COVID Vaccines Analysis

Phase 4: Development Part 2

Objective:

The goal is to advance the COVID vaccine analysis through exploratory data analysis (EDA), statistical examination, and visualization. The focus is on uncovering patterns, trends, and relationships within the data to inform evidence-based insights. By employing a combination of statistical tools and visualizations, the aim is to provide a comprehensive understanding of vaccination dynamics, identify global trends, and contribute valuable insights for strategic decision-making.

1. Importing Required Libraries

import numpy as np
import pandas as pd
import seaborn as sns
import matplotlib.pyplot as plt
import plotly.express as px
import plotly.graph_objs as go
from plotly.offline import init_notebook_mode, iplot, plot

2. Loading Data Sets

Read data from CSV files

df_manufacturer = pd.read_csv('/content/drive/MyDrive/Naan
Mudalvan/country_vaccinations_by_manufacturer.csv')
df_vaccinations = pd.read_csv('/content/drive/MyDrive/Naan
Mudalvan/country_vaccinations.csv')

Columns Present in Given Data Sets

df_manufacturer.columns

Output:

['location', 'date', 'vaccine', 'total_vaccinations']

df vaccinations.columns

```
Output:
```

```
['country', 'iso_code', 'date', 'total_vaccinations', 'people_vaccinated', 'people_fully_vaccinated', 'daily_vaccinations_raw', 'daily_vaccinations', 'total_vaccinations_per_hundred', 'people_vaccinated_per_hundred', 'people_fully_vaccinated_per_hundred', 'daily_vaccinations_per_million', 'vaccines', 'source_name', 'source_website']
```

Shape of DataFrames

df_manufacturer.shape

Output:

(35623, 4)

df_vaccinations.shape

Output:

(86512, 15)

Information about Given Data Sets

df manufacturer.info()

Output:

