scientific data

Check for updates

OPEN Automatic question answering for multiple stakeholders, the epidemic DATA DESCRIPTOR question answering dataset

Travis R. Goodwin¹¹[∞], Dina Demner-Fushman¹, Kyle Lo², Lucy Lu Wang², Hoa T. Dang³ & Ian M. Soboroff³

One of the effects of COVID-19 pandemic is a rapidly growing and changing stream of publications to inform clinicians, researchers, policy makers, and patients about the health, socio-economic, and cultural consequences of the pandemic. Managing this information stream manually is not feasible. Automatic Question Answering can quickly bring the most salient points to the user's attention. Leveraging a collection of scientific articles, government websites, relevant news articles, curated social media posts, and questions asked by researchers, clinicians, and the general public, we developed a dataset to explore automatic Question Answering for multiple stakeholders. Analysis of questions asked by various stakeholders shows that while information needs of experts and the public may overlap, satisfactory answers to these questions often originate from different information sources or benefit from different approaches to answer generation. We believe that this dataset has the potential to support the development of question answering systems not only for epidemic questions, but for other domains with varying expertise such as legal or finance.

Background & Summary

The COVID-19 pandemic has highlighted two major problems with access to reliable domain-specific information: (1) information providers are rapidly generating reams of potentially vital information that might never reach their intended audience, and (2) information seekers struggle to find answers to their specific and changing information needs in the rapidly growing and changing body of communications. These two problems are effectively two sides of the same coin, and making information easier to discover reduces the potential for vital information to remain overlooked by its intended audience. Information Retrieval (IR) systems (e.g., search engines such as Google, Bing, or PubMed) are now an integral part of information seeking and one of the main tools used to satisfy information needs. The pandemic exposed a major problem when relying on information retrieval systems under stressful and fast-changing circumstances: existing solutions retrieve hundreds or thousands of documents -- many of which may not be relevant to the user's specific information need. This needlessly increases the time and mental effort required by information seekers who must read through these documents to find answers to their information needs. Question Answering is a research area aiming to alleviate the problem of finding pertinent information from thousands of documents by bringing the most salient points of relevant documents to focus immediately, saving time and highlighting information that may have been overlooked.

To this end, we propose a new question answering dataset: Epidemic QA (EPIC-QA), intended to evaluate automatic approaches for answering ad-hoc questions about the disease COVID-19, its causal virus SARS-CoV-2, related coronaviruses, and the recommended response to the pandemic. While COVID-19 has been an impetus for a large body of emergent scientific research and inquiry, the response to COVID-19 raises questions for the general public. The rapid increase in coronavirus literature and evolving guidelines on community response creates a challenging burden not only for the scientific and medical communities but also the general public to stay up-to-date on the latest developments. Consequently, the goal of the dataset is to evaluate systems on their ability to provide timely and accurate expert-level answers as expected by the scientific and medical communities as well as answers in suitable language for the general public. The EPIC-QA dataset consists of (a) questions collected from various stakeholders who need information on COVID-19, (b) documents

¹National Library of Medicine, Bethesda, MD, USA. ²Allen Institute for AI, Seattle, WA, USA. ³National Institute of Standards and Technology, Gaithersburg, MD, USA. [™]e-mail: travis.goodwin@nih.gov

How does the vaccine work?

SARS-CoV-2 mRNA vaccines have shown remarkable clinical efficacy, but questions remain about the nature and kinetics of T cell priming. We performed longitudinal antigen-specific T cell analyses on healthy SARS-CoV-2-naive and recovered individuals prior to and following mRNA prime and boost vaccination. Vaccination induced rapid antigen-specific CD4' T cell responses in naive subjects after the first dose, whereas CD8' T cell responses developed gradually and were variable in magnitude. Vaccine-induced Th1 and Tfh cell responses following the first dose correlated with post-boost CD8' T cells and neutralizing antibodies, respectively. Integrated analysis revealed coordinated immune responses with distinct trajectories in SARS-CoV-2-naive and recovered individuals. Last, whereas booster vaccination improved T cell responses in SARS-CoV-2-naive subjects, the second dose had little effect in SARS-CoV-2-recovered individuals.				
Painter, Mark M et al. "Rapid induction of antigen-specific CD4 ⁺ T cells is associated with coordinated humoral and cellular immunity to SARS-CoV-2 mRNA vaccination." <i>Immunity</i> .				
COVID-19 vaccines help our bodies develop immunity to the virus that causes COVID-19 without us having to get the illness. Different types of vaccines work in different ways to offer protection. But with all types of vaccines, the body is left with a supply of "memory". Trymphocytes as well as B-lymphocytes that will remember how to fight that virus in the future.				

"Understanding How COVID-19 Vaccines Work." Online: cdc.gov

Fig. 1 Examples of different answers appropriate for experts and the general public, respectively.

- including scientific research articles suited for experts as well as relevant news, social media, and pages from authoritative websites intended for the general public, and (c) answers to these questions extracted from documents in the dataset and systematically annotated and judged by experts.

Only a handful of publicly available health-related QA datasets exist. In terms of information retrieval, CORD-19¹ is the most widely used document collection, containing documents from PubMed, research articles from the World Health Organization (WHO), pre-prints from bioRxiv, medRxiv, and arXiv, and SemanticScholar. CORD-19, however, does not provide questions or answers. In terms of COVID question answering, CovidQA² provides 124 question–article–answer triplets from 85 articles; COVID-Q³ consists of 1,690 questions (without answers) annotated into 15 categories and 207 clusters; Medical Question Pairs (MQP)⁴ contains 3,048 pairs of medical questions annotated as being similar or dissimilar to COVID-19 FAQs; the Synergy Task of the 2021 BioASQ Challenge (http://www.bioasq.org/) included four rounds of binary, factoid, list, and summary questions based on the CORD-19 collection; and the 2021 TREC Health Misinformation Track (https://trec-health-misinfo.github.io/) includes topics focused on COVID-19 with annotations indicating whether documents contradict the topic's answer. For a detailed review of additional natural language processing resources for COVID-19, see Chen *et al.* (2021)⁵. By contrast, EPIC-QA, which incorporates both public-facing and expert-level documents, is the only dataset exploring the difference between answers suitable for the general public and those appropriate for experts. It is the only dataset designed to include a wide variety of unique possible answers for each question.

