Angel U. Ortega Computer Vision – Lab 4 Version 1.0 11/26/2016

Table of Contents

1.	IN	FRODUCTION	3
1	.1.	LAB OVERVIEW	3
1	.2.	REFERENCES	
2.	PR	OPOSED SOLUTION DESIGN AND IMPLEMENTATION	3
2	.1.	P1	3
2	.2.	P2	3
2	.3.	P3	4
2	.4.	P4	4
2	.5.	P5	4
2	.6.	P6	4
3.	EX	PERIMENTAL RESULTS AND CONCLUSIONS	4
3	.1.	P1	4
3	.2.	P2	
3	.3.	P3	8
3	.4.	P4	9
3	.5.	P5	10
3	.6.	P6	11
4.	AP	PENDIX	11
4	.1.	P1 CODE	11
4	.2.	P2 CODE	
4	.3.	P3 CODE	
4	.4.	P4 CODE	
4	.5.	P5 CODE	
4	.6.	P6 CODE	
EN	D OI	F DOCUMENT	22

Computer Vision – Lab 4	Angel U. Ortega	Date	Page
		11/26/2016 5:49 PM	ii

1. Introduction

1.1. Lab Overview

The lab is composed of five problems, P1 - P6. The problem statement is as follows:

- P1. Pixel clustering. Suppose you want to represent a color image as accurately as possible but have a limit on the number of colors you can display. If you have k colors available you could cluster the {r, g, b} values of the pixels in the image into k clusters, and then replace each pixel value by the mean of the cluster it is assigned to. Implement this and experiment with various values of k.
- P2. Image clustering. Cluster the MNIST dataset using pixel intensities as features. Observe if images contained in the same cluster represent the same digits. More formally, partition the dataset into k clusters (experiment with k = 10 and k = 20) and determine the fraction of the instances in each cluster that belong to the majority class in that cluster.
- P3. Image clustering. Cluster the CIFAR-10 dataset using color histograms as features. As in the previous question, observe if images contained in the same cluster represent objects of the same class.
- P4. Image clustering. Cluster the CIFAR-10 dataset histograms of gradients as features. As in the previous question, observe if images contained in the same cluster represent objects of the same class.
- P5. Image classification. Download the scikit machine learning toolbox and classify the MNIST dataset using the algorithm and feature set of your choice. Make sure you don't use the same examples for training and testing as this will give you overoptimistic results. Extra credit points will be given to the best results obtained in the class. For reference, state of the art error rates for this dataset are around 1%.
- P6. Image classification. Classify the CIFAR-10 dataset using the algorithm and feature set of your choice. Make sure you don't use the same examples for training and testing. Extra credit points will be given to the best results obtained in the class. For reference, state of the art error rates for this dataset are around 5%.

1.2. References

- [1] https://drive.google.com/open?id=0B_kWRxLZdmeJWERtV2ExMFhNdnc
- [2] http://johnloeber.com/docs/kmeans.html
- [3] http://stackoverflow.com/questions/31528800/how-to-implement-zca-whitening-python

2. Proposed Solution Design and Implementation

2.1. P1

For this problem, I investigated the best tool to conduct kmeans to retrieve the closest color average given k colors. I decided on using CV2. The program retrieves image as an array, and applies kmeans with the number of colors desired. The results are then used to modify a copy of the original image using only the k colors retrieved by kmeans. The new image is then saved in a specified directory.

2.2. P2

For this problem, I had to first extract the dataset MNIST. I utilized the code provided by Mario Landa during class for this purpose. I utilized the cv2.kmeans method like in P1 to cluster the images. As instructed by the problem statement I utilized k=10 and k=20 to compare results. The results can be seen in Section 3.2. To compare accuracy of my solution

Computer Vision – Lab 4	Angel U. Ortega	Date	Page	
		11/26/2016 5:49 PM	3	

to another approach, I modified John Loeber's kmeans implementation [2] to cluster MNIST dataset. The results from both programs are compared in Section 3.2.

2.3. P3

For this problem, I had to first extract the dataset CIFAR-10. I utilized the code provided in class for this purpose. To be able to use color histograms as features, I had to implement a method (getCH) to get the color histogram of an image. Then I read in all images, retrieve the color histogram from each, and store the resultant in a different object. Next, I use kmeans from the previous question but using the color histograms as clustering features. Finally, I implemented two methods to visualize my results: getAccuracy and plotData. The getAccuracy method returns the calculated accuracy for each cluster. The plotData method gets the resultant from the getAccuracy method, and plots it. Results for P3 can be seen in Section 3.3.

