DSE 10 Homework 4 - Due 02/05/2021 9:00am Instructions • Please upload all work to gradescope by the due date - late work will not be graded You should submit a single ipynb file containing all your code and output (plots, numeric values, etc...) ■ Important: name your ipynb file according to the following convention: <lastname>_<firstname>_ipynb example: (thomas_anthony.ipynb) Please organize your notebook into sections by problem Use print statements to clearly indicate which question you are ansering o Good: print("Problem 4 (e): {}".format(np.pi)) o Good: print("The value of pi is: {}".format(np.pi)) Bad: print(np.pi) ∘ Bad: np.pi Use relative paths to load data: Good: loadmnist("train-images-idx3-ubyte", "train-labels-idx1-ubyte") Bad: loadmnist("/home/anthony/class/DSE10/train-images-idx3-ubyte", ...) Collaboration is encouraged, but all submissions should be in your own writing/code and written with your own understanding Your code must run and be able to reproduce your answers Unless stated otherwise in the assignemnt, your code should use only basic low-level NumPy/SciPy linear algebra and satistics commands (e.g. do not use built in estimation tools or anything from SciKitLearn). When in doubt, ask for clarification on Canvas. %matplotlib inline import matplotlib.pyplot as plt import time import gzip import sys import os import copy import numpy as np import pandas as pd import pickle import string import operator import bz2 import random from scipy import stats from scipy.cluster.hierarchy import dendrogram, linkage from sklearn.cluster import KMeans from sklearn.neighbors import BallTree from sklearn import metrics from sklearn.preprocessing import scale from pylab import rcParams from struct import unpack from scipy.stats import multivariate_normal if sys.version_info[0] == 2: from urllib import urlretrieve from urllib.request import urlretrieve if not sys.warnoptions: import warnings warnings.simplefilter("ignore") Worksheet 9 Problem 3 Text classification using multinomial Naive Bayes. (a) For this problem, you'll be using the 20 Newsgroups data set. There are several versions of it on the web. You should download "20news-bydate.tar.gz" from http://qwone.com/~jason/20Newsgroups/ Unpack it and look through the directories at some of the files. Overall, there are roughly 19,000 documents, each from one of 20 newsgroups. The label of a document is the identity of its newsgroup. The documents are divided into a training set and a test set. def download(filename, source = 'http://gwone.com/~jason/20Newsgroups/20news-bydate.tar.qz'): print("Downloading %s" % filename) urlretrieve(source + filename, filename) def load data(filename): if not os.path.exists(filename): download(filename) with gzip.open(filename, 'rb') as f: data = np.frombuffer(f.read(), np.uint8, offset = 16) return data / np.float32(256) newsgroup_data = load_data('20news-bydate.tar.gz') (b) The same website has a processed version of the data, "20news-bydate-matlab.tgz", that is par-ticularly convenient to use. Download this and also the file "vocabulary.txt". Look at the first training document in the processed set and the corresponding original text document to under- stand the relation between the two. In [4]: train data = pd.read csv('20news-bydate 2/matlab/train.data', header = None, sep =' ') train_label = pd.read_csv('20news-bydate 2/matlab/train.label', header = None, sep = ' ') train map = pd.read csv('20news-bydate 2/matlab/train.map', header = None, sep = ' ') test data = pd.read