INFORMATION RETRIEVAL ASSIGNMENT-2

Group No. 78

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Importing necessary libraries

```
In [1]: #Installing necessary libraries
        import os
        import glob
        import nltk
        from nltk.tokenize import RegexpTokenizer
        from nltk.corpus import stopwords
        from nltk.stem import WordNetLemmatizer
        from nltk.stem.porter import PorterStemmer
        import re
        import string
        import json
        import pandas as pd
        import numpy as np
        import math
        from textblob import TextBlob, Word
        import joblib
        import matplotlib.pyplot as plt
        from nltk.tokenize import word tokenize
        from pandas.core.dtypes.cast import dict compat
        nltk.download('punkt')
        nltk.download('wordnet')
        nltk.download('stopwords')
        lema = WordNetLemmatizer()
        [nltk_data] Downloading package punkt to
        [nltk data]
                      C:\Users\bhava\AppData\Roaming\nltk data...
        [nltk_data] Package punkt is already up-to-date!
        [nltk data] Downloading package wordnet to
                        C:\Users\bhava\AppData\Roaming\nltk data...
        [nltk_data]
        [nltk data] Package wordnet is already up-to-date!
```

QUESTION 1

Methodology:

Read files

- To read files present in "Humor, Hist, Media, Food" folder
 path = "/content/drive/MyDrive/Datasets/Humor, Hist, Media, Food"
- To open a particular file

```
rf = open(path,'r',errors = 'ignore')
```

To read content of file

```
rf = rf.read()
```

a) Preprocessing steps for the given data:

- (i) Converting the text to lower case
- (ii) Word Tokenization
- (iii) Removing the stop words from tokens
- (iv) Removing the punctuation marks from tokens
- (v) Removing the blank spaces tokens

```
[ ] In = len(string.punctuation)
    #Cleaning the data

def pre_process(content):
    #Convert the text to lower case
    content = content.lower()
    #Remove punctuation marks from tokens
    content = content.translate(str.maketrans(string.punctuation, " "*ln,''))
    #Perform word tokenization
    ctokens = word_tokenize(content)
    #Remove stopwords from tokens and do lemmatization
    #Checking length, if length = 1
    ctokens = [lema.lemmatize(s) for s in ctokens if s not in stopwords.words('english') and s.isalpha and len(s)>1]
    return ctokens
```

b)Performing intersection and union between document and query

```
In [7]: function to make set of the document token and query token and perform intersection and union between the query and each document

af jaccard_coeff(tdoc,tquery):
    set1 = set(tdoc)
    set2 = set(tquery)
    un = set1.union(set2)
    it = set1.union(set2)
    it = set1.intersection(set2)
    #Jaccard Coefficient = Intersection of (doc,query) / Union of (doc,query)
    jc = len(it)/len(un)
    return jc
```

c) Implementation of Jaccard co-efficient and reporting top 5 relevant documents

```
In [8]: #Function to Report the top 5 relevant documents based on the value of the Jaccard coefficient
def top_doc(pquery):
    js = []
    for i in dc:
        js.append([jaccard_coeff(dc[i],pquery),i])
        #Sorting based on the coefficient value
    js.sort(key=lambda x:x[0])
    return js
```

Output:

```
In [9]: #Take input query from user
        print("Enter Query:")
        query = input()
        #Pre process the input query
        pquery = pre process(query)
        print(pquery)
        js = top_doc(pquery)
        #reporting top 5
        for i in range(5):
            print(document[js[i][1]])
        Enter Query:
        100 west by 50 north
        ['100', 'west', '50', 'north']
        1st aid.txt
        abbott.txt
        acetab1.txt
        aclamt.txt
        acronym.lis
```

To calculate Jaccard coefficient:

- Create tokens for queries and documents after pre-processing
- Converting them to sets
- Jaccard coefficient = Intersection of (doc, query)/Union of (doc, query)
- Sort values in descending order
- Reporting files having top 5 jaccard coefficients

Ranked-Information Retrieval and Evaluation

```
f = freq()
print(len(f))
```

To find the unique words from the documents

```
: unq = []
#finding unique words present
for lt in dc:
    list1 = list(set(dc[lt]))
    unq.extend(list1)
    unq = list(set(unq))
print(len(unq))
```

To find the document frequency

```
doc_f = doc_freq()
print(len(doc_f))
```

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Finding the inverse document frequency

```
#Function to compute inverse document frequency
def inverse_df():
    idf = {}
    total_doc = len(dc)
    for t in doc_f:
        idf[t] = math.log(total_doc/doc_f[t]+1)
    return idf
```