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 35623 entries, 0 to 35622
Data columns (total 4 columns):
  Column
                    Non-Null Count Dtype
--- -----
                     -----
0 location
                   35623 non-null object
1 date
                    35623 non-null object
2
   vaccine
                     35623 non-null object
   total_vaccinations 35623 non-null int64
dtypes: int64(1), object(3)
memory usage: 1.1+ MB
```

df_vaccinations.info()

Output:

Rang	ss 'pandas.core.frame.DataFrame'> eIndex: 86512 entries, 0 to 86511 columns (total 15 columns):					
#	Column	Non-Null Count	Dtype			
0	country	86512 non-null	object			
1	iso_code	86512 non-null	object			
2	date	86512 non-null	object			
3	total_vaccinations	43607 non-null	float64			
4	people_vaccinated	41294 non-null	float64			
5	people_fully_vaccinated	38802 non-null	float64			
6	daily_vaccinations_raw	35362 non-null	float64			
7	daily_vaccinations	86213 non-null	float64			
8	total_vaccinations_per_hundred	43607 non-null	float64			
9	people_vaccinated_per_hundred	41294 non-null	float64			
10	people_fully_vaccinated_per_hundred	38802 non-null	float64			
11	daily_vaccinations_per_million	86213 non-null	float64			
12	vaccines	86512 non-null	object			
13	source_name	86512 non-null	object			
14	source_website	86512 non-null	object			
dtypes: float64(9), object(6)						
memory usage: 9.9+ MB						

Vaccines Manufactured on a Particular Date

$$\label{eq:df_manufacturer} \begin{split} df_manufacturer &= df_manufacturer[df_manufacturer.date == '2022-02-04'] \\ df_manufacturer.head() \end{split}$$

Output:

	location	date	vaccine	total_vaccinations
2305	Argentina	2022-02-04	CanSino	468481
2306	Argentina	2022-02-04	Moderna	5318406
2307	Argentina	2022-02-04	Oxford/AstraZeneca	25606912
2308	Argentina	2022-02-04	Pfizer/BioNTech	11225368
2309	Argentina	2022-02-04	Sinopharm/Beijing	27396208

Country-Wise Vaccination Status on a Particular Date

$$\label{eq:df_vaccinations} \begin{split} df_vaccinations = df_vaccinations[df_vaccinations.date == '2022-02-04'] \\ df_vaccinations.head() \end{split}$$

	country	iso_code	date total_vaccination	ns people_vaccinat	ed people_fully_vaccinated	daily_vaccinations_ra	<pre>daily_vaccinations</pre>	total_vaccinations_per_hundred	people_vaccinated_per_hundred people	e_fully_vaccinated_per_hundred	daily_vaccinations_per_million	vaccines	source_name	source_
347	Afghanistan	AFG	2022- 02-04 Na	aN Na	aN NaN	Nati	12299.0	NaN	NaN	NaN	309.0	Johnson&Johnson, Oxford/AstraZeneca, Pfizer/Bi	World Health Organization	https://covid19
784	Abania	ALB	2022- 02-04 Na	an Na	IN NaN	Nat	16144.0	NaN	NaN	NaN	5619.0	Oxford/AstraZeneca, Pfizer/BioNTech, Sinovac,	Ministry of Health	https://shendetesia.gov.al/v an
1204	Algeria	DZA	2022- 02-04 Na	ıN Nı	aN NaN	Nat	16222.0	NaN	NaN	NaN	364.0		World Health Organization	https://covid19
1613	Andorra	AND	2022- 02-04 Na	aN Na	aN NaN	Nat	126.0	NaN	NaN	NaN	1629.0	Moderna, Oxford/AstraZeneca, Pfizer/BioNTech	World Health Organization	https://covid19
1991	Angola	AGO	2022- 02-04 Na	aN Na	aN NaN	Nat	83460.0	NaN	NaN	NaN	2460.0	Oxford/AstraZeneca	World Health Organization	https://covid19

3. Preprocessing Data

I. Null Values:

- Null values, often represented as NaN (Not a Number) in Python, indicate missing or undefined data.
- It's crucial to identify and handle null values, as they can affect the accuracy of our analysis or machine learning models.

Common methods to handle null values include:

- a) Removing Rows: If a small percentage of rows have null values and removing them won't significantly impact our analysis.
- b) Imputation: Fill null values with the mean, median, or mode of the respective column.
- c) Forward/Backward Fill: Use the values from the previous or next row to fill null values.
- d) Interpolation: Estimate missing values based on the values of other rows using methods like linear interpolation.

II. Missing Values:

- Missing values can occur due to various reasons such as data collection errors or intentional gaps.
- Techniques for handling missing values are similar to those for null values.

III. Outliers:

- Outliers are data points that deviate significantly from the rest of the data.
- Identifying outliers is crucial for accurate analysis and modeling.

Common methods for detecting outliers include:

a) Visual Inspection: Plotting the data and looking for points that deviate from the overall pattern.

- b) Statistical Methods: Using measures like Z-scores or IQR (Interquartile Range) to identify values that are significantly different from the mean or median.
- c) Machine Learning Models: Some models are sensitive to outliers, so detecting them during model training can be beneficial.

IV. Removing Duplicates:

- Duplicate values in a dataset can arise from data entry errors or other issues.
- Removing duplicates ensures that each data point is unique.
- In Python, you can use the `drop_duplicates` method for DataFrames in pandas to remove duplicate rows.
- Columns can also be checked for duplicates using methods like `duplicated()`.

Checking for Missing Values

df_manufacturer.isna().sum()

Output:

location 0
date 0
vaccine 0
total_vaccinations 0
dtype: int64

df_vaccinations.isna().sum()

Output:

country	0
iso_code	0
date	0
total_vaccinations	112
people_vaccinated	119
people_fully_vaccinated	115
daily_vaccinations_raw	132
daily_vaccinations	0
total_vaccinations_per_hundred	112
people_vaccinated_per_hundred	119
people_fully_vaccinated_per_hundred	115
daily_vaccinations_per_million	0
vaccines	0
source_name	0
source_website	0
dtype: int64	

Dropping Missing Values

df_vaccinations.isna().sum()

Output:

```
country
                                             0
                                             0
iso_code
date
                                             0
total_vaccinations
                                           112
people_vaccinated
                                           119
people_fully_vaccinated
daily_vaccinations_raw
                                           115
                                           132
daily vaccinations
                                             0
total_vaccinations_per_hundred
people_vaccinated_per_hundred
                                           112
                                           119
people_fully_vaccinated_per_hundred
                                           115
daily_vaccinations_per_million
vaccines
source_name
source_website
dtype: int64
```

df vaccinations =

 $df_vaccinations.drop(df_vaccinations[df_vaccinations.total_vaccinations.isna()].index)$

df_vaccinations =

 $df_vaccinations.drop(df_vaccinations[df_vaccinations.people_vaccinated.isna()].index)$

df_vaccinations =

df vaccinations.drop(df vaccinations[df vaccinations.daily vaccinations raw.isna()].index)

Checking for Null Values

df_vaccinations.isnull().sum()

Output:

```
country
iso_code
date
total_vaccinations
people_vaccinated
people_fully_vaccinated
daily_vaccinations_raw
daily_vaccinations
total_vaccinations_per_hundred
people_vaccinated_per_hundred
people_fully_vaccinated_per_hundred
daily_vaccinations_per_million
vaccines
                                      0
                                      0
source_name
source_website
dtype: int64
```

Filling Mean Values

```
df_vaccinations = df_vaccinations.fillna(df_vaccinations.mean())
df_vaccinations.isnull().sum()
```

Output:

```
country
iso code
date
total vaccinations
people vaccinated
people_fully_vaccinated
                                     0
daily_vaccinations_raw
                                      0
daily_vaccinations
total_vaccinations_per_hundred
people_vaccinated_per_hundred
people_fully_vaccinated_per_hundred
daily_vaccinations_per_million
vaccines
source name
                                      0
source_website
dtype: int64
```

Checking for Duplicated Records

```
duplicate_rows = df_vaccinations[df_vaccinations.duplicated()]
print(len(duplicate_rows))
print(duplicate_rows)
Output:
```

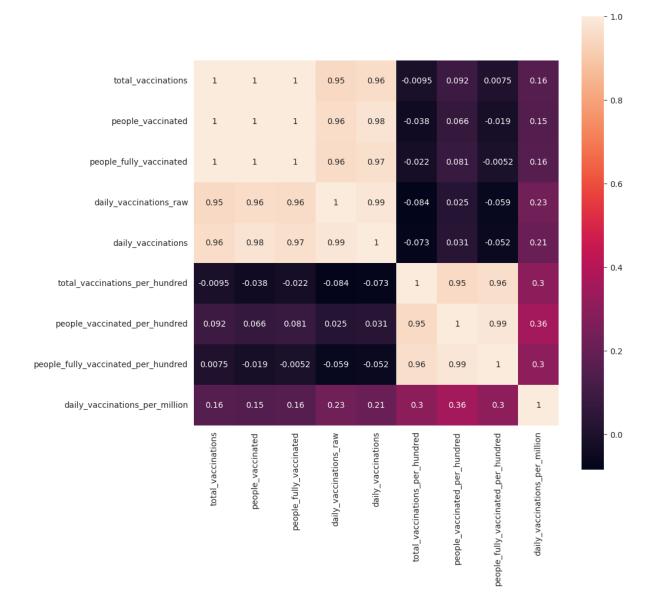
```
0
Empty DataFrame
Columns: [country, iso_code, date, total_vaccinations,
Index: []
```

4. Visualization and Statistical Analysis of Data

Visualization techniques, such as heatmaps, bar plots, and histograms, are employed to illustrate correlations, top countries in vaccination utilization, and distribution of daily vaccinations. The analysis go through statistical summaries of key attributes, including total vaccinations, people vaccinated, and daily vaccinations. Additionally, country-specific analyses, such as preferred vaccines in India and daily vaccinations per million in top countries, offer targeted insights.

Heatmap Visualization to Check Correlation Between Attributes

plt.subplots(figsize=(10, 10))
sns.heatmap(df_vaccinations.corr(), annot=True, square=True)
plt.show()



Top Countries in Vaccination Utilization

df_vaccinations["Total_vaccinations_count"] =
df_vaccinations.groupby("country").total_vaccinations.tail(1)