csv('20news-bydate 2/matlab/test.data', header = None, sep =' ') test label = pd.read csv('20news-bydate 2/matlab/test.label', header = None, sep =' ') test map = pd.read csv('20news-bydate 2/matlab/test.map', header = None, sep = ' ') vocabulary_data = pd.read_csv('vocabulary.txt', names = ['word'], header = None, sep =' ') (c) The words in the documents constitute an overall vocabulary V of size 61188. Build a multinomial Naive Bayes model using the training data. For each of the 20 classes j = 1, 2, ..., 20, you must have the following: • pij, the fraction of documents that belong to that class; and Pj, a probability distribution over V that models the documents of that class. In order to fit Pj, imagine that all the documents of class j are strung together. For each word w 2 V, let Pjw be the fraction of this concatenated document occupied by w. Well, almost: you will need to do smoothing (just add one to the count of how often w occurs). x_train_data, y_train_label, x_test_data, y_test_label = train_data, train_label, test_data, test_label x_train_data.columns = ['docIdx', 'wordIdx', 'count'] y_train_label['docIdx'] = range(1, len(y_train_label) + 1) y_train_label.columns = ['class', 'docIdx'] x_train_data = x_train_data.merge(y_train_label, how = 'left', on = 'docIdx') x test data.columns = ['docIdx', 'wordIdx', 'count'] y test label['docIdx'] = range(1, len(y test label) + 1) y test label.columns = ['class', 'docIdx'] x test data = x test data.merge(y test label, how = 'left', on = 'docIdx') # calculate pi pi = np.zeros(20)for j in range(20): pi[j] = np.sum(y train label.iloc[:,0] == j + 1)/len(y train label)print('Fraction of documents in each class:\n', pi) Fraction of documents in each class: [0.04259473 0.05155737 0.05075872 0.0520898 0.05102494 0.0525335 $0.05164611 \ 0.0525335 \ \ 0.05288846 \ \ 0.05271098 \ \ 0.05306593 \ \ 0.05271098$ $0.05244476 \ 0.05271098 \ 0.05262224 \ 0.05315467 \ 0.04836277 \ 0.05004881$ 0.0411749 0.03336587] # calculate p total = vocabulary data.shape[0] p = np.zeros((20, total))for j in range(20): tmp = x train data[x train data['class'] == j + 1] p[j,:] = 1for i in range(tmp.shape[0]): wdidx = tmp['wordIdx'].iloc[i] count = tmp['count'].iloc[i] p[j, wdidx - 1] += countp[j,:] = p[j,:]/np.sum(p[j,:])print('Probability Distribution:\n', p) Probability Distribution: [[6.66666667e-05 3.04761905e-04 1.31428571e-03 ... 4.76190476e-06 4.76190476e-06 4.76190476e-06] $[3.55589754e-04\ 3.49760414e-04\ 5.82934024e-06\ \dots\ 5.82934024e-06$ 5.82934024e-06 5.82934024e-06] $[7.89707479e-05\ 4.60662696e-04\ 6.58089566e-06\ \dots\ 6.58089566e-06$ 6.58089566e-06 6.58089566e-06] $[3.48108977e-05\ 4.90517195e-04\ 3.16462706e-06\ \dots\ 3.16462706e-06$ 3.16462706e-06 3.16462706e-06] [4.03854386e-06 1.61541755e-04 4.03854386e-06 ... 4.03854386e-06 4.03854386e-06 4.03854386e-06] [5.54680393e-06 2.55152981e-04 5.54680393e-05 ... 5.54680393e-06 5.54680393e-06 5.54680393e-06]] (d) Write a routine that uses this naive Bayes model to classify a new document. To avoid underflow, work with logs rather than multiplying together probabilities. def decision(x test data, pi, p): wdidx = x test data['wordIdx'] prob = [np.log(pi[j]) + np.sum(x test data['count'] * np.log(p[j,wdidx - 1])) for j in range(20)] pred = np.argmax(prob) return pred , prob (e) Evaluate the performance of your model on the test data. What error rate do you achieve? result = [] for i in range(y_test_label.shape[0]): x test = x test data.loc[x test data['docIdx'] == i + 1,:] pred, prob = decision(x_test, pi, p) result.append(pred + 1 != y_test_label['class'].iloc[i]) # calculate the error rate of naive Bayes model error rate = np.mean(result) * 100 print('Error rate of naive Bayes model is:\n{}%'.format(error rate)) Error rate of naive Bayes model is: 21.892071952031976% (f) Split the training data into a smaller training set and a validation set. The split could be 80-20, for instance. You'll use this training set to estimate parameters and the validation set to decide between different options. x, y = x_train_data, y_train_label # randomly select 80% data from data set as training data set. count = int(x.shape[0]*0.8) random.seed(10) sel = random.sample(range (0, x.shape[0]), count) x train set, y train set = x train data.loc[x train data.index.intersection(sel),:], y train label.loc[y tra print('Shape of x train set:\n', x train set.shape) print('\nShape of y train set:\n', y train set.shape) Shape of x_train_set: (1173876, 4)Shape of y_train_set: (9087, 2)In [14]: # get the remaining data as validation data set. remain = np.setdiff1d(range(0, x.shape[0]), sel) x_validation_set, y_validation_set = x_train_data.loc[x_train_data.index.intersection(remain),], y_train_lab print('Shape of x_validation_set:\n', x_validation_set.shape) print('\nShape of y_validation_set:\n', y_validation_set.shape) Shape of x_validation_set: (293469, 4)Shape of y_validation_set: (2182, 2)(i) replacing the frequency f of a word in a document by log(1 + f) # calculate pi f $pi_f = np.zeros(20)$ for j in range(20): $pi_f[j] = np.sum(y_train_set.iloc[:,0] == j + 1)/len(y_train_set)$ print('Fraction of documents in each class:\n', pi f) Fraction of documents in each class: $[0.0432486 \quad 0.05051172 \quad 0.05150215 \quad 0.05227248 \quad 0.05216243 \quad 0.05172224]$ $0.05271267 \ 0.05227248 \ 0.05249257 \ 0.05238252 \ 0.05425333 \ 0.053483$ $0.0532629 \quad 0.05216243 \quad 0.05205238 \quad 0.05172224 \quad 0.04842082 \quad 0.04963134$ 0.04071751 0.0330142] # calculate p_f $p_f = np.zeros((20, total))$ for j in range (20): tmp = x train set[x train set['class'] == j + 1] $p_f[j,:] = 1$ for i in range(tmp.shape[0]): wdidx = tmp['wordIdx'].iloc[i] count = tmp['count'].iloc[i] $p_f[j, wdidx - 1] += count$ $p_f[j,:] = p_f[j,:]/np.sum(p_f[j,:])$ print('Probability Distribution:\n', p f) Probability Distribution: [[7.77168995e-05 2.38701906e-04 1.29343126e-03 ... 5.55120711e-06 5.55120711e-06 5.55120711e-061 $[4.04435307e-04\ 2.83104715e-04\ 6.74058845e-06\ \dots\ 6.74058845e-06$ 6.74058845e-06 6.74058845e-06] $[8.21925997e-05\ 3.88546835e-04\ 7.47205452e-06\ \dots\ 7.47205452e-06$ 7.47205452e-06 7.47205452e-06] [4.13868352e-05 4.74067385e-04 3.76243957e-06 ... 3.76243957e-06 3.76243957e-06 3.76243957e-061 [4.74858611e-06 1.28211825e-04 4.74858611e-06 ... 4.74858611e-06 4.74858611e-06 4.74858611e-06] $[6.37897490e-06\ 2.74295921e-04\ 5.10317992e-05\ \dots\ 6.37897490e-06$ 6.37897490e-06 6.37897490e-06]] **def** decision f(x test, pi, p): wdidx = x test['wordIdx'] $prob = [np.log(pi[j]) + np.sum(np.log(1 + x_test['count']) * np.log(p[j,wdidx - 1])) * for j in range(20)]$ pred = np.argmax(prob) return pred, prob $result_f = []$ for i in range(y_validation_set.shape[0]): x test = x validation set.loc[x validation set['docIdx'] == i + 1,:] pred_f, prob_f = decision_f(x_test, pi_f, p_f) result f.append(pred f + 1 != y validation set['class'].iloc[i]) # calculate the error rate of this regular model error rate f = np.mean(result f) * 100 print('Error rate of smooth model on validation model