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Date: 12/7/2022

Lab Tutorial

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Covid 19 Surveillance Data

Objectives:

In this lab, you will:

- Get the dataset from Google Drive using wget
- Upload the dataset to the tmp folder
- Create a database within beeline
- Create tables based on the data using HiveQL commands
- Download the data to the local computer
- Use Excel and PowerBI for visualization of the data

Platform Specifications:

- Oracle Cloud
- CPU Speed: 1995.309 MHz:
- # of CPU Cores: 32
- # of nodes: 3
- Total Memory Size: 58GB

1. open a shell terminal – git bash, minty, putty etc- and run the ssh command to connect to the Hadoop Cloud.

```
$ssh yourusername@ipaddress [144.24.14.145]
```

2. Download the covid19 dataset file using wget

```
wget --load-cookies /tmp/cookies.txt
"https://docs.google.com/uc?export=download&confirm=$(wget --quiet --save-cookies
/tmp/cookies.txt --keep-session-cookies --no-check-certificate
'https://docs.google.com/uc?export=download&id=1s-9aKPqcQq8id8oGgW6HBCQzuXKBR1x
O' -O- | sed -rn
's/.*confirm=([0-9A-Za-z_]+).*/\1\n/p')&id=1s-9aKPqcQq8id8oGgW6HBCQzuXKBR1xO" -O
covid19data.csv
```

```
-bash-4.2$ wget --load-cookies /tmp/cookies.txt "https://docs.google.com/uc?export=download&confirm=$(wget --quiet --save-cookies /tmp/cookies.txt --keep-session-cookies --no-check-certificate 'https://docs.google.com/uc?export=download&id=FILEID' -O- | sed -rn 's/.*confirm=([0-9A-Za-z_]+).*/\1\n/p')&id=1s-9aKPqcQq8id8oGgW6HBCQzuXKBR1xO" -O covid19data.csv && rm -rf /tmp/cookies.txt
--2022-12-07 19:33:20-- https://docs.google.com/uc?export=download&confirm=8id=1s-9aKPqcQq8id8oGgW6HBCQzuXKBR1xO
Resolving docs.google.com (docs.google.com)... 142.250.188.238, 2607:f8b0:4007:80f::200e
Connecting to docs.google.com (docs.google.com)[142.250.188.238]:443... connected.
HTTP request sent, awaiting response... 303 See Other
Location: https://doc-0s-0k-docs.googleusercontent.com/docs/securesc/ha0ro937gcuc717deffksulhg5h7mbp1/965h5kf02121rjd4h34tg9i6bm7ruiq1/1670441550000/13778178939762846769/"1s-9aKPqcQq8id8oGgW6HBCQzuXKBR1xO?e=download&uiid=aea2c644-1db7-4551-ba45-6c29b64399ef [following]
Warning: redirect not supported in HTTP.
--2022-12-07 19:33:20-- https://doc-0s-0k-docs.googleusercontent.com/docs/securesc/ha0ro937gcuc717deffksulhg5h7mbp1/965h5kf02121rjd4h34tg9i6bm7ruiq1/1670441550000/13778178939762846769/"1s-9aKPqcQq8id8oGgW6HBCQzuXKBR1xO?e=download&uiid=aea2c644-1db7-4551-ba45-6c29b64399ef
Resolving doc-0s-0k-docs.googleusercontent.com (doc-0s-0k-docs.googleusercontent.com)... 142.250.68.65, 2607:f8b0:4007:818::2001
Connecting to doc-0s-0k-docs.googleusercontent.com (doc-0s-0k-docs.googleusercontent.com)[142.250.68.65]:443... connected.
HTTP request sent, awaiting response... 200 OK
Length: 2148939052 (2.06) [application/octet-stream]
Saving to: 'covid19data.csv'

100%[=====] 2,148,939,052 187MB/s in 11s

2022-12-07 19:33:32 (183 MB/s) - 'covid19data.csv' saved [2148939052/2148939052]
```

3. You have to upload the files to hdfs folder coviddata. Run the following HDFS commands to create and list coviddata directory in HDFS:

```
$ hdfs dfs -mkdir tmp/covid19data
$ hdfs dfs -put covid19data.csv tmp/covid19data/
```

```
-bash-4.2$ hdfs dfs -mkdir tmp/covid19data/
-bash-4.2$ hdfs dfs -put covid19data.csv tmp/covid19data/
-bash-4.2$ hdfs dfs -ls tmp/covid19data/
Found 1 items
-rw-r--r-- 3 pilabac hdfs 2148939052 2022-12-07 19:38 tmp/covid19data/covid19data.csv
-bash-4.2$ |
```

