Technical Report of the Document Clustering Project with Python

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Section 1: General Objectives of the Chapter

In this chapter, we aim to examine the application of data mining in textual datasets. By merging the concepts of textual data with clustering, we will arrive at a broader topic known as Document Clustering. In fact, we will utilize data mining algorithms with the approach that our data unlike usual- is text and, generally, documents. In text mining, we aim to extract hidden, valuable, and meaningful insights and relationships from seemingly ordinary and worthless textual data.

Section 2: Dataset characteristics

The data used in this project is inspired by a well-known travel agency company called TripAdvisor. This dataset includes 1,850 textual data entries (documents), each containing reviews and ratings provided by the site -based on the opinions of previous travelers- regarding each hotel. Since this number of documents is large for our work and slows down execution, the first 100 documents from this collection have been selected as the input and raw data. This selection does not follow any specific order and can be purely random as well.

Section 3: Programming Language and IDE Description

The programming languages R and Python are the most well-known and powerful languages for data science. In this project, we have used *Python 3.5*; there is a powerful distribution of Python called *Anaconda*, which has been specifically designed and developed for such projects. This distribution includes various packages and libraries for implementing data mining techniques.

Both the Anaconda environment and all Python IDEs can be used to develop and execute the program code. In this project, we utilized the popular Python programming IDE called *PyCharm Community*, version 2016.1.4. The only remaining point is that we must first provide the path of the Anaconda interpreter to the software through $File \rightarrow Settings \rightarrow Project$ Interpreter so that its libraries become operational and ready to import and use.

Section 4: Data Cleansing and Clustering Phase

In general, the program consists of 4 functions, which are illustrated in the figure below:

```
#def tokenize_and_stem_and_removestopwords(input_text):...
#def k_means(tfidfvector, numofclusters):...
#def show_result(result_array, clustercounts):...
#def draw_shape(a, b, c):...
```

Initially, the following code snippet is executed:

```
1
      import pandas as pd
 2
      import nltk
 3
      from nltk.stem.porter import *
 4
      import string
 5
      from sklearn.feature extraction.text import TfidfVectorizer
 6
      from sklearn.cluster import KMeans
 7
      import os
      import matplotlib.pyplot as plt
 9
      from sklearn.manifold import MDS
10
      from sklearn.metrics.pairwise import cosine similarity
11
      from scipy.cluster.hierarchy import ward, dendrogram
12
      path = "C:/Users/vaio/Desktop/ramin-project/source/TripAdvisor_First_100_Hotels"
13
14
      token dict = {}
15
      stopwords = nltk.corpus.stopwords.words('english')
16
      stemmer = PorterStemmer()
     document names = []
```

In the above figure, all the necessary libraries and packages for the project are first imported. Then, in lines 12 to 17, we specify the path of the dataset and prepare the groundwork for the body of the codes.

Note: To run the program, you must change the given directory shown as the "path" variable to the existing path on your hard drive.

Next, the following section will be run and executed:

```
120
        # iterate throw data collection
121
        for subdir, dirs, files in os.walk(path):
122
            for file in files:
123
                file path = subdir + os.path.sep + file
124
                shakes = open(file path, encoding="utf8")
                new path = file path.replace(file path, file path[72:90])
125
126
                document names.append(new path)
127
                text = shakes.read()
128
                lowers = text.lower()
129
                no punctuation = lowers.translate(string.punctuation)
130
                token dict[file] = no punctuation
```

In this section, we read the files in the initial path one by one, convert the text inside them to lowercase, and store the result in *token_dict* which is a list object. Additionally, we will store a list of the names of each document in the *document names* list object.

