

TEXT CLUSTERING

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Introduction

A text cluster is formed by grouping a number of unlabeled texts together in such a way that texts in one cluster are more likely to be more similar than those in other clusters. so, the assignment goes this way we take five different samples of Gutenberg digital books, which are all of five different genres and of five different authors, that are semantically different. Secondly, preprocess and clean the data and create random samples of 200 documents from each book. Thirdly, applying different transformers such as BOW and TF-IDF also uses other features LDA and Word-Embedding. Fourthly, use clustering algorithms such as K-means, EM, and Hierarchical then Calculate Kappa against true authors, Consistency, Coherence, and Silhouette for each model. Finally, Compare and decide which clustering result is the closest to the human labels

The Goal

The overall aim is to produce similar clusters and compare them; analyze the pros and cons of algorithms, and Compare and decide which clustering result is the closest to the human labels

Methodology

1. Selecting the Books

We picked five books with different authors and different genres from the Gutenberg Digital Library.











2. Preprocessing and Data Cleansing

• Import essential libraries

Before starting we imported libraries for many reasons that help us to read, clean, model, evaluate the data, and visualization. we use stopword, pandas, Word2Vec, Dictionary, plt, Svm, and WordNetLemmatizer, and many of them are in the code.

Read the data

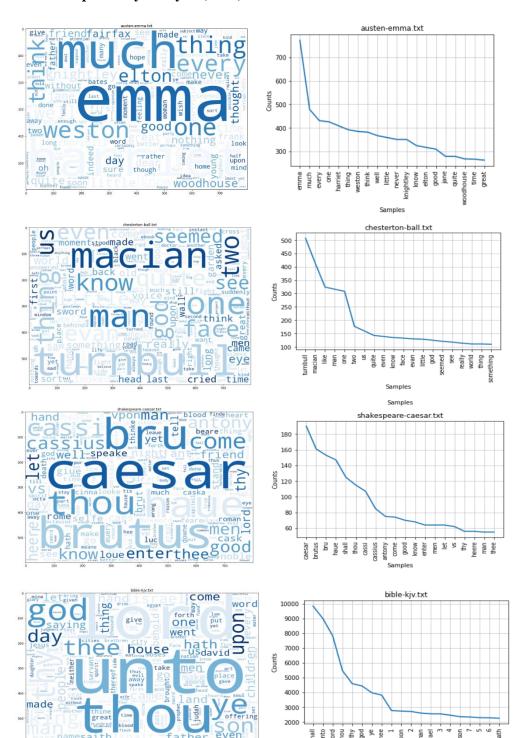
In this stage we read the list of books from nltk.corpus.gerenberg.words, the function takes the list of books and returns a list of book words using nltk. Corpus.

```
# list of books of the same geners (fiction book)
list_of_books=["austen-emma.txt", 'milton-paradise.txt', 'bible-kjv.txt', 'chesterton-ball.txt', 'shakespeare-caesar.txt']
# list of authors
list_of_authors=["Jane Austen", "JOHN MILTON", "King James Version", "G.K. Chesterton ", "Maria Edgeworth"]
```

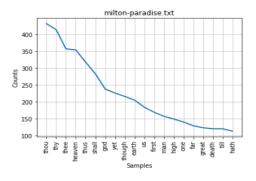
Clean the Data

It is a very important step to clean the raw text with different methods and make sure that is not contain anything that makes mis cluster or noise. We started by removing punctuations and any not needed numbers by using RegexpTokenizer from the nltk library and removing stop words, which are the noise in the text. After that, we create random samples of 200 partitions from each book, we prepared the records of 150 words records for each document. Then we performed labeling and indexing and created a Data frame. Finally, we normalize text to words by using Lemmatisation which reduces words to their word root.

• Exploratory Analysis (EDA)







We clearly can see different word distribution in the 5 book plots...

3. Transformation

We have to convert that text into some numbers or number vectors. And we applied four types of transformations (BOW, TF-IDF, LDA, and Word-Embedding).

