DSE 210: final

Q4, Q6, Q7, Q8

```
In [2]: %pylab inline
   import numpy as np
```

import sklearn as sk

Populating the interactive namespace from numpy and matplotlib

Q4

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/ (http://qwone.com/~jason/20Newsgroups/) The same website has a processed version of the data, "20news-bydate-matlab.tgz", that is particularly 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 understand the relation between the two. The words in the documents constitute an overall vocabulary V of size 61188. Build a Bernoulli Naive Bayes model using the training data. 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.

In [2]: !curl -0 http://qwone.com/~jason/20Newsgroups/20news-bydate-matlab.tgz
!tar xzvf 20news-bydate-matlab.tgz

%	Total	%	Receive	d %)	Kferd	_	•		Time Spent		Current Speed
0	0	0	0	0	0	0	0	::	::	::	0
0	0	0	0	0	0	0	0	::	0:00:01	::	0
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100	7398k	100	7398k	0	0	1673k	0	0:00:04	0:00:04	::	1678k

20news-bydate/matlab/

20news-bydate/matlab/train.data

20news-bydate/matlab/train.label

20news-bydate/matlab/train.map

20news-bydate/matlab/test.data

20news-bydate/matlab/test.label

20news-bydate/matlab/test.map

0 [main] tar 14144 find_fast_cwd: WARNING: Couldn't compute FAST_CWD poin ter. Please report this problem to the public mailing list cygwin@cygwin.com 3/17/2017 fin:

```
In [3]: data train = np.loadtxt('20news-bydate/matlab/train.data',dtype='int')
         Y train = np.loadtxt('20news-bydate/matlab/train.label',dtype='int')
         data test = np.loadtxt('20news-bydate/matlab/test.data',dtype='int')
         Y test = np.loadtxt('20news-bydate/matlab/test.label',dtype='int')
         max_doc,max_word,max_count = np.amax(data_train, axis=0)
         max\_word = 61188
         X train = np.zeros((max doc,max word), dtype=np.dtype('b'))
         for d in data train:
             doc, word = d[0], d[1]
             X_{train[doc-1][word-1]} = 1
         max_doc_test,dummy1,dummy2 = np.amax(data_test, axis=0)
         X_test = np.zeros((max_doc_test,max_word), dtype=np.dtype('b'))
         for d in data test:
             doc, word = d[0], d[1]
             X_{\text{test[doc-1][word-1]}} = 1
         print X_train.shape
         print X_test.shape
         class names = []
         with open('20news-bydate/matlab/train.map') as f:
             class_names = [1.split(' ')[0] for 1 in iter(f)]
         (11269L, 61188L)
         (7505L, 61188L)
In [14]: # (a) Evaluate the performance of your model on the test data. What error rate do
         from sklearn.naive bayes import BernoulliNB
         clf = BernoulliNB()
         clf.fit(X train, Y train)
         Y predict = clf.predict(X test)
```

error rate is: 0.37601598934

errors = np.sum(Y_test != Y_predict)

