# 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.

## 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/autoreload-of-modules-in-ipython
%load_ext autoreload
%autoreload 2
```

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

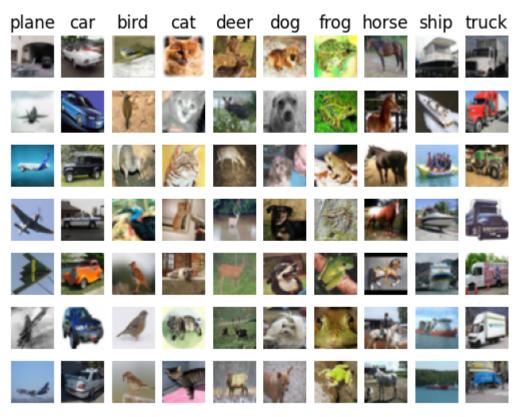
```
In [27]: # Set the path to the CIFAR-10 data
    cifar10_dir = 'C:/Users/jessi/OneDrive/Desktop/classes/c147/HW2_code/cifar-10-batch
    X_train, y_train, X_test, y_test = load_CIFAR10(cifar10_dir)

# 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 [28]: # 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', '
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 [29]: # 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)
```

# K-nearest neighbors

(5000, 3072) (500, 3072)

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

```
In [30]: # Import the KNN class
from nndl import KNN

In [31]: # 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)
```

#### Questions

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

#### **Answers**

- (1) knn.train() is setting the training data to the first 500 examples in the CIFAR-10 dataset.
- (2) Pro: It is simple and fast. Con: It is memory intensive because we need to store all the data.

#### **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.

```
In [32]: # Implement the function compute_distances() in the KNN class.
# Do not worry about the input 'norm' for now; use the default definition of the no
# in the code, which is the 2-norm.
# You should only have to fill out the clearly marked sections.

import time
time_start =time.time()

dists_L2 = knn.compute_distances(X=X_test)

print('Time to run code: {}'.format(time.time()-time_start))
print('Frobenius norm of L2 distances: {}'.format(np.linalg.norm(dists_L2, 'fro')))
```

Time to run code: 25.525479793548584

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: ~7906696

#### 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.

```
In [33]: # Implement the function compute_L2_distances_vectorized() in the KNN class.
# In this function, you ought to achieve the same L2 distance but WITHOUT any for L
# Note, this is SPECIFIC for the L2 norm.

time_start =time.time()
dists_L2_vectorized = knn.compute_L2_distances_vectorized(X=X_test)
print('Time to run code: {}'.format(time.time()-time_start))
print('Difference in L2 distances between your KNN implementations (should be 0): {
Time to run code: 0.27126359939575195
```

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.

#### 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%.

# **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.

#### Create training and validation folds

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

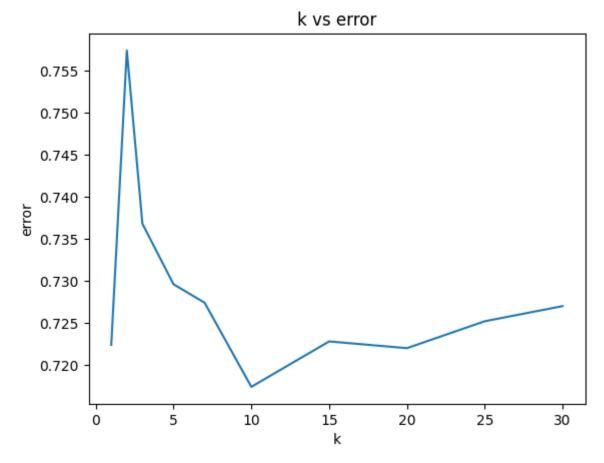
```
# Create the dataset folds for cross-valdiation.
In [35]:
       num_folds = 5
       X_train_folds = []
       y_train_folds = []
       X_val_folds = []
       y_val_folds = []
        # YOUR CODE HERE:
       # Split the training data into num_folds (i.e., 5) folds.
       # X_train_folds is a 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]
        # ------ #
       num_training = 5000
        cv_idx = np.arange(num_training)
        np.random.shuffle(cv_idx)
       fold_size = num_training // num_folds
       for i in np.arange(num_folds):
           val_idx = cv_idx[i*fold_size:(i+1)*fold_size]
```

#### 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.

