Atelier 4: Clustering

L'objectif de cet atelier est de detecter le plagiarisme dans une collection de documents texte en regroupant dans les mêmes clusters les documents texte qui était recopier à partir du même document original. On va reprendre le même dataset des ateliers 1 et 2.

1. Preparation du DataSet

Écrire le code permettant de récupérer le dataset du plagiarisme traité dans les ateliers précédents et appliquer les prétraitements nécessaires.

```
import alob
from nltk.stem import WordNetLemmatizer
from nltk.tokenize import word tokenize
from nltk.corpus import stopwords
import string, re
import pandas as pd
import numpy as np
path = 'c:/Users/dscon/Documents/COURS
UM6P/S3/TEXT-MINING/final df.csv'
data = pd.read csv(path)
df_content = data[['User', 'Task', 'UResponse', 'OResponse']]
stop words = set(stopwords.words('english'))
def get_corpus(text):
    '''Uttiliser la librairie glob pour recuperer les documents dans
    liste de listes dont les sous listes representent le contenu des
documents
    du corpus'''
    '''Appliquer la segmentation, le cleaning et la lemmatisation'''
    text = text.lower()
    text = text.translate(str.maketrans('', '', string.punctuation))
    text = re.sub(r'\d+', '', text)
    words = word tokenize(text)
    words = [word for word in words if word not in stop words]
    return ' '.join(words)
df content['ProcessedUserContent'] =
df content['UResponse'].apply(get corpus)
df content['ProcessedOriginalContent'] =
df content['OResponse'].apply(get corpus)
```

```
C:\Users\dscon\AppData\Local\Temp\ipykernel 9352\42808191.py:29:
SettingWithCopyWarning:
A value is trying to be set on a copy of a slice from a DataFrame.
Try using .loc[row indexer,col indexer] = value instead
See the caveats in the documentation:
https://pandas.pydata.org/pandas-docs/stable/user guide/indexing.html#
returning-a-view-versus-a-copy
  df content['ProcessedUserContent'] =
df content['UResponse'].apply(get corpus)
C:\Users\dscon\AppData\Local\Temp\ipykernel 9352\42808191.py:30:
SettingWithCopyWarning:
A value is trying to be set on a copy of a slice from a DataFrame.
Try using .loc[row indexer,col indexer] = value instead
See the caveats in the documentation:
https://pandas.pydata.org/pandas-docs/stable/user guide/indexing.html#
returning-a-view-versus-a-copy
  df content['ProcessedOriginalContent'] =
df content['OResponse'].apply(get corpus)
df content.head()
   User Task
                                                      UResponse \
             Inheritance is a basic concept of Object-Orien...
  Aq0p
             Inheritance is a basic concept in object orien...
1
  g0pB
2
  g0pC
             inheritance in object oriented programming is ...
3
           a Inheritance in object oriented programming is ...
  q0pD
4 g0pE
          a In object-oriented programming, inheritance is...
                                           OResponse \
  In object-oriented programming, inheritance is...
1
  In object-oriented programming, inheritance is...
2
  In object-oriented programming, inheritance is...
  In object-oriented programming, inheritance is...
  In object-oriented programming, inheritance is...
                                ProcessedUserContent \
  inheritance basic concept objectoriented progr...
  inheritance basic concept object oriented prog...
1
  inheritance object oriented programming new cl...
  inheritance object oriented programming way fo...
4 objectoriented programming inheritance way for...
                            ProcessedOriginalContent
  objectoriented programming inheritance way for...
  objectoriented programming inheritance way for...
1
  objectoriented programming inheritance way for...
  objectoriented programming inheritance way for...
  objectoriented programming inheritance way for...
```

2. Vectorisation

2.1 TF-IDF

Définir le traitement permettant de récupérer la représentation vectorielle TFIDF des différents documents du corpus.

```
from sklearn.feature_extraction.text import TfidfVectorizer

def vectorsTFIDF(corpus_lemetized):
    vectorizer = TfidfVectorizer()
    tfidf_matrix = vectorizer.fit_transform(corpus_lemetized)
    return tfidf_matrix, vectorizer.get_feature_names_out()
```

