

COVID-SEE

Scientific Evidence Explorer for COVID-19

—
Simon Šuster



Stream 4: Real-time clinical decision support

17/7/2020



Team

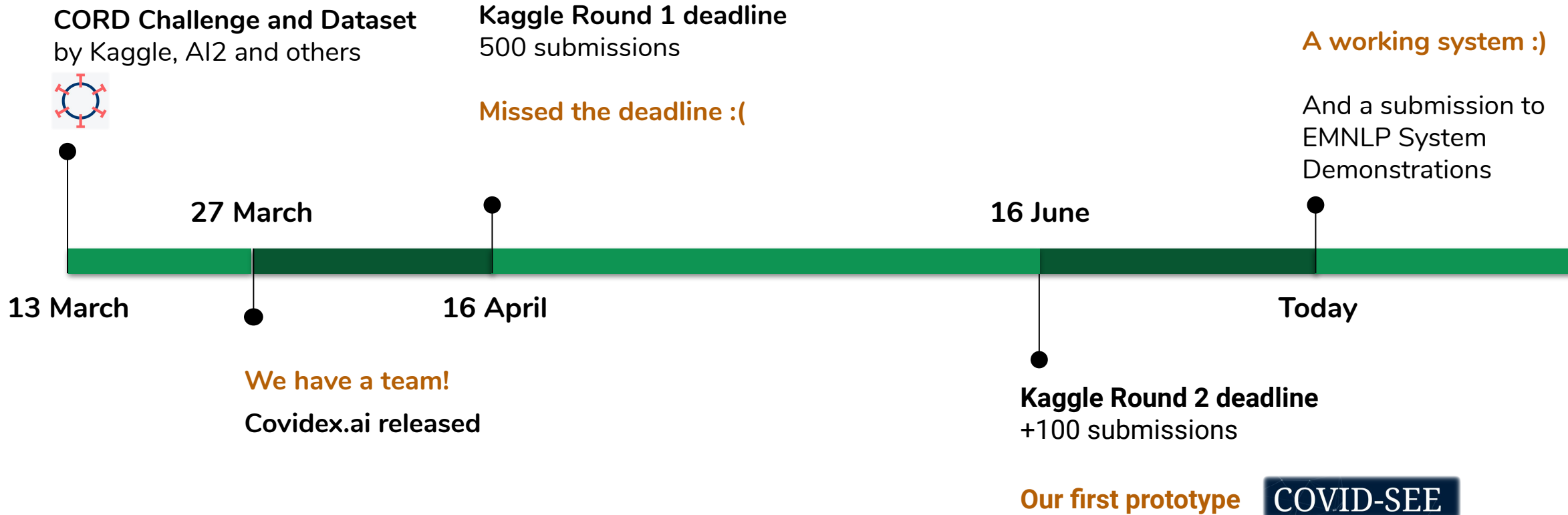
Led by **Karin Verspoor**

Core team: **Simon Suster, Yulia Otmakhova, Zenan Zhai, Biaooyan Fang**

Help from colleagues **Jey-Han Lau** and **Tim Baldwin**, and external collaborators **Antonio Jimeno Yepes** and **David Martinez** (IBM Research Melbourne)