```
country
India 1.687048e+09
United States 5.469684e+08
                3.677782e+08
Pakistan
                1.823960e+08
Vietnam
                1.816654e+08
Mexico
                 1.685357e+08
Germany
                 1.666940e+08
Russia
                 1.553786e+08
Turkey
                 1.427355e+08
United Kingdom 1.384598e+08
```

Name: Total_vaccinations_count, dtype: float64

 $\mathbf{x} =$

df_vaccinations.groupby("country")["Total_vaccinations_count"].mean().sort_values(ascending =False).head(10)

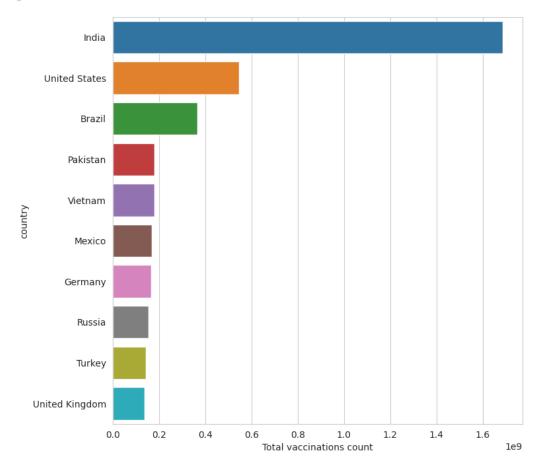
sns.set_style("whitegrid")

plt.figure(figsize=(8, 8))

ax = sns.barplot(x=x.values, y=x.index)

ax.set_xlabel("Total vaccinations count")

plt.show()



Fully Vaccinated Count

df_vaccinations["Full_vaccinations_count"] =
df_vaccinations.groupby("country").people_fully_vaccinated.tail(1)

country India 724768356.0 United States 213893460.0 Brazil 150682483.0 Pakistan 84731497.0 Mexico 77478070.0 Vietnam 74187748.0 Russia 70232028.0 Germany 61873548.0 Iran 54405243.0 Turkey 52489431.0

Name: Full_vaccinations_count, dtype: float64

 $\mathbf{x} =$

df_vaccinations.groupby("country")["Full_vaccinations_count"].mean().sort_values(ascending=False).head(10)

