

Text Classification on COVID-19: a Transformer-based Approach

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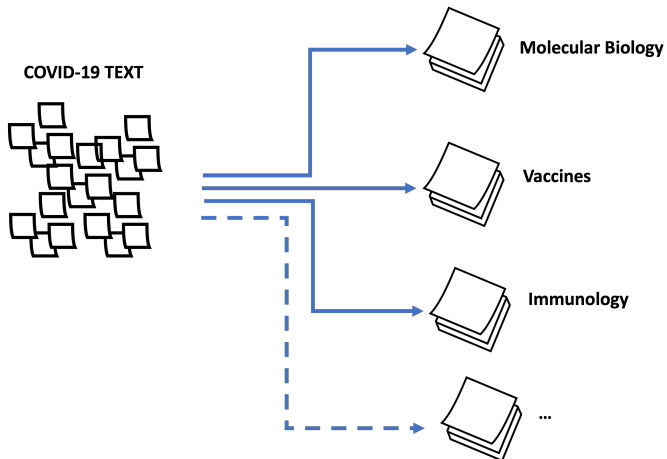
▶ Background

▶ Methodology

▶ Results

▶ Summary

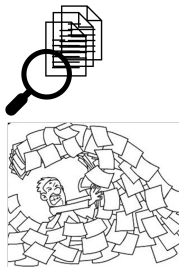
Text classification aims to assign labels or tags to textual units, such as sentences, queries, and documents.



The **COVID-19 pandemic** has led to an **unprecedented amount of scientific publications**, growing at a pace never seen before.



IDENTIFY



SELECT AND ASSESS

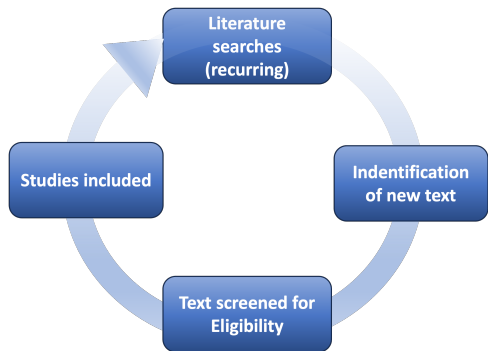


SYNTHESIZE AND FILTER

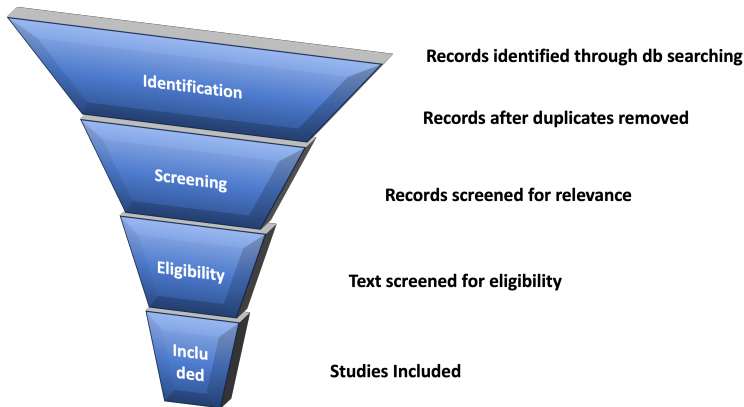


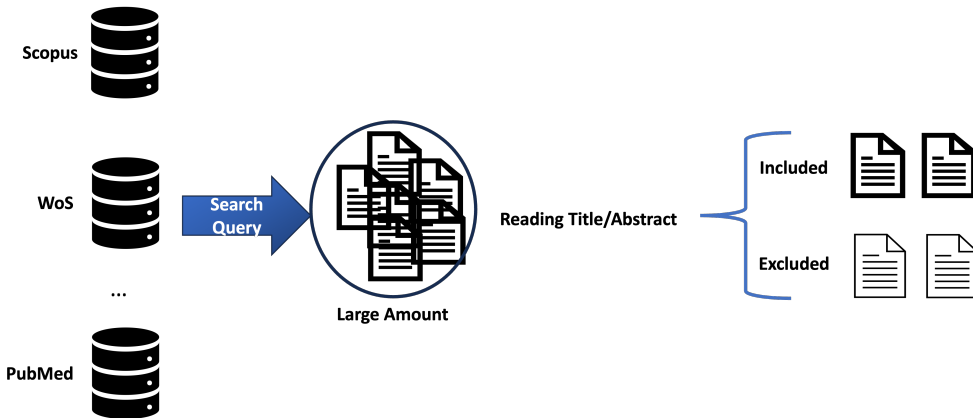
| PRO | CONS |
|---|---|
| Trasparent | Time consuming and resource intensive |
| Replicable & Updatable | Unpredictable |
| Reduce bias | Dependent on primary studies |
| Provide best evidence for decision making | Findings not directly relevant |
| Findings easily assessable | Commissioning is costly |
| Can improve future research design | Difficult to locate relevant unpublished research |

