knn nosol

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0.1 This is the k-nearest neighbors workbook for ECE C147/C247 Assignment #2

Please follow the notebook linearly to implement k-nearest neighbors.

Please print out the workbook entirely when completed.

The goal of this workbook is to give you experience with the data, training and evaluating a simple classifier, k-fold cross validation, and as a Python refresher.

0.2 Import the appropriate libraries

```
import numpy as np # for doing most of our calculations
import matplotlib.pyplot as plt# for plotting
from utils.data_utils import load_CIFAR10 # function to load the CIFAR-10_
dataset.

# Load matplotlib images inline
%matplotlib inline

# These are important for reloading any code you write in external .py files.
# see http://stackoverflow.com/questions/1907993/
dautoreload-of-modules-in-ipython
%load_ext autoreload
%autoreload 2
```

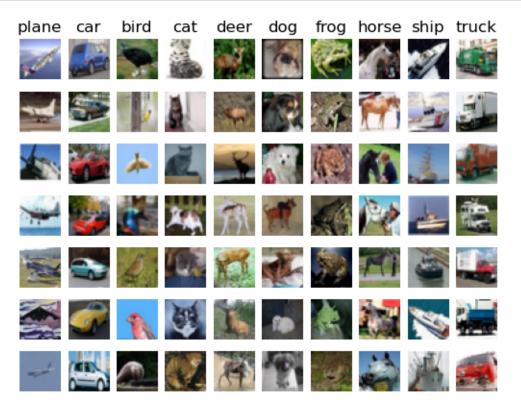
Training data shape: (50000, 32, 32, 3)

Training labels shape: (50000,)

Test data shape: (10000, 32, 32, 3) Test labels shape: (10000,)

```
[6]: # Visualize some examples from the dataset.
     # We show a few examples of training images from each class.
     classes = ['plane', 'car', 'bird', 'cat', 'deer', 'dog', 'frog', 'horse',

     ⇔'ship', 'truck']
     num_classes = len(classes)
     samples_per_class = 7
     for y, cls in enumerate(classes):
         idxs = np.flatnonzero(y_train == y)
         idxs = np.random.choice(idxs, samples_per_class, replace=False)
         for i, idx in enumerate(idxs):
             plt_idx = i * num_classes + y + 1
             plt.subplot(samples_per_class, num_classes, plt_idx)
             plt.imshow(X_train[idx].astype('uint8'))
             plt.axis('off')
             if i == 0:
                 plt.title(cls)
     plt.show()
```



```
[8]: # Subsample the data for more efficient code execution in this exercise
num_training = 5000
mask = list(range(num_training))
X_train = X_train[mask]
y_train = y_train[mask]

num_test = 500
mask = list(range(num_test))
X_test = X_test[mask]
y_test = y_test[mask]

# Reshape the image data into rows
X_train = np.reshape(X_train, (X_train.shape[0], -1))
X_test = np.reshape(X_test, (X_test.shape[0], -1))
print(X_train.shape, X_test.shape)
```

(5000, 3072) (500, 3072)

1 K-nearest neighbors

In the following cells, you will build a KNN classifier and choose hyperparameters via k-fold cross-validation.

```
[15]: # Import the KNN class
from nndl import KNN
```

```
[17]: # Declare an instance of the knn class.
knn = KNN()

# Train the classifier.
# We have implemented the training of the KNN classifier.
# Look at the train function in the KNN class to see what this does.
knn.train(X=X_train, y=y_train)
```

1.1 Questions

- (1) Describe what is going on in the function knn.train().
- (2) What are the pros and cons of this training step?

