# Text Clustering Example

- This example shows text Clustering as an example of unsupervised machine learning task.
- The main steps are:
  - Explore dataset
  - Data Preparation
  - Feature Engineering
  - Model Training
  - Performance Assessment
- Clustering: grouping similar objects together based on their inherent attributes.
- Being unsupervised appraoch there is no need of prelabelled datasets.
- Examples: KMeans, PAM, DBSCAN, Spectral clustering, etc.
- The choice of the algorithm mainly depends on the problem domain and the algorithm property. For example, to use KMeans we should have pre-knowledge of the number of clusters to obtain. However, if we do not know the number of clusters already, then DBSCAN is better.

#### **Problem Definition**

- As discussed above, the problem in context is unsupervised learning problem.
- We will use BCC news datasets available from <a href="http://mlg.ucd.ie/files/datasets/bbc-fulltext.zip">http://mlg.ucd.ie/files/datasets/bbc-fulltext.zip</a>. A total of 2225 documents with five different news categories (i.e. business, entertainment, politics, sport, and tech) are available in the dataset.

```
import numpy as np
import pandas as pd
from sklearn.cluster import KMeans
from sklearn.feature_extraction.text import TfidfVectorizer
from sklearn.decomposition import PCA
import matplotlib.pyplot as plt
```

#### 1. Load and explore data

• The dataset contains 5 different categories in 5 different folders.

```
In [2]: from sklearn.datasets import load_files

# for reproducibility
random_state = 22

DATA_DIR = "./data/bbc/"
data = load_files(DATA_DIR, encoding="utf-8", decode_error="replace", random_state=random_state)
df = pd.DataFrame(list(zip(data['data'], data['target'])), columns=['text', 'label'])
```

It is necessary to do an exploratory data analysis in order to gain some insights from the data.

How is the data like?

```
In [3]: df.head()

Out[3]: text label

O Jones files Conte lawsuit\n\nMarion Jones has ... 3

1 English clubs make Euro history\n\nAll four of... 3

2 Tarantino 'to make Friday sequel'\n\nDirector ... 1

3 Lesotho textile workers lose jobs\n\nSix forei... 0

4 Celtic make late bid for Bellamy\n\nNewcastle ... 3

how many 'class label' present in the dataset??

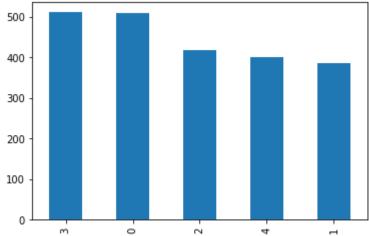
In [4]: df['label'].unique()
```

Is the dataset balanced??

Out[4]: array([3, 1, 0, 4, 2])

If the dataset is balanced then, the dataset contains an approximately equal portion of each class.

```
In [5]: df['label'].value_counts().plot(kind='bar')
Out[5]: <AxesSubplot:>
```



## 2. Feature Engineering

• we will use TFIDF approach.

tfidf

Out[7]:

- TfidfVectorizer will convert all the documents to a matrix of TF-IDF features.
- With Tfidfvectorizer we will calculate the word counts, IDF and TFIDF values in a go.

```
In [6]: tfidf_vectorizer = TfidfVectorizer(stop_words="english")
    tfidf_vectorizer.fit(df.text.values)
    features = tfidf_vectorizer.transform(df.text.values)
```

Lets check some values from the first document

```
In [7]: # get the first vector out (for the first document)
    first_vector_tfidfvectorizer=features[0]

# place tf-idf values in a pandas data frame
    feature_df = pd.DataFrame(first_vector_tfidfvectorizer.T.todense(), index=tfidf_vectorizer.get_feature_names
    feature_df.sort_values(by=["tfidf"],ascending=False).head()
```

```
      conte
      0.673159

      jones
      0.292039

      marion
      0.213606

      doping
      0.181978
```

