## k-Nearest Neighbor (kNN) exercise

Complete and hand in this completed worksheet (including its outputs and any supporting code outside of the worksheet) with your assignment submission. For more details see the <u>assignments page (https://compsci682-fa19.github.io/assignments2019/assignment1)</u> on the course website.

The kNN classifier consists of two stages:

- · During training, the classifier takes the training data and simply remembers it
- During testing, kNN classifies every test image by comparing to all training images and transfering the labels of the k most similar training examples
- · The value of k is cross-validated

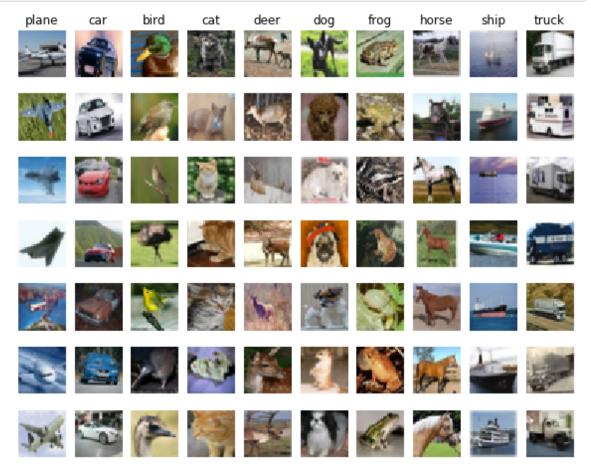
In this exercise you will implement these steps and understand the basic Image Classification pipeline, cross-validation, and gain proficiency in writing efficient, vectorized code.

```
In [121]:
          # Run some setup code for this notebook.
          from future import print function
          import random
          import numpy as np
          from cs682.data utils import load CIFAR10
          import matplotlib.pyplot as plt
          # This is a bit of magic to make matplotlib figures appear inline in
           the notebook
          # rather than in a new window.
          %matplotlib inline
          plt.rcParams['figure.figsize'] = (10.0, 8.0) # set default size of pl
          plt.rcParams['image.interpolation'] = 'nearest'
          plt.rcParams['image.cmap'] = 'gray'
          # Some more magic so that the notebook will reload external python mo
          dules;
          # see http://stackoverflow.com/questions/1907993/autoreload-of-module
          s-in-ipython
          %load ext autoreload
          %autoreload 2
```

The autoreload extension is already loaded. To reload it, use: %reload ext autoreload

```
# Load the raw CIFAR-10 data.
In [122]:
          cifar10 dir = 'cs682/datasets/cifar-10-batches-py'
          # Cleaning up variables to prevent loading data multiple times (which
          may cause memory issue)
          try:
             del X_train, y_train
             del X test, y_test
             print('Clear previously loaded data.')
          except:
             pass
          X_train, y_train, X_test, y_test = load_CIFAR10(cifar10_dir)
          # As a sanity check, we print out the size of the training and test d
          ata.
          print('Training data shape: ', X_train.shape)
          print('Training labels shape: ', y_train.shape)
          print('Test data shape: ', X_test.shape)
          print('Test labels shape: ', y_test.shape)
          Clear previously loaded data.
          Training data shape: (50000, 32, 32, 3)
          Training labels shape: (50000,)
          Test data shape: (10000, 32, 32, 3)
          Test labels shape: (10000,)
```

```
In [111]: # 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', 'hor
           se', 'ship', 'truck']
num_classes = len(classes)
           samples_per_class = 7
           for y, cls \overline{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()
```



```
# Subsample the data for more efficient code execution in this exerci
In [123]:
           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_{\text{test}} = X_{\text{test}}[mask]
           y_{\text{test}} = y_{\text{test}}[mask]
In [124]: # Reshape the image data into rows
           X_train = np.reshape(X_train, (X_train.shape[0], -1))
           X_{\text{test}} = \text{np.reshape}(X_{\text{test}}, (X_{\text{test.shape}}[0], -1))
           print(X train.shape, X test.shape)
           (5000, 3072) (500, 3072)
In [125]: from cs682.classifiers import KNearestNeighbor
           # Create a kNN classifier instance.
           # Remember that training a kNN classifier is a noop:
           # the Classifier simply remembers the data and does no further proces
           classifier = KNearestNeighbor()
           classifier.train(X train, y train)
```

We would now like to classify the test data with the kNN classifier. Recall that we can break down this process into two steps:

- 1. First we must compute the distances between all test examples and all train examples.
- 2. Given these distances, for each test example we find the k nearest examples and have them vote for the label

Lets begin with computing the distance matrix between all training and test examples. For example, if there are **Ntr** training examples and **Nte** test examples, this stage should result in a **Nte** x **Ntr** matrix where each element (i,j) is the distance between the i-th test and j-th train example.