1.2 Answers

- (1) In the function knn.train(), the input features (X) and their corresponding labels (y) are stored in memory. Each row of X will represent a data point in which the dimensions correspond to the features. The points are plotted in the space as vectors and stored.
- (2) The training step of the KNN is is very simple since storing the input data (X) and the labels corresponding to them (y) without needing to build a complex function. Additionally, the

model allows for minimal computation during training so it is easy handle updates to the dataset. However, storing all of the points and their corresponding features and labels means that it requires a lot of space and memory to store all of the data points. Plus, KNN will be sensistive to noisy data and irrelevant features so it will be less accurate on the predictions unless the data is preprocessed before training.

1.3 KNN prediction

In the following sections, you will implement the functions to calculate the distances of test points to training points, and from this information, predict the class of the KNN.

Time to run code: 14.366737127304077 Frobenius norm of L2 distances: 7906696.077040902

Really slow code Note: This probably took a while. This is because we use two for loops. We could increase the speed via vectorization, removing the for loops.

If you implemented this correctly, evaluating np.linalg.norm (dists_L2, 'fro') should return: $\sim\!7906696$

1.3.1 KNN vectorization

The above code took far too long to run. If we wanted to optimize hyperparameters, it would be time-expensive. Thus, we will speed up the code by vectorizing it, removing the for loops.

```
print('Difference in L2 distances between your KNN implementations (should be<sub>□</sub> ⇒0): {}'.format(np.linalg.norm(dists_L2 - dists_L2_vectorized, 'fro')))
```

```
Time to run code: 0.17363977432250977

Difference in L2 distances between your KNN implementations (should be 0): 0.0
```

Speedup Depending on your computer speed, you should see a 10-100x speed up from vectorization. On our computer, the vectorized form took 0.36 seconds while the naive implementation took 38.3 seconds.

1.3.2 Implementing the prediction

Now that we have functions to calculate the distances from a test point to given training points, we now implement the function that will predict the test point labels.

0.726

If you implemented this correctly, the error should be: 0.726.

This means that the k-nearest neighbors classifier is right 27.4% of the time, which is not great, considering that chance levels are 10%.

2 Optimizing KNN hyperparameters

In this section, we'll take the KNN classifier that you have constructed and perform cross-validation to choose a best value of k, as well as a best choice of norm.

2.0.1 Create training and validation folds

First, we will create the training and validation folds for use in k-fold cross validation.

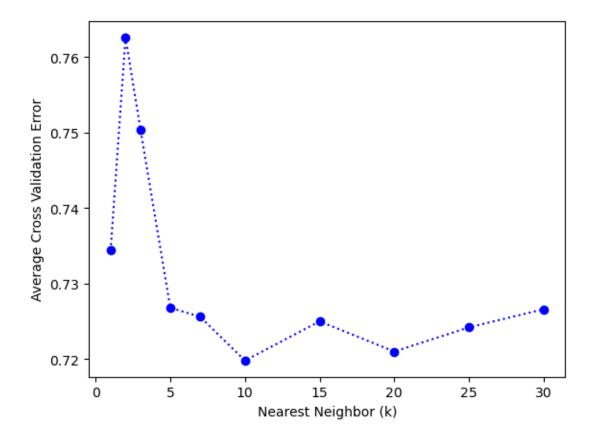
```
[34]: # Create the dataset folds for cross-valdiation.
    num_folds = 5
    X_train_folds = []
    y_train_folds = []
              ______ #
    # YOUR CODE HERE:
       Split the training data into num_folds (i.e., 5) folds.
       X_train_folds is a list, where X_train_folds[i] contains the
         data points in fold i.
       y_train_folds is also a list, where y_train_folds[i] contains
         the corresponding labels for the data in X_train_folds[i]
    # ----- #
    batch = X_train.shape[0]//num_folds
    X_train_folds = np.split(X_train,num_folds)
    y_train_folds = np.split(y_train, num_folds)
    # ----- #
    # END YOUR CODE HERE
    # ----- #
```

