x test, 1)
           # print(idx)
           y test = []
           for i in np.arange(len(x test)):
             y test.append(y train[idx[i]])
           return np.array(y test)
         def nearest neaighbour classifier Faiss LSH(x train, y train, x test):
           dim = x train.shape[1]
           index = faiss.IndexLSH(dim, dim)
           index.add(x train)
           dist, idx = index.search(x test,1)
           # print(idx)
           y test = []
           for i in np.arange(len(x test)):
             y test.append(y train[idx[i]])
           return np.array(y test)
         nsample = [100, 1000, 10000, 60000]
         acc exact = []
         timing exact = []
         for s in nsample:
           start time = time.time()
           trained_y_test = nearest_neaighbour_classifier_Faiss(x_train[:s], y_train[
           # end time = time.time()
           # timing exact.append((str(end time - start time)))
           wrong values = 0
           for i in range(len(trained y test)):
             if(trained y test[i] != y test[i]):
               wrong values += 1
           sample = 100
           error = (wrong_values/ sample) * 100
           acc exact.append(error)
           end time = time.time()
           timing_exact.append((str(end_time - start_time)))
           print(str(error) + " error found when sample was " + str(s) + " and wrong
         acc lsh = []
         timing lsh = []
         for s in nsample:
           start time = time.time()
           trained y test = nearest neaighbour classifier Faiss LSH(x train[:s], y tr
           # end time = time.time()
           # timing lsh.append(str(end time - start time))
           wrong values = 0
           for i in range(len(trained y test)):
             if(trained_y_test[i] != y_test[i]):
               wrong values += 1
           sample = 100
```

```
error = (wrong values/ sample) * 100
 acc lsh.append(error)
 end time = time.time()
 timing lsh.append(str(end time - start time))
 print(str(error) + " error found when sample was " + str(s) + " and wrong
# TO DO
acc exact = np.array(acc exact)
acc lsh = np.array(acc lsh)
timing exact = np.array(timing exact)
timing lsh = np.array(timing lsh)
plt.semilogx(nsample, (acc exact), label='exact')
plt.semilogx(nsample, (acc lsh), label='lsh')
plt.legend(), plt.ylabel('% error'), plt.xlabel('# training samples')
plt.show()
plt.semilogx(nsample, timing exact, label='exact')
plt.semilogx(nsample, timing lsh, label='lsh')
plt.legend(), plt.ylabel('time'), plt.xlabel('# training samples')
plt.show()
```

- 51.0 error found when sample was 100 and wrong values were 51with time required for execution 0.002027750015258789
- 19.0 error found when sample was 1000 and wrong values were 19with time required for execution 0.009075641632080078
- 7.0000000000000 error found when sample was 10000 and wrong values were 7w ith time required for execution 0.1337108612060547
- 0.0 error found when sample was 60000 and wrong values were 0with time required for execution 1.5055036544799805
- 55.000000000000 error found when sample was 100 and wrong values were 55 w ith time required for execution 0.1817176342010498
- 19.0 error found when sample was 1000 and wrong values were 19 with time required for execution 0.17548036575317383
- 4.0 error found when sample was 10000 and wrong values were 4 with time required for execution 0.3151075839996338
- $1.0~{
  m error}$  found when sample was  $60000~{
  m and}$  wrong values were  $1~{
  m with}$  time required for execution 1.7116515636444092



In your report, indicate which label is most often confused with '3' when using the full training set and exact search for 1-NN.

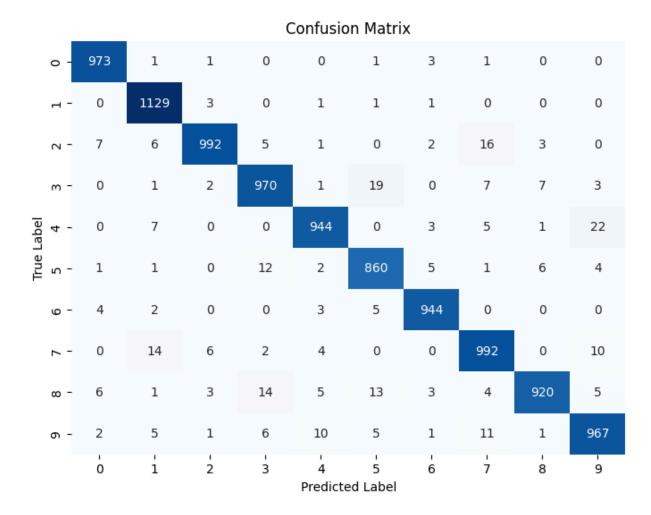
```
In [69]: # Confusion matrix
import sklearn as sns
from sklearn.metrics import confusion_matrix
```