While there is overlap in the types of questions asked by different stakeholders, the answers to such questions should vary based on the background knowledge of the user. For example, consider the simple question illustrated in Fig. 1, *How does the vaccine work?* In the eyes of an expert, an answer should indicate the exact mechanism and pathways involved. By contrast, for a member of the general public, this information is too involved and may cause confusion; instead, a more appropriate answer would provide a more general overview of the role of the vaccine and a high-level description of its mechanism. In the context of the rapidly accelerating knowledge of COVID-19, managing this duality between expert- and general- level information is even more important. It is our hope that the EPIC-QA dataset will stimulate research in automatic question answering not only to support providing high-quality timely information needs in the face of varying levels of expertise for other domains, such as general healthcare, finance, or legal.

Methods

Question answering systems traditionally involve two main steps: an *information retrieval* step, in which relevant documents are retrieved for a given question, and an *answer extraction* step, in which relevant passages or answers are identified within the retrieved documents and extracted as-is or used to synthesize an answer. Recently, *end-to-end* question answering systems have been developed in which (deep) neural networks are trained to jointly identify relevant passages and extract answers from those passages. Consequently, the EPIC-QA dataset includes two collections intended to facilitate research on (a) answer extraction or (b) end-to-end question answering, respectively. The questions in the answer-extraction dataset are adapted from the queries of the TREC-COVID retrieval shared task (https://ir.nist.gov/trec-covid), and are associated with the set of relevancy annotations from that task. For the end-to-end dataset, we introduce a novel set of questions, and provide relevant documents and answer annotations without associated retrieval relevance rankings. These two collections can be used separately or in concert to study the relationship between document retrieval and answer extraction. Both collections consist of three components:

- 1. Two sets of manually-produced questions, one asked by the general public as well as another asked by experts pertaining to COVID-19;
- 2. A collection of documents relevant to COVID-19 including published and pre-print biomedical research articles as well as documents obtained from government websites, news, and social media, automatically parsed into sentences; and

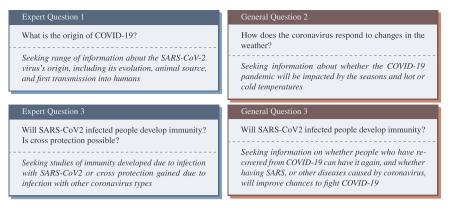


Fig. 2 Example questions included in EPIC-QA, shown above their corresponding backgrounds.

3. A diverse list of manually-produced answers for each question as well as a set of relevant document excerpts with manual sentence-level annotations indicating which answers, if any, are evidenced by each sentence.

To facilitate answer-extraction research, the questions in the answer-extraction dataset are additionally associated with a set of relevant documents as might be retrieved by an optimal search engine, while the questions in the end-to-end dataset are not. We detail the process by which each of these components was produced.

Generating questions. Two sets of questions are included: one for expert-level questions and one for the general public. By design, many of the expert and consumer questions overlap in focus to facilitate comparison between both types of stakeholders. For the answer extraction dataset, we adapted the topics evaluated in the fourth round of TREC-COVID (https://ir.nist.gov/covidSubmit). The majority of these questions originated from users' interactions with MedlinePlus. Additional scientific questions were developed based on group discussions from the National Institutes of Health (NIH) special interest group on COVID-19, questions asked by Oregon Health & Science University clinicians, and responses to a public call for questions. The TREC-COVID topics included expert-level background descriptions of the user's information need; to facilitate public-facing question answering we produced public-friendly backgrounds for each question. We provide the document-level relevance judgments produced during the TREC-COVID evaluation for these questions. The answer extraction collection includes 45 expert questions and 42 general public questions with document-level relevance judgments. For the end-to-end collections, a new set of 30 expert and 30 general public questions were developed, none of which were evaluated in TREC-COVID; consequently, these questions do not come with document-level relevance judgments. Example questions are provided in Fig. 2. While these may seem like a small number of questions (87 answer extraction questions in total and 60 end-to-end questions in total), each question is associated with multiple answers, and 45-60 passages hand-annotated at the sentence level to indicate which answers, if any, are included in each sentence.

Collecting documents. Each document in EPIC-QA includes explicitly pre-defined *contexts* (a generalization of paragraphs or sections) and sentence boundaries. To support both levels of expertise, the document collections consists of two parts: scientific and medical research, and public-facing online documents.

Scientific and research documents. We adapt the collection of biomedical articles released for the COVID-19 Open Research Dataset Challenge (CORD-19)⁶. The dataset was created by the Allen Institute for AI in partnership with the Chan Zuckerberg Initiative, Georgetown University's Center for Security and Emerging Technology, Microsoft Research, and the National Library of Medicine – National Institutes of Health, in coordination with The White House Office of Science and Technology Policy. The CORD-19 collection includes a subset of articles in PubMed Central (PMC) as well as pre-prints from bioRxiv and medRxiv. Contexts in this collection correspond to automatically identified paragraphs in the articles' abstracts, or main texts. We include two snapshots of CORD-19: a snapshot from June 19, 2020 to be used with the answer extraction collection, and a snapshot from October 22, 2020, to be used for the end-to-end collection.

