

COVID19 Data Analysis from Google Open Data using Hive and Tableau

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Objectives

- Upload data and analyze via wget
- Create a table and test data via Hive
- Clean and test data via Pig
- Store and Export data into Hadoop then Local Machine
- Visualizations and Analysis using PowerBI

Platform Specifications

- **Hostname:** bigdaiun0.sub03291929060.trainingvcn.oraclevcn.com
- **OS:** Oracle Linux Server 7.9
- **Storage:** 120 GB
- **Memory Size:** 32 GB
- **CPU:** AMD EPYC 7J13 64-Core Processor @ 2.45 Ghz
- **Cluster Version:** Hadoop 3.1.2
- **Cluster Number of Nodes:** 3

Prerequisites

- **Operating Systems**
 - Windows 10 or 11
 - MacOS
 - Linux
- **Shell terminal**
 - For windows, download and install Git Bash: <https://git-scm.com/downloads>
 - On Linux or Mac computer, use **terminal**
- **Microsoft Excel**
- **Microsoft PowerBI**
- **OpenRefine**

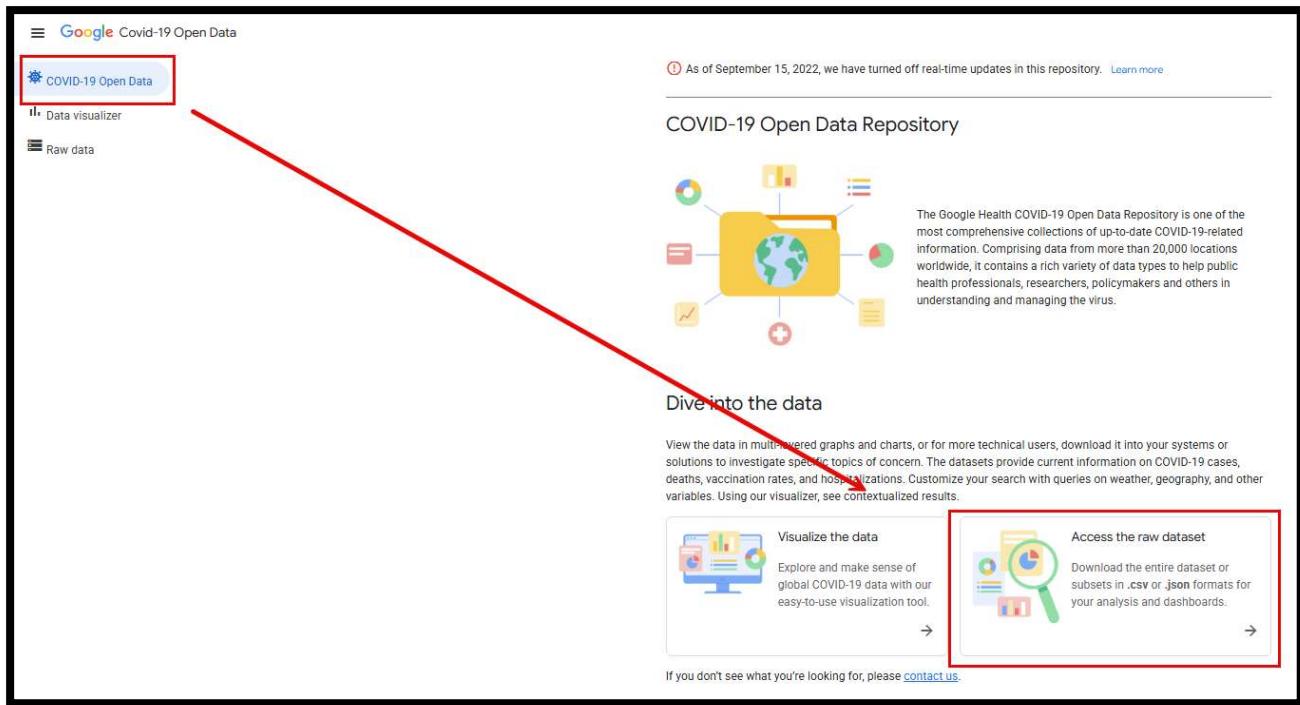
Introduction

This project demonstrates a complete big data analytic workflow using Google COVID-19 Open Data. The process includes data acquisition, data cleaning, distributed storage, table creation using Hive, aggregation of large datasets, and visualization of key results. By following this step-by-step tutorial, we apply core concepts from CIS 4560 related to big data processing, data warehousing, and analytical reporting.

Step 1: Download datasets from Google Covid-19 Open Data

The purpose of this step is to acquire raw COVID-19 data from a reliable and publicly available source. Google COVID-19 Open Data provides standardized datasets that include epidemiological, demographic, geographic, and vaccination information across multiple regions. Downloading these datasets serves as the foundation for all subsequent data processing, analysis, and visualization tasks in this project.

Go to <https://health.google.com/covid-19/open-data/>



Download the csv files of the following tables:

- Index
- Demographics
- Epidemiology
- Geography
- Health

- Hospitalizations
- Vaccinations
- By age
- By sex

Download the data					
Table	Indexed by ¹	Content	Source ²	Download	
Annotated	[key] [date]	Flat, compressed table with records from (almost) all other tables joined by date and/or key; see below for more details	All tables below	.csv	
Index	[key]	Various names and codes, useful for joining with other datasets	Wikidata, DataCommons, Eurostat	.csv	.json
Demographics	[key]	Various (current ³) population statistics	Wikidata, DataCommons, WorldBank, WorldPop, Eurostat	.csv	.json
Economy	[key]	Various (current ³) economic indicators	Wikidata, DataCommons, Eurostat	.csv	.json
Epidemiology	[key] [date]	COVID-19 cases, deaths, recoveries and tests	Various ⁴	.csv	.json
Emergency declarations	[key] [date]	Government emergency declarations and mitigation policies	LawAtlas Project	.csv	
Geography	[key]	Geographical information about the region	Wikidata	.csv	.json
Health	[key]	Health indicators for the region	Wikidata, WorldBank, Eurostat	.csv	.json
Hospitalizations	[key] [date]	Information related to patients of COVID-19 and hospitals	Various ⁴	.csv	.json
Mobility	[key] [date]	Various metrics related to the movement of people. To download or use the data, you must agree to the Google Terms of Service	Google	.csv	.json
Search trends	[key] [date]	Trends in symptom search volumes due to COVID-19. To download or use the data, you must agree to the Google Terms of Service	Google	.csv	
Vaccination access	[place_id]	Metrics quantifying access to COVID-19 vaccination sites. To download or use the data, you must agree to the Google Terms of Service	Google	.csv	
Vaccination search	[key] [date]	Trends in Google searches for COVID-19 vaccination information. To download or use the data, you must agree to the Google Terms of Service	Google	.csv	
Vaccinations	[key] [date]	Trends in persons vaccinated and population vaccination rate regarding various Covid-19 vaccines. To download or use the data, you must agree to the Google Terms of Service	Google	.csv	
Government response	[key] [date]	Government interventions and their relative stringency	University of Oxford	.csv	.json
Weather	[key] [date]	Dated meteorological information for each region	NOAA	.csv	
WorldBank	[key]	Latest record for each indicator from WorldBank for all reporting countries	WorldBank	.csv	.json
By age	[key] [date]	Epidemiology and hospitalizations data stratified by age	Various ⁴	.csv	.json
By sex	[key] [date]	Epidemiology and hospitalizations data stratified by sex	Various ⁴	.csv	.json

Clean tables using OpenRefine

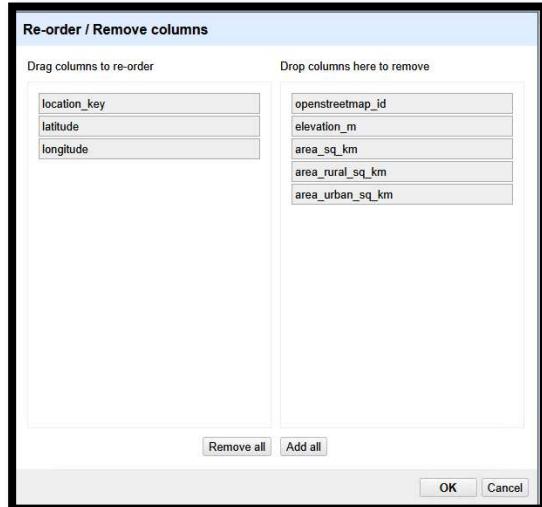
OpenRefine geography csv [Permalink](#)