4. Open hive

\$ beeline

```
-bash-4.2$ beeline
SLF4J: Class path contains multiple SLF4J bindings.
SLF4J: Found binding in [jar:file:/usr/odh/1.1.2/hive/lib/log4j-slf4j-impl-2.17.1.jar!/org/slf4j/impl/StaticLoggerBinder.class]
SLF4J: Found binding in [jar:file:/usr/odh/1.1.2/hadoop/lib/slf4j-log4j12-1.7.25.jar!/org/slf4j/impl/StaticLoggerBinder.class]
SLF4J: See http://www.slf4j.org/codes.html#multiple_bindings for an explanation.
SLF4J: Actual binding is of type [org.apache.logging.slf4j.Log4jLoggerFactory]
Connecting to jdbc:hive2://bigdaiwn0.sub02180640120.trainingvcn.oraclevcn.com:2181,bigdaimn0.sub02180640120.trainingvcn.oraclevcn.com:2181/default;password=pilabac;serviceDiscoveryMode=zooKeeper;user=pilabac;zooKeeperNamespace=hiveserver2
22/12/07 19:43:42 [main-EventThread]: ERROR impls.EnsembleTracker: Invalid config event received: {server.1=bigdaimn0.sub02180640120.trainingvcn.oraclevcn.com:2888:3888:participant, version=0, server.3=bigdaiwn0.sub02180640120.trainingvcn.oraclevcn.com:2888:3888:participant, server.2=bigdaiun0.sub02180640120.trainingvcn.oraclevcn.com:2888:3888:participant}
22/12/07 19:43:42 [main-EventThread]: ERROR impls.EnsembleTracker: Invalid config event received: {server.1=bigdaimn0.sub02180640120.trainingvcn.oraclevcn.com:2888:3888:participant, version=0, server.3=bigdaiwn0.sub02180640120.trainingvcn.oraclevcn.com:2888:3888:participant, server.2=bigdaiun0.sub02180640120.trainingvcn.oraclevcn.com:2888:3888:participant}
22/12/07 19:43:42 [main]: INFO jdbc.HiveConnection: Connected to bigdaiun0.sub02180640120.trainingvcn.oraclevcn.com:10000
Connected to: Apache Hive (version 3.1.2)
Driver: Hive JDBC (version 3.1.2)
Transaction isolation: TRANSACTION_REPEATABLE_READ
Beeline version 3.1.2 by Apache Hive
0: jdbc:hive2://bigdaiwn0.sub02180640120.traig> |
```

5. Create your own database and use that database

\$ create database Covid19;
\$ use Covid19;

```
0: jdbc:hive2://bigdaiwn0.sub02180640120.traig> use covid19;
INFO : Compiling command(queryId=hive_20221207195141_6f7b24f9-3e2d-45c4-80c0-665e3464f455): use covid19
INFO : Concurrency mode is disabled, not creating a lock manager
INFO : Semantic Analysis Completed (retrial = false)
INFO : Returning Hive schema: Schema(fieldSchemas:null, properties:null)
INFO : Completed compiling command(queryId=hive_20221207195141_6f7b24f9-3e2d-45c4-80c0-665e3464f455); Time taken: 0.03 seconds
INFO : Concurrency mode is disabled, not creating a lock manager
INFO : Executing command(queryId=hive_20221207195141_6f7b24f9-3e2d-45c4-80c0-665e3464f455): use covid19
INFO : Starting task [Stage-0:DDL] in serial mode
INFO : Completed executing command(queryId=hive_20221207195141_6f7b24f9-3e2d-45c4-80c0-665e3464f455); Time taken: 0.216 seconds
INFO : OK
INFO : Concurrency mode is disabled, not creating a lock manager
No rows affected (0.26 seconds)
0: jdbc:hive2://bigdaiwn0.sub02180640120.traig> |
```

6. Create external table “Covid19Data”

- - create the covid19 table on comma-separated covid19data
CREATE EXTERNAL TABLE IF NOT EXISTS covid19 (case_months string,
res_state string,
state_fips_code string,
res_country string,
county_fips_county string,
age_group string,
sex string,
race string,
ethnicity string,
case_positive_specimen_interval int,
case_onset_interval int,
process string,

```
exposure_yn string,  
current_status string,  
symptom_status string,  
hosp_yn string,  
icu_yn string,  
death_yn string,  
underlying_conditions_yn string)  
row format delimited fields terminated by ","  
stored as textfile location '/user/ufong9/tmp/covid19data'  
tblproperties ('skip.header.line.count' = '1');
```

```

0: jdbc:hive2://bigdaiwn0.sub02180640120.trai> describe formatted covid19;
INFO : Compiling command(queryId=hive_20221207195421_3b6645da-a0a6-498b-be0e-f6a4c47690eb): describe formatted covid19
INFO : Concurrency mode is disabled, not creating a lock manager
INFO : Semantic Analysis Completed (retrial = false)
INFO : Returning Hive schema: Schema(FieldSchemas:[FieldSchema(name:col_name, type:string, comment:from deserializer), FieldSchema(n
me:data_type, type:string, comment:from deserializer), FieldSchema(name:comment, type:string, comment:from deserializer)], properties
null)
INFO : Completed compiling command(queryId=hive_20221207195421_3b6645da-a0a6-498b-be0e-f6a4c47690eb); Time taken: 0.046 seconds
INFO : Concurrency mode is disabled, not creating a lock manager
INFO : Executing command(queryId=hive_20221207195421_3b6645da-a0a6-498b-be0e-f6a4c47690eb): describe formatted covid19
INFO : Starting task [Stage-0:DDL] in serial mode
INFO : Completed executing command(queryId=hive_20221207195421_3b6645da-a0a6-498b-be0e-f6a4c47690eb); Time taken: 0.28 seconds
INFO : OK
INFO : Concurrency mode is disabled, not creating a lock manager

```

col_name	data_type	comment
# col_name	data_type	comment
case_months	string	
res_state	string	
state_fips_code	string	
res_country	string	
county_fips_county	string	
age_group	string	
sex	string	
race	string	
ethnicity	string	
case_positive_specimen_interval	int	
case_onset_interval	int	
process	string	
exposure_yn	string	
current_status	string	
symptom_status	string	
hosp_yn	string	
icu_yn	string	
death_yn	string	
underlying_conditions_yn	string	
	NULL	NULL
# Detailed Table Information	NULL	NULL
Database:	covid19	NULL
OwnerType:	USER	NULL
Owner:	tfong9	NULL
CreateTime:	Tue Dec 06 19:25:44 GMT 2022	NULL
LastAccessTime:	UNKNOWN	NULL
Retention:	0	NULL
Location:	hdfs://bigdaiwn0.sub02180640120.trainingvcn.oraclevcn.com:8020/user/tfong9/tmp/coviddata	NULL
Table Type:	EXTERNAL_TABLE	NULL
Table Parameters:	NULL	NULL
	EXTERNAL	TRUE
	bucketing_version	2
	numFiles	1
	skip.header.line.count	1
	totalSize	2148939052
	transient_lastDdlTime	1670354744
	NULL	NULL
# Storage Information	NULL	NULL
SerDe Library:	org.apache.hadoop.hive.serde2.lazy.LazySimpleSerDe	NULL
InputFormat:	org.apache.hadoop.mapred.TextInputFormat	NULL
OutputFormat:	org.apache.hadoop.hive.q1.io.HiveIgnoreKeyTextOutputFormat	NULL
Compressed:	No	NULL
Num Buckets:	-1	NULL
Bucket Columns:	[]	NULL
Sort Columns:	[]	NULL
Storage Desc Params:	NULL	NULL
	field.delim	,
	serialization.format	,