2.4. P4

For this problem, I used same program to extract dataset as in P3. For P3, I utilized the implementation I had for HOG from Lab 2. I started by reading in the images from the dataset and retrieving the histogram of gradients from each. I saved the result in a different object and used kmeans to cluster the array containing the histograms of gradients. I used the getAccuracy and plotData methods from the previous problem to visualize accuracy. Results for P4 can be seen in Section 3.4.

2.5. P5

For this problem, I utilized the KNN classifier with k = 10 and k = 20. I used pixel intensities as features for the classification. I first had to load the training and testing sets. I used the code provided in class for this purpose. I used a minkowski metric of 2 for the Euclidean standard. I fit the classifier with training set, then retrieve the score of both training set and the testing set. Results for P5can be seen in Section 3.5

2.6. P6

For this problem, I used a multi-layer perceptron classifier. I used pixel intensities and histograms of gradients as features. I used batch 1 as training and batch 3 as testing from the CIFAR-10 dataset. For this problem, I used both grayscale and color images. I utilized ZCA whitening normalization for pixel intensity features. For ZCA, I followed Timshel's implementation [3]. For HOG, I utilized the same implementation from P4. I applied whitening normalization and histogram of gradients to bot training and testing sets; I fit the resultants into two MLP classifiers. Results for P6 can be seen in Section 3.6.

3. Experimental Results and Conclusions

All images used for this lab can be accessed at: https://drive.google.com/open?id=0B_kWRxLZdmeJWERtV2ExMFhNdnc [1]

3.1. P1

Figures 1A and 1B show results for P1. As expected, the greater the number of colors retrieved from original image, the more accurate the pixel cluster image is to the original. For example, in Figure 1A, the original image is a picture of Mount Fuji. With k=1, the result is a single-color image. With k=2, we can start to see the separation between sky and ground, but Mount Fuji is completely lost in the background. At k=5, we can clearly see Mount Fuji, the sky, separation between horizon and foreground, and details in the flowers. There is also a color variation between the area that encased the flowers in k=2, and the same area in k=5. In k=10 we can see a lot more detail on the flowers as well as the buildings and a greater variation in the sky colors. Finally, in k=25, the image is very detailed, with at least 5 different shades of blue making up the sky. The same behavior can be seen in Figure 1B. The k=1 image is a single-color image. With k=2, the image is well defined, with a lot of contrast. As k increases, the details become more prominent in the image.

Computer Vision – Lab 4	Angel U. Ortega	Date	Page	
		11/26/2016 5:49 PM	4	



Figure 1A: From top to bottom and left to right - Original Image, and Pixel Clustering Images (k = 1, 2, 5, 10, 25)



Figure 1B: From top to bottom and left to right - Original Image, and Pixel Clustering Images (k = 1, 2, 5, 10, 25)

3.2. P2

Figures 2 and 3 show the results from my implementation of k-means and Loeber's modified implementation from pixel intensity image clustering of MNIST using k=10 and k=20. In comparison, my implementation receives almost identical results. Figure 3 shows the resulting images using k=10. Certain resulting images seem very accurate, like "0" and "8". In comparison, however, others seem very blurry and like a mixture of multiple digits "3", "4", and "5". Finally, some digits are repeated, like "9", which might be a result of the program clustering similar digits together like "7" and "9".

Computer Vision – Lab 4	Angel U. Ortega	Date	Page	
		11/26/2016 5:49 PM	6	

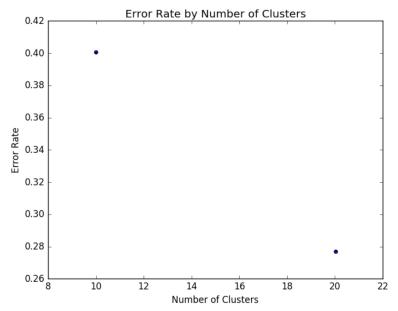


Figure 2: P2 implementation results

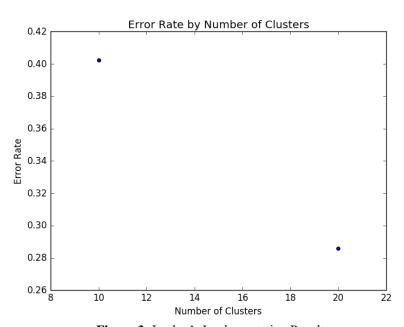


Figure 3: Loeber's Implementation Results



Figure 3: Image results for P2 using k = 10.