22,130 rows

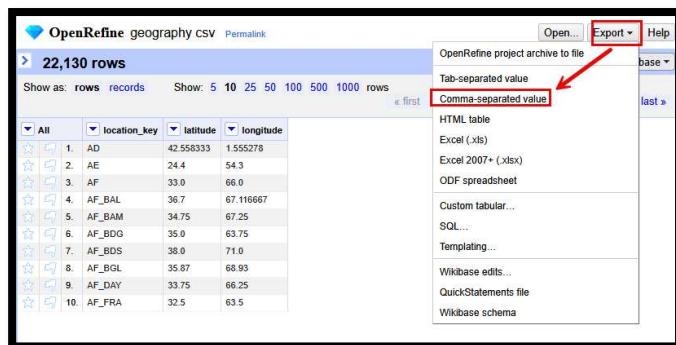
Show as: rows records Show: 5 10 25 50 100 500 1000 rows

All	location_key	openstreetmap_id	latitude	longitude	elevation_m	area_sq_km	area_rural_sq_km	area_urban_sq_km
1.	AD	9407	42.558333	1.555278		470		
2.	AE	307763	24.4	54.3		83600	70575	8568
3.	AF	303427	33.0	66.0		652860		
4.	AF_BAL	1674795	36.7	67.116667	340	16186		
5.	AF_BAM		34.75	67.25	3042	14175		
6.	AF_BDG		35.0	63.75	1589	20591		
7.	AF_BDS		38.0	71.0	3669	44059		
8.	AF_BGL		35.87	68.93	2013	21118		
9.	AF_DAY		33.75	66.25	2200	18088		
10.	AF_FRA	1674802	32.5	63.5	1081	48470		

Selected the fields I want to drop



Export the file into a csv

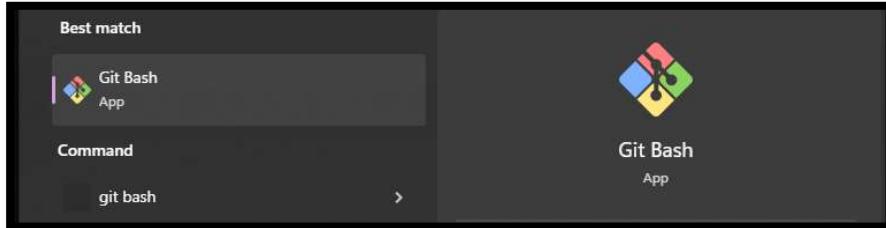


Do this with all the csv files

Step 2: Load Data in Oracle Linux Server

This step transfers the cleaned datasets from the local machine to the Oracle Linux Server environment. Uploading the files to the cluster allows the data to be processed using distributed computing tools such as Hadoop and Hive. Secure Shell (SSH) is used to establish a remote connection to the server and manage files directly within the big data environment.

Open **Git Bash** terminal on your local computer



In the shell terminal, use ssh to connect to the Oracle cluster

```
ssh username@ip
```

Download the following file “covid19_tables.zip” into oracle big data server

```
 wget https://github.com/chica-94/COVID19-Data-Analysis/raw/refs/heads/main/covid19\_tables.zip
ls -lha covid19_tables.zip
```

```
-bash-4.2$ wget https://github.com/chica-94/COVID19-Data-Analysis/raw/refs/heads/main/covid19_tables.zip
--2025-12-03 04:36:33-- https://github.com/chica-94/COVID19-Data-Analysis/raw/refs/heads/main/covid19_tables.zip
Resolving github.com (github.com)... 140.82.113.4
Connecting to github.com (github.com)|140.82.113.4|:443... connected.
HTTP request sent, awaiting response... 302 Found
Location: https://media.githubusercontent.com/media/chica-94/COVID19-Data-Analysis/refs/heads/main/covid19_tables.zip [following]
--2025-12-03 04:36:33-- https://media.githubusercontent.com/media/chica-94/COVID19-Data-Analysis/refs/heads/main/covid19_tables.zip
Resolving media.githubusercontent.com (media.githubusercontent.com)... 185.199.110.133, 185.199.111.133, 185.199.108.133, ...
Connecting to media.githubusercontent.com (media.githubusercontent.com)|185.199.110.133|:443... connected.
HTTP request sent, awaiting response... 200 OK
Length: 140991586 (134M) [application/zip]
Saving to: 'covid19_tables.zip'

100%[=====] 140,991,586 47.6MB/s  in 2.8s

2025-12-03 04:36:36 (47.6 MB/s) - 'covid19_tables.zip' saved [140991586/140991586]

-bash-4.2$ ls -lha covid19_tables.zip
-rw-rw-r-- 1 bcnica bcnica 135M Dec  3 04:36 covid19_tables.zip
-bash-4.2$
```

Unzip tables folder

```
unzip covid19_tables.zip
ls -lha covid19_tables
```

```

-bash-4.2$ 
-bash-4.2$ unzip covid19_tables.zip
Archive: covid19_tables.zip
  creating: covid19_tables/
  inflating: covid19_tables/age.csv
  inflating: covid19_tables/demographics.csv
  inflating: covid19_tables/epidemiology.csv
  inflating: covid19_tables/gender.csv
  inflating: covid19_tables/geography.csv
  inflating: covid19_tables/health.csv
  inflating: covid19_tables/hospitalizations.csv
  inflating: covid19_tables/index.csv
  inflating: covid19_tables/vaccinations.csv
-bash-4.2$ 
-bash-4.2$ ls -lha covid19_tables
total 942M
drwxrwxrwx  2 bchica bchica 4.0K Dec  3 04:41 .
drwx----- 10 bchica bchica 4.0K Dec  3 04:50 ..
-rw-rw-rw-  1 bchica bchica 212M Dec  3 00:14 age.csv
-rw-rw-rw-  1 bchica bchica 1.4M Dec  2 18:52 demographics.csv
-rw-rw-rw-  1 bchica bchica 457M Dec  2 20:17 epidemiology.csv
-rw-rw-rw-  1 bchica bchica 142M Dec  2 19:53 gender.csv
-rw-rw-rw-  1 bchica bchica 662K Dec  2 19:05 geography.csv
-rw-rw-rw-  1 bchica bchica 91K Dec  2 18:45 health.csv
-rw-rw-rw-  1 bchica bchica 55M Dec  2 18:41 hospitalizations.csv
-rw-rw-rw-  1 bchica bchica 1.1M Dec  2 18:42 index.csv
-rw-rw-rw-  1 bchica bchica 75M Dec  2 19:00 vaccinations.csv
-bash-4.2$ 

```

Step 3: Upload datasets to HDFS

Hadoop Distributed File System (HDFS) is used to store large datasets across multiple nodes for efficient access and fault tolerance. In this step, directories are created for each dataset, and the CSV files are uploaded into HDFS. Organizing the data into separate directories ensures structured storage and prepares the datasets for table creation in Hive.

Create a directory rating to put the files to HDFS. Then upload the covid data to the new directory in HDFS. Once the file is added, verify *all* files are uploaded to **covid19** directory (table):

```

cd covid19_tables

hdfs dfs -mkdir tmp/group3_covid19/index
hdfs dfs -mkdir tmp/group3_covid19/hospitalizations
hdfs dfs -mkdir tmp/group3_covid19/health
hdfs dfs -mkdir tmp/group3_covid19/vaccinations
hdfs dfs -mkdir tmp/group3_covid19/demographics
hdfs dfs -mkdir tmp/group3_covid19/geography
hdfs dfs -mkdir tmp/group3_covid19/gender
hdfs dfs -mkdir tmp/group3_covid19/epidemiology
hdfs dfs -mkdir tmp/group3_covid19/age
hdfs dfs -mkdir tmp/group3_covid19/gender
hdfs dfs -mkdir /tmp/group3_covid19/aggregated
hdfs dfs -mkdir /tmp/group3_covid19/top10_countries/

hdfs dfs -ls tmp/group3_covid19/