```
In [36]: 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.
        errors = np.zeros(len(ks))
        for k in range(len(ks)):
           fold_errors = np.empty(num_folds)
           for i in np.arange(num_folds):
              knn.train(X=X_train_folds[i], y=y_train_folds[i])
              dists_L2_vectorized = knn.compute_L2_distances_vectorized(X=X_val_folds[i])
              y_pred = knn.predict_labels(dists=dists_L2_vectorized, k=ks[k])
              error = np.sum(y_pred != y_val_folds[i]) / len(y_pred)
              fold_errors[i] = error
           errors[k] = np.mean(fold_errors)
        print("best error was: ", str(np.min(errors)), "with k=", str(ks[np.argmin(errors)]
       fig1 = plt.figure()
        plt.plot(ks, errors)
        plt.title("k vs error")
        plt.xlabel("k")
       plt.ylabel("error")
       plt.show()
        pass
        # END YOUR CODE HERE
```

```
# ======== #
print('Computation time: %.2f'%(time.time()-time_start))
```



Computation time: 28.10

# **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?

#### **Answers:**

- (1) The best value was k=10
- (2) Th cross-validation error for k = 10 was 0.7174.

#### 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.

```
In [47]: 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]
        names = ["L1_norm", "L2_norm", "Linf_norm"]
        # YOUR CODE HERE:
        # 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.
        errors = np.empty(len(norms))
        for n in range(len(norms)):
           fold_errors = np.empty(num_folds)
           for i in np.arange(num_folds):
              knn.train(X=X_train_folds[i], y=y_train_folds[i])
              dist = knn.compute_distances(X=X_val_folds[i], norm=norms[n])
              y_pred = knn.predict_labels(dists=dist, k=10)
              error = np.sum(y_pred != y_val_folds[i]) / len(y_pred)
              fold_errors[i] = error
           errors[n] = np.mean(fold_errors)
        fig2 = plt.figure()
        plt.bar(names, errors)
        plt.title("loss type vs error")
        plt.xlabel("loss")
        plt.ylabel("error")
        plt.show()
        pass
        # ------ #
        # END YOUR CODE HERE
        print('Computation time: %.2f'%(time.time()-time_start))
```

# 0.8 - 0.7 - 0.6 - 0.5 - 0.3 - 0.2 - 0.1 - 0.0 - 0.1 - 0.5 -

Computation time: 561.75

```
In [48]: print("best error was:", str(np.min(errors)), "with norm =", str(names[np.argmin(er
best error was: 0.6876 with norm = L1_norm
```

#### **Questions:**

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

#### **Answers:**

- (1) L1-norm had the best cross-validation error
- (2) 0.6876 with norm =  $L1_norm$

# 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.

Error rate achieved: 0.722

## **Question:**

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

#### **Answer:**

My error improved by 0.726-0.722 = 0.004. Lol. Less than half a percent.

```
import numpy as np
import pdb
class KNN(object):
 def init (self):
   pass
 def train(self, X, y):
   Inputs:
   - X is a numpy array of size (num examples, D)
   - y is a numpy array of size (num examples, )
   self.X train = X
   self.y train = y
 def compute distances(self, X, norm=None):
   Compute the distance between each test point in X and each training point
   in self.X train.
   Inputs:
   - X: A numpy array of shape (num test, D) containing test data.
   - norm: the function with which the norm is taken.
   Returns:
   - dists: A numpy array of shape (num test, num train) where dists[i, j]
    is the Euclidean distance between the ith test point and the jth training
    point.
   ,,,,,,
   if norm is None:
    norm = lambda x: np.sqrt(np.sum(x**2))
     \#norm = 2
   num test = X.shape[0]
   num train = self.X train.shape[0]
   dists = np.zeros((num test, num train))
   # print(num test)
   # print(X[0].shape)
   # print(num train)
   # print(self.X train[0].shape)
   for i in np.arange(num test):
     for j in np.arange(num train):
      # YOUR CODE HERE:
        Compute the distance between the ith test point and the jth
         training point using norm(), and store the result in dists[i, j].
       # ------ #
      dists[i,j] = norm(self.X train[j]-X[i])
      pass
       # ----- #
       # END YOUR CODE HERE
       # ------ #
   return dists
 def compute L2 distances vectorized(self, X):
   Compute the distance between each test point in X and each training point
   in self.X train WITHOUT using any for loops.
```

