Group 27

Team Members:

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Topic Name:

"Disease Symptom Prediction and its Data Analytics"

Involved Components:

- 1)Indexing
- 2)Searching Component
- 3)Refining
- 4)Feedback
- 5)Assessment Components

1)Indexing:

```
file_name="data2.csv"

# Read File
df=pd.read_csv(file_name)

documentname_list=list(df['label_dis'])
print(documentname_list)
df=df.iloc[:,1:]
columnsName=list(df.columns)

# print(columnsName)

documentname_list=list(documentname_list)

# df
Num_of_diseases=len(df)
Num_of_symptoms=len(columnsName)
print(Num_of_diseases,Num_of_symptoms)
```

['hypertensive disease', 'diabetes', 'depression to disease', 'pneumonia', 'failure heart congest olesterolemia', 'infection', 'infection urinary iency renal', 'confusion', 'degenerative polyamalignant neoplasm', 'acquired immuno-deficient disease', 'septicemia', 'systemic infection', bolism pulmonary', 'epilepsy', 'cardiomyopathy disease', 'psychotic disorder', 'hyperlipidemian prostatic hypertrophy'. 'kidney failure acut

We took Indexed data online and found the TF, IDF, and TF_IDF score.

2) Searching Component

```
In [59]: user symptoms = str(input(" enter symptoms separated by comma")).lower().split(',')
          print()
          processed_user_symptoms=[]
          for sym in user_symptoms:
              sym=sym.strip()
             sym=sym.replace('-','')
sym=sym.replace("'",'')
             sym = ' '.join([lemmatizer.lemmatize(word) for word in splitter.tokenize(sym)])
             processed user symptoms.append(sym)
          processed user symptoms
          print(processed_user_symptoms)
          user_symptoms=processed_user_symptoms
          enter symptoms separated by commafever, cold, sneeze, head ache
          ['fever', 'cold', 'sneeze', 'head ache']
In [60]: symptoms matched = set()
          for idx, data in enumerate(symptoms dataset):
              data split=data.split()
              for user_sym in user_symptoms:
                 count=0
                 for symp in data_split:
                      if symp in user_sym.split():
                  if count/len(data_split)>0.5:
                      symptoms_matched.add(data)
          symptoms_matched = list(symptoms_matched)
          symptoms matched
Out[60]: ['sneeze', 'ache', 'fever']
```

Enter the query in comma-separated values, then the list gets matched with the symptoms dataset and returns the symptoms_matched list

3) Refining Searches

['chill', 'cough', 'pain', 'diarrhea', 'vomiting', 'tachypnea', 'apyrexial', 'shortness of breath', 'unresponsiveness', 'night sweat', 'rale', 'decreased body weight', 'pleuritic pain', 'spontaneous rupture of membranes', 'pain abdominal', 'nausea', 'pro ductive cough', 'prunitus', 'swelling', 'lethangy', 'decreased translucency,' (siterses respiratory', 'feeling suicidal', 'pati ent non compliance', 'lesion', 'haemorrhage', 'hypotension', 'agitation', 'rhonchus', 'asthenia', 'haemoptysis', 'hypotonic', 'muscle hypotonia', 'erythema', 'redness', 'hallucinations auditory', 'mental status changes', 'abscess bacterial', 'sore to touch', 'bradycardia', 'throat sore', 'abdominal tenderness', 'unsteady gait', 'gurgle', 'transaminitis', 'debilitation', 'irritable mood', 'mass of body structure', 'hyponatremia', 'difficulty passing urine', 'hemodynamically stable', 'dysuri a', 'breech presentation', 'cyanosis', 'chest tightness', 'hyperkalemia', 'malise', 'anorexia', 'frail', 'dyspnea', 'sensory d iscomfort', 'snuffle', 'wheezing', 'blackout', 'headache', 'scratch marks', 'ecchymosis', 'bedridden', 'facial paresis', 'synco pe', 'unconscious state', 'extreme exhaustion', 'hemiplegia', mediastinal shift', 'acites', 'distended abdomen', 'lung nodul e', 'metastatic lesion', 'gravida o', 'drowsiness', 'suicidal', 'withdraw', 'worry', 'green 'sputum', 'thicken', 'consciousness clear', 'hematuria', 'hyperacusis', 'pain chest', 'hepatosplenomegaly', 'tremor', 'urgency of micturition', 'egophony', 'fremit us', 'non-productive cough', 'splenomegaly', 'labored breathing', 'myalgia', 'scleral icterus', 'symptom aggravating factors', 'indifferent mood', 'dizziness', 'arthralgia', 'macule', 'painful swallowing', 'photophobia', 'monocytosis', 'posterior rhinorr hea', 'fall', 'clonus', 'seizure', 'stupor', 'asterixis', 'heavy feeling', 'macerated skin', 'mass in breast', 'paraparesis', 'sleepy', 'verbally abusive behavior', 'pain foot', 'prostate tender', 'urinary hesitation', 'makening early', 'nausea and vom iting', 'tenesmus', 'urg