72012900534670193896

Figure 4: Image results for P2 using k = 20.

Computer Vision – Lab 4	Angel U. Ortega	Date	Page
		11/26/2016 5:49 PM	7

3.3. P3

Figures 5 and 6 shows the accuracy results for P3. From the results, I would conclude that color histograms do not make for a good feature to cluster CIFAR-10 images. Apart from the single almost perfect accuracy of one cluster, the other clusters receive very low results. Proportionally, using k = 10 gave better results than using k = 20.

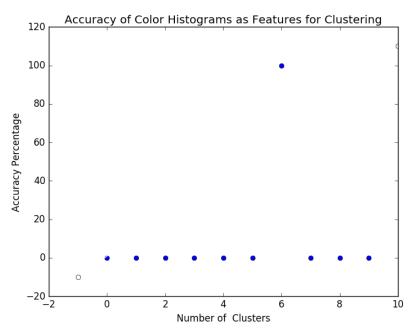


Figure 5: Accuracy of color histograms as features for clustering using k = 10

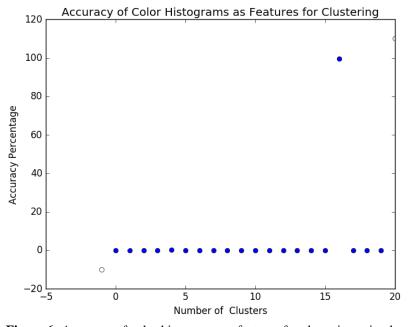


Figure 6: Accuracy of color histograms as features for clustering using k = 20

Computer Vision – Lab 4	Angel U. Ortega	Date	Page	
		11/26/2016 5:49 PM	8	

Figure:

Figures 7 and 8 show the accuracy results of using HOG as features for image clustering CIFAR-10. In comparison to color histograms as features for clustering, HOG had better accuracy results. While certain clusters were still very low with near-zero accuracy, half of the clusters had accuracy above 20% for k = 10. Like in P3, using k = 10 gave better results, on average.

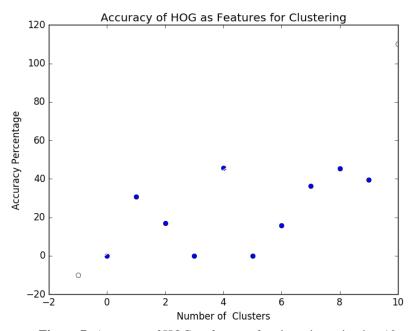


Figure 7: Accuracy of HOG as features for clustering using k = 10

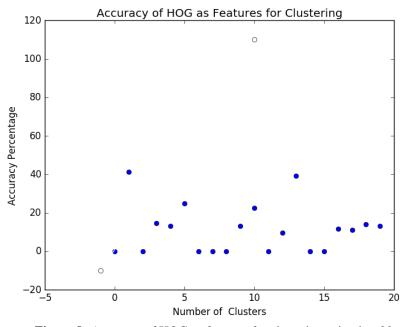


Figure 8: Accuracy of HOG as features for clustering using k = 20

Computer Vision – Lab 4	Angel U. Ortega	Date	Page
		11/26/2016 5:49 PM	9

3.5. P5

Figures 9 and 10 show the results for P5. I used k = 10 and k = 20 for the nearest neighbor classification. The blue dot in each graph represents the training set accuracy. The red dot represents the testing set accuracy. As expected, in both, the training had higher accuracy than the testing. A smaller number of neighbors returned a better accuracy.