Then it's time to execute the following code snippet:

```
flag = True
141
       ans 1 = input("Please enter number of clusters(k for k-means): ")
142
       while flag:
144
               val = int(ans 1)
145
              flag = False
146
           except ValueError:
               print ("That's not an int!")
147
148
       tfidf = TfidfVectorizer(tokenizer=tokenize and stem and removestopwords, stop words='english', ngram_range=(1, 2))
149
       tfs = tfidf.fit transform(token dict.values())
```

With the execution of the above code, the user is first asked about setting the parameter K, which represents the number of clusters. Then, using the definition made in line 133, the initial settings for creating the *tf-idf matrix* for the tokens are established. An important point is considering the parameter ngram range=(1,2), which indicates that we also want to obtain the 1-grams and 2-grams of each token (Following RapidMiner parameters). The tokenizer parameter specifies how the tokens are extracted, which is set to the tokeniz and stem and removestopwords(), which we will explain further. Finally, we assign *token dict* for execution to *tfidf* (line 135).

Now, the function <code>tokeniz_and_stem_and_removestopwords()</code> is executed. Its internal body is shown in the figure below. As the name suggests, this function takes text as input and performs tokenization, stemming, and the removal of stopwords in the English language. Ultimately, the output of this function consists of words ready for data mining. For this section, we have utilized the <code>nltk library</code>.

```
def tokenize and stem and removestopwords(input text): # Tokenize, Stem and remove stopwords
22
          tokens = [word for sent in nltk.sent tokenize(input text) for word in nltk.word tokenize(sent)]
23
          filtered stopwords = [w for w in tokens if w not in stopwords]
24
          filtered tokens = []
25
          for token in filtered_stopwords:
26
               if re.search('[a-zA-Z]', token):
27
                  filtered tokens.append(token)
28
          stems = [stemmer.stem(t) for t in filtered tokens]
29
          return stems
```

The *stopwords* and *stemmer* variables that were initialized earlier were used here (in the figure above). In this function, the operations of token extraction (line 22), removal of frequent words (line 23), removal of all characters except for the English alphabet (lines 25 to 27), and finally stemming (line 28) are performed. The output of this function is stored in the variable *tfs* after being converted to *tf-idf* numerical vectors.

Now the following snippet will be executed, The $k_means()$ function is called with the parameters tfs and ans_1 (which represents the number of clusters). In this function, the main operation of clustering the documents is performed. The input parameters of this function are the tf-idf matrix of the documents and the number of clusters. The internal body of this function is illustrated in the figure below.

```
136    result = k_means(tfs, ans_1)
```

```
32
       def k means (tfidfvector, numofclusters):
           cluster id = []
33
           doc id per cluster = []
34
           km = KMeans(n clusters=numofclusters)
35
           km.fit(tfidfvector)
36
37
           clusters = km.labels .tolist()
           for k in range(0, numofclusters):
38
39
               cluster id.append([])
               doc id per cluster.append([])
40
           for k in range(0, len(clusters)):
41
42
               for i in range(0, numofclusters):
                   if clusters[k] == i:
43
44
                       cluster id[i].append(clusters[k])
45
                       doc id per cluster[i].append(k)
           return clusters, cluster id, doc id per cluster
46
```

Now it is time to display the results obtained from the clustering in the output. To do this, we pass the output of the k_means function to the $show_result()$ (result, ans_1) function. Additionally, the number of clusters is also one of the input parameters of this function (line 151). The internal body of this function is shown below.

```
51
      def show result(result array, clustercounts):
52
          print("**There are ", len(result array[0]), " documents in TripAdvisor data collection**")
53
          print("**The data collection is clustered in ", clustercounts, "clusters as shown in below**")
54
          print("##########")
55
          for j in range(0, len(result_array[1])):
              print("There are: ", len(result array[1][j]), " documents in cluster: ", j)
56
              print ("Here we can see names of the documents:")
57
58
              print()
59
              for m in result array[2][j]:
                  print(document names[m])
60
61
              print()
              print("======"")
62
```

After executing the above function, we can observe the clustering broken down by the name and the number of documents present in each cluster. We will see this result in the results analysis section.