```

```

-bash-4.2$ cd covid19_tables/
-bash-4.2$ hdfs dfs -mkdir /tmp/group3_covid19/
-bash-4.2$ hdfs dfs -mkdir /tmp/group3_covid19/index
-bash-4.2$ hdfs dfs -mkdir /tmp/group3_covid19/hospitalizations
-bash-4.2$ hdfs dfs -mkdir /tmp/group3_covid19/health
-bash-4.2$ hdfs dfs -mkdir /tmp/group3_covid19/vaccinations
-bash-4.2$ hdfs dfs -mkdir /tmp/group3_covid19/demographics
-bash-4.2$ hdfs dfs -mkdir /tmp/group3_covid19/geography
-bash-4.2$ hdfs dfs -mkdir /tmp/group3_covid19/gender
-bash-4.2$ hdfs dfs -mkdir /tmp/group3_covid19/epidemiology
-bash-4.2$ hdfs dfs -mkdir /tmp/group3_covid19/age
-bash-4.2$ hdfs dfs -mkdir /tmp/group3_covid19/gender
mkdir: `/tmp/group3_covid19/gender': File exists
-bash-4.2$ hdfs dfs -ls /tmp/group3_covid19/
Found 9 items
drwxr-xr-x - bchica hdfs 0 2025-12-03 04:56 /tmp/group3_covid19/age
drwxr-xr-x - bchica hdfs 0 2025-12-03 04:56 /tmp/group3_covid19/demographics
drwxr-xr-x - bchica hdfs 0 2025-12-03 04:56 /tmp/group3_covid19/epidemiology
drwxr-xr-x - bchica hdfs 0 2025-12-03 04:56 /tmp/group3_covid19/gender
drwxr-xr-x - bchica hdfs 0 2025-12-03 04:56 /tmp/group3_covid19/geography
drwxr-xr-x - bchica hdfs 0 2025-12-03 04:56 /tmp/group3_covid19/health
drwxr-xr-x - bchica hdfs 0 2025-12-03 04:56 /tmp/group3_covid19/hospitalizations
drwxr-xr-x - bchica hdfs 0 2025-12-03 04:56 /tmp/group3_covid19/index
drwxr-xr-x - bchica hdfs 0 2025-12-03 04:56 /tmp/group3_covid19/vaccinations
-bash-4.2$ 

```

Upload the covid19 datasets into their respective directories in HDFS.

```

hdfs dfs -put index.csv tmp/group3_covid19/index
hdfs dfs -put hospitalizations.csv tmp/group3_covid19/hospitalizations
hdfs dfs -put health.csv tmp/group3_covid19/health
hdfs dfs -put vaccinations.csv tmp/group3_covid19/vaccinations
hdfs dfs -put demographics.csv tmp/group3_covid19/demographics
hdfs dfs -put geography.csv tmp/group3_covid19/geography
hdfs dfs -put gender.csv tmp/group3_covid19/gender
hdfs dfs -put epidemiology.csv tmp/group3_covid19/epidemiology
hdfs dfs -put age.csv tmp/group3_covid19/age

```

Ensure that all files are loaded to hdfs and are in their designated directories

```

hdfs dfs -ls -R tmp/group3_covid19/

```

```

-bash-4.2$ 
-bash-4.2$ hdfs dfs -put index.csv /tmp/group3_covid19/index
-bash-4.2$ hdfs dfs -put hospitalizations.csv /tmp/group3_covid19/hospitalizations
-bash-4.2$ hdfs dfs -put health.csv /tmp/group3_covid19/health
-bash-4.2$ hdfs dfs -put vaccinations.csv /tmp/group3_covid19/vaccinations
-bash-4.2$ hdfs dfs -put demographics.csv /tmp/group3_covid19/demographics
-bash-4.2$ hdfs dfs -put geography.csv /tmp/group3_covid19/geography
-bash-4.2$ hdfs dfs -put gender.csv /tmp/group3_covid19/gender
-bash-4.2$ hdfs dfs -put epidemiology.csv /tmp/group3_covid19/epidemiology
-bash-4.2$ hdfs dfs -put age.csv /tmp/group3_covid19/age
-bash-4.2$ hdfs dfs -ls -R /tmp/group3_covid19/
ls: `/tmp/group3_covid19/': No such file or directory
-bash-4.2$ hdfs dfs -ls -R /tmp/group3_covid19/
lrwxr-xr-x - bchica mors 0 2025-12-03 04:56 /tmp/group3_covid19/age
-rw-r--r-- 3 bchica hdfs 221791739 2025-12-03 04:59 /tmp/group3_covid19/age/age.csv
lrwxr-xr-x - bchica hdfs 0 2025-12-03 04:59 /tmp/group3_covid19/demographics
-rw-r--r-- 3 bchica hdfs 1375489 2025-12-03 04:59 /tmp/group3_covid19/demographics/demographics.csv
lrwxr-xr-x - bchica hdfs 0 2025-12-03 04:59 /tmp/group3_covid19/geography
-rw-r--r-- 3 bchica hdfs 478654081 2025-12-03 04:59 /tmp/group3_covid19/geography/geography.csv
lrwxr-xr-x - bchica hdfs 0 2025-12-03 04:59 /tmp/group3_covid19/gender
-rw-r--r-- 3 bchica hdfs 148284695 2025-12-03 04:59 /tmp/group3_covid19/gender/gender.csv
lrwxr-xr-x - bchica hdfs 0 2025-12-03 04:59 /tmp/group3_covid19/health
-rw-r--r-- 3 bchica hdfs 676897 2025-12-03 04:59 /tmp/group3_covid19/health/health.csv
lrwxr-xr-x - bchica hdfs 0 2025-12-03 04:59 /tmp/group3_covid19/hospitalizations
-rw-r--r-- 3 bchica hdfs 56630483 2025-12-03 04:58 /tmp/group3_covid19/hospitalizations/hospitalizations.csv
lrwxr-xr-x - bchica hdfs 0 2025-12-03 04:58 /tmp/group3_covid19/index
-rw-r--r-- 3 bchica hdfs 1114818 2025-12-03 04:58 /tmp/group3_covid19/index/index.csv
lrwxr-xr-x - bchica hdfs 0 2025-12-03 04:59 /tmp/group3_covid19/vaccinations
-rw-r--r-- 3 bchica hdfs 78235179 2025-12-03 04:59 /tmp/group3_covid19/vaccinations/vaccinations.csv
-bash-4.2$ 

```

Step 4: Create Tables Using Beeline

The purpose of this step is to create external Hive tables that map directly to the datasets stored in HDFS. Using Beeline and HiveQL commands, each dataset is structured into a table with defined schemas. External tables are used so that the underlying data remains in HDFS while still allowing SQL-based querying and analysis.

Now that the data is loaded in HDFS, we can now use Hive to create a table and query the data using hiveql commands

Open the Hive shell environment using **beeline**.

```
beeline
```

Use your database to create tables

```
use username;
```

Paste the following HiveQL code to create an external table called which will contain all the data from each dataset uploaded to HDFS. We start by applying a “DROP” command to remove any existing table that may come into conflict

Create **covid19_index** table

```
DROP TABLE IF EXISTS covid19_index;

CREATE EXTERNAL TABLE IF NOT EXISTS covid19_index(
    location_key STRING,
    country_code STRING,
    country_name STRING,
    subregion1_name STRING,
    subregion2_name STRING)
ROW FORMAT DELIMITED FIELDS TERMINATED BY ','
STORED AS TEXTFILE LOCATION '/tmp/group3_covid19/index/'
TBLPROPERTIES ("skip.header.line.count"="1");
```

Create **covid19_hospitalizations** table

```
DROP TABLE IF EXISTS covid19_hospitalizations;

CREATE EXTERNAL TABLE IF NOT EXISTS covid19_hospitalizations(
    event_date DATE,
    location_key STRING,
    new_hospitalized_patients INT,
    cumulative_hospitalized_patients INT,
    new_intensive_care_patients INT,
    cumulative_intensive_care_patients INT)
ROW FORMAT DELIMITED FIELDS TERMINATED BY ',' ROW FORMAT DELIMITED
FIELDS TERMINATED BY ','
STORED AS TEXTFILE LOCATION '/tmp/group3_covid19/hospitalizations/'
TBLPROPERTIES ("skip.header.line.count"="1");
```

Create covid19_health table

```
DROP TABLE IF EXISTS covid19_health;

CREATE EXTERNAL TABLE IF NOT EXISTS covid19_health(
    location_key STRING,
    adult_male_mortality_rate DOUBLE,
    adult_female_mortality_rate DOUBLE,
    life_expectancy DOUBLE,
    diabetes_prevalence DOUBLE,
    health_expenditure_usd DOUBLE,
    out_of_pocket_health_expenditure_usd DOUBLE)
ROW FORMAT DELIMITED FIELDS TERMINATED BY ','
STORED AS TEXTFILE LOCATION '/tmp/group3_covid19/health/'
TBLPROPERTIES ("skip.header.line.count""1");
```