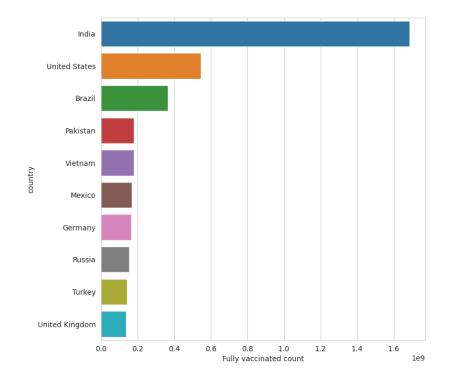
sns.set_style("whitegrid")

plt.figure(figsize=(8, 8))

ax = sns.barplot(x=x.values, y=x.index)

ax.set_xlabel("Fully vaccinated count")

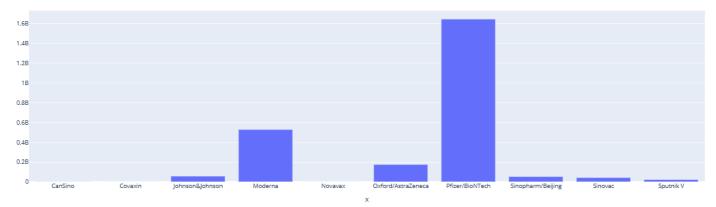
plt.show()



Most Commonly Used Vaccines in the World

total = df_manufacturer.groupby('vaccine').sum()

px.bar(x=total.index, y=total['total_vaccinations'], title='Most Used Vaccine in the World')



People Vaccinated per Hundred for the Date 2022-02-04

df_vaccinations = df_vaccinations[df_vaccinations['date'] == '2022-02-04']

df_vaccinations = df_vaccinations.sort_values(by='people_vaccinated_per_hundred',

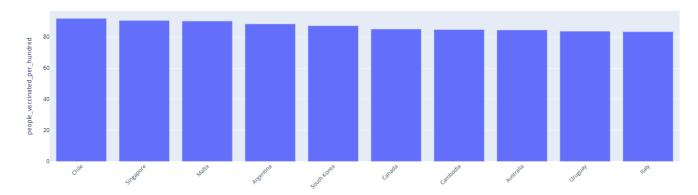
ascending=False)

fig = px.bar(df_vaccinations.head(10), x='country', y='people_vaccinated_per_hundred',

title='People Vaccinated per Hundred for the Date 2022-02-04')

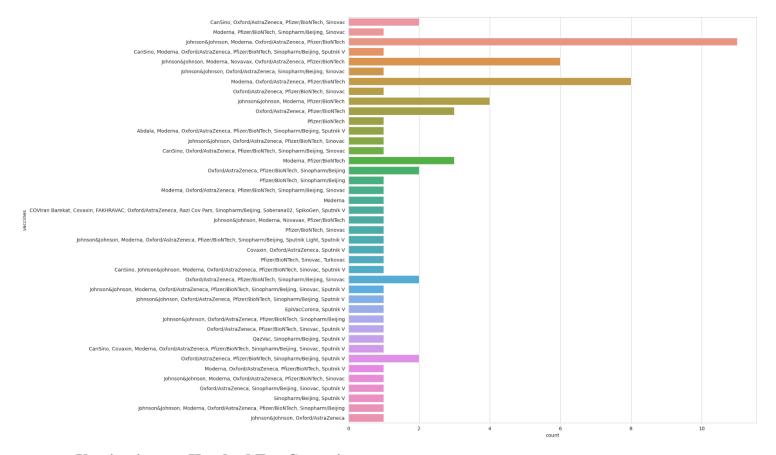
fig.update_layout(xaxis_tickangle=-45)

fig.show()



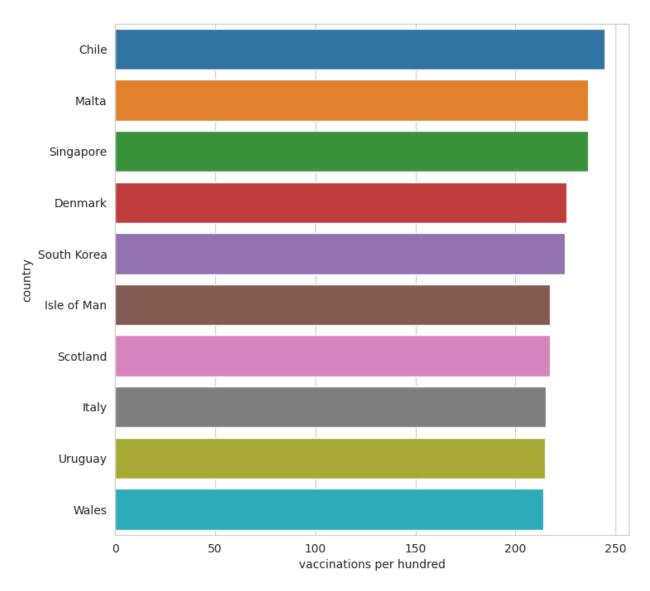
Type of Vaccine Utilized vs Count

plt.figure(figsize=(15, 15))
sns.countplot(y="vaccines", data=df_vaccinations)
plt.show()



Vaccination per Hundred Top Countries

