#### Homework 1

Data Mining Technology for Business and Society
Deadline: 28 April 2020 23:59 (Rome Time Zone)
Having TWO students per group is RECOMMENDED.

Having one student per group is allowed only if coordination is not possible.

The total length of the report cannot exceed 5 pages.

It is forbidden to submit software written with Python-Notebook.

Only ".py" software is considered as a valid solution .

The software must be commented .

Data and software are available at:

http://www.diag.uniroma1.it/~fazzone/Teaching/Data\_Mining\_Technology\_for\_Business\_and\_Society 2019 2020/DMT4BaS 2019 2020.html

The homework is composed of two parts: Search-Engine Evaluation and Near-Duplicates-Detection.

### Part 1

You have to index two collections of documents and improve the search-engines performance by changing their configurations. Using the provided sets of queries, and the associated Ground-Truths. For this part of the homework you must use the Whoosh API.

#### The Two Collections of Documents

The two different collections of documents are: Cranfield\_DATASET and Time\_DATASET .

They consist of:

- ..) a set of html documents.
- ..) a set of queries.
- ..) a set of relevant documents identifiers for each of a subset of the guery set the Ground-Truth.

The ground truth does not provide the set of relevant documents for all queries in the query set.

# **Documents, Queries and Ground-Truth**

The documents to index are stored in html files and they are composed of two fields: content and title (please, open them with a text-editor and not with a browser). The content of the "title" field is located between the "<title>" tags and the content of the "content" field is located between the "<body>" tags. The document-id is the integer number at the end of the html file name. For instance, for the Cranfield\_DATASET, the file with name "\_\_\_\_\_\_42.html" contains the document with ID " 42", title " the gyroscopic effect of a rigid rotating propeller..." and content " in many wing vibration analyses it is found necessary...". All documents are stored inside the "DMT/HW\_1/part\_1/<COLLECTION\_NAME>/DOCUMENTS" directories.

Queries are stored in the "DMT/HW\_1/part\_1/<COLLECTION\_NAME>/<COLLECTION\_NAME>\_Queries.tsv" file and the ground-truth is stored inside the "

DMT/HW 1/part 1/<COLLECTION NAME>/<COLLECTION NAME> Ground Truth.tsv" file.

These two files are linked by the "Query id" field value.

An Important consideration for Time DATASET.

The content of the field "title" is not informative . The content of this field must not be taken into consideration.

## **Evaluation Metrics**

For each configuration, you must provide the following "MRR table":

Search Engine Configuration	MRR
conf_x	?.???
conf_y	?.???
conf_z	?.???
•••	?.???

Only for the Top-5 configurations in the "MRR table" (the ones with the best five MRR values), you must provide the following information:

# .) "R-Precision distribution table", with the following format:

Search Engine Configurati on	Mean (R-Precisio n_Distrbuti on)	min(R-Preci sion_Distrb ution)	1°_quartile (R-Precisio n_Distrbuti on)	MEDIAN(R-Pr ecision_Dis trbution)	3°_quartile (R-Precisio n_Distrbuti on)	MAX(R-Preci sion_Distrb ution)
conf_w	?.???	?.???	?.???	?.???	?.???	?.???
conf_t	?.???	?.???	?.???	?.???	?.???	?.???
conf_z	?.???	?.???	?.???	?.???	?.???	?.???
	?.???	?.???	?.???	?.???	?.???	?.???

- .) The "P@k plot", where:
- .) the x axis represents the considered values for k: you must consider k  $\epsilon$  {1, 3, 5, 10}
- .) the y axis represents the average (correctly normalized) P@k over all provided queries.
- .) Each curve represents one of the Top-5 search engine configurations (according to the "MRR table").
- .) The "nDCG@k plot ", where:
- .) the x axis represents the considered values for k: you must consider k  $\in$  {1, 3, 5, 10}
- .) the y axis represents the average nDCG over all provided queries.
- .) Each curve represents one of the Top-5 search engine configurations (according to the "MRR table").

#### Information to Provide in the Report

For both Cranfield\_DATASET and Time\_DATASET, you have to provide in the report the following information:

- .) Number of indexed documents and the number of queries.
- .) Number of queries in the Ground-Truth.
- .) A schematic description of all tested search engine configurations.
- .) The "MRR table" for all tested search engine configurations.
- .) The set of all Top-5 search engine configurations according to the "MRR table".
- .) The "R-Precision distribution table" with data from the Top-5 search engine configurations according to the "MRR table".
- .) The "P@k plot" with data from the Top-5 search engine configurations according to the "MRR table".
- .) The "nDCG@k plot" with data from the Top-5 search engine configurations according to the "MRR table". You must provide all this information in at most three pages.