Finding the Frequency and its length

```
from pandas.core.dtypes.cast import dict_compat
#Finding frequency
def freq():
    f = {}
    for i in dc:
        f[i] = {}
        for t in dc[i]:
            if t in f[i]:
                 f[i][t] += 1
        else:
        f[i][t] = 1
    return f
```

```
f = freq()
print(len(f))

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```

e) Implementation of TF-IDF matrix and finding top 5 relevant documents

f) Implementation of top 5 Weighting Schemas The five Weighting Schemas are:

- Binary weighting scheme
- Raw count weighting scheme
- Term frequency weighting scheme
- Log normalization weighting scheme
- Double normalization weighting scheme

Binary weighting scheme

```
#Function to compute Binary weighting scheme
def binary_tf():
    btf = {}
    for di in f:
        btf[di]={}
        for w in f[di]:
            btf[di][w] = 1
    return btf

|: btf = binary_tf()
    print(btf[1])

{'herbalherb1st': 1, 'aidcalendulacomfreyremediessickmedicine' ointment': 1, 'use': 1, 'minor': 1, 'cut': 1, 'graz': 1, 'ree': 1, 'damage': 1, 'external': 1, 'blood': 1, 'vessel': 1,
```

Raw count weighting scheme

```
#Function to compute Raw count weighting scheme
def rawcount_tf():
    rctf = {}
    for di in f:
        rctf[di]={}
        for w in f[di]:
            rctf[di][w]=f[di][w]
    return rctf
rctf = rawcount_tf()
print(len(rctf[1]))
```

Term frequency weighting scheme

```
#Function to compute term frequency weighting scheme
def term_freq():
    tf = {}
    for di in f:
        tf[di]={}
        s=sum(f[di].values())
        for w in f[di]:
            tf[di][w] = f[di][w]/s
    return tf

tf = term_freq()
print(len(tf[1]))
```

Log normalization weighting scheme

```
#Function to compute Log normalization weighting scheme
def log_tf():
    ltf={}
    for di in f:
        ltf[di]={}
        for w in f[di]:
            ltf[di][w]=math.log(1+f[di][w])
    return ltf
```

```
ltf = log_tf()
print(len(ltf[1]))
```

Double normalization weighting scheme

```
#Function to compute Double normalization weighting scheme

def double_tf():
    dntf = {}
    for di in f:
        dntf[di] = {}
        m=max(f[di].values())
        for w in f[di]:
            dntf[di][w] = 0.5+0.5*(f[di][w]/m)
    return dntf

dntf = double_tf()
print(len(dntf[1]))
```

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Output

```
print("Enter Input Query:")
query = input()
pquery = pre process(query)
print(pquery)
#binary weighting scheme
print("\nbinary ")
bd = iquery(pquery,btfidf)
print(*bd)
#raw count weighting scheme
print("\nraw count ")
rcd = iquery(pquery,rctfidf)
print(*rcd)
#term frequency weighting scheme
print("\nterm frequency ")
tfd = iquery(pquery,tfidf)
print(*tfd)
#log normalization weighting scheme
print("\nlog norm ")
ld = iquery(pquery,ltfidf)
print(*ld)
#double normalization weighting scheme
print("\ndouble norm ")
dnd = iquery(pquery,dntfidf)
print(*dnd)
```

```
Enter Input Query:
once upon a time.
['upon', 'time']
binary
adt_miam.txt allusion all_grai amazing.epi ambrose.bie ayurved.txt
raw count
mlverb.hum practica.txt barney.txt humor9.txt manners.txt xibovac.txt
term frequency
timetr.hum ookpik.hum sysman.txt corporat.txt trukdeth.txt yuppies.hum
log norm
barney.txt mindvox practica.txt quack26.txt humor9.txt jokes1.txt
double norm
ookpik.hum jokes1.txt trukdeth.txt ambrose.bie mindvox flux_fix.txt
```

Pros and Cons

Binary:

Advantage: This scheme is the simplest one to compute, as for this only presence and absence of words matters

Disadvantages: does not consider frequency od word

Raw Count:

Advantage: raw count of words in a document is determined, and comparatively more relevant documents are retrieved.

Disadvantage: Large documents are favored more

Term Frequency

Advantage: Reduced the bias caused by the length of documents Disadvantage: more storage space

Log normalization

Advantage: Reduced computational power

Disadvantage: This is similar to raw count prefers large documents over small ones which is not always the case in reality.

Double normalization

Advantage: Reduced computational power and also considers the term frequency that is normalized using document length

Disadvantage: In the denominator, we have a frequency of terms that is maximum in the document. There can occur a case in which that term is not relevant which gives wrong results

QUESTION 2

Methodology:

Read files

- To read files present in "Humor, Hist, Media, Food" folder

 path = "/content/drive/MyDrive/Datasets/Humor, Hist, Media, Food"
- To open a particular file