```
Inputs:
 - X: A numpy array of shape (num test, D) containing test data.
 Returns:
 - dists: A numpy array of shape (num test, num train) where dists[i, j]
   is the Euclidean distance between the ith test point and the jth training
 num test = X.shape[0]
 num_train = self.X train.shape[0]
 dists = np.zeros((num_test, num_train))
 # ============= #
 # YOUR CODE HERE:
   Compute the L2 distance between the ith test point and the jth
    training point and store the result in dists[i, j]. You may
   NOT use a for loop (or list comprehension). You may only use
   numpy operations.
   HINT: use broadcasting. If you have a shape (N,1) array and
 #
   a shape (M,) array, adding them together produces a shape (N, M)
   array.
 # ----- #
 # print(X.shape)
 # print(self.X train.shape)
 # diff = self.X train - X.T
 norm train = self.X train**2
 norm train = np.sum(norm train, axis=1)
 term2 = ( X @ self.X train.T )
 norm test = X**2
 norm_test = np.sum(norm_test,axis=1)
 sum norms = norm test.reshape(len(norm test), 1) + norm train
 \#sum norms = (np.reshape(norm\ test,\ (len(norm\ test),1)) + norm\ train).T
 # print(norm_train.shape)
 # print(term2.shape)
 # print(norm test.shape)
 # print(sum norms.shape)
 # print(term2.shape)
 dists = np.sqrt(sum norms - 2*term2)
 pass
 # END YOUR CODE HERE
 return dists
def predict labels(self, dists, k=1):
 Given a matrix of distances between test points and training points,
 predict a label for each test point.
 Inputs:
```

- dists: A numpy array of shape (num\_test, num\_train) where dists[i, j] gives the distance betwen the ith test point and the jth training point. Returns: - y: A numpy array of shape (num test,) containing predicted labels for the test data, where y[i] is the predicted label for the test point X[i]. num test = dists.shape[0] y pred = np.zeros(num test) for i in np.arange(num test): # A list of length k storing the labels of the k nearest neighbors to # the ith test point. closest y = []# ============ # # YOUR CODE HERE: Use the distances to calculate and then store the labels of the k-nearest neighbors to the ith test point. The function numpy.argsort may be useful. # After doing this, find the most common label of the k-nearest neighbors. Store the predicted label of the ith training example as y pred[i]. Break ties by choosing the smaller label. # ----- # sortedDists = np.argsort(dists[i]) closest y = self.y train[sortedDists[:k]] y pred[i] = np.argmax(np.bincount(closest y))

# ----- #

# ------ #

return y pred

# END YOUR CODE HERE

pass

# This is the softmax workbook for ECE C147/C247 Assignment #2

Please follow the notebook linearly to implement a softmax classifier.

Please print out the workbook entirely when completed.

The goal of this workbook is to give you experience with training a softmax classifier.

```
In [80]: import random
    import numpy as np
    from utils.data_utils import load_CIFAR10
    import matplotlib.pyplot as plt

%matplotlib inline
%load_ext autoreload
%autoreload 2
```

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

```
In [81]: def get_CIFAR10_data(num_training=49000, num_validation=1000, num_test=1000, num_de
             Load the CIFAR-10 dataset from disk and perform preprocessing to prepare
             it for the linear classifier. These are the same steps as we used for the
             SVM, but condensed to a single function.
             # Load the raw CIFAR-10 data
             cifar10_dir = 'C:/Users/jessi/OneDrive/Desktop/classes/c147/HW2_code/cifar-10-b
             X_train, y_train, X_test, y_test = load_CIFAR10(cifar10_dir)
             # subsample the data
             mask = list(range(num_training, num_training + num_validation))
             X_val = X_train[mask]
             y_val = y_train[mask]
             mask = list(range(num training))
             X_train = X_train[mask]
             y_train = y_train[mask]
             mask = list(range(num_test))
             X_{\text{test}} = X_{\text{test}}[mask]
             y_test = y_test[mask]
             mask = np.random.choice(num_training, num_dev, replace=False)
             X_dev = X_train[mask]
             y_dev = y_train[mask]
             # Preprocessing: reshape the image data into rows
             X_train = np.reshape(X_train, (X_train.shape[0], -1))
             X_val = np.reshape(X_val, (X_val.shape[0], -1))
             X_test = np.reshape(X_test, (X_test.shape[0], -1))
             X_dev = np.reshape(X_dev, (X_dev.shape[0], -1))
             # Normalize the data: subtract the mean image
```