Create covid19_vaccinations table

```
DROP TABLE IF EXISTS covid19_vaccinations;

CREATE EXTERNAL TABLE IF NOT EXISTS covid19_vaccinations(
    event_date DATE,
    location_key STRING,
    new_persons_fully_vaccinated INT,
    cumulative_persons_fully_vaccinated INT)
ROW FORMAT DELIMITED FIELDS TERMINATED BY ','
STORED AS TEXTFILE LOCATION '/tmp/group3_covid19/vaccinations/'
TBLPROPERTIES ("skip.header.line.count""1");
```

Create covid19_geography table

```
DROP TABLE IF EXISTS covid19_geography;

CREATE EXTERNAL TABLE IF NOT EXISTS covid19_geography(
    location_key STRING,
    latitude DOUBLE,
    longitude DOUBLE)
ROW FORMAT DELIMITED FIELDS TERMINATED BY ','
STORED AS TEXTFILE LOCATION '/tmp/group3_covid19/geography/'
TBLPROPERTIES ("skip.header.line.count""1");
```

Create **covid19_demographics** table

```
DROP TABLE IF EXISTS covid19_demographics;

CREATE EXTERNAL TABLE IF NOT EXISTS covid19_demographics(
    location_key STRING,
    population INT,
    population_male INT,
    population_female INT,
    population_age_00_09 INT,
    population_age_10_19 INT,
    population_age_20_29 INT,
    population_age_30_39 INT,
    population_age_40_49 INT,
    population_age_50_59 INT,
    population_age_60_69 INT,
    population_age_70_79 INT,
    population_age_80_and_older INT)
ROW FORMAT DELIMITED FIELDS TERMINATED BY ','
STORED AS TEXTFILE LOCATION '/tmp/group3_covid19/demographics/'
TBLPROPERTIES ("skip.header.line.count""1");
```

Create **covid19_gender** table

```
DROP TABLE IF EXISTS covid19_gender;

CREATE EXTERNAL TABLE IF NOT EXISTS covid19_gender(
    event_date DATE,
    location_key STRING,
    new_deceased_male INT,
    cumulative_deceased_male INT,
    new_deceased_female INT,
    cumulative_deceased_female INT,
    new_recovered_male INT,
    cumulative_recovered_male INT,
    new_recovered_female INT,
    cumulative_recovered_female INT)
ROW FORMAT DELIMITED FIELDS TERMINATED BY ','
STORED AS TEXTFILE LOCATION '/tmp/group3_covid19/gender/'
TBLPROPERTIES ("skip.header.line.count""1");
```

Create **covid19_epidemiology** table

```
DROP TABLE IF EXISTS covid19_epidemiology;

CREATE EXTERNAL TABLE IF NOT EXISTS covid19_epidemiology(
    event_date DATE,
    location_key STRING,
    new_confirmed INT,
    cumulative_confirmed INT,
    new_deceased INT,
    cumulative_deceased INT,
    new_recovered INT,
    cumulative_recovered INT)
ROW FORMAT DELIMITED FIELDS TERMINATED BY ','
STORED AS TEXTFILE LOCATION '/tmp/group3_covid19/epidemiology/'
TBLPROPERTIES ("skip.header.line.count""1");
```

Create covid19_age table

```
DROP TABLE IF EXISTS covid19_age;

CREATE EXTERNAL TABLE IF NOT EXISTS covid19_age(
    event_date DATE,
    location_key STRING,
    new_recovered_age_0-9 INT,
    new_recovered_age_10-19 INT,
    new_recovered_age_20-29 INT,
    new_recovered_age_30-39 INT,
    new_recovered_age_40-49 INT,
    new_recovered_age_50-59 INT,
    new_recovered_age_60-69 INT,
    new_recovered_age_70-79 INT,
    new_recovered_age_80-89 INT,
    new_recovered_age_90-99 INT,
    cumulative_recovered_age_0-9 INT,
    cumulative_recovered_age_10-19 INT,
    cumulative_recovered_age_20-29 INT,
    cumulative_recovered_age_30-39 INT,
    cumulative_recovered_age_40-49 INT,
    cumulative_recovered_age_50-59 INT,
    cumulative_recovered_age_60-69 INT,
    cumulative_recovered_age_70-79 INT,
    cumulative_recovered_age_80-89 INT,
    cumulative_recovered_age_90-99 INT )
ROW FORMAT DELIMITED FIELDS TERMINATED BY ','
STORED AS TEXTFILE LOCATION '/tmp/group3 covid19/age/'
```

Then, in the Hive shell, make sure that the tables **covid19** are created and is shown:

```
: jdbc:hive2://bigdatuno.sub03291929060.tra!> show tables;
INFO : Compiling command(queryId=hive_20251203061324_7d68ac0c-b);
INFO : Concurrency mode is disabled, not creating a lock manager
INFO : Semantic Analysis Completed (retrial = false)
INFO : Returning Hive schema: Schema(fieldschemas:[FieldSchema()]
INFO : Completed compiling command(queryId=hive_20251203061324_7d68ac0c-b);
INFO : Concurrency mode is disabled, not creating a lock manager
INFO : Executing command(queryId=hive_20251203061324_7d68ac0c-b);
INFO : Starting task [Stage-0:DDL] in serial mode
INFO : Completed executing command(queryId=hive_20251203061324_7d68ac0c-b);
INFO : OK
INFO : Concurrency mode is disabled, not creating a lock manager
+-----+
| tab_name |
+-----+
| covid19_age |
| covid19_demographics |
| covid19_epidemiology |
| covid19_gender |
| covid19_geography |
| covid19_health |
| covid19_hospitalizations |
| covid19_index |
| covid19_vaccinations |
+-----+
```

Step 4: Create external aggregate table

An aggregated table is created to combine multiple COVID-19 datasets into a single unified structure. This table enables comprehensive analysis by integrating epidemiology, hospitalizations, vaccinations, health indicators, demographics, and geographic data. Aggregating the data simplifies querying and improves analytical efficiency.

Paste the following HiveQL code to create a table called “**covid19_aggregated**,” to store data from the **covid19** tables.

```
DROP TABLE IF EXISTS covid19_aggregated;

CREATE EXTERNAL TABLE IF NOT EXISTS covid19_aggregated(
    event_date DATE, location_key STRING,country_code STRING,country_name
    STRING, subregion1_name STRING,subregion2_name STRING, new_confirmed INT,
    cumulative_confirmed INT,new_deceased INT,cumulative_deceased
    INT,new_recovered INT,cumulative_recovered INT,new_hospitalized_patients
    INT,cumulative_hospitalized_patients INT,new_intensive_care_patients
    INT,cumulative_intensive_care_patients INT,new_persons_fully_vaccinated
    INT,cumulative_persons_fully_vaccinated INT,adult_male_mortality_rate
    DOUBLE,adult_female_mortality_rate DOUBLE,life_expectancy
    DOUBLE,diabetes_prevalence DOUBLE,health_expenditure_usd
    DOUBLE,out_of_pocket_health_expenditure_usd DOUBLE,population
    INT,population_male INT,population_female INT,population_age_00_09
    INT,population_age_10_19 INT,population_age_20_29 INT,population_age_30_39
    INT,population_age_40_49 INT,population_age_50_59 INT,population_age_60_69
    INT,population_age_70_79 INT,population_age_80_and_older INT,latitude
    DOUBLE,longitude DOUBLE,new_deceased_male INT,cumulative_deceased_male
    INT,new_deceased_female INT,cumulative_deceased_female
    INT,new_recovered_male INT,cumulative_recovered_male
    INT,new_recovered_female INT,cumulative_recovered_female
    INT,new_recovered_age_0-9 INT,new_recovered_age_10-19
    INT,new_recovered_age_20-29 INT,new_recovered_age_30-39
    INT,new_recovered_age_40-49 INT,new_recovered_age_50-59
    INT,new_recovered_age_60-69 INT,new_recovered_age_70-79
    INT,new_recovered_age_80-89 INT,new_recovered_age_90-99
    INT,cumulative_recovered_age_0-9 INT,cumulative_recovered_age_10-19
    INT,cumulative_recovered_age_20-29 INT,cumulative_recovered_age_30-39
    INT,cumulative_recovered_age_40-49 INT,cumulative_recovered_age_50-59
    INT,cumulative_recovered_age_60-69 INT,cumulative_recovered_age_70-79
    INT,cumulative_recovered_age_80-89 INT,cumulative_recovered_age_90-99 INT)
ROW FORMAT DELIMITED FIELDS TERMINATED BY ','
STORED AS TEXTFILE LOCATION '/tmp/group3_covid19/aggregated';
```

