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.

We thank Serena Yeung & Justin Johnson for permission to use code written for the CS 231n class (cs231n.stanford.edu). These are the functions in the cs231n folders and code in the jupyer notebook to preprocess and show the images. The classifiers used are based off of code prepared for CS 231n as well.

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

```
In [108]: import numpy as np # for doing most of our calculations
import matplotlib.pyplot as plt# for plotting
from cs23ln.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 [109]: # Set the path to the CIFAR-10 data
    cifar10_dir = './cifar-10-batches-py' # You need to update this line
    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 [110]: # 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 [111]: # 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)
```

K-nearest neighbors

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

```
In [6]: # Import the KNN class
    from nndl import KNN

In [7]: # 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) Describe what is going on in the function knn.train().
- (2) What are the pros and cons of this training step?

Answers

- (1) The knn.train() function caches the entire data set provided by the x and y parameters.
- (2) The pro of this training step is that it is fast and simple, with constant time complexity. The con is that it is memory intensive because all the training data must be stored, scaling with the training data with space complexity of O(N).

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 [12]: # Implement the function compute_distances() in the KNN class.
# Do not worry about the input 'norm' for now; use the default definition of the norm
# 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: 53.55126595497131
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 [73]: # 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 loops.
# 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): {}'.format(np.1)

Time to run code: 0.3632807731628418
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

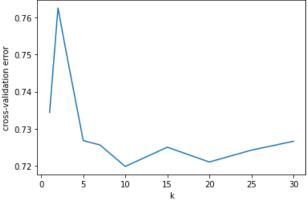
First, we will create the training and validation folds for use in k-fold cross validation.

```
In [155]: # 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]
       # ----- #
       X train folds = np.split(X train, num folds)
       y train folds = np.split(y train, num folds)
       # END YOUR CODE HERE
        ______#
```

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 [163]: 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.
         fold_size = len(X_train) // num_folds
         training fold size = int((num folds-1) * fold size)
         x size = np.array(X train folds).shape[2]
         errors = []
         for k val in ks:
             error_total = 0
             for fold_idx in range(num_folds):
                # train on training fold
                X_training_fold = np.array(X_train_folds[:fold_idx] + X_train_folds[fold_idx+1:]).reshape
                y_training_fold = np.array(y_train_folds[:fold_idx] + y_train_folds[fold_idx+1:]).flatten
                knn.train(X=X_training_fold, y=y_training_fold)
                # compute L2 distance and predict y on test fold
                dists = knn.compute L2 distances vectorized(X=X train folds[fold idx])
                y pred = knn.predict labels(dists, k=k val)
                # add error to error total
                error_total += np.mean(y_train_folds[fold_idx] != y_pred)
             error_avg = error_total / num_folds
            print('k={}: error={}'.format(k_val, round(error_avg, 4)))
            errors.append(error avg)
         plt.plot(ks, errors)
         plt.xlabel('k')
         plt.ylabel('cross-validation error')
         # END YOUR CODE HERE
         # ----- #
         print('Computation time: %.2f'%(time.time()-time start))
         k=1: error=0.7344
         k=2: error=0.7626
         k=3: error=0.7504
         k=5: error=0.7268
         k=7: error=0.7256
         k=10: error=0.7198
         k=15: error=0.725
         k=20: error=0.721
         k=25: error=0.7242
         k=30: error=0.7266
         Computation time: 51.17
           0.76
           0.75
```



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:

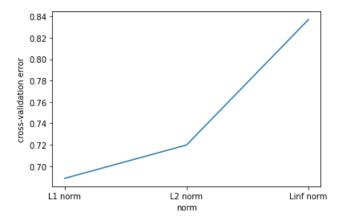
- (1) The best $\, \mathbf{k} \,$ value is 10 since it had the lowest error.
- (2) The cross-validation error for k=10 is 0.7198.