```
In [70]: timing_exact = []
    start_time = time.time()
    size = len(x_test)
    trained_y_test = nearest_neaighbour_classifier_Faiss(x_train, y_train, x_test end_time = time.time()
    timing_exact.append((str(end_time - start_time)))
    wrong_values = 0
    for i in range(size):
        if(trained_y_test[i] != y_test[i]):
            wrong_values += 1
        error = (wrong_values/ size) * 100
# acc_exact.append(int(error))
    print(str(error) + " error found when sample was " + str(s) + " and wrong values/ size) * 100
```

 $3.09~{\rm error}$  found when sample was  $60000~{\rm and}$  wrong values were  $309~{\rm with}$  time r equired for execution 33.41194415092468

```
import seaborn as sns
cm = confusion_matrix(y_test, trained_y_test)

# Step 2: Visualize the confusion matrix
plt.figure(figsize=(8, 6))
sns.heatmap(cm, annot=True, fmt='d', cmap='Blues', cbar=False)
plt.title('Confusion Matrix')
plt.xlabel('Predicted Label')
plt.ylabel('True Label')
plt.show()
```



8 is mostly confused as 3

# Part 3: Temperature Regression

Include all your code used for part 3 in this section.

```
In [72]: from google.colab import files
    uploaded = files.upload()
```

Choose Files No file chosen

Upload widget is only available when the cell

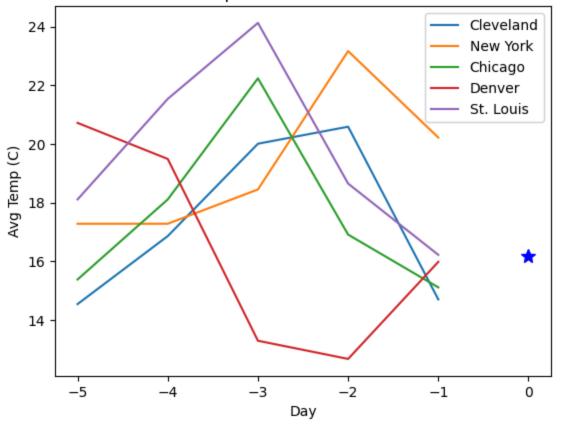
has been executed in the current browser session. Please rerun this cell to enable. Saving temperature\_data.npz to temperature\_data.npz

```
import numpy as np
from google.colab import drive
%matplotlib inline
from matplotlib import pyplot as plt
from sklearn.linear_model import Ridge
from sklearn.linear_model import Lasso

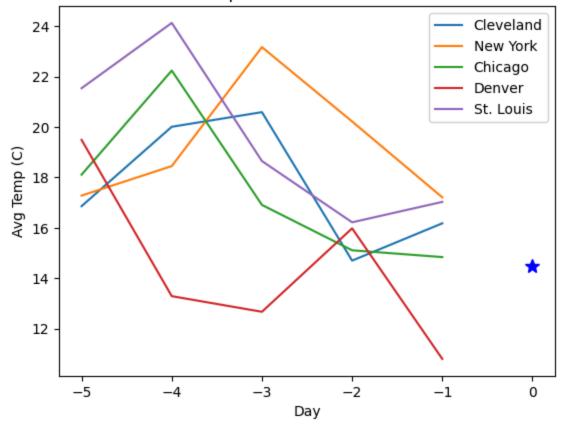
# load data (modify to match your data directory or comment)
def load_temp_data():
    # drive.mount('/content/drive')
```