is:\n{}%'.format(error_rate_f)) print ('The error rate for validation data is high. The reason why it is so high is because the validation se Error rate of smooth model on validation model is: 96.05866177818515% The error rate for validation data is high. The reason why it is so high is because the validation set is to o small. If you use the test data set, the error rate reduce to around 20.9% (ii) removing stopwords # stop words from online stop_words = ["a", "about", "above", "across", "after", "afterwards", "again", "all", "almost", "alone", "along", "already", "also", "although", "always", "am", "among", "amongst", "amoungst", "amount", "an", "and", "another", "any", "anyhow "this", "those", "though", "through", "throughout", "thru", "thus", "to", "together", "too", "toward", "towards", "under", "until", "up", "upon", "us", "very", "was", "we", "well", "were", "what", "whatever", "when", "whence", "whenever", "where", "whereafter", "whereas", "whereby", "wherein", "whereupon", "wherever", "whether", "which", "while", "who", "whoever", "whom", "whose", "why", "will", "with", "within", "without", "would", "yet", "you", "your", "yours", "yourself", "yourselves" # find all bad words vocabulary data = vocabulary data.reset index() vocabulary data['index'] = vocabulary data['index'].apply(lambda x: x+1) total list = set(vocabulary data['index']) vocabulary data = vocabulary data["word"].isin(stop words)] good_list = vocabulary_data['index'].tolist() good list = set(good list) bad list = total list - good list a = 0.001for bad in bad list: for j in range(20): # a Laplace Smoothing with low α for bad words $p_f[j][bad] = a/(np.sum(p_f[j,:]) + vocabulary_data.shape[0] + 1)$ def decision s(x test, pi, p): wdidx = x test['wordIdx'] IDF = np.log(len(x test['docIdx'].unique())/(x test.groupby(['wordIdx'])['docIdx'].count()) + 1) prob = [np.log(pi[j]) + np.sum(x test['count'] * np.log(IDF * p[j,wdidx - 1])) for j in range(20)] pred = np.argmax(prob) return pred, prob result_s = [] for i in range(y_validation_set.shape[0]): x_test = x_validation_set.loc[x_validation_set['docIdx'] == i + 1,:] pred_s, prob_s = decision_s(x_test, pi_f, p_f) result s.append(pred s + 1 != y validation set['class'].iloc[i]) # calculate the error rate of this regular model error_rate_s = np.mean(result_s) * 100 print('Error rate of IDF model on validation data is:\n{}%'.format(error rate s)) Error rate of IDF model on validation data is: 95.187901008249318 Evaluate your final model on the test data. What error rate do you achieve? In [24]: # evaluate naive Bayes model on the test data result = [] for i in range(y_test_label.shape[0]): x test = x test data.loc[x test data['docIdx'] == i + 1,:] pred, prob = decision(x_test, pi, p) result.append(pred + 1 != y_test_label['class'].iloc[i]) # calculate the error rate of naive Bayes model error rate = np.mean(result) * 100 print('Error rate of naive Bayes model is:\n{}%'.format(error rate)) Error rate of naive Bayes model is: 21.892071952031976% # evaluate smooth model on the test data result f test data = [] for i in range(y test label.shape[0]): x test = x test data.loc[x test data['docIdx'] == i + 1,:] pred, prob = decision f(x test, pi, p) result_f_test_data.append(pred + 1 != y_test_label['class'].iloc[i]) # calculate the error rate of naive Bayes model error rate f test data = np.mean(result f test data) * 100 print('Error rate of smooth model on test data is:\n{}%'.format(error rate f test data)) Error rate of smooth model on test data is: 