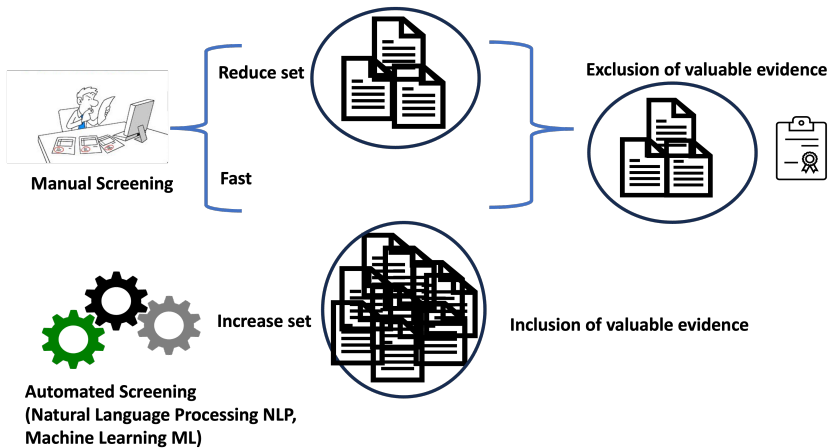
- Allows updating information as soon as new evidence becomes available.
- Living evidence can narrow the gap between knowledge and practice.
- Maintaining living evidence systems requires ongoing manual curation by skilled human resources.
- Title and/or abstract screening is still one of the most time-consuming tasks.



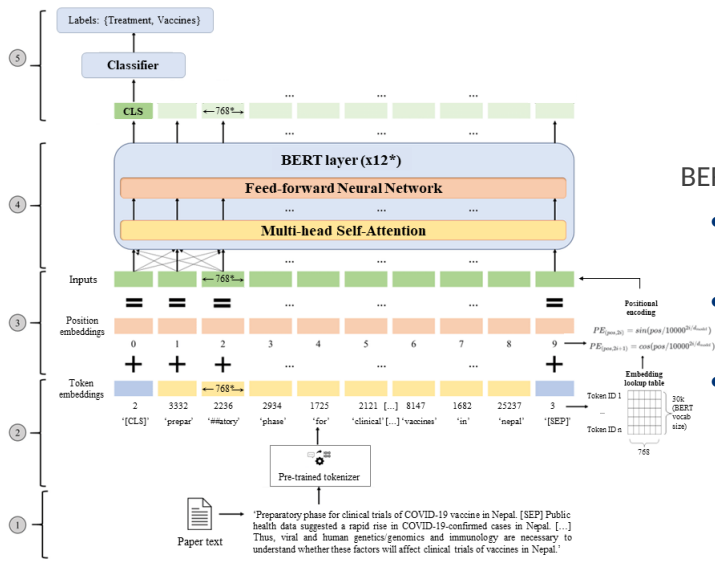
The **Preferred Reporting Items for Systematic reviews and Meta-Analyses (PRISMA)** is a reporting **guideline** that comprises a checklist of 27 items and a **four-phase** flow diagram recommended **for** reporting in **SR**.







- To streamline the **screening phase**, we have employed pre-trained transformer models, such as BioBERT and PubMedBERT derived from Bidirectional Encoder Representations from Transformers (BERT).
- Transformers are based on **pre-training** and **self-attention**:
 - Pre-training:
 - cheap training time and data size for fine tuning.
 - mitigates overfitting problem.
 - Self-attention:
 - considers global dependencies between words, keeping track of the whole input sequence.
- BERT is the precursor pre-trained model.
- Further BERT-based models have been defined according to the specific corpus domain, such as biomedical and COVID-19.



BERTbase:

- 12 layers of transformers
- 768 hidden size
- 12 self-att.

RQ1 : Can transformer-based models successfully integrate into the living systematic review process on COVID-19 research?

