CS6370: Natural Language Processing Project

Release Date: 21st March 2024 Deadline:

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General Instructions:

- 1. The template for the code (in Python) is provided in a separate zip file. You are expected to fill in the template wherever instructed. Note that any Python library, such as nltk, stanfordcorenlp, spacy, etc, can be used.
- 2. A folder named 'Roll_number.zip' that contains a zip of the code folder and your responses to the questions (a PDF of this document with the solutions written in the text boxes) must be uploaded on Moodle by the deadline.
- 3. Any submissions made after the deadline will not be graded.
- 4. Answer the theoretical questions concisely. All the codes should contain proper comments.
- 5. For questions involving coding components, paste a screenshot of the code.
- 6. The institute's academic code of conduct will be strictly enforced.

The first assignment in the NLP course involved building a basic text processing module that implements sentence segmentation, tokenization, stemming / lemmatization, stopword removal, and some aspects of spell check. This module involves implementing an Information Retrieval system using the Vector Space Model. The same dataset as in Part 1 (Cranfield dataset) will be used for this purpose. The project is split into two components - the first is a *warm-up* component comprising of Parts 1 through 4 that would act as a precursor for the second and main component, where you improve over the basic IR system.

Consider the following three documents:

d₁: Herbivores are typically plant eaters and not meat eaters

d₂: Carnivores are typically meat eaters and not plant eaters

d₃: Deers eat grass and leaves

1. Assuming {are, and, not} as stop words, arrive at an inverted index representation for the above documents.

```
Herbivores \rightarrow d<sub>1</sub>

typically \rightarrow d<sub>1</sub>, d<sub>2</sub>

plant \rightarrow d<sub>1</sub>, d<sub>2</sub>

eaters \rightarrow d<sub>1</sub>, d<sub>2</sub>

meat \rightarrow d<sub>1</sub>, d<sub>2</sub>

Carnivores \rightarrow d<sub>2</sub>

Deers \rightarrow d<sub>3</sub>

eat \rightarrow d<sub>3</sub>

grass \rightarrow d<sub>3</sub>

leaves \rightarrow d<sub>3</sub>
```

2. Construct the TF-IDF term-document matrix for the corpus $\{d_1, d_2, d_3\}$.

	Counts, tfi				Weights, $w_i = tf_i \times IDF_i$		
Terms	d ₁	d ₂	d ₃	IDF _i =log ₁₀ (D/df _i)	dı	d ₂	d ₃
Herbivore	1	0	0	0.4771	0.4771	0	0
typically	1	1	0	0.1761	0.1761	0.1761	0
plant	1	1	0	0.1761	0.1761	0.1761	0
eaters	2	2	0	0.1761	0.3522	0.3522	0
meat	1	1	0	0.1761	0.1761	0.1761	0
Carnivore	0	1	0	0.4771	0	0.4771	0
Deers	0	0	1	0.4771	0	0	0.4771
eat	0	0	1	0.4771	0	0	0.4771
grass	0	0	1	0.4771	0	0	0.4771
leaves	0	0	1	0.4771	0	0	0.4771

3. Suppose the query is "plant eaters," which documents would be retrieved based on the inverted index constructed before?

Documents: d_1 , d_2

4. Find the cosine similarity between the query and each of the retrieved documents. Is the result desirable? Why?

Cosine Similarity calculations:

 $Q = \{0, 0, 0.1761, 0.1761, 0, 0, 0, 0, 0, 0, 0\}$

|Q| = 0.2490

 $|D_1| = 0.6669$

 $|D_2| = 0.6669$

 $|D_3| = 0.9542$

 $Q \cdot D_1 = 0.0930$

 $Q \cdot D_2 = 0.0930$

 $Q \cdot D_3 = 0$

 $sim(Q, D_1) = 0.56$

 $sim(Q, D_2) = 0.56$

 $sim(Q, D_3) = 0$

Ranking documents:

D₁/D₂ then followed by D₃ (tie between first two documents)

Is the ordering desirable? If no, why not?:

No. We are unable to determine which document to be prioritised among D_1 and D_2