2.2 SVD

Définir le traitement permettant de récupérer la représentation vectorielle des différents documents du corpus en se basant sur l'approche SVD.

```
import numpy as np
from sklearn.decomposition import TruncatedSVD
from sklearn.preprocessing import Normalizer
from sklearn.pipeline import make pipeline
from sklearn.decomposition import TruncatedSVD
from sklearn.preprocessing import Normalizer
from sklearn.pipeline import make pipeline
def build cooccurrence matrix(corpus, window size=2):
    tokeniser corpus = [doc.split() for doc in corpus]
    vocab = list(word for doc in tokeniser corpus for word in doc)
    vocab index = {word : i for word, i in enumerate(vocab)}
    vocab size = len(vocab)
    matrix cooccurence = np.zeros((vocab size, vocab size),
dtype=np.float32)
    for doc in tokeniser corpus:
        for idx, word in enumerate(doc):
            if word in vocab index:
                word idx = vocab index[word]
                start = max(0, idx - window size)
                end = \min(len(word), idx + \overline{window size+1})
                for neighbor idx in range(start, end):
                    neighbor word = doc[neighbor idx]
                    if neighbor word in vocab_index:
                        neighbor word idx = vocab index[neighbor word]
```

```
matrix cooccurence[word idx,
neighbor word idx] += 1
    return matrix cooccurence, vocab index
def docVectSVD(corpus lemetized, n components=100, window size=2):
    ''' recuperer le vocabulaire du corpus'''
    ''' creer la matrice des cooccurrences des termes'''
    ''' Appliquer la reduction SVD'''
    ''' genenrer le dictionaire terme/vecteur'''
    ''' Definir les traitement permettant de recuperer la
representation
        vectorielle des documents du corpus en se basant sur l'apprche
SVD
        (la moyenne des vecteurs composant chaque document).'''
    cooccurrence matrix, vocab to index =
build cooccurrence matrix(corpus lemetized, window size)
    svd = TruncatedSVD(n components=n components)
    term vectors = svd.fit transform(cooccurrence matrix)
    index to vocab = {idx:word for word, idx in
vocab to index.items()}
    # term vector dict = {index to vocab[idx]: term vectors[idx] for
idx in range(len(index to vocab))}
    term vector dict = {index to vocab.get(idx, f"missing {idx}"):
term vectors[idx] for idx in range(len(term vectors))}
    doc vectors = []
    for doc in corpus lemetized:
        words = doc.split()
        word vectors = [term vector dict[word] for word in words if
word in term vector dict]
        if word vectors:
            doc vector = np.mean(word_vectors, axis=0)
        else:
            doc vector = np.zeros(n components)
        doc vectors.append(doc vector)
    return term vector dict, doc vectors
```

2.3 Word2Vec

Definir le traitement permettant de récupérer la representation vectorielle des differents documents du corpus en se basant sur l'approche word2vec.

Tester avec le pretrained glove-wiki-gigaword-50 et avec un apprentissage à partir du corpus.

```
from gensim.downloader import load
def pretrainedDocWV(corpus lemetized):
    glove model = load('glove-wiki-gigaword-50')
    doc_vectors = []
    for doc in corpus lemetized:
        # Tokeniser le document
        words = doc.split()
        word vectors = [glove model[word] for word in words if word in
glove model]
        if word vectors:
            doc vector = np.mean(word vectors, axis=0)
            doc vectors.append(doc vector)
        else:
            doc vectors.append(np.zeros(glove model.vector size))
    return np.array(doc vectors)
from gensim.models import Word2Vec
def propreDocWV(corpus lemmatized, vector size=50, epochs=10):
    ''' Récupérer la représentation vectorielle des documents du
corpus
       en entraînant un modèle Word2Vec sur notre propre corpus.
    1.1.1
    # Tokeniser le corpus pour Word2Vec
    tokenized corpus = [doc.split() for doc in corpus lemmatized]
    # Créer et entraîner le modèle Word2Vec
    model = Word2Vec(sentences=tokenized corpus,
vector size=vector size, window=5, min count=1, workers=4)
    # Initialisation de la liste pour stocker les vecteurs des
documents
    doc vectors = []
    for doc in tokenized corpus:
        # Récupérer les vecteurs pour chaque mot, ignorer les mots
absents du vocabulaire
        word vectors = [model.wv[word] for word in doc if word in
model.wvl
```

```
if word_vectors:
    # Calculer la moyenne des vecteurs des mots pour obtenir
le vecteur du document
    doc_vector = np.mean(word_vectors, axis=0)
    doc_vectors.append(doc_vector)
    else:
        # En cas d'absence de mots valides, on peut ajouter un
vecteur nul ou le gérer autrement
        doc_vectors.append(np.zeros(vector_size))
return np.array(doc_vectors)
```