Run the following HiveQL at the query editor to see how the dataset looks like

```
select * from covid19 limit 10;
```

covid19_case_months	covid19_res_state	covid19_state_fips_code	covid19_res_country	covid19_county_fips_code	covid19_age_group	covid19_sex	covid19_race	covid19_ethnicity	covid19_case_positive_specimen_interval	covid19_case_onset_interval
covid19_process	covid19_exposure_yn	covid19_current_status	covid19_symptom_status	covid19_hosp_yn	covid19_icu_yn	covid19_death_yn	covid19_underlying_conditions_yn			
2021-12	CA	06	VENTURA	06111	18 to 49 years	Female	White	Non-Hispanic/Latino	NULL	NULL
Missing	Missing	Laboratory-confirmed case	Unknown	Missing	Missing	Missing				
2021-09	TX	48	TARRANT	48439	18 to 49 years	Male	White	Non-Hispanic/Latino	NULL	NULL
Missing	Missing	Laboratory-confirmed case	Missing	Missing	Missing	Missing				
2022-01	MA	25	MIDDLESEX	25017	18 to 49 years	Female	Unknown	Unknown	0	NULL
Missing	Missing	Laboratory-confirmed case	Missing	Missing	Missing	Missing				
2020-12	NY	36	KINGS	36047	65+ years	Female	White	Non-Hispanic/Latino	0	0
Missing	Missing	Laboratory-confirmed case	Symptomatic	Missing	Missing	Missing				
2022-01	NJ	34	ESSEX	34013	0 - 17 years	Male	White	Non-Hispanic/Latino	0	NULL
Missing	Missing	Laboratory-confirmed case	Missing	No	Missing	No				
2022-06	CA	06	SACRAMENTO	06067	18 to 49 years	Female	Unknown	Non-Hispanic/Latino	NULL	NULL
Missing	Missing	Laboratory-confirmed case	Unknown	Missing	Missing	Missing				
2021-12	NJ	34	OCEAN	34029	50 to 64 years	Female	White	Non-Hispanic/Latino	0	NULL
Missing	Missing	Laboratory-confirmed case	Missing	No	Missing	No				
2021-09	NY	36	MONROE	36055	0 - 17 years	Female	Black	Non-Hispanic/Latino	NULL	0
Missing	Missing	Laboratory-confirmed case	Symptomatic	No	Missing	No				
2021-07	FL	12	PALM BEACH	12099	18 to 49 years	Male	Black	Non-Hispanic/Latino	NULL	0
Missing	Missing	Laboratory-confirmed case	Symptomatic	No	Missing	No				
2022-05	FL	12	PINELLAS	12103	18 to 49 years	Male	White	Non-Hispanic/Latino	0	NULL
Missing	Missing	Laboratory-confirmed case	Missing	Missing	Missing	Missing				

7. Create external table "patient_profile"

-- create the patient_profile table on comma-seperated covid19data

```
CREATE EXTERNAL TABLE IF NOT EXISTS patient_profile(month STRING, age_group
STRING, sex STRING, race STRING, ethnicity STRING, res_state STRING,
underlying_conditions STRING, death STRING)
row format delimited fields terminated by ","
STORED AS TEXTFILE LOCATION '/user/ufong9/tmp/covid19data';
```

insert overwrite table patient_profile

```
Select case_months, age_group, sex, race, ethnicity, res_state, underlying_conditions_yn,
death_yn
from covid19;
```

Now run the following HiveQL at the query editor to see how the dataset looks like

```
Select * from patient_profile limit 10;
```

patient_profile.month	patient_profile.age_group	patient_profile.sex	patient_profile.race	patient_profile.ethnicity	patient_profile.res_state
2021-12	18 to 49 years	Female	White	Non-Hispanic/Latino	CA
2021-09	18 to 49 years	Male	White	Non-Hispanic/Latino	TX
2022-01	18 to 49 years	Female	Unknown	Unknown	MA
2020-12	65+ years	Female	White	Non-Hispanic/Latino	NY
2022-01	0 - 17 years	Male	White	Non-Hispanic/Latino	NJ
2022-06	18 to 49 years	Female	Unknown	Non-Hispanic/Latino	CA
2021-12	50 to 64 years	Female	White	Non-Hispanic/Latino	NJ
2021-09	0 - 17 years	Female	Black	Non-Hispanic/Latino	NY
2021-07	18 to 49 years	Male	Black	Non-Hispanic/Latino	FL
2022-05	18 to 49 years	Male	White	Non-Hispanic/Latino	FL

10 rows selected (0.341 seconds)
0: jdbc:hive2://bigdaiwn0.sub02180640120.traig> |