Public-facing documents. We include a subset of the articles used by the Consumer Health Information Question Answering (CHIQA) service of the U.S. National Library of Medicine (NLM)⁷. This collection includes authoritative articles from the Centers for Disease Control and Prevention (CDC); the Genetic and Rare Disease Information Center (GARD); the Genetics Home Reference (GHR); Medline Plus; the National Institute of Allergy and Infectious Diseases (NIAID); and the World Health Organization (WHO). Contexts in this collection correspond to paragraphs or sections as indicated by the HTML markup of the document. All articles were filtered for COVID-19 content using the terms in Fig. 3. In the end-to-end collection, we also included 265 Reddit threads as well as a subset of the CommonCrawl News Crawl (CCNC) from January 1 through April 30, 2020, as used in the TREC Health Misinformation Track (https://trec-health-misinfo.github.io/). To avoid misinformation, we only considered Reddit threats from the heavily-moderated /r/askscience community

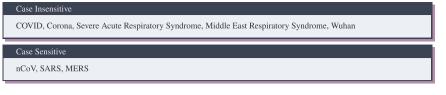


Fig. 3 Terms used to test documents for COVID-19 (note: while terms like Corona or Wuhan may appear under-specified, all documents originate from healthcare-focused collections resulting in few false positive matches).

Which COVID-19 vaccine trials were paused and what were the health safety concerns?

Expert Nuggets		General Nuggets
 trial pause, early stopping, COVID trial, BCG vaccine, COVID-19 vaccine, chloroquine, hydroxychloroquine, lopinavir-rotinavir, patient safety, 	 cardiac toxicity, lack of benefit, trial phase, personal protective equipment, transverse myelitis, neurologic symptoms, failure to reduce mortality, review of data 	 trial pause, early termination, patient safety, no effect on mortality, lack of eligible patients, remdesivir, hydroxychloroquine, COVID-19 vaccine

Fig. 4 Examples of information "nuggets" or answers produced during answer key generation.

.....

that were tagged as pertaining to COVID-19, Medicine, Biology, or the Human Body (and which also contain the COVID-19 terms in Fig. 3). Likewise, to avoid irrelevant pages and misinformation, CCNC documents were filtered by top-level domain (i.e., ".gov" or ".edu"), using the top 100 first-level domains as measured by SALSA⁸, PageRank⁹, and HITS¹⁰; remaining pages were filtered for COVID-19 content using the terms in Fig. 3.

Producing answers. When producing answers for the EPIC-QA dataset, our intent was to explore the landscape of answers asserted in the document collection. Thus, we consider any statement that answers the question as an "answer" regardless of whether or not the answer is factually accurate at the time the document was authored or given current information. The answers in this dataset are intended as an intermediary step where-in one would like to explore all answers provided by the document collection – both correct answers as well as incorrect answers that people may have discovered on their own. Answers were produced by applying multiple automatic question answering systems to the questions in EPIC-QA to identify contexts that may answer the question (see *Technical Validation* for more details). These contexts were pooled at a depth of five for expert questions and a depth of eight for questions from the general public, resulting in roughly 45–60 contexts per question. Human assessors read these retrieved contexts and judged them. Specifically, 17 medical indexers with five to thirty years of experience at the National Library of Medicine analyzed the pooled contexts in two rounds: (1) an answer-key generation round, followed by (2) a sentence-level answer annotation round. Annotations were performed as part of the medical indexers' official duties at the National Library of Medicine.

Answer key generation. In the first round, assessors had access to a pool of contexts identified for a given question as well as an ad-hoc search engine over the document collection. The role of the answer key generation round was for assessors to explore the contexts identified by automatic question answering systems as well as the document collection to determine a set of atomic "facts" that answer the question based on the question's information need or background (see Fig. 2). As in Voorhees $(2005)^{11}$, we refer to these facts as *nuggets*, such that each nugget indicates a fact for which the assessor could make a binary decision as to whether a context or sentence contained that fact. Example nuggets that were produced for the question "Which COVID-19 vaccine trials were paused and what were the health safety concerns?" are illustrated in Fig. 4: *lack of benefit, patient safety*, etc. The primary role of this round is to create an answer key for the question comprised of nuggets identified from pooled contexts or the assessor's own ad-hoc search of the collection. The search engine was provided to help assessors explore the topic and identify nuggets that the assessor may be aware of from their prior knowledge of the question but which may not have been included in any of the pooled contexts. Assessors were not expected to exhaustively identify every possible nugget; rather the intent was for them to identify important (at the discretion of the assessor) nuggets that they feel should be included in the answer key based on their understanding of the question.

Sentence-level answer annotation. In the second round, the answer key (list of nuggets) was fixed. Assessors were given the same set of pooled contexts used in round one. This time, they were asked to annotate all sentences in each context indicating which nugget(s) (if any) are addressed by each sentence. For example, for the question "Which COVID-19 vaccine trials were paused and what were the health safety concerns?" the sentence "[I]n the light of a recent publication...on the lack of safety and efficacy of HCQ in the treatment for COVID-19 patients, the Executive Group of the Solidarity Trial decided to implement a temporary pause of the HCQ arm

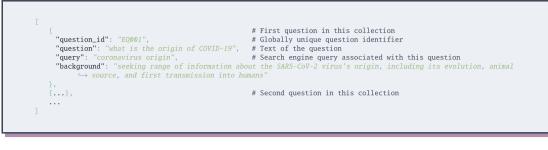


Fig. 5 JSON Schema for questions.