1. Implement the retrieval component of the IR system in the template provided. Use the TF-IDF vector representation for representing documents.

```
u,s,vt = np.linalq.svd(tfidf); # u: txt, s: txd, vt = dxd
s_size = np.size(s) # 1400 = d
s_values = np.zeros([len_corpus, len_docs]) # txd
s_values[:s_size, :s_size] = np.diag(s) # dxd is diagonal
us = np.dot(u,s_values) # txd
tfidf1 = np.dot(us, vt) # txd
            totalSum = s.sum()
currSum = 0
count = 0
for elem in s:
   if (currSum/totalSum) > 0.65:
        break
            #print(count)
syvalues_k = s_values[:count, :count] # sxs
u_k = u|: :count| # txs
vt_k = vt|:count, :| # sxd
us_k = np.dot(u_k,s_values_k) # txs
tfidf_k = np.dot(us_k, vt_k) # txd
                     ine_sims = []
DS_doc in DS_docs_matrix: # 1xs
cosine_sim = spatial.distance.cosine(DS_query, DS_doc)
if np.linalg.norm(DS_doc) == 0:
    cosine_sim = 0
    else:
    cosine_sim = np.dot(DS_query, DS_doc)/((np.linalg.norm(DS_query)) *
.linalg.norm(DS_doc)))
            return cosine sims
```

1. Implement the following evaluation measures in the template provided (i). Precision@k, (ii). Recall@k, (iii). F_{0.5} score@k, (iv). AP@k, and (v) nDCG@k.

Precision@k: Computation of precision of the Information Retrieval System at a given value of ${\bf k}$ for a single query $\label{lem:condition} $$\operatorname{retAndRelevant} = \operatorname{list(set(query_doc_IDs_ordered[:k]) \& set(true_doc_IDs))$$ precision = \operatorname{len(retAndRelevant)} / k$$ return precision$ def meanPrecision(self, doc_IDs_ordered, query_ids, qrels, k): Computation of precision of the Information Retrieval System at a given value of k, averaged over all the queries arg1 : list A list of lists of integers where the ith sub-list is a list of IDs of documents in their predicted order of relevance to the ith query arg2 : list A list of IDs of the queries for which the documents are ordered 3: list A list of dictionaries containing document-relevance judgements - Refer cran_grels.json for the structure of each float The mean precision value as a number between 0 and 1 qRelsDict = {} for qRel in qrels: if qRel["query_num"] in qRelsDict: qRelsDict[qRel["query_num"]].append(int(qRel["id"])) else: qRelsDict[qRel["query_num"]] = [int(qRel["id"])] for i in range(0, len(query_ids)): totalPrecision = totalPrecision + self.queryPrecision(doc_IDs_ordered[i], query_ids[i], list(qRelsDict[str(query_ids[i])]), k) meanPrecision = totalPrecision / len(query_ids) return meanPrecision

Recall@k:

```
Computation of recall of the Information Retrieval System at a given value of k for a single query
        #Filt in code here
retAndRelevant = list(set(query_doc_IDs_ordered[:k]) & set(true_doc_IDs))
recall = len(retAndRelevant) / len(true_doc_IDs)
return recall
def meanRecall(self, doc_IDs_ordered, query_ids, qrels, k):
        Computation of recall of the Information Retrieval System at a given value of k, averaged over all the queries
      arg1 : list

A list of lists of integers where the ith sub-list is a list of IDs of documents in their predicted order of relevance to the ith query arg2 : list

A list of IDs of the queries for which the documents are ordered arg3 : list

A list of dictionaries containing document-relevance judgements - Refer cran_qrels.json for the structure of each dictionary arg4 : int

The k value
        totalRecall = 0
qRelsDict = {}
for qRel in qrels:
   if qRel["query_num"] in qRelsDict:
        qRelsDict[qRel["query_num"]].append(int(qRel["id"]))
        for i in range(0, len(query_ids)):
    totalRecall = totalRecall + self.queryRecall(doc_IDs_ordered[i], query_ids[i], list(qRelsDict[str(query_ids[i])]), k)
meanRecall = totalRecall / len(query_ids)
return meanRecall
```

F_{0.5} score@k:

```
Computation of fscore of the Information Retrieval System at a given value of k for a single query
        precision = self.queryPrecision(query_doc_IDs_ordered, query_id, true_doc_IDs, k)
recall = self.queryRecall(query_doc_IDs_ordered, query_id, true_doc_IDs, k)
if precision == 0 and recall == 0:
    reture
        return 0

fscore = (2*precision*recall)/(precision+recall)
return fscore
def meanFscore(self, doc_IDs_ordered, query_ids, qrels, k):
        Computation of fscore of the Information Retrieval System at a given value of k,\ \mbox{averaged} over all the queries
       arg1 : list

A list of lists of integers where the ith sub-list is a list of IDs of documents in their predicted order of relevance to the ith query arg2 : list

A list of IDs of the queries for which the documents are ordered arg3 : list

A list of dictionaries containing document-relevance judgements - Refer cran_qrels.json for the structure of each dictionary arg4 : int
        totalFScore = 0
qRelsDict = {}
for qRel in qrels:
    if qRel["query_num"] in qRelsDict:
        qRelsDict[qRel["query_num"]].append(int(qRel["id"]))
                 qnctsDict(qRel["query_num"]] = [int(qRel["id"])]
qRelsDict[qRel["query_num"]] = [int(qRel["id"])]
        for i in range(0, len(query_ids)):
    totalFScore = totalFScore + self.queryFscore(doc_IDs_ordered[i], query_ids[i], list(qRelsDict[str(query_ids[i])]), k)
    meanFScore = totalFScore / len(query_ids)
    return meanFScore
```

