knn

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1 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 assignments page 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 [1]: # Run some setup code for this notebook.
        import random
        import numpy as np
        from cs231n.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 plots
        plt.rcParams['image.interpolation'] = 'nearest'
        plt.rcParams['image.cmap'] = 'gray'
        # Some more magic so that the notebook will reload external python modules;
        # see http://stackoverflow.com/questions/1907993/autoreload-of-modules-in-ipython
        %load_ext autoreload
        %autoreload 2
In [2]: # Load the raw CIFAR-10 data.
        cifar10_dir = 'cs231n/datasets/cifar-10-batches-py'
        # Cleaning up variables to prevent loading data multiple times (which may cause memory i
```

```
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 data.
        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)
Training data shape: (50000, 32, 32, 3)
Training labels shape: (50000,)
Test data shape: (10000, 32, 32, 3)
Test labels shape: (10000,)
In [3]: # 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()
```



```
In [4]: # 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)
In [5]: from cs231n.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 processing
        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 $Nte \times Ntr$ matrix where each element (i,j) is the distance between the i-th test and j-th train example.

Note: For the three distance computations that we require you to implement in this notebook, you may not use the np.linalg.norm() function that numpy provides.

First, open cs231n/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 [6]: # Open cs231n/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 [7]: # We can visualize the distance matrix: each row is a single test example and
        # its distances to training examples
        plt.imshow(dists, interpolation='none')
        plt.show()
           0
        250
        500
                       1000
                                   2000
                                                3000
                                                            4000
                                                                        5000
```

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: Bright rows indicate high distances which mean image are far away from other images.

```
In [8]: # 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:

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

Inline Question 2

We can also use other distance metrics such as L1 distance. For pixel values $p_{ij}^{(k)}$ at location (i, j) of some image I_k ,

the mean μ across all pixels over all images is

$$\mu = \frac{1}{nhw} \sum_{k=1}^{n} \sum_{i=1}^{h} \sum_{j=1}^{w} p_{ij}^{(k)}$$

And the pixel-wise mean μ_{ij} across all images is

$$\mu_{ij} = \frac{1}{n} \sum_{k=1}^{n} p_{ij}^{(k)}.$$

The general standard deviation σ and pixel-wise standard deviation σ_{ij} is defined similarly.

Which of the following preprocessing steps will not change the performance of a Nearest Neighbor classifier that uses L1 distance? Select all that apply. 1. Subtracting the mean μ ($\tilde{p}_{ij}^{(k)} = p_{ij}^{(k)} - \mu$.) 2. Subtracting the per pixel mean μ_{ij} ($\tilde{p}_{ij}^{(k)} = p_{ij}^{(k)} - \mu_{ij}$.) 3. Subtracting the mean μ and dividing by the standard deviation σ . 4. Subtracting the pixel-wise mean μ_{ij} and dividing by the pixel-wise standard deviation σ_{ij} . 5. Rotating the coordinate axes of the data.

\$Your Answer:

1, 2, 3, 5

Your Explanation:

 L_1 -distance (Manhattan distance) between images I_l and I_m :

$$L_1(I_l, I_m) = \sum_{i=1}^h \sum_{j=1}^w |p_{ij}^{(l)} - p_{ij}^{(m)}|$$

1.

$$\tilde{L}_1(I_l, I_m) = \sum_{i=1}^h \sum_{j=1}^w |(p_{ij}^{(l)} - \mu) - (p_{ij}^{(m)} - \mu)| = L_1(I_l, I_m),$$

The results are the same with each other and this step will not change the performance.

2.

$$ilde{L}_1(I_l,I_m) = \sum_{i=1}^h \sum_{j=1}^w |(p_{ij}^{(l)} - \mu_{ij}) - (p_{ij}^{(m)} - \mu_{ij})| = L_1(I_l,I_m),$$

This step will not change the performance as well.

3.

$$ilde{L}_1(I_l,I_m) = \sum_{i=1}^h \sum_{j=1}^w |rac{p_{ij}^{(l)} - \mu}{\sigma} - rac{p_{ij}^{(m)} - \mu}{\sigma}| = rac{1}{\sigma} L_1(I_l,I_m)$$

All distances between images are shrinked by a factor of σ , but it will not not change the performance of classification, because all distances are changed in the same time.

4.

$$\tilde{L}_{1}(I_{l},I_{m}) = \sum_{i=1}^{h} \sum_{j=1}^{w} |\frac{p_{ij}^{(l)} - \mu_{ij}}{\sigma_{ij}} - \frac{p_{ij}^{(m)} - \mu_{ij}}{\sigma_{ij}}| = \sum_{i=1}^{h} \sum_{j=1}^{w} |\frac{p_{ij}^{(l)} - p_{ij}^{(m)}}{\sigma_{ij}}| \neq L_{1}(I_{l},I_{m}),$$

This step will change results.