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 [161]: 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]
         norm_labels = ['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.
         k best = 10
         errors = []
         for norm_idx, norm in enumerate(norms):
            error_total = 0
            for fold idx in range(num folds):
                # train on training fold
                X training fold = np.array(X train folds[:fold idx] + X train folds[fold idx+1:]).reshape
                y_training_fold = np.array(y_train_folds[:fold_idx] + y_train_folds[fold_idx+1:]).flatten
                knn.train(X=X_training_fold, y=y_training_fold)
                # compute distance and predict y on test fold
                dists = knn.compute_distances(X=X_train_folds[fold_idx], norm=norm)
                y pred = knn.predict labels(dists, k=k best)
                # add error to error total
                error_total += np.mean(y train folds[fold_idx] != y pred)
            error_avg = error_total / num_folds
            print('norm={}: error={}'.format(norm_labels[norm_idx], round(error_avg, 4)))
            errors.append(error_avg)
         plt.plot(norm labels, errors)
         plt.xlabel('norm')
         plt.ylabel('cross-validation error')
         # END YOUR CODE HERE
         # ----- #
         print('Computation time: %.2f'%(time.time()-time start))
         norm=L1 norm: error=0.6886
```

norm=L1 norm: error=0.6886 norm=L2 norm: error=0.7198 norm=Linf norm: error=0.837 Computation time: 1129.76



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 achieved the best cross-validation error.
- (2) Cross-validation error for L1 norm and k=10 is 0.6886.

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: 0.716

Question:

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

Answer:

Error improved by 0.01 (around 1.3%), since error with k=10 and L1-norm is 0.716, and error with k=1 and L2-norm was 0.726.

```
In [ ]:
```

```
import numpy as np
import pdb
This code was based off of code from cs231n at Stanford University, and modified for ECE C147/
C247 at UCLA.
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.
    - 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))
    for i in np.arange(num_test):
      for j in np.arange(num_train):
               # ========
                                        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])
               # 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.
    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.
```

```
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)
 # Let A = X, B = self.X_train
 \# (A-B)**2 = A**2 - 2AB + B**2
 # Have: A = (500, 3072), B = (5000, 3072)
 # Want: A**2 = (500, 1), B**2 = (5000,), AB = (500, 5000)
 # (500,) array with each row being the sum of a testing data point's coordinates
  a_squared = np.sum(np.square(X), axis=1)
  # Reshape to (500,1) array
 a_squared = a_squared.reshape((a_squared.shape[0], 1))
 # (5000,) array with each row being the sum of a test data point's coordinates squared b_squared = np.sum(np.square(self.X_train), axis=1)
  \#(500,1) array + (5000,) array = (500, 5000) array that holds A**2 + B**2 for each
 # pair of testing points and training points
 dists = a_squared + b_squared
 \# (500, 3072).T * (5000, 3072) = (500, 3072) * (3072, 5000) = (500, 5000)
 a_times_b = np.dot(X, self.X_train.T)
 dists -= 2*a_times_b
 dists = np.sqrt(dists)
     # 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.
  - 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.
  - 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
```

return y_pred

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

Please follow the notebook linearly to implement a linear support vector machine.

Please print out the workbook entirely when completed.

We thank Serena Yeung & Justin Johnson for permission to use code written for the CS 231n class (cs231n.stanford.edu). These are the functions in the cs231n folders and includes code to preprocess and show the images. The classifiers used are based off of code prepared for CS 231n as well.

The goal of this workbook is to give you experience with training an SVM classifier via gradient descent.

Importing libraries and data setup

```
In [79]: import numpy as np # for doing most of our calculations
    import matplotlib.pyplot as plt# for plotting
    from cs23ln.data_utils import load_CIFAR10 # function to load the CIFAR-10 dataset.
    import pdb

# 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 [80]: # Set the path to the CIFAR-10 data
    cifar10 dir = './cifar-10-batches-py' # You need to update this line
```