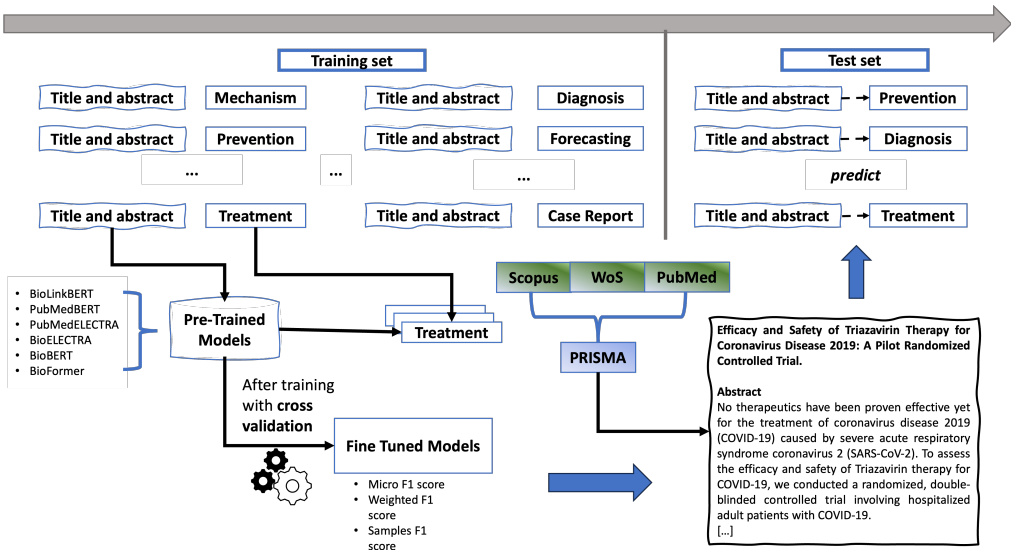
RQ2 : Are the provided labels enough to classify documents?

▶ Background

▶ **Methodology**

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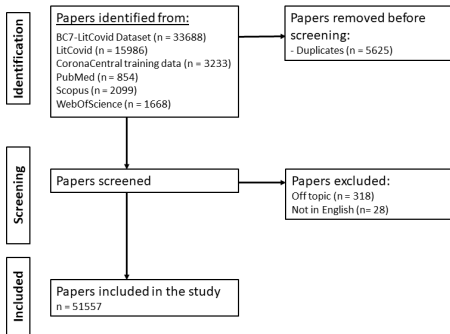
e.g. Scopus search query keywords in title:

- artificial intelligence OR ... OR deep learning OR computer vision
- sars-cov-2 OR corona OR covid-19 OR coronavirus OR covid
- imaging OR image OR screen OR screening OR scan OR medical OR medical imaging OR medical image OR chest OR pulmonary OR lung OR simulation OR modeling OR tomographic OR therapy OR clinical OR gamma-ray OR radiation OR imaging systems OR ... OR quantitative imaging

From the titles and abstracts

- Identified almost 58k
- Included 51k
- Excluded only 5k

Identification of studies: PRISMA Flowchart

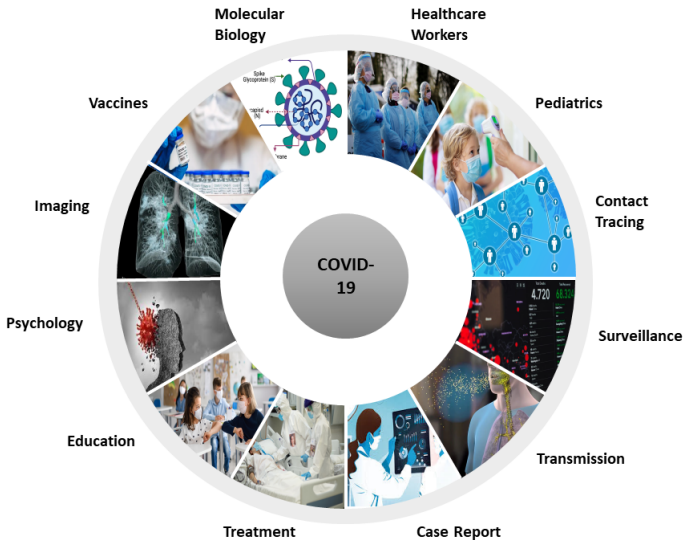


Search queries performed in July 2023.

| Corpus | Training set | Test set | Total | Labeled |
|---------------|--------------|----------|-------|---------|
| BC7-LitCovid | 29436 | 2293 | 31729 | ✓ |
| LitCovid | 14781 | 1205 | 15986 | ✓ |
| CoronaCentral | 1308 | 200 | 1508 | ✓ |
| PubMed | 0 | 718 | 718 | |
| Scopus | 0 | 708 | 708 | |
| WebOfScience | 0 | 908 | 908 | |
| Total | 45525 | 6032 | 51557 | |

| Model name | Source | Instances | Words |
|--------------------|----------------------|-----------|-------|
| PubMedBERT | PubMed | 14M | 3.2B |
| BioLinkBERT | PubMed | 14M | 3.2B |
| PubMedELECTRA | PubMed | 14M | 3.2B |
| BioELECTRA | PubMed | 22M | 4.2B |
| | PubMed Central (PMC) | 3.2M | 9.6B |
| BioBERT | PubMed | N/A | 4.5B |
| | PubMed Central (PMC) | | 13.5B |
| BioFormer-LitCovid | PubMed | 33M | N/A |
| | PubMed Central (PMC) | 1M | |
| | LitCovid | 164K | |