AP@k:

```
Computation of average precision of the Information Retrieval System at a given value of k for a single query (the average of precision@i values for i such that the ith document is truly relevant)
                avgPrecision = -1
               retervantcount = 0
count = 0
totalPrecision = 0
for predictedRes in query_doc_IDs_ordered[:k]:
    count = count + 1
    if predictedRes in true_doc_IDs:
        relevantCount = relevantCount + 1
        totalPrecision = totalPrecision + (relevantCount/count)
avgPrecision = totalPrecision/(len(true_doc_IDs))
return avgPrecision
        def meanAveragePrecision(self, doc_IDs_ordered, query_ids, q_rels, k):
                Computation of MAP of the Information Retrieval System at given value of k, averaged over all the queries
               arg1: list
A list of lists of integers where the ith sub-list is a list of IDs of documents in their predicted order of relevance to the ith query arg2: list
A list of IDs of the queries arg3: list
A list of dictionaries containing document-relevance judgements - Refer cran_qrels.json for the structure of each dictionary arg4: int
                float
The MAP value as a number between 0 and 1
                meanAveragePrecision = -1
                totalAveragePrecision = 0
               totalAveragePrecision = 0
qRelsDict = {}
for qRel in q_rels:
    if qRel["query_num"] in qRelsDict:
        qRelsDict[qRel["query_num"]].append(int(qRel["id"]))
                       qnetsbirt(q
else:
    qRelsDict[qRel["query_num"]] = [int(qRel["id"])]
```

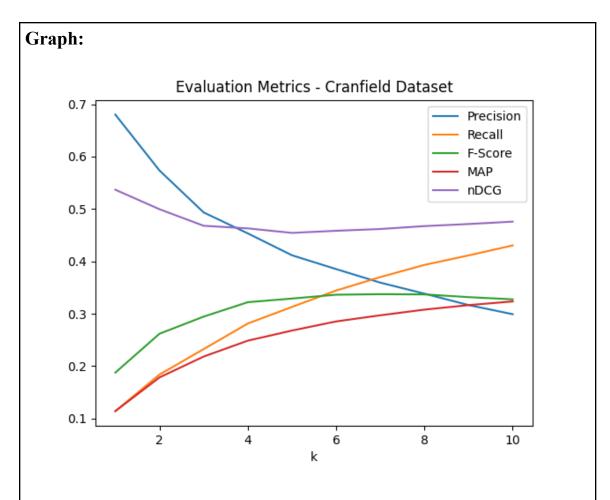
nDCG@k:

```
def queryNDCG(self, query_doc_IDs_ordered, query_id, true_doc_IDs, k):
          Computation of nDCG of the Information Retrieval System at given value of k for a single query
          for predictedRes in query_doc_IDs_ordered[:k]:
    if predictedRes in true_doc_IDs[0]:
        dcg = dcg + (true_doc_IDs[1][true_doc_IDs[0].index(predictedRes)]/math.log(count+1,2))
    count = count + 1
         count = count + 1
idcg = 0
count = 1
for idealRes in true_doc_IDs[0][:k]:
    idcg = idcg + (true_doc_IDs[1][count-1]/math.log(count+1,2))
    count = count+1
nDCG = dcg/idcg
          return nDCG
def meanNDCG(self, doc_IDs_ordered, query_ids, qrels, k):
          Computation of nDCG of the Information Retrieval System at a given value of \mathbf{k}\text{,} averaged over all the queries
          A list of IDs of the queries for which the documents are ordered

A list of IDs of which the documents are ordered
                   3: list
A list of dictionaries containing document-relevance
judgements - Refer cran_qrels.json for the structure of each
dictionary
                    The mean nDCG value as a number between 0 and 1
          meanNDCG = -1
         totalNDCG = 0
qRelsDict = {}
for qRel in qrels:
    if qRel["query_num"] in qRelsDict:
        qRelsDict[qRel["query_num"]][0].append(int(qRel["id"]))
        qRelsDict[qRel["query_num"]][1].append(5 - int(qRel["position"]))
        queristrict[query_num ]] [[int(qRel["id"])], [5 - int(qRel["position"])]

for i in range(0, len(query_ids)):
    sortedIndex = np.argsort(list(qRelsDict[str(query_ids[i])]) [1])
    docIdList = []
    relList = []
    trueDocIds = []
    for ind in sortedIndex[::-1]:
        docIdList.append(list(qRelsDict[str(query_ids[i])]) [0][ind])
        relList.append(list(qRelsDict[str(query_ids[i])]) [0][ind])
        relList.append(list(qRelsDict[str(query_ids[i])]) [1][ind])
    trueDocIds.append(docIdList)
    trueDocIds.append(relList)
    totaINDCG = totaINDCG / self.queryNDCG(doc_IDs_ordered[i], query_ids[i], trueDocIds, k)
meanNDCG = totalNDCG / len(query_ids)
return meanNDCG
```