All Symptoms List got from First Query.

The whole list of symptoms of the "diseases having at least one symptom in symptoms_matched list".

Actual Refining:

```
# refine the input of symptoms by adding the more Symptoms to it
          input2=input("enter some more symptoms from printed above Symptoms:")
          arr3=input2.split(",")
          for i in arr3:
              finalSymptoms.append(i)
          finalSymptoms
         enter some more symptoms from printed above Symptoms:chill,cough,pain
Out[65]: ['sneeze',
           'ache',
           'fever',
           'chill',
           'cough',
           'pain',
           'diarrhea',
           'vomiting',
           'tachypnea',
           'apyrexial',
           'shortness of breath',
           'unresponsiveness',
           'night sweat',
           'chill',
           'cough',
           'pain']
```

Enter some more symptoms (query terms)matching your need those get added to the final symptoms list.

4) Feedback

Before Feedback:

```
In [69]:
         print(" Disease based on Cosine Similarity ")
         print(cosine_similiarity_docs1)
         cosine_similiarity_docs1_sorted = dict(sorted(cosine_similiarity_docs1.items(),
                                              key=lambda kv: kv[1], reverse=True))
         print(cosine similiarity docs1 sorted)
         print()
         j = 0
         cosine similiarity docs1 index mapping = {}
         for key in cosine similiarity docs1 sorted:
           print(f" {diseases[key]} =====> {round(cosine_similiarity_docs1_sorted[key], 2)}"
           cosine_similiarity_docs1_index_mapping[j] = diseases[key]
          Disease based on Cosine Similarity
         {40: 0.2586992556490109, 12: 0.2845617966292759, 114: 0.2677458320613382, 115: 0.26
         0.2983775039229011, 26: 0.2983775039229011, 124: 0.2422478978250813, 29: 0.22580634
         {24: 0.2983775039229011, 25: 0.2983775039229011, 26: 0.2983775039229011, 12: 0.2845
         0.2677458320613382, 40: 0.2586992556490109, 124: 0.2422478978250813, 29: 0.22580634
          cardiomyopathy ====> 0.3
          cellulitis ====> 0.3
          cholecystitis ====> 0.3
          asthma ====> 0.28
          overload fluid ====> 0.27
          pancreatitis ====> 0.27
          dehydration ====> 0.26
          pneumonia aspiration ====> 0.24
          chronic kidney failure ====> 0.23
          chronic obstructive airway disease ====> 0.23
```

Document retrieval("Disease Name") based on cosine similarity score before the pseudo relevance feedback.