```
df_vaccinations["Total_vaccinations_per_hundred"] =
df_vaccinations.groupby("country").total_vaccinations_per_hundred.tail(1)
x =
df_vaccinations.groupby("country")["Total_vaccinations_per_hundred"].mean().sort_values(asc ending=False).head(10)
plt.figure(figsize=(8, 8))
ax = sns.barplot(x=x.values, y=x.index)
ax.set_xlabel("Vaccinations per hundred")
plt.show()
```

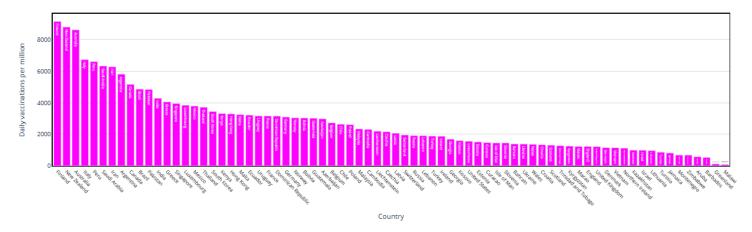


Country-Wise Daily Vaccination per Million

```
def trace_bar(data, feature, title, xlab, ylab, color):
    data = data.sort_values(feature, ascending=False)
    trace = go.Bar(
        x=data['country'],
        y=data[feature],
        marker=dict(color=color),
        text=data['country']
)
```

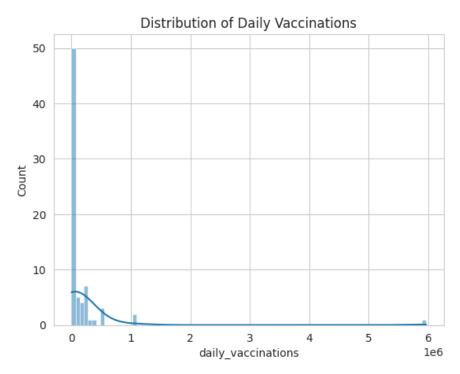
```
data = [trace]
layout = dict(
  title=title,
  xaxis=dict(
     title=xlab,
     showticklabels=True,
     tickangle=45,
     zeroline=True,
     zerolinewidth=1,
     zerolinecolor='grey',
     showline=True,
     linewidth=2,
    linecolor='black',
     mirror=True,
     tickfont=dict(size=10, color='black'),
  ),
  yaxis=dict(
     title=ylab,
    gridcolor='lightgrey',
     zeroline=True,
     zerolinewidth=1,
     zerolinecolor='grey',
     showline=True,
    linewidth=2,
    linecolor='black',
     mirror=True
  ),
  plot_bgcolor='rgba(0, 0, 0, 0)',
  paper_bgcolor='rgba(0, 0, 0, 0)',
  hovermode='closest'
)
```

fig = dict(data=data, layout=layout)
iplot(fig)
trace_bar(df_vaccinations, 'daily_vaccinations_per_million', 'Daily Vaccinations per Million per
Country', 'Country', 'Daily Vaccinations per Million', 'blue')



Distribution of Daily Vaccine

sns.histplot(df_vaccinations['daily_vaccinations'], kde=True)
plt.title("Distribution of Daily Vaccinations")
plt.show()



Statistical Analysis of Given Data Sets

Descriptive Statistics:

- *Mean:* The average of a set of values.
- *Median:* The middle value of a sorted dataset.
- Standard Deviation: A measure of the amount of variation or dispersion in a set of values.

Total Vaccinations Statistical Analysis

df_manufacturer['total_vaccinations'].describe()

Output:

	total_vaccinations				
count	1.600000e+02				
mean	1.574315e+07				
std	5.730594e+07				
min	0.000000e+00				
25%	2.378710e+05				
50%	1.569373e+06				
75%	9.042346e+06				
max	5.821192e+08				

df_vaccinations.describe()

Output:

	total_vaccinations	people_vaccinated	people_fully_vaccinated	daily_vaccinations_raw	daily_vaccinations	total_vaccinations_per_hundred	people_vaccinated_per_hundred	people_fully_vaccinat
count	7.400000e+01	7.400000e+01	7.400000e+01	7.400000e+01	7.400000e+01	74.000000	74.000000	
mean	7.341445e+07	3.528194e+07	3.046394e+07	2.092764e+05	1.883525e+05	156.099054	66.902568	
std	2.094558e+08	1.149868e+08	8.904082e+07	6.877380e+05	7.106291e+05	55.557095	18.881493	
min	6.936300e+04	2.667600e+04	2.607400e+04	0.000000e+00	7.000000e+00	9.610000	7.670000	
25%	3.041524e+06	1.438836e+06	1.296678e+06	3.582500e+03	3.229500e+03	119.397500	57.685000	
50%	1.300016e+07	6.382784e+06	5.586156e+06	3.372050e+04	2.401850e+04	166.300000	72.355000	
75%	6.246783e+07	2.616142e+07	2.915660e+07	1.294548e+05	1.485200e+05	198.355000	80.177500	
max	1.687048e+09	9.487174e+08	7.247684e+08	5.530743e+06	5.964928e+06	244.500000	91.900000	

People Fully Vaccinated Statistical Analysis

df_manufacturer['people_fully_vaccinated'].describe()