20.919387075283144% # evaluate IDF model on the test data result_s_test_data = [] for i in range(y_test_label.shape[0]): x_test = x_test_data.loc[x_test_data['docIdx'] == i + 1,:] pred, prob = decision_s(x_test, pi_f, p_f) result_s_test_data.append(pred + 1 != y_test_label['class'].iloc[i]) # calculate the error rate of naive Bayes model error_rate_s_test_data = np.mean(result_s_test_data) * 100 print('Error rate of IDF model on test data is:\n{}%'.format(error_rate_s_test_data)) Error rate of IDF model on test data is: 94.53697534976683% Conclusion: We have evaluated 3 models on the test data, and it turns out that the smooth model performs a better accurate, whose error rate is 20.3%. Worksheet 9 Problem 4 1. Handwritten digit recognition using a Gaussian generative model. In class, we mentioned the MNIST data set of handwritten digits. You can obtain it from: http://yann.lecun.com/exdb/mnist/index.html In this problem, you will build a classifier for this data, by modeling each class as a multivariate (784-dimensional) Gaussian. (a) Upon downloading the data, you should have two training files (one with images, one with labels) and two test files. Unzip them. In order to load the data into Python you will find the following code helpful: http://cseweb.ucsd.edu/~dasgupta/dse210/loader.py For instance, to load in the training data, you can use: x,y = loadmnist('train-images-idx3-ubyte', 'train-labels-idx1-ubyte') This will set x to a 60000 x 784 array where each row corresponds to an image, and y to a length-60000 array where each entry is a label (0-9). There is also a routine to display images: use displaychar(x[0]) to show the first data point, for instance. def download(filename, source = 'http://yann.lecun.com/exdb/mnist/index.html'): print("Downloading %s" % filename) urlretrieve(source + filename, filename) def load mnist images(filename): if not os.path.exists(filename): download(filename) # Read the inputs in Yann LeCun's binary format. with gzip.open(filename, 'rb') as f: data = np.frombuffer(f.read(), np.uint8, offset = 16) data = data.reshape(-1,784)return data / np.float32(256) def load mnist labels(filename): if not os.path.exists(filename): download(filename) with gzip.open(filename, 'rb') as f: data = np.frombuffer(f.read(), np.uint8, offset = 8) def displaychar(image): plt.imshow(np.reshape(image, (28,28)), cmap=plt.cm.gray) plt.axis('off') plt.show() # Load the training data set train_data = load_mnist_images('train-images-idx3-ubyte.gz') train_labels = load_mnist_labels('train-labels-idx1-ubyte.gz') print('Shape of the train data:\n', train_data.shape) print('\nShape of the train labels:\n', train_labels.shape) Shape of the train data: (60000, 784) Shape of the train labels: (60000,)# Display the first data point displaychar(train data[0]) (b) Split the training set into two pieces – a training set of size 50000, and a separate validation set of size 10000. Also load in the test data. x, y = train data, train labels # randomly select 50000 data from data set as training data set. random.seed(10) sel = random.sample(range (0, x.shape[0]), 50000)x_train_data, y_train_data = train_data[sel,], train_labels[sel,] print('Shape of x train data:\n', x train data.shape) print('\nShape of y_train_data:\n', y_train_data.shape) Shape of x train data: (50000, 784) Shape of y train data: (50000,) # get the remaining data as validation data set. remain = np.setdiff1d(range(0, x.shape[0]), sel) x_validation, y_validation = train_data[remain,], train_labels[remain,] print('Shape