

# Covid-19 Vaccines Analysis

## Importing Necessary Libraries

```
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
import plotly.express as px
import plotly.graph_objects as go
from sklearn import metrics
from sklearn.metrics import mean_squared_error
from sklearn.metrics import r2_score
```

## Data Loading and Extraction

```
df = pd.read_csv('country_vaccinations.csv')
df0 = pd.read_csv('country_vaccinations.csv')
df.head()
```

	country	iso_code	date	total_vaccinations
people_vaccinated \				
0	Afghanistan	AFG	2021-02-22	0.0
0.0				
1	Afghanistan	AFG	2021-02-23	NaN
NaN				
2	Afghanistan	AFG	2021-02-24	NaN
NaN				
3	Afghanistan	AFG	2021-02-25	NaN
NaN				
4	Afghanistan	AFG	2021-02-26	NaN
NaN				

	people_fully_vaccinated	daily_vaccinations_raw	daily_vaccinations
\			
0	NaN	NaN	NaN
1	NaN	NaN	1367.0
2	NaN	NaN	1367.0
3	NaN	NaN	1367.0
4	NaN	NaN	1367.0

	total_vaccinations_per_hundred	people_vaccinated_per_hundred	\
0	0.0	0.0	
1	NaN	NaN	
2	NaN	NaN	
3	NaN	NaN	
4	NaN	NaN	

	people_fully_vaccinated_per_hundred	daily_vaccinations_per_million	\
0	NaN	NaN	
1	NaN	34.0	
2	NaN	34.0	
3	NaN	34.0	
4	NaN	34.0	

	vaccines	\
0	Johnson&Johnson, Oxford/AstraZeneca, Pfizer/Bi...	
1	Johnson&Johnson, Oxford/AstraZeneca, Pfizer/Bi...	
2	Johnson&Johnson, Oxford/AstraZeneca, Pfizer/Bi...	
3	Johnson&Johnson, Oxford/AstraZeneca, Pfizer/Bi...	
4	Johnson&Johnson, Oxford/AstraZeneca, Pfizer/Bi...	

	source_name	source_website
0	World Health Organization	<a href="https://covid19.who.int/">https://covid19.who.int/</a>
1	World Health Organization	<a href="https://covid19.who.int/">https://covid19.who.int/</a>
2	World Health Organization	<a href="https://covid19.who.int/">https://covid19.who.int/</a>
3	World Health Organization	<a href="https://covid19.who.int/">https://covid19.who.int/</a>
4	World Health Organization	<a href="https://covid19.who.int/">https://covid19.who.int/</a>

```
df.isnull().sum()
```

country	0
iso_code	0
date	0
total_vaccinations	42905
people_vaccinated	45218
people_fully_vaccinated	47710
daily_vaccinations_raw	51150
daily_vaccinations	299
total_vaccinations_per_hundred	42905
people_vaccinated_per_hundred	45218
people_fully_vaccinated_per_hundred	47710
daily_vaccinations_per_million	299
vaccines	0

```

source_name      0
source_website   0
dtype: int64

df1 = df.copy()
df1.info()

<class 'pandas.core.frame.DataFrame'>
RangeIndex: 86512 entries, 0 to 86511
Data 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  datetime64[ns]
3   total_vaccinations                          86512 non-null  float64
4   people_vaccinated                          86512 non-null  float64
5   people_fully_vaccinated                    86512 non-null  float64
6   daily_vaccinations_raw                     86512 non-null  float64
7   daily_vaccinations                         86512 non-null  float64
8   total_vaccinations_per_hundred             86512 non-null  float64
9   people_vaccinated_per_hundred              86512 non-null  float64
10  people_fully_vaccinated_per_hundred         86512 non-null  float64
11  daily_vaccinations_per_million              86512 non-null  float64
12  vaccines                                    86512 non-null  object
13  source_name                                86512 non-null  object
14  source_website                             86512 non-null  object

dtypes: datetime64[ns](1), float64(9), object(5)
memory usage: 9.9+ MB

```

# Handling Missing Values

```
df.fillna({'total_vaccinations': 0,  
'people_vaccinated': 0,  
'people_fully_vaccinated':0,  
'daily_vaccinations_raw':0,  
'daily_vaccinations':0,  
'total_vaccinations_per_hundred': 0,  
'people_vaccinated_per_hundred': 0,  
'people_fully_vaccinated_per_hundred' :0,  
'daily_vaccinations_per_million':0}, inplace=True)
```

```
df.isnull().sum()
```

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

```
df['date'] = pd.to_datetime(df['date'])  
df
```

	country	iso_code	date	total_vaccinations
people_vaccinated \				
0	Afghanistan	AFG	2021-02-22	0.0
0.0				
1	Afghanistan	AFG	2021-02-23	0.0
0.0				
2	Afghanistan	AFG	2021-02-24	0.0
0.0				
3	Afghanistan	AFG	2021-02-25	0.0
0.0				
4	Afghanistan	AFG	2021-02-26	0.0
0.0				
...	...	...	...	...
...				
86507	Zimbabwe	ZWE	2022-03-25	8691642.0
4814582.0				