```
mean_image = np.mean(X_train, axis = 0)
     X_train -= mean_image
     X val -= mean image
     X_test -= mean_image
     X_dev -= mean_image
     # add bias dimension and transform into columns
     X_train = np.hstack([X_train, np.ones((X_train.shape[0], 1))])
     X_val = np.hstack([X_val, np.ones((X_val.shape[0], 1))])
     X_test = np.hstack([X_test, np.ones((X_test.shape[0], 1))])
     X_dev = np.hstack([X_dev, np.ones((X_dev.shape[0], 1))])
     return X_train, y_train, X_val, y_val, X_test, y_test, X_dev, y_dev
 # Invoke the above function to get our data.
 X_train, y_train, X_val, y_val, X_test, y_test, X_dev, y_dev = get_CIFAR10_data()
 print('Train data shape: ', X_train.shape)
 print('Train labels shape: ', y_train.shape)
 print('Validation data shape: ', X_val.shape)
 print('Validation labels shape: ', y_val.shape)
 print('Test data shape: ', X_test.shape)
 print('Test labels shape: ', y_test.shape)
 print('dev data shape: ', X_dev.shape)
 print('dev labels shape: ', y_dev.shape)
Train data shape: (49000, 3073)
Train labels shape: (49000,)
Validation data shape: (1000, 3073)
Validation labels shape: (1000,)
Test data shape: (1000, 3073)
Test labels shape: (1000,)
dev data shape: (500, 3073)
dev labels shape: (500,)
```

# Training a softmax classifier.

The following cells will take you through building a softmax classifier. You will implement its loss function, then subsequently train it with gradient descent. Finally, you will choose the learning rate of gradient descent to optimize its classification performance.

```
In [82]: from nndl import Softmax
In [83]: # Declare an instance of the Softmax class.
# Weights are initialized to a random value.
# Note, to keep people's first solutions consistent, we are going to use a random s

np.random.seed(1)

num_classes = len(np.unique(y_train))
num_features = X_train.shape[1]

softmax = Softmax(dims=[num_classes, num_features])
```

#### **Softmax loss**

2.3277607028048966

#### **Question:**

You'll notice the loss returned by the softmax is about 2.3 (if implemented correctly). Why does this make sense?

#### **Answer:**

 $ln(10) \sim = 2.3$ . Since there are 10 classes, there is a 1/10 chance the classifier guesses the class correctly since the weight vector was randomized.

#### Softmax gradient

```
In [86]: ## Calculate the gradient of the softmax loss in the Softmax class.
    # For convenience, we'll write one function that computes the loss
    # and gradient together, softmax.loss_and_grad(X, y)
    # You may copy and paste your loss code from softmax.loss() here, and then
    # use the appropriate intermediate values to calculate the gradient.

loss, grad = softmax.loss_and_grad(X_dev,y_dev)

# Compare your gradient to a gradient check we wrote.
    # You should see relative gradient errors on the order of 1e-07 or less if you impl softmax.grad_check_sparse(X_dev, y_dev, grad)

