

COVID-QA: A Question Answering Dataset for COVID-19

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Abstract

We present COVID-QA, a Question Answering dataset consisting of 2,019 question/answer pairs annotated by volunteer biomedical experts on scientific articles related to COVID-19. To evaluate the dataset we compared a RoBERTa base model fine-tuned on SQuAD with the same model trained on SQuAD and our COVID-QA dataset. We found that the additional training on this domain-specific data leads to significant gains in performance. Both the trained model and the annotated dataset have been open-sourced at: <https://github.com/deepset-ai/COVID-QA>.

1 Summary

We selected 147 scientific articles mostly related to COVID-19 from the CORD-19 ([The White House Office of Science and Technology Policy, 2020](#) (accessed May 9, 2020)) collection to be annotated by 15 experts. Although the annotators were volunteers, it was required that all have at least a Master’s degree in biomedical sciences. The annotation team was led by a medical doctor (G.A.R.) who vetted the volunteer’s credentials and manually verified each question/answer pair produced. We used an existing, web-based annotation tool that had been created by deepset¹ and is available at their Neural Search framework haystack². The annotations were created in SQuAD ([Rajpurkar et al., 2016](#)) style fashion where annotators mark text as answers and formulate corresponding questions. COVID-QA differs from SQuAD in that answers come from longer texts (6118.5 vs 153.2 tokens), answers are generally longer (13.9 vs. 3.2 words) and it does not contain n-way annotated development nor test sets. We chose a RoBERTa-base architecture ([Liu et al., 2019](#)) and fine-tuned it on the

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¹<https://deepset.ai/>

²<https://github.com/deepset-ai/haystack>

Model	Exact Match	F1
Baseline	21.84	49.43
COVID-QA model	25.90	59.53

Table 1: Performance comparison. The scores for the COVID-QA model are averaged across the 5 folds.

SQuAD dataset as a baseline model. We continued training the baseline model using our COVID-QA annotations in 5-fold cross validation manner.

Table 1 shows the performance of the baseline model vs. the model finetuned on COVID-QA. Finetuning the model on COVID-QA results in significant improvement across both metrics though the overall scores are pretty low compared to SQuAD. We hypothesize the low scores relate to more complex question/answer pairs on much longer documents and the lack of multiple annotations per question. Without n-way annotations it is difficult to assess the human benchmark and thereby the true difficulty of the dataset. We might add n-way annotations in a future version of the dataset.

References

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