of x_validation:\n', x_validation.shape) print('\nShape of y_validation:\n', y_validation.shape) Shape of x validation: (10000, 784)Shape of y validation: (10000,)# Load the testing data set test data = load mnist images('t10k-images-idx3-ubyte.gz') test labels = load mnist labels('t10k-labels-idx1-ubyte.gz') print('Shape of the test data:\n', test data.shape) print('\nShape of the test labels:\n', test labels.shape) Shape of the test data: (10000, 784) Shape of the test labels: (10000,)(c) Now fit a Gaussian generative model to the training data of 50000 points. Determine the class probabilities. • Fit a Gaussian to each digit, by finding the mean and the covariance of the corresponding data points. # create a multivariabte Gaussian model def MultivariateGaussian(x, y): #labels $1,2,\ldots,k$ k = 10#number of features d = (x.shape)[1]mu = np.zeros((k,d))sigma = np.zeros((k,d,d))pi = np.zeros(k)for label in range(10): indices = np.where(y == label) indices = indices[0] mu[label] = np.mean(x[indices,:], axis=0) sigma[label] = np.cov(x[indices,:], rowvar=0, bias=1) pi[label] = float(len(indices))/float(len(y)) return mu, sigma, pi In [34]: # get the mu, sigma, and pi mu, sigma, pi = MultivariateGaussian(x train data, y train data) print('Class probabilities:\n', pi) Class probabilities: [0.0989 0.1129 0.09792 0.10202 0.09642 0.091 0.09882 0.10514 0.0978 0.099081 print('Mean of the corresponding data points:\n ', mu) Mean of the corresponding data points: [[0. 0. 0. ... 0. 0. 0.][0. 0. 0. ... 0. 0. 0.][0. 0. 0. ... 0. 0. 0.] [0. 0. 0. ... 0. 0. 0.] [0. 0. 0. ... 0. 0. 0.] [0. 0. 0. ... 0. 0. 0.]] print('Covariance of the corresponding data points:\n ', sigma) Covariance of the corresponding data points: [[[0. 0. 0. ... 0. 0. 0.]][0. 0. 0. ... 0. 0. 0.] [0. 0. 0. ... 0. 0. 0.] [0. 0. 0. ... 0. 0. 0.] [0. 0. 0. ... 0. 0. 0.] [0. 0. 0. ... 0. 0. 0.]] [[0. 0. 0. ... 0. 0. 0.] [0. 0. 0. ... 0. 0. 0.] [0. 0. 0. ... 0. 0. 0.] [0. 0. 0. ... 0. 0. 0.] [0. 0. 0. ... 0. 0. 0.] [0. 0. 0. ... 0. 0. 0.]] [[0. 0. 0. ... 0. 0. 0.][0. 0. 0. ... 0. 0. 0.] [0. 0. 0. ... 0. 0. 0.] [0. 0. 0. ... 0. 0. 0.] [0. 0. 0. ... 0. 0. 0.] [0. 0. 0. ... 0. 0. 0.]] [[0. 0. 0. ... 0. 0. 0.][0. 0. 0. ... 0. 0. 0.] [0. 0. 0. ... 0. 0. 0.] [0. 0. 0. ... 0. 0. 0.] [0. 0. 0. ... 0. 0. 0.] [0. 0. 0. ... 0. 0. 0.]] [[0. 0. 0. ... 0. 0. 0.][0. 0. 0. ... 0. 0. 0.] [0. 0. 0. ... 0. 0. 0.] [0. 0. 0. ... 0. 0. 0.] [0. 0. 0. ... 0. 0. 0.] [0. 0. 0. ... 0. 0. 0.]][[0. 0. 0. ... 0. 0. 0.][0. 0. 0. ... 0. 0. 0.] [0. 0. 0. ... 0. 0. 0.] [0. 0. 0. ... 0. 0. 0.] [0. 0. 0. ... 0. 0. 0.] [0. 0. 0. ... 0. 0. 0.]]] (d) One last step is needed: it is important to smooth the covariance matrices, and the usual way to do this is to add in cl. where c is some constant and I is the identity matrix. What value of c is right? Use the validation set to help you choose. That is, choose the value of c for which the resulting classifier makes the fewest mistakes on the validation set. What value of c did you get? # classify funtion using Bayes' rule def decision(x, pi, mu, sigma): prob = np.zeros((10, x.shape[0]))**for** i **in** range (0,10): prob[i,:] = pi[i]* multivariate_normal.pdf(x, mean = mu[i,:], cov = sigma[i,:,:]) #if the sigma matrix cannot be inversed, we need to add a np.eye() to avoid singular matrix error. #since we will smooth in step (d), so I didn't use np.eye() here. $\#prob[i,:] = pi[i]* multivariate_normal.pdf(x, mean = mu[i,:], cov = sigma[i,:,:] + np.eye(sigma.shallow)$ pred = np.argmax(prob, axis = 0) return