CS5304 - Machine Learning - Assignment 1

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For this assignment, you will be required to classify documents from the RCV1 dataset, which is an archive of over 800k manually categorized newswire stories made available by Reuters, Ltd. for research purposes [1]. The first 23,149 samples will be used for training, 10k are randomly chosen from the remaining for validation, and 50k will be randomly chosen for testing. The ids used for validation are provided in validation.txt and the ids for testing are hidden. Use scikit-learn [2] to load the data.

[1]: http://www.jmlr.org/papers/volume5/lewis04a/lewis04a.pdf (http://www.jmlr.org/papers/volume5/lewis04a/lewis04a.pdf)

[2]: http://scikit-learn.org/stable/datasets/rcv1.html (http://scikit-learn.org/stable/datasets/rcv1.html)

1. k-Nearest-Neighbors and Naive Bayes

1a. (3 Points) Classification with kNN.

Use the KNeighborsClassifier from scikit- learn to classify documents in the validation set. Only consider the labels from labels.txt. Find a good value for K by exploring a few options. Wrap your classifier in a Python class called CS5304KNNClassifier and it should have a train and predict method. We will be calling these methods during grading. You'll also need to submit a file called ks.txt that includes the value for K you've chosen for each label. For this exercise, only use the first 1000 training examples.

```
In [5]: def load_training_data():
             data = fetch rcv1(subset='train')
             return data.data, data.target.toarray(), data.sample_id
 In [6]: def load validation data(path to ids):
             data = fetch_rcv1(subset='test')
             ids = pd.read_csv(path_to_ids, names=['id'], dtype=np.int32)
             mask = np.isin(data.sample id, ids['id'])
             validation data = data.data[mask]
             validation_target = data.target[mask].toarray()
             validation ids = data.sample id[mask]
             return validation_data, validation_target, validation_ids
In [54]: class CS5304BaseClassifier(object):
             def __init__(self):
                 pass
             def train(self, x, y):
                 raise NotImplementedError
             def predict(self, x):
                 raise NotImplementedError
In [52]:
         from sklearn.neighbors import KNeighborsClassifier
         class CS5304KNNClassifier:
             def init (self, n neighbors=3):
                 super(CS5304KNNClassifier, self).__init__()
                 self.n neighbors = n neighbors
             def train(self, X train, y train):
                 self.model = KNeighborsClassifier(n neighbors=self.n neighbors)
                 self.model.fit(X train, y train)
             def predict(self, X test):
                 if self.model:
                     return self.model.predict(X test)
                 else:
                     raise "No model has been trained yet"
             def get score(self, X test, y test):
                 if self.model:
                     return self.model.score(X test, y test)
                 else:
                     raise "No model has been trained yet"
 In [9]: train data, train target, train ids = load training data()
```

eval data, eval target, eval ids = load validation data('validation.txt'

X, y = train_data[:1000, :], train target[:1000, :]

Choose K

In order to choose an ideal k for each label, we run a 5-fold cross validation on training data for each label. For each label, we tried different odd number k, ranging from 1 to 41. For each k value, we got 5 accuracy score for each fold and took an average. Then, we compared the average accuracy score for each k, picked the one with highest accuracy, and stored it for each k.

```
In [165]:
          #cross validation (5-fold) on training set to choose k
          def choose k(X, y, label):
              k_and_accuracy = dict()
              for k in range(1, 41, 2):
                  my knn = CS5304KNNClassifier(k)
                  avg = 0.0
                  for train_index, test_index in kf.split(X):
                      X_train, X_test = X[train_index], X[test_index]
                      y_train, y_test = y[train_index], y[test_index]
                      my_knn.train(X_train, y_train)
                      output = my_knn.predict(X_test)
                      check output(output, y test)
                      avg += my_knn.get_score(X_test, y_test)
                  k_and_accuracy[k] = avg / kf.get_n_splits()
                  #print ("The average accuracy for k =", k, "is", avg / kf.get n
          splits())
              max accuracy = max(k and accuracy.values())
              optimal_k = [k for k, v in k_and_accuracy.items() if v == max accura
          cy][0]
              print ("Optimal k for label", label, "is ",
                     optimal k, " and accuracy is", max accuracy)
              return label, optimal k
          #check output dimension
          def check output(output, y):
              assert type(output) == np.ndarray
              assert output.ndim == 1
              assert output.shape[0] == y.shape[0]
```

