Initial Experiments on the DRSM Corpus

Aakash Bhatnagar*, Nidhir Bhavsar*, Tirthankar Ghosal†
*Navarachna University, India
†Oak Ridge National Laboratory, US

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Abstract

This report documents the initial experiments we perform on the Disease Research State Model (DRSM) corpus. DRSM is a dataset containing abstract of rare disease articles. Here, 'rare disease' are referred to those diseases that affects less than 200,000 people in the USA. However, there are less than 5% approved therapies. The number of approved/possible therapies would increase if we can leverage the scientific knowledge encoded in rare disease research papers.

The DRSM corpus [3] consists of six distinct labels:

- Clinical characteristics or disease pathology

- Disease mechanism

- Therapeutics in the clinic

- Patient-based therapeutics

- Other

- Irrelevant

These labels represents the state of the research of each disease. As these classes are manually annotated, it takes a good amount of time to curate thousands of instances. So far, the DRSM corpus have a total of 8919 instances. The dataset is imbalanced as the number of instances are not evenly distributed amongst all classes. For example, the 'Disease mechanism' class contains only 108 instances, which makes a ML model prone to overfit and produce unstable results. Similarly 'Therapeutics in the clinic and 'Irrelevant' classes also have fewer instances.

Since manual curation is time consuming, we tried to automate the exercise of 'abstract classification'. We experimented with variations of transformer-based neural models. We chose three BERT[5] models that are trained on large volume of biomedical articles.

1. BioBERT [8]

2. PubMedBERT [7]

3. SPECTER [4].

Along with these models, we experimented with dual-attention and label-wise-attention models. Dual attention[6] is a mechanism of applying self-attention twice. Using dual-attention model increased our scores significantly as seen in table1. Also to further improve our results, we added a label-wise-attention(LWAN)[1]

network that gives special attention to each word according to the labels. We adopted the same approach in the 'LitCovid' track of the BioCreative Challenge [2] to perform multi-label classification of COVID-19 articles. Our code is publicly available here[1].

	Precision	Recall	F1-score
clinical characteristics or disease pathology	0.9296	0.9392	0.9344
other	0.9269	0.9664	0.9462
disease mechanism	0.6154	0.4706	0.5333
therapeutics in the clinic	0.8333	0.3846	0.5263
irrelevant	0.7857	0.8627	0.8224
patient-based therapeutics	0.9180	0.8571	0.8865
micro avg	0.9161	0.9058	0.9109
macro avg	0.8348	0.7468	0.7749
weighted avg	0.9138	0.9058	0.9061
sample avg	0.9021	0.9058	0.9033

Table 1: precision, recall and f1-score of the best performing system i.e. Specter dual-attention LWAN

As we mention earlier, the DRSM corpus suffers from class imbalance. E.g., after splitting the training and testing set to 80:20, we had only 17 instances of the 'disease mechanism' class. We are further experimenting to tackle the class imbalance problem.

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