

The Problem.

Historically the vast amount of knowledge that experts publish has been increasing in such a pace that keeping up to date and having a full perspective, even in particular topics, has become quite challenging.

Such is the case of the current COVID-19 pandemic where there are so many clinical notes, experiments, expert observations around the world that doctors, researchers, and public authorities struggle to explore pieces of related but not explicitly connected knowledge concerning to their respective duties.

HOW WE TACKLE IT

A EXPLORATION ENVIRONMENT. We propose a smart literature analysis environment, which includes several NLP-powered components to enable a more efficient reading process. The following two strategies are the core of our environment.

TRANSVERSAL READING. We propose a semantically-guided transversal reading. We believe that this type of reading can significantly benefit the process of grasping the prominent opinion and state-of-the-art of a particular aspect. Our strategy to provide this feature was to interlink all semantically related sentences by semantic-textual-similarity (STS).

SEMANTIC ENRICHMENT. We enrich the literature with named-entity recognition and disambiguation (NERD), using the major life science databases as entity sources, enable named-entity searches, provide network-graphs of the most interconnected publications and, an interactive tool to highlight the most central statements within an article.



A smart literature exploration environment for COVID-19 literature.

<http://covid19.ccg.unam.mx:82/>



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Methodology.

NAMED ENTITY RECOGNITION AND DISAMBIGUATION (NERD). These capabilities are provided by OGER, a state-of-the-art biomedical NER annotator which in turn depends on the Bio Term Hub (BTH). BTH is a combined terminological resource created by dynamically sourcing entity names and their identifiers from reference databases.



The OntoGene's Biomedical Entity Recogniser (OGER) is a RESTful web service implemented on top of the BTH which allows a remote user to batch annotate a collection of documents.

SEMANTIC TEXTUAL SIMILARITY (STS). Our approach to measure STS is representing the sentences as embeddings and then use the cosine between two embeddings as their semantic similarity.

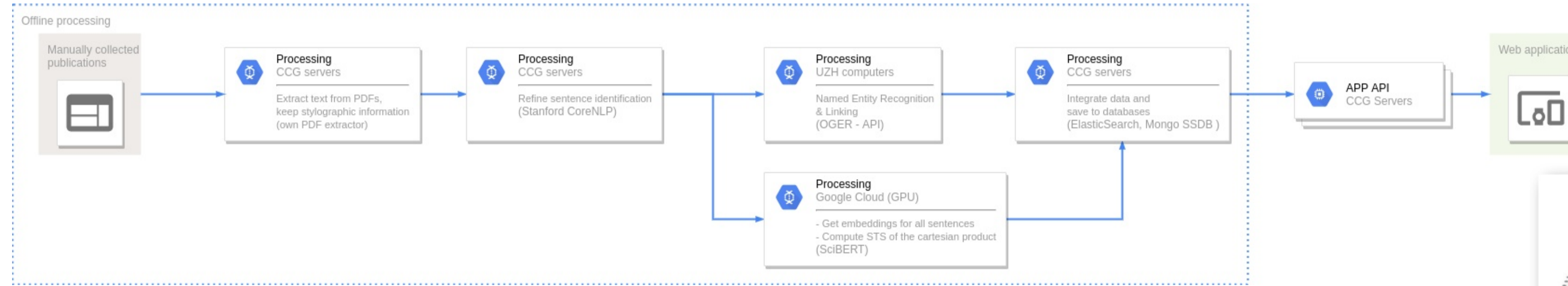
To compute the embeddings we used SciBERT, an unsupervised transformer language model pre-trained in the scientific literature. First, we map tokens to embeddings and then apply mean pooling to get fixed-sized sentence vectors.

Due to the lack of STS corpora specific to the COVID-19 literature we did not apply any fine-tuning..

Results.