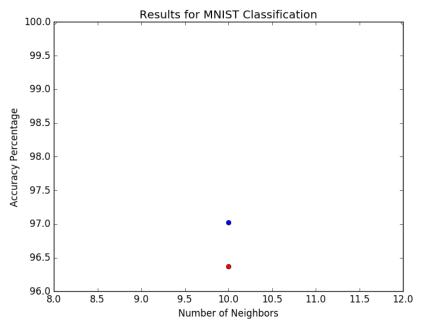


Figure 9: Results for P5 with 10 neighbors

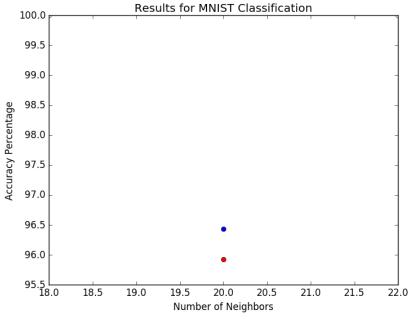


Figure 10: Results for P5 with 20 neighbors

Computer Vision – Lab 4	Angel U. Ortega	Date	Page	
		11/26/2016 5:49 PM	10	

3.6. P6

Accuracy of classification for CIFAR-10 was much lower than for MNIST. Pixel intensity as features had a much lower accuracy than HOG. Pixel intensity returned accuracy of approximately 11%. In comparison, using HOG returned accuracy of about 25%.

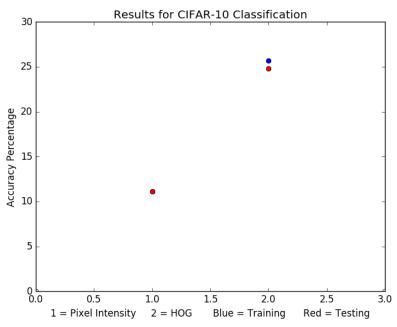


Figure 11: Results for P6

4. Appendix

4.1. P1 Code

```
# -*- coding: utf-8 -*-
"""

Created on Wed Oct 19 16:07:54 2016

@author: auort

Use k means clustering to posterize an image with n number of colors

"""

import numpy as np
import cv2

def getarray(img):
    array = img.reshape((-1,3))
    array = np.float32(array) #cv2.kmeans needs float array, soarray must be converted
    return array

def getimage(array, label, img):
    array = np.uint8(array)#convert from float to int
    result = array[label.flatten()]
```

Computer Vision – Lab 4	Angel U. Ortega	Date	Page
		11/26/2016 5:49 PM	11

```
resultimage = result.reshape((img.shape))#convert to original shape
    return resultimage
def posterize(imagename, numcolors, savedirectory):
    img = cv2.imread(imagename)
    array = getarray(img)
    criteria
               = (cv2.TERM CRITERIA EPS + cv2.TERM CRITERIA MAX ITER,
                                                                                     10,
1.0) #stop iteration if epsilon is reached or if max iterations has occured; max
iterations = 10, epsilon = 1.0
    ret, label, result=cv2.kmeans(array, numcolors, None, criteria,
                                                                                     10.
cv2.KMEANS RANDOM CENTERS)
    resultimage = getimage(result, label, img)
    cv2.imwrite(savedirectory, resultimage)
    cv2.imshow('posterized image using k means with k=' + str(numcolors) +' of
'+ imagename, resultimage)
    cv2.waitKey(0)
    cv2.destroyAllWindows()
posterize('fujismall.jpg', 1, "fuji1.jpg")
posterize('fujismall.jpg', 2, 'fuji2.jpg')
posterize('fujismall.jpg', 5, 'fuji5.jpg')
posterize('fujismall.jpg', 10, 'fuji10.jpg')
posterize('fujismall.jpg', 25, 'fuji25.jpg')
```

4.2. P2 Code

```
# -*- coding: utf-8 -*-
Created on Tue Nov 8 09:44:32 2016
@author: auort
from pylab import *
import numpy as np
import cv2
from loadMNIST import load mnist
import matplotlib.pyplot as plt
def classify digit(digit, labelled centroids):
    print "classify digit called!"
    mindistance = float("inf")
    for (label, centroid) in labelled centroids:
        distance = np.linalq.norm(centroid - digit)
        if distance < mindistance:</pre>
            mindistance = distance
            closest_centroid label = label
    return closest centroid label
def get error rate(digits, labelled centroids):
    print "get error rate called!"
```