Below shows the optimal k for each label and its corresponding accuracy score from cross validation.

```
In [166]: from sklearn.model selection import KFold
         kf = KFold(n splits=5)
         #getting k for each binary classifer and store them in a dictionary
         k for labels = dict()
         for label in load labels('labels.txt'):
             k, v = choose_k(X, y[:, label], label)
             k \text{ for labels}[k] = v
         k_for_labels
         Optimal k for label 4 is 27
                                    and accuracy is 0.940000000000001
         Optimal k for label 5 is 35 and accuracy is 0.952
         Optimal k for label 7 is 27 and accuracy is 0.942
         Optimal k for label 33 is 3 and accuracy is 0.859
         Optimal k for label 70 is 5 and accuracy is 0.891
         Optimal k for label 83 is 7 and accuracy is 0.959
         Optimal k for label 95 is 17 and accuracy is 0.955000000000001
         Optimal k for label 98 is 3 and accuracy is 0.96
         Optimal k for label 102 is 39 and accuracy is 0.923
Out[166]: {4: 27, 5: 35, 7: 27, 33: 3, 59: 7, 70: 5, 83: 7, 95: 17, 98: 3, 102: 3
         9}
```

Then, we run KNN with the validation set with the k we picked above. Below shows the accuracy score for the validation set with label and k indicated on KNN model.

```
Validation Result for KNN

For label 4 and k = 27 accuracy: 0.9221

For label 5 and k = 35 accuracy: 0.9518

For label 7 and k = 27 accuracy: 0.909

For label 33 and k = 3 accuracy: 0.8101

For label 59 and k = 7 accuracy: 0.8907

For label 70 and k = 5 accuracy: 0.9

For label 83 and k = 7 accuracy: 0.9533

For label 95 and k = 17 accuracy: 0.9649

For label 98 and k = 3 accuracy: 0.9537

For label 102 and k = 39 accuracy: 0.9126
```

1b. (3 Points) Classification with NB.

Use the BernoulliNB from scikit-learn to classify documents in the validation set. Only consider the labels from labels.txt. Wrap your classifier in a Python class called CS5304NBClassifier and it should have a train and predict method. Use all the training examples.

```
In [55]: from sklearn.naive bayes import BernoulliNB
         class CS5304NBClassifier(CS5304BaseClassifier):
             def __init__(self):
                 super(CS5304NBClassifier, self). init ()
             def train(self, X_train, y_train):
                  self.model = BernoulliNB().fit(X train, y train)
             def predict(self, X test):
                 if self.model:
                      return self.model.predict(X test)
                 else:
                     raise "No model has been trained yet"
             def get score(self, X test, y test):
                 if self.model:
                      return self.model.score(X test, y test)
                 else:
                      raise "No model has been trained yet"
```

Below shows the accuracy score for the validation set with label and k indicated on Bernoulli NB Classifier.

```
Validation result for Bernoulli NB Classifier
For label 4 , the accuracy on validation set is:
                                                  0.8109
For label 5 , the accuracy on validation set is:
                                                  0.8824
For label 7 , the accuracy on validation set is:
                                                  0.9212
For label 33 , the accuracy on validation set is: 0.7931
For label 59 , the accuracy on validation set is:
                                                  0.9115
For label 70 , the accuracy on validation set is:
                                                   0.8991
For label 83 , the accuracy on validation set is:
                                                   0.9033
For label 95 , the accuracy on validation set is:
                                                   0.9634
For label 98 , the accuracy on validation set is:
                                                   0.9735
For label 102, the accuracy on validation set is: 0.9018
```

1c. (1 Points) Data Analysis.

Which label is the most difficult to classify? Why do you think this is the case?

Answer

From both of the validation results above (KNN and Bernoulli_NB), we can see that despite of different classification method, the class with label 33 has the lowest accuracy (0.8101 for KNN and 0.7931 for Bernoulli_NB), thereby the most difficult one to classify.

By examing the prior distribution on both the training (first 1,000 and total of 23,149) and the validation set, we can see that label 33 takes up about 50% of the total numbers while all the other labels are not. We suspect this is the reason that it is the most difficult to classify.

Since there are a lot of data points with label 33, we expect the variance within the class to be high. We can also confirm that by looking at the best k chosen by cross validation, even though there are many true labels of 33, the best k (k = 3) is surprisingly small. This is probably resulted from the variance within class is high. Thus, a higher value of K would lead to worse performance for label 33.