8. Now run the following HiveQL at the Query editor to see the number of cases

```
select case_months, count(sex) as number_of_cases
from covid19
group by case_months;
```

case_months	number_of_cases
2022-01	2648280
2020-05	153218
2020-04	210489
2021-12	1412096
2022-03	175036
2020-03	110842
2020-07	345947
2020-12	985109
2022-04	269741
2020-09	183285
2021-07	299067
2022-02	423524
2020-11	598596
2021-01	1024501
2021-03	380664
2021-06	77038
2022-10	109129
2021-05	122949
2022-09	221328
2021-09	426063
2020-01	548
2020-10	259393
2022-07	598233
2022-05	699285
2022-06	519352
2021-11	307190
2022-08	451627
2020-02	1227
2021-02	391593
2021-08	585600
2021-10	295130
2020-06	213922
2021-04	274905
2020-08	225092

9. Now download data into your PC

- - download to local file

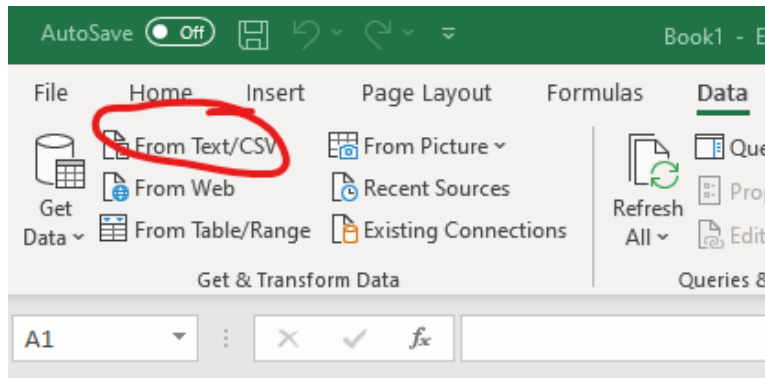
```
hdfs dfs -get tmp/covid19data/000000_0
```

- - download file to your PC

```
scp tfong9@144.24.14.145:/home/tfong9/000000_0 covid19data.csv
```

10. Loading Data into and Visualizing using Power Map in Excel

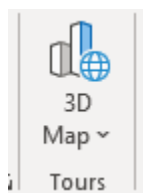
Import your covid19data.csv into Excel. You should then be able to open 3d maps for visualization.



Rename the table columns into the following: date, age, gender, race, ethnicity, state, underlying condition, and death

	A	B	C	D	E	F	G	H
1	Date	Age	Gender	Race	Ethnicity	State	Underlying conditions	Death

After renaming the columns, highlight all your columns and select the 3D maps under the "insert" tab



11. You need to select the properties and values in the layer as follows:

- For Location, add state
- For Size, click on gender (size may also appear as "height" in the beginning)
- For Category, add Age

Location 100%

☒ State State/Province ✕

+ Add Field

Size

Gender (Count - Not Blank) ▼ ✕

+ Add Field

Category

Age ✕

Time

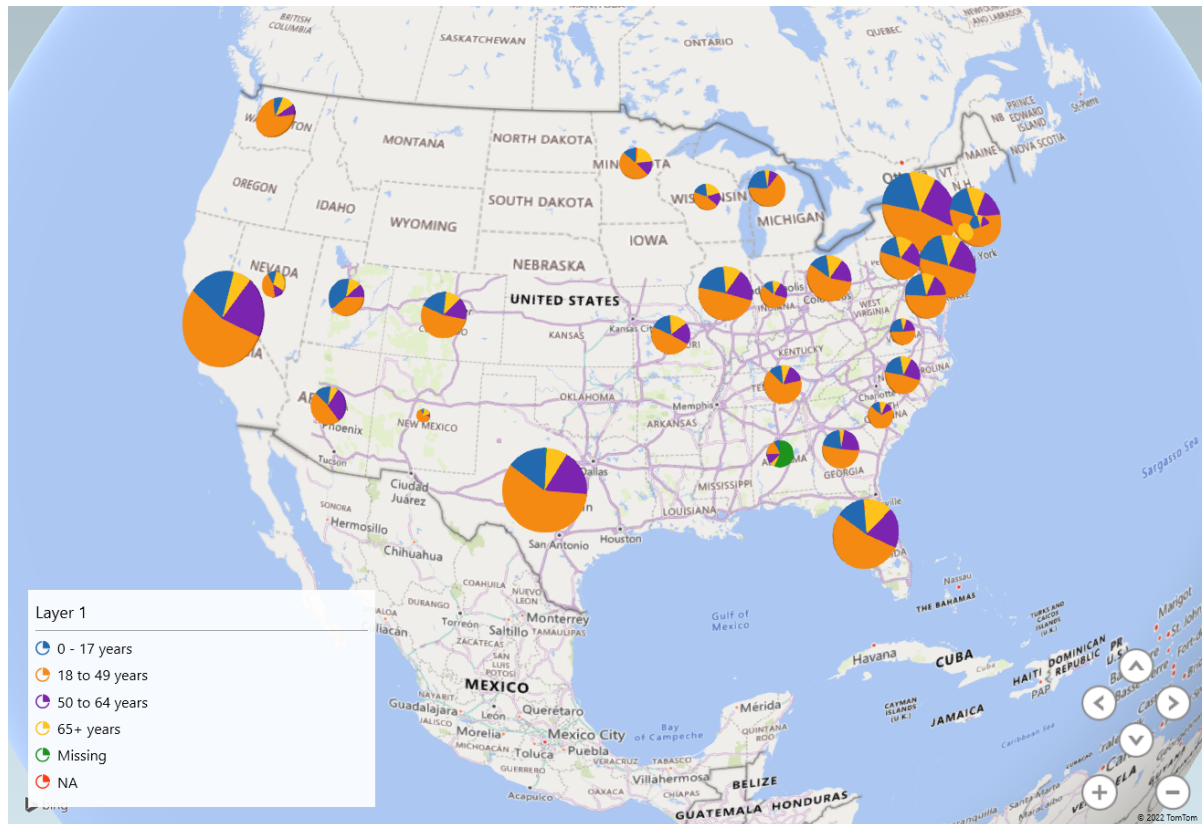
+ Add Field

▼ **Filters**

+ Add Filter

After you should change the graph to pie graphs





Afterwards, you should change the Category from **Age Groups** to **Underlying Conditions**

Category

+ Add Field

covid19data

Age

Date

Death

Ethnicity

Gender

Race

State

Underlying conditions

Add a filter to the underlying condition, and make sure to check the boxes there are only those who said yes or no.