```
count 7.400000e+01
mean 3.046394e+07
std 8.904082e+07
min 2.607400e+04
25% 1.296678e+06
50% 5.586156e+06
75% 2.915660e+07
max 7.247684e+08
Name: people_fully_vaccinated, dtype: float64
```

Daily Vaccinations Statistical Analysis

df_manufacturer['daily_vaccinations'].describe()

Output:

```
count 7.400000e+01
mean 1.883525e+05
std 7.106291e+05
min 7.000000e+00
25% 3.229500e+03
50% 2.401850e+04
75% 1.485200e+05
max 5.964928e+06
Name: daily_vaccinations, dtype: float64
```

Total Vaccinations in Country Statistical Analysis

 $df_vaccinations['total_vaccinations'].describe()$

Output:

```
count 7.40000e+01
mean 7.341445e+07
std 2.094558e+08
min 6.936300e+04
25% 3.041524e+06
50% 1.300016e+07
75% 6.246783e+07
max 1.687048e+09
Name: total_vaccinations, dtype: float64
```

People Fully Vaccinated in Country Statistical Analysis

df_vaccinations['people_fully_vaccinated'].describe()

```
count
       7.400000e+01
mean
       3.046394e+07
std
       8.904082e+07
min
        2.607400e+04
25%
        1.296678e+06
50%
       5.586156e+06
75%
        2.915660e+07
max
        7.247684e+08
```

Name: people_fully_vaccinated, dtype: float64

Most Used Vaccine in the World

df_manufacturer['vaccine'].value_counts()

Output:

Pfizer/BioNTech	39		
Moderna	35		
Johnson&Johnson	33		
Oxford/AstraZeneca	30		
Novavax	8		
Sinovac	6		
Sinopharm/Beijing	4		
CanSino	2		
Sputnik V	2		
Covaxin	1		
Name: vaccine, dtype:	int64		

Daily Vaccinations per Million Top Countries

 $\label{lem:country} df_vaccinations.groupby("country")["daily_vaccinations_per_million"].mean().sort_values(ascending=False).head(20)$

```
country
Finland 9154.0

New Zealand 8800.0

Australia 8621.0

Italy 6733.0

Peru 6609.0
Saudi Arabia 6330.0
Iran
                  6280.0
                5814.0
Argentina
Canada
                 5165.0
Brazil
                4864.0
Pakistan
                4841.0
India
                 4281.0
Greece 4055.0
Singapore 3951.0
Luxembourg 3845.0
Mexico
                  3792.0
Thailand 3718.0
South Korea 3447.0
Kenya 3315.0
Hong Kong 3293.0
Name: daily_vaccinations_per_million, dtype: float64
```

Preferred Vaccine in India

```
x = df_vaccinations[df_vaccinations["country"] == "India"]
z = x.vaccines.value_counts()
c = list(z.index)
print(c)
Output:
    ["Covaxin, Oxford/AstraZeneca, Sputnik V"]
```

Outcome:

1. Data Understanding:

• The documentation begins with importing necessary libraries and loading the data sets, providing insight into the tools and datasets used.

2. Exploratory Data Analysis (EDA):

• The EDA section explores the structure and content of the data, presenting key information such as columns, shapes, and data types.

• It also includes the preprocessing steps, checking and handling missing values, and ensuring data quality.

3. Visualization:

- Various visualizations are included to provide a graphical representation of the data. This includes heatmaps, bar plots, and charts showcasing top countries in vaccination, preferred vaccines, and more.
- The visualizations aim to make complex data more understandable and highlight trends and patterns.

4. Statistical Analysis:

- Statistical analysis is performed on key attributes, providing summary statistics such as mean, standard deviation, and quartiles.
- This section enables a quantitative understanding of the data distribution and central tendencies.

5. Insights and Interpretation:

- Throughout the documentation, insights are provided, such as the top countries in vaccination, most commonly used vaccines, and statistical summaries.
- These insights aid in drawing meaningful conclusions from the data, supporting decision-making processes.

6. Customization:

 The documentation is designed to be customizable based on specific requirements. Users can adapt and extend the documentation to suit their analysis goals or share it with others to facilitate collaboration.

Implementation - Phase 4 of COVID Vaccine Analysis:

https://colab.research.google.com/drive/18iN5o_u16y5msl0uB-RTOKWbWn3n8W33?usp=sharing