After executing the above function, most of the clustering work has been completed. The only remaining part is the section below, which begins with a question for the user.

```
ans_2 = input("Do you want to see clustering visual shapes? y or n: ")
if ans_2 == "y":
    draw_shape(result[0], document_names, tfs, int(ans_1))
```

Two different types of charts have been prepared for the graphical representation of the document clustering. If the user wishes to view them, they should answer "y" to the above question and press the Enter key. In this case, the last function of this project, namely the <code>draw_shape()</code> function, will be called with the input parameters <code>result[0] = clusters</code>, <code>document_names</code>, <code>tfs</code>, and the number of clusters (<code>ans_1</code>). The first part of the internal body of this function is shown in the figure below.

```
65
       def draw shape (a, b, c, d):
           dist = 1 - cosine similarity(c)
67
           mds = MDS(n components=2, dissimilarity="precomputed", random state=1)
68
           pos = mds.fit transform(dist)
69
           xs, ys = pos[:, 0], pos[:, 1]
           cluster colors = []
70
           cluster names = []
71
           prefix = "cluster "
72
73
           for i in range (0, d):
74
              cluster_colors.append(colorize())
75
              cluster names.append(prefix+str(i))
76
           df = pd.DataFrame(dict(x=xs, y=ys, label=a, title=b))
77
           groups = df.groupby('label')
78
           fig, ax = plt.subplots(figsize=(17, 9)) # set size
79
           ax.margins(0.05) # Optional, just adds 5% padding to the autoscaling
```

```
80
            for name, group in groups:
 81
                ax.plot(group.x, group.y, marker='o', linestyle='', ms=12,
 82
                        label=cluster names[name], color=cluster colors[name], mec='none')
 83
                ax.set aspect('auto')
                ax.tick_params(
 84
                    axis='x', # changes apply to the x-axis
 85
 86
                    which='both', # both major and minor ticks are affected
 87
                    bottom='off', # ticks along the bottom edge are off
 88
                    top='off', # ticks along the top edge are off
                    labelbottom='off')
 89
 90
                ax.tick params (
 91
                    axis='y', # changes apply to the y-axis
 92
                    which='both', # both major and minor ticks are affected
                    left='off', # ticks along the bottom edge are off
 93
 94
                    top='off', # ticks along the top edge are off
 95
                    labelleft='off')
 96
            ax.legend(numpoints=1) # show legend with only 1 point
 97
            # add label in x,y position with the label as the film title
 98
            for i in range (len (df)):
 99
                ax.text(df.ix[i]['x'], df.ix[i]['y'], df.ix[i]['title'], size=5)
100
            plt.show()
101
            plt.close()
```

By executing the above section and with the help of the *matplotlib.pyplot library*, which is specifically for drawing various charts, we can observe the clustering results graphically, differentiated by color (each document in a cluster has a unique color). An important point is that a different color and name are generated for the number of clusters that the user provided in the first step of the program (parameter k). Lines 70 to 75 contain the remaining lines of the program related to chart drawing settings, which we will skip explaining. In the next section, we will see this chart.

The second part of this function is related to drawing the *hierarchical clustering* chart of the documents, where we do not focus on the number of clusters but rather perform clustering based on the similarity obtained between the documents (line 66 – cosine similarity). The result of this section is the well-known Dendrogram chart. The code for this part is also shown in the figure below.

An important point in this section is the name in line 106, which is set for the parameter *labels = document_names*. This ensures that the names of the documents are displayed as labels on the leaves of the chart. This chart is also drawn with the help of the *matplotlib.pyplot library*. In the next section, we will also see the result of this part.

```
103
            # dendrogram plot
104
           linkage matrix = ward(dist) # define the linkage matrix using ward clustering pre-computed distances
            fig, ax = plt.subplots(figsize=(15, 20)) # set size
105
            ax = dendrogram(linkage matrix, orientation="top", labels=document names, leaf font size=6)
106
107
           plt.tick params (
108
               axis='x', # changes apply to the x-axis
109
               which='both', # both major and minor ticks are affected
110
               bottom='on', # ticks along the bottom edge are off
               top='on', # ticks along the top edge are off
111
112
                labelbottom='on')
113
            plt.show()
114
           plt.close()
115
           return True
```

Section Five: Displaying and Analyzing Results

Before starting this section, it is essential to mention that since the K-means algorithm operates randomly at its initial stage and selects starting points without a specific order, different output results may be observed each time the program is executed.