Good! The distance matrices are the same

5. Rotating the coordinate axes will not change the performance.

```
In [10]: # Now lets speed up distance matrix computation by using partial vectorization
         # with one loop. Implement the function compute_distances_one_loop and run the
         # code below:
         dists_one = classifier.compute_distances_one_loop(X_test)
         # To ensure that our vectorized implementation is correct, we make sure 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 norm. In case
         # you haven't seen it before, the Frobenius norm of two matrices is the square
         # root of the squared sum of differences of all elements; in other words, reshape
         # the matrices into vectors and compute the Euclidean distance between them.
         difference = np.linalg.norm(dists - dists_one, ord='fro')
         print('One loop difference was: %f' % (difference, ))
         if difference < 0.001:
             print('Good! The distance matrices are the same')
         else:
             print('Uh-oh! The distance matrices are different')
One loop difference was: 0.000000
```

```
In [11]: # 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 before:
         difference = np.linalg.norm(dists - dists_two, ord='fro')
         print('No loop 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')
No loop difference was: 0.000000
Good! The distance matrices are the same
In [12]: # Let's compare how fast the implementations are
         def time_function(f, *args):
             11 11 11
             Call a function f with args and return the time (in seconds) that it took to execut
             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 vectorized implementat
         # NOTE: depending on what machine you're using,
         # you might not see a speedup when you go from two loops to one loop,
         # and might even see a slow-down.
Two loop version took 35.683620 seconds
One loop version took 38.481179 seconds
No loop version took 0.231135 seconds
```

1.0.1 Cross-validation

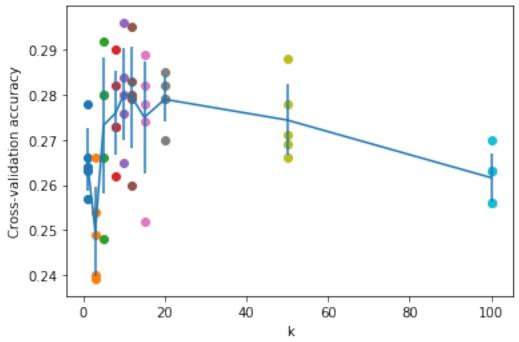
We have implemented the k-Nearest Neighbor classifier but we set the value k = 5 arbitrarily. We will now determine the best value of this hyperparameter with cross-validation.

```
In \lceil 13 \rceil: num folds = 5
       k_{choices} = [1, 3, 5, 8, 10, 12, 15, 20, 50, 100]
       X_train_folds = []
       v_train_folds = []
       # TODO:
       # Split up the training data into folds. After splitting, X_train_folds 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_folds[i].
       # Hint: Look up the numpy array_split function.
       # *****START OF YOUR CODE (DO NOT DELETE/MODIFY THIS LINE)*****
       X_train_folds = np.array(np.array_split(X_train, num_folds))
       y_train_folds = np.array(np.array_split(y_train, num_folds))
       # *****END OF YOUR CODE (DO NOT DELETE/MODIFY THIS LINE) ****
       # A dictionary holding the accuracies for different values of k that we find
       # when running cross-validation. After running cross-validation,
       # k_to_accuracies[k] should be a list of length num_folds giving the different
       # 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 each
       # 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 data and the #
       # last fold as a validation set. Store the accuracies for all fold and all
       # values of k in the k_to_accuracies dictionary.
       # *****START OF YOUR CODE (DO NOT DELETE/MODIFY THIS LINE)****
       for n in k_choices:
           accuracies = []
           for i in range(num_folds):
              m = np.array([j for j in range(5) if j!=i])
              X_train = np.vstack(X_train_folds[m])
              y_train = np.vstack(y_train_folds[m]).flatten()
              classifier.train(X_train,y_train)
              y_pred = classifier.predict(X_train_folds[i], n, num_loops=0)
              num_correct = np.sum(y_pred == y_train_folds[i])
```

```
accuracies.append(num_correct/len(y_pred))
             k_to_accuracies[n] = accuracies
         # *****END OF YOUR CODE (DO NOT DELETE/MODIFY THIS LINE)****
         # 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 = 0.239000
k = 3, accuracy = 0.249000
k = 3, accuracy = 0.240000
k = 3, accuracy = 0.266000
k = 3, accuracy = 0.254000
k = 5, accuracy = 0.248000
k = 5, accuracy = 0.266000
k = 5, accuracy = 0.280000
k = 5, accuracy = 0.292000
k = 5, accuracy = 0.280000
k = 8, accuracy = 0.262000
k = 8, accuracy = 0.282000
k = 8, accuracy = 0.273000
k = 8, accuracy = 0.290000
k = 8, accuracy = 0.273000
k = 10, accuracy = 0.265000
k = 10, accuracy = 0.296000
k = 10, accuracy = 0.276000
k = 10, accuracy = 0.284000
k = 10, accuracy = 0.280000
k = 12, accuracy = 0.260000
k = 12, accuracy = 0.295000
k = 12, accuracy = 0.279000
k = 12, accuracy = 0.283000
k = 12, accuracy = 0.280000
k = 15, accuracy = 0.252000
k = 15, accuracy = 0.289000
k = 15, accuracy = 0.278000
k = 15, accuracy = 0.282000
k = 15, accuracy = 0.274000
k = 20, accuracy = 0.270000
k = 20, accuracy = 0.279000
k = 20, accuracy = 0.279000
```

```
k = 20, accuracy = 0.282000
k = 20, accuracy = 0.285000
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 [14]: # 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 deviation
         accuracies_mean = np.array([np.mean(v) for k,v in sorted(k_to_accuracies.items())])
         accuracies_std = np.array([np.std(v) for k,v in sorted(k_to_accuracies.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()
```

Cross-validation on k



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 decision boundary of the k-NN classifier is linear. 2. The training error of a 1-NN will always be lower than that of 5-NN. 3. The test error of a 1-NN will always be lower than that of a 5-NN. 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: 2 and 4.
Your Explanation:

- *statement 1 is incorrect, because the boundary is non-linear.*
- statement 2 is correct, because training error of 1-NN is always zero. If given a point x, the nearest neighbor will be the exact same point and thus the error will be 0.
- *statement 3 is incorrect, because it's too absolute that 1-NN will work better.*
- statement 4 is correct, test time complexity is depend on the training set length.