```
In [80]: # Set the path to the CIFAR-10 data
    cifar10_dir = './cifar-10-batches-py' # You need to update this line
    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 [81]: # 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 [82]: # Split the data into train, val, and test sets. In addition we will
         # create a small development set as a subset of the training data;
         # we can use this for development so our code runs faster.
         num training = 49000
         num validation = 1000
         num_test = 1000
         num dev = 500
         # Our validation set will be num_validation points from the original
         # training set.
         mask = range(num_training, num_training + num_validation)
         X_val = X_train[mask]
         y_val = y_train[mask]
         # Our training set will be the first num train points from the original
         # training set.
         mask = range(num_training)
         X_train = X_train[mask]
         y_train = y_train[mask]
         # We will also make a development set, which is a small subset of
         # the training set.
         mask = np.random.choice(num training, num dev, replace=False)
         X_dev = X_train[mask]
         y_dev = y_train[mask]
         # We use the first num test points of the original test set as our
         # test set.
         mask = range(num test)
         X test = X test[mask]
         y_test = y_test[mask]
         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, 32, 32, 3)
         Train labels shape: (49000,)
         Validation data shape: (1000, 32, 32, 3)
         Validation labels shape: (1000,)
         Test data shape: (1000, 32, 32, 3)
         Test labels shape: (1000,)
         Dev data shape: (500, 32, 32, 3)
         Dev labels shape: (500,)
In [83]: # Preprocessing: reshape the image data into rows
         X_train = np.reshape(X_train, (X_train.shape[0], -1))
         X \text{ val} = \text{np.reshape}(X \text{ val}, (X \text{ val.shape}[0], -1))
         X_test = np.reshape(X_test, (X_test.shape[0], -1))
         X \text{ dev} = \text{np.reshape}(X \text{ dev}, (X \text{ dev.shape}[0], -1))
         # As a sanity check, print out the shapes of the data
         print('Training data shape: ', X_train.shape)
         print('Validation data shape: ', X_val.shape)
         print('Test data shape: ', X_test.shape)
         print('dev data shape: ', X dev.shape)
         Training data shape: (49000, 3072)
         Validation data shape: (1000, 3072)
         Test data shape: (1000, 3072)
         dev data shape: (500, 3072)
```

```
In [84]: # Preprocessing: subtract the mean image
    # first: compute the image mean based on the training data
    mean_image = np.mean(X_train, axis=0)
    print(mean_image[:10]) # print a few of the elements
    plt.figure(figsize=(4,4))
    plt.imshow(mean_image.reshape((32,32,3)).astype('uint8')) # visualize the mean image
    plt.show()
[130.64189796 135.98173469 132.47391837 130.05569388 135.34804082
```

131.75402041 130.96055102 136.14328571 132.47636735 131.48467347]

0 -5 -10 -15 -20 -25 --

10

```
In [85]: # second: subtract the mean image from train and test data
X_train -= mean_image
X_val -= mean_image
X_test -= mean_image
X_dev -= mean_image
```

```
In [86]: # third: append the bias dimension of ones (i.e. bias trick) so that our SVM
# only has to worry about optimizing a single weight matrix W.

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))])

print(X_train.shape, X_val.shape, X_test.shape, X_dev.shape)
```

(49000, 3073) (1000, 3073) (1000, 3073) (500, 3073)

30

Question:

30

(1) For the SVM, we perform mean-subtraction on the data. However, for the KNN notebook, we did not. Why?

Answer:

(1) We perform mean-subtraction for SVM because we would like to center the data (the pixel values) around zero to allow features to more equally influence where the decision boundary is drawn to separate the data, i.e. more fairly contribute to the values of the weights during training. On the other hand, subtracting the mean in KNN does not change distances between points, therefore which k points are nearest to the tested point stays the same, so the predictions are the same.

Training an SVM

The following cells will take you through building an SVM. 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 [87]: from nndl.svm import SVM

In [88]: # Declare an instance of the SVM class.
# Weights are initialized to a random value.
# Note, to keep people's initial solutions consistent, we are going to use a random seed.

np.random.seed(1)

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

svm = SVM(dims=[num_classes, num_features])
```

SVM loss

```
In [89]: ## Implement the loss function for in the SVM class(nndl/svm.py), svm.loss()
    loss = svm.loss(X_train, y_train)
    print('The training set loss is {}.'.format(loss))

# If you implemented the loss correctly, it should be 15569.98
```

The training set loss is 15569.97791541019.

SVM gradient

```
In [90]: ## Calculate the gradient of the SVM class.
         # For convenience, we'll write one function that computes the loss
         # and gradient together. Please modify svm.loss and grad(X, y).
         # You may copy and paste your loss code from svm.loss() here, and then
           use the appropriate intermediate values to calculate the gradient.
         loss, grad = svm.loss and grad(X dev,y dev)
         # Compare your gradient to a numerical gradient check.
         \# You should see relative gradient errors on the order of 1e-07 or less if you implemented the gr
         svm.grad_check_sparse(X_dev, y_dev, grad)
         numerical: -0.888162 analytic: -0.888162, relative error: 3.716477e-07
         numerical: -2.394414 analytic: -2.394414, relative error: 9.713307e-09
         numerical: -3.320923 analytic: -3.320922, relative error: 8.933639e-08
         numerical: 16.351982 analytic: 16.351981, relative error: 1.437057e-08
         numerical: 0.405991 analytic: 0.405991, relative error: 5.224500e-07
         numerical: -2.866195 analytic: -2.866195, relative error: 5.677315e-08
         numerical: 3.430926 analytic: 3.430926, relative error: 2.426969e-08
         numerical: -16.097096 analytic: -16.097096, relative error: 8.285457e-10
         numerical: 2.020525 analytic: 2.020525, relative error: 2.228789e-08
         numerical: -11.892703 analytic: -11.892702, relative error: 1.038786e-08
```