86508	Zimbabwe	ZWE	2022-03-26	8791728.0
4886242.0				
86509	Zimbabwe	ZWE	2022-03-27	8845039.0
4918147.0				
86510	Zimbabwe	ZWE	2022-03-28	8934360.0
4975433.0				
86511	Zimbabwe	ZWE	2022-03-29	9039729.0
5053114.0				

	people_fully_vaccinated	daily_vaccinations_raw
daily_vaccinations \		
0	0.0	0.0
0.0		
1	0.0	0.0
1367.0		
2	0.0	0.0
1367.0		
3	0.0	0.0
1367.0		
4	0.0	0.0
1367.0		

...	...	...
...		
86507	3473523.0	139213.0
69579.0		
86508	3487962.0	100086.0
83429.0		
86509	3493763.0	53311.0
90629.0		
86510	3501493.0	89321.0
100614.0		
86511	3510256.0	105369.0
103751.0		

	total_vaccinations_per_hundred	people_vaccinated_per_hundred \
0	0.00	0.00
1	0.00	0.00
2	0.00	0.00
3	0.00	0.00
4	0.00	0.00
...	...	...
86507	57.59	31.90

86508	58.25	32.38
86509	58.61	32.59
86510	59.20	32.97
86511	59.90	33.48
people_fully_vaccinated_per_hundred		
daily_vaccinations_per_million \		
0	0.00	
0.0		
1	0.00	
34.0		
2	0.00	
34.0		
3	0.00	
34.0		
4	0.00	
34.0		
...	...	
...		
86507	23.02	
4610.0		
86508	23.11	
5528.0		
86509	23.15	
6005.0		
86510	23.20	
6667.0		
86511	23.26	
6874.0		
vaccines \		
0	Johnson&Johnson, Oxford/AstraZeneca, Pfizer/Bi...	
1	Johnson&Johnson, Oxford/AstraZeneca, Pfizer/Bi...	
2	Johnson&Johnson, Oxford/AstraZeneca, Pfizer/Bi...	
3	Johnson&Johnson, Oxford/AstraZeneca, Pfizer/Bi...	
4	Johnson&Johnson, Oxford/AstraZeneca, Pfizer/Bi...	
...	...	
86507	Oxford/AstraZeneca, Sinopharm/Beijing, Sinovac...	
86508	Oxford/AstraZeneca, Sinopharm/Beijing, Sinovac...	
86509	Oxford/AstraZeneca, Sinopharm/Beijing, Sinovac...	
86510	Oxford/AstraZeneca, Sinopharm/Beijing, Sinovac...	
86511	Oxford/AstraZeneca, Sinopharm/Beijing, Sinovac...	
source_name \		
0	World Health Organization	
1	World Health Organization	

```

2      World Health Organization
3      World Health Organization
4      World Health Organization
...
86507      Ministry of Health
86508      Ministry of Health
86509      Ministry of Health
86510      Ministry of Health
86511      Ministry of Health

                                source_website
0      https://covid19.who.int/
1      https://covid19.who.int/
2      https://covid19.who.int/
3      https://covid19.who.int/
4      https://covid19.who.int/
...
86507  https://www.arcgis.com/home/webmap/viewer.html...
86508  https://www.arcgis.com/home/webmap/viewer.html...
86509  https://www.arcgis.com/home/webmap/viewer.html...
86510  https://www.arcgis.com/home/webmap/viewer.html...
86511  https://www.arcgis.com/home/webmap/viewer.html...

```

[86512 rows x 15 columns]

*#Checking For Duplicate values*

```

df.drop_duplicates(inplace=True)
df[df.duplicated()]

```

Empty DataFrame

Columns: [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]  
Index: []

All the Duplicates values are dropped.

```

df.info()

<class 'pandas.core.frame.DataFrame'>
RangeIndex: 86512 entries, 0 to 86511
Data 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	
	datetime64[ns]			
3	total_vaccinations	86512	non-null	float64
4	people_vaccinated	86512	non-null	float64
5	people_fully_vaccinated	86512	non-null	float64
6	daily_vaccinations_raw	86512	non-null	float64
7	daily_vaccinations	86512	non-null	float64
8	total_vaccinations_per_hundred	86512	non-null	float64
9	people_vaccinated_per_hundred	86512	non-null	float64
10	people_fully_vaccinated_per_hundred	86512	non-null	float64
11	daily_vaccinations_per_million	86512	non-null	float64
12	vaccines	86512	non-null	object
13	source_name	86512	non-null	object
14	source_website	86512	non-null	object

dtypes: datetime64[ns](1), float64(9), object(5)

memory usage: 9.9+ MB

*#Dropping unwanted*