Underlying conditions $\Sigma \rightarrow$  

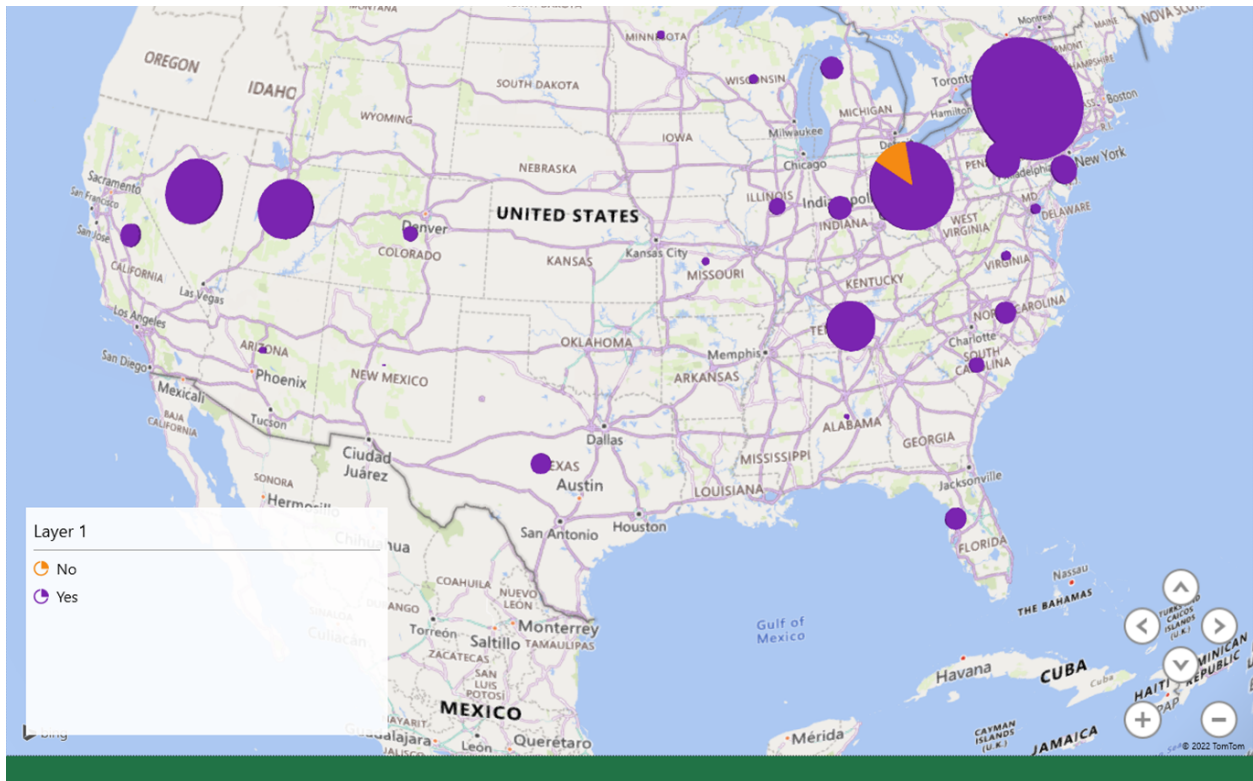
Filtered

☒ (All)