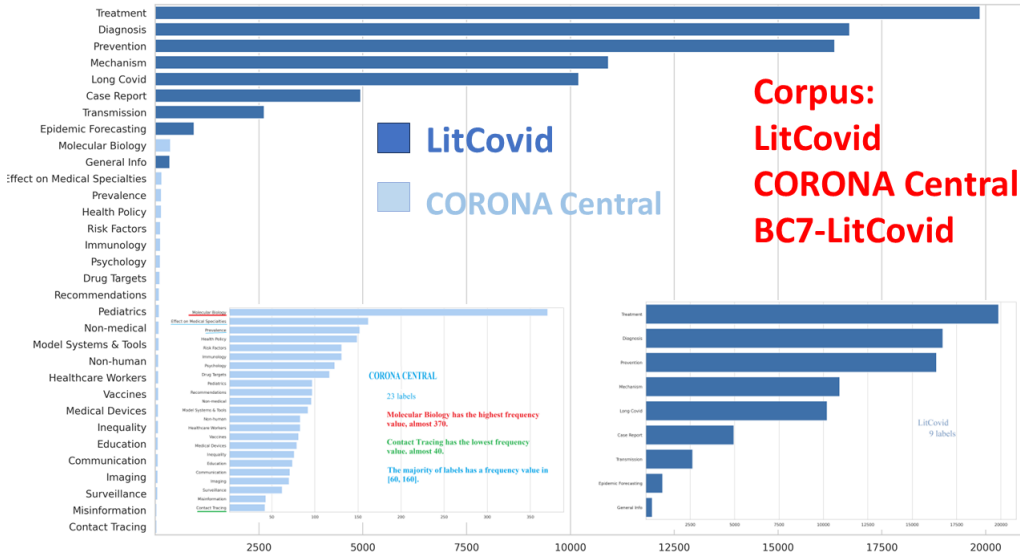
| Size | Model | Hidden Size | Att. Heads | Hidden Layers | Parameters |
|-------|--|-------------|------------|---------------|------------|
| Large | PubMedBERT-Large, BioLinkBERT-Large, PubMedELECTRA-Large, BioBERT-Large | 1024 | 16 | 24 | 340M |
| Base | PubMedBERT-Base, BioLinkBERT-Base, PubMedELECTRA-Base, BioBERT-Base, BioELECTRA | 768 | 12 | 12 | 110M |
| Tiny | BioFormer-LitCovid | 512 | 8 | 8 | 14.5M |



Instance Total Label Distribution in the Training Set

2 Methodology

Label distribution



Performance F1 Score Metric Methods for Model Comparisons

2 Methodology

- **Micro average** computes a global average F1 score by considering total true positives, false negatives, and false positives. Does not separate each class.
- **Macro-average** computes the arithmetic mean of all the per-class F1 scores.
- **Weighted average** computes the weighted average of all the per-class F1 score. The weight W depends on the number of instances of each class.
$$F1_{class1} \times W1 + F2_{class2} \times W2 + \dots + FN_{classN} \times WN$$
- **Samples average** computes F1 score metric for each instance, and returns their average (specifically meaningful for multilabel classification).