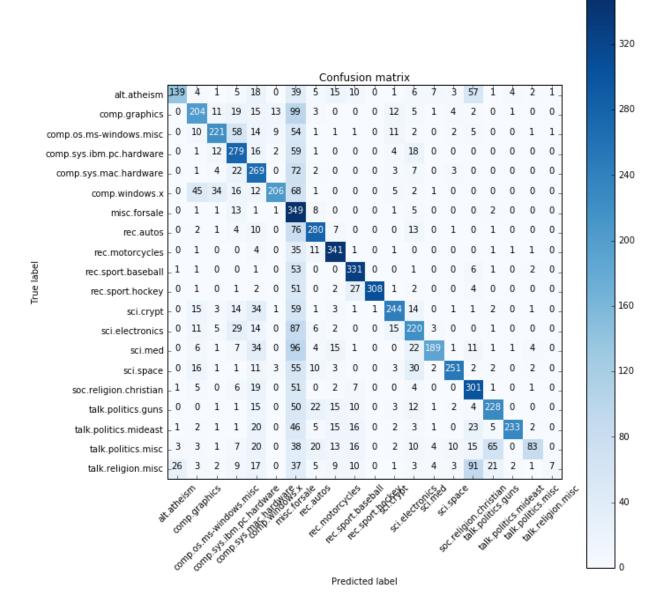
print "error rate is:", float(errors) / len(Y test)

```
In [10]: # (b) Evaluate your final model on the test data. Construct a confusion matrix.
         import itertools
         from sklearn.metrics import confusion matrix
         def plot_confusion_matrix(cm, classes,
                                    normalize=False,
                                    title='Confusion matrix',
                                    cmap=plt.cm.Blues):
              .. .. ..
             This function prints and plots the confusion matrix.
             Normalization can be applied by setting `normalize=True`.
             plt.imshow(cm, interpolation='nearest', cmap=cmap)
             plt.title(title)
             plt.colorbar()
             tick_marks = np.arange(len(classes))
             plt.xticks(tick_marks, classes, rotation=45)
             plt.yticks(tick_marks, classes)
             if normalize:
                  cm = cm.astype('float') / cm.sum(axis=1)[:, np.newaxis]
                  print("Normalized confusion matrix")
             else:
                  print('Confusion matrix, without normalization')
             print(cm)
             thresh = cm.max() / 2.
             for i, j in itertools.product(range(cm.shape[0]), range(cm.shape[1])):
                  plt.text(j, i, cm[i, j],
                           horizontalalignment="center",
                           color="white" if cm[i, j] > thresh else "black")
             plt.tight layout()
             plt.ylabel('True label')
             plt.xlabel('Predicted label')
         # Compute confusion matrix
         cnf matrix = confusion matrix(Y test, Y predict)
         np.set_printoptions(precision=2)
         # Plot non-normalized confusion matrix
         plt.figure(figsize=(10, 10))
         plot confusion matrix(cnf matrix, classes=class names)
         plt.show()
         Confusion matrix, without normalization
         [[139
                 4
                     1
                         5
                            18
                                     39
                                          5 15
                                                10
                                                      0
                                                          1
                                                              6
                                                                  7
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                                                                         57
                                                                              1
                                                                                   4
             2
                 1]
            0 204 11 19 15 13
                                     99
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                                                         12
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          0
                10 221 58 14
                                     54
                                          1
                                              1
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                                    59
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```

1

0

	0	0]																
[0	45 0]	34	16	12	206	68	1	0	0	0	5	2	1	0	0	0	0
[0 0	1 0]	1	13	1	1	349	8	0	0	0	1	5	0	0	0	2	0
[0 0	2 0]	1	4	10	0	76	280	7	0	0	0	13	0	1	0	1	0
[0 1	1 0]	0	0	4	0	35	11	341	1	0	1	0	0	0	0	1	1
[1 2	1 0]	0	0	1	0	53	0	0	331	0	0	1	0	0	6	1	0
[0 0	1 0]	0	1	2	0	51	0	2	27	308	1	2	0	0	4	0	0
[0 1	15 0]	3	14	34	1	59	1	3	1	1	244	14	0	1	1	2	0
[0 0	11 0]	5	29	14	0	87	6	2	0	0	15	220	3	0	0	1	0
[9 4	6 0]	1	7	34	0	96	4	15	1	0	0	22	189	1	11	1	1
[0	16 0]	1	1	11	3	55	10	3	0	0	3	30	2	251	2	2	0
[1 1	5 0]	0	6	19	0	51	0	2	7	0	0	4	0	0	301	1	0
[0 0	0 0]	1	1	15	0	50	22	15	10	0	3	12	1	2	4	228	0
[1 2	2 0]	1	1	20	0	46	5	15	16	0	2	3	1	0	23	5	233
[3 83	3 0]	1	7	20	0	38	20	13	16	0	2	10	4	10	15	65	0
[26 1	3 7]]	2	9	17	0	37	5	9	10	0	1	3	4	3	91	21	2

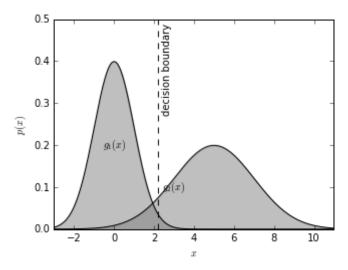


Q6

Urn A contains a Gaussian pdf: N(0, σ =1); Urn B contains another Gaussian N(5, σ =2); We draw a number and it is X=2.5. P(UrnA)=2P(UrnB) Determine the likelihood of (UrnA | X=2.5)? Determine a decision boundary (Urn A vs Urn B) for this problem.