2.0.2 Optimizing the number of nearest neighbors hyperparameter.

In this section, we select different numbers of nearest neighbors and assess which one has the lowest k-fold cross validation error.

```
[67]: time start =time.time()
     ks = [1, 2, 3, 5, 7, 10, 15, 20, 25, 30]
     # ----- #
     # YOUR CODE HERE:
       Calculate the cross-validation error for each k in ks, testing
        the trained model on each of the 5 folds. Average these errors
        together and make a plot of k vs. cross-validation error. Since
       we are assuming L2 distance here, please use the vectorized code!
        Otherwise, you might be waiting a long time.
     # ----- #
     def CV_error(num_training, num_of_folds, k_i, norm_i = None):
        cv total error = 0
        for i in range(num_of_folds):
            X_fold_validation = X_train_folds[i]
            y_fold_validation = y_train_folds[i]
            X_fold_train = np.vstack(np.delete(X_train_folds, i, axis = 0))
            y_fold_train = np.hstack(np.delete(y_train_folds, i, axis = 0))
```

```
knn.train(X=X_fold_train, y = y_fold_train)
             if(norm_i):
                dist = knn.compute_distances(X = X_fold_validation, norm = norm_i)
                dist = knn.compute_L2_distances_vectorized(X=X_fold_validation)
            test_fold_number = num_training/num_of_folds
             y_pred = knn.predict_labels(dists = dist, k = k_i)
             wrong_cases = test_fold_number - np.count_nonzero(y_pred ==_
      →y_fold_validation)
             cv_total_error += wrong_cases/test_fold_number
         return cv_total_error/num_of_folds
     avg_errors = [CV_error(num_training,num_folds, ks[k]) for k in range(len(ks))]
      [print("k = {}, Average cross validation error: {}".format(
         ks[k], avg_errors[k])) for k in range(len(ks))]
     # Plotting the results
     fig, ax = plt.subplots(1,1)
     ax.plot(ks, avg_errors, "bo:")
     ax.set_xlabel("Nearest Neighbor (k)")
     ax.set_ylabel("Average Cross Validation Error")
     # ----- #
     # END YOUR CODE HERE
     # ----- #
     print('Computation time: %.2f'%(time.time()-time_start))
     k = 1, Average cross validation error: 0.7344
     k = 2, Average cross validation error: 0.7626000000000002
     k = 3, Average cross validation error: 0.750400000000001
     k = 5, Average cross validation error: 0.726799999999999
     k = 7, Average cross validation error: 0.7256
     k = 10, Average cross validation error: 0.7198
     k = 15, Average cross validation error: 0.725
     k = 20, Average cross validation error: 0.721
     k = 25, Average cross validation error: 0.7242
     k = 30, Average cross validation error: 0.7266
     Computation time: 23.21
[68]: plt.show()
```



2.1 Questions:

- (1) What value of k is best amongst the tested k's?
- (2) What is the cross-validation error for this value of k?

2.2 Answers:

- (1) Based on the printed results for each value of k, the best value of k is ten as it has the lowest cross validation error out of all of the values.
- (2) The average cross validation error for when k is 10 is 0.7198 or 71.98%