### 3. Model Building

- We will check a version of KMeans algorithm, i.e. Mini-batch KMeans (MB KMeans).
- MB KMeans is similar to the standard KMeans. But in the MB KMeans the computationly expensive step is performed on a random sample only instead of the entire data.
- We will set the number of clusters = 5 because our dataset contains categories.
- But, if we do not have knowledge of the different labels then we may have to using domain knowledge and other ways to get the optimal numbers of clusters.

```
In [8]:
         mkmeans = MiniBatchKMeans(n clusters=5, random state=random state)
         mkmeans.fit(features)
Out[8]: MiniBatchKMeans(n clusters=5, random state=22)
        lets check the predictions useing labels attribute of the model.
         mkmeans.predict(features)
In [9]:
         print (mkmeans.labels )
         print ("\n")
         print(mkmeans.cluster_centers_)
        [4 2 4 ... 4 4 1]
         [[1.34473036e-04\ 1.09906331e-02\ 0.00000000e+00\ \dots\ 0.00000000e+00]
          0.00000000e+00 0.0000000e+00]
          [1.68041680e-04 \ 1.05754629e-02 \ 0.00000000e+00 \ \dots \ 0.00000000e+00
          0.00000000e+00 0.0000000e+00]
          [2.47895846e-04 2.12690116e-03 0.00000000e+00 ... 0.00000000e+00
```

```
4.60818743e-04 0.00000000e+001
          [0.000000000e+00\ 5.00195763e-03\ 0.00000000e+00\ \dots\ 0.00000000e+00
           0.00000000e+00 0.0000000e+001
          [4.67908112e-06 1.15377847e-02 4.45013653e-05 ... 4.09596975e-05
        Check KMeans
In [10]:
          kmeans = KMeans(n clusters=5, init='k-means++', max iter=100, n init=1)
          kmeans.fit(features)
          kmeans.predict(features)
          print (mkmeans.labels )
          print ("\n")
          print(mkmeans.cluster centers )
         [4 2 4 ... 4 4 1]
         [1.34473036e-04\ 1.09906331e-02\ 0.00000000e+00\ \dots\ 0.00000000e+00
           0.00000000e+00 0.00000000e+001
          [1.68041680e-04 1.05754629e-02 0.00000000e+00 ... 0.00000000e+00
           0.00000000e+00 0.0000000e+00]
          [2.47895846e-04 2.12690116e-03 0.00000000e+00 ... 0.00000000e+00
           4.60818743e-04 0.00000000e+00]
          [0.00000000e+00 5.00195763e-03 0.0000000e+00 ... 0.00000000e+00
           0.00000000e+00 0.0000000e+00]
          [4.67908112e-06 1.15377847e-02 4.45013653e-05 ... 4.09596975e-05
           0.00000000e+00 4.63992050e-04]]
In [11]: # lets order the centroids
          order centroids = kmeans.cluster centers .argsort()[:, ::-1]
          order centroids
Out[11]: array([[29039, 5711, 27997, ..., 18590, 18591,
                                                              01,
                 [22845, 12210, 16592, ..., 16826, 16827,
                 [17704, 22845, 15294, ..., 15629, 15628, 14562],
                 [11464, 9609, 22845, ..., 16998, 16999, 14562],
                 [10650, 3844, 3182, ..., 13381, 13382, 14562]])
In [12]: # lets check the features
          terms = tfidf_vectorizer.get feature names()
          terms[23361]
Out[12]: 'seed'
```

```
In [13]: # lets see the centroids into which clusters they belongs
          for i in range(5):
              print("\nCluster %d:" % i)
              for ind in order_centroids[i, :10]:
                  print('%s' % terms[ind])
         Cluster 0:
         yukos
         china
         virus
         oil
         microsoft
         russian
         spyware
         gazprom
         security
         windows
         Cluster 1:
         said
         growth
         market
         year
         economy
         sales
         bank
         company
         prices
         economic
         Cluster 2:
         mr
         said
         labour
         people
         election
         blair
         government
         party
         brown
         minister
         Cluster 3:
         game
```

```
england
said
win
cup
club
match
team
players
play
Cluster 4:
film
best
awards
award
music
band
year
said
films
```