2. Assume that for a given query, the set of relevant documents is as listed in incran_qrels.json. Any document with a relevance score of 1 to 4 is considered as relevant. For each query in the Cranfield dataset, find the Precision, Recall, F-score, average precision, and nDCG scores for k = 1 to 10. Average each measure over all queries and plot it as a function of k. The code for plotting is part of the given template. You are expected to use the same. Report the graph with your observations based on it.



Observation:

We compared Precision@k and nDCG@k for values k=1 to k=10 to compare our approach with other approaches mentioned in this work. We chose to use Precision and nDCG measures, since it is search engine application and in this case, precision is a bit more important than recall. It is not required for the system to retrieve all relevant documents. Instead it should return the most relevant documents in the first few results. nDCG is another good measure to compare because it also takes into account the graded relevance of the document to the query. Since we have access to the query-document relevance judgements for this dataset, we can leverage these scores to calculate nDCG scores.

3. Using the time module in Python, report the run time of your IR system.

Total time taken = 6 minutes 20 seconds

1. What are the limitations of such a Vector space model? Provide examples from the cranfield dataset that illustrate these shortcomings in your IR system.

Limitations:

- 1. word orders are not taken into account
- 2. Every coordinate is assumed to be independent of each other.

Examples from your results:

[Refer to the doc: Here]

Part 4: Improving the IR system

Based on the factual record of actual retrieval failures you reported in the assignment, you can develop hypotheses that could address these retrieval failures. You may have to identify the implicit assumptions made by your approach that may have resulted in undesirable results. To realize the improvements, you can use any method(s), including hybrid methods that combine knowledge from linguistic, background, and introspective sources to represent documents. Some examples taught in class are Latent Semantic Analysis (LSA) and Explicit Semantic Analysis (ESA).

You can also explore ways in which a search engine could be improved in aspects such as its efficiency of retrieval, robustness to spelling errors, ability to auto-complete queries, etc.

You are also expected to test these hypotheses rigorously using appropriate hypothesis testing methods. As an outcome of your work, you should be able to make a statement of structure similar to what was presented in the class:

An algorithm A_1 is better than A_2 with respect to the evaluation measure E in task T on a specific domain D under certain assumptions A.

Note that, unlike the assignment, the scope of this component is open-ended and not restricted to the ideas mentioned here. For each method, the final report must include a critical analysis of results; methods can be combined to come up with improvisations. It is advised that such hybrid methods are well founded on principles and not just ad hoc combinations (an example of an ad hoc approach is a simple convex combination of three methods with parameters tuned to give desired improvements).

You could either build on the template code given earlier for the assignment or develop from scratch as demanded by your approach. Note that while you are free to use any datasets to experiment with, the Cranfield dataset will be used for evaluation. The project will be evaluated based on the rigor in

methodology and depth of understanding, in addition to the quality of the report and your performance in Viva.

Your project report (for Part 4) should be well structured and should include the following components.

- 1. An introduction to the problem setting,
- 2. The limitations of the basic VSM with appropriate examples from the dataset(s),
- 3. Your proposed approach(es) to address these issues,
- 4. A description of the dataset(s) used for experimentations,
- 5. The results obtained with a comparative study of your approach has improved the IR system, both qualitatively and quantitatively.

The latex template for the final report will be uploaded on Moodle. You are instructed to follow the template strictly.