#### Part 2

You have to find, in an approximated way, all near-duplicate documents inside the following dataset: /DMT/HW\_1/part\_2/dataset/250K\_lyrics\_from\_MetroLyrics.csv.

The dataset contains data on 250K songs.

Two songs are considered near-duplicates if, and only if, the Jaccard similarity between their associated sets of shingles computed only on their lyrics is  $\geq 0.89$ .

To complete this part of the homework, you have to use the Near\_Duplicates\_Detection\_Tool that is entirely contained inside the directory " DMT/HW\_1/part\_2/tools ". The file

"DMT/HW\_1/part\_2/script\_for\_testing.txt" contains a short description and an example on how to run the Near Duplicates Detection Tool. Moreover, the file

"DMT/HW\_1/part\_2/dataset/1K\_\_test\_sets\_for\_LSH.tsv" contains a representation of 1000 documents as sets of shingle\_IDs and can be used only for testing the

Near\_Duplicates\_Detection\_Tool. For creating hash functions you can use the following software: "DMT/HW\_1/part\_2/hash\_functions\_creator.py".

# **Details on Shingling**

For representing a song as a set of shingles identifiers in a correct way, you have to assign a natural number IDENTIFIER to each distinct shingle you generated by processing all 250K documents . I suggest you use as shingle identifier a natural number that spans from 0 to the number of distinct shingles you generated minus one: 0, 1, 2, 3, ... , number\_of\_all\_observed\_distinct\_shingles-1 . Before shingling a document, it is required to remove punctuations and convert all words in lower-case, moreover, stopword removal, stemming and lemmatization are forbidden . The length of each shingle must be 3 . You have to shingle only the lyric of the song.

## **Details on Sketching**

Constraint 1: Each set of shingles, that represents an original document, must be sketched in a Min-Hashing sketch with a length of at most 300.

# **Details on LSH**

Constraint 2 : The probability to have as a near-duplicate candidate a pair of documents with Jaccard=0.89 must be > 0.97 .

## <u>Information to Provide in the Report</u>

You have to provide in the report the following information:

- .) The number of rows and the number of bands that you chose.
- .) The probability to have False-Negatives, in the set of candidate pairs, for the following Jaccard values: 0.89, 0.9, 0.95 and 1.
- .) The probability to have False-Positives, in the set of candidate pairs, for the following Jaccard values: 0.85, 0.8, 0.75, 0.7, 0.65, 0.6, 0.55 and 0.5.
- .) How did you reduce the probability to have False-Negatives?
- .) The Execution-Time of the Near-Duplicates-Detection tool.
- .) The number of Near-Duplicates couples you found.
- .) The number of Near-Duplicates couples you found with an approximated Jaccard similarity value of at least 0.89, 0.90, 0.91, 0.92, 0.93, 0.94, 0.95, 0.96, 0.97, 0.98, 0.99, 1.

You must provide all these information in at most two pages.

# Where/What To Send

At the end of the process, you have to create a zip file with ONLY the following data:

- 1. The software for addressing Part\_1: /DMT\_2020/HW\_1/part\_1/sw/ ( .py files ).
- 2. The software for addressing Part\_2: /DMT\_2020/HW\_1/part\_2/sw/ ( .py files ).
- 3. The tsv file you created for addressing Part\_2 that contains the sets of shingles identifies: /DMT 2020/HW 1/part 2/data/ (.tsv files).

4. The tsv file containing the Near-Duplicates you found for Part\_2:

/DMT\_2020/HW\_1/part\_2/data/ (.tsv files).

5. The final report in PDF: /DMT\_2020/HW\_1/report.pdf.

The name of the zip file must have this format:

DMT\_2020\_\_HW\_1\_\_StudentID\_StudentName\_StudentSurname\_StudentID\_StudentName\_StudentSurname .zip

Finally you must send the ".zip" file to fazzone@diag.uniroma1.it with the following email subject :

 ${\tt DMT\_2020\_HW\_1\_StudentID\_StudentName\_StudentSurname\_StudentID\_StudentName\_StudentSurname\_StudentID\_StudentName\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurname\_StudentSurn$