```
df1 = df1.drop(['vaccines', 'source_name', 'source_website']), axis=1)
df1.info()
```

<class 'pandas.core.frame.DataFrame'>

RangeIndex: 86512 entries, 0 to 86511

Data columns (total 12 columns):

#	Column	Non-Null Count	Dtype
---	-----	-----	-----
0	country	86512 non-null	object
1	iso_code	86512 non-null	object
2	date	86512 non-null	
	datetime64[ns]		
3	total_vaccinations	86512 non-null	float64
4	people_vaccinated	86512 non-null	float64



5	people_fully_vaccinated	86512	non-null	float64
6	daily_vaccinations_raw	86512	non-null	float64
7	daily_vaccinations	86512	non-null	float64
8	total_vaccinations_per_hundred	86512	non-null	float64
9	people_vaccinated_per_hundred	86512	non-null	float64
10	people_fully_vaccinated_per_hundred	86512	non-null	float64
11	daily_vaccinations_per_million	86512	non-null	float64

dtypes: datetime64[ns](1), float64(9), object(2)  
memory usage: 7.9+ MB

```
from sklearn.preprocessing import LabelEncoder
le=LabelEncoder()
df2['country']=le.fit_transform(df2['country'])
df2['iso_code']=le.fit_transform(df2['iso_code'])

df2['vaccines']=le.fit_transform(df2['vaccines'])

df2['source_name']=le.fit_transform(df2['source_name'])

df2['source_website']=le.fit_transform(df2['source_website'])
# df2['date'] = df2['date'].str.replace('-', ' ')
```

## Machine Learning in Python

### Testing and Training

```
x=df2[['country', 'iso_code', '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']]
y=df2[['total_vaccinations']]

from sklearn.model_selection import train_test_split
x_train, x_test, y_train, y_test = train_test_split(x,y,
```

```
random_state=42)
x_train.shape, x_test.shape, y_train.shape, y_test.shape

((64884, 13), (21628, 13), (64884, 1), (21628, 1))
```

### Linear Regression Model

```
from sklearn.linear_model import LinearRegression
LR=LinearRegression()
LR.fit(x_train, y_train)
y_pred_LR=LR.predict(x_test)
y_pred_LR

array([[ -1.59766070e+07],
       [ -9.63050813e+06],
       [ -3.70713648e+07],
       ...,
       [  1.09876415e+08],
       [ -1.43512829e+07],
       [ -2.66557595e+06]])

# Model Evaluation
print('R^2:', metrics.r2_score(y_test, y_pred_LR))
print('MAE:', metrics.mean_absolute_error(y_test, y_pred_LR))
print('MSE:', metrics.mean_squared_error(y_test, y_pred_LR))
print('RMSE:', np.sqrt(metrics.mean_squared_error(y_test, y_pred_LR)))