```
# datadir = "/content/drive/My Drive/CS441/24FA/hw1/"
           T = np.load('temperature data.npz')
           xt train, yt train, xt val, yt val, xt test, yt test, dates train, dates v
           T['x train'], T['y train'], T['x val'], T['y val'], T['x test'], T['y test
           return (xt_train, yt_train, xt_val, yt_val, xt_test, yt_test, dates_train,
         # plot one data point for listed cities and target date
         def plot temps(x, y, cities, feature_to_city, feature_to_day, target_date):
           nc = len(cities)
           ndays = 5
           xplot = np.array([-5, -4, -3, -2, -1])
           yplot = np.zeros((nc,ndays))
           for f in np.arange(len(x)):
             for c in np.arange(nc):
               if cities[c] == feature to city[f]:
                 yplot[feature to day[f]+ndays,c] = x[f]
           plt.plot(xplot,yplot)
           plt.legend(cities)
           plt.plot(0, y, 'b*', markersize=10)
           plt.title('Predict Temp for Cleveland on ' + target date)
           plt.xlabel('Day')
           plt.ylabel('Avg Temp (C)')
           plt.show()
         # load temp data()
         # plot temps()
In [74]: # load data (use xt and yt so that we aren't replacing the MNIST variables)
         (xt train, yt train, xt val, yt val, xt test, yt test, dates train, dates va
         ''' Data format:
               x train, y train: features and target value for each training sample (
               x val, y val: features and target value for each validation sample (us
               x test, y test: features and target value for each test sample (used t
               dates xxx: date of the target value for the corresponding sample
               feature to city: maps from a feature number to the city
               feature to day: maps from a feature number to a day relative to the ta
               Note: 361 is the temperature of Cleveland on the previous day
         f = 361
         print('Feature {}: city = {}, day= {}'.format(f,feature to city[f], feature
         baseline rmse = np.sqrt(np.mean((yt val[1:]-yt val[:-1])**2)) # root mean sq
         print('Baseline - prediction using previous day: RMSE={}'.format(baseline rm
         # plot first two x/y for val
         plot temps(xt val[0], yt val[0], ['Cleveland', 'New York', 'Chicago', 'Denvϵ
         plot temps(xt val[1], yt val[1], ['Cleveland', 'New York', 'Chicago', 'Denve
        Feature 361: city = Cleveland, day= -1
        Baseline - prediction using previous day: RMSE=3.460601246750482
```

## Predict Temp for Cleveland on 2018-09-27



# Predict Temp for Cleveland on 2018-09-28



**KNN Regression**: Perform 5-NN regression, reporting RMSE for two variants:

- 1. Original features
- Normalize the features by subtracting the previous day's Cleveland temperature. I.e., if previous day's Cleveland temperature is c, features are X, and value to predict is y, then predict y\_query-c = NN(X\_query, X-c, y-c)

For these experiments, train on  $(x_{train}, y_{train})$  and test on  $(x_{test}, y_{test})$ . To validate your method, if you set K=3, you should get an RMSE of 3.314 for the original features.

```
In [77]: # K-NN Regression
         def regress KNN(x train, y train, x test, K=5):
           Predict the target value for each data point in X tst using a
           K-nearest neighbor regressor based on (X trn, y trn), with L2 distance.
           Input: X trn[i] is the ith training data. y trn[i] is the ith training lat
           Output: return y pred, where y pred[i] is the predicted ith test value
           # TO DO
           index = faiss.IndexFlatL2(x train.shape[1]) # set for exact search
           index.add(x train)
           dist, idx = index.search(x test,K)
           res = np.zeros(x test.shape[0])
           for i in np.arange(len(x test)):
             ind = idx[i]
             values = []
             for j in range(K):
               values.append(y train[ind[j]])
             \# avg = np.mean(y train[idx[i]])
             avg = np.mean(values)
             res[i] = avg
           return res
         def normalize features(x, y, fnum):
           ''' Normalize the features in x and y.
               For each data sample i:
                 x2[i] = x[i]-x[i,fnum]
                 y2[i] = y[i]-x[i,fnum]
           # TO DO
           temp x = x.copy()
           temp y = y.copy()
           for i in np.arange(len(x)):
             for j in np.arange(len(x[i])):
               temp x[i,j] = x[i,j] - x[i,fnum]
             temp y[i] = y[i] - x[i,fnum]
           return temp x, temp y
```

```
# print(len(xt test))
# KNN with original features
yt pred = regress KNN(xt train, yt train, xt test,K)
rmse = np.sqrt(np.mean((yt test - yt pred) ** 2))
print("RMSE with original features: " + str(rmse))
# TO DO
# KNN with normalized features
fnum = 361 # previous day temp in Cleveland
norm xt train, norm yt train = normalize features(xt train, yt train, fnum)
norm xt test, norm yt test = normalize features(xt test, yt test, fnum)
norm yt pred = regress KNN(norm xt train, norm yt train, norm xt test,K)
norm rmse = np.sqrt(np.mean((norm yt test - norm yt pred) ** 2))
print("RMSE with normalized features: " + str(norm rmse))
# print(len(feature to city))
# print(feature to city[:30])
# print(feature to day[:100])
# xt train, yt train, xt val, yt val, xt test, yt test, dates train, dates v
```

RMSE with original features: 3.1846378844060936 RMSE with normalized features: 2.8999927784510935

### Part 5: Stretch Goals

Include all your code used for part 5 in this section. You can copy-paste code from parts 1-3 if it is re-usable.

Compare K-NN on the MNIST classification for N=1, 3, 5, 11, 25. For these tests, use  $x_{train}[:50000]$  as a training set and  $x_{train}[50000:]$  as a validation set. Report error on the validation set for all parameters. Then performance on the test set for the best parameter using the full training set.

When K is greater than 1, return the most common label of the nearest samples. If there is a tie, return the most common label with the closest sample.