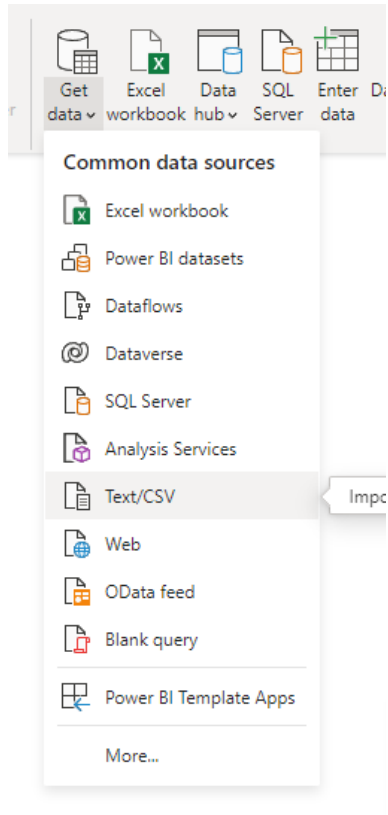
☒ No

☒ Yes

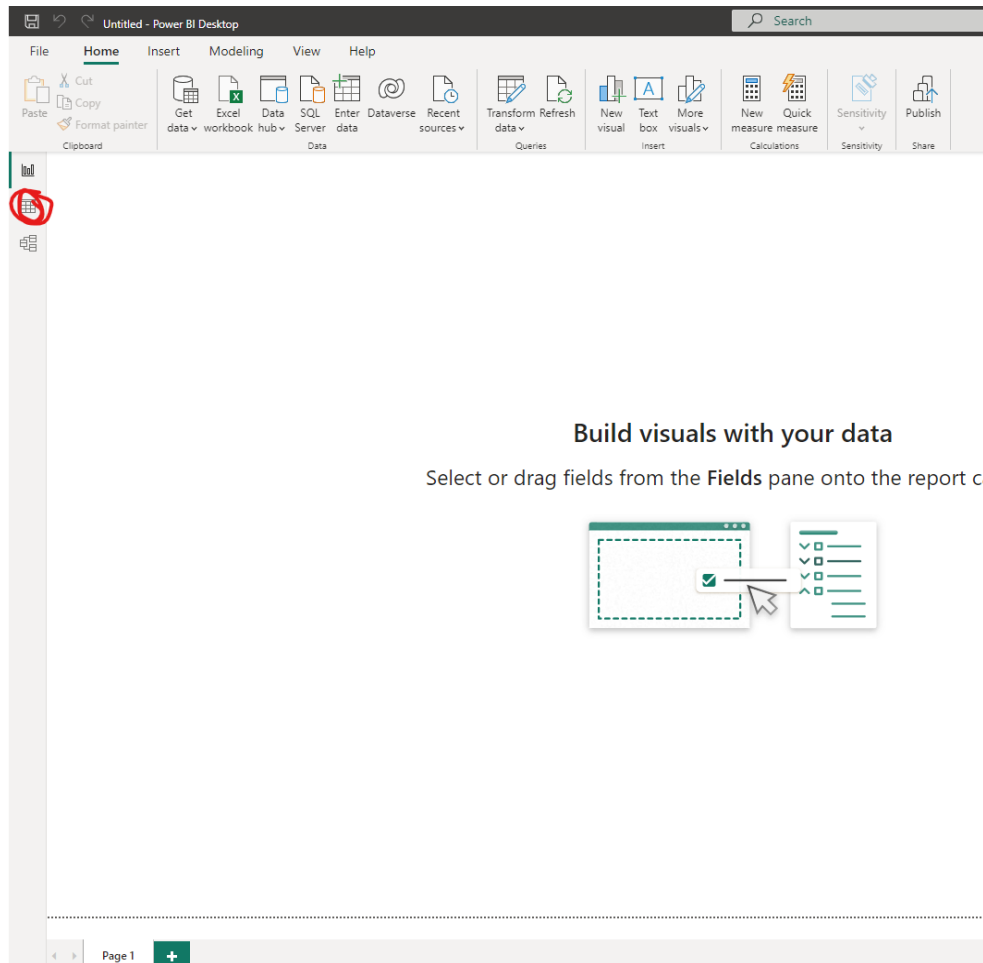


12. Loading Data into and Visualizing using PowerBI Desktop (You have to use PowerBI Desktop)

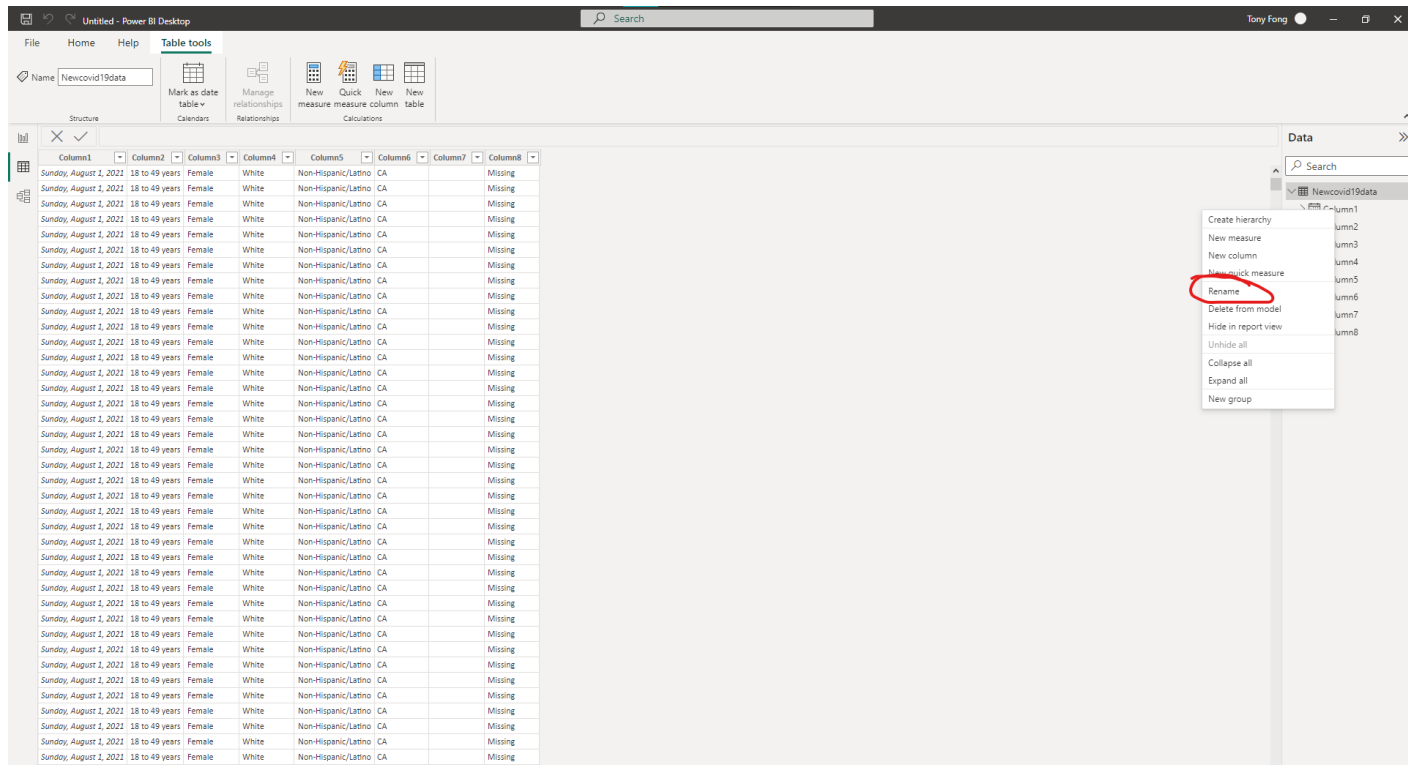
Open PowerBI desktop and load your data into PowerBI