R^2: 0.6376719832015039
MAE: 22774656.10221463
MSE: 1.0086802875248108e+16
RMSE: 100433076.59953521
```

$R^2$ : It is a measure of the linear relationship between X and Y. It is interpreted as the proportion of the variance in the dependent variable that is predictable from the independent variable.

Adjusted  $R^2$ : The adjusted R-squared compares the explanatory power of regression models that contain different numbers of predictors.

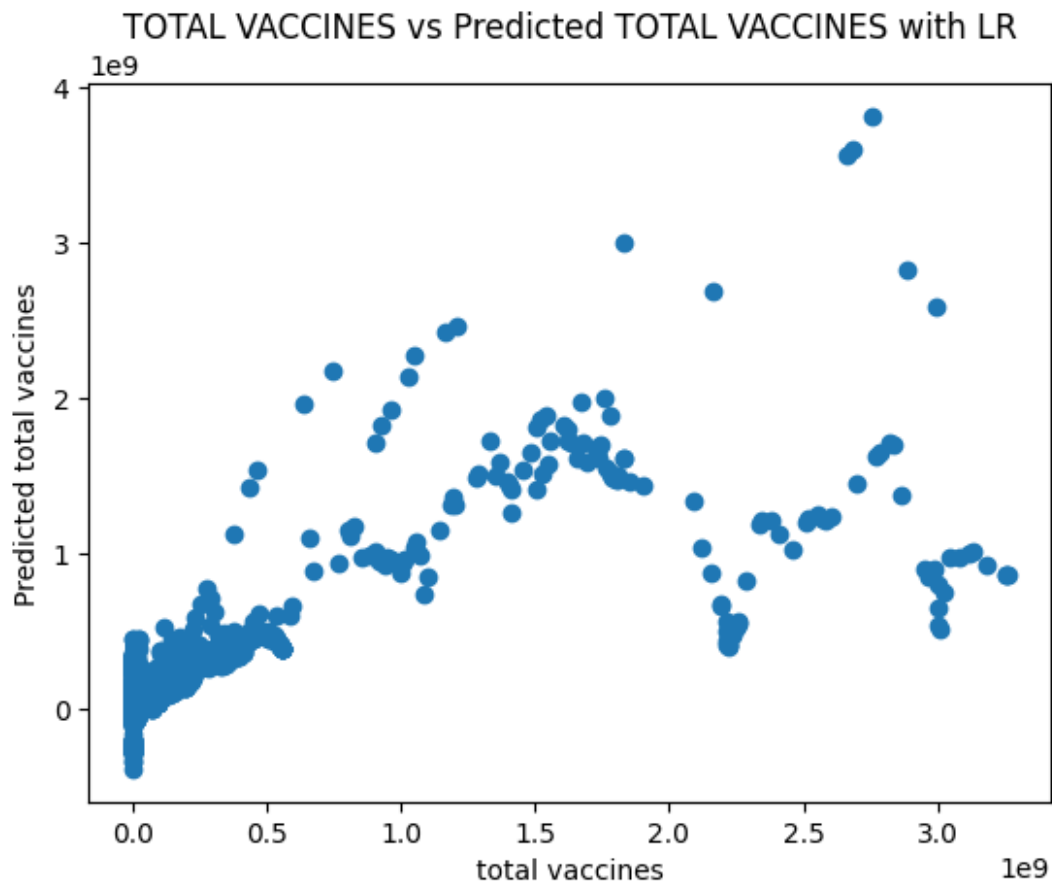
MAE: It is the mean of the absolute value of the errors. It measures the difference between two continuous variables, here actual and predicted values of y.

MSE: The mean square error (MSE) is just like the MAE, but squares the difference before summing them all instead of using the absolute value.

RMSE: The mean square error (MSE) is just like the MAE, but squares the difference before summing them all instead of using the absolute value.

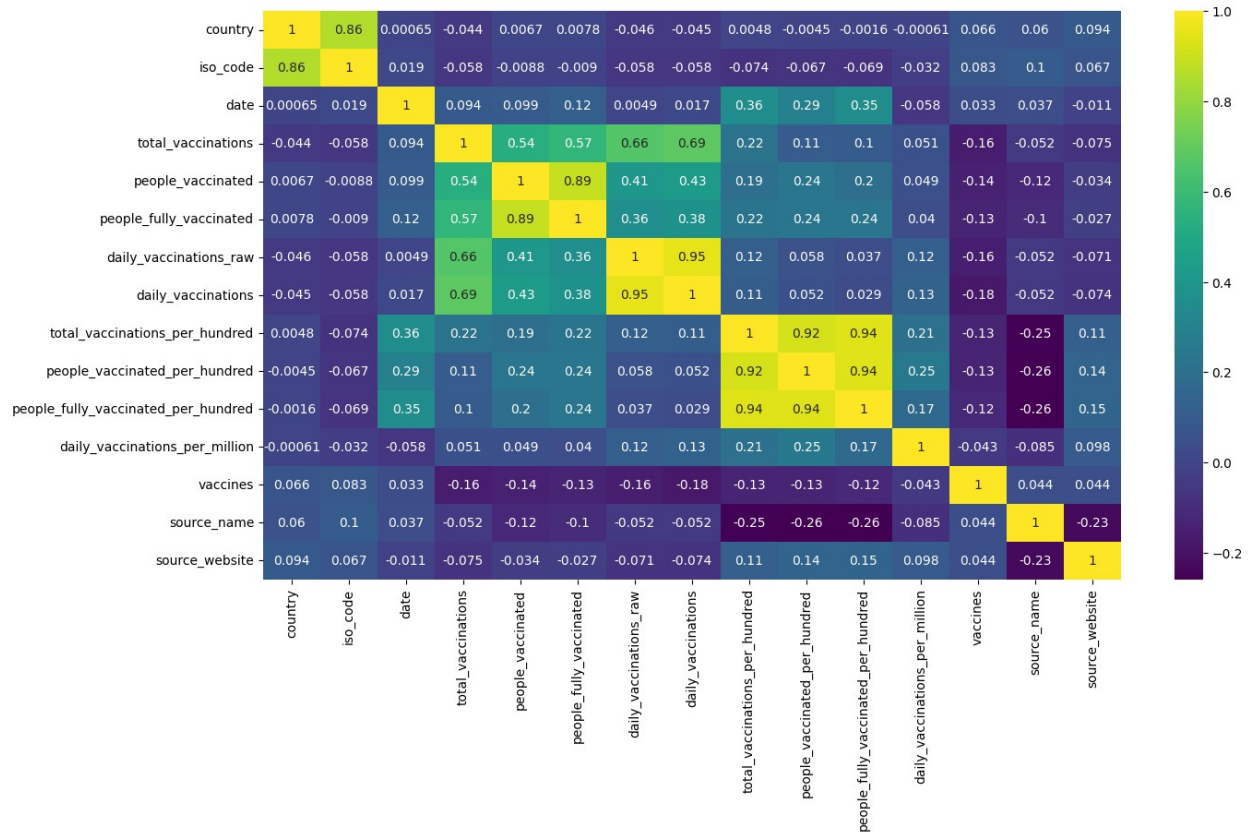
```
plt.scatter(y_test, y_pred_LR)
plt.xlabel("total vaccines")
```

```
plt.ylabel("Predicted total vaccines")
plt.title("TOTAL VACCINES vs Predicted TOTAL VACCINES with LR")
plt.show()
```



```
corr = df2.corr()
plt.figure(figsize=(15,8))
sns.heatmap(corr, cmap='viridis', annot=True)
```

<Axes: >



## Exploratory Data Analysis