<pre>"document_id": "0hldozml", "metadata": {</pre>	# 40-character SHA1 of the URL or document
"title": "SARS-CoV-2 amino acid substitution	ns widely spread in the human population are mainly located in highly
<pre></pre>	# URL(s) associated with the document
	0!\u00ed!az, Ivan; Clilverd, Hepzibar; Darwich, Laila; Mateu, Enric"
"contexts": [<pre># List of context(s) in the document</pre>
 → study the introduction and evolut → identified and analysed the amino ↔ the pandemic. Eight mutations hav ↔ segments of the structural protei ↔ they might have a functional impa ↔ success in the SARS-CoV-2 virus p ↔ absolutely prevalent, two mutatio 	<pre># Globally unique context identifier # Name of the section (if any) containing the context hrome coronavirus 2 (SARS-CoV-2) pandemic offers a unique opportunity to ion of a pathogen into a completely na\u00efve human population. We acid mutations that gained prominence worldwide in the early months of e been identified along the viral genome, mostly located in conserved ns and showing low variability among coronavirus, which indicated that ct. At the moment of writing this paper, these mutations present a varied opulation; ranging from a change in the spike protein that becomes ns in the nucleocapsid protein showing frequencies around 25%, to a at nearly fades out after reaching a frequency of 20%.", # Inclusive character start offset of the sentence</pre>
"end": 197, "sentence_id": "0hldozml-C000-S000",	<pre># Exclusive character end offset of the sentence # Globally unique identifier for the sentence</pre>
<pre>}, { "start": 198, "end": 319, "sentence_id": "Ohldozml-C000-S001", }.</pre>	# Second sentence in the context
{},]	# Third sentence in the context
}, {},	# Second context in the document

Fig. 6 JSON Schema for full-text documents. Note: the full-text of the document is provided within the list of contexts of that document.

within the trial as a precaution, while the safety data is being reviewed" was annotated as containing the expert nuggets: *COVID trial, trial pause, lack of benefit, hydroxychloroquine, patient safety,* and *review of the data.*

Data Records

The Epidemic Question Answering (EPIC-QA) dataset is available through the Open Science Foundation (OSF) at https://doi.org/10.17605/OSF.IO/VNYK8¹². The questions in the EPIC-QA dataset are provided in four files following the JSON format illustrated in Fig. 5:

- ae/expert_questions.json contains 45 expert-level questions produced for the answer extraction collection; and
- ae/general_questions.json contains 42 questions from the general public produced for the answer extraction collection.
- e2e/expert_questions.json contains 30 expert-level questions produced for the end-to-end collection; and
- e2e/general_questions.json contains 30 questions from the general public produced for the end-to-end collection.

[{ "question_id": "EQ001",	# First question in the collection # Globally unique question identifier for this set of answers
"nuggets": [{ "nugget_id": "EQ001-N00", "nugget": "emergence"	<pre># First nugget for this question # Globally unique nugget identifier # Description of this nugget</pre>
<pre>}, { "nugget_id": "EQ001-N01", "nugget": "species of origin"</pre>	<pre># Second nugget for this question</pre>
}, {}, 	
],	
"annotations": [{ "sentence_id": "0hldozml-C001-S000", "nugget_ids": [<pre># First sentence-level nugget annotation for this question # Globally unique sentence identifier</pre>
"EQ001-N00"]	<pre># Nugget(s) associated with this sentence</pre>
<pre>}, { "sentence_id": "0hldozml-C001-S002", "nugget_ids": [</pre>	# Second sentence-level nugget annotation for this question
}, {}, 	# Third sentence-level nugget annotation for this question
<pre>}, { "question_id": "EQ002", "nuggets": [], "annotations": []</pre>	# Second question in the collection
}, 1	

Fig. 7 JSON Schema for answers.

.....

The documents included in the EPIC-QA dataset are provided separately for the answer extraction and end-to-end collections. For each collection, each document is provided as a separate JSON file, adhering to the JSON schema illustrated in Fig. 6:

- ae/cord19/ contains 129,069 JSON files, each corresponding to a biomedical research article from a June 19, 2020 snapshot of CORD-19; and
- ae/general/ contains 925 JSON files, each corresponding to a government or health agency website crawled on June 10, 2020.
- e2e/cord19/ contains 236,035 JSON files, each corresponding to a biomedical research article from an October 22, 2020 snapshot of CORD-19;
- e2e/ask_science/ is initially empty (users retain copyright on each of their comments and posts on Reddit, preventing us from redistributing their data directly), however, it can be populated using the Python script scripts/populate_reddit.py to create 263 JSON files, each corresponding to a post made between December 01 2019 and October 29, 2020 to the /r/askscience community on Reddit tagged by community moderators as corresponding to COVID-19, health, medicine, or biology, which contained one of the COVID-19 keywords shown in Fig. 3
- e2e/ccns_trec/ contains 114,645 JSON files, each corresponding to an HTML-parsed website included in the CommonCrawl News Subset used by the TREC Misinformation Track containing one of the COVID-19 keywords shown in Fig. 3; and
- e2e/chqa contains 2,739 JSON files, each corresponding to an October 9, 2020 HTML-parsed snapshot of a web page affiliated with the NIH, CDC, or WHO containing one of the COVID-19 keywords shown in Fig. 3.

Finally, answers are provided in four files, corresponding to the answer extraction and end-to-end collections of expert and public questions. For each question: each answer (i.e., nugget) is associated with the set of sentence-IDs that nugget was identified within, as shown by the JSON schema in Fig. 7. The answer extraction collection additionally includes document-level relevance judgments in trec_eval (https://github.com/ usnistgov/trec_eval) format, as illustrated in Fig. 8. Note: due to time and budget restrictions, only 21 expert and 17 general public questions were fully-annotated with answers in the answer extraction collection. Finally, EPIC-QA also includes the rankings and answers provided by all participants of the associated EPIC-QA evaluation at the Text Analysis Conference (TAC, see *Technical Validation* for details), as well as an "ideal" ranking of answers based on our assessor's answer annotations.