Now you need to populate data from the **covid19 tables** to **covid19_aggregated** using **SELECT statement** as follows:

```
INSERT OVERWRITE TABLE covid19_aggregated
SELECT
    epi.event_date,epi.location_key, idx.country_code, idx.country_name, idx.subregion1_name,
    idx.subregion2_name, epi.new_confirmed, epi.cumulative_confirmed, epi.new_deceased,
    epi.cumulative_deceased, epi.new_recovered, epi.cumulative_recovered, hosp.new_hospitalized_patients, hosp.cumulative_hospitalized_patients, hosp.new_intensive_care_patients,
    hosp.cumulative_intensive_care_patients, vac.new_persons_fully_vaccinated, vac.cumulative_persons_fully_vaccinated, health.adult_male_mortality_rate, health.adult_female_mortality_rate, health.life_expectancy, health.diabetes_prevalence, health.health_expenditure_usd, health.out_of_pocket_health_expenditure_usd, demo.population, demo.population_male, demo.population_female, demo.population_age_00_09, demo.population_age_10_19, demo.population_age_20_29, demo.population_age_30_39, demo.population_age_40_49, demo.population_age_50_59, demo.population_age_60_69, demo.population_age_70_79, demo.population_age_80_and_older, geo.latitude, geo.longitude, gender.new_deceased_male, gender.cumulative_deceased_male, gender.new_deceased_female, gender.cumulative_deceased_female, gender.new_recovered_male, gender.cumulative_recovered_male, gender.new_recovered_female, gender.cumulative_recovered_female, age.new_recovered_age_0_9, age.new_recovered_age_10_19, age.new_recovered_age_20_29, age.new_recovered_age_30_39, age.new_recovered_age_40_49, age.new_recovered_age_50_59, age.new_recovered_age_60_69, age.new_recovered_age_70_79, age.new_recovered_age_80_89, age.new_recovered_age_90_99, age.cumulative_recovered_age_0_9, age.cumulative_recovered_age_10_19, age.cumulative_recovered_age_20_29, age.cumulative_recovered_age_30_39, age.cumulative_recovered_age_40_49, age.cumulative_recovered_age_50_59, age.cumulative_recovered_age_60_69, age.cumulative_recovered_age_70_79, age.cumulative_recovered_age_80_89, age.cumulative_recovered_age_90_99
    FROM covid19_epidemiology epi
    LEFT JOIN covid19_index idx ON epi.location_key = idx.location_key
    LEFT JOIN covid19_hospitalizations hosp ON epi.location_key = hosp.location_key AND epi.event_date = hosp.event_date
    LEFT JOIN covid19_vaccinations vac ON epi.location_key = vac.location_key AND epi.event_date = vac.event_date
    LEFT JOIN covid19_health health ON epi.location_key = health.location_key
    LEFT JOIN covid19_demographics demo ON epi.location_key = demo.location_key
    LEFT JOIN covid19_geography geo ON epi.location_key = geo.location_key
    LEFT JOIN covid19_gender gender ON epi.location_key = gender.location_key AND epi.event_date = gender.event_date
    LEFT JOIN covid19_age age ON epi.location_key = age.location_key AND epi.event_date = age.event_date;
```

Check if the table **covid19_aggregated** tables are shown

```
show tables;
```

Query the content of the **covid19_aggregated** table to ensure that the table has been created successfully.

```
SELECT * FROM covid19_aggregated LIMIT 3;
```

Step 5: Data Cleaning and Preparation

This step focuses on refining the dataset to support meaningful analysis. By querying the aggregated table, the top ten countries with the highest cumulative confirmed COVID-19 cases are identified. Reducing the dataset to the most relevant records improves performance and allows the analysis to focus on the most impacted regions.

Query the aggregated tables to find the top 10 countries with the most confirmed COVID-19 cases

```
SELECT country_name, MAX(cumulative_confirmed) AS total_cases
FROM covid19_aggregated
WHERE cumulative_confirmed IS NOT NULL
GROUP BY country_name
ORDER BY total_cases DESC
LIMIT 10;
```

country_name	total_cases
United States of America	92440495
India	44516479
Brazil	34568833
France	33766090
Germany	32604993
South Korea	24264470
United Kingdom	23554971
Italy	22114423
Russia	20265004
Japan	19868288

In the Hive shell, use the following code to create an empty table called **covid19_top10_countries**

```
DROP TABLE IF EXISTS covid19_top10_countries;

CREATE EXTERNAL TABLE IF NOT EXISTS covid19_top10_countries (
    location_key STRING, country_code STRING, country_name STRING, subregion1_name
    STRING, subregion2_name STRING, cumulative_confirmed INT, cumulative_deceased INT,
    cumulative_recovered INT, cumulative_hospitalized_patients INT,
    cumulative_intensive_care_patients INT, cumulative_persons_fully_vaccinated INT,
    adult_male_mortality_rate DOUBLE, adult_female_mortality_rate DOUBLE,
    life_expectancy DOUBLE, diabetes_prevalence DOUBLE, health_expenditure_usd DOUBLE,
    out_of_pocket_health_expenditure_usd DOUBLE, population INT, population_male INT,
    population_female INT, population_age_00_09 INT, population_age_10_19 INT,
    population_age_20_29 INT, population_age_30_39 INT, population_age_40_49 INT,
    population_age_50_59 INT, population_age_60_69 INT, population_age_70_79 INT,
    population_age_80_and_older INT, latitude DOUBLE, longitude DOUBLE,
    cumulative_deceased_male INT, cumulative_deceased_female INT,
    cumulative_recovered_male INT, cumulative_recovered_female INT,
    cumulative_recovered_age_0_9 INT, cumulative_recovered_age_10_19 INT,
    cumulative_recovered_age_20_29 INT, cumulative_recovered_age_30_39 INT,
    cumulative_recovered_age_40_49 INT, cumulative_recovered_age_50_59 INT,
    cumulative_recovered_age_60_69 INT, cumulative_recovered_age_70_79 INT,
    cumulative_recovered_age_80_89 INT, cumulative_recovered_age_90_99 INT)
ROW FORMAT DELIMITED
FIELDS TERMINATED BY ','
STORED AS TEXTFILE
LOCATION '/tmp/qgroup3 covid19/top10_countries/';
```

Copy and paste the following Hive code to Hive shell in order to create a table **covid19_top10_countries**. It will be data from the top 10 countries with the most confirmed cases from the **covid19_aggregated** tables

```
WITH top10 AS (
    SELECT country_name, MAX(cumulative_confirmed) AS total_cases
    FROM covid19_aggregated
    WHERE cumulative_confirmed IS NOT NULL
    GROUP BY country_name
    ORDER BY total_cases DESC
    LIMIT 10)
INSERT OVERWRITE TABLE covid19_top10_countries
SELECT
    a.location_key, a.country_code, a.country_name, a.subregion1_name, a.subregion2_name,
    MAX(a.cumulative_confirmed) AS cumulative_confirmed, MAX(a.cumulative_deceased) AS
    cumulative_deceased, MAX(a.cumulative_recovered) AS cumulative_recovered,
    MAX(a.cumulative_hospitalized_patients) AS cumulative_hospitalized_patients,
    MAX(a.cumulative_intensive_care_patients) AS cumulative_intensive_care_patients,
    MAX(a.cumulative_persons_fully_vaccinated) AS cumulative_persons_fully_vaccinated,
    MAX(a.adult_male_mortality_rate), MAX(a.adult_female_mortality_rate),
    MAX(a.life_expectancy), MAX(a.diabetes_prevalence), MAX(a.health_expenditure_usd),
    MAX(a.out_of_pocket_health_expenditure_usd), MAX(a.population),
    MAX(a.population_male), MAX(a.population_female), MAX(a.population_age_00_09),
    MAX(a.population_age_10_19), MAX(a.population_age_20_29),
    MAX(a.population_age_30_39), MAX(a.population_age_40_49),
    MAX(a.population_age_50_59), MAX(a.population_age_60_69),
    MAX(a.population_age_70_79), MAX(a.population_age_80_and_older), a.latitude,
    a.longitude, MAX(a.cumulative_deceased_male), MAX(a.cumulative_deceased_female),
    MAX(a.cumulative_recovered_male), MAX(a.cumulative_recovered_female),
    MAX(a.cumulative_recovered_age_0_9), MAX(a.cumulative_recovered_age_10_19),
    MAX(a.cumulative_recovered_age_20_29), MAX(a.cumulative_recovered_age_30_39),
    MAX(a.cumulative_recovered_age_40_49), MAX(a.cumulative_recovered_age_50_59),
    MAX(a.cumulative_recovered_age_60_69), MAX(a.cumulative_recovered_age_70_79),
    MAX(a.cumulative_recovered_age_80_89), MAX(a.cumulative_recovered_age_90_99)
FROM covid19_aggregated a
JOIN top10 t ON a.country_name = t.country_name
WHERE a.subregion1_name IS NOT NULL
GROUP BY a.location_key,a.country_code,a.country_name,a.subregion1_name,a.subregion2_name;
```