2.4 Doc2Vec

Définir le traitement permettant de récupérer la representation vectorielle doc2vec des différents documents du corpus

```
from gensim.models.doc2vec import Doc2Vec, TaggedDocument
def vectorsDoc2Vec(corpus lemetized):
    ''' Récupérer la représentation vectorielle des documents du
corpus en utilisant Doc2Vec '''
    # Préparation les données pour Doc2Vec
    tagged data = [TaggedDocument(words=doc.split(), tags=[str(i)])
for i, doc in enumerate(corpus lemetized)]
    # modèle Doc2Vec
    model = Doc2Vec(vector size=50, min count=2, epochs=40)
    # vocabulaire
    model.build vocab(tagged data)
    # Entrainement du modèle
    model.train(tagged data, total examples=model.corpus count,
epochs=model.epochs)
    # vecteurs de documents
    document vectors = [model.infer vector(doc.words) for doc in
tagged data]
    return document vectors
doc vectors = vectorsDoc2Vec(df content['ProcessedUserContent'])
doc vectors
```

```
[array([-2.343715 , -0.9948454 , -2.5225084 , -0.15528658, 1.7676123]
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0.50456285.
```

```
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```

```
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```

```
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```
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```

```
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        -1.285949 , 0.90032095, 1.7048161 , -0.81052756, 0.8475076
        0.15654308, -0.4568725, 0.27824828, -1.1970198, 1.0450124
        0.4968546 , 1.2717084 , 0.56202954 , 1.2864543 , -1.3672763
        0.8171574 , 0.85772043, 1.1604904 , 0.66688347, -0.8432483
        2.514248 , -0.28266758, -1.0973277 , 0.31491902, 2.1787488
        -0.28987047, 0.24312526, -2.0917003 , 1.9592978 , 0.8550908
],
      dtvpe=float32),
array([ 1.1912045 , -0.34570098, -0.25651047, 0.70430315,
0.57890016,
        -0.46226636, 0.10987736, 1.2249973, -1.175947, 0.2287073
        0.65393496, 0.4651022, 0.0518361, 0.15478814, -1.4077591
        0.6351686 , -1.4742026 , 0.4894086 , -1.2268775 , -0.0022603
        -1.1458106 , 1.0612313 , 1.0004092 , -0.4600927 ,
0.49444303,
        -0.06273472, -0.7924119 , -1.0166361 , -0.26313242, 0.7834526
        0.53286314, 1.1753328, -1.0009428, 1.5952307, -
0.34678784,
        0.22882892, 0.27295727, 0.80460215, -0.21784651, -1.6407375
        1.9385422 , 0.38117686, -0.18998569, 0.01863312, 1.2542734
```