A vectorized version of SVM

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

```
In [91]: import time
```

```
In [97]: ## Implement sym.fast_loss_and_grad which calculates the loss and gradient
    # Standard loss and gradient
    tic = time.time()
    loss, grad = sym.loss_and_grad(X_dev, y_dev)
    toc = time.time()
    print('Normal loss / grad_norm: {} / {} computed in {}s'.format(loss, np.linalg.norm(grad, 'fro'))

    tic = time.time()
    loss_vectorized, grad_vectorized = sym.fast_loss_and_grad(X_dev, y_dev)
    toc = time.time()
    print('Vectorized loss / grad: {} / {} computed in {}s'.format(loss_vectorized, np.linalg.norm(grad)
    # The losses should match but your vectorized implementation should be much faster.
    print('difference in loss / grad: {} / {}'.format(loss_loss_vectorized, np.linalg.norm(grad) - grad)
    # You should notice a speedup with the same output, i.e., differences on the order of le-12
```

```
Normal loss / grad_norm: 15380.605267476883 / 1985.4908887207862 computed in 0.0620729923248291 s

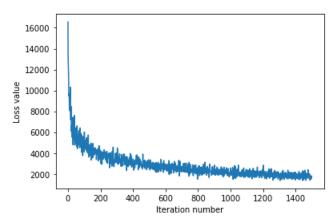
Vectorized loss / grad: 15380.60526747687 / 1985.4908887207862 computed in 0.007290124893188476 6s

difference in loss / grad: 1.2732925824820995e-11 / 4.032768305804647e-12
```

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 16557.38000190916
iteration 100 / 1500: loss 4701.089451272714
iteration 200 / 1500: loss 4017.333137942788
iteration 300 / 1500: loss 3681.922647195363
iteration 400 / 1500: loss 2732.6164373988995
iteration 500 / 1500: loss 2786.637842464506
iteration 600 / 1500: loss 2837.0357842782673
iteration 700 / 1500: loss 2206.2348687399317
iteration 800 / 1500: loss 2269.0388241169803
iteration 900 / 1500: loss 2543.23781538592
iteration 1000 / 1500: loss 2566.692135726827
iteration 1000 / 1500: loss 2182.068905905164
iteration 1200 / 1500: loss 1861.1182244250456
iteration 1300 / 1500: loss 1982.9013858528251
iteration 1400 / 1500: loss 1927.520415858212
That took 8.007238864898682s
```



Evaluate the performance of the trained SVM on the validation data.

```
In [99]: ## Implement sym.predict() and use it to compute the training and testing error.

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

training accuracy: 0.28530612244897957
validation accuracy: 0.3
```

Optimize the SVM

Note, to make things faster and simpler, we won't do k-fold cross-validation, but will only optimize the hyperparameters on the validation dataset (X_val, y_val).

```
In [106]: # ========= #
         # YOUR CODE HERE:
           Train the SVM with different learning rates and evaluate on the
             validation data.
             - 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.
            Note: You do not need to modify SVM class for this section
         learning_rates = [1e-4, 0.001, 0.01, 0.05, 0.1, 0.25, 0.5, 1]
         val accuracies = []
         for i, rate in enumerate(learning_rates):
            svm = SVM()
            svm.train(X train, y train, learning rate=rate, num iters=1500, verbose=False)
            predictions = svm.predict(X val)
            accuracy = np.mean(np.equal(predictions, y_val))
            val accuracies.append(accuracy)
            print('rate={}: accuracy={}, error={}'.format(rate, accuracy, 1-accuracy))
         best_rate = learning_rates[np.argmax(val_accuracies)]
         best_accuracy = np.max(val_accuracies)
         svm.train(X train, y train, learning rate=best rate, num iters=1500, verbose=False)
         test_error = 1 - np.mean(np.equal(svm.predict(X_test), y_test))
         print()
        print('Best learning rate:', best_rate)
        print('Best validation accuracy:', best accuracy)
        print('Test error with best learning rate:', test_error)
         # ------ #
         # END YOUR CODE HERE
         rate=0.0001: accuracy=0.279, error=0.721
         rate=0.001: accuracy=0.306, error=0.694
         rate=0.01: accuracy=0.31, error=0.69
         rate=0.05: accuracy=0.329, error=0.671
         rate=0.1: accuracy=0.322, error=0.6779999999999999
         rate=0.25: accuracy=0.305, error=0.695000000000001
         rate=1: accuracy=0.259, error=0.741
         Best learning rate: 0.05
         Best validation accuracy: 0.329
         Test error with best learning rate: 0.677
```

In []:

```
import numpy as np
import pdb
This code was based off of code from cs231n at Stanford University, and modified for ECE C147/
C247 at UCLA.
class SVM(object):
       _init__(self, dims=[10, 3073]):
    self.init_weights(dims=dims)
  def init_weights(self, dims):
       Initializes the weight matrix of the SVM. 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) # (10, 3037)
  def loss(self, X, y):
   Calculates the SVM 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
   # compute the loss and the gradient
   num_classes = self.W.shape[0] # 10
   num_train = X.shape[0] # 49000
    loss = 0.0
    for i in np.arange(num_train):
    # YOUR CODE HERE:
            Calculate the normalized SVM loss, and store it as '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 j in range(num_classes):
            if y[i] != j:
               a_j = np.matmul(self.W[j], X[i]) # score of incorrect class j
               a_y = np.matmul(self.W[y[i]], X[i]) # score of correct class y[i]
               loss += max(0, 1 + a_j - a_y)
    loss /= num_train
                         # 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.
   # compute the loss and the gradient
   num_classes = self.W.shape[0]
    num_train = X.shape[0]
    loss = 0.0
```

```
grad = np.zeros_like(self.W)
    for i in np.arange(num_train):
    # YOUR CODE HERE:
           Calculate the SVM loss and the gradient. Store the gradient in
       the variable grad.
    # =
        for j in range(num_classes):
            if y[i] != j:
                a_j = np.matmul(self.W[j], X[i]) # score of incorrect class j
                a_y = np.matmul(self.W[y[i]], X[i]) # score of correct class y[i]
                z_j = 1 + a_j - a_y
loss += max(0, z_j)
                if z_j > 0:
                    grad[j] += X[i]
                    grad[y[i]] -= X[i]
   # END YOUR CODE HERE
    loss /= num_train
   grad /= num_train
    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.
    loss = 0.0
   grad = np.zeros(self.W.shape) # initialize the gradient as zero
   # ----- #
    # YOUR CODE HERE:
       # Calculate the SVM loss WITHOUT any for loops.
   num_train = X.shape[0]
   # Scores w*x for each training image: (500,3073) * (3073,10) = (500,10) with row per training
image
   WX = np.matmul(X, self.W.T)
   # Extract w*x for each training point's correct classification
   A_y = WX[np.arange(num_train), y].reshape(num_train, 1)
```

```
# Calculate L_i = summation(max(0, 1 + WX- A_y)) across all i, including correct predictions
 L = np.maximum(np.zeros((num_train, 1)), 1 + WX - A_y)
 \# Since loss for correct class predictions are currently 1, we need to set them to 0
 L[np.arange(num_train), y] = 0
  loss += np.sum(L) / num_train
 # END YOUR CODE HERE
 # YOUR CODE HERE:
     # Calculate the SVM grad WITHOUT any for loops.
 # Store whether each value in L is > 0
 positivity = 1*(L>0)
 # Gradient w.r.t. w for correct class is negative summation of x_i for incorrectly classified
 positivity[np.arange(num_train), y] = -np.sum(positivity, axis=1)
 grad = np.matmul(positivity.T, X) / num_train
               ______ #
 # 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.
 - 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.</pre>
 - 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: (dim, batch_size)
# - 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.
                       ._____ #
```

```
idxs = np.random.choice(num_train, batch_size, replace=True)
  X_{batch} = X[idxs]
  y_batch = y[idxs]
  # END YOUR CODE HERE
  # evaluate loss and gradient
  loss, grad = self.fast_loss_and_grad(X_batch, y_batch)
  loss_history.append(loss)
                 # YOUR CODE HERE:
     Update the parameters, self.W, with a gradient step
  self.W -= learning_rate * grad
      # 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.