You need to rename the columns by After loading your data onto PowerBI Desktop, go to your data as shown below:

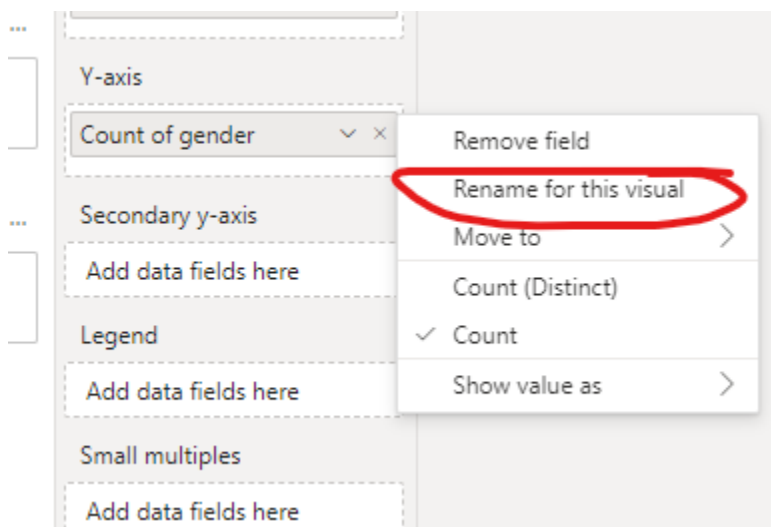


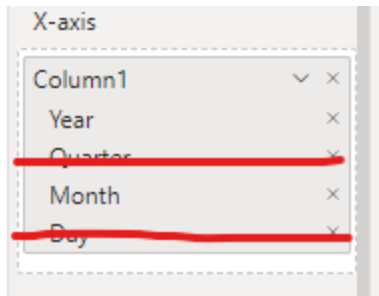
Rename each column to the corresponding names in this order: **(Date, Age, Gender, Race, Ethnicity, State, Underlying condition, Death)**



Note: Circled red is the Rename option

Next you need to select the properties and values in the layer as follows **to find the number of people who contracted covid by year and month**. First select your visualization types as a line chart. Next, format your visual, rename “count of gender” to “# of people who contracted covid19” and turn on the data label.

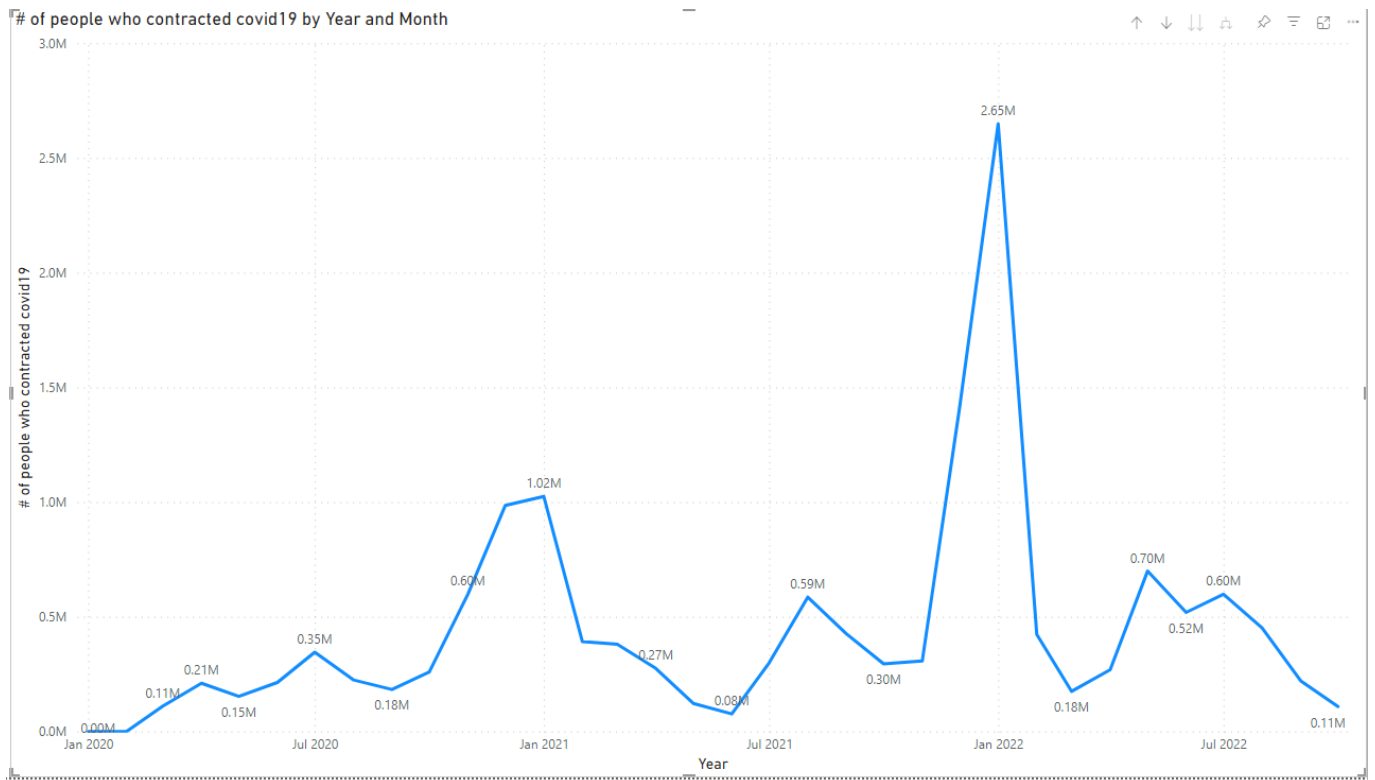




Note: Remove Quarter and Day from the date column

The screenshot shows the Power BI interface with the following components:

- Filters:** Search bar, Filters on this visual, Filters on this page, Filters on all pages.
- Visualizations:** Build visual, X-axis (Column1, Year, Month), Y-axis (# of people who cont...), Secondary y-axis, Legend, Small multiples, Tooltips, Drill through, Cross-report, Keep all filters, Add drill-through fields here.
- Fields:** Search bar, Newcovid19data, age, Column1, death, ethnicity, gender, race, state, underlying condi...
- Visual format:** Search bar, Visual, General, X-axis, Y-axis, Secondary y-axis, Legend, Small multiples, Gridlines, Zoom slider, Lines, Markers, Data labels (highlighted with a red circle), Series labels, Plot area background.