```
0.41197807, 0.45282733, -1.0811417, 0.8924423,
0.211401151,
      dtype=float32),
array([ 1.0550765 , 0.4016326 , -0.30028322, 0.5347264 , -1.1303015
        -1.0288426 , -0.519207 , 1.0358164 , -0.54820114,
0.58313096,
        0.35589936, 0.26630768, 0.99990636, 1.0799222, -
0.78849274,
        1.1046107 , 0.04086257, 0.14473897, -0.48449004, 0.506392
        -1.1240592 , -0.14958838, 0.69233704, -0.699772 ,
0.75741994,
        0.94913304, -0.36347622, 0.16773498, -0.7488279, 1.2519944
        0.23210037, 0.29153818, 0.22593394, -0.02435908, -1.418425
        0.51458573, 0.8857249, -0.10821696, 0.5814505, -1.0855784
        0.92153007, 0.01124167, -0.9127518, 0.7143649,
0.91121846,
        -0.49709 , 0.46249372, -1.2843794 , 0.48400363, -
0.10517471],
      dtype=float32)]
```

3. Clustering avec K-means

3.1 k-means avec TFIDF

Ci-dessous un exemple d'application du k-means en se basant sur le representation TFIDF.

```
#K-MEANS CLUSTERING

#fit(X[, y, sample_weight]) : Compute k-means clustering.
#predict(X[, sample_weight]) : Predict the closest cluster
each sample in X belongs to.
#fit_predict(X[, y, sample_weight]) :Compute cluster centers and
predict cluster index for each sample.

#fit(X[, y, sample_weight]) : Compute k-means clustering.
#transform(X) :Transform X to a cluster-
distance space.
#fit_transform(X[, y, sample_weight]):Compute clustering and transform
X to cluster-distance space.

#score(X[, y, sample_weight]) : Opposite of the value of X on
```

```
the K-means objective.
from sklearn.decomposition import PCA
from sklearn.cluster import KMeans
import numpy as np
#initialisation automatique
#km = KMeans(n clusters=5, init='k-means++', n init=50)
#predetermination des centroids
centroids=np.array([[-0.10512687, -0.35341345],[-0.24220548,
[0.32397296], [-0.03986034, [0.09208828], [0.72163033, [0.16627089],
[0.72163033, 0.16627089]])
km = KMeans(n_clusters=5, init=centroids,n init=50)
print("Clustering sparse data with %s" % km)
# vect=vectorsTFIDF(get_corpus('/Users/rachad/Desktop/corpus/*'))
vects, feature names = vectorsTFIDF([" ".join(doc.split()) for doc in
df_content['ProcessedUserContent']])
pca = PCA(n components=2)
reduced vects = pca.fit transform(vects.toarray())
km.fit(reduced vects)
y means = km.predict(reduced vects)
print("Cluster assignments:", y_means)
Clustering sparse data with KMeans(init=array([[-0.10512687, -
0.353413451,
      [-0.24220548,
                   0.323972961,
      [-0.03986034, 0.09208828],
      [ 0.72163033, 0.16627089],
      [ 0.72163033, 0.16627089]]),
      n clusters=5, n init=50)
c:\Users\dscon\anaconda3\Lib\site-packages\sklearn\cluster\
_kmeans.py:1414: RuntimeWarning: Explicit initial center position
passed: performing only one init in KMeans instead of n init=50.
 super(). check params vs input(X, default n init=10)
2 2 2 2 2 2 2 2 2 2 2 2 2 2
c:\Users\dscon\anaconda3\Lib\site-packages\sklearn\cluster\
kmeans.py:1429: UserWarning: KMeans is known to have a memory leak on
Windows with MKL, when there are less chunks than available threads.
```

```
You can avoid it by setting the environment variable OMP_NUM_THREADS=1. warnings.warn(
```

Affichage du cluster de chaque document

```
dict(zip(np.arange(0,100),y_means[:10]))
{0: 2, 1: 2, 2: 1, 3: 1, 4: 1, 5: 1, 6: 2, 7: 1, 8: 2, 9: 2}
```

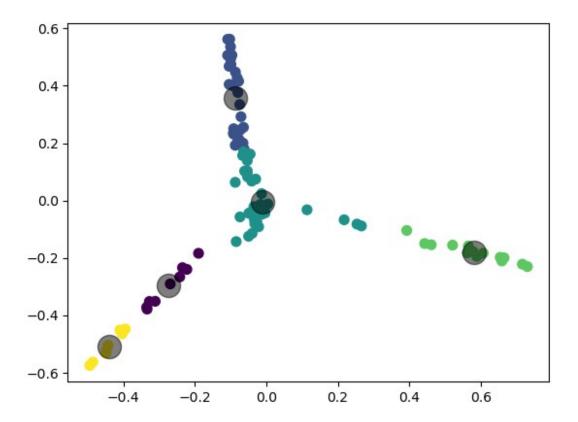
Le score du modele

