# COVID-19 CASE SURVEILLANCE DATA

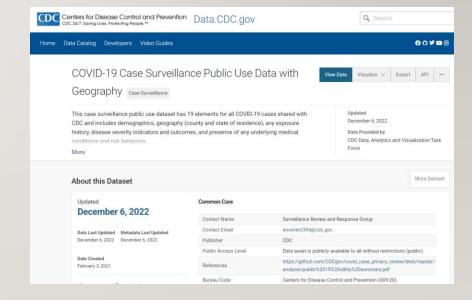
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**GROUP 3** 

# BACKGROUND INFORMATION ABOUT THE DATASET

- The dataset was from the Center for Disease and Control Prevention (CDC), and it shows data on Covid-19 cases.
- Link to Data Set: <a href="https://data.cdc.gov/Case-Surveillance/COVID-I9-Case-Surveillance-Public-Use-Data-with-Ge/n8mc-b4w4">https://data.cdc.gov/Case-Surveillance-Public-Use-Data-with-Ge/n8mc-b4w4</a>
- Data Set size: I I.2 GB



## HOW UNIQUE IS OUR DATASET?

- Our dataset is unique because we aren't just looking at the number of Covid-19 cases all around the country, but we are looking at if those with underlying conditions were more affected and had harsher outcomes.
- We are also looking for the average age group that was affected the most by Covid-19, and draw up conclusions as to why this happened.

#### **CLUSTER SPECIFICATIONS**

- Run the command Hadoop version to show the version of your Hadoop (Our version is 3.1.2)
- Run the command **Iscpu** to get the CPU specifications (The number of CPUs used is 8, and the CPU speed is 1995.309 MHZ)
- Run the command yarn node -list -all to get the total number of nodes used (Total number of nodes is 3)
- Run the command **free** –**h** to get the memory specifications

### # CPUs and CPU Speed

#### jmarti168@144.24.14.145 narti168@144.24.14.145's password: ast login: Tue Dec 6 11:23:11 2022 from 38-34-104-182.starry-inc.net x86\_64 32-bit, 64-bit Little Endian Bvte Order: n-line CPU(s) list: 0-7 Thread(s) per core: ore(s) per socket: Socket(s): NUMA node(s): GenuineIntel CPU family: Intel(R) Xeon(R) Platinum 8167M CPU @ 2.00GHz odel name:

1995.309

## Memory Size

-bash-4.29	free -h						
	total	used	free	shared	buff/cache	available	
Mem:	58G	19G	16G	2.7G	22G	35G	
Swap:	8.0G	22M	8.0G				
-bash-4.29	5						

### Hadoop Version

```
-bash-4.2$ hadoop version
Hadoop 3.1.2
Source code repository ssh://git@bitbucket.oci.oraclecorp.com:7999/bdcs/a
Compiled by root on 2022-10-26T22:15Z
Compiled with protoc 2.5.0
From source with checksum b367ca15864aef16725a3035859c9ece
This command was run using /usr/odh/1.1.2/hadoop/hadoop-common-3.1.2.jar
-bash-4.2$
```