3.1.BOW

Bag of words is a Natural Language Processing technique of text modelling. In technical terms, we can say that it is a method of feature extraction with text data. This approach is a simple and flexible way of extracting features from documents. A bag of words is a representation of text that describes the occurrence of words within a document.



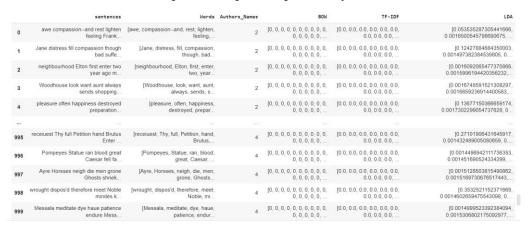
3.2.TF-IDF

Term Frequency: In document d, the frequency represents the number of instances of a given word t. Therefore, we can see that it becomes more relevant when a word appears in the text, which is rational. Since the ordering of terms is not significant, we can use a vector to describe the text in the bag of term models. For each specific term in the paper, there is an entry with the value being the term frequency.



3.3 Latent Dirichlet Allocation (LDA)

Latent Dirichlet allocation is one of the most popular methods for performing topic modeling. Each document consists of various words and each topic can be associated with some words. The aim behind the LDA is to find topics that the document belongs to, on the basis of words contains in it. It assumes that documents with similar topics will use a similar group of words. This enables the documents to map the probability distribution over latent topics and topics are probability distribution.



3.4 Doc2Vec

Doc2Vec is an unsupervised algorithm that learns embeddings from variable-length pieces of texts, such as sentences, paragraphs, and documents.



4. Modeling (Clustering algorithms)

We applied 3 unsupervised clustering algorithms:

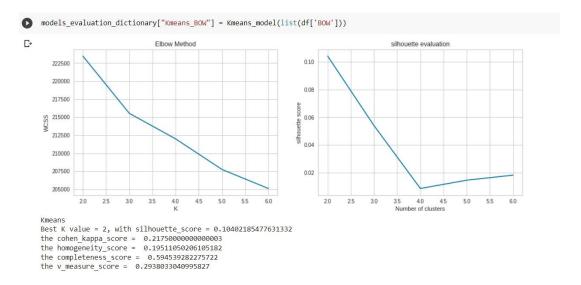
- K-Means
- EM
- Hierarchical

4.1 K-Mean

Finding groups in the data is the objective of the K-Means algorithm, with the variable K indicating how many groups there are in total. The algorithm assigns via an iterative process. Based on the above features, each point of data belongs to one of the K groups. Data Based on feature similarity, points are grouped. We employ the El-Bow Method to optimize the number of clusters while implementing the K-Means Model with one of each transformation approach, even though we only require 5 clusters

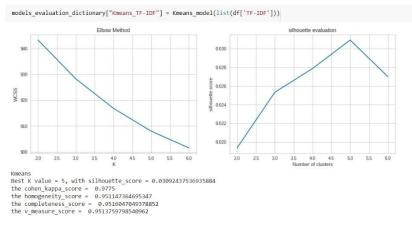
a) K-Mean with BOW

Here we applied the k-mean to the BOW transformer, from using the Elbow method the best number of clusters is 5



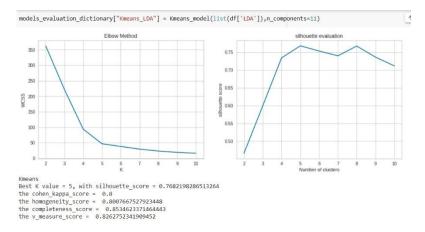
b) K-Mean with TF-IDF

We applied K-mean with TF-IDF, we get the best K is 5 clusters and the silhouette score are 0.03.