 - 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
 class.
 y_pred = np.zeros(X.shape[1])
              ______#
 # YOUR CODE HERE:
   Predict the labels given the training data with the parameter self.W.
 y_pred = np.argmax(np.matmul(X, self.W.T), axis=1)
 # END YOUR CODE HERE
         ______#
 return y_pred
```

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.

We thank Serena Yeung & Justin Johnson for permission to use code written for the CS 231n class (cs231n.stanford.edu). These are the functions in the cs231n folders and code in the jupyer notebook to preprocess and show the images. The classifiers used are based off of code prepared for CS 231n as well.

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

```
In [1]: import random
    import numpy as np
    from cs231n.data_utils import load_CIFAR10
    import matplotlib.pyplot as plt

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

```
In [3]: def get CIFAR10 data(num training=49000, num validation=1000, num test=1000, num dev=500):
            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 = './cifar-10-batches-py' # You need to update this line
            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_test = X_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 \text{ test} = \text{np.reshape}(X \text{ test.} shape[0], -1))
            X \text{ dev} = \text{np.reshape}(X \text{ dev}, (X \text{ 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 \text{ dev} = \text{np.hstack}([X \text{ dev}, \text{np.ones}((X \text{ 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 [4]: from nndl import Softmax
In [5]: # 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 seed.

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

```
In [7]: ## Implement the loss function of the softmax using a for loop over
    # the number of examples
loss = softmax.loss(X_train, y_train)
```

```
In [8]: print(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:

Since the weights have been initialized at random, there is around a 1/10 chance that the classifier predicts an input image's class correctly. Therefore, the softmax function will return approximatly 1/10, of which we take the log for the loss function and get -log(1/10) which is around 2.3.

Softmax gradient

```
In [13]: ## 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 implemented the gr
         softmax.grad_check_sparse(X_dev, y_dev, grad)
         numerical: -1.002885 analytic: -1.002885, relative error: 2.129165e-09
         numerical: 0.069499 analytic: 0.069499, relative error: 2.258907e-07
         numerical: 0.913719 analytic: 0.913719, relative error: 3.455543e-09
         numerical: 0.561850 analytic: 0.561850, relative error: 3.574384e-10
         numerical: -0.243901 analytic: -0.243901, relative error: 2.155266e-07
         numerical: 1.106812 analytic: 1.106812, relative error: 2.926994e-08
         numerical: 0.357237 analytic: 0.357237, relative error: 6.844256e-08
         numerical: -2.172427 analytic: -2.172427, relative error: 5.036179e-09
         numerical: -1.649068 analytic: -1.649068, relative error: 5.232190e-09
         numerical: -1.095710 analytic: -1.095710, relative error: 4.729251e-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.

```
In [14]:
        import time
In [28]: ## 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.norm(grad, 'fro')
         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.linalg.norm(gr
         # The losses should match but your vectorized implementation should be much faster.
         print('difference in loss / grad: {} /{} '.format(loss - loss_vectorized, np.linalg.norm(grad - g
         # You should notice a speedup with the same output.
```

Normal loss / grad_norm: 2.3422220643301808 / 282.0786751924475 computed in 0.10903811454772949 s
Vectorized loss / grad: 2.3422220643301817 / 282.0786751924475 computed in 0.0261843204498291s difference in loss / grad: -8.881784197001252e-16 /2.466349875728883e-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).

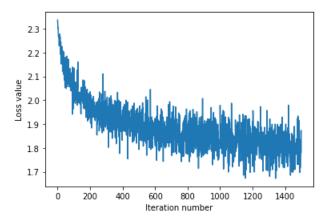
Question:

How should the softmax gradient descent training step differ from the sym training step, if at all?

Answer:

The gradient descent training step is the same between the two models because both essentially calculate the gradient and take a step in the direction of steepest descent. The results are different due to different loss functions, but the overall training steps should be the same.

```
iteration 0 / 1500: loss 2.336592660663754
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 1.8293892468827635
iteration 900 / 1500: loss 1.899215853035748
iteration 1000 / 1500: loss 1.9783503540252299
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 15,612247228622437s
```



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

```
In [30]: ## 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)), ))

training accuracy: 0.3811428571428571
```

Optimize the softmax classifier

validation accuracy: 0.398

You may copy and paste your optimization code from the SVM here.