numerical: -0.955958 analytic: -0.955958, relative error: 1.324181e-08
numerical: 0.532381 analytic: 0.532381, relative error: 3.267644e-08
numerical: 0.054286 analytic: 0.054286, relative error: 1.088763e-07
numerical: 0.886285 analytic: 0.886285, relative error: 2.604743e-08
numerical: -0.294172 analytic: -0.294172, relative error: 1.099968e-07
numerical: 2.912753 analytic: 2.912753, relative error: 7.731013e-09
```

numerical: -0.399253 analytic: -0.399253, relative error: 1.410311e-07 numerical: -1.660508 analytic: -1.660508, relative error: 5.876757e-09 numerical: 0.156205 analytic: 0.156205, relative error: 2.717069e-10 numerical: -1.590637 analytic: -1.590637, relative error: 2.307966e-08

#### A vectorized version of Softmax

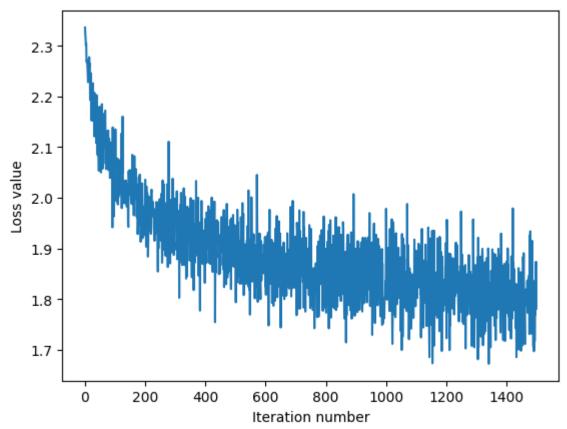
To speed things up, we will vectorize the loss and gradient calculations. This will be helpful for stochastic gradient descent.

```
import time
In [87]:
In [88]: ## Implement softmax.fast_loss_and_grad which calculates the loss and gradient
              WITHOUT using any for loops.
         # Standard Loss and gradient
         tic = time.time()
         loss, grad = softmax.loss_and_grad(X_dev, y_dev)
         toc = time.time()
         print('Normal loss / grad_norm: {} / {} computed in {}s'.format(loss, np.linalg.nor
         tic = time.time()
         loss_vectorized, grad_vectorized = softmax.fast_loss_and_grad(X_dev, y_dev)
         toc = time.time()
         print('Vectorized loss / grad: {} / {} computed in {}s'.format(loss_vectorized, np.
         # The losses should match but your vectorized implementation should be much faster.
         print('difference in loss / grad: {} /{} '.format(loss - loss_vectorized, np.linalg
         # You should notice a speedup with the same output.
       Normal loss / grad_norm: 2.320174544164465 / 336.4140707952609 computed in 0.0187957
       2868347168s
       Vectorized loss / grad: 2.3201745441644626 / 336.4140707952609 computed in 0.0030055
       04608154297s
       difference in loss / grad: 2.220446049250313e-15 /2.9133394406250957e-13
```

### Stochastic gradient descent

We now implement stochastic gradient descent. This uses the same principles of gradient descent we discussed in class, however, it calculates the gradient by only using examples from a subset of the training set (so each gradient calculation is faster).

```
iteration 0 / 1500: loss 2.3365926606637544
iteration 100 / 1500: loss 2.0557222613850827
iteration 200 / 1500: loss 2.0357745120662813
iteration 300 / 1500: loss 1.9813348165609888
iteration 400 / 1500: loss 1.9583142443981614
iteration 500 / 1500: loss 1.8622653073541355
iteration 600 / 1500: loss 1.8532611454359382
iteration 700 / 1500: loss 1.8353062223725827
iteration 800 / 1500: loss 1.8293892468827642
iteration 900 / 1500: loss 1.8992158530357484
iteration 1000 / 1500: loss 1.97835035402523
iteration 1100 / 1500: loss 1.8470797913532633
iteration 1200 / 1500: loss 1.8411450268664082
iteration 1300 / 1500: loss 1.7910402495792102
iteration 1400 / 1500: loss 1.8705803029382257
That took 3.442908763885498s
```



#### Evaluate the performance of the trained softmax classifier on the validation data.