```
rf = open(path,'r',errors = 'ignore')
```

• To read content of file

```
rf = rf.read()
```

• To read "assignment-2.txt" file

```
data =
pd.read_csv("/content/drive/MyDrive/IR-assignment-2-data.txt", sep='
', header=None)
```

To convert the text file to csv file

```
data.to csv('Q2.csv', index = None)
```

```
data = pd.read_csv("/content/drive/MyDrive/IR-assignment-2-data.txt", sep=' ', header=None)
data.to_csv('Q2.csv', index = None)
```

a) Preprocessing steps for the given data:

- (i) Converting the text to lower case
- (ii) Word Tokenization
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#Perform word tokenization
ctokens = word_tokenize(content)
#Remove stopwords from tokens and do lemmatization
#Checking length, if length = 1
ctokens = [lema.lemmatize(s) for s in ctokens if s not in stopwords.words('english') and s.isalpha and len(s)>1]
return ctokens
```

b) qid:4 queries consideration

```
In [47]: #Select rows with qid:4
    fdata = df[df[1] == 'qid:4']

    total_files = 1
    temp = []
    r = fdata[0].unique()
    for i in r:
        temp.append(len(fdata[fdata[0] == i]))
        total_files = total_files * math.factorial(len(fdata[fdata[0] == i]))

#Sort the data on the basis of relevance judgement label
    final_data = fdata.sort_values(by = 0, ascending = False )
    final_data = final_data.reset_index(drop=True)
```

```
In [53]: #Select rows with qid:4
fdata = df[df[1] == 'qid:4']

total_files = 1
temp = []
r = fdata[0].unique()
for i in r:
    temp.append(len(fdata[fdata[0] == i]))
    total_files = total_files * math.factorial(len(fdata[fdata[0] == i]))

#Sort the data on the basis of relevance judgement label
final_data = fdata.sort_values(by = 0, ascending = False )
final_data = final_data.reset_index(drop=True)
```

c) File arrangement

d) Implementation of DCG

Firstly we will sort the data before the DCG Calculation

```
In [62]: #Sorting data in ascending order
d1 = fdata.sort_values(by=0, ascending=False)
```

```
In [67]: #Function to compute DCG
def DCG_Calculation(datadup,n):
    datadup = np.asfarray(datadup)[:n]
    dcg_val = datadup[0] + np.sum(datadup[1:] / np.log2(np.arange(2, datadup.size + 1)))
    return dcg_val
```

e) Max DCG Calculation

The Maximium DCG is : 20.989750804831452

f) Implementation of nDCG

```
In [69]: #computing nDCG
def NDCG_Calculation(r, k):
    denominator = DCG_Calculation(d1[0], k)
    numerator = DCG_Calculation(r, k)
    value = numerator/denominator
    return value
```

g) Implementation of nDCG at 50

```
In [70]: #nDCG value at 50
nDCG_value_50 = NDCG_Calculation(fdata[0],50)
print("The value of nDCG at 50 is : ",nDCG_value_50 )
The value of nDCG at 50 is : 0.35210427403248856
```

h) Implementation of nDCG for whole data set

```
In [71]: #nDCG value for complete dataset
    nDCG_value = NDCG_Calculation(fdata[0],len(fdata))
    print("The value of nDCG for all is : ",nDCG_value )

The value of nDCG for all is : 0.5979226516897828
```

i) Obtaining values from TF-IDF and storing them into a list

```
In [73]: #To store the values obtained from TF-IDF into a list
    p=0
    value1 = fdata[76]
    value2 = fdata[0]
    value11 = value1.tolist()
    value21 = value2.tolist()
    for dup_list in value11:
        if dup_list[0] == "7" and dup_list[1] == "5" and dup_list[2] == ":":
        i1 = dup_list[3:]
        value11[p] = i1
        p = p+1
    value11f = [float(i) for i in value11]
```