As for Naive Bayes Classifier, it employs a very simple linear hypothesis function and it also assumes the feature to be independent. Therefore, NB suffers from high bias, as it could not sufficiently represent many complex situations (data is very spread out in our case). The simplicity of the model leads to the low performance on this particular label.

```
In [441]: #check prior distribution for each classifier in the training set
          # (first 1000 samples from training data)
          print ("Prior distribution for each classifier in the first 1,000 traini
          ng samples")
          for label in load_labels('labels.txt'):
              print (label, ": ", np.sum(train_target[:1000, label])/train_target
          [:1000, label].shape[0])
          Prior distribution for each classifier in the first 1,000 training samp
          les
          4: 0.2
          5: 0.146
          7: 0.058
          33: 0.536
          59: 0.151
          70:
               0.276
          83: 0.064
          95: 0.072
          98: 0.084
          102: 0.239
In [442]:
         #check prior distribution for training set (all samples from training da
          ta)
          print ("Prior distribution for each classifier in all training samples")
          for label in load_labels('labels.txt'):
              print (label, ": ", np.sum(train_target[:, label])/train_target[:,la
          bel].shape[0])
          Prior distribution for each classifier in all training samples
          4: 0.18052615663743574
          5: 0.1022074387662534
          7: 0.08337293187610696
          33: 0.46593805347963196
          59: 0.14899131711953
          70: 0.3010929197805521
          83 : 0.07114778176163117
          95 : 0.06894466283640761
          98: 0.10976716056849108
```

102: 0.254093049375783

```
In [443]: #check prior distribution for validation set (all samples from validatio
          n set)
          print ("Prior distribution for each classifier for all samples")
          for label in load_labels('labels.txt'):
              print (label, ": ", np.sum(eval_target[:, label])/eval_target[:,labe
          1].shape[0])
          Prior distribution for each classifier for all samples
               0.1947
          5:
               0.107
               0.0922
          33: 0.4804
          59:
                0.1421
                0.2935
          70:
                0.0703
          95:
                0.0677
          98:
                0.1059
```

2. k-Means

102: 0.2578

2a. (6 Points) Classification with k-Means.

Use the KMeans from scikit-learn to classify documents in the validation set. Only consider the first 10 labels. Wrap this kMeans classifier in CS5304KMeansClassifier with a train and predict method, which will be called during grading. Use all the training examples.

Hint: Find the centroids for the training data. Pass these to scikit-learn's KMeans as the value for init.

```
In [ ]: from sklearn.cluster import KMeans
        import scipy
        class CS5304KMeansClassifier():
            def __init__(self, n_clusters = 2):
                self.n_clusters = n_clusters
            def train(self, X, y):
                init centroids = self.get_init_centroid(X,y)
                self.model = KMeans(n_clusters=self.n_clusters,
                                     init=init centroids, n init=1).fit(X)
            def predict(self, X):
                if self.model:
                    return self.model.predict(X)
                else:
                    raise "No model has been trained yet"
            def get init centroid(self, X, y):
                true set = [i for i, j in enumerate(y) if j == 1]
                false_set = [i for i, j in enumerate(y) if j == 0]
                true_cluster = X[true_set]
                false_cluster = X[false_set]
                true_centroid = X[true_set].mean(axis = 0)
                false centroid = X[false set].mean(axis = 0)
                return np.concatenate((false_centroid, true_centroid), axis = 0)
            def get score(self, X, y):
                if self.model:
                    n samples = X.shape[0]
                    y hat = np.array(self.model.predict(X)).flatten()
                    y true label = np.array(y).flatten()
                    count = len([i for i in range(n samples)
                                  if y_hat[i] == y_true_label[i]])
                    return count / float(n_samples)
                else:
                    raise "No model has been trained yet"
            def get final centroid(self):
                return self.model.cluster centers
In [ ]: for label in load_labels('labels.txt'):
            kmeans classifier = CS5304KMeansClassifier()
            kmeans classifier.train(train data, train target[:, label])
```

```
print ("For label", label, ", the accuracy score is:",
          kmeans_classifier.get_score(eval_data, eval_target[:, label
]))
```

2b. (1 Point) Visualizing Clusters.

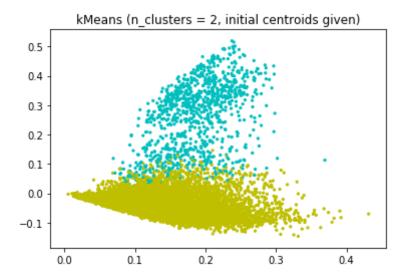
Visualize the predictions from KMeans for label 33 with a 2-dimensional plot. You'll need to choose a dimensionality reduction technique (such as TruncatedSVD). For plotting, use matplotlib.

```
In [11]:
         %matplotlib inline
         import matplotlib
         import matplotlib.pyplot as plt
In [12]:
         kmeans classifier = CS5304KMeansClassifier()
         kmeans_classifier.train(train_data, train_target[:, 33])
         kmeans_classifier.get_score(eval_data, eval_target[:, 33])
Out[12]: 0.6193
In [13]:
         predicted_33 = kmeans_classifier.predict(eval_data)
In [14]:
         from sklearn.decomposition import TruncatedSVD
         from sklearn.random projection import sparse random matrix
         svd = TruncatedSVD(n components=2, random state=42)
         eval_data_truncated = svd.fit_transform(eval_data)
In [35]: def get_color(y):
             color = []
             for label in y:
                 if label == 1:
                      color.append('c')
                 else:
                      color.append('y')
             return color
In [36]: def draw plot(X, Y):
             color_array = get_color(y)
             plt.scatter(X[:,0], X[:,1],color=color array, s=5)
```