Web developer: **Shevon Mendis**

Timeline



Goals

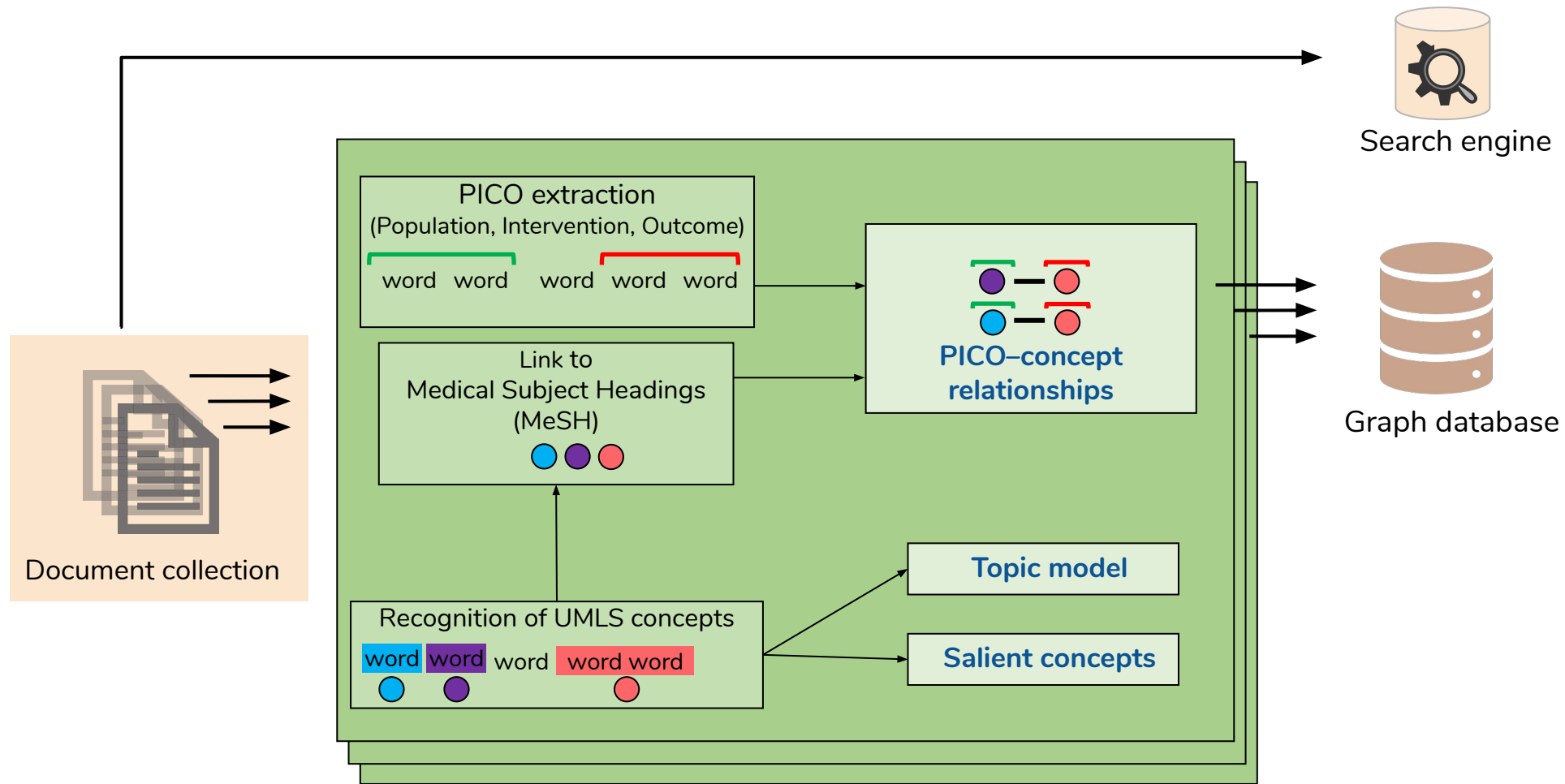
A large number of systems perform lookup

- information retrieval and QA with free-text queries (cord19.vespa.ai, discovid.ai, covid19.mendel.ai)
- semantic search ([IBM COVID-19 Navigator](#), [DOC Search](#), [Trialstreamer](#))