Feedback:

```
#pseudo relevance feedback
In [70]:
          diseases
         dd keys=list(cosine similiarity docs1 sorted.keys())
          dd=[]
          print(dd keys)
          j=0
          for i in dd keys:
              if j<5:
                  dd.append(i)
                  j+=1
          print(dd)
         symp=[]
          for i in dd:
              for col in columnsName:
                  if(df.loc[i,col]!=0):
                   symp.append(col)
         symp=list(set(symp))
          symp
          [24, 25, 26, 12, 114, 115, 40, 124, 29, 30]
          [24, 25, 26, 12, 114]
Out[70]:
          ['tachypnea',
           'muscle hypotonia',
           'pruritus'.
```

Feedback considering Top 5 as relevant documents. And computing the Cosine similarity between query(final symptoms list) and symptoms in Top5 diseases.

After FeedBack:

```
In [/1]:
             cosine similiarity docs3=cosine similarity(count, symp)
             # In[108]:
             print("Top most 10 Refined Diseases on Cosine Similarity")
             print()
             print(cosine similiarity docs3)
             cosine similiarity docs3 sorted = dict(sorted(cosine similiarity docs3.items(),
                                                     key=lambda kv: kv[1], reverse=True))
             print(cosine similiarity docs3 sorted)
             print()
             cosine_similiarity_docs3_index_mapping = {}
             for key in cosine_similiarity_docs3_sorted:
               print(f" {diseases[key]} =======> {round(cosine_similiarity_docs3_sorted[key], 2)}")
               cosine_similiarity_docs3_index_mapping[j] = diseases[key]
             Top most 10 Refined Diseases on Cosine Similarity
             {12: 0.5458296576564219, 48: 0.2007299652295673, 114: 0.6825490747441342, 115: 0.6825490
             0.5943095890082317, 25: 0.5943095890082317, 26: 0.5943095890082317, 27: 0.25084640023732
             {114: 0.6825490747441342, 115: 0.6825490747441342, 24: 0.5943095890082317, 25: 0.5943095
             0.5458296576564219, 53: 0.27728771385986933, 27: 0.250846400237328, 125: 0.2454419099685
              overload fluid =====> 0.68
              pancreatitis ======> 0.68
              cardiomyopathy ======> 0.59
              cellulitis ======> 0.59
              cholecystitis ======> 0.59
              asthma ======> 0.55
```

Document retrieval("Disease Name") based on cosine similarity score after the pseudo relevance feedback.

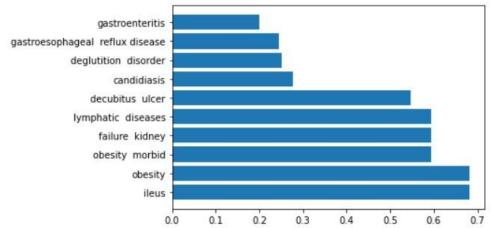
5)Assessment Components Accuracy:

```
In [38]: #Accuracy
         # diseases
         dd_keys=list(cosine_similiarity_docs1_sorted.keys())
         print(dd keys)
         for i in dd keys:
            dd2.append(diseases[i])
         # print(dd2)
         dd_keys=list(cosine_similiarity_docs3_sorted.keys())
         print(dd keys)
         for i in dd keys:
           dd3.append(diseases[i])
         # print(dd3)
         #Accuracy
         def common(a,b):
             c = [value for value in a if value in b]
             return c
         d=common(dd2,dd3)
         print("Accuracy Before and after Pseudo Relevance Feedback is :",(len(d)/10) *100)
         [109, 110, 60, 93, 83, 18, 36, 138, 96, 19]
         [83, 109, 110, 60, 93, 36, 18, 39, 64, 63]
         Accuracy Before and after Pseudo Relevance Feedback is : 70.0
```

Finding accuracy between the documents retrieved before and after the relevance feedback.

Plotting:

```
In [72]: # plotting in graph
    diseases_lst = dd3
    data = list(cosine_similiarity_docs3_sorted.values())
    x = dd3
    y = list(cosine_similiarity_docs3_sorted.values())
    plt.barh(x, y)
    plt.show()
    fig = plt.figure(figsize =(10, 7))
    plt.pie(data, labels = diseases_lst)
    plt.show()
```



Plotting the diseases in a Bar Graph with respective cosine similarity scores