```
#score(X[, y, sample_weight]): Opposite of the value of X on the K-
means objective.

km.score(reduced_vects)
-1.2091161826384558
```

Visialisation des clusters

```
from matplotlib import pyplot as plt
plt.scatter(reduced_vects[:,0], reduced_vects[:,1], c=y_means, s=50,
cmap='viridis')
centers = km.cluster_centers_
plt.scatter(centers[:,0], centers[:,1], c='black', s=300, alpha=0.5);
plt.show()
```



```
import numpy as np
cluster_0= [i for i, x in enumerate(y_means) if x == 0]
cluster_1= [i for i, x in enumerate(y_means) if x == 1]
cluster_2= [i for i, x in enumerate(y_means) if x == 2]
cluster_3= [i for i, x in enumerate(y_means) if x == 3]
cluster_4= [i for i, x in enumerate(y_means) if x == 4]
```

Affichage des clusters

```
print(type(y_means))
print(cluster_0)
print(cluster_1)
print(cluster_2)
print(cluster_3)
print(cluster_4)

<class 'numpy.ndarray'>
[57, 61, 62, 63, 64, 65, 68, 70, 74]
[2, 3, 4, 5, 7, 10, 11, 14, 16, 76, 77, 78, 80, 81, 82, 83, 84, 85, 86, 87, 88, 89, 90, 91, 92, 93, 94]
[0, 1, 6, 8, 9, 12, 13, 15, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30, 31, 32, 33, 34, 35, 36, 37, 49, 52, 54, 55, 60, 75, 79]
```

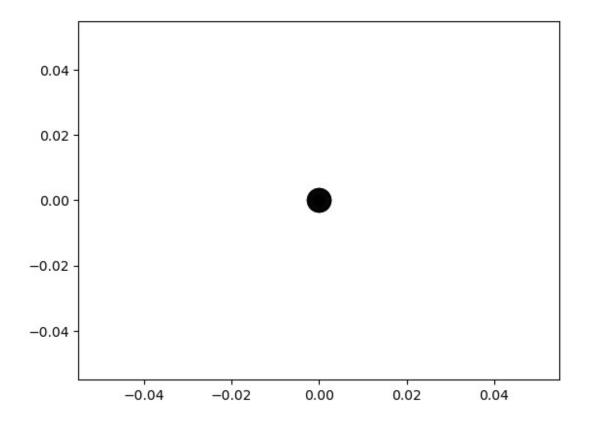
```
[38, 39, 40, 41, 42, 43, 44, 45, 46, 47, 48, 50, 51, 53, 56]
[58, 59, 66, 67, 69, 71, 72, 73]
```

Classer les documents dans chaque cluster en fonction de leurs similarités avec leur centroid (5 classe:....) Comparer les resultats obtenus avec les annotation preetablies

3.2 k-means avec SVD