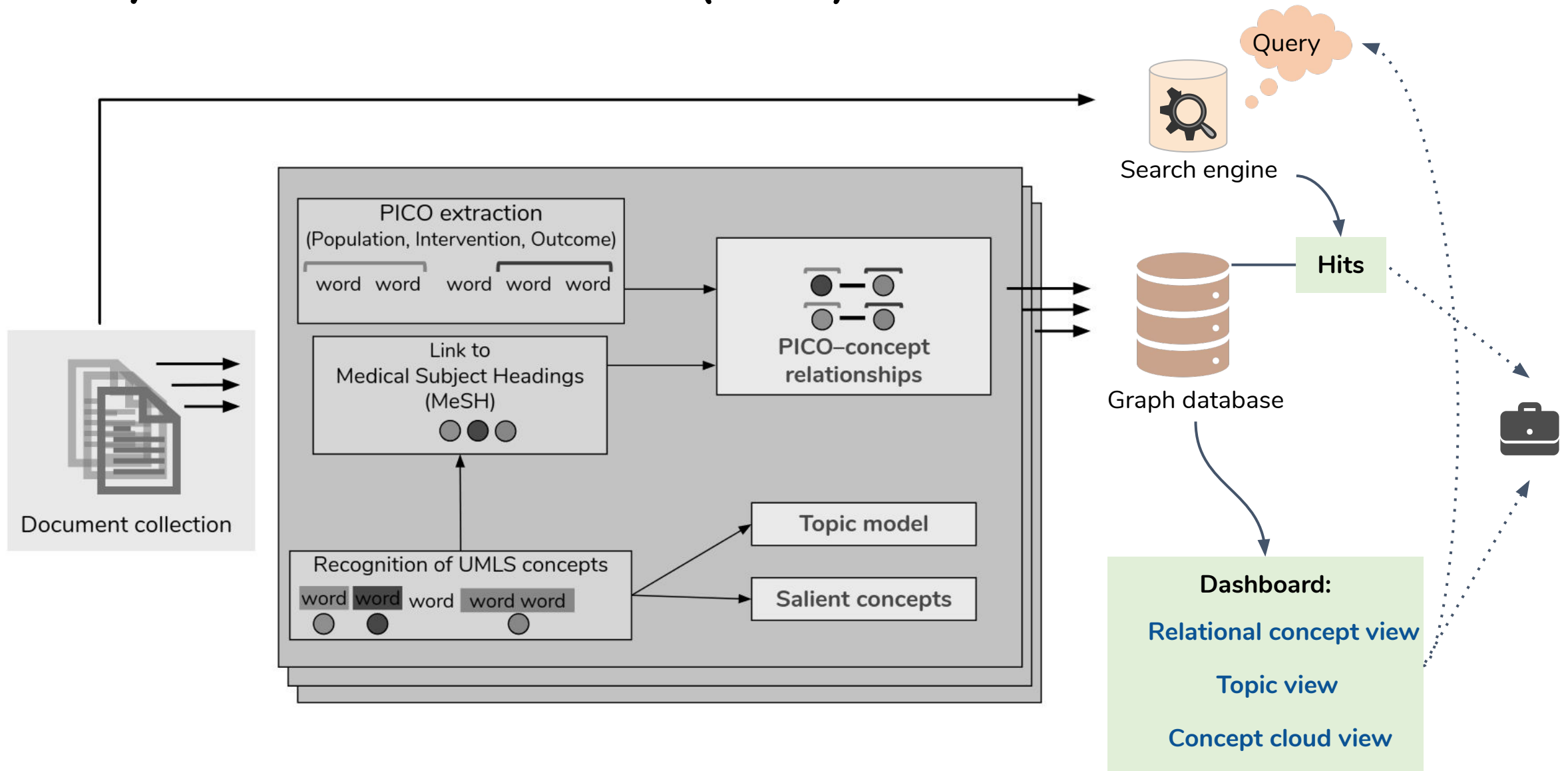
We focus on interactive exploration of literature (White and Roth, 2009; Pang et al., 2015)

- initiate search with a tentative query in natural language
- explore different views of the retrieved documents and obtain cues about next steps
- high-level (current collection) and low-level (document) views
- progressively build up your briefcase

System overview (1/2)



System overview (2/2)



Information retrieval

Use Anserini (Zhang et al., 2020) in Python, builds on Lucene search

- index from CORD-19 abstracts
- retrieved documents ranked with BM25 scoring function

Simple search, but in top 30% of submissions to TREC-COVID shared task (Roberts et al., 2020)

Possible extensions with neural re-ranking



COVID-SEE

🔍 Search literature...