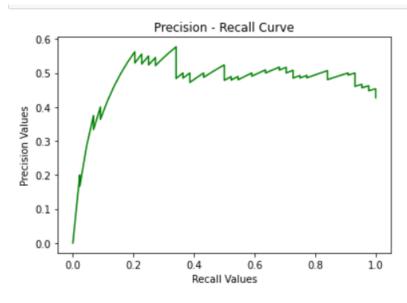
k) Plotting precision and recall curve

```
In [80]: #Declaring the required variables and the precision and recall lists
    total=0
    relevant_present=0
    relevant_doc=0
    list_prevision=[]
    list_recall=[]
```

```
In [79]: #Calculation of precision and recall values
         sorted_rel = [m for n,m in sorted(zip(xmlf,xol),reverse=True)]
         sorted_rel2 = [(n,m) for n,m in sorted(zip(xmlf,xol),reverse=True)]
         for m in sorted_rel:
             if m == 0:
                 relevant_doc = relevant_doc
             else:
                 relevant doc = relevant doc + 1
         for m in sorted rel:
             total= total + 1
             if m == 0:
                relevant present = relevant present
             else:
                 relevant_present = relevant_present + 1
             list prevision.append(relevant present/total)
             list_recall.append(relevant_present/relevant_doc)
         #Plotting Precision - Recall Graph
         plt.title("Precision - Recall Curve")
         plt.plot(list_recall, list_prevision, color="green")
         plt.xlabel("Recall Values")
         plt.ylabel("Precision Values")
         plt.show()
```

Output Graph:

Distinctive Sawtooth shape curve



QUESTION 3

Methodology:

- To read files present in "Humor, Hist, Media, Food" folder path = "/content/drive/MyDrive/20 newsgroups.zip"
- To open a particular file

 rf = open(path, 'r', errors = 'ignore')
- To read content of file

```
[26] #Load the file
    file = "/content/drive/MyDrive/20_newsgroups.zip"
    #Defining labels
    label = ['comp.graphics', 'rec.sport.hockey', 'sci.med', 'sci.space', 'talk.politics.misc']
    # opening the zip file in read mode
    with ZipFile(file, 'r') as zip:
        zip.printdir()
        zip.extractall()
```

Read files

- a) Preprocessing steps for the given data:
 - (i) Converting the text to lower case
 - (ii) Word Tokenization

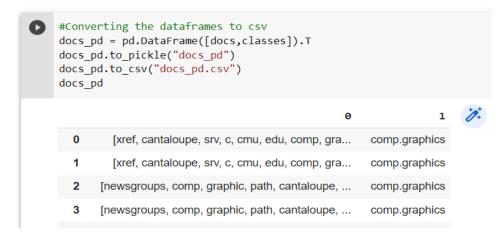
- (iii) Removing the stop words from tokens
- (iv) Removing the punctuation marks from tokens
- (v) Removing the blank spaces tokens

```
[ ] In = len(string.punctuation)
    #Cleaning the data
    def pre_process(content):
        #Convert the text to lower case
        content = content.lower()
        #Remove punctuation marks from tokens
        content = content.translate(str.maketrans(string.punctuation, " "*ln,''))
        #Perform word tokenization
        ctokens = word_tokenization
        ctokens = word_tokenize(content)
        #Remove stopwords from tokens and do lemmatization
        #Checking length, if length = 1
        ctokens = [lema.lemmatize(s) for s in ctokens if s not in stopwords.words('english') and s.isalpha and len(s)>1]
        return ctokens
```

```
[12] #performing preprocessing and saving the data
docs = []
    # word_list={}
    for path in file_list:
        file = open(path, 'r', encoding='cp1250')
        text = file.read().strip()
        x=pre_process(text)
        file.close()
        docs.append(x)
```

'docs_pd = pd.DataFrame([docs,classes]).T\nprint(docs_pd)\ndocs_pd[0] = pre_pro

Converting the data frames to CSV:



b) Implementation of TF-ICF

```
#Class for implementing Naive Bayes algorithm with TF-ICF
class NaiveBayes_tf_icf:
  #Function to predict
  def predict(self,X_test):
    predc = []
    for i in range(len(X_test)):
        classes_words_probability = []
        for 1 in label:
            words_probability = 0
            for word in X_test[i]:
                fr, cn = self._word_freq(word, 1)
                pp = (fr+1) /(cn+len(self._unique_words))
                words_probability += np.log(pp)
            classes_words_probability.append(words_probability)
        predc.append(label[np.argmax(classes_words_probability)])
    return predc
  #Function to compute confusion matrix
  def confusion_matrix(self, ypred, ytest):
    matrix= np.zeros((len(label), len(label))).astype(int)
    for i in range(len(ypred)):
        matrix[label.index(ypred[i])][label.index(ytest[i])]+= 1
    return matrix
```

```
#Function to compute accuracy
def calculate_accuracy(self, ypred, ytest):
 return len([1 for i in range(len(ypred)) if ypred[i] == ytest[i]])/len(ypred)
#Function to compute word frequency
def _word_freq(self, word, label):
 try:
     return self._word_freq_per_class[label, word], self._number_words_perclass[label]
 except:
     return 0, self. number words perclass[label]
#Calculate tf-icf
def _calculate_tf_icf(self):
 self._tf_icf = {}
 c = Counter(self._word_list)
 for i in set(self._word_list):
   tf = c[i]
   icf = np.log(len(self._m_dict)/self._class_word[i]+1)
   self._tf_icf[i] = tf*icf
```

```
#Function to fit the data
def fit(self,X train,y train, k):
  words = X_train
  self. N = len(words)
  classes = y_train
  self. m dict = {}
  for i in range(self._N):
    if classes[i] in self._m_dict.keys():
        self._m_dict[classes[i]] = self._m_dict[classes[i]] + words[i]
    else:
        self. m dict[classes[i]] = words[i]
  #Listing words containing multiple occurence of same word
  self._word_list = []
  for i in self._m_dict:
      self. word list = self. word list + self. m dict[i]
  #Count of word per class
  self._class_word = {}
  for i in self. m dict:
    l=self._m_dict[i]
    for j in set(1):
      if j not in self._class_word.keys():
        self._class_word[j] = 1
      else:
        self. class word[j] += 1
  self._calculate_tf_icf()
  sorted_x = sorted(self._tf_icf.items(), key = operator.itemgetter(1), reverse=True)
```

```
#considering top k features
self._unique_words = [i[0] for i in sorted_x[:int(len(sorted_x)*k)]]
self._word_freq_per_class = {}
self._number_words_perclass = {}
for i in label:
   freq_list= Counter(self._m_dict[i])
    for j in self._unique_words:
        self._word_freq_per_class[i,j] = freq_list[j]
        if i in self._number_words_perclass.keys():
            self._number_words_perclass[i] = self._number_words_perclass[i] +freq_list[j]
        else:
            self._number_words_perclass[i] = freq_list[j]
self._freq_train = {}
for i in y train:
  if i not in self._freq_train.keys():
    self._freq_train[i] = 1
  else:
    self. freq train[i] += 1
```

C) Training the data with Naive Bayes Classifier

```
_{268}^{\checkmark} [24] #50:50, 70:30, and 80:20 training and testing split ratios
        ratio = [0.5,0.7,0.8]
        naive_tf_dict = []
        for i in range(3):
          train = docs_pd.sample(frac=ratio[i],random_state=42)
          xtrain, ytrain = train[0].tolist(),train[1].tolist()
          test = docs_pd.sample(frac=1,random_state=42).drop(train.index)
          xtest,ytest = test[0].tolist(),test[1].tolist()
          nb = NaiveBayes_tf_icf()
          #Fitting the xtrain and ytrain
          #Taking k as 500
          k = 500
          nb.fit(xtrain, ytrain, k)
          ypred = nb.predict(xtest)
          r = int(ratio[i]*100)
          accuracy = nb.calculate_accuracy(ypred, ytest)*100
          naive_tf_dict.append(accuracy)
          print("At ratio {}:{}".format(r,100-r))
          print("\n")
          print("Confusion Matrix is given by: \n",nb.confusion_matrix(ypred,ytest))
          print("\n")
          print("Accuracy is {:.2f} \n".format(accuracy))
```

<u>Output</u>

Accuracy and Confusion matrix at the ratio of 50 and 50:

Accuracy is 97.60

Accuracy is 98.20

At ratio 50:50

Accuracy and Confusion matrix at the ratio of 70 and 30 :

Accuracy and Confusion matrix at the ratio of 80 and 20:

Analysis

Naive Bayes using TF-ICF has less noise due to feature selection which only selects the important words whose TF-ICF value is more than other words.

Naive Bayes with TF-ICf performs better than the normal Naive Bayes because of top k feature selection.