```
centroids=np.array([[-0.10512687, -0.35341345],[-0.24220548,
[0.32397296], [-0.03986034, [0.09208828], [0.72163033, [0.16627089],
[0.72163033, 0.16627089]])
km = KMeans(n clusters=5, init=centroids, n init=50)
print("Clustering sparse data with %s" % km)
# vect=vectorsTFIDF(get corpus('/Users/rachad/Desktop/corpus/*'))
vects SVD, feature names SVD = docVectSVD([" ".join(doc.split()) for
doc in df content['ProcessedUserContent']], n components=50,
window size=4)
vects SVD matrix = np.array(list(vects SVD.values()),
dtype=np.float64) # Ensure float64
pca = PCA(n components=2)
reduced vects SVD = pca.fit transform(vects SVD matrix)
km.fit(reduced vects SVD)
v means SVD = km.predict(reduced vects SVD)
print("Cluster assignments:", y means SVD)
Clustering sparse data with KMeans(init=array([[-0.10512687, -
0.35341345],
       [-0.24220548, 0.32397296],
       [-0.03986034, 0.09208828],
       [ 0.72163033, 0.16627089],
       [ 0.72163033, 0.16627089]]),
       n clusters=5, n init=50)
Cluster assignments: [0 0 0 ... 0 0 0]
c:\Users\dscon\anaconda3\Lib\site-packages\sklearn\decomposition\
truncated svd.py:275: RuntimeWarning: invalid value encountered in
divide
  self.explained variance ratio = exp var / full var
c:\Users\dscon\anaconda3\Lib\site-packages\sklearn\decomposition\
_pca.py:653: RuntimeWarning: invalid value encountered in divide
  explained_variance_ratio_ = explained_variance_ / total_var
c:\Users\dscon\anaconda3\Lib\site-packages\sklearn\cluster\
kmeans.py:1414: RuntimeWarning: Explicit initial center position
passed: performing only one init in KMeans instead of n init=50.
  super(). check params vs input(X, default n init=10)
c:\Users\dscon\anaconda3\Lib\site-packages\sklearn\base.py:1473:
```

```
ConvergenceWarning: Number of distinct clusters (1) found smaller than
n clusters (5). Possibly due to duplicate points in X.
  return fit method(estimator, *args, **kwargs)
print(f'dict : {dict(zip(np.arange(0,100),y means))}')
score = km.score(reduced vects SVD)
print("Score:", score)
plt.scatter(reduced vects SVD[:,0], reduced vects SVD[:,1],
c=y means SVD, s=50, cmap='viridis')
centers = km.cluster centers
plt.scatter(centers[:,0],centers[:,1],c='black',s=300,alpha=0.5);
plt.show()
dict: {0: 2, 1: 2, 2: 1, 3: 1, 4: 1, 5: 1, 6: 2, 7: 1, 8: 2, 9: 2,
10: 1, 11: 1, 12: 2, 13: 2, 14: 1, 15: 2, 16: 1, 17: 2, 18: 2, 19: 2,
20: 2, 21: 2, 22: 2, 23: 2, 24: 2, 25: 2, 26: 2, 27: 2, 28: 2, 29: 2,
30: 2, 31: 2, 32: 2, 33: 2, 34: 2, 35: 2, 36: 2, 37: 2, 38: 3, 39: 3,
40: 3, 41: 3, 42: 3, 43: 3, 44: 3, 45: 3, 46: 3, 47: 3, 48: 3, 49: 2,
50: 3, 51: 3, 52: 2, 53: 3, 54: 2, 55: 2, 56: 3, 57: 0, 58: 4, 59: 4,
60: 2, 61: 0, 62: 0, 63: 0, 64: 0, 65: 0, 66: 4, 67: 4, 68: 0, 69: 4,
70: 0, 71: 4, 72: 4, 73: 4, 74: 0, 75: 2, 76: 1, 77: 1, 78: 1, 79: 2,
80: 1, 81: 1, 82: 1, 83: 1, 84: 1, 85: 1, 86: 1, 87: 1, 88: 1, 89: 1,
90: 1, 91: 1, 92: 1, 93: 1, 94: 1}
Score: -0.0
```



3.3 k-means avec Word2Vec

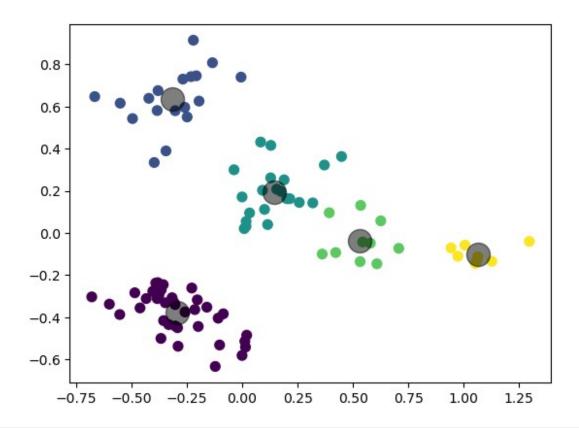
pretrainedDocWV