```
In [90]: ## Implement softmax.predict() and use it to compute the training and testing error

y_train_pred = softmax.predict(X_train)
print('training accuracy: {}'.format(np.mean(np.equal(y_train,y_train_pred), )))
y_val_pred = softmax.predict(X_val)
print('validation accuracy: {}'.format(np.mean(np.equal(y_val, y_val_pred)), ))
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training accuracy: 0.3811428571428571

validation accuracy: 0.398

# Optimize the softmax classifier

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In [91]: np.finfo(float).eps
Out[91]: 2.220446049250313e-16
In [95]:
       # YOUR CODE HERE:
        # Train the Softmax classifier with different learning rates and
            evaluate on the validation data.
          Report:
            - The best learning rate of the ones you tested.
            - The best validation accuracy corresponding to the best validation error.
          Select the SVM that achieved the best validation error and report
            its error rate on the test set.
        val_accs = []
        lr = [0.001, 1e-4, 1e-5, 1e-6, 1e-7, 1e-7, 1e-8, 1e-9]
        for r in range(len(lr)):
           softmax = Softmax(dims=[num_classes, num_features])
           softmax.train(X_train, y_train, learning_rate=lr[r],num_iters=1500)
           y_val_pred = softmax.predict(X_val)
           error = np.sum(y_val_pred != y_val) / len(y_val)
           val accs.append(error)
           print('validation error: {}'.format(error))
        lowest_error = np.min(val_accs)
        best_lr = lr[np.argmin(val_accs)]
        print("best error was: ", str(np.min(val_accs)), "with learning rate =", str(lr[np.
        # END YOUR CODE HERE
        validation error: 0.913
      validation error: 0.722
      validation error: 0.669
      validation error: 0.581
      validation error: 0.606
      validation error: 0.606
      validation error: 0.7
      validation error: 0.862
      best error was: 0.581 with learning rate = 1e-06
In [93]: #Testing on test data#
        softmax.train(X_train, y_train, learning_rate=best_lr)
        y_test_pred = softmax.predict(X_test)
        error = np.sum(y test pred != y test) / len(y test pred)
        print("Test set error:", str(error))
        print("Test set accuracy:", str(1-error))
      Test set error: 0.645
      Test set accuracy: 0.355
```

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class Softmax(object):
 def init (self, dims=[10, 3073]):
   self.init weights(dims=dims)
 def init weights(self, dims):
   Initializes the weight matrix of the Softmax classifier.
   Note that it has shape (C, D) where C is the number of
   classes and D is the feature size.
   self.W = np.random.normal(size=dims) * 0.0001
 def loss(self, X, y):
   Calculates the softmax loss.
   Inputs have dimension D, there are C classes, and we operate on minibatches
   of N examples.
   Inputs:
   - X: A numpy array of shape (N, D) containing a minibatch of data.
   - y: A numpy array of shape (N,) containing training labels; y[i] = c means
     that X[i] has label c, where 0 \le c < C.
   Returns a tuple of:
   - loss as single float
   # Initialize the loss to zero.
   loss = 0.0
   # ------ #
   # YOUR CODE HERE:
     Calculate the normalized softmax loss. Store it as the variable loss.
      (That is, calculate the sum of the losses of all the training
   # set margins, and then normalize the loss by the number of
     training examples.)
   for i in range(len(X)): #for each example
     term1 = 0
     for c in np.unique(y): #for each class
      term1 += np.exp(self.W[c] @ X[i])
     loss += np.log(term1) - self.W[y[i]] @ X[i]
   loss /= len(y)
   pass
   # END YOUR CODE HERE
   # ----- #
   return loss
 def loss and grad(self, X, y):
   Same as self.loss(X, y), except that it also returns the gradient.
   Output: grad -- a matrix of the same dimensions as W containing
    the gradient of the loss with respect to W.
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import numpy as np