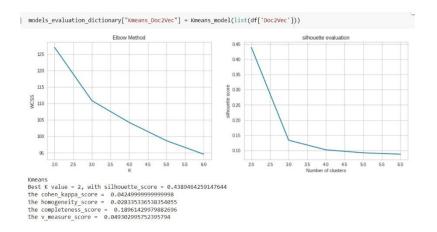
c) K-Mean with LDA

We applied K-mean with LDA, we get the best K is 5 clusters and the silhouette score are 0.76



d) K-Mean with Doc2Vic

We applied K-mean with LDA, we get the best K is 2 clusters and the silhouette score are 0.43



4.2 EM

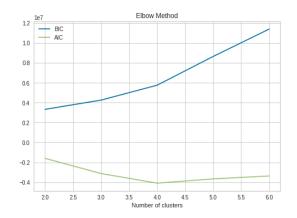
Similar to k-means but instead of using k points, we use k Gaussian mixture distributions. The model then tries to find the best parameters' values for the Gaussian that fit the data well.

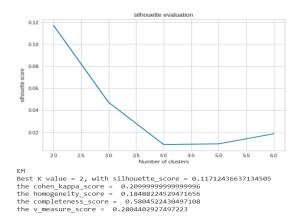
Elbow method measures

AIC stands for Akaike information criterion for the current model on the input X. BIC stands for Bayesian information criterion for the current model on the input X.

a) EM with BOW

The shape of the input data for the model is (1000, 12315) and it takes too much time and memory storage for training and evaluation so we used the PCA dimensionality reduction technique to reduce the number of features from 12315 to 1000.



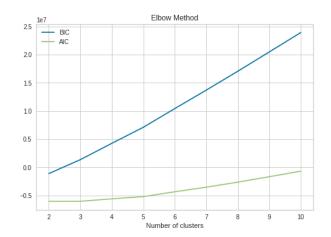


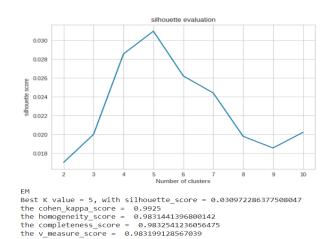
The model has the highest silhouette value at k=2 which is far from reality. From the Elbow method the AIC state that the best value is 4 but on the other hand it is the worst value when it comes to the silhouette. Also, the Kappa the other measures scores are very small near 0.

b) EM with TF-IDF

We encountered the same problem as with BOW so we used the PCA dimensionality reduction technique to reduce the number of features from 12315 to 1000.

The Elbow method with BIC and AIC doesn't show anything the BIC is increasing all the time and there's no elbow here. There may be a slight elbow for AIC at a number of clusters = 3

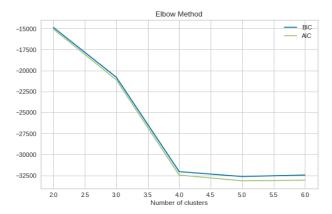


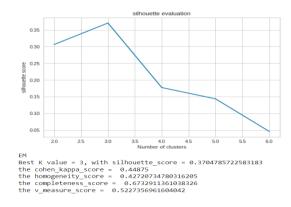


It is totally clear from the silhouette evaluation that the best number of clusters is 5 and it gives very good values of the Kappa = 0.99 meaning that the predicted clustering labels are almost equal to the actual labels, homogeneity, completeness, and V-measure scores which are so close to 1 so the clusters are well separated.

c) EM with LDA

We hadn't any problem with the computational power here as the number of features is really small (5). From the Elbow method's perspective, the best value for the number of clusters = 4 using both AIC and BIC measures.

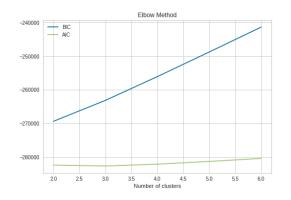


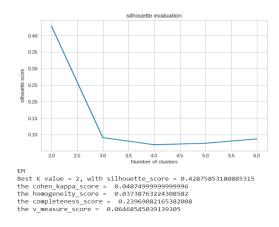


Here it is different, using the silhouette the best number of clusters is 3. In that case, the Kappa score and the other measures are on average near 0.5.

d) EM with Doc2Vec:

The Elbow method with BIC and AIC doesn't show anything the BIC is increasing all the time and there's no elbow here. There may be a slight elbow for AIC when the number of clusters = 3





The best number of clusters = 2 measured from the highest silhouette values. These are very bad scores the Kappa score = 0.04 which is a very small value meaning that the predicted labels are so far from the actual labels. Also, the other measures' values are very small meaning that the model doesn't separate the clusters well

4.3 **Hierarchical clustering:** is a general family of clustering algorithms that build nested clusters by merging or splitting them successively. This hierarchy of clusters is represented as a tree (or dendrogram). The root of the tree is the unique cluster that gathers all the samples, the leaves being the clusters with only one sample.