Search >

We found 22 potential matches:

Filter

Year Published:

- 2005 - 2009 (4)
- 2010 - 2014 (1)
- 2015 - 2019 (4)
- 2020 (22)

Source:

- Elsevier (5)
- MedRxiv (15)
- Medline (12)
- PMC (14)
- WHO (5)

Clear All

- Estimate the incubation period of coronavirus 2019 (COVID-19)**

Authors: Han, Henry
Journal: N/A, 2020

Motivation: Wuhan pneumonia is an acute infectious disease caused by the 2019 novel coronavirus (COVID-19). It is being treated as a Class A infectious disease though it was classified as Class B according to the Infectious Disease Prevention Act of China. Accurate estimation of the incubation period...
[+]
- Estimation of the incubation period of SARS-CoV-2 in Vietnam**

Authors: Bui, L. V.; Nguyen, H. T.; Levine, H.; Nguyen, H.; Nguyen, T. A.; Nguyen, T. P.; Nguyen, T.; Do, T. T. T.; Tuan, N. P.; Bui, H. M.
Journal: N/A, 2020

Objective: To estimate the incubation period of Vietnamese confirmed COVID-19 cases. Methods: Only confirmed COVID-19 cases who are Vietnamese and locally infected with available data on date of symptom onset and clearly defined window of possible SARS-CoV-2 exposure were included. We used three parametric...
[+]
- Transmission of corona virus disease 2019 during the incubation period may lead to a quarantine loophole**

Authors: Xia, Wei; Liao, Jiaqiang; Li, Chunhui; Li, Yuanyuan; Qian, Xi; Sun, Xiaojie; Xu, Hongbo; Mahai, Gaga; Zhao, Xin; Shi, Lisha; Liu, Juan; Yu, Ling; Wang, Meng; Wang, Qianqian; Namat, Asmagvl; Li, Ying; Qu, Jingyu; Liu, Q; Lin, Xiaofang; Cao, Shuting; Huan, Shu; Xiao, Jiyong; Ruan, Fengyu; Wang, Hanjin; Xu, Qing; Ding, Xingjuan; Fang, Xingjie; Qiu, Feng; Ma, Jiaolong; Zhang, Yu; Wang, Aizhen; Xing, Yuling; Xu, Shunqing
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Transmission of corona virus disease 2019 during the incubation period may lead to a quarantine loophole
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Journal: N/A
Source: MedRxiv
License: medrxiv
Year: 2020

Background: The ongoing outbreak of novel corona virus disease 2019 (COVID-19) in Wuhan, China, is arousing international concern. This study evaluated whether and when the infected but asymptomatic cases during the incubation period could infect others. Methods: We collected data on demographic characteristics, exposure history, and symptom onset day of the confirmed cases, which had been announced by the Chinese local authorities. We evaluated the potential of transmission during the incubation period in 50 infection clusters, including 124 cases. All the secondary cases had a history of contact with their first-generation cases prior to symptom onset. Results: The estimated mean incubation period for COVID-19 was 4.9 days (95% confidence interval [CI], 4.4 to 5.4) days, ranging from 0.8 to 11.1 days (2.5th to 97.5th percentile). The observed mean and standard

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3 Articles Selected Collect Export

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Collect Selected Articles ✕

Collection:

Name:

Collect

Collections



Selection Manager

Selected Collection: incubation

No. of Articles(Total): 3

No. of Articles(Selected): 0

Move

Delete

Export

Explorer



Active
Collection



incubation



New
Briefcase

Collection Viewer



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Transmission of corona virus disease 2019 during the incubation period may lead to a quarantine loophole