```
In [16]:
         import numpy as np
         from scipy.stats import norm
         p X A = norm(0, 1).pdf(2.5)
         p_X_B = norm(5, 2).pdf(2.5)
         posterior_AX = p_X_A * 2/3
         posterior_BX = p_X_B * 1/3
         likelihood_A_X = posterior_AX / (posterior_AX + posterior_BX)
         print "likehood of (UrnA | X=2.5) is ", likelihood_A_X
         # Compute the two PDFs
         x = np.linspace(-3, 12, 1000)
         pdf1 = norm(0, 1).pdf(x)
         pdf2 = norm(5, 2).pdf(x)
         x_{bound} = x[np.where(2*pdf1 < pdf2)][0]
         # Plot the pdfs and decision boundary
         fig = plt.figure(figsize=(5, 3.75))
         ax = fig.add_subplot(111)
         ax.plot(x, pdf1, '-k', lw=1)
         ax.fill_between(x, pdf1, color='gray', alpha=0.5)
         ax.plot(x, pdf2, '-k', lw=1)
         ax.fill_between(x, pdf2, color='gray', alpha=0.5)
         # plot decision boundary
         ax.plot([x bound, x bound], [0, 0.5], '--k')
         ax.text(x bound + 0.2, 0.49, "decision boundary",
                  ha='left', va='top', rotation=90)
         ax.text(0, 0.2, '$g 1(x)$', ha='center', va='center')
         ax.text(3, 0.1, '$g_2(x)$', ha='center', va='center')
         ax.set xlim(-3, 11)
         ax.set_ylim(0, 0.5)
         ax.set xlabel('$x$')
         ax.set ylabel('$p(x)$')
         plt.show()
         print "decision boundary is: ", x_bound
```

likehood of (UrnA | X=2.5) is 0.277387907451



decision boundary is: 2.1951951952

Q7

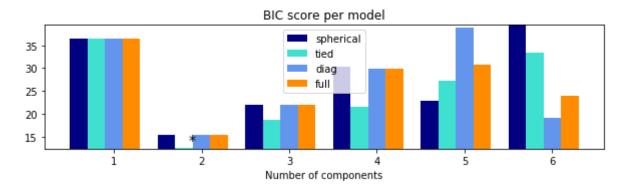
Consider the following observations: X=(-0.1,-0.2, 0.1, 0.2, 0, 0.1, -0.1, 0, -0.05, 0.1, 1.05, 1.1, 0.9, 0.8, 0.9, 1, 1.2, 1.1,1.2, .9) Cluster this data into two classes using the K-means algorithm. What are the cluster centers?

Q8

Construct a Gaussian Mixture Model for the above data. What is your estimate for the number of mixtures?

```
In [14]: import itertools
         from sklearn import mixture
         # Number of samples per component
         lowest_bic = np.infty
         bic = []
         n_components_range = range(1, 7)
         cv_types = ['spherical', 'tied', 'diag', 'full']
         for cv_type in cv_types:
             for n_components in n_components_range:
                  # Fit a Gaussian mixture with EM
                  gmm = mixture.GaussianMixture(n_components=n_components,
                                                covariance_type=cv_type)
                  gmm.fit(X)
                 bic.append(gmm.bic(X))
                  if bic[-1] < lowest_bic:</pre>
                      lowest_bic = bic[-1]
                      best_gmm = gmm
         bic = np.array(bic)
         color_iter = itertools.cycle(['navy', 'turquoise', 'cornflowerblue',
                                        'darkorange'])
         clf = best_gmm
         bars = []
         # Plot the BIC scores
         plt.figure(figsize=(10, 5))
         spl = plt.subplot(2, 1, 1)
         for i, (cv_type, color) in enumerate(zip(cv_types, color_iter)):
             xpos = np.array(n_components_range) + .2 * (i - 2)
             bars.append(plt.bar(xpos, bic[i * len(n_components_range):
                                            (i + 1) * len(n_components_range)],
                                  width=.2, color=color))
         plt.xticks(n components range)
         plt.ylim([bic.min() * 1.01 - .01 * bic.max(), bic.max()])
         plt.title('BIC score per model')
         xpos = np.mod(bic.argmin(), len(n_components_range)) + .65 +\
              .2 * np.floor(bic.argmin() / len(n_components_range))
         plt.text(xpos, bic.min() * 0.97 + .03 * bic.max(), '*', fontsize=14)
         spl.set_xlabel('Number of components')
         spl.legend([b[0] for b in bars], cv types)
```

Out[14]: <matplotlib.legend.Legend at 0xc958208>



Based on BIC score above, 2 mixtures seem to be good estimate