Query from the table and to see if it has the correct data and values:

```
SELECT * FROM covid19_top10_countries LIMIT 10;
```

In the Beeline shell, you need to check if the table **covid19_top10_countries** shows your database and hdfs directory

```
describe formatted covid19_top10_countries;
```

Then, in the hive shell, you need to check if the tables **covid19_aggregated** and **covid19_top10_countries** are shown:

```
show tables;
```

tab_name
covid19_age
covid19_aggregated
covid19_demographics
covid19_epidemiology
covid19_gender
covid19_geography
covid19_health
covid19_hospitalizations
covid19_index
covid19_top10_countries
covid19_vaccinations

Compare the number of records between the tables **covid19_aggregated** and **covid19_top10_countries**. You will see the table was reduced from **12,525,825** to **10,660**.

```
SELECT COUNT(*) FROM covid19_aggregated;  
SELECT COUNT(*) FROM covid19_top10_countries;
```

covid19_records
12525825

covid19_top10_records
10660

Step 6: Export Cleaned Table

After identifying the top ten countries, the cleaned dataset is exported from HDFS to a single CSV file. The (-getmerge) command is used to combine multiple output files into one consolidated file, which is then downloaded to the local machine. This exported dataset is used for visualization and further analysis.

After the **covid19_top10_countries** table is created, listing the contents inside the **/tmp/group3_covid19/top10_countries/** will show 49 files

```
hdfs dfs -ls /tmp/group3_covid19/top10_countries/
```

-bash-4.2\$ pwd
/home/bchica
-bash-4.2\$ cd covid19_tables/
-bash-4.2\$ hdfs dfs -ls /tmp/group3_covid19/top10_countries/
found 49 items
-rw-r--r-- 3 bchica hdfs 41387 2025-12-03 21:53 /tmp/group3_covid19/top10_countries/000000_0
-rw-r--r-- 3 bchica hdfs 42593 2025-12-03 21:53 /tmp/group3_covid19/top10_countries/000001_0
-rw-r--r-- 3 bchica hdfs 41729 2025-12-03 21:53 /tmp/group3_covid19/top10_countries/000002_0
-rw-r--r-- 3 bchica hdfs 43412 2025-12-03 21:53 /tmp/group3_covid19/top10_countries/000003_0
-rw-r--r-- 3 bchica hdfs 47807 2025-12-03 21:53 /tmp/group3_covid19/top10_countries/000004_0
-rw-r--r-- 3 bchica hdfs 45334 2025-12-03 21:53 /tmp/group3_covid19/top10_countries/000005_0
-rw-r--r-- 3 bchica hdfs 44734 2025-12-03 21:53 /tmp/group3_covid19/top10_countries/000006_0
-rw-r--r-- 3 bchica hdfs 43000 2025-12-03 21:53 /tmp/group3_covid19/top10_countries/000007_0
-rw-r--r-- 3 bchica hdfs 44016 2025-12-03 21:53 /tmp/group3_covid19/top10_countries/000008_0
-rw-r--r-- 3 bchica hdfs 43231 2025-12-03 21:53 /tmp/group3_covid19/top10_countries/000009_0
-rw-r--r-- 3 bchica hdfs 35980 2025-12-03 21:53 /tmp/group3_covid19/top10_countries/000010_0
-rw-r--r-- 3 bchica hdfs 49569 2025-12-03 21:53 /tmp/group3_covid19/top10_countries/000011_0
-rw-r--r-- 3 bchica hdfs 48585 2025-12-03 21:53 /tmp/group3_covid19/top10_countries/000012_0
-rw-r--r-- 3 bchica hdfs 47181 2025-12-03 21:53 /tmp/group3_covid19/top10_countries/000013_0
-rw-r--r-- 3 bchica hdfs 41618 2025-12-03 21:53 /tmp/group3_covid19/top10_countries/000014_0

Merge the 49 output files to a file named **covid19_top10_countries.csv** using the following:

```
hdfs dfs -getmerge /tmp/group3_covid19/top10_countries/0000*  
covid19_top10_countries.csv
```

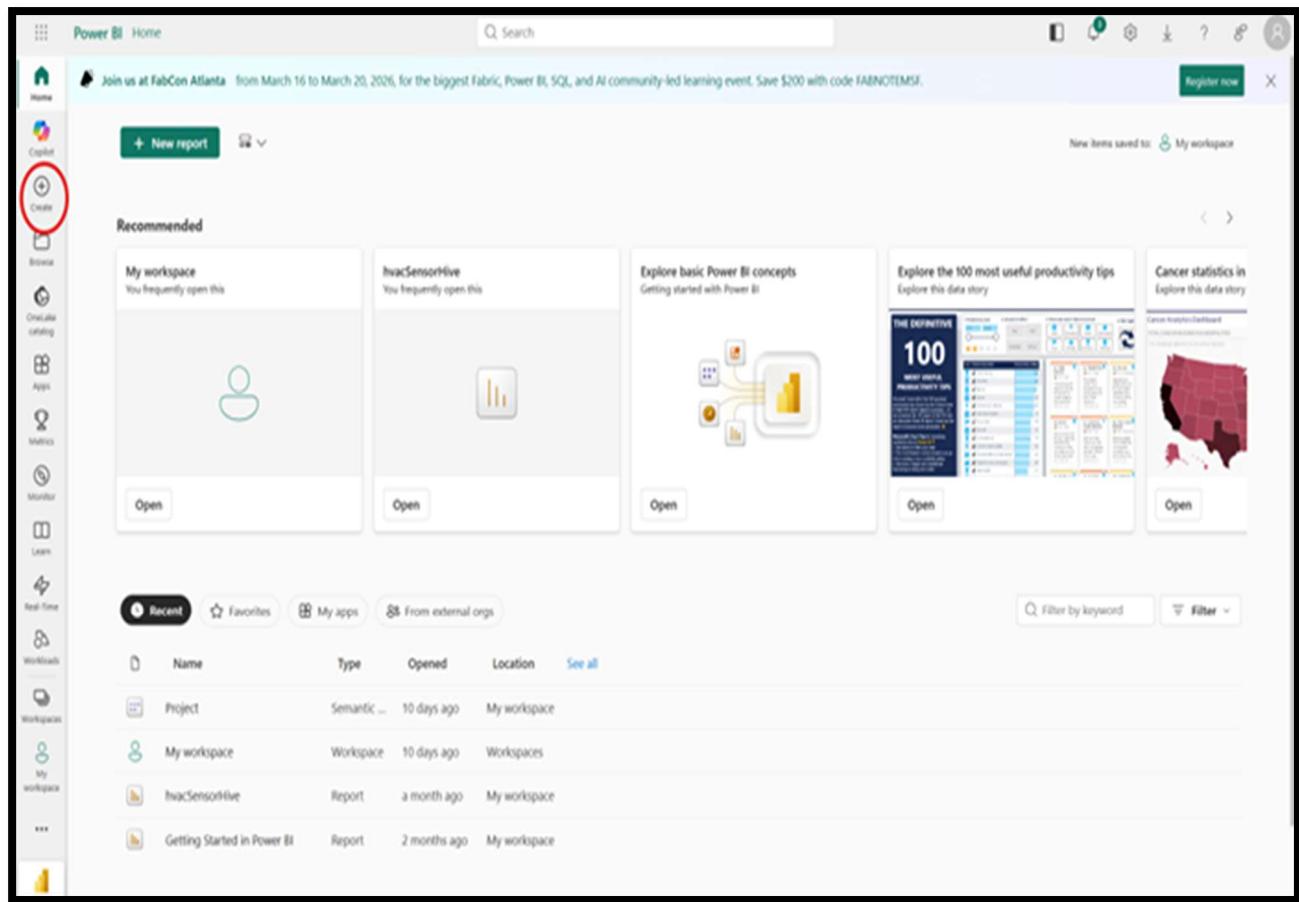
Logout of the cluster and download the csv file into your local machine

```
scp username@ip:/home/bchica/covid19_tables/covid19_top10_countries.csv  
~/Downloads/
```