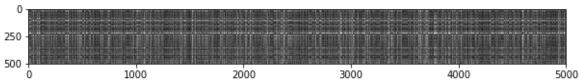
First, open cs682/classifiers/k\_nearest\_neighbor.py and implement the function compute\_distances\_two\_loops that uses a (very inefficient) double loop over all pairs of (test, train) examples and computes the distance matrix one element at a time.

```
In [126]: # Open cs682/classifiers/k_nearest_neighbor.py and implement
# compute_distances_two_loops.

# Test your implementation:
dists = classifier.compute_distances_two_loops(X_test)
print(dists.shape)

(500, 5000)
```

```
In [115]: # We can visualize the distance matrix: each row is a single test exa
    mple and
    # its distances to training examples
    plt.imshow(dists, interpolation='none')
    plt.show()
```



**Inline Question #1:** Notice the structured patterns in the distance matrix, where some rows or columns are visible brighter. (Note that with the default color scheme black indicates low distances while white indicates high distances.)

- What in the data is the cause behind the distinctly bright rows?
- · What causes the columns?

## Your Answer:

- 1. Distinctly bright rows means that their distance from other points is huge. It is possible that certain points in testing set are really far from the training data points, which is why their distance is very high. These images must be very different from most of the training set, making them outliers.
- 2. Bright columns are caused by images which are outliers in training set.

```
In [127]: # Now implement the function predict_labels and run the code below:
    # We use k = 1 (which is Nearest Neighbor).
    y_test_pred = classifier.predict_labels(dists, k=1)

# Compute and print the fraction of correctly predicted examples
    num_correct = np.sum(y_test_pred == y_test)
    accuracy = float(num_correct) / num_test
    print('Got %d / %d correct => accuracy: %f' % (num_correct, num_test, accuracy))
```

Got 137 / 500 correct => accuracy: 0.274000

You should expect to see approximately 27% accuracy. Now lets try out a larger k, say k = 5:

```
In [128]: y_test_pred = classifier.predict_labels(dists, k=5)
    num_correct = np.sum(y_test_pred == y_test)
    accuracy = float(num_correct) / num_test
    print('Got %d / %d correct => accuracy: %f' % (num_correct, num_test, accuracy))
```

Got 139 / 500 correct => accuracy: 0.278000

You should expect to see a slightly better performance than with k = 1.

**Inline Question 2** We can also other distance metrics such as L1 distance. The performance of a Nearest Neighbor classifier that uses L1 distance will not change if (Select all that apply.):

- 1. The data is preprocessed by subtracting the mean.
- 2. The data is preprocessed by subtracting the mean and dividing by the standard deviation.
- 3. The coordinate axes for the data are rotated.
- None of the above.

Your Answer: 1,2 - TRUE

*Your explanation*: Boundaries for nearest neighbour classifer (NNC) that uses L1 distance are in the shape of a rotated square.

- 1. When any constant is removed from all the points in the data, their relative positions remain unchanged, hence L1 distance between them remain unchanged. hence the performance remains the same.
- 2. When constant is removed and divided by a single number, their L1 distance is compressed by that factor but their relative distance remain unchanged. Hence, the performance remains the same.
- 3. When axes are totated, L1 distance between the points changes unequally, resulting in a different performance. So, this is not true.

```
In [129]:
          # Now lets speed up distance matrix computation by using partial vect
          # with one loop. Implement the function compute distances one loop an
          d run the
          # code below:
          dists one = classifier.compute distances one loop(X test)
          # To ensure that our vectorized implementation is correct, we make su
          re that it
          # agrees with the naive implementation. There are many ways to decide
          whether
          # two matrices are similar; one of the simplest is the Frobenius nor
          m. In case
          # you haven't seen it before, the Frobenius norm of two matrices is t
          he square
          # root of the squared sum of differences of all elements; in other wo
          rds, reshape
          # the matrices into vectors and compute the Euclidean distance betwee
          n them.
          difference = np.linalg.norm(dists - dists one, ord='fro')
          print('Difference was: %f' % (difference, ))
          if difference < 0.001:</pre>
              print('Good! The distance matrices are the same')
          else:
              print('Uh-oh! The distance matrices are different')
```