Computer Vision – Lab 4	Angel U. Ortega	Date	Page
		11/26/2016 5:49 PM	12

```
classified incorrect = 0
    for (label, digit) in digits:
        classified label = classify digit(digit, labelled centroids)
        if classified label != label:
            classified incorrect +=1
    error rate = classified incorrect / float(len(digits))
    return error rate
trainImages, trainLabels = load mnist('training', digits=[0,1,2,3,4,5,6,7,8,9])
trainImages = np.float32(trainImages)
trainLabels = trainLabels.flatten()
# Define criteria = ( type, max iter = 10 , epsilon = 1.0 )
criteria = (cv2.TERM CRITERIA EPS + cv2.TERM CRITERIA MAX ITER, 10, 1.0)
# Set flags (Just to avoid line break in the code)
flags = cv2.KMEANS RANDOM CENTERS
error_rates = \{x: None for x in range(10,,10)+[100]\}
k = 10
# Apply KMeans
compactness,labels,centers = cv2.kmeans(trainImages,k,None,criteria,10,flags)
error rate = get error rate(centers, labels)
error rates[k] = error rate
k = 20
# Apply KMeans
compactness,labels,centers = cv2.kmeans(trainImages,k,None,criteria,10,flags)
error rate = get error rate(centers, labels)
error rates[k] = error rate
# Show the error rates
x axis = sorted(error rates.keys())
y axis = [error rates[key] for key in x axis]
plt.figure()
plt.title("Error Rate by Number of Clusters")
plt.scatter(x axis, y axis)
plt.xlabel("Number of Clusters")
plt.ylabel("Error Rate")
plt.show()
111
i = 0
for i in range(k):
   img1 = (centers[i,:])
    img1 = np.reshape(img1, (28,28))
    cv2.imwrite("centroidsB20 %d" % (i,) + ".png", img1)
    figure()
    gray()
    imshow(img1)
```

Computer Vision – Lab 4	Angel U. Ortega	Date	Page	
		11/26/2016 5:49 PM	13	

```
# -*- coding: utf-8 -*-
Created on Thu Nov 17 14:20:08 2016
@author: auort
import random
from base64 import b64decode
from json import loads
import numpy as np
import matplotlib.pyplot as plt
import time
def parse(x):
   print "parse called!"
    to parse the digits file into tuples of
    (labelled digit, numpy array of vector representation of digit)
    11 11 11
   digit = loads(x)
    array = np.fromstring(b64decode(digit["data"]),dtype=np.ubyte)
    array = array.astype(np.float64)
   return (digit["label"], array)
with open ("digits.base64.json", "r") as f:
    digits = map(parse, f.readlines())
# pick a ratio for splitting the digits list into a training and a validation
set.
ratio = int(len(digits)*0.25)
validation = digits[:ratio]
training = digits[ratio:]
def display digit(digit, labeled = True, title = ""):
    print "display digit called!"
    11 11 11
    graphically displays a 784x1 vector, representing a digit
    if labeled:
        digit = digit[1]
    image = digit
    plt.figure()
    fig = plt.imshow(image.reshape(28,28))
    fig.set cmap('gray r')
    fig.axes.get xaxis().set visible(False)
    fig.axes.get yaxis().set visible(False)
    if title != "":
        plt.title("Inferred label: " + str(title))
# writing Lloyd's Algorithm for K-Means clustering.
# (This exists in various libraries, but it's good practice to write by hand.)
def init centroids(labelled data,k):
   print "init centroids called!"
```

Computer Vision – Lab 4	Angel U. Ortega	Date	Page
		11/26/2016 5:49 PM	14

```
randomly pick some k centers from the data as starting values for centroids.
    Remove labels.
    return map(lambda x: x[1], random.sample(labelled data,k))
def sum cluster(labelled cluster):
   print "sum cluster called!"
   sum = labelled cluster[0][1].copy()
    for (label, vector) in labelled cluster[1:]:
        sum += vector
    return sum
def mean cluster (labelled cluster):
    print "mean cluster called!"
    sum of points = sum cluster(labelled cluster)
    mean of points = sum of points * (1.0 / len(labelled cluster))
    return mean of points
def form clusters(labelled data, unlabelled centroids):
    print "form clusters called!"
    # enumerate because centroids are arrays which are unhashable,
    centroids indices = range(len(unlabelled centroids))
    # initialize an empty list for each centroid. The list will contain
    # all the datapoints that are closer to that centroid than to any other.
    # That list is the cluster of that centroid.
    clusters = {c: [] for c in centroids indices}
    for (label, Xi) in labelled data:
        # for each datapoint, pick the closest centroid.
        smallest distance = float("inf")
        for cj index in centroids indices:
            cj = unlabelled centroids[cj index]
            distance = np.linalg.norm(Xi - cj)
            if distance < smallest distance:
                closest centroid index = cj index
                smallest distance = distance
        # allocate that datapoint to the cluster of that centroid.
        clusters[closest centroid index].append((label,Xi))
    return clusters.values()
def move centroids (labelled clusters):
    print "move centroids called!"
    returns a list of centroids corresponding to the clusters.
   new centroids = []
    for cluster in labelled clusters:
        new centroids.append(mean cluster(cluster))
    return new centroids
             repeat until convergence (labelled data,
def
                                                       labelled clusters,
unlabelled centroids):
    print "repeat until convergence called!"
    previous \max \text{ difference} = 0
   while True:
```