```
df0.rename(columns =
{'total_vaccinations_per_hundred': 'total_vaccinations_percent',
'people_fully_vaccinated_per_hundred': 'people_fully_vaccinated_percent',
'people_vaccinated_per_hundred': 'people_vaccinated_percent'},
inplace=True)
df0.columns
Index(['country', 'iso_code', 'date', 'total_vaccinations',
      'people_vaccinated', 'people_fully_vaccinated',
      'daily_vaccinations_raw', 'daily_vaccinations',
      'total_vaccinations_percent', 'people_vaccinated_percent',
      'people_fully_vaccinated_percent',
      'daily_vaccinations_per_million',
      'vaccines', 'source_name', 'source_website'],
      dtype='object')
```

Basic info about Dataset

```

print('Data point starts from:',df0.date.min(),'\n')
print('Data point ends at:',df0.date.max(),'\n')
print('Total no of Countries in the data
set:',len(df0.country.unique()),'\n')
print('Total no of unique Vaccine Schemes in the data
set:',len(df0.vaccines.unique()),'\n')

```

Data point starts from: 2020-12-02

Data point ends at: 2022-03-29

Total no of Countries in the data set: 223

Total no of unique Vaccine Schemes in the data set: 84

```
df0.country.unique()
```

```

array(['Afghanistan', 'Albania', 'Algeria', 'Andorra', 'Angola',
      'Anguilla', 'Antigua and Barbuda', 'Argentina', 'Armenia',
      'Aruba',
      'Australia', 'Austria', 'Azerbaijan', 'Bahamas', 'Bahrain',
      'Bangladesh', 'Barbados', 'Belarus', 'Belgium', 'Belize',
      'Benin',
      'Bermuda', 'Bhutan', 'Bolivia', 'Bonaire Sint Eustatius and
      Saba',
      'Bosnia and Herzegovina', 'Botswana', 'Brazil',
      'British Virgin Islands', 'Brunei', 'Bulgaria', 'Burkina Faso',
      'Burundi', 'Cambodia', 'Cameroon', 'Canada', 'Cape Verde',
      'Cayman Islands', 'Central African Republic', 'Chad', 'Chile',
      'China', 'Colombia', 'Comoros', 'Congo', 'Cook Islands',
      'Costa Rica', 'Cote d'Ivoire', 'Croatia', 'Cuba', 'Curacao',
      'Cyprus', 'Czechia', 'Democratic Republic of Congo', 'Denmark',
      'Djibouti', 'Dominica', 'Dominican Republic', 'Ecuador',
      'Egypt',
      'El Salvador', 'England', 'Equatorial Guinea', 'Estonia',
      'Eswatini', 'Ethiopia', 'Faeroe Islands', 'Falkland Islands',
      'Fiji', 'Finland', 'France', 'French Polynesia', 'Gabon',
      'Gambia',
      'Georgia', 'Germany', 'Ghana', 'Gibraltar', 'Greece',
      'Greenland',
      'Grenada', 'Guatemala', 'Guernsey', 'Guinea', 'Guinea-Bissau',
      'Guyana', 'Haiti', 'Honduras', 'Hong Kong', 'Hungary',
      'Iceland',
      'India', 'Indonesia', 'Iran', 'Iraq', 'Ireland', 'Isle of Man',
      'Israel', 'Italy', 'Jamaica', 'Japan', 'Jersey', 'Jordan',
      'Kazakhstan', 'Kenya', 'Kiribati', 'Kosovo', 'Kuwait',
      'Kyrgyzstan', 'Laos', 'Latvia', 'Lebanon', 'Lesotho',
      'Liberia',
      'Libya', 'Liechtenstein', 'Lithuania', 'Luxembourg', 'Macao',

```

```

'Madagascar', 'Malawi', 'Malaysia', 'Maldives', 'Mali',
'Malta',
'Mauritania', 'Mauritius', 'Mexico', 'Moldova', 'Monaco',
'Mongolia', 'Montenegro', 'Montserrat', 'Morocco',
'Mozambique',
'Myanmar', 'Namibia', 'Nauru', 'Nepal', 'Netherlands',
'New Caledonia', 'New Zealand', 'Nicaragua', 'Niger',
'Nigeria',
'Niue', 'North Macedonia', 'Northern Cyprus', 'Northern
Ireland',
'Norway', 'Oman', 'Pakistan', 'Palestine', 'Panama',
'Papua New Guinea', 'Paraguay', 'Peru', 'Philippines',
'Pitcairn',
'Poland', 'Portugal', 'Qatar', 'Romania', 'Russia', 'Rwanda',
'Saint Helena', 'Saint Kitts and Nevis', 'Saint Lucia',
'Saint Vincent and the Grenadines', 'Samoa', 'San Marino',
'Sao Tome and Principe', 'Saudi Arabia', 'Scotland', 'Senegal',
'Serbia', 'Seychelles', 'Sierra Leone', 'Singapore',
'Sint Maarten (Dutch part)', 'Slovakia', 'Slovenia',
'Solomon Islands', 'Somalia', 'South Africa', 'South Korea',
'South Sudan', 'Spain', 'Sri Lanka', 'Sudan', 'Suriname',
'Sweden',
'Switzerland', 'Syria', 'Taiwan', 'Tajikistan', 'Tanzania',
'Thailand', 'Timor', 'Togo', 'Tokelau', 'Tonga',
'Trinidad and Tobago', 'Tunisia', 'Turkey', 'Turkmenistan',
'Turks and Caicos Islands', 'Tuvalu', 'Uganda', 'Ukraine',
'United Arab Emirates', 'United Kingdom', 'United States',
'Uruguay', 'Uzbekistan', 'Vanuatu', 'Venezuela', 'Vietnam',
'Wales', 'Wallis and Futuna', 'Yemen', 'Zambia', 'Zimbabwe'],
dtype=object)

```