13. You need to select the properties and values in the layer as follows to find the count of death by state and filter death to only those who said yes. Then show value as percent of grand total. Also, make sure to change the visualization type to stacked bar graph.

Filters

Search

Filters on this visual

%GT Count of death is (All)

Count of death is (All)

state is (All)

Add data fields here

Filters on this page

death is Yes

Filter type

Basic filtering

Search

☒ Select all

Missing9518734

NA142293

No4081658

Unknown1152069

Yes105245

☐ Require single selection

Add data fields here

Filters on all pages

Add data fields here

Visualizations

Build visual

state

X-axis

%GT Count of death

Legend

Add data fields here

Small multiples

Add data fields here

Tooltips

Add data fields here

Drill through

Cross-reportOff

Keep all filtersOn

Add drill-through fields here

state

X-axis

%GT Count of death

Legend

Add data fields here

Small multiples

No calculation

Percent of grand total

Add data fields here

Drill through

Cross-reportOff

Keep all filtersOn

Remove field

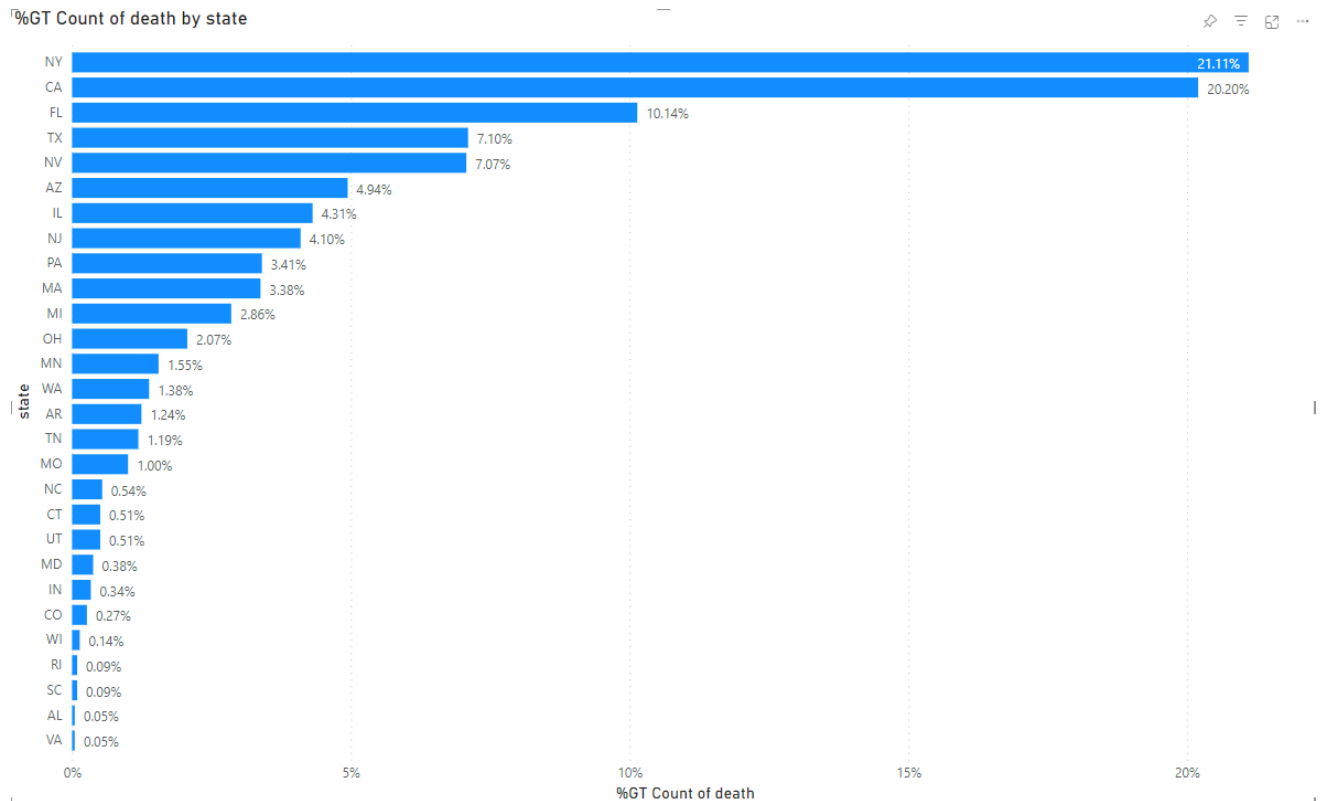
Rename for this visual

Move to

Count (Distinct)

Count

Show value as



14. You need to select the properties and values in the layer as follows to find **the count of death by gender**. Then switch the visual type to pie chart and put the gender as the legend and values as death. Then add the following filters as shown in the picture below:

References:

<https://data.cdc.gov/Case-Surveillance/COVID-19-Case-Surveillance-Public-Use-Data-with-Ge/n8mc-b4w4>

https://github.com/mike0nthemic/G5_Big_Data_4560.git