```
centroids=np.array([[-0.10512687, -0.35341345],[-0.24220548,
0.32397296],[-0.03986034, 0.09208828],[0.72163033, 0.16627089]],
[0.72163033, 0.16627089]])
km = KMeans(n_clusters=5, init=centroids,n_init=50)
print("Clustering sparse data with %s" % km)

# vect=vectorsTFIDF(get_corpus('/Users/rachad/Desktop/corpus/*'))
vects_DWV = pretrainedDocWV([" ".join(doc.split()) for doc in
df_content['ProcessedUserContent']])
vects_DWV_matrix = np.array(list(vects_DWV), dtype=np.float64) #
Ensure float64
pca = PCA(n_components=2)

reduced_vects_DWV = pca.fit_transform(vects_DWV_matrix)
km.fit(reduced_vects_DWV)
y_means_DWV = km.predict(reduced_vects_DWV)
print("Cluster assignments:", y_means_DWV)
```

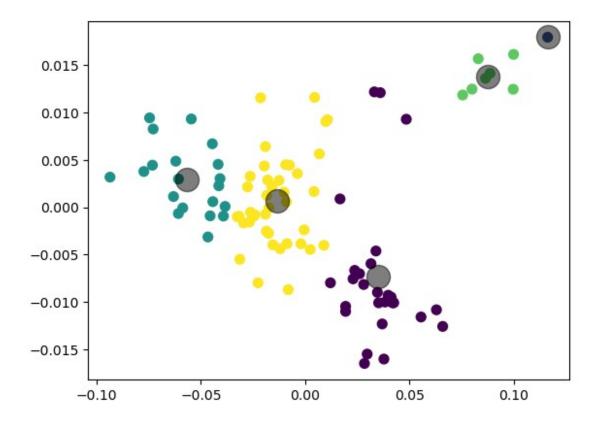
```
score = km.score(reduced vects DWV)
print("Score:", score)
plt.scatter(reduced_vects DWV[:,0], reduced vects DWV[:,1],
c=v means DWV, s=50, cmap='viridis')
centers = km.cluster centers
plt.scatter(centers[:,0],centers[:,1],c='black',s=300,alpha=0.5);
plt.show()
Clustering sparse data with KMeans(init=array([[-0.10512687, -
0.35341345],
      [-0.24220548, 0.32397296],
      [-0.03986034, 0.09208828],
      [ 0.72163033, 0.16627089],
      [ 0.72163033, 0.16627089]]),
      n_clusters=5, n init=50)
c:\Users\dscon\anaconda3\Lib\site-packages\sklearn\cluster\
kmeans.py:1414: RuntimeWarning: Explicit initial center position
passed: performing only one init in KMeans instead of n init=50.
 super(). check params vs input(X, default n init=10)
c:\Users\dscon\anaconda3\Lib\site-packages\sklearn\cluster\
kmeans.py:1429: UserWarning: KMeans is known to have a memory leak on
Windows with MKL, when there are less chunks than available threads.
You can avoid it by setting the environment variable
OMP NUM THREADS=1.
 warnings.warn(
1 1 2 1 1 1 1 1 1 1 1 2
4 4
Score: -3.084462097492615
```



propreDocWV

```
centroids=np.array([[-0.10512687, -0.35341345],[-0.24220548,
0.32397296], [-0.03986034, 0.09208828], [0.72163033, 0.16627089],
[0.72163033, 0.16627089]])
km = KMeans(n clusters=5, init=centroids, n init=50)
print("Clustering sparse data with %s" % km)
# vect=vectorsTFIDF(get corpus('/Users/rachad/Desktop/corpus/*'))
vects PDWV = propreDocW\overline{V}(["".join(doc.split())] for doc in
df content['ProcessedUserContent']])
vects PDWV matrix = np.array(list(vects PDWV), dtype=np.float64) #
Ensure float64
pca = PCA(n components=2)
reduced vects PDWV = pca.fit transform(vects PDWV matrix)
km.fit(reduced vects PDWV)
y means PDWV = km.predict(reduced vects PDWV)
print("Cluster assignments:", y means PDWV)
score = km.score(reduced vects PDWV)
```