Difference was: 0.000000 Good! The distance matrices are the same

```
In [130]:
          # Now implement the fully vectorized version inside compute distances
          _no_loops
          # and run the code
          dists two = classifier.compute distances no loops(X test)
          # check that the distance matrix agrees with the one we computed befo
          difference = np.linalq.norm(dists - dists two, ord='fro')
          print('Difference was: %f' % (difference, ))
          if difference < 0.001:</pre>
              print('Good! The distance matrices are the same')
          else:
              print('Uh-oh! The distance matrices are different')
          Difference was: 0.000000
          Good! The distance matrices are the same
In [131]:
          # Let's compare how fast the implementations are
          def time function(f, *args):
              Call a function f with args and return the time (in seconds) that
          it took to execute.
              import time
              tic = time.time()
              f(*args)
              toc = time.time()
              return toc - tic
          two_loop_time = time_function(classifier.compute_distances_two_loops,
          X test)
          print('Two loop version took %f seconds' % two loop time)
          one loop time = time function(classifier.compute distances one loop,
          X test)
          print('One loop version took %f seconds' % one loop time)
          no loop time = time function(classifier.compute distances no loops, X
          _test)
          print('No loop version took %f seconds' % no loop time)
          # you should see significantly faster performance with the fully vect
          orized implementation
          Two loop version took 17.916950 seconds
```

Two loop version took 17.916950 seconds One loop version took 42.243923 seconds No loop version took 0.161431 seconds

## **Cross-validation**

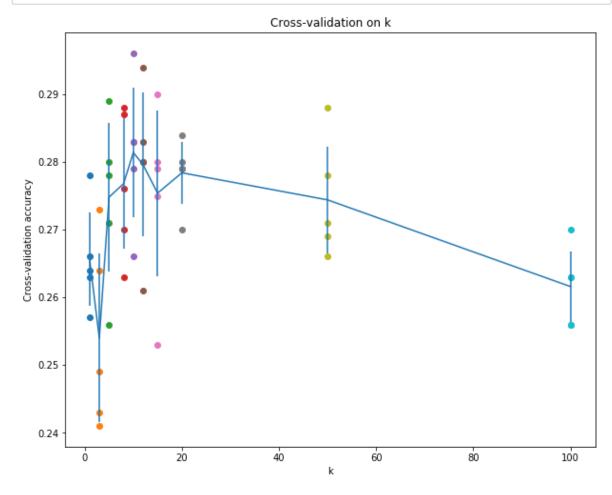
We have implemented the k-Nearest Neighbor classifier but we set the value k = 5 arbitrarily. We will now determine the best value of this hyperparameter with cross-validation.

```
In [132]:
       num folds = 5
        k \text{ choices} = [1, 3, 5, 8, 10, 12, 15, 20, 50, 100]
        X train folds = []
        y train folds = []
        ##########
        # TODO:
        # Split up the training data into folds. After splitting, X train fol
        ds and
        # y train folds should each be lists of length num folds, where
        # y train folds[i] is the label vector for the points in X train fold
        s[i].
        # Hint: Look up the numpy array split function.
        ##########
        # Your code
       X train folds = np.split(X train, num folds)
        y_train_folds = np.split(y_train, num_folds)
        ##########
        #
                                 END OF YOUR CODE
        ##########
        # A dictionary holding the accuracies for different values of k that
        # when running cross-validation. After running cross-validation,
        # k to accuracies[k] should be a list of length num folds giving the
        # accuracy values that we found when using that value of k.
        k to accuracies = {}
        ##########
        # TODO:
        # Perform k-fold cross validation to find the best value of k. For ea
        # possible value of k, run the k-nearest-neighbor algorithm num folds
        times,
        # where in each case you use all but one of the folds as training dat
        a and the #
        # last fold as a validation set. Store the accuracies for all fold an
        d all
        # values of k in the k to accuracies dictionary.
        ##########
        # Your code
```

```
for k in k choices:
   classifier = KNearestNeighbor()
   accuracies = []
   for fold in np.arange(num folds):
      X train current = np.zeros((0,X train folds[0].shape[1]), dty
pe = int)
      y train current = np.zeros((0,), dtype = int)
       for i in np.arange(num folds):
          if i != fold:
             X train current = np.concatenate((X train current, X
train folds[i]), axis = 0)
             y_train_current = np.concatenate((y_train_current, y_
train folds[i]), axis = 0)
       classifier.train(X_train_current, y_train_current)
       y test pred = classifier.predict(X train folds[fold], k=k)
       num_correct = np.sum(y_test_pred == y_train_folds[fold])
       accuracy = float(num correct) / y train folds[fold].shape[0]
       accuracies.append(accuracy)
   k to accuracies[k] = accuracies
###########
#
                             END OF YOUR CODE
##########
# Print out the computed accuracies
for k in sorted(k_to_accuracies):
   for accuracy in k to accuracies[k]:
       print('k = %d, accuracy = %f' % (k, accuracy))
```