Question ID	Judgment Round	Document ID	Relevance
1	1.50	0ti403i4	0
1	3.50	0v5wo0ty	1
1	1	0xhho1sh	2
1	5	0xruezf2	1

Fig. 8 Example format for document-level relevance judgments provided for the answer extraction collection. Question ID indicates which question the relevance judgment was made for; Judgment Round refers to the round of the TREC-COVID evaluation in which the judgment was made (this may be safely ignored for EPIC-QA); Document ID indicates which document was judged; and Relevance indicates the relevance of that document, with 0 for not relevant, 1 for partially relevant, and 2 for fully relevant.

.....

Technical Validation

To validate the EPIC-QA dataset, we organized a community evaluation at the 2020 Text Analysis Conference (TAC), in which participants were provided with the dataset (without answers) and asked to submit ranked lists of up to 1,000 relevant *passages* for each question, where a passage was defined as a contiguous sequence of sentences from a single context. We first evaluated the answer extraction collection, and provided the judgments produced from that evaluation as additional data for participants of the later end-to-end evaluation.

We encouraged teams to explore diversity in their answers rather than returning many passages providing the most common or obvious answer. To this end, we validated EPIC-QA using a modified version of Normalized Discounted Cumulative Gain¹³ (NDCG) which we refer to as the Normalized Discount Novelty Score (NDNS). Importantly, while the cumulative gain in NDCG can be computed for a document independently of the other retrieved documents, the Novelty Score (NS) measures the information in a passage that has not been seen previously in the ranked list of passages. Formally, we define the novelty score, NS, of passage *p* as:

$$NS(p) = \frac{n_a \cdot (n_a + 1)}{n_a + n_s} \quad DNS(p_1, \dots, p_l) = \sum_{r=1}^l \frac{NS(p_r)}{\log_2(r+1)} \quad NDNS(p) = \frac{DNS(p)}{DNS(\hat{p})}$$

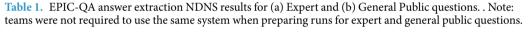
where n_a is the number of *novel* answers in passage p and n_s is the number of sentences in the passage (where an answer is considered novel if it was not included in any earlier-retrieved passages for the question.). This metric ensures that identified passages must be brief (i.e., they must express a novel nugget in as few sentences as possible) and they should not contain sentences with only nuggets provided in previous answers. As in NDCG, we (A) compute the discounted cumulative novelty score, DNS, by adjusting the novelty score, NS, of each answer retrieved up to rank l using a logarithmic reduction factor, and (b) normalize the DNS of ranking $a = a_1, \dots, a_l$ by the DNS of the optimal or ideal ranking of possible answers, \hat{a} that could have been retrieved for that question (based on all the relevance judgments produced for that question). In our evaluation, we used beam-search with a width of 10 to determine the ideal ranking of answers. We report three variations of NDNS which differ in how sentences are counted when calculating NS:

- 1. *Exact*, which prioritizes passages that contain as many novel nuggets in as few sentences as possible, such that *n_e* is exactly equal to the number of sentences in the passage;
- 2. *Partial*, in which the number of sentences used to express novel nuggets is not considered, i.e., *n_s* is the number of sentences if all sentences with novel nuggets were merged into a single sentence before counting; and
- 3. *Relaxed*, in which the number of sentences used to express novel or previously seen nuggets is not considered, i.e., *n_s* is the number of sentences if all sentences containing novel nuggets were merged into a single sentence, and all sentences containing previously seen nuggets were merged into a separate single sentence before counting.

The evaluation script is included with the dataset.

Answer Extraction Validation. A total of 17 submissions were received from eight teams for the expert questions, and 10 submissions from six teams for the general public questions, as shown in Table 1. We observed that the ixa and IBM submissions exhibited the best performance for expert questions while the HLTRI submissions obtained the highest performance followed by IBM for the questions from the general public. The ixa runs used a neural language model powered by SciBERT¹⁴ fine-tuned on the SQuAD 2.0¹⁵ general domain question answering data; their second and third runs additionally fine-tuned on QuAC¹⁶, a dialogue-based question answering dataset involving crowd workers and a teacher discussing a passage of text. Interestingly, their third and best-performing run relied on their own information retrieval system rather than the relevance labels produced during TREC-COVID. IBM's three runs ignored the included relevance labels, using three different ensembles of elastic search with passage retrieval and/or neural re-ranking approaches. Their final runs were a combination of their ensemble information retrieval scores and scores from RoBERTa¹⁷ fine-tuned on Natural Questions¹⁸, a general domain question answering dataset derived from Google search queries and Wikipedia passages. The HLTRI runs relied on a BioBERT¹⁹-based neural passage re-ranker applied to sentence shingles (runs of multiple adjacent sentences) using the TREC-COVID relevance labels. Overall, while the relevance judgments did help

	(a) Expert			(b) General Public			
System	Relaxed	Partial	Exact	Relaxed	Partial	Exact	
CORONAWHY							
Run 1	_	-	-	0.059	0.059	0.043	
covidbert							
Run 1	0.143	0.142	0.131	0.364	0.364	0.389	
Run 2	0.149	0.149	0.165	0.281	0.276	0.257	
Dindadiel	Dindadiel						
Run 1	0.148	0.144	0.158	-	—	-	
HLTRI							
Run 1	0.302	0.295	0.327	0.488	0.482	0.475	
IBM							
Run 1	0.294	0.293	0.325	0.398	0.396	0.413	
Run 2	0.293	0.293	0.306	0.374	0.372	0.39	
Run 3	0.294	0.294	0.327	0.394	0.394	0.409	
ixa							
Run 1	0.276	0.277	0.3	-	—	-	
Run 2	0.303	0.304	0.338	—	—	-	
Run 3	0.305	0.307	0.341	—	—	-	
nlm_lhc_qa							
Run 1	0.113	0.111	0.109	0.278	0.272	0.253	
Run 2	0.134	0.132	0.131	—	—	-	
UPC_USMBA							
Run 1	0.226	0.218	0.215	0.315	0.307	0.302	
Run 2	0.204	0.198	0.212	0.307	0.299	0.286	
vigicovid							
Run 1	0.191	0.192	0.192	-	_	-	
Run 2	0.266	0.267	0.297	-	—	-	
Run 3	0.263	0.265	0.289	—	_	-	



.....

with answering questions for the general public, their impact was less clear for expert-level questions, suggesting that documents may be relevant from an information retrieval perspective despite not having explicitly extraditable answers to the question.