2.2.1 Optimizing the norm

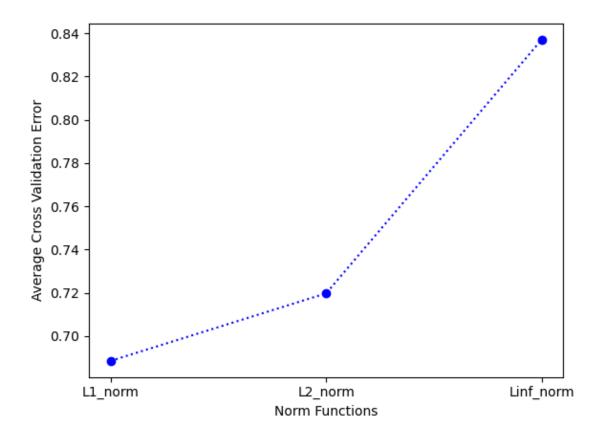
Next, we test three different norms (the 1, 2, and infinity norms) and see which distance metric results in the best cross-validation performance.

```
[79]: time_start =time.time()

L1_norm = lambda x: np.linalg.norm(x, ord=1)
L2_norm = lambda x: np.linalg.norm(x, ord=2)
Linf_norm = lambda x: np.linalg.norm(x, ord= np.inf)
```

```
norms = [L1_norm, L2_norm, Linf_norm]
     # ----- #
        Calculate the cross-validation error for each norm in norms, testing
        the trained model on each of the 5 folds. Average these errors
        together and make a plot of the norm used vs the cross-validation error
        Use the best cross-validation k from the previous part.
       Feel free to use the compute_distances function. We're testing just
        three norms, but be advised that this could still take some time.
       You're welcome to write a vectorized form of the L1- and Linf- norms
       to speed this up, but it is not necessary.
     num_of_folds = num_folds
     k=10
     average_error = [CV_error(num_training, num_of_folds, k_i = k, norm_i = norm)_
      ofor norm in norms]
     normalization_types = ["L1_norm", "L2_norm", "Linf_norm"]
     [print("k = {}, Average cross validation error: {}".

¬format(normalization types[k], avg errors[k])) for k in range(len(norms))]
     # END YOUR CODE HERE
     print('Computation time: %.2f'%(time.time()-time_start))
    k = L1_norm, Average cross validation error: 0.7344
    k = L2_norm, Average cross validation error: 0.7626000000000002
    k = Linf_norm, Average cross validation error: 0.750400000000001
    Computation time: 380.45
[90]: fig, ax = plt.subplots(1, 1)
     ax.plot(average error, "bo:")
     ax.set_xlabel("Norm Functions")
     ax.set_xticks(np.arange(3), normalization_types)
     ax.set_ylabel("Average Cross Validation Error")
     plt.show()
```



2.3 Questions:

- (1) What norm has the best cross-validation error?
- (2) What is the cross-validation error for your given norm and k?

2.4 Answers:

- (1) The L1 norm has the best cross validation error out of the 3 types of norms possible.
- (2) The cross validation error for my L1_norm was 0.7344 or 73.44%

3 Evaluating the model on the testing dataset.

Now, given the optimal k and norm you found in earlier parts, evaluate the testing error of the k-nearest neighbors model.

```
for your optimal hyperparameters found by 5-fold cross-validation.
# Optimal k = 10, L1 Norm
optimal_norm = lambda x: np.linalg.norm(x, ord= 1)
knn.train(X = X_train, y = y_train)
test_dists = knn.compute_distances(X = X_test, norm = optimal_norm)
test_predictions = knn.predict_labels(dists = test_dists, k = 10)
total_incorrect = num_test - np.count_nonzero(test_predictions == y_test)
total_error = total_incorrect/num_test
# Naive baseline: k = 1, L2-norm
baseline norm = lambda x: np.linalg.norm(x, ord=2)
baseline dists = knn.compute distances(X=X test, norm=baseline norm)
baseline predictions = knn.predict_labels(dists=baseline_dists, k=1)
print(f"Baseline error (k=1, L2-norm): {baseline_error}")
print(f"Optimal error (k=10, L1-norm): {total_error}")
print(f"Improvement in error: {error_improvement}")
# ----- #
# END YOUR CODE HERE
print('Error rate achieved: {}'.format(error))
```

Baseline error (k=1, L2-norm): 0.726 Optimal error (k=10, L1-norm): 0.722 Improvement in error: 0.004000000000000036

Error rate achieved: 1

3.1 Question:

How much did your error improve by cross-validation over naively choosing k = 1 and using the L2-norm?

3.2 Answer:

There wasn't that much of an improvement. The baseline error rate was actually higher than the optimal error rate, but only by a marginal number of 0.0004. So in this case, naively choosing k=1 and L2 norm isn't much of an improvement for our error.