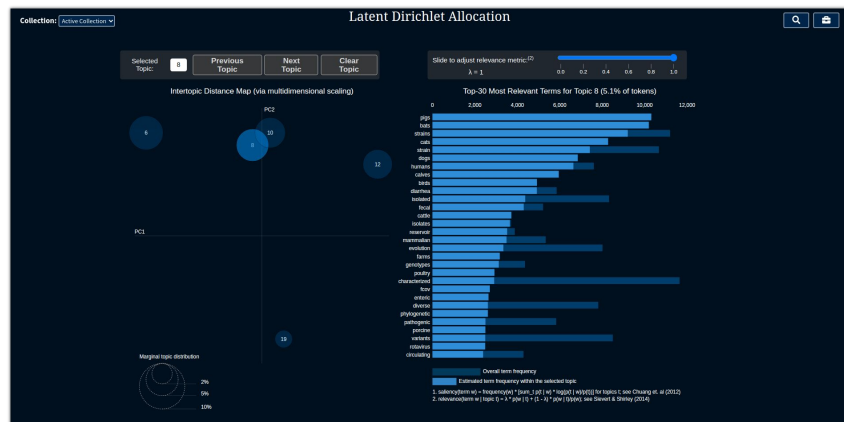
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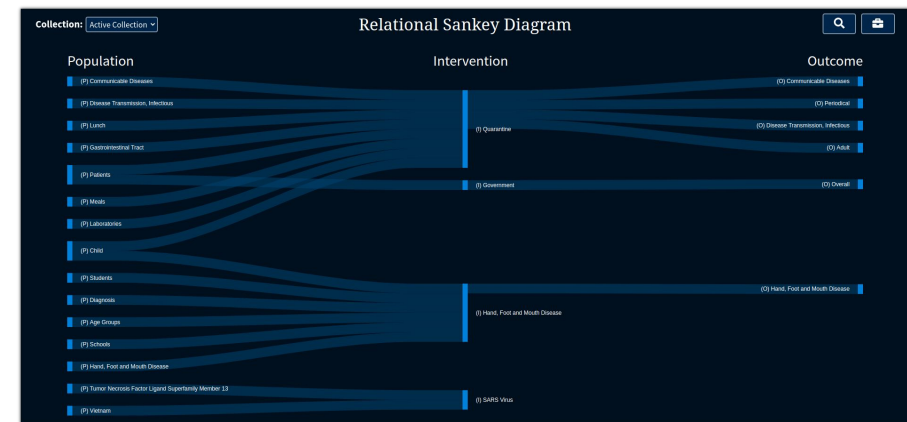
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Dashboard

1. Relational concept view



2. Topic view



3. Concept cloud view



1. Relational concept view

Identify spans describing populations, interventions/comparators and outcomes (PICO) & recognise MeSH terms

“cumulative COVID-19-related ^{Hospitalization} *hospitalization* and ^{Mortality} *death rates*”

OUTCOME

PICO concepts: *Hospitalization*_{OUTCOME}, *Mortality*_{OUTCOME}

Relate Population-Intervention (or Intervention-Outcome) concepts if they occur in the same abstract

- BiLSTM-CRF labeler from 5,000 annotated PubMed abstracts (EBM-NLP, Nye et al., 2018)
- Test set F1: 0.69 for all PICO labels
- MetaMap (metamap.nlm.nih.gov) to extract MeSH terms