Below is the 2-dimensional plot to visualize predictions from KMeans for label 33.

```
In [37]: draw_plot(eval_data_truncated, predicted_33)
   plt.title("kMeans (n_clusters = 2, initial centroids given)")
```

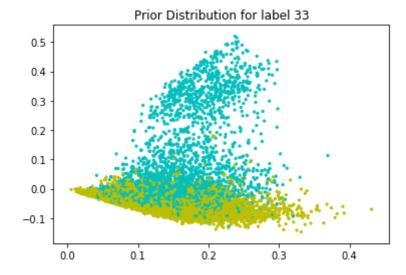
Out[37]: Text(0.5,1,'kMeans (n_clusters = 2, initial centroids given)')



Below is the 2-dimensional plot to visualize distribution for label 33 in the evaluation dataset.

```
In [38]: draw_plot(eval_data_truncated, eval_target[:, 33])
   plt.title("Prior Distribution for label 33")
```

Out[38]: Text(0.5,1,'Prior Distribution for label 33')



2c. (1 Point) Data Analysis.

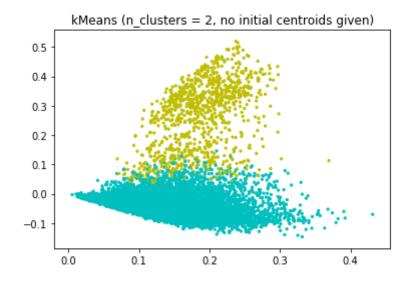
Train a new KMeans model for label 33. This time, initialize the centroids randomly (if you use the scikit-learn method fit, then don't provide a value for y). Visualize the clusters using the same dimensions as from 2b. Do the clusters look similar as when you provided the labels? What do the clusters look like if you use 3 or 4 clusters instead of 2? Explain why.

Answer

Comparing the clusters using the same dimensions as from 2b, we can see that the two clusters looks very similar and almost identical (the points are cluttering with each other so it is hard to tell, but it should be the same).

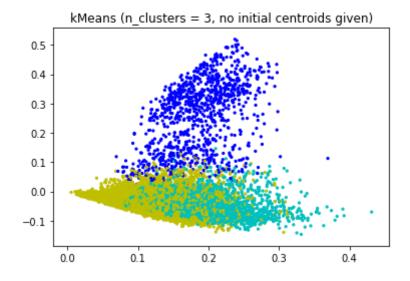
This make senses. If we refer back to the prior distribution we get from 1c, we can see that almost half of the data points (4804) in the evaluation set is labeled 33. So even if we don't give the classifier with an initial centroid, the algorithm will adjust parameters to reduce the loss functions at each step. In this case, to reduce the loss when the number of cluster is 2, it is not surprising to see that the algorithm converges to the same local minimum regardless of what initilization condition is given.

If using 3 or 4 clusters instead of two, the additional clusters look like they are dividing on top of the result from using 2 clusters only. When using 3 clusters, the bottom half is further divided into two. When using 4 clusters, the top half is further divided into two. This can also be explained by the reason mentioned above. When giving the k-means classifier more clusters, the algorithm is still looking to reduce the loss functions at each step. Therefore, it makes sense that the clusters with 3 and 4 clusters still look similar to the ones with 2 clusters only.



```
In [42]: kMeans_random_3_cluster = KMeans(n_clusters=3).fit(train_data)
    random_predicted_3_cluster = kMeans_random_3_cluster.predict(eval_data)
```

Out[48]: Text(0.5,1,'kMeans (n_clusters = 3, no initial centroids given)')



In [46]: kMeans_random_4_cluster = KMeans(n_clusters=4).fit(train_data)
 random_predicted_4_cluster = kMeans_random_4_cluster.fit_predict(eval_data)

Out[56]: Text(0.5,1,'kMeans (n_clusters = 4, no initial centroids given)')