k = 1, accuracy = 0.263000 k = 1, accuracy = 0.257000 k = 1, accuracy = 0.264000 k = 1, accuracy = 0.278000 k = 1, accuracy = 0.266000 k = 3, accuracy = 0.241000 k = 3, accuracy = 0.249000 k = 3, accuracy = 0.243000 k = 3, accuracy = 0.273000 k = 3, accuracy = 0.264000 k = 5, accuracy = 0.256000 k = 5, accuracy = 0.271000 k = 5, accuracy = 0.280000 k = 5, accuracy = 0.289000 k = 5, accuracy = 0.278000 k = 8, accuracy = 0.263000 k = 8, accuracy = 0.287000 k = 8, accuracy = 0.276000 k = 8, accuracy = 0.288000 k = 8, accuracy = 0.270000 k = 10, accuracy = 0.266000 k = 10, accuracy = 0.296000 k = 10, accuracy = 0.279000 k = 10, accuracy = 0.283000 k = 10, accuracy = 0.283000 k = 12, accuracy = 0.261000 k = 12, accuracy = 0.294000 k = 12, accuracy = 0.280000 k = 12, accuracy = 0.283000 k = 12, accuracy = 0.280000 k = 15, accuracy = 0.253000 k = 15, accuracy = 0.290000 k = 15, accuracy = 0.279000 k = 15, accuracy = 0.280000 k = 15, accuracy = 0.275000 k = 20, accuracy = 0.270000 k = 20, accuracy = 0.279000 k = 20, accuracy = 0.279000 k = 20, accuracy = 0.280000 k = 20, accuracy = 0.284000 k = 50, accuracy = 0.271000 k = 50, accuracy = 0.288000 k = 50, accuracy = 0.278000 k = 50, accuracy = 0.269000 k = 50, accuracy = 0.266000 k = 100, accuracy = 0.256000 k = 100, accuracy = 0.270000 k = 100, accuracy = 0.263000 k = 100, accuracy = 0.256000 k = 100, accuracy = 0.263000

```
In [133]:
          # plot the raw observations
          for k in k choices:
              accuracies = k_to_accuracies[k]
              plt.scatter([k] * len(accuracies), accuracies)
          # plot the trend line with error bars that correspond to standard dev
          iation
          accuracies mean = np.array([np.mean(v) for k,v in sorted(k to accurac
          ies.items())])
          accuracies_std = np.array([np.std(v) for k,v in sorted(k_to_accuracie
          s.items())])
          plt.errorbar(k_choices, accuracies_mean, yerr=accuracies_std)
          plt.title('Cross-validation on k')
          plt.xlabel('k')
          plt.ylabel('Cross-validation accuracy')
          plt.show()
```



```
In [134]: # Based on the cross-validation results above, choose the best value
    for k,
    # retrain the classifier using all the training data, and test it on
        the test
    # data. You should be able to get above 28% accuracy on the test dat
    a.
    best_k = 10

classifier = KNearestNeighbor()
    classifier.train(X_train, y_train)
    y_test_pred = classifier.predict(X_test, k=best_k)

# Compute and display the accuracy
    num_correct = np.sum(y_test_pred == y_test)
    accuracy = float(num_correct) / num_test
    print('Got %d / %d correct => accuracy: %f' % (num_correct, num_test, accuracy))
```

Got 141 / 500 correct => accuracy: 0.282000

**Inline Question 3** Which of the following statements about k-Nearest Neighbor (k-NN) are true in a classification setting, and for all k? Select all that apply.

- 1. The training error of a 1-NN will always be better than that of 5-NN.
- 2. The test error of a 1-NN will always be better than that of a 5-NN.
- 3. The decision boundary of the k-NN classifier is linear.
- 4. The time needed to classify a test example with the k-NN classifier grows with the size of the training set.
- 5. None of the above.

Your Answer: 1,4 - TRUE

## Your explanation:

- 1. Training error of 1-NN is zero since we are seeing the difference between the same set of points. This will always be better than or equal to that of 5-NN. (Equality in very special case, where all the points of each class are closer to themselves than the other class). So, this is TRUE.
- 2. Test error of 1-NN can be better or worse than 5-NN depending on the case. So, this is FALSE.
- 3. Boundary is not linear. Consider three concentric circles of points. Here, the boundaries would most probably be circular. Hence, FALSE.
- 4. Time need to classify a test example is the time needed to evaluate the distance of the test example from each of the training example and taking the least k-distances. This computation time increases with the number of training examples. So, this is TRUE.