```
In [97]: from collections import Counter
# Stretch: KNN classification (Select K)
def classifier_KNN(x_train, y_train, x_test, K=5):
    index = faiss.IndexFlatL2(x_train.shape[1]) # set for exact search
    index.add(x_train)
    dist, idx = index.search(x_test,K)
    res = np.zeros(x_test.shape[0])
    for i in np.arange(len(x_test)):
        ind = idx[i]
        values = []
```

```
for j in range(K):
               values.append(y train[ind[j]])
             # most common element = max(set(values), key=values.count)
             most common element = Counter(values).most common(1)[0][0]
             # max occurrences = values.count(most common element)
             res[i] = most common element
           return res
         N = [1,3,5,11,25]
         result = []
         for i in N:
          trained y test = classifier KNN(x train[:50000], y train[:50000], x train[
           test_y_train = y_train[50000:]
           wrong values = 0
           for j in range(len(trained y test)):
             if(trained y test[j] != test y train[j]):
               wrong values += 1
           sample = len(test y train)
           error = (wrong values/ sample) * 100
           print( "Error for K = " + str(i) + " is " + str(error) )
           result.append(error)
        Error for K = 1 is 3.04
        Error for K = 3 is 2.85
        Error for K = 5 is 3.02
        Error for K = 25 is 4.3
In [100... best result = min(result)
         index_best_result = result.index(best_result)
         best N = N[index best result]
         print("Best K is " + str(best_N))
         trained y test = classifier KNN(x train, y train, x test, best N)
         wrong values = 0
         for i in range(len(trained y test)):
           if(trained_y_test[i] != y_test[i]):
             wrong values += 1
         sample = len(trained y test)
         error = (wrong values/ sample) * 100
         print("Error for best K is " + str(error))
        Best K is 3
        Error for best K is 2.83
```

Compare K-NN on the temperature regression dataset for N=1, 3, 5, 11, 25 using both feature types. Report all results on the validation set, and then run the single best setting on the test set, using all training samples.

```
In [101... # Stretch: KNN regression (Select K)
# original features
N = [1,3,5,11,25]
result = []
for i in N:
    yt_pred = regress_KNN(xt_train, yt_train, xt_val, i)
    rmse = np.sqrt(np.mean((yt_val - yt_pred) ** 2))
```

```
print("RMSE with original features: " + str(rmse) + " for N=" + str(i))
           result.append(rmse)
        RMSE with original features: 4.334638061394246 for N=1
        RMSE with original features: 3.2477289008934584 for N=3
        RMSE with original features: 3.1233409141099897 for N=5
        RMSE with original features: 3.0008389705781706 for N=11
        RMSE with original features: 3.0281505759415834 for N=25
In [102... best result = min(result)
         index best result = result.index(best result)
         best N = N[index best result]
         print("Best K is " + str(best N))
         yt pred = regress KNN(xt_train, yt_train, xt_test, best_N)
         rmse = np.sqrt(np.mean((yt test - yt pred) ** 2))
         print("Best RMSE with original features " + str(rmse) + " for N=" + str(best
         result.append(rmse)
         # yt pred = regress KNN(xt_train, yt_train, xt_test, 25)
         # rmse = np.sqrt(np.mean((yt test - yt pred) ** 2))
         # print("Best RMSE with original features " + str(rmse) + " for N=" + str(25
         # result.append(rmse)
        Best K is 11
        Best RMSE with original features 3.051018719912078 for N=11 on test set
In [90]: # with Normalized features
         norm xt train, norm yt train = normalize features(xt train, yt train, fnum)
         norm xt test, norm yt test = normalize features(xt test, yt test, fnum)
         norm xt val, norm yt val = normalize features(xt val, yt val, fnum)
In [94]: N = [1,3,5,11,25]
         result = []
         for i in N:
           norm yt pred = regress KNN(norm xt train, norm yt train, norm xt val, i)
           rmse = np.sqrt(np.mean((norm_yt_val - norm_yt_pred) ** 2))
           print("RMSE with normalized features: " + str(rmse) + " for N=" + str(i))
           result.append(rmse)
        RMSE with normalized features: 3.9396003204553005 for N=1
        RMSE with normalized features: 3.2584032096117133 for N=3
        RMSE with normalized features: 3.080241273239787 for N=5
        RMSE with normalized features: 2.917200409530882 for N=11
        RMSE with normalized features: 2.9156171084845264 for N=25
In [95]: best result = min(result)
         index best result = result.index(best result)
         best N = N[index best result]
         print("Best K is " + str(best N))
         norm yt pred = regress KNN(norm xt train, norm yt train, norm xt test, best
         rmse = np.sqrt(np.mean((norm yt test - norm yt pred) ** 2))
         print("Best RMSE with normalized features " + str(rmse) + " for N=" + str(be
         result.append(rmse)
        Best K is 25
        Best RMSE with normalized features 2.771407614735461 for N=25on test set
```

For K-means (on MNIST), does running with multiple re-runs or running a single run longer tend to provide lower RMSE? For this, you can use the FAISS Kmeans function. Syntax is below.