End-to-End Validation. For the end-to-end questions, a total of 16 submissions were received from seven teams for expert questions, and 12 submissions from five teams were received for questions from the general public, as shown in Table 2. For expert questions, the HLTRI runs generally exhibited the best performance, though Yastil_R's first run outperformed HLTRI's first run. The HLTRI runs relied on a BM25-based information retrieval step²⁰, followed by a BERT-based²¹ re-ranker, and a final-step inspired by recognizing question entailment²². The first run from Yastil_R relied on an information retrieval step informed by BM25, DeepCT²³, ColBERT²⁴, and docTTTTTquery²⁵, followed by an answer-extraction step relying on BERT fine-tuned on MS-MARCO²⁶. When looking at the general public, the top performance was exhibited by h2100, whose runs involved a BM25-based retrieval step followed by point-wise and pair-wise re-ranking using T5, and a final sentence-re-ranking step using maximal marginal relevance (MMR) to improve diversity.

When comparing between the answer extraction and end-to-end validation results, it is clear that teams generally performed better on end-to-end despite end-to-end being a more challenging problem. This is likely explained by the fact that the answer extraction phase of the TAC challenge was designed and promoted as a preliminary data-collection round, while the end-to-end questions were presented as the primary round for evaluation. It is our belief that the higher performance on the end-to-end evaluation suggests that teams spent more of their research effort on that task than on the answer extraction task.

Overall, teams were able to produce satisfactory answers using a variety of approaches based on the EPIC-QA data. Interestingly, the changing in ranking between expert and public performance also suggests that one-size does not fit all, and that systems need to better account for varying stakeholders, such as experts and the general public.

Answer Validation. Fig. 9 provides the top scoring expert and general answers extracted by the top performing expert (hltri's third run) and public (h20100's second run) systems developed using the EPIC-QA dataset. Overall, we can see that systems developed using this dataset obtained reasonable results: they were able to identify diverse, concise, and reasonably complete answers that can satisfy experts and the general public.

	(a) Expert			(b) General Public			
System	Relaxed	Partial	Exact	Relaxed	Partial	Exact	
h2oloo							
Run 1	0.388	0.34	0.341	0.407	0.359	0.361	
Run 2	0.39	0.344	0.344	0.414	0.366	0.368	
Run 3	0.376	0.337	0.338	0.382	0.338	0.339	
HLTRI							
Run 1	0.408	0.359	0.36	0.346	0.304	0.305	
Run 2	0.413	0.363	0.364	0.353	0.312	0.313	
Run 3	0.421	0.37	0.371	0.363	0.316	0.317	
IBM							
Run 1	0.353	0.315	0.315	0.267	0.249	0.245	
Run 2	0.367	0.331	0.329	0.282	0.268	0.264	
Run 3	0.354	0.328	0.327	0.278	0.268	0.263	
nlm_lhc_qa	l						
Run 1	0.209	0.223	0.219	0.183	0.186	0.184	
vigicovid	vigicovid						
Run 1	0.359	0.329	0.329	-	—	-	
Run 2	0.374	0.336	0.334	—	—	-	
Run 3	0.391	0.345	0.344	-	_	-	
UPC_USM	BA						
Run 1	0.148	0.126	0.127	0.033	0.03	0.03	
Run 2	—	—	-	0.175	0.176	0.172	
Yastil_R							
Run 1	0.41	0.361	0.362	-	_	-	
Run 2	0.385	0.338	0.339	-	_	-	

 Table 2.
 EPIC-QA end-to-end NDNS results for (a) Expert and (b) General Public questions.
 Note: teams

 were not required to use the same system when preparing runs for expert and general public questions.
 Image: Comparing runs for expert and general public questions.

What measures could mitigate resurgence of COVID-19?

... There appears to be a strong body of evidence, and also guidance from respected international authorities to support the use of simple, society wide home-made face masks and the use of better protection for healthcare workers exposed to the risk of COVID-19. **Hopefully, combined with ongoing social distancing measures, and availability of** widespread testing facilities to support isolated quarantine, such an approach might mitigate against the resurgence of COVID-19 as lockdown measures are relaxed.

... There are six key actions that we recommend. First, expand, train and deploy your health care and public health workforce; Second, implement a system to find every suspected case at community level; Third, ramp up the production, capacity and availability of testing; Fourth, identify, adapt and equip facilities you will use to treat and isolate patients; Fifth, develop a clear plan and process to quarantine contacts; And sixth, refocus the whole of government on suppressing and controlling COVID-19. These measures are the best way to suppress and stop transmission, so that when restrictions are lifted, the virus doesn't resurge...

Are there alternative treatments for COVID-19?

Exper

... Use of Chinese herbal products for treating viruses is not guided by viral pathology, rather herbs are prescribed by herbalists according to Chinese diagnostic patterns (inspection, listening, smelling, inquiry, and palpitation) [12, 13]. **The implications of medicating with herbal-based formulas is serious and dangerous because there is no scientific evidence suggesting that these alternative remedies can prevent or cure COVID-19.** There are several adverse effects noted with herbal medications, such as hepatotoxicity, and there have been numerous reports of toxic contaminants, including pesticides and heavy metals [14]...