1. Hierarchical clustering with BOW transformation:

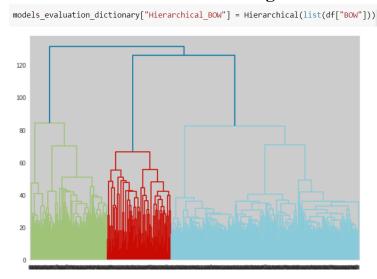


Figure 2 Hierarchical Cluster with BOW

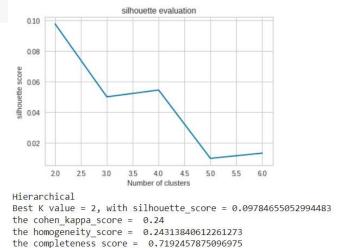


Figure 1 Hierarchical Cluster Evaluation with BOW

2. Hierarchical clustering with TFIDF transformation:

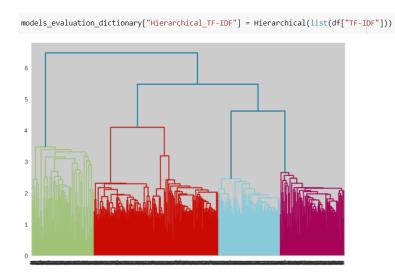
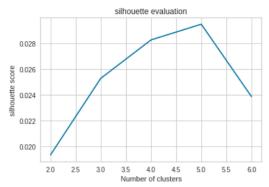


Figure 4:Hierarchical Clustering with TFIDF Transformer



the v_measure_score = 0.36342299788918747

Hierarchical
Best K value = 5, with silhouette_score = 0.029476961617526044
the cohen_kappa_score = 0.96125
the homogeneity_score = 0.9228519183948832
the completeness_score = 0.9233909187014164
the v_measure_score = 0.9231213398690822

Figure 3: Hierarchical Clustering Evaluation TFIDF Transformation

3. Hierarchical clustering with LDA transformation:

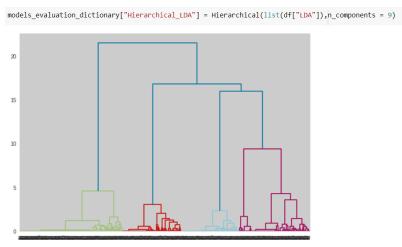
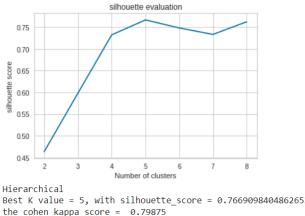


Figure 5: Hierarchical Clustering with LDA Transformation



Hierarchical
Best K value = 5, with silhouette_score = 0.7669098404862652
the cohen_kappa_score = 0.79875
the homogeneity_score = 0.8245917470247597
the completeness_score = 0.9035198851541059
the v_measure_score = 0.8622533714809469

Figure 6: Hierarchical Clustering Evaluation with LDA Transformation

4. Hierarchical clustering with Doc2Vec transformation:

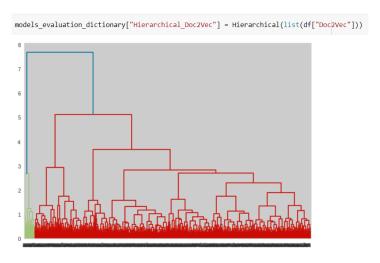
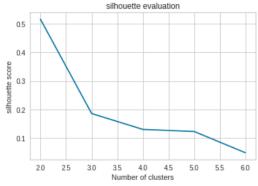