```
kmeans = faiss.Kmeans(x_train.shape[1], 30, niter=10,
nredo=1, seed=int(t))
kmeans.train(x_train)
dist, idx = kmeans.index.search(x_train, 1)
rmse = np.sqrt(np.sum(dist) / x_train.shape[0])
```

Compare (niter=10, nredo=5) vs. (niter=50, nredo=1) for K=30. Repeat this test five times and report the mean and standard deviation of the RMSE. Compare (niter=4, nredo=5) vs. (niter=20, nredo=1) for K=30. Repeat this test five times and report the mean and standard deviation of the RMSE.

```
In [103... # Stretch: K-means (more iters vs redos)
         K=30
         test 1 = [[10,5],[50,1],[4,5],[20,1]]
         for i in range(4):
           rmse list = []
           for j in range(5):
             kmeans = faiss.Kmeans(x train.shape[1], 30, niter=test 1[i][0], nredo=te
             kmeans.train(x train)
             dist, idx = kmeans.index.search(x train, 1)
             rmse = np.sqrt(np.sum(dist) / x train.shape[0])
             rmse list.append(rmse)
           mean = np.mean(rmse list)
           std = np.std(rmse list)
           print("For niter = " + str(test_1[i][0]) + " and nredo = " + str(test_1[i]
        For niter = 10 and nredo = 5 mean = 5.784186645165795 standard deviation =
        0.008603519764584013
        For niter = 50 and nredo = 1 mean = 5.779994090042927 standard deviation =
        0.008189570735299564
        For niter = 4 and nredo = 5 \text{ mean} = 5.826142983081921 \text{ standard deviation} = 0.
        006489017796755849
        For niter = 20 and nredo = 1 mean = 5.786202065039114 standard deviation =
        0.01073526294213036
 In [ ]: # from https://gist.github.com/jonathanagustin/b67b97ef12c53a8dec27b343dca4a
         # For use in Colab. For local, just use jupyter nbconvert directly
         import os
         # @title Convert Notebook to PDF. Save Notebook to given directory
         NOTEBOOKS DIR = "/content/drive/My Drive/CS441/hw1" # @param {type:"string"}
         NOTEBOOK NAME = "CS441 HW1 Solution.ipynb" # @param {type:"string"}
         from google.colab import drive
         drive.mount("/content/drive/", force_remount=True)
```

```
NOTEBOOK_PATH = f"{NOTEBOOKS_DIR}/{NOTEBOOK_NAME}"

assert os.path.exists(NOTEBOOK_PATH), f"NOTEBOOK_NOT FOUND: {NOTEBOOK_PATH}"
!apt install -y texlive-xetex texlive-fonts-recommended texlive-plain-generi
!jupyter nbconvert "$NOTEBOOK_PATH" --to pdf > /dev/null 2>&1
NOTEBOOK_PDF = NOTEBOOK_PATH.rsplit('.', 1)[0] + '.pdf'
assert os.path.exists(NOTEBOOK_PDF), f"ERROR MAKING PDF: {NOTEBOOK_PDF}"
print(f"PDF_CREATED: {NOTEBOOK_PDF}")
```

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