Computer Vision – Lab 4	Angel U. Ortega	Date	Page	
		11/26/2016 5:49 PM	15	

```
unlabelled old centroids = unlabelled centroids
        unlabelled centroids = move centroids(labelled clusters)
        labelled clusters = form clusters(labelled data, unlabelled centroids)
                        =
                               map(lambda
                                                      b:
                                                               np.linalg.norm(a-
        differences
                                                a,
b), unlabelled old centroids, unlabelled centroids)
        max difference = max(differences)
        difference change
                                                             abs((max difference-
previous max difference)/np.mean([previous max difference,max difference]))
100
        previous max difference = max_difference
        if np.isnan(difference change):
            break
    return labelled clusters, unlabelled_centroids
def cluster(labelled data, k):
    print "cluster called!"
    runs k-means clustering on the data. It is assumed that the data is
labelled.
    centroids = init centroids(labelled data, k)
    clusters = form clusters(labelled data, centroids)
    final clusters, final centroids = repeat until convergence (labelled data,
clusters, centroids)
    return final clusters, final centroids
def assign labels to centroids(clusters, centroids):
    print "assign labels to centroids called!"
    labelled centroids = []
    for i in range(len(clusters)):
        labels = map(lambda x: x[0], clusters[i])
        # pick the most common label
        most common = max(set(labels), key=labels.count)
        centroid = (most common, centroids[i])
        labelled centroids.append(centroid)
    return labelled centroids
def classify digit(digit, labelled centroids):
    print "classify digit called!"
    mindistance = float("inf")
    for (label, centroid) in labelled centroids:
        distance = np.linalg.norm(centroid - digit)
        if distance < mindistance:</pre>
            mindistance = distance
            closest centroid label = label
    return closest centroid label
def get error rate(digits, labelled centroids):
    print "get error rate called!"
    classified incorrect = 0
    for (label, digit) in digits:
        classified label = classify digit(digit, labelled centroids)
        if classified label != label:
```

Computer Vision – Lab 4	Angel U. Ortega	Date	Page
		11/26/2016 5:49 PM	16

```
classified incorrect +=1
    error rate = classified incorrect / float(len(digits))
    return error rate
error rates = \{x: None for x in range(5,10,10)+[100]\}
for k in range (5, 10, 10):
    print "K: %s" % k
    start time = time.time()
    trained clusters, trained centroids = cluster(training, k)
    labelled centroids
                            =
                                    assign labels to centroids (trained clusters,
trained centroids)
    error rate = get error rate(validation, labelled centroids)
    error rates[k] = error rate
    end time = time.time()
    run time = end time - start time
    print "Runtime: %s seconds" % run time
print "error_rates:"
print error rates
# Show the error rates
x axis = sorted(error rates.keys())
y axis = [error rates[key] for key in x axis]
plt.figure()
plt.title("Error Rate by Number of Clusters")
plt.scatter(x axis, y axis)
plt.xlabel("Number of Clusters")
plt.ylabel("Error Rate")
plt.show()
```

4.3. P3 Code

```
# -*- coding: utf-8 -*-
Created on Thu Nov 17 14:15:54 2016
@author: auort
import matplotlib.pyplot as plt
import numpy as np
def getAccuracy(derivLabels,k,origLabels):
   print(origLabels)
   data = np.zeros(k)
   for i in range (0, k):
        acc = [1 if derivLabels[x] == i else 0 for x in derivLabels]
        print(derivLabels)
        acc = np.multiply(acc,origLabels)
        print(acc)
        acc = [x for x in acc if x>0]
        data[i]=[i,acc]
   return data
def plotData(data, boundaries, gtitle, xlbl, ylbl):
   plt.plot(*zip(*data), marker='o', color='b', ls='')
   plt.plot(*zip(*boundaries), marker='o', color='w')
```