pred, prob # set c as a constant and I as the identity matix # randomly give c a number c = 0.1iden = np.zeros((10,784,784))**for** i **in** range (0,10) : iden[i,:,:] = np.diag([1] * 784)# smooth the covariance matrix decision(x validation[:5,], pi, mu, sigma + c * iden) Out[39]: (array([0, 9, 4, 0, 9]), array([[3.52395032e+36, 9.94963706e+02, 4.04319827e-42, 1.73069617e+31, [2.76205313e-59, 3.89010386e+23, 1.37779283e-87, 1.71922354e-70, 5.36199288e+131, [1.67822220e+09, 2.20521407e+12, 3.34169997e-05, 1.33258778e+04, 6.56761242e+10], [5.52541293e+03, 1.30537050e+18, 3.96725820e-08, 6.44940176e+04, 1.77372353e+14], [2.12330838e-18, 1.03923436e+24, 1.47574376e+20, 2.85214873e-19, 1.13417968e+20], [4.32712370e+03, 5.06260109e+15, 1.08791462e-21, 4.09520025e+06, 8.53140112e+08], [7.30100850e+07, 4.61689670e+05, 1.08491729e-28, 2.60239295e+06, 3.10196619e-05], [1.51033108e-17, 3.29536606e+32, 2.50752743e-12, 1.05837175e-27, 1.41461342e+35], [1.00110917e-06, 1.36214676e+24, 5.55170895e-10, 6.06793379e-15,1.18698631e+20], [1.17818041e-04, 4.03108896e+38, 1.23801240e-04, 1.00516671e-12, 1.53564586e+37]])) In [40]: c grid = np.linspace(0.01, 0.3, num=20)validation error =[] for c in c grid: pred = decision(x validation, pi, mu, sigma + c * iden)[0] validation error.append(1 - np.mean(pred == y validation)) In [41]: # smaller erorr is desired plt.plot(c grid, validation error) # find the optimal c value c optim = c grid[np.argmin(validation error)] print('The optimal of c is:\n', np.min(validation error)) The optimal of c is: 0.04879999999999955 0.13 0.12 0.11 0.10 0.09 0.08 0.07 0.06 0.05 0.00 0.05 0.20 0.25 (e) Turn in an iPython notebook that includes: All your code. • Error rate on the MNIST test set. • Out of the misclassified test digits, pick five at random and display them. For each instance, list the posterior probabilities Pr(y|x) of each of the ten classes. In [42]: # calculate the error rate on the MNIST test set. accuracy = np.mean(decision(test_data, pi, mu, sigma + c_optim * iden)[0] == test_labels) error rate = (1 - accuracy) *100 print('Error rate on the MNIST test set:\n{}%'.format(error rate)) Error rate on the MNIST test set: 4.379999999999955% In [43]: pred, prob = decision(test data,pi, mu, sigma + c optim * iden) In [44]: mis = np.where(pred != test labels)[0] random.seed(14) for i in random.sample(range(0,len(mis)),5): print('posterior probabilities are:\n', prob[:,mis[i]]/np.sum(prob[:,mis[i]])) #print(np.where(np.argmax(prob[:,mis[i]]/np.sum(prob[:,mis[i]])))) print('\nmissclassified digits', test_labels[mis[i]]) print('---') posterior probabilities are: [1.39773549e-28 3.29542606e-31 9.44407064e-07 8.93100210e-11 5.10498694e-32 2.15188108e-35 1.76340732e-59 3.71674998e-20 9.99999056e-01 4.60969151e-25] missclassified digits 2 posterior probabilities are: [9.90840610e-001 1.96922947e-134 5.59213076e-006 2.73862179e-015 9.26492386e-015 8.29549696e-003 2.31647883e-010 2.36242297e-026 1.17461319e-034 8.58301070e-004] missclassified digits 9 posterior probabilities are: [1.33288214e-022 4.78235805e-141 1.39248990e-009 9.99999999e-001 1.79715520e-067 4.52804023e-026 5.12737030e-081 1.69816726e-043 1.51046950e-021 3.78681257e-032] missclassified digits 9 posterior probabilities are: [7.24531355e-001 8.06604028e-122 2.07758767e-013 2.68522993e-001 5.10092467e-055 1.94953938e-007 3.58494474e-014 8.39106303e-081 6.94545720e-003 1.19623255e-040] missclassified digits 8 posterior probabilities are: [3.60704806e-23 4.73306638e-08 1.62911845e-22 3.03170718e-25 