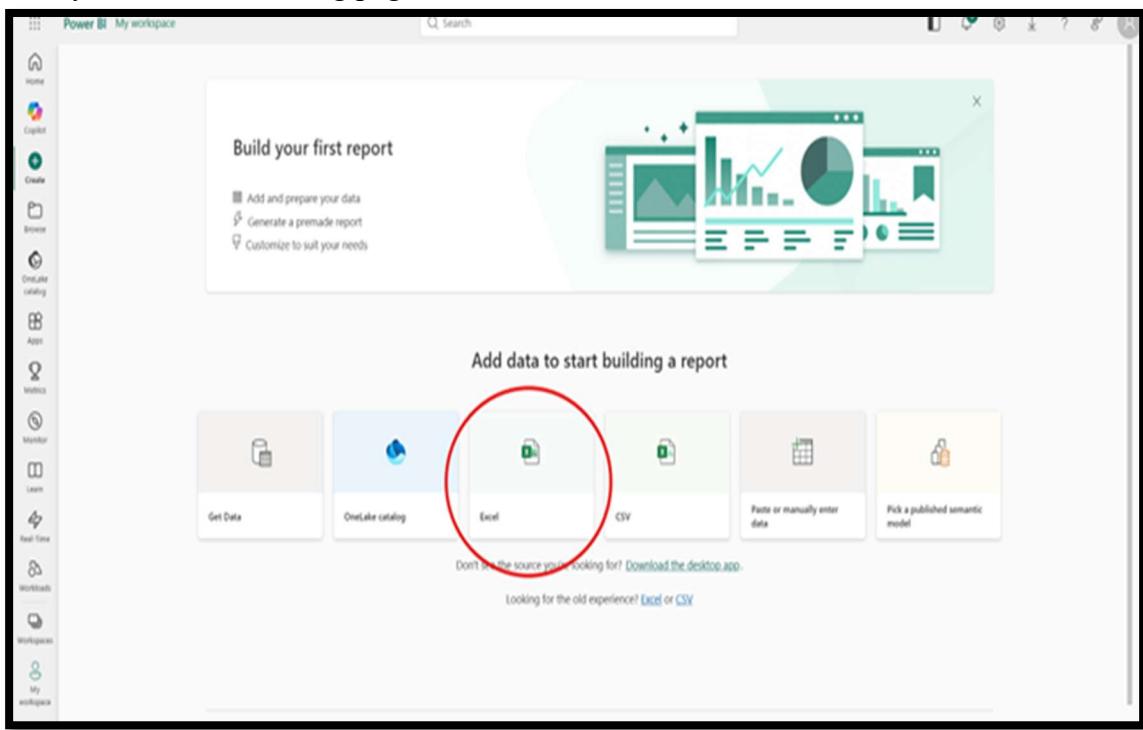
Step 7: Visualization using PowerBI

Power BI is used to visualize the results of the aggregated COVID-19 analysis. A bubble map is created to display the top ten countries by cumulative confirmed COVID-19 cases, where bubble size represents the relative magnitude of total confirmed cases. This visualization provides a clear geographic comparison of COVID-19 impact among the most affected countries and supports the findings derived from the Hive-based analysis.

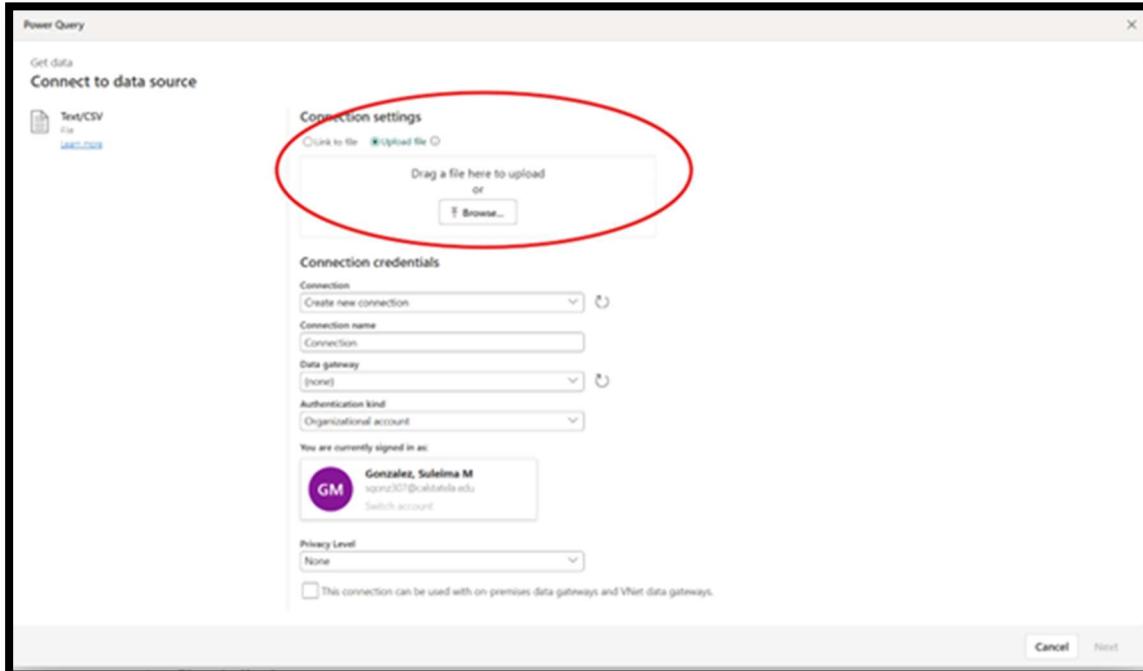
Open your MS Power BI Desktop at your local computer. Select **Create**

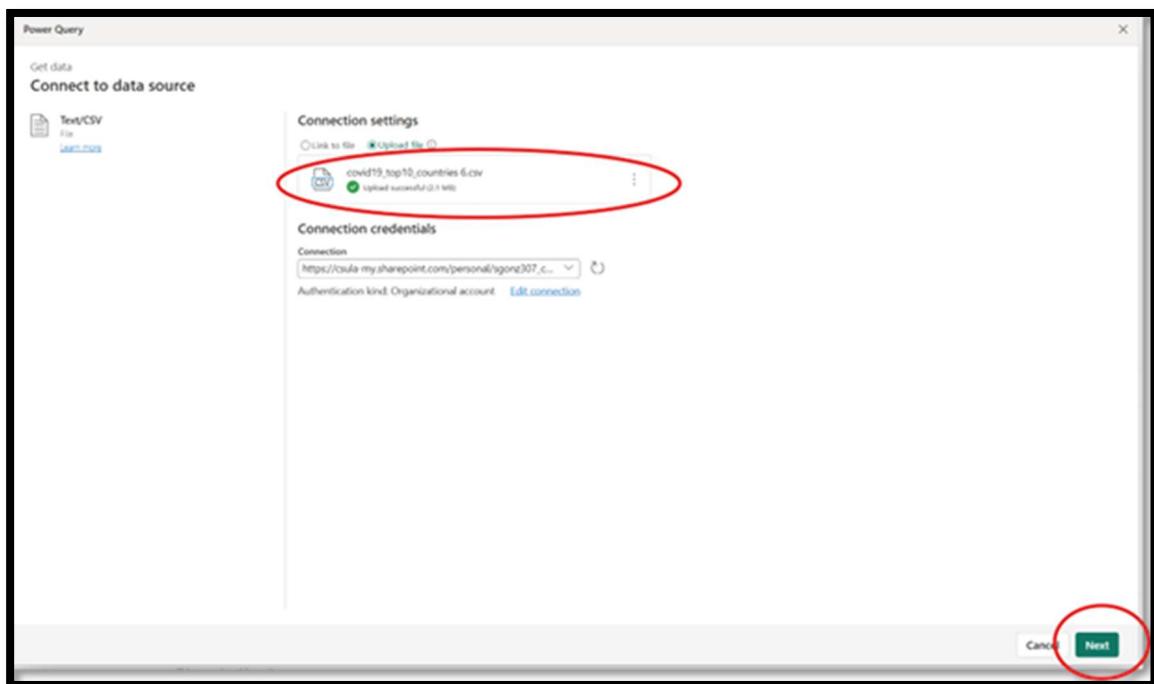


Once you see the following page, select **csv**



Select the **Browse** button. Choose the file you want to use on the File Explorer, then hit **Next**





