Modern data analytics COVID-19 analysis

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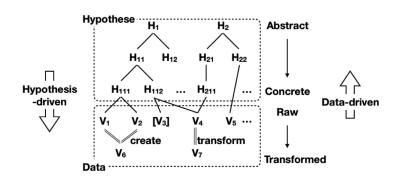
KUL

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Methodology

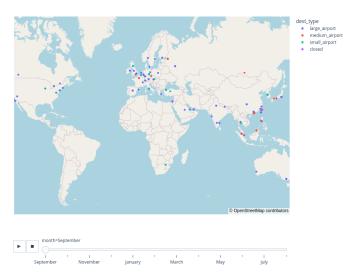


Data sources

- Economical information (Bureau of Economic Analysis, 2022)
- **③** The CORD-19 dataset (Lucy et al., 2021)

Graph mining

Destination of Chinese international flights



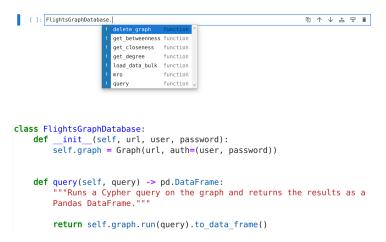
Graph mining

- Degree $c_D(i) = \sum_{j}^{N} x_{ij}$ where i, j are nodes and x is the adjacency matrix.
- Betweenness $c_B(i) = \sum_{s,t \in i} \frac{\sigma(s,t|i)}{\sigma(s,t)}$ where i is a node, s and t are source and target nodes, $\sigma(s,t)$ is the number of shortest paths between the source and target and $\sigma(s,t|i)$ is the number of shortest paths passing through the node i.
- Closeness $c_C(i) = \left[\sum_{j=1}^{N} d(i,j)\right]^{-1}$ where d is a distance metric.

(Brandes, 2008; Opsahl et al., 2010)



Graph mining - the code



Graph mining - degree centrality

The degree of the most popular American airports over time (weekly)



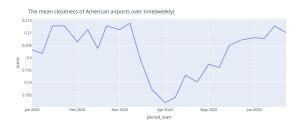
Graph mining - betweenness centrality

The betweenness of the most popular American airports over time (weekly)



Graph mining - closeness centrality

```
def get_closeness(self) -> pd.DataFrame:
    # First, create a projection
    self.query("CALL gds.graph.drop('flights', false)")
    self.query("CALL gds.graph.project('flights', 'Airport', 'FLIGHT') YIELD "")
    # Then, calculate the centrality for each node
    result = self.query("""CALL gds.beta.closeness.stream('flights')
    YIELD noded, score MATCH (n:Airport)
    WHERE ID(n) = nodeId RETURN n.name, n.code, score"")
    # Finally, sorting the results by degree
    result = result.sort_values("score", ascending=False)
    return result
```



Text mining - preprocessing

```
def preprocess text(self, series: pd.Series) -> pd.Series:
    """Applies some preprocessing steps to a collection of documents."""
    # Remove stopwords
    series = [gensim.parsing.preprocessing.remove stopwords(i)
              for i in seriesl
    # Stem
    series = gensim.parsing.porter.PorterStemmer().stem documents(series)
    # Remove numeric
    series = [gensim.parsing.preprocessing.strip numeric(i)
              for i in series]
    # Remove punctuation
    series = [gensim.parsing.preprocessing.strip punctuation(i)
              for i in seriesl
    # Remove special characters
    series = [re.sub("\\W+", " ", i) for i in series]
    # Remove short words
    series = pd.Series([gensim.parsing.preprocessing.strip short(i)
                        for i in series])
    return series
```

Text mining - latent dirichet allocation

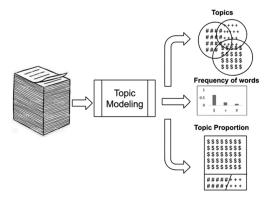
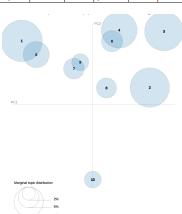


Figure: The area of topic modelling is concerned with making inferences from a huge collection of documents in terms of topics, frequency of words and topic proportions. Taken from Chauchan & Shah (2022)

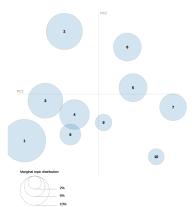
Text mining - before the pandemic

Topic 1	Topic 2	Topic 3	Topic 4	Topic 5	Topic 6	Topic 7	Topic 8	Topic 9	Topic 10
cell	health	patient	cov	cell	infect	immun	vaccin	calv	der
protein	diseas	respiratori	detect	protein	respiratori	vaccin	viru	protein	die
viru	develop	studi	viru	activ	viral	cell	infect	viru	und
viral	model	infect	sequenc	lung	group	respons	influenza	structur	model
infect	public	associ	human	express	studi	protein	studi	studi	ein
activ	emerg	children	respiratori	induc	viru	antigen	result	result	network
express	data	clinic	virus	airwai	associ	antibodi	ibv	effect	von
rna	infecti	influenza	pcr	result	virus	specif	virus	test	mit
virus	studi	case	infect	increas	caus	express	protect	differ	bei
host	risk	hospit	sars	mice	patient	infect	effect	domain	cell



Text mining - during pandemic

Topic 1	Topic 2	Topic 3	Topic 4	Topic 5	Topic 6	Topic 7	Topic 8	Topic 9	Topic 10
health	patient	covid	test	patient	cov	protein	health	mask	der
covid	covid	model	covid	cell	sars	cov	studi	effect	die
studi	hospit	vaccin	model	covid	covid	cell	mental	covid	und
research	risk	case	data	diseas	infect	sars	depress	cov	text
data	studi	studi	method	treatment	patient	ace	associ	sars	studi
care	clinic	infect	detect	immun	antibodi	viral	anxieti	aerosol	infect
pandem	sever	number	perform	activ	respiratori	bind	women	patient	activ
provid	associ	effect	result	studi	coronaviru	covid	social	studi	gene
develop	group	data	studi	clinic	sever	viru	covid	infect	effect
social	patients	time	sampl	associ	studi	drug	effect	risk	model



```
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```

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