

CovidGraph: Integrating COVID-19 Data

CovidGraph: In 2020 SARS-COV-2 began to impact life across the globe on a scale unbeknown to humanity. Over the last two years fast and extensive research in that field generated a vast amount of knowledge about the virus. Research-wise, COVID-19 has been encountered with publications, patents, genome analysis, simulation studies for spread prediction, health studies and the extension of ontology information. One factor for fast and reliable research is commitment to the FAIR guiding principles [1]. CovidGraph offers findable accessible interoperable and re-usable COVID-19 data obtained, integrated and connected from open data resources. Data sets from the aforementioned domains are stored in a graph database to offer researchers quick and efficient access to information about COVID-19 (Fig.1). The connections within CovidGraph allow for new types of queries across previously disconnected aspects of the disease.

Domains: CovidGraph comprises information about publications from the COVID-19 Open Research Dataset [2], information about patents [3] and clinical trials [4]. Biomedical entities (e.g. genes, transcripts and proteins) are integrated from a variety of well-established databases [5]. Statistical data is imported from Johns Hopkins University [6]. Simulation models in standard format [7], including a Covid-19 model collection, are integrated from a domain-specific graph database (MaSyMoS, [8]).

CovidGraph – Domains & Connections BioBERT (NLP+NER) BioMedical Patents x 32,080 21,000 Hetionet IOI LENS.ORG 125,000 128,053 488,971 47,000 410,000 Systems Biology Models [MaSyMoS] Papers x 128,053 Statistical & Geographic Clinical Trials x 1,700 NIH U.S. National Library of Medicine ClinicalTrials.gov Pub Med bioRχiv medRχiv JOHNS HOPKINS Figure 1: Domains included in Covidgraph

CovidGraph – How to explore the data?

CovidGraph offers several interfaces for data exploration. The Visual Graph Explorer (Fig. 2) provides predefined views for an intuitive keyword-based graph exploration without prior knowledge of database query languages. SemSpect [9] (Fig. 3) supports drag & drop, expand and filter data items and automatic grouping of similar data items. Hence the graph can easily be traversed and visual representations can be created without detailed knowledge of the data model. Neo4j Bloom (Fig. 4) is an application for graph exploration. It offers semi-natural language queries, rule-based styling and search for phrases.

Interfaces COVID GRAPH Genes imely. In addition, ACE2 is a potential Figure 2: Visual Graph Explorer by yWorks Figure 3: SemSpect by derivo Is there a model for a corona virus gene? Is there a model for a corona virus gene? MASYMOS_ANNOTA... 1 MASYMOS_RESOUR... 1 MASYMOS_SBML_M... 1 Figure 4: Bloom by Neo4J

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