General

... Rigorous testing of herbal and other traditional remedies is needed, say experts. The U.S. National Institutes of Health has warned against alternative medicine — including certain herbal therapies and teas — for treating or preventing COVID-19, saying there was no evidence they work and some may be unsafe. "It's the responsibility of those who make a herbal drink to show the scientific evidence that their claims are valid," said Dr. Stephen Barrett, a retired psychiatrist who runs Quackwatch, a website about unproven medical therapies. ...

Fig. 9 The top answers retrieved by the best-performing automatic systems for two questions in EPIC-QA. The specific answer passage identified by the system is typeset in bold.

This indicates that the EPIC-QA dataset can be used to train reliable question answering systems to answer epidemic-related questions posed by different stakeholders.

Impressions. We presented the Epidemic Question Answering (EPIC-QA) dataset, designed to evaluate and draw attention to the important problem of answering questions about COVID-19 from emergent literature, as well as to explore the differences between answers expected by different stakeholders. The EPIC-QA dataset consists of two parts, intended to facilitate answer extraction or end-to-end question answering research, respectively, with each part including (a) two sets of questions collected from two types of stakeholders (experts and the general public), (b) manually-produced answers to these questions extracted from documents in the dataset and systematically hand-annotated and judged by human experts, and (c) a document collection including scientific research articles suited for experts as well as relevant news, social media, and pages from authoritative websites intended for the general public. Technical validation accomplished through a community evaluation at the Text Analysis Conference (TAC) indicates that systems can discover useful answers in the collections, and that answers (and supporting passages) expected by different stakeholders can vary substantially. We believe that our technical validation demonstrates the importance of exploring the diverse landscape of answers available online for health questions and shows the importance of accounting for varying levels of understanding when identifying satisfactory answers to health questions, and hope that this dataset will be of value to researchers exploring answer diversity, multiple stakeholders, or open-ended healthcare question answering.

Usage Notes

We have provided instructions for how to process this dataset in the README file provided with the dataset. Descriptions of automatic systems developed using this dataset are available at https://tac.nist.gov.

Code availability

The code used to prepare the EPIC-QA dataset is provided at https://github.com/h4ste/epic_qa, and a Python script for computing the evaluation metrics reported in the technical validation section of this manuscript is provided with the dataset.

Received: 21 February 2022; Accepted: 5 July 2022; Published online: 21 July 2022

References

- Wang, L. L. et al. CORD-19: The COVID-19 open research dataset. In Proceedings of the 1st Workshop on NLP for COVID-19 at ACL 2020 (Association for Computational Linguistics, Online, 2020).
- 2. Tang, R. et al. Rapidly bootstrapping a question answering dataset for covid-19 (2020). 2004.11339
- 3. Wei, J., Huang, C., Vosoughi, S. & Wei, J. What are people asking about COVID-19? a question classification dataset. In *Proceedings* of the 1st Workshop on NLP for COVID-19 at ACL 2020 (Association for Computational Linguistics, Online, 2020).
- McCreery, C. H., Katariya, N., Kannan, A., Chablani, M. & Amatriain, X. Effective Transfer Learning for Identifying Similar Questions: Matching User Questions to COVID-19 FAQs, 3458–3465 (Association for Computing Machinery, New York, NY, USA, 2020).
- Chen, Q. *et al.* Artificial intelligence in action: Addressing the covid-19 pandemic with natural language processing. *Annual Review of Biomedical Data Science* 4, 313–339, https://doi.org/10.1146/annurey-biodatasci-021821-061045 (2021). PMID: 34465169.
 Wang, L. L. *et al.* Cord-19: The covid-19 open research dataset. *ArXiv* abs/2004.10706 (2020).
- Demner-Fushman, D., Mrabet, Y. & Ben Abacha, A. Consumer health information and question answering: helping consumers find answers to their health-related information needs. *Journal of the American Medical Informatics Association* 27, 194–201, https://doi. org/10.1093/jamia/ocz152 (2019). https://academic.oup.com/jamia/article-pdf/27/2/194/32500415/ocz152.pdf.
- Lempel, R. & Moran, S. Salsa: The stochastic approach for link-structure analysis. ACM Trans. Inf. Syst. 19, 131–160, https://doi. org/10.1145/382979.383041 (2001).
- 9. Page, L., Brin, S., Motwani, R. & Winograd, T. The pagerank citation ranking: Bringing order to the web. Tech. Rep., *Stanford InfoLab* (1999).
- 10. Kleinberg, J. M. Authoritative sources in a hyperlinked environment. Journal of the ACM (JACM) 46, 604-632 (1999).
- Voorhees, E. Using question series to evaluate question answering system effectiveness. In Proceedings of Human Language Technology Conference and Conference on Empirical Methods in Natural Language Processing, 299–306 (Association for Computational Linguistics, Vancouver, British Columbia, Canada, 2005).
- Goodwin, T. & Demner, D. Epidemic Question Answering (EPIC-QA), Open Science Framework, https://doi.org/10.17605/OSF.IO/ VNYK8 (2022).
- Järvelin, K. & Kekäläinen, J. Cumulated gain-based evaluation of ir techniques. ACM Trans. Inf. Syst. 20, 422–446, https://doi. org/10.1145/582415.582418 (2002).
- Beltagy, I., Lo, K. & Cohan, A. SciBERT: A pretrained language model for scientific text. In Proceedings of the 2019 Conference on Empirical Methods in Natural Language Processing and the 9th International Joint Conference on Natural Language Processing (EMNLP-IJCNLP), 3615–3620, https://doi.org/10.18653/v1/D19-1371 (Association for Computational Linguistics, Hong Kong, China, 2019).
- Rajpurkar, P., Jia, R. & Liang, P. Know what you don't know: Unanswerable questions for SQuAD. In Proceedings of the 56th Annual Meeting of the Association for Computational Linguistics (Volume 2: Short Papers), 784–789, https://doi.org/10.18653/v1/P18-2124 (Association for Computational Linguistics, Melbourne, Australia, 2018).
- Choi, E. et al. QuAC: Question answering in context. In Proceedings of the 2018 Conference on Empirical Methods in Natural Language Processing, 2174–2184, https://doi.org/10.18653/v1/D18-1241 (Association for Computational Linguistics, Brussels, Belgium, 2018).
- 17. Liu, Y. et al. Ro{bert}a: A robustly optimized {bert} pretraining approach (2020).
- Kwiatkowski, T. et al. Natural questions: A benchmark for question answering research. Transactions of the Association for Computational Linguistics 7, 452–466, https://doi.org/10.1162/tacl_a_00276 (2019).
- Lee, J. et al. BioBERT: a pre-trained biomedical language representation model for biomedical text mining. Bioinformatics 36, 1234–1240, https://doi.org/10.1093/bioinformatics/btz682 (2019). https://academic.oup.com/bioinformatics/article-pdf/36/4/1234/32527770/btz682.pdf.
- 20. Robertson, S. E. et al. Okapi at trec-3. Proceedings of the Third Text REtrieval Conference (TREC 1995) 109 (1995).
- Devlin, J., Chang, M.-W., Lee, K. & Toutanova, K. BERT: Pre-training of deep bidirectional transformers for language understanding. In Proceedings of the 2019 Conference of the North American Chapter of the Association for Computational Linguistics: Human Language Technologies, Volume 1 (Long and Short Papers), 4171–4186, https://doi.org/10.18653/v1/N19-1423 (Association for Computational Linguistics, Minneapolis, Minnesota, 2019).
- 22. Abacha, A. B. & Demner-Fushman, D. A question-entailment approach to question answering. BMC bioinformatics 20, 1–23 (2019).