```
In [31]: np.finfo(float).eps
Out[31]: 2.220446049250313e-16
In [36]: | # ============= #
        # 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.
        learning_rates = [1e-8, 1e-7, 1e-6, 1e-5, 1e-4, 1e-3, 1e-1]
        val_accuracies = []
        sm = Softmax()
        for i, rate in enumerate(learning_rates):
           sm.train(X_train, y_train, learning_rate=rate, num_iters=1500, verbose=False)
           predictions = sm.predict(X_val)
           accuracy = np.mean(np.equal(predictions, y_val))
           val accuracies.append(accuracy)
           print('rate={}: accuracy={}, error={}'.format(rate, accuracy, 1-accuracy))
        best_rate = learning_rates[np.argmax(val_accuracies)]
        best_accuracy = np.max(val_accuracies)
        sm.train(X_train, y_train, learning_rate=best_rate, num_iters=1500, verbose=False)
        test_error = 1 - np.mean(np.equal(sm.predict(X_test), y_test))
        print()
        print('Best learning rate:', best_rate)
        print('Best validation accuracy:', best_accuracy)
        print('Test error with best learning rate:', test_error)
        # ------ #
        # END YOUR CODE HERE
        # ------ #
        rate=1e-08: accuracy=0.308, error=0.692
        rate=1e-07: accuracy=0.386, error=0.614
        rate=1e-06: accuracy=0.412, error=0.5880000000000001
        rate=1e-05: accuracy=0.291, error=0.7090000000000001
        rate=0.0001: accuracy=0.283, error=0.717000000000001
        rate=0.001: accuracy=0.087, error=0.913
        rate=0.1: accuracy=0.087, error=0.913
        Best learning rate: 1e-06
        Best validation accuracy: 0.412
        Test error with best learning rate: 0.617
In [ ]:
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```
import numpy as np
class Softmax(object):
       _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:
   -\dot{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 <= 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.)
   num_train = X.shape[0]
    for i in range(num_train):
        A_i = np.matmul(self.W, X[i])
        log_summed_e_A = np.log(np.sum(np.exp(A_i)))
        A_y = np.matmul(self.W[y[i]], X[i])
        loss += log_summed_e_A - A_y
    loss /= num_train
   # 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.
   # 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.
```

```
num_train = X.shape[0]
   num_classes = self.W.shape[0]
   for i in range(num_train):
       A_i = np.matmul(self.W, X[i])
       e_A = np.exp(A_i)
       summed_e_A = np.sum(e_A)
       log_summed_e_A = np.log(summed_e_A)
       A_y = np.matmul(self.W[y[i]], X[i])
       loss += log_summed_e_A - A_y
       for j in range(num_classes):
           grad[j] += X[i] * e_A[j] / summed_e_A
       grad[y[i]] -= X[i]
   loss /= num_train
   grad /= num_train
   # 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.
   grad = np.zeros(self.W.shape) # initialize the gradient as zero
   # YOUR CODE HERE:
         Calculate the softmax loss and gradient WITHOUT any for loops.
   num_train = X.shape[0]
   # Loss
   A = np.matmul(X, self.W.T)
   e_A = np.exp(A)
   summed_e_A = np.sum(e_A, axis=1)
   log_summed_e_A = np.log(summed_e_A)
   pred = np.sum(np.multiply(self.W[y], X), axis=1)
```

```
loss_matrix = log_summed_e_A - pred
 loss = np.sum(loss_matrix) / num_train
 # Gradient
 S = e_A / summed_e_A.reshape(num_train, 1)
 S[np.arange(num_train), y] -= 1
 S = np.matmul(S.T, X)
 grad = S / num_train
 # 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 optimizingbatch_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: (dim, batch_size)
            - 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.
   idxs = np.random.choice(num_train, batch_size, replace=True)
   X  batch = X[idxs]
   y_batch = y[idxs]
   # END YOUR CODE HERE
   # evaluate loss and gradient
   loss, grad = self.fast_loss_and_grad(X_batch, y_batch)
   loss_history.append(loss)
   # YOUR CODE HERE:
       Update the parameters, self.W, with a gradient step
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```
self.W -= learning_rate * grad
   # 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.
 - 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
 class.
 y_pred = np.zeros(X.shape[1])
 # YOUR CODE HERE:
   Predict the labels given the training data.
 y_pred = np.argmax(np.matmul(X, self.W.T), axis=1)
 # END YOUR CODE HERE
```

return y_pred