▶ Background

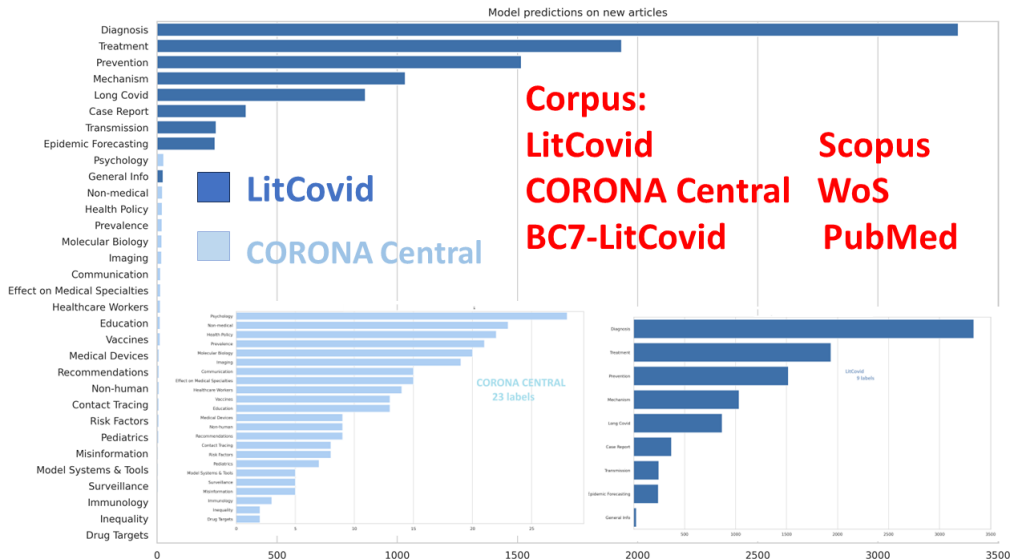
▶ Methodology

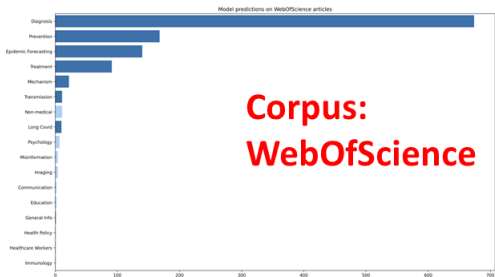
▶ **Results**

▶ Summary

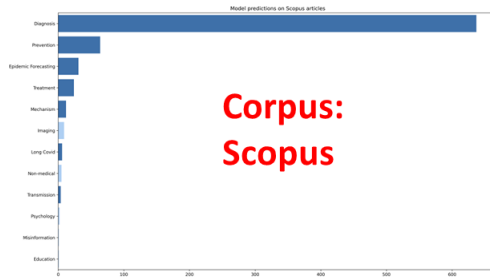
Model Labels Prediction in the Test Set

3 Results



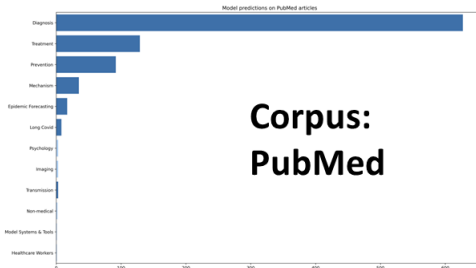


**Corpus:
WebOfScience**



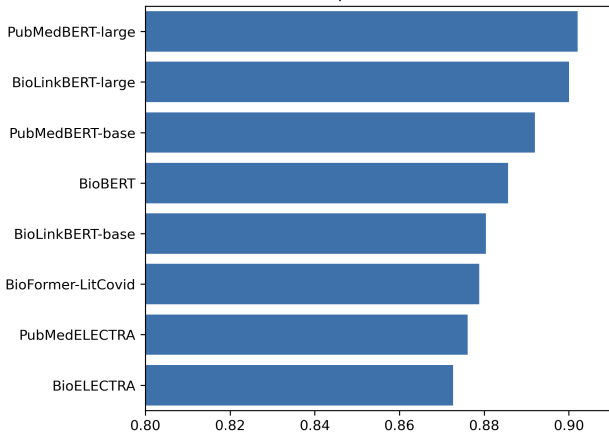
**Corpus:
Scopus**

**WebOfScience
and Scopus
have the same
first 5 labels.**



**Corpus:
PubMed**

Samples F1 scores



The **PubMedBERT-large** model got the best performance.

| Method | F1 |
|----------|--------|
| Micro | 0.8967 |
| Weighted | 0.9010 |
| Samples | 0.9021 |

▶ Background

▶ Methodology

▶ Results

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- RQ1: Can transformer-based models successfully integrate into the living systematic review process on COVID-19 research?
 - Yes, our results show that transformer-based models can be integrated to classify **papers on COVID-19** achieving up to **0.9021** of F1-value with the **PubMedBERT-large** model.
 - Model is **domain specific**.
- Transformers have a positive impact on **speed**, **efficiency** and **customizability** of systematic reviews.
- RQ2: Are the provided labels enough to classify documents? Our research indicates that the labels extracted from LitCovid and Corona Central provide only broad information (such as diagnosis or treatment) regarding the content of publications.

- Addition of labels to enable a more specific classification.
 - e.g. Treatment. Which kind? Which organ is involved?
 - e.g. Diagnosis. Which method? Which part of the body?
- How to organize labels:
 - Flat Text Classification: classes are defined independently of one another.
 - Hierarchical Text Classification: pre-defined classes are organized in a tree-like structure.
- Update extraction results.
- Explorer other models.

Any Questions?