```

# All the different kinds of vaccines

```

```

df0.vaccines.unique()

```

```

array(['Johnson&Johnson', 'Oxford/AstraZeneca', 'Pfizer/BioNTech',
'Sinopharm/Beijing',
'Oxford/AstraZeneca', 'Pfizer/BioNTech', 'Sinovac', 'Sputnik V',
'Oxford/AstraZeneca', 'Sinopharm/Beijing', 'Sinovac', 'Sputnik V',
'Moderna', 'Oxford/AstraZeneca', 'Pfizer/BioNTech',
'Oxford/AstraZeneca', 'Oxford/AstraZeneca', 'Pfizer/BioNTech',
'Oxford/AstraZeneca', 'Pfizer/BioNTech', 'Sputnik V',
'CanSino', 'Moderna', 'Oxford/AstraZeneca', 'Pfizer/BioNTech',
'Sinopharm/Beijing', 'Sputnik V',
'Moderna', 'Oxford/AstraZeneca', 'Sinopharm/Beijing', 'Sinovac',
'Sputnik V',
'Pfizer/BioNTech',
'Johnson&Johnson', 'Moderna', 'Novavax', 'Oxford/AstraZeneca',
'Pfizer/BioNTech',
'Johnson&Johnson', 'Oxford/AstraZeneca', 'Pfizer/BioNTech',
'Johnson&Johnson', 'Moderna', 'Oxford/AstraZeneca', 'Pfizer/BioNTech',

```

Sinopharm/Beijing, Sputnik Light, Sputnik V',  
    'Johnson&Johnson, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech,  
Sinopharm/Beijing, Sinovac',  
    'Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing',  
    'Sinopharm/Beijing, Sputnik V',  
    'Johnson&Johnson, Moderna, Oxford/AstraZeneca,  
Pfizer/BioNTech',  
    'Johnson&Johnson, Oxford/AstraZeneca, Pfizer/BioNTech,  
Sinovac',  
    'Moderna, Oxford/AstraZeneca, Pfizer/BioNTech,  
Sinopharm/Beijing',  
    'Johnson&Johnson, Oxford/AstraZeneca, Pfizer/BioNTech,  
Sinopharm/Beijing, Sputnik V',  
    'Moderna, Pfizer/BioNTech',  
    'Covaxin, Johnson&Johnson, Moderna, Oxford/AstraZeneca,  
Pfizer/BioNTech, Sinovac',  
    'Johnson&Johnson, Oxford/AstraZeneca',  
    'Johnson&Johnson, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech,  
Sinopharm/Beijing',  
    'Johnson&Johnson, Oxford/AstraZeneca, Sinopharm/Beijing',  
    'Sinopharm/Beijing',  
    'Johnson&Johnson, Oxford/AstraZeneca, Sinopharm/Beijing,  
Sinovac',  
    'Covaxin, Oxford/AstraZeneca',  
    'CanSino, Oxford/AstraZeneca, Pfizer/BioNTech, Sinovac',  
    'CanSino, Sinopharm/Beijing, Sinopharm/Wuhan, Sinovac, ZF2001',  
    'Johnson&Johnson, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech,  
Sinovac',  
    'Covaxin, Oxford/AstraZeneca, Sinopharm/Beijing',  
    'Moderna, Oxford/AstraZeneca, Sinopharm/Beijing, Sputnik V',  
    'Abdala, Soberana Plus, Soberana02',  
    'Johnson&Johnson, Moderna, Pfizer/BioNTech',  
    'Johnson&Johnson, Oxford/AstraZeneca, Pfizer/BioNTech,  
Sinopharm/Beijing, Sinovac, Sputnik V',  
    'Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing,  
Sinovac',  
    'Covaxin, Johnson&Johnson, Oxford/AstraZeneca,  
Sinopharm/Beijing, Sinovac',  
    'Johnson&Johnson, Pfizer/BioNTech',  
    'Pfizer/BioNTech, Sinopharm/Beijing, Sputnik V',  
    'Oxford/AstraZeneca, Sputnik V', 'Moderna',  
    'Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sputnik V',  
    'Oxford/AstraZeneca, Sinopharm/Beijing',  
    'Moderna, Oxford/AstraZeneca, Pfizer/BioNTech,  
Sinopharm/Beijing, Sputnik V',  
    'Johnson&Johnson, Moderna',  
    'Johnson&Johnson, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech,  
Sputnik V',  
    'Pfizer/BioNTech, Sinovac',

'Johnson&Johnson, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sputnik V',  
'Covaxin, Oxford/AstraZeneca, Sputnik V',  
'Johnson&Johnson, Moderna, Novavax, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sinovac',  
'COVIran Barekat, Covaxin, FAKHRAVAC, Oxford/AstraZeneca, Razi Cov Pars, Sinopharm/Beijing, Soberana02, SpikoGen, Sputnik V',  
'Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sputnik V',  
'QazVac, Sinopharm/Beijing, Sputnik V',  
'Johnson&Johnson, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sinovac, Sputnik Light, Sputnik V',  
'Johnson&Johnson, Moderna, Novavax, Pfizer/BioNTech',  
'Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sinovac, Sputnik V',  
'Pfizer/BioNTech, Sinopharm/Beijing',  
'CanSino, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sinovac',  
'CanSino, Johnson&Johnson, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinovac, Sputnik V',  
'Abdala, Johnson&Johnson, Oxford/AstraZeneca, Pfizer/BioNTech, Soberana02, Sputnik Light, Sputnik V',  
'Oxford/AstraZeneca, Pfizer/BioNTech, Sinovac',  
'CanSino, Covaxin, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sinovac, Sputnik V',  
'Johnson&Johnson, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sinovac, Sputnik Light, Sputnik V',  
'Covaxin, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sinovac, Sputnik V',  
'EpiVacCorona, Sputnik V',  
'Johnson&Johnson, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sinovac, Sputnik V',  
'Pfizer/BioNTech, Sputnik V',  
'Oxford/AstraZeneca, Sinopharm/Beijing, Sputnik V',  
'Moderna, Pfizer/BioNTech, Sinopharm/Beijing, Sinovac',  
'Johnson&Johnson, Moderna, Novavax, Oxford/AstraZeneca, Pfizer/BioNTech, Sputnik V',  
'Johnson&Johnson, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sinovac',  
'Johnson&Johnson, Oxford/AstraZeneca, Sinopharm/Beijing, Sinovac, Sputnik Light, Sputnik V',  
'Medigen, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech',  
'Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinovac, Sputnik V',  
'Johnson&Johnson, Pfizer/BioNTech, Sinopharm/Beijing',  
'Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing, Sinovac',  
'Pfizer/BioNTech, Sinovac, Turkovac',  
'EpiVacCorona, Oxford/AstraZeneca, QazVac, Sinopharm/Beijing,