#### # of Nodes

```
-bash-4.2$ yarn node -list -all 22/12/07 02:04:21 INFO client.RMPro 22/12/07 02:04:21 INFO client.AHSPr Total Nodes:3

Node-Id

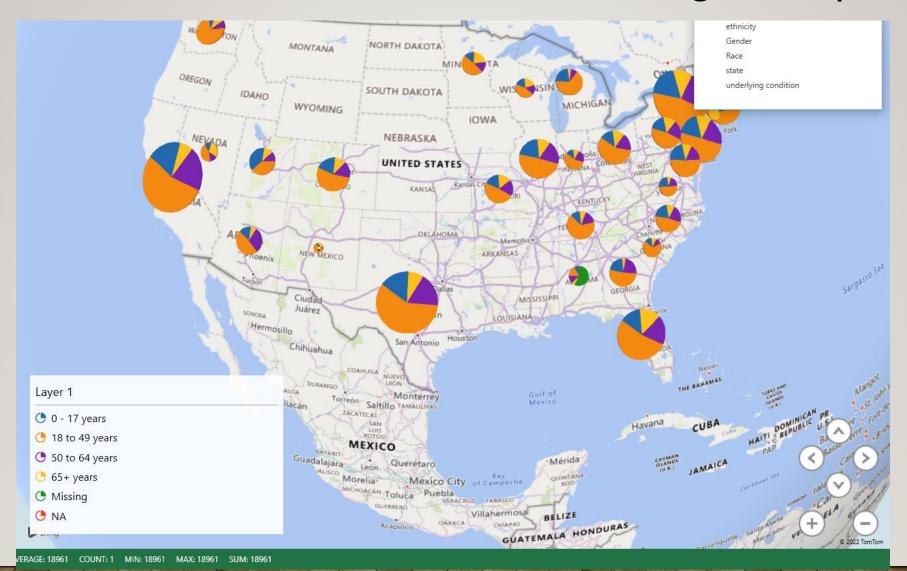
Node-S bigdaiwn1.sub02180640120.trainingvc bigdaiwn0.sub02180640120.trainingvc bigdaiwn2.sub02180640120.trainingvc -bash-4.2$
```

#### UPLOADING THE DATASET TO HDFS

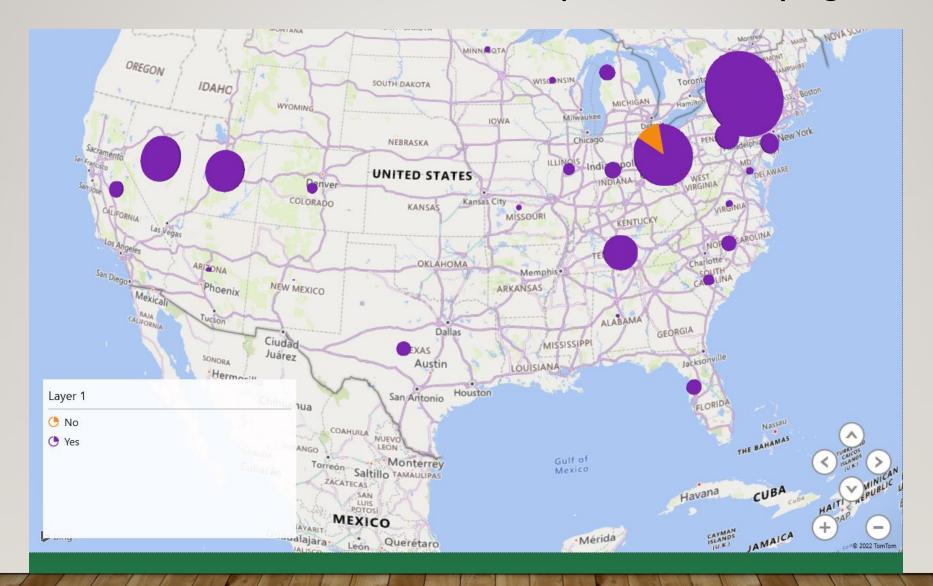
- We found two ways to upload the dataset to HDFS:
- I. Using the command **split -I [number] [filename]** and upload to Google Drive
- 2. Uploading the dataset tsv or csv file to Google Drive (since they provide 15Gb of storage data) and running the command below to upload it onto

HDFS: wget --load-cookies /tmp/cookies.txt
"https://docs.google.com/uc?export=download&confirm=\$(wget -quiet --save-cookies /tmp/cookies.txt --keep-session-cookies --nocheck-certificate
'https://docs.google.com/uc?export=download&id=FILEID'-O- |
sed -rn 's/.\*confirm=([0-9A-Za-z\_]+).\*/\1\n/p')&id=1s9aKPqcQq8id8oGgW6HBCQzuXKBR1xO" -O covid19data.csv &&
rm -rf /tmp/cookies.txt