Extractive summarization

- Select the sentences based in their centrality within the publication content
- Strategy inspired on the Textual Energy¹ of the sentence interactions



Browse articles through semantically similar sentences

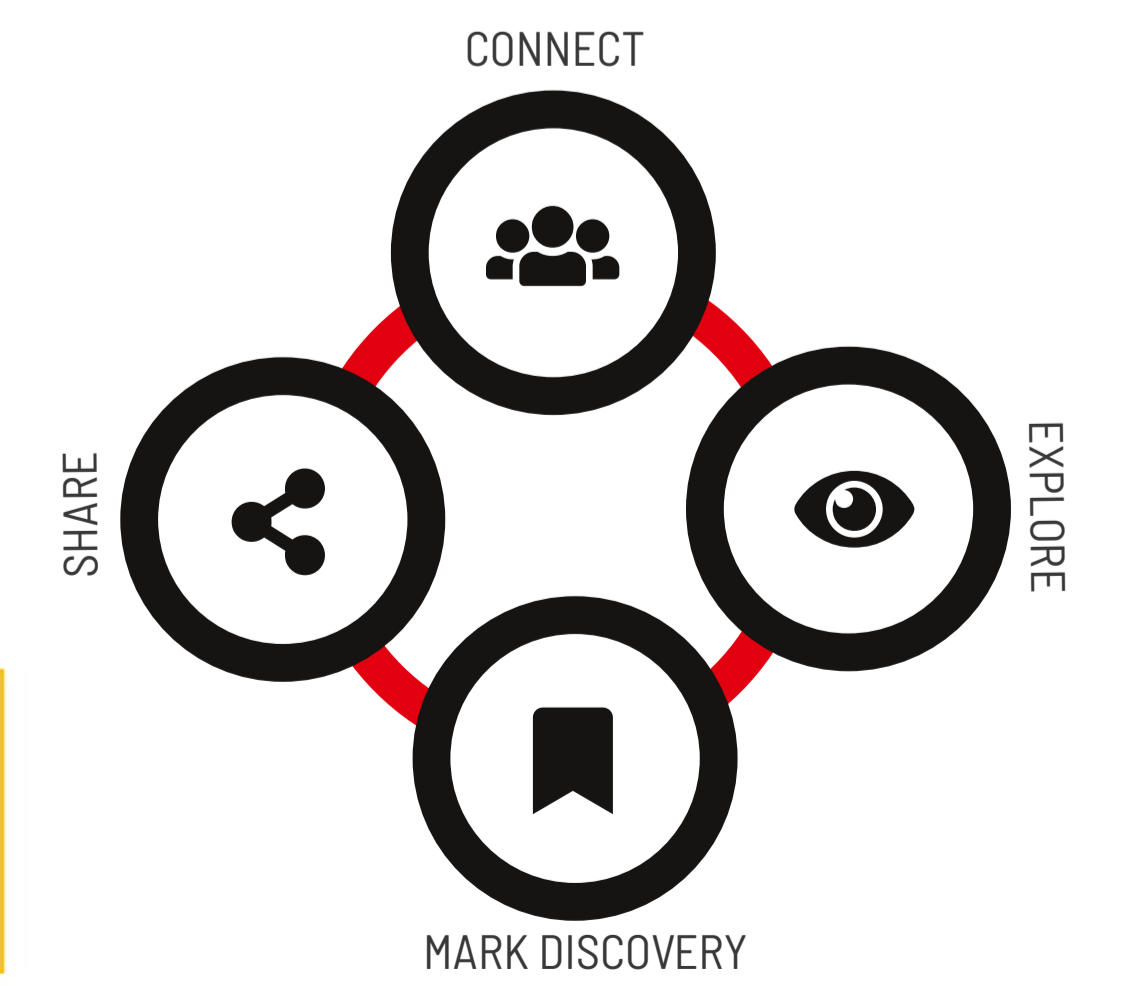
	Pearson	Spearman	Model
0	0.645213	0.620046	Distilled Bert
1	0.686426	0.742812	SciBert
2	0.428662	0.487770	InterSent GloVe 6B
3	0.468821	0.574450	InterSent GloVe RegulonLit
4	0.53823	0.583925	InterSent GloVe-840B

- Offline unsupervised STS (SciBert)
- Indirect validations on a Microbial Transcriptional Regulation corpus

- Interactive directed graph based on STS
- Edges between 2 publications are the sum of the STSs among their sentences

Searches on content and metadata

- Uses the ElasticSearch engine
- OGER annotations are transformed to ES annotation format
- User queries are transformed to ES_span-near queries



DISCOVERY PATHS