Collection: Active Collection ▾

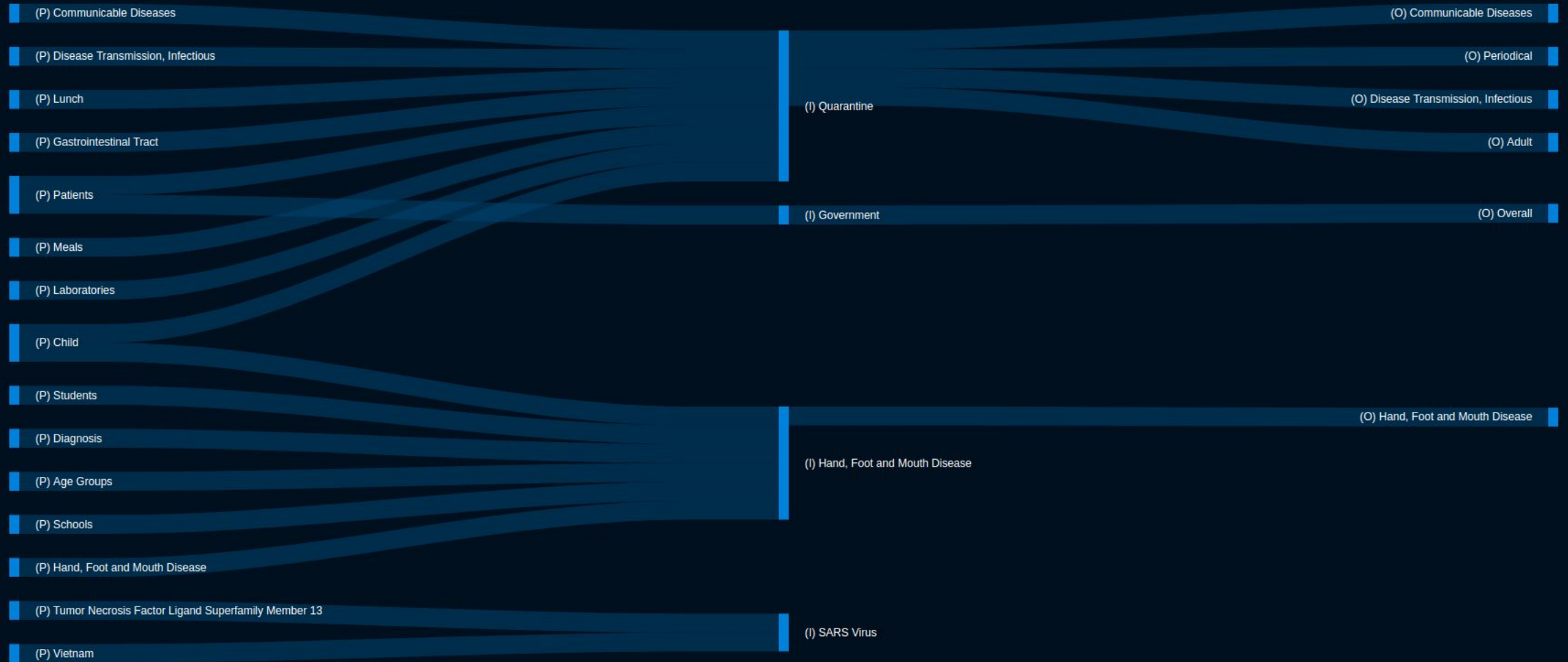
Relational Sankey Diagram



Population

Intervention

Outcome



(P) Disease Transmission, Infectious → (I) Quarantine



Related Literature:

Is a 14-day quarantine period optimal for effectively controlling coronavirus disease 2019 (COVID-19)?

Authors: Jiang, Xue; Niu, Yawei; Li, Xiong; Li, Lin; Cai, Wenxiang; Chen, Yucan; Liao, Bo; Wang, Edwin

Journal: N/A, 2020

Background The outbreak of a new coronavirus (SARS-CoV-2) disease (Covid-19) has become pandemic. To be more effectively controlling the disease, it is critical to set up an optimal quarantine period so that about 95% of the cases developing symptoms will be retained for isolation. At the moment, the...

Currently stored in:

Add to

2. Topic view

Provide a more thematic summary of the current collection

- LDA (Blei et al., 2013) with 20 topics
- Topic model as a mixture over documents, topics and terms
- Shown as a two-dimensional map ([pyLDAvis](#); Sivert and Shirley, 2004)
- We use concept strings as terms