```

Sputnik V, ZF2001',
'Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing,
Sinopharm/Wuhan, Sputnik V',
'Moderna, Oxford/AstraZeneca, Pfizer/BioNTech, Sinovac, Sputnik
Light, Sputnik V, ZF2001',
'Abdala, Sinopharm/Beijing, Sinovac, Soberana02, Sputnik Light,
Sputnik V',
'Abdala, Moderna, Oxford/AstraZeneca, Pfizer/BioNTech,
Sinopharm/Beijing, Sputnik V',
'Johnson&Johnson, Oxford/AstraZeneca, Sinovac'], dtype=object)

```

*# Here we are creating `country\_data` which store basic info about a country, like the vaccine scheme it uses, total # vaccinations completed and its percentage with the population*

```

country_data = df0.copy()
cols = ['country', 'total_vaccinations', 'iso_code', 'vaccines',
'total_vaccinations_percent']

country_data =
country_data[cols].groupby('country').max().sort_values('total_vaccina
tions', ascending=False)
country_data.reset_index(inplace = True)

country_data.columns = ['Country', 'Total Vaccinations', 'iso_code',
'Vaccines', 'Total Vaccinations Percentage']
country_data

```

	Country	Total Vaccinations	iso_code	\
0	China	3.263129e+09	CHN	
1	India	1.834501e+09	IND	
2	United States	5.601818e+08	USA	
3	Brazil	4.135596e+08	BRA	
4	Indonesia	3.771089e+08	IDN	
...	...	...	...	
218	Falkland Islands	4.407000e+03	FLK	
219	Montserrat	4.211000e+03	MSR	
220	Niue	4.161000e+03	NIU	
221	Tokelau	1.936000e+03	TKL	
222	Pitcairn	9.400000e+01	PCN	

	Vaccines	\
0	CanSino, Sinopharm/Beijing, Sinopharm/Wuhan, S...	
1	Covaxin, Oxford/AstraZeneca, Sputnik V	
2	Johnson&Johnson, Moderna, Pfizer/BioNTech	
3	Johnson&Johnson, Oxford/AstraZeneca, Pfizer/Bi...	
4	Johnson&Johnson, Moderna, Novavax, Oxford/Astr...	
...	...	
218	Oxford/AstraZeneca	
219	Oxford/AstraZeneca	

```

220
221
222
Pfizer/BioNTech
Pfizer/BioNTech
Oxford/AstraZeneca

```

```

Total Vaccinations Percentage
0 225.94
1 131.66
2 168.72
3 193.26
4 136.45
.. ..
218 124.91
219 84.54
220 257.81
221 141.52
222 200.00

```

```
[223 rows x 5 columns]
```

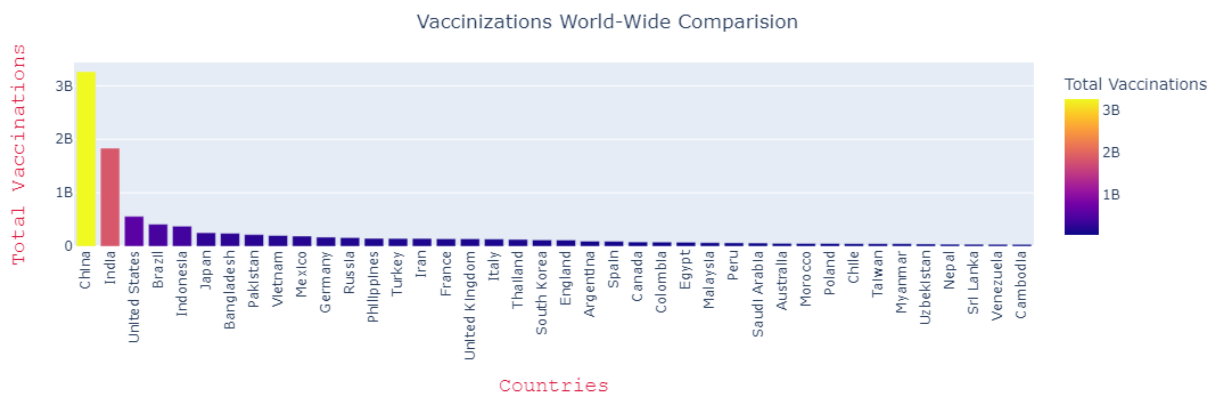
```

fig = px.bar(country_data[:40], x = 'Country', y = 'Total
Vaccinations', color = 'Total Vaccinations')

fig.update_layout(title = dict(text = 'Vaccinizations World-Wide
Comparision', x=0.5, y=0.95))
fig.update_xaxes(title = 'Countries', title_font = dict(size=18,
family='Courier', color='crimson'), tickangle=-90)
fig.update_yaxes(title = 'Total Vaccinations', title_font =
dict(size=18, family='Courier', color='crimson'))

fig.show()

```



From the plot, some interesting facts stand out:

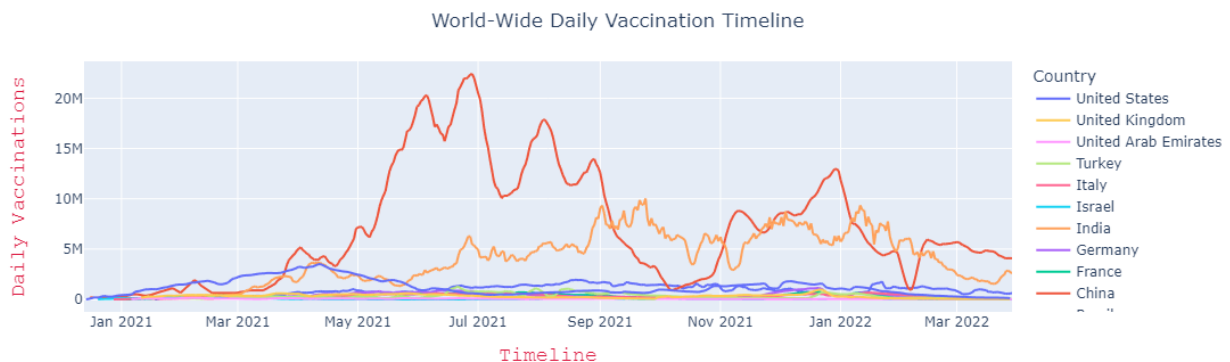
- The United States, despite having the highest number of people affected by Covid-19, has the highest number of vaccinated people.
- China, from where the virus started spreading, is at second.

- India, who has been supplying vaccines to the world is at 3th position.
- UK, where we have found a new variant strain of the virus, is right next.
- Following that, we have Israel, UAE, Brazil, Germany and others

```
top_countries =
['USA', 'CHN', 'GBR', 'IND', 'ISR', 'ARE', 'BRA', 'DEU', 'TUR', 'ITA', 'FRA']
fig = px.line(df0[df0.iso_code.isin(top_countries)], x='date',
y='daily_vaccinations', color='country')

fig.update_layout(title = dict(text = 'World-Wide Daily Vaccination
Timeline', x=0.5, y=0.95),
                    legend = dict(title = 'Country', traceorder =
'reversed'))
fig.update_xaxes(title = 'Timeline', title_font = dict(size=18,
family='Courier', color='crimson'))
fig.update_yaxes(title = 'Daily Vaccinations', title_font =
dict(size=18, family='Courier', color='crimson'))

fig.show()
#Country wise daily vaccination
```



From the plot, we can deduce:

- The Line plot for China is composed entirely of straight lines. This can be attributed to the CCP which tries to restrict flow of information in and out of China. Thus, information from China usually comes in intervals and can be taken with a grain of salt.
- Comparatively, the plot of vaccinations in the USA is better plotted. We can also see that while the USA was heavily affected by the virus, its vaccination drive is highly effective.
- Others like the UK have a steady increase in Daily Vaccinations and India, while supplying to many countries, maintains a respectable 3th position.

```
fig = px.treemap(country_data, path = ['Vaccines', 'Country'], values
= 'Total Vaccinations', height = 650,
                    custom_data = ['Country', 'Vaccines', 'Total
```

