Assignment 3

Assignment 3

Documents All Pairs Similarity

```
(usefull for ex. in duplicate search)
or span removal
```

- To Be Delivered:
 - Sequential implementation (Python-Numpy/Java/Scala)
 - Parallel implementation
 - MapReduce / Apache Spark (Python/Java/Scala)
 - Report discussing performance figures of the proposed parallel implementation
 - varying datasets (and samples), similarity thresholds
 - uarying number of workers waching
 - max 2 pages

```
(LOOK AT THE PREVIOUS REPORT
```

Documents All Pairs Similarity

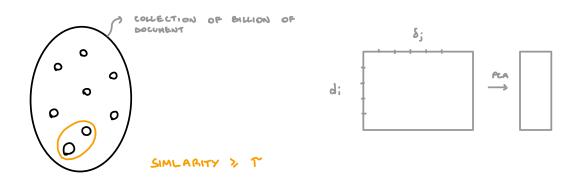
- A document is a vector d of N elements
 - N is the number of distinct words in the corpus (the lexicon)
 - d[i]: stores the frequency of the term i in document d (tf(i)).
 Then d is normalized (divided by its L₂ norm)
 - additionally you can use tf-idf:

$$tf-idf(i) = tf(i) \cdot \ln \frac{N_{docs}}{df(i)}$$

- There are many ways to measure similarity
 - Cosine:

$$s(a,b) = \sum_{i=1...N} a[i] \cdot b[i]$$

DOCUMENT ALL PAIRS SIMILARITY SEARCH



What is a document?

Can be any object - I need to find a repr.

For movies dataset I have: m; 34 44 U;

Similarity can be computed using COSINE SIMILARITY that is basically a model dot product (di.dj) the lidilly score of a doc if a append to result

I want the
EXACT similarity
(but I can compare it
with approximations)

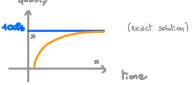
- cluster ooks and inside cluster
 do search (run alg. on clusters)
 (can be done on clusters in
 parallel)
- . L5H

(investigate the tradeoff error - speed)

· PLA or JOHNSON - LINDENSTROUD

I can plot the threadoff comparison





using for ex. Jackaro sin. CERNEEN

RESULTS (exact and approx.)

COMPUTING THE EXACT SOLUTION:

For each
$$d_i$$
:

For each d_i :

 $compute sim (d_i, d_i)$



THERE ARE SOME TECHNIQUES THAT SPEED UP THE COMPUTATION BUT NOT REDUCE COMPLEXITY

WE CAN ALSO SORT BY DOC LENGTH AND SKIP ALL THE DOC
SHORTER THAN A TRESHOLD BECAUSE THE DOT PROD SUEALY DON'T REACH THE VALUE
(SKIP A LOT OF DOC IN ONE PASS WITH SORTING, RATHER THEN SKIPPING THEM ONE AT A TIME)

Sequential Algorithm

Given the minimum required similarity threshold

```
\square SIM_DOCS = 0
```

- lacksquare For-each document $d_{_{\it I}}$ in the corpus D :
 - lacksquare For-each document d_2 in the corpus D :
 - \Box if $d_1!=d_2$ and $s(d_1, d_2)>=threshold:$
 - \square SIM_DOCS += 1
- Note: usually you are interested to the similar document pairs, rather than to the number of similar document
- Try your optimizations!!
- Datasets:
 - https://github.com/beir-cellar/beir
 - ☐ https://grouplens.org/datasets/movielens/

Deadline and Evaluation

- Delivery before May 26:
 - point if positively evaluated, +0.5 if sufficient, re-submit if insufficient
- Or, delivery at written exam
 - +1 point if positively evaluated, +0 if sufficient, exam
 not passed if insufficient
- Positive Evaluation means:
 - good report, good code, good analysis