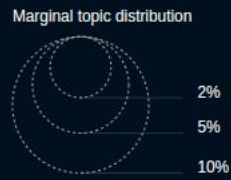
Latent Dirichlet Allocation



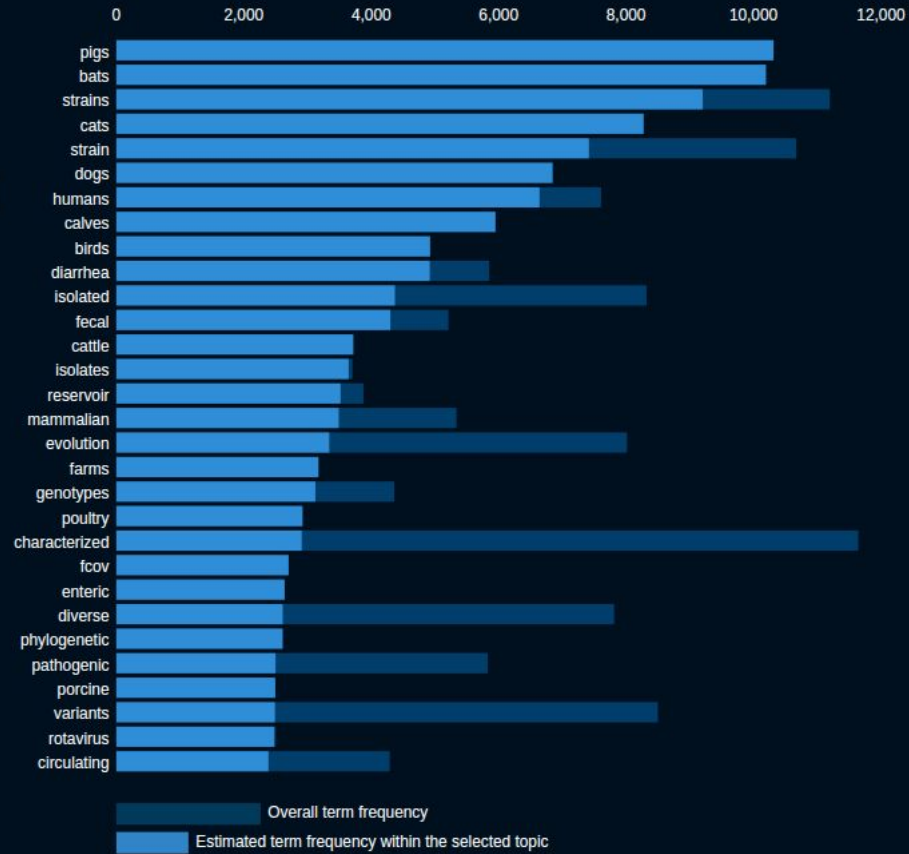
Selected Topic: **8** Previous Topic Next Topic Clear Topic

Slide to adjust relevance metric:⁽²⁾ $\lambda = 1$

Intertopic Distance Map (via multidimensional scaling)



Top-30 Most Relevant Terms for Topic 8 (5.1% of tokens)



1. saliency(term w) = frequency(w) * [sum_t p(t | w) * log(p(t | w)/p(t))]; see Chuang et. al (2012)
 2. relevance(term w | topic t) = $\lambda * p(w | t) + (1 - \lambda) * p(w) / p(w)$; see Sievert & Shirley (2014)

3. Concept cloud view

- 20 most representative concepts per document
- Concept distributions compared with a log-likelihood test (Rayson and Garside, 2000)

Term Clouds

Current Article:

Transmission characteristics of the COVID-19 outbreak in China: a study driven by data



public health major contact with isolate lead symptoms
estimated quarantine threat day diagnosed isolation
excluded ineffective mean incubation period recommendations province
description contact

Where to go from here

- Implement semantic search using concepts and PICO elements
- Recommend articles which have similar topic distributions
- Allow varying concept granularity, visualise how concepts are embedded in the hierarchy
- Evaluate with a user study to better understand exploration and design

COVID-SEE: <http://covid-see.cis.unimelb.edu.au>

- Designed a tool to help scientists navigate the literature on COVID-19
- Used NLP methods (ontology concept recognition, PICO element detection, topic modelling) to structure the key biomedical information
- To facilitate exploration and discovery of novel information, COVID-SEE exposes a user to visual overviews of the content of a document collection

References

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Paul Rayson and Roger Garside. 2000. Comparing corpora using frequency profiling. *The workshop on comparing corpora*, pages 1–6.

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