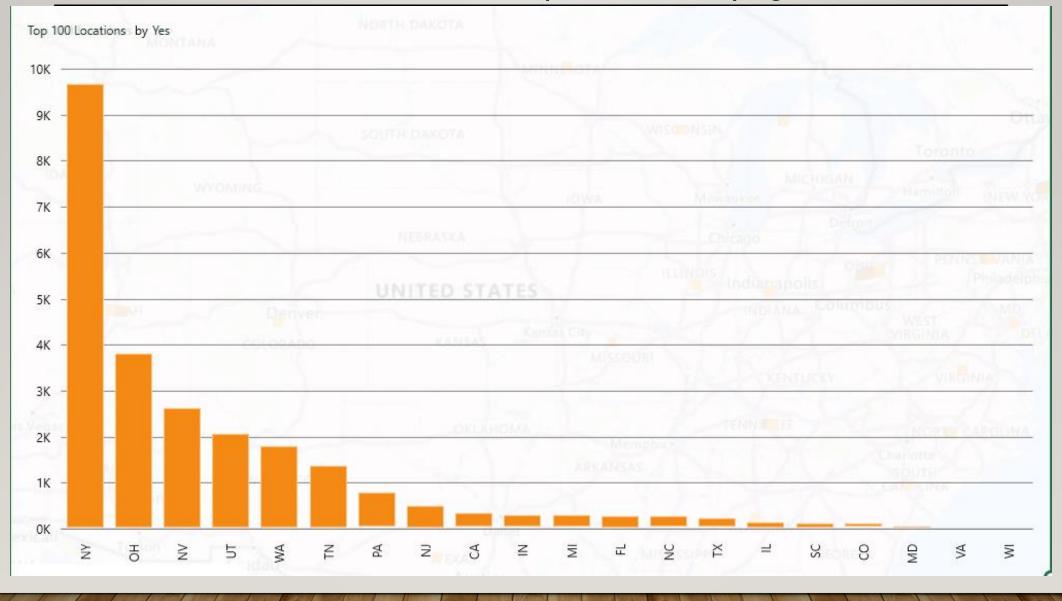
# Visualization of Covid-19 Cases Per Age Group



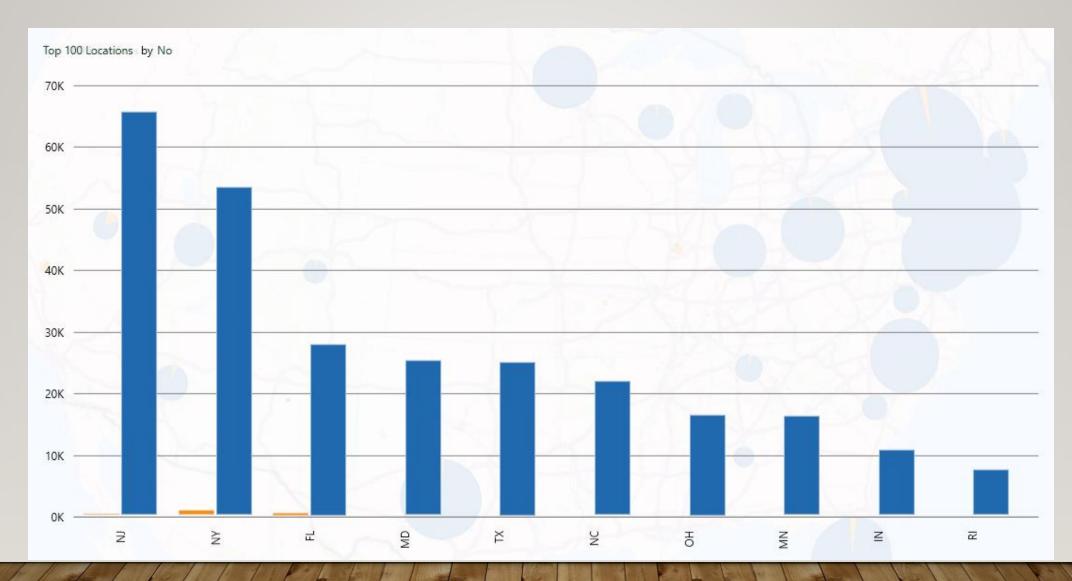
# Visualization of Covid-19 Cases With Reported Underlying Conditions