```
# Initialize the loss and gradient to zero.
   loss = 0.0
   grad = np.zeros like(self.W)
   # ----- #
   # YOUR CODE HERE:
     Calculate the softmax loss and the gradient. Store the gradient
     as the variable grad.
   N = len(X)
   scores = np.exp(X.dot(self.W.T))
   probs = scores/np.sum(scores, axis=1, keepdims=True)
   for i in range(N):
    loss += -np.log(probs[i, y[i]])
    for c in range(self.W.shape[0]):
      grad[c] += X[i] * probs[i, c]
      if c == y[i]:
                                 #if the model guesses correctly, make gradient small
        grad[c] -= X[i]
   loss /= N
   grad /= N
   # END YOUR CODE HERE
   return loss, grad
 def grad check sparse(self, X, y, your grad, num checks=10, h=1e-5):
   sample a few random elements and only return numerical
   in these dimensions.
   for i in np.arange(num_checks):
    ix = tuple([np.random.randint(m) for m in self.W.shape])
    oldval = self.W[ix]
    self.W[ix] = oldval + h # increment by h
    fxph = self.loss(X, y)
    self.W[ix] = oldval - h # decrement by h
    fxmh = self.loss(X, y) # evaluate f(x - h)
    self.W[ix] = oldval # reset
    grad numerical = (fxph - fxmh) / (2 * h)
    grad analytic = your grad[ix]
    rel_error = abs(grad_numerical - grad_analytic) / (abs(grad_numerical) +
abs(grad analytic))
    print('numerical: %f analytic: %f, relative error: %e' % (grad numerical, grad analytic,
rel_error))
 def fast loss and grad(self, X, y):
   A vectorized implementation of loss and grad. It shares the same
   inputs and ouptuts as loss_and_grad.
   11 11 11
   grad = np.zeros(self.W.shape) # initialize the gradient as zero
   # ----- #
   # YOUR CODE HERE:
     Calculate the softmax loss and gradient WITHOUT any for loops.
   # ----- #
   N = len(X)
   scores = np.exp(X.dot(self.W.T))
   probs = scores/np.sum(scores, axis=1, keepdims=True)
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loss = -np.sum(np.log(probs[np.arange(N), y])) / N
 probs[np.arange(N), y] -= 1
 probs /= N
 grad = probs.T.dot(X)
 pass
 # =================== #
 # END YOUR CODE HERE
 # ------ #
 return loss, grad
def train(self, X, y, learning rate=1e-3, num iters=100,
        batch size=200, verbose=False):
 Train this linear classifier using stochastic gradient descent.
 Inputs:
 - X: A numpy array of shape (N, D) containing training data; there are N
   training samples each of dimension D.
 - y: A numpy array of shape (N,) containing training labels; y[i] = c
  means that X[i] has label 0 <= c < C for C classes.
 - learning rate: (float) learning rate for optimization.
 - num iters: (integer) number of steps to take when optimizing
 - batch size: (integer) number of training examples to use at each step.
 - verbose: (boolean) If true, print progress during optimization.
 Outputs:
 A list containing the value of the loss function at each training iteration.
 num train, dim = X.shape
 num classes = np.max(y) + 1 # assume y takes values 0...K-1 where K is number of classes
 self.init weights (dims=[np.max(y) + 1, X.shape[1]]) # initializes the weights of self.W
 # Run stochastic gradient descent to optimize W
 loss_history = []
 for it in np.arange(num iters):
   X  batch = None
   y batch = None
   # ============= #
   # YOUR CODE HERE:
     Sample batch size elements from the training data for use in
      gradient descent. After sampling,
       - X batch should have shape: (batch size, dim)
   #
       - y_batch should have shape: (batch_size,)
     The indices should be randomly generated to reduce correlations
      in the dataset. Use np.random.choice. It's okay to sample with
     replacement.
   # ----- #
   batch index = np.random.choice(num train, batch size)
   X_batch = X[batch index]
   y_batch = y[batch_index]
   pass
   # END YOUR CODE HERE
   # ----- #
   # evaluate loss and gradient
   loss, grad = self.fast loss and grad(X batch, y batch)
   loss history.append(loss)
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# ============= #
  # YOUR CODE HERE:
  # Update the parameters, self.W, with a gradient step
  # ============= #
  self.W -= grad * learning rate
  pass
  # ------ #
  # END YOUR CODE HERE
  # ----- #
  if verbose and it % 100 == 0:
   print('iteration {} / {}: loss {}'.format(it, num iters, loss))
 return loss history
def predict(self, X):
 Inputs:
 - X: N x D array of training data. Each row is a D-dimensional point.
 Returns:
 - y pred: Predicted labels for the data in X. y pred is a 1-dimensional
 array of length N, and each element is an integer giving the predicted
 mmm
 y_pred = np.zeros(X.shape[1])
 # ----- #
 # YOUR CODE HERE:
  Predict the labels given the training data.
 # ----- #
 scores = X.dot(self.W.T)
 y pred = np.argmax(scores, axis = 1)
 # END YOUR CODE HERE
 # ----- #
 return y pred
```