INFORMATION RETRIEVAL

HOMEWORK EXERCISES LOS. NEURAL IR

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GENERL INSTRUCTIONS

- You are going to do a practical exercise with ColBERT reranking in PyTerrier (A Python framework for performing information retrieval experiments, building on Terrier)
- Submit only the added code snippets and requested output as PDF, don't submit the whole notebook
- If you need help, you can contact the TAs at ircourse@liacs.leidenuniv.nl
- The goal of this exercise are
 - 1. to learn to recognize the steps of a retrieval and ranking pipeline;
 - 2. to understand the data structures necessary for the pipeline;
 - 3. to learn to interpret model output



EXERCISE PREPARATIONS

Preparation:

- Start a new Python notebook on Google colab
- Make sure that the notebook uses a GPU (under Resources -> Change runtime type)
- 3. Follow the steps in this tutorial (3.2 only):

 https://github.com/terrier-org/ecir2021tutorial/blob/main/notebooks/notebook3.2.ipynb
 and make sure everything runs and you get output



EXERCISE 1

- Let's first collect the results
- In the Experiment function, ColBERT is compared to DPH, the model that is used as default first-stage retriever in Terrier [1].
- 1. Show the results table with DPH and DPH >> ColBERT as two rows
- Look up in the PyTerrier documentation [2] how you can replace DPH by BM25. Adapt this in your code and show the result table for BM25 and BM25 >> ColBERT
- 3. Which do you prefer, DPH+ColBERT, or BM25+ColBERT?



EXERCISE 1 - SOLUTIONS

1. Results table with DPH and DPH >> ColBERT

	name	map	ndcg	ndcg_cut.10	P.10	mrt
0	DPH	0.068056	0.165653	0.609058	0.658	49.933852
1	DPH >> ColBERT	0.074727	0.172074	0.689785	0.750	689.051942



EXERCISE 1 - SOLUTIONS

```
br = pt.BatchRetrieve(index, wmodel="BM25") % 100
pipeline = br >> pt.text.get_text(dataset, 'abstract') >> colbert
pt.Experiment(
    [br, pipeline],
    topics,
    qrels,
    names=['BM25', 'BM25 >> ColBERT'],
    eval_metrics=["map", "ndcg", 'ndcg_cut.10', 'P.10', 'mrt']
)
```

2. Results table with BM25 and B25 >> ColBERT

	name	map	ndcg	ndcg_cut.10	P.10	mrt
0	BM25	0.077892	0.177767	0.644374	0.692	49.320280
1	BM25 >> ColBERT	0.081463	0.180422	0.666929	0.738	688.697242



EXERCISE 1 - SOLUTIONS

- It depends on the use case:
 - MAP and nDCG on the full ranking are higher for BM25. Thus, a user who inspects all 100 results will get more relevant information with BM25+ColBERT than with DPH+ColBERT
 - nDCG@10 and P@10 are higher for DPH, indicating a higher effectiveness by BM25 for the top-10 retrieved documents. Thus, a user who only inspects the top-10 will get more relevant information with DPH+ColBERT than with BM25+ColBERT



EXERCISE 2

The test collection used in the tutorial is TREC-COVID https://ir.nist.gov/covidSubmit/index.html

We are going to explore the topic collection a bit.

- Add code to your notebook to print the first 5 topics from the test collection.
- 2. Look up in the PyTerrier documentation [3] how you can view the results per query. Show the query (id and content) with the lowest nDCG@10 and the queries with the highest nDCG@10.



EXERCISE 2 - SOLUTIONS

1. topics.iloc[0:5]

index	qid	query	
0	1	what is the origin of covid 19	
1	2	how does the coronavirus respond to changes in the weather	
2	3	will sars cov2 infected people develop immunity is cross protection possible	
3	4	what causes death from covid 19	
4	5	what drugs have been active against sars cov or sars cov 2 in animal studies	



EXERCISE 2 - SOLUTIONS

Result per query:

```
br = pt.BatchRetrieve(index, wmodel="BM25") % 100
pipeline = br >> pt.text.get_text(dataset, 'abstract') >> colbert
result = pt.Experiment(
    [pipeline],
    topics,
    qrels,
    names=['BM25 >> ColBERT'],
    eval_metrics=['ndcg_cut.10'],
    perquery=True
)
result.sort_values('value')
```

- qid 31 has the lowest nDCG@10: 0.151771
- qids 42, 20, 40, 22, 38 have the highest nDCG@10: 1.00



EXERCISE 2 - SOLUTIONS

- > 31: how does the coronavirus differ from seasonal flu
- > 42 : does vitamin d impact covid 19 prevention and treatment
- 20 : are patients taking angiotensin converting enzyme inhibitors ace at increased risk for covid 19
- ➤ 40 : what are the observed mutations in the sars cov 2 genome and how often do the mutations occur
- 22 : are cardiac complications likely in patients with covid 19
- > 38: what is the mechanism of inflammatory response and pathogenesis of covid 19 cases



EXERCISE 3

- 1. Run the retrieval pipeline for the query with the lowest nDCG@10 and one of the queries with the highest nDCG@10. What is the highest ranked document? Is it relevant or not?
- 2. Output the attention matrix for the document retrieved in first position for the query with the lowest nDCG@10 and for the document retrieved in first position for the query with the highest nDCG@10. What does it tell you?

EXERCISE 3 - SOLUTIONS

1. Irrelevant for the query (31) with the lowest nDCG (label 0)

```
res1 = pipeline(topics.iloc[30:31])
res1.merge(dataset.get_qrels(), how='left')

qid query docno score rank label iteration

0 31 how does the coronavirus differ from seasonal flu luloic87 21.411621 0 0.0 4
```

Relevant for the query (42) with the highest nDCG (label 2)

```
res1 = pipeline(topics.iloc[41:42])
res1.merge(dataset.get_qrels(), how='left')

qid query docno score rank label iteration

0 42 does vitamin d impact covid 19 prevention and ... wr9hkvd3 24.390995 0 2.0 5
```

EXERCISE 3 - SOLUTIONS

2. Attention analysis

```
text = dataset.irds_ref().docs_store().get('luloic87').abstract[:300] + '...' # truncate text
colbert_factory.explain_text('how does the coronavirus differ from seasonal flu', text)
```

```
text = dataset.irds_ref().docs_store().get('wr9hkvd3').abstract[:300] + '...' # truncate text
colbert_factory.explain_text('does vitamin d impact covid 19 prevention and treatment', text)
```



p q 31 (left): the highest attention to the general tokens 'corona virus'; there is no relation to flu in the abstract

q 42 (right): the highest attention to the relevant tokens 'vitamin d' and 'influences'