Figure 7: Hierarchical Clustering with Doc2Vec Transformation



Number of clusters

Hierarchical

Best K value = 2, with silhouette_score = 0.5166573524475098

the cohen_kappa_score = 0.027499999999997

the homogeneity_score = 0.01664093466641861

the completeness_score = 0.16919938937369638

the v_measure_score = 0.03030166890537523

Figure 8: Hierarchical Clustering Evaluation with Doc2Vec

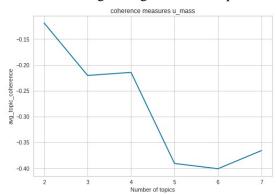
5. Evaluation

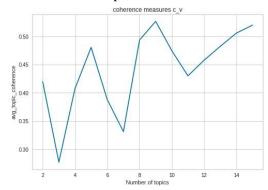
We try 6 different measurements from each model to each transformer:

- Silhouette
- Cohen's Kappa
- Coherence
- Homogeneity score
- Completeness score
- V measure score

Coherence

It measures how the words for each topic are related to that topic and related to each other. We used it to get insights and to help us select the best number of topics.





• Silhouette score

Measures how each cluster's points are close to each other and separated from the other clusters. The best value is 1 and the worst value is -1. Values near 0 indicate overlapping clusters. Negative values generally indicate that a sample has been assigned to the wrong cluster, as a different cluster is more similar.

Cohen kappa score

Measures the similarity between the true labels and the predicted labels.

A kappa statistic is a number between -1 and 1. The maximum value means complete agreement; zero or lower means chance agreement.

Homogeneity score:

A clustering result satisfies homogeneity if all of its clusters contain only data points that are members of a single class.

The result is between 0 and 1. Score = 1 for perfectly homogeneous labeling.

• Completeness score:

A clustering result satisfies completeness if all the data points that are members of a given class are elements of the same cluster.

The result is between 0 and 1. Score = 1 for perfectly complete labeling.

• V measure score

Measures the harmonic mean between homogeneity and completeness. The result is between 0 and 1. Score = 1 for perfectly complete labeling.

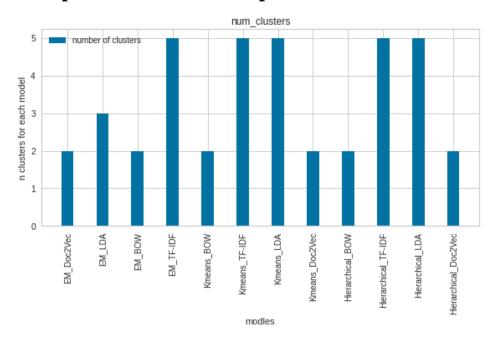
In general, we select the best number of clusters for each model based on the highest **silhouette score**

	Silhouette	Cohen's Kappa	Homogeneity score	Completeness score	V measure score
K-Mean with BOW	0.10	0.21	0.19	0.59	0.29
K-Mean with TF-IDF	0.03	0.97	0.95	0.95	0.95
K-Mean with LDA	0.76	0.80	0.80	0.85	0.85
K-Mean with Doc2Vec	0.43	0.04	0.02	0.18	0.04

	Silhouette	Cohen's Kappa	Homogeneity score	Completeness score	V measure score
EM with					
BOW	0.11	0.20	0.18	0.58	0.28
EM with					
TF-IDF	0.03	0.99	0.98	0.98	0.98
EM with				0.67	
LDA	0.37	0.44	0.42		0.52
EM with					
Doc2Vec	0.42	0.04	0.03	0.23	0.06

	Silhouette	Cohen's Kappa	Homogeneity score	Completeness score	V measure score
Hierarchical with BOW	0.097	0.24	0.243	0.71	0.36
Hierarchical with TF-IDF	0.029	0.96	0.92	0.92	0.92
Hierarchical with LDA	0.76	0.79	0.82	0.90	0.86
Hierarchical with Doc2Vec	0.51	0.027	0.016	0.169	0.03

6. Comparison and the champion Model



From the human perspective, we know we know that the best number of clusters is 5 which equals the number of books. We eliminated all the models other than that. Now we have 5 models to compare (EM-TFI-DF, Kmeans_TF-IDF, Kmeans_LDA, Hirarchical_TF-IDF, Hirarchical_LDA).