```
print("Score:", score)
plt.scatter(reduced vects_PDWV[:,0], reduced_vects_PDWV[:,1],
c=y means PDWV, s=50, cmap='viridis')
centers = km.cluster centers
plt.scatter(centers[:,0],centers[:,1],c='black',s=300,alpha=0.5);
plt.show()
Clustering sparse data with KMeans(init=array([[-0.10512687, -
0.353413451.
       [-0.24220548, 0.32397296],
       [-0.03986034, 0.09208828],
       [ 0.72163033, 0.16627089],
       [ 0.72163033,
                    0.16627089]]),
      n clusters=5, n init=50)
4 4 4 4 4 4 2 0 2 2 2 4 4
4 0 0 4 0 0 0 0 0 0 0 0 4 0 0 4 0 2 4 0 4 3 3 2 4 4 0 0 0 3 3 4 3 4 1
3 3
2 2 0 0 4 2 0 4 4 0 4 0 0 4 4 0 0 0 4 4 2]
Score: -0.018221913539229657
c:\Users\dscon\anaconda3\Lib\site-packages\sklearn\cluster\
kmeans.py:1414: RuntimeWarning: Explicit initial center position
passed: performing only one init in KMeans instead of n init=50.
  super(). check params vs input(X, default n init=10)
c:\Users\dscon\anaconda3\Lib\site-packages\sklearn\cluster\
kmeans.py:1429: UserWarning: KMeans is known to have a memory leak on
Windows with MKL, when there are less chunks than available threads.
You can avoid it by setting the environment variable
OMP NUM THREADS=1.
 warnings.warn(
```



3.4 k-means avec Doc2Vec

```
centroids=np.array([[-0.10512687, -0.35341345],[-0.24220548,
[0.32397296], [-0.03986034, [0.09208828], [0.72163033, [0.16627089],
[0.72163033, 0.16627089]])
km = KMeans(n clusters=5, init=centroids, n init=50)
print("Clustering sparse data with %s" % km)
# vect=vectorsTFIDF(get corpus('/Users/rachad/Desktop/corpus/*'))
vects_D2V = vectorsDoc2Vec([" ".join(doc.split()) for doc in
df content['ProcessedUserContent']])
vects D2V matrix = np.array(list(vects D2V), dtype=np.float64) #
Ensure float64
pca = PCA(n components=2)
reduced vects D2V = pca.fit transform(vects D2V matrix)
km.fit(reduced vects D2V)
y means D2V = km.predict(reduced vects D2V)
print("Cluster assignments:", y means D2V)
score = km.score(reduced vects D2V)
print("Score:", score)
```

```
plt.scatter(reduced vects D2V[:,0], reduced vects D2V[:,1],
c=y means D2V, s=50, cmap='viridis')
centers = km.cluster centers
plt.scatter(centers[:,0],centers[:,1],c='black',s=300,alpha=0.5);
plt.show()
Clustering sparse data with KMeans(init=array([[-0.10512687, -
0.35341345],
      [-0.24220548,
                   0.323972961,
      [-0.03986034,
                   0.092088281,
      [ 0.72163033,
                   0.16627089],
      [ 0.72163033, 0.16627089]]),
      n clusters=5, n init=50)
3 3 3 3 3 3 3 3 3 3 3 3
2 2
2 2 0 0 0 2 2 2 0 0 0 0 2 0 0 2 2 0 0 2 2]
Score: -126.77455046640674
c:\Users\dscon\anaconda3\Lib\site-packages\sklearn\cluster\
kmeans.py:1414: RuntimeWarning: Explicit initial center position
passed: performing only one init in KMeans instead of n init=50.
 super(). check params vs input(X, default n init=10)
c:\Users\dscon\anaconda3\Lib\site-packages\sklearn\cluster\
kmeans.pv:1429: UserWarning: KMeans is known to have a memory leak on
Windows with MKL, when there are less chunks than available threads.
You can avoid it by setting the environment variable
OMP NUM THREADS=1.
 warnings.warn(
```