lets do the prediction and check the results

### 4. Visualization

- Lets plot the features in a 2D space.
- The dimension of features got from TfldfVectorizer is large ( > 10,000)so we will do the dimensionality reduction using Principal Component Analysis.
- Here, PCA will transform the high dimensional features into 2 dimensions.

```
In [14]: def plot_clusters(reduced_features, reduced_cluster_centers, cls):
    plt.scatter(reduced_features[:,0], reduced_features[:,1], c=cls.predict(features))
    plt.scatter(reduced_cluster_centers[:, 0], reduced_cluster_centers[:,1], marker='x', s=150, c='b')

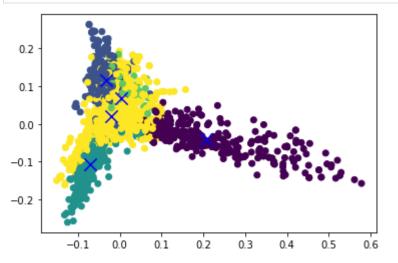
# reduce the features to 2D
pca = PCA(n_components=2, random_state=random_state)
reduced_features = pca.fit_transform(features.toarray())

# reduce the cluster centers to 2D
reduced_cluster_centers_mk = pca.transform(mkmeans.cluster_centers_)
reduced_cluster_centers_k = pca.transform(kmeans.cluster_centers_)
```

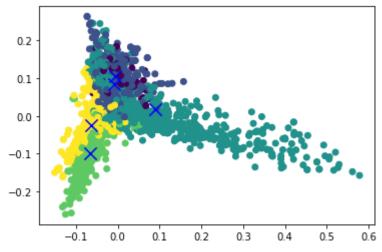
lets visualize using scatter plot.

- In the plot:
  - X = first dimension
  - Y = second dimension
- Different colors represent the cluster members.

```
In [15]: # MK Means
plot_clusters(reduced_features, reduced_cluster_centers_mk,mkmeans )
```



```
In [16]: # K Means
plot_clusters(reduced_features, reduced_cluster_centers_k, kmeans)
```



#### 5. Evaluation

• Let us check the predictions.

• The prediction label is [2] i.e. cluster 2. It is related with technology and also the examined text is also technology related. This prediction is correct.

• Further, we ahve to test the classifier with other text as well.

```
In [18]: print("Prediction using Mini Batch KMeans")
X = tfidf_vectorizer.transform(["The new social network is popular in cell phones"])
predicted = mkmeans.predict(X)
print(predicted)
```

```
Prediction using Mini Batch KMeans
[1]

Is the prediction accurate??

In [ ]:
```

#### Other formal methods for evaluation:

- Since we have a labelled dataset is easier than the dataset without labelled dataset.
- homogeneity\_score: values ranges from 0 and 1 where 1 indicates ideally homogeneous labeling.

```
In [19]: from sklearn.metrics import homogeneity_score
    hs_mk = homogeneity_score(df.label, mkmeans.predict(features))
    hs_k = homogeneity_score(df.label, kmeans.predict(features))

print("Homogenity Mini KMeans: %f"%hs_mk)
print("Homogenity KMeans: %f"%hs_k)
```

Homogenity Mini KMeans: 0.573055 Homogenity KMeans: 0.619430

- In absence of labelled dataset, Cluster evaluation can be done using other metrics like Silhouette Coefficient (SC).
- SC uses mean intra-cluster distance (a) and the mean nearest-cluster distance (b) for each sample.
- Here, b is the distance between a sample and the nearest cluster that the sample is not a part of.
- The SC for a sample is (b a) / max(a,b).
  - best value = 1
  - worst value = -1

silhouette KMeans: 0.009264

- near 0 values means overlapping clusters.
- Negative values means bad clustering.

```
In [20]: from sklearn.metrics import silhouette_score
    ss_mk = silhouette_score(features, labels=mkmeans.predict(features))
    ss_k = silhouette_score(features, labels=kmeans.predict(features))

print("silhouette Mini KMeans: %f"%ss_mk)
print("silhouette KMeans: %f"%ss_k)

silhouette Mini KMeans: 0.009695
```

## References:

- https://sanjayasubedi.com.np/nlp/nlp-with-python-document-clustering/
- https://www.kaggle.com/jbencina/clustering-documents-with-tfidf-and-kmeans
- http://kavita-ganesan.com/extracting-keywords-from-text-tfidf/
- https://towardsdatascience.com/applying-machine-learning-to-classify-an-unsupervised-text-document-e7bb6265f52

In [ ]:
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