- Dai, Z. & Callan, J. Context-Aware Term Weighting For First Stage Passage Retrieval, 1533–1536 (Association for Computing Machinery, New York, NY, USA, 2020).
- 24. Khattab, O. & Zaharia, M. ColBERT: Efficient and Effective Passage Search via Contextualized Late Interaction over BERT, 39–48 (Association for Computing Machinery, New York, NY, USA, 2020).
- 25. Nogueira, R., Lin, J. & Epistemic, A. From doc2query to doctttttquery. Online preprint (2019).
- 26. Nguyen, T. et al. MS MARCO: A human generated machine reading comprehension dataset. In Besold, T. R., Bordes, A., d'Avila Garcez, A. S. & Wayne, G. (eds.) Proceedings of the Workshop on Cognitive Computation: Integrating neural and symbolic approaches 2016 co-located with the 30th Annual Conference on Neural Information Processing Systems (NIPS 2016), Barcelona, Spain, December 9, 2016, vol. 1773 of CEUR Workshop Proceedings (CEUR-WS.org, 2016).

Acknowledgements

The content of this manuscript is solely the responsibility of the authors and does not necessarily represent the official views of the National Institutes of Health (NIH) nor the National Institute of Standards and Technology (NIST). Certain commercial equipment, instruments, or materials are identified in this paper in order to specify the experimental procedure adequately. Such identification is not intended to imply recommendation or endorsement by NIST or NIH, nor is it intended to imply that the materials or equipment identified are necessarily the best available for the purpose. This work was supported by the intramural research program at the U.S. National Library of Medicine, National Institutes of Health. We would like to thank Sam Skjonsberg and Mark Neumann for their help with the annotation platform, Rob Guzman for leading the annotation effort as well as Annemieke van der Sluijs, Cathy Smith, David Cissel, Deborah Whitman, Dorothy Trinh, Farzaneh Badiei, Funmi Akhigbe, Janice Ward, Jenny Rewolinski, Keiko Sekiya, Melanie Huston, Naim Rogers, Nick Miliaras, Oleg Rodionov, Olga Printseva, Rob Guzman, and Susan Schmidt for creating the answer keys and their annotation efforts. Finally, we thank Audrey Tong for her preliminary review of the manuscript.

Author contributions

T.R.G. and D.D.F. originated the study, drafted the manuscript, analyzed the results, collected the non-CORD-19 data, developed the evaluation tools, and were the primary organizers for the associated shared task at TAC. K.L. and L.L.W. prepared the adjusted CORD-19 snapshots used in EPIC-QA and developed the annotation interface. H.T.D. and I.M.S. organized TAC, handled team submissions and pooling, and advised on the role of nuggets in the evaluation. All authors reviewed the manuscript and co-organized the shared task at TAC.

Funding

Open Access funding provided by the National Institutes of Health (NIH).

Competing interests

The authors declare no competing interests.

Additional information

Correspondence and requests for materials should be addressed to T.R.G.

Reprints and permissions information is available at www.nature.com/reprints.

Publisher's note Springer Nature remains neutral with regard to jurisdictional claims in published maps and institutional affiliations.

Open Access This article is licensed under a Creative Commons Attribution 4.0 International License, which permits use, sharing, adaptation, distribution and reproduction in any medium or format, as long as you give appropriate credit to the original author(s) and the source, provide a link to the Creative Commons license, and indicate if changes were made. The images or other third party material in this article are included in the article's Creative Commons license, unless indicated otherwise in a credit line to the material. If material is not included in the article's Creative Commons license and your intended use is not permitted by statutory regulation or exceeds the permitted use, you will need to obtain permission directly from the copyright holder. To view a copy of this license, visit http://creativecommons.org/licenses/by/4.0/.

This is a U.S. Government work and not under copyright protection in the US; foreign copyright protection may apply 2022