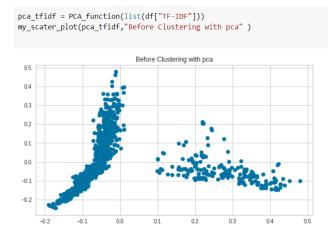


Figure 10 Before Clustering with PCA



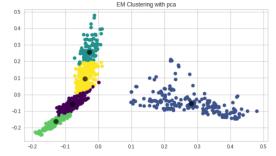
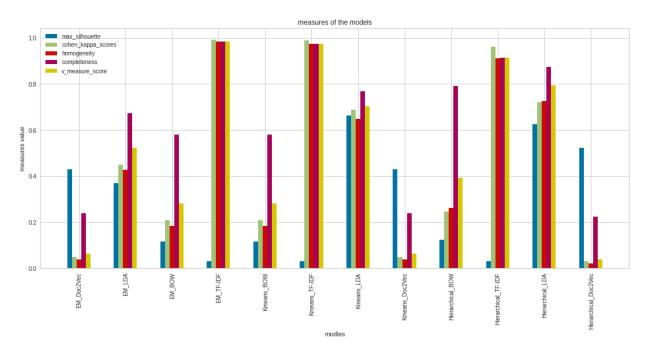


Figure 9 EM Clustering with PCA

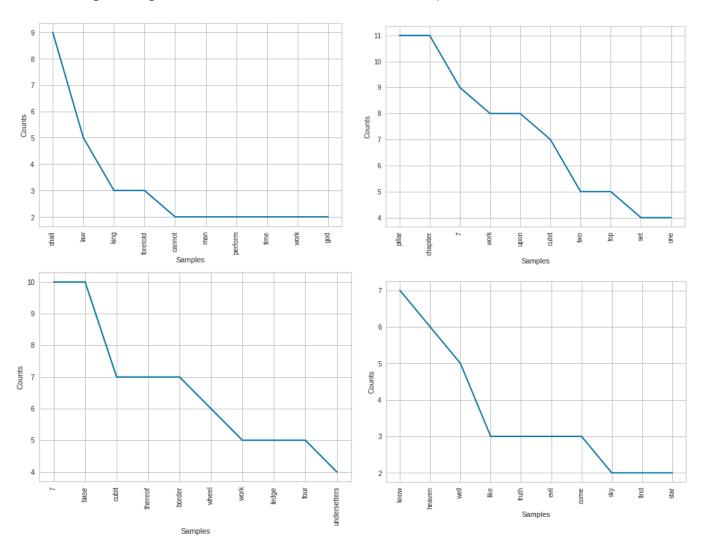
Now we move to our measures:



It is obvious now that the champion model is EM with TF-IDF (EM_TFI-DF). Although for that model we've applied dimensionality reduction with PCA to reduce the number of features from 12315 to only 1000. It has the highest Kappa score meaning that the predicted clustering labels are almost equal to the actual labels, also homogeneity, completeness, and V-measure scores are so close to 1 meaning that the clusters are well separated and with no overlapping.

7. Error Analysis

For our champion, we tracked all the records that were predicted in a wrong cluster with respect to the actual true clustering labels. We found only 4 records then, and we tried to find the 10 most frequent words in those records. Those words are the main cause of the wrong labeling and if we removed them the model would perform better.



We noticed that some of these words are just numbers written in words like (one, two, and four), others have a high frequency in more than one record like (7, and work) so these words should be filtered in the preprocessing step.

Also if we used the original set of features without dimensionality reduction the model would perform better. Although, it will require more computational power (memory storage particularly).

References:

https://scikit-learn.org/stable/modules/generated/sklearn.metrics.silhouette_score.html
https://scikit-learn.org/stable/modules/generated/sklearn.metrics.cohen_kappa_score.html
https://scikit-learn.org/stable/modules/generated/sklearn.metrics.homogeneity_score.html
https://scikit-learn.org/stable/modules/generated/sklearn.metrics.completeness_score.html
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