Computer Vision – Lab 4	Angel U. Ortega	Date	Page
		11/26/2016 5:49 PM	17

```
plt.title(gtitle)
    plt.xlabel(xlbl)
    plt.ylabel(ylbl)
    plt.show()
# -*- coding: utf-8 -*-
Created on Thu Nov 17 15:29:11 2016
@author: auort
from loadCIFAR2 import load cifar
import numpy as np
import cv2
import plot
def getCH(Im,n):
    colorHistogram = np.zeros((3,n))
    barSize = 255/(n-1)
    for i in range(len(Im)):
        for j in range(len(Im[i])):
            for k in range(3):
                index = Im[i][j][k]/barSize
                colorHistogram[k][index] = colorHistogram[k][index]+1
    return colorHistogram.flatten()
                                               load cifar("color", "data batch 1",
images,
                trainLabels
"C:/Users/auort/Desktop/CV_L4/RCode/cifar-10-batches-py/")
bins = 16
dataCH = np.zeros((len(trainLabels),bins*3))
for i in trainLabels:
    dataCH[i] = getCH(images[i], bins)
dataCH = np.float32(dataCH)
# Define criteria = ( type, max iter = 10 , epsilon = 1.0 )
criteria = (cv2.TERM CRITERIA EPS + cv2.TERM CRITERIA MAX ITER, 10, 1.0)
# Set flags (Just to avoid line break in the code)
flags = cv2.KMEANS RANDOM CENTERS
k = 20
# Apply KMeans
compactness,labels,centers = cv2.kmeans(dataCH,k,None,criteria,10,flags)
labels = labels.flatten()
data = plot.getAccuracy(labels,k,trainLabels)
title = "Accuracy of Color Histograms as Features for Clustering"
xlabel = "Number of Clusters"
ylabel = "Accuracy Percentage"
plot.plotData(data,[[-1,0],[k+1,100]], title, xlabel, ylabel)
```

4.4. P4 Code

```
Created on Sat Nov 19 18:06:43 2016
@author: auort
from PIL import Image
from collections import *
from loadCIFAR2 import load cifar
import plot
from HOG import imHOG
import numpy as np
import cv2
             trainLabels =
                                          load cifar("blackWhite", "data batch 1",
images,
"C:/Users/auort/Desktop/CV L4/RCode/cifar-10-batches-py/")
div,bins = 4,4
histData = np.zeros((len(trainLabels), div*div, bins))
for i in range(len(images)):
   histData[i] = imHOG(images[i], div, bins)
histData = np.float32(histData)
criteria = (cv2.TERM CRITERIA EPS + cv2.TERM CRITERIA MAX ITER, 10, 1.0)
flags = cv2.KMEANS RANDOM CENTERS
k = 10
compactness,labels,centers = cv2.kmeans(histData,k,None,criteria,10,flags)
labels = labels.flatten()
data = plot.getAccuracy(labels, k, trainLabels)
title = "Accuracy of Color Histograms as Features for Clustering"
xlabel = "Number of Clusters"
vlabel = "Accuracy Percentage"
plot.plotData(data,[[-1,0],[k+1,100]], title, xlabel, ylabel)
```

4.5. P5 Code

```
# -*- coding: utf-8 -*-
"""
Created on Sat Nov 19 14:48:28 2016

@author: auort
"""
import numpy as np
from loadMNIST import load_mnist
from sklearn.neighbors import KNeighborsClassifier
#from collections import *
import matplotlib.pyplot as plt

def train():
    trainIm, trainLb = load_mnist('training', digits=[0,1,2,3,4,5,6,7,8,9])
    trainIm = np.float32(trainIm).reshape((len(trainIm), -1))
    trainLb = trainLb.flatten()
    return trainIm, trainLb
```