You will see the following page in PowerBI to preview your dataset

File origin	Delimiter	Data type detection						
65001: Unicode (UTF-8)	Comma	Based on first 200 rows						
# location_key # country_code # country_name # subregion1_name # subregion2_name # cumulative_confirmed # cumulative_deceased # cumulative_recovered # cumulative_hospitalized_patients # cumulative_intensive_care_patients # cumulative								
BR_AL_270530 BR	Brazil	Alegrete	Minas do Negrito	447 10	0	17	11	1N
BR_AL_270530 BR	Brazil	Alegrete	Nova Leao	277 14	0	13	5	1N
BR_AL_MC2 BR	Brazil	Alegrete		129062 2162	0	5864	1602	1N
BR_AP BR	Brazil	Amapá		178204 2159	0	3528	1207	501812
BR_BA_290330 BR	Brazil	Bahia	Belo Campo	1764 22	0	26	11	1N
BR_BA_290330 BR	Brazil	Bahia	Caxolinha	1337 2	0	41	2	1N
BR_BA_291230 BR	Brazil	Bahia	Ibirataia	2607 50	0	36	15	1N
BR_BA_291430 BR	Brazil	Bahia	Imarata	636 17	0	16	7	1N
BR_BA_291530 BR	Brazil	Bahia	Itambe	2731 25	0	45	13	1N
BR_BA_291830 BR	Brazil	Bahia	Kapicuri	2055 27	0	50	25	1N
BR_CE_230130 BR	Brazil	Ceará	Baixo	922 18	0	15	11	1N
BR_CE_230730 BR	Brazil	Ceará	Madalena	1777 39	0	59	31	1N
BR_COFOR BR	Brazil	Ceará		291527 11491	0	19238	4959	1N
BR_ES_320630 BR	Brazil	Espírito Santo	Alfredo Chaves	5136 47	0	13	10	1N
BR_ES_320930 BR	Brazil	Espírito Santo	Iara	6687 96	0	31	17	1N
BR_ES_320332 BR	Brazil	Espírito Santo	Maratábas	73609 247	0	66	34	1N
BR_ES_320430 BR	Brazil	Espírito Santo	Ponto Belo	2126 17	0	5	2	1N
BR_ES_320460 BR	Brazil	Espírito Santo	São Mateus	34454 258	0	87	38	1N
BR_GO_520330 BR	Brazil	Goiás	Bom Jesus de Goiás	5950 125	0	236	71	1N
BR_GO_520430 BR	Brazil	Goiás	Cage	3003 46	0	63	27	1N
BR_GO_520630 BR	Brazil	Goiás	Coronel	809 12	0	24	12	1N
BR_GO_520830 BR	Brazil	Goiás	Fornos	722 20	0	36	16	1N
BR_GO_520930 BR	Brazil	Goiás	Goiânia	8298 98	0	139	91	1N
BR_GO_521130 BR	Brazil	Goiás	Itaúba	2246 41	0	67	34	1N
BR_MT_520130 BR	Brazil	Mato Grosso	Alto Rio Paranaíba	1416 151	0	41	14	1N

On this page, select **Create a report** -> **Create**

The screenshot shows the Power Query Editor interface. On the left, there's a 'Queries [1]' pane with a single query named 'covid19_top10_countries'. The main area displays a table with columns: location_key, country_code, country_name, subregion1_name, subregion2_name, cumulative_confirmed, cumulative_deceased, and cumulative_recovered. The data consists of 27 rows of Brazilian cities and their COVID-19 statistics. On the right, there's a 'Query settings' pane with sections for 'Properties' (Name: covid19_top10_countries) and 'Applied steps' (Source, Promoted, etc.). At the bottom right of the main area, there's a green 'Create a report' button with a dropdown arrow, which is circled in red.

Click on the map icon to begin setting up the visualization

The screenshot shows the Power BI workspace. On the left, there's a navigation bar with icons for Home, Create, Visualizations, Apps, Metrics, Monitor, Learn, and Workspaces. The 'My workspace' section is selected. In the center, there's a map visualization with several gray dots representing data points. To the right of the map is a 'Visualizations' pane containing various icons for different types of visualizations like charts, tables, and maps. One specific map icon is highlighted with a red circle. Below the map icon is a 'Filters' pane with sections for Filters on this visual, Filters on this page, and Filters on all pages. To the right of the filters is a 'Data' pane listing numerous fields from the 'covid19_top10_countries' query, such as adult_male_mortality_rate, country_code, country_name, cumulative_confirmed, cumulative_deceased, etc.

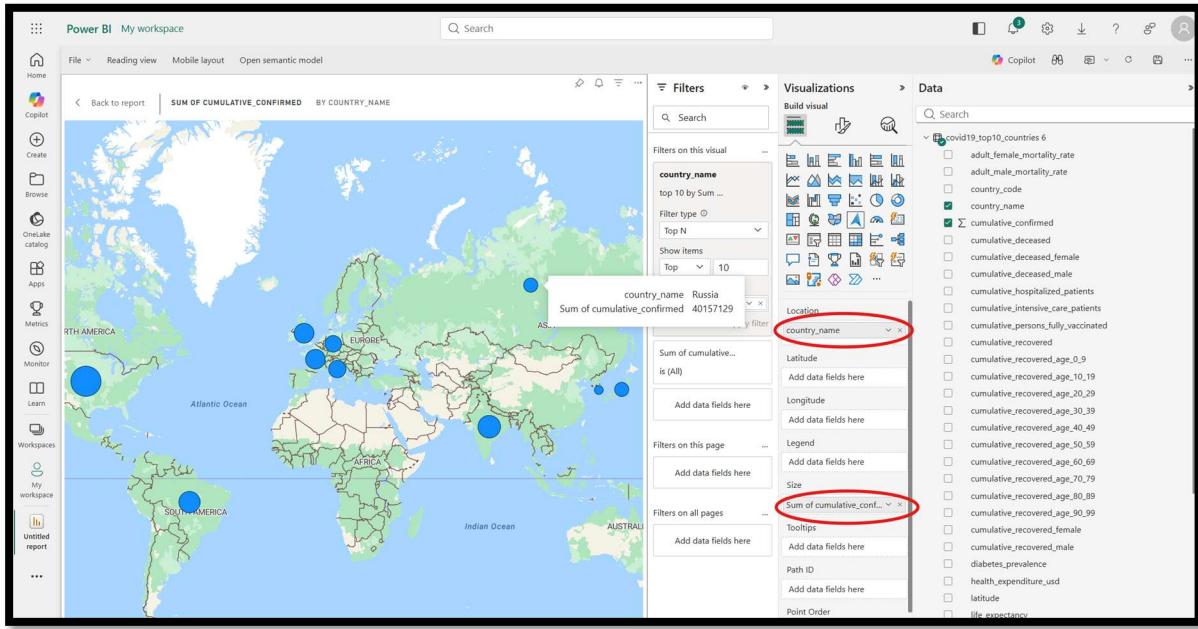
Drag the **country_name** to the filters on this visual

The screenshot shows a Power BI workspace with a map visualization on the left. The map displays several locations with bubbles of varying sizes. On the right, the 'Filters' pane is open, showing the 'Filters on this visual' section. A filter for 'country_name' is applied, set to '(All)'. Below it, the 'Filter type' dropdown is set to 'Top N', which is highlighted with a red circle. Under 'Top N', there is a list of countries with their cumulative confirmed cases: Brazil (5615), France (114), and others. The 'Data' pane on the far right lists various COVID-19 metrics, with 'country_name' also circled in red.

On Filter type click Top N, then on show items click Top then next to it change it to **10** then drag the **cumulative_confirmed** to By value, then click **Apply filter**

This screenshot shows the same Power BI interface after applying the changes described in the text. In the 'Filters' pane, the 'Filter type' is now set to 'Top N', and the 'Show items' dropdown has 'Top' selected with the value '10' entered. Below this, the 'By value' dropdown is expanded, showing 'Sum of cumulative_co...' followed by a dropdown arrow. The 'Data' pane on the right still shows the 'cumulative_confirmed' metric circled in red.

Drag the **country_name** field to the **Location** area and the **cumulative_confirmed** field to the **Size** area



References

1. **COVID19-Data-Analysis: Final Project for CIS 4560. COVID-19 data analysis using HDFS Hive** - <https://github.com/chica-94/COVID19-Data-Analysis>