Final for the Biomedical Information Retrieval Course

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Environment

- Python3
- Flask
- nltk
- gensim

Github

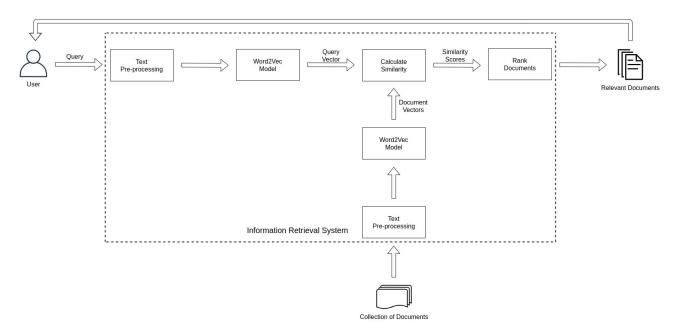
https://github.com/yyyyuwen/BIR-course-final

功能

- Data pre-Processing
 - 1. 讀檔:

讀取關鍵字為covid跟pneumonia跟heart disease的文章各1000 篇

- 2. 將每一個word做前處理: Stop Word -> Lemmatizer
- Search & rank by cosine similarity



- 文本聚類 (text clustering)

- tf-idf計算每篇文章的詞向量
- Kmeans進行文本聚類分析
- · TSNE降維與視覺化
 - Word2Vec

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Cluster 1: heart, ventricular, echocardiography, coronary, myocardial Cluster 2: cov, test, assay, sample, rt, detection Cluster 3: covid19, sars, pcr, coronavirus, infection Cluster 4: pneumoniae, mycoplasma, infection, pathogen, case Cluster 5: diagnosis, lung, clinical, chest, pulmonary, bacterial
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將全部的文章集合起來做關鍵字的cosine similarity,為了分析,我們取了heart, coronavirus, pneumonia, fever, cough,經過TSNE降維可以看出來covid與pneumonia的距離比跟heart的還要近,因此可以推測出covid與pneumonia的關聯度較高。