Computer Vision – Lab 4	Angel U. Ortega	Date	Page
		11/26/2016 5:49 PM	19

```
def test():
   testIm, testLb = load mnist('testing', digits=[0,1,2,3,4,5,6,7,8,9])
   testIm = np.float32(testIm).reshape((len(testIm), -1))
   testLb = testLb.flatten()
   return testIm, testLb
def plotResult(training, testing, boundaries, gtitle, xlbl, ylbl):
   plt.plot(*zip(*training), marker='o', color='b', ls='')
   plt.plot(*zip(*testing), marker='o', color='r', ls='')
   plt.plot(*zip(*boundaries), marker='o', color='w')
   plt.title(gtitle)
   plt.xlabel(xlbl)
   plt.ylabel(ylbl)
   plt.show()
def classify(k):
   trainImages, trainLabels = train()
   testImages, testLabels = test()
   classifier = KNeighborsClassifier(n neighbors=k,p=2)
   classifier.fit(trainImages[:30000], trainLabels[:30000])
   training = [[1,classifier.score(trainImages, trainLabels)]]
   testing = [[1,classifier.score(testImages, testLabels)]]
   boundaries = [[k-2,100],[k+2,100]]
   title = "Accuracy of Classification"
   xlabel = "Number of Neighbors
   ylabel = "Accuracy Percentage"
   plotResult(training, testing, boundaries, title, xlabel, ylabel)
classify(10)
classify(20)
```

4.6. P6 Code

```
Created on Mon Nov 21 09:53:28 2016
@author: auort
11 11 11
import numpy as np
from loadCIFAR2 import load cifar
from sklearn.neural network import MLPClassifier
from HOG import imHOG
import matplotlib.pyplot as plt
import plot
def plotResult(training, testing, boundaries, gtitle, xlbl, ylbl):
    plt.plot(*zip(*training), marker='o', color='b', ls='')
    plt.plot(*zip(*testing), marker='o', color='r', ls='')
    plt.plot(*zip(*boundaries), marker='o', color='w')
    plt.title(gtitle)
    plt.xlabel(xlbl)
    plt.ylabel(ylbl)
    plt.show()
```

```
def zca whitening matrix (X):
    11 11 11
    Function to compute ZCA whitening matrix (aka Mahalanobis whitening).
    INPUT: X: [M x N] matrix.
       Rows: Variables
        Columns: Observations
    OUTPUT: ZCAMatrix: [M x M] matrix
    sigma = np.cov(X, rowvar=True)
    U,S,V = np.linalq.svd(sigma)
    epsilon = 1e-5
    ZCAMatrix = np.dot(U, np.dot(np.diag(1.0/np.sqrt(S + epsilon)), U.T)) # [M x
M]
    return ZCAMatrix
               trainLabels =
                                        load cifar("color", "data batch 1",
trainImages,
C:/Users/auort/Desktop/CV L4/RCode/cifar-10-batches-py/")
trainImages = np.float32(trainImages).reshape((len(trainImages), -1))
zxtrain = zca whitening matrix(trainImages)
trainImagesX = np.dot(zxtrain, trainImages)
trainLabels = np.array(trainLabels).flatten()
testImages,
                testLabels
                                        load cifar("color", "data batch 3",
C:/Users/auort/Desktop/CV L4/RCode/cifar-10-batches-py/")
testImages = np.float32(testImages).reshape((len(testImages), -1))
zxtest = zca whitening matrix(testImages)
testImagesX = np.dot(zxtest, testImages)
testLabels = np.array(testLabels).flatten()
classNorm = MLPClassifier()
classNorm.fit(trainImagesX, trainLabels)
#predictedLabels = classNorm.predict(testImages)
scoreNormTrain = classNorm.score(trainImagesX, trainLabels)
scoreNormTest = classNorm.score(testImagesX, testLabels)
                                   load cifar("blackWhite","data batch 1",
hogImages,
              hogLabels
                            =
C:/Users/auort/Desktop/CV L4/RCode/cifar-10-batches-py/")
hogTestImages, hogTestLabels = load cifar("blackWhite","data batch 3",
C:/Users/auort/Desktop/CV L4/RCode/cifar-10-batches-py/")
div.bins = 4.4
histData = np.zeros((len(trainLabels),div*div*bins))
histTestData = np.zeros((len(trainLabels),div*div*bins))
for i in range(len(hogImages)):
    histData[i] = imHOG(hogImages[i], div, bins).flatten()
    histTestData[i] = imHOG(hogTestImages[i], div, bins).flatten()
```

Computer Vision – Lab 4	Angel U. Ortega	Date	Page	
		11/26/2016 5:49 PM	21	

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