After executing the program, we can observe the following result, which is part of the output of this stage (here we consider K=5):

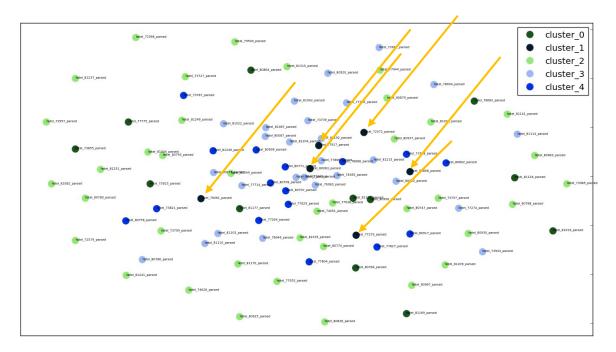
```
D:\Anaconda3\python.exe C:/Users/vaio/ramin/__init__.py
Please enter number of clusters (k for k-means): 5
**There are 100 documents in TripAdvisor data collection**
**The data collection is clustered in 5 clusters as shown in below**
***************
There are: 12 documents in cluster: 0
Here we can see names of the documents:
hotel_73855_parsed
hotel_73923_parsed
hotel_77775_parsed
hotel_78682_parsed
hotel_80784_parsed
hotel_80864_parsed
hotel_80990_parsed
hotel_81019_parsed
hotel_81126_parsed
hotel_81165_parsed
hotel_81169_parsed
hotel_81177_parsed
There are: 6 documents in cluster: 1
Here we can see names of the documents:
hotel_72572_parsed
hotel_76061_parsed
hotel_77270_parsed
hotel_77917_parsed
hotel_80083_parsed
hotel 80808 parsed
There are: 39 documents in cluster: 2
Here we can see names of the documents:
hotel_72579_parsed
hotel_72586_parsed
hotel_73727_parsed
```

Here, we can see the names of the documents within each cluster in a separate manner. For example, in cluster number 0, there are 12 documents, and their names are as follows from top to bottom:

```
hotel_73855_parsed
hotel_73923_parsed
hotel_77775_parsed
hotel_78682_parsed
hotel_80784_parsed
hotel_80864_parsed
hotel_80990_parsed
hotel_81019_parsed
hotel_81126_parsed
hotel_81165_parsed
hotel_81169_parsed
hotel_81177_parsed
```

For another example, we will focus on the documents within cluster number 1. This cluster contains 6 documents, which are highlighted in the first above figure. Now, let's move on to displaying the charts.

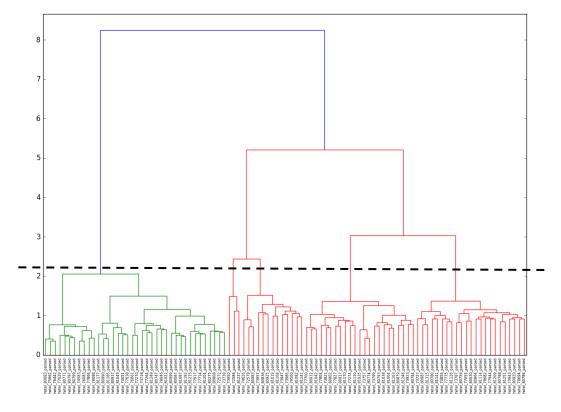
To view the charts, the user must respond with "y" to the question asked by the program. In this case, the chart will appear first. However, due to the large number of documents, it can be somewhat difficult to observe and read the names of the documents. In the chart below, we can see the same 6 documents related to cluster 1, which are shown in black.



The selection of colors is also random. After closing the above chart window, the Dendrogram related to this clustering will appear, which we can see in the figure below.

Unfortunately, even with zooming in on the figure, the names of the documents are not clearly visible due to the large number of leaves. Ultimately, this issue will be resolved when the user runs the program. It can be observed that, in general, the documents are divided into two clusters (at the highest level of clustering).

To achieve clustering with k=5, we need to cut the chart at the section indicated by the dashed line in the figure.



All the program codes along with their output are available in a separate file as plain text.