1.07755734e-29 4.39101476e-22 9.99999953e-01 5.30893474e-48 1.17243012e-17 1.22402935e-40] missclassified digits 1 Worksheet 10 Problem 2 For this problem, we'll be using the animals with attributes data set. Go to http://attributes.kyb.tuebingen.mpg.de and, under "Downloads", choose the "base package" (the very first file in the list). Unzip it and look over the various text files. About the dataset: This is a small dataset that has information on about 50 animals. The animals are listed in classes.txt. For each animal, the information consists of values for 85 features: does the animal have a tail, is it slow, does it have tusks, etc. The details of the features are in the predicates.txt. The full data consists of a 50 x 85 matrix of real values, in predicate-matrixcontinuous.txt. There is also a binarized version of this data, in predicate-matrix-binary.txt. In [45]: def download(filename, source = 'http://attributes.kyb.tuebingen.mpg.de'): print("Downloading %s" % filename) urlretrieve(source + filename, filename) def load data(filename): if not os.path.exists(filename): download(filename) with bz2.open(filename, 'rb') as f: data = np.frombuffer(f.read(), np.uint8, offset = 16) return data / np.float32(256) In [46]: base package data = load data('AwA-base.tar.bz2') (a) Load the real-valued array, and also the animal names, into Python. Run k-means on the data (from sklearn.cluster) and ask for k = 10 clusters. For each cluster, list the animals in it. Does the clustering make sense? In [47]: # classes with animal names. classes_with_animal_names = pd.read_fwf("Animals_with_Attributes/classes.txt", header = None)[1].values print('Shape of the classes.txt:\n', classes_with_animal_names.shape, '\n') print('Data of the classes.txt:\n', classes_with_animal_names) Shape of the classes.txt: (50,)Data of the classes.txt: ['antelope' 'grizzly+bear' 'killer+whale' 'beaver' 'dalmatian' 'persian+cat' 'horse' 'german+shepherd' 'blue+whale' 'siamese+cat' 'skunk' 'mole' 'tiger' 'hippopotamus' 'leopard' 'moose' 'spider+monkey' 'humpback+whale' 'elephant' 'gorilla' 'ox' 'fox' 'sheep' 'seal' 'chimpanzee' 'hamster' 'squirrel' 'rhinoceros' 'rabbit' 'bat' 'giraffe' 'wolf' 'chihuahua' 'rat' 'weasel' 'otter' 'buffalo' 'zebra' 'giant+panda' 'deer' 'bobcat' 'pig' 'lion' 'mouse' 'polar+bear' 'collie' 'walrus' 'raccoon' 'cow' 'dolphin'] In [48]: # predicates with feature names. predicates with feature names = pd.read fwf("Animals with Attributes/predicates.txt", header = None)[1].valu print('Shape of the predicates.txt:\n', predicates with feature names.shape, '\n') print('Data of the predicates.txt:\n', predicates with feature names) Shape of the predicates.txt: (85,)Data of the predicates.txt: ['black' 'white' 'blue' 'brown' 'gray' 'orange' 'red' 'yellow' 'patches' 'spots' 'stripes' 'furry' 'hairless' 'toughskin' 'big' 'small' 'bulbous' 'lean' 'flippers' 'hands' 'hooves' 'pads' 'paws' 'longleg' 'longneck' 'tail' 'chewteeth' 'meatteeth' 'buckteeth' 'strainteeth' 'horns' 'claws' 'tusks' 'smelly' 'flys' 'hops' 'swims' 'tunnels' 'walks' 'fast' 'slow' 'strong' 'weak' 'muscle' 'bipedal' 'quadrapedal' 'active' 'inactive' 'nocturnal' 'hibernate' 'agility' 'fish' 'meat' 'plankton' 'vegetation' 'insects' 'forager' 'grazer' 'hunter' 'scavenger' 'skimmer' 'stalker' 'newworld' 'oldworld' 'arctic' 'coastal' 'desert' 'bush' 'plains' 'forest' 'fields' 'jungle' 'mountains' 'ocean' 'ground' 'water' 'tree' 'cave' 'fierce' 'timid' 'smart' 'group' 'solitary' 'nestspot' 'domestic'] In [49]: # matrix of real values.