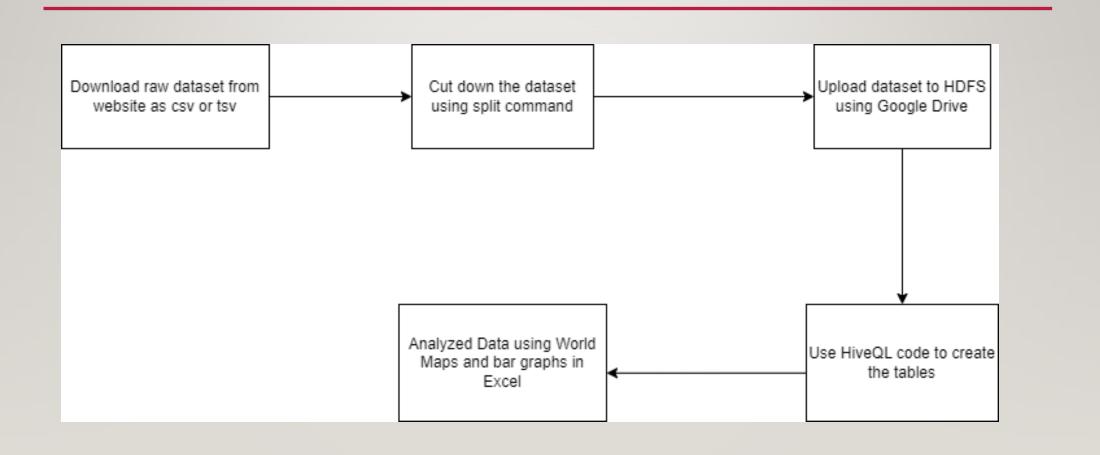
#### Visualization of Covid-19 Cases With Reported Underlying Conditions



#### Visualization of Covid-19 Cases Per State and Death Count



# **WORKFLOW CHART**



# UNDERSTANDING EPIDEMIC DATA AND STATISTICS: A CASE STUDY OF COVID-19

- Tries to interpret the number of people being infected in each country
- Evaluate how effective the policies each country implements in lowering cases
- Examines covid confirmed cases, recovered, and deaths
- Each country's policies and reactions to the outbreak determine how fast it spread.

# DATA MINING AND ANALYSIS OF SCIENTIFIC RESEARCH DATA RECORDS ON COVID-19 MORTALITY, IMMUNITY, AND VACCINE DEVELOPMENT

- Purpose of the research was to investigate and determine early warning systems developed in previous epidemic responses to contain the virus from spreading.
- Examined Covid-19 scientific literature regarding Covid-19 mortality, vaccines, and immunity via data mining.
- Bibliometric analysis was done using the Web of Science Analysis Results tool to search the most dominant keywords and related concepts with Covid-19.
- Factorial analysis was done using R Studio to examine the correlation between different concepts (mortality, immunity, & vaccine development) as well as generate visualizations such as tree maps and conceptual structure maps.

# **GITHUB LINK**

GitHub Link:

https://github.com/mikeOnthemic/G5\_Big\_Data\_4560

#### **WORK CITED**

- Hoseinpour Dehkordi, A., Alizadeh, M., Derakhshan, P., Babazadeh, P., & Jahandideh, A. (2020). Understanding epidemic data and statistics: A case study of COVID-19. Journal of medical virology, 92(7), 868–882.
   https://doi.org/10.1002/jmv.25885
- Radanliev, P., De Roure, D., & Walton, R. (2020). Data mining and analysis of scientific research data records on Covid-19 mortality, immunity, and vaccine development-In the first wave of the Covid-19 pandemic. *Diabetes & Metabolic Syndrome: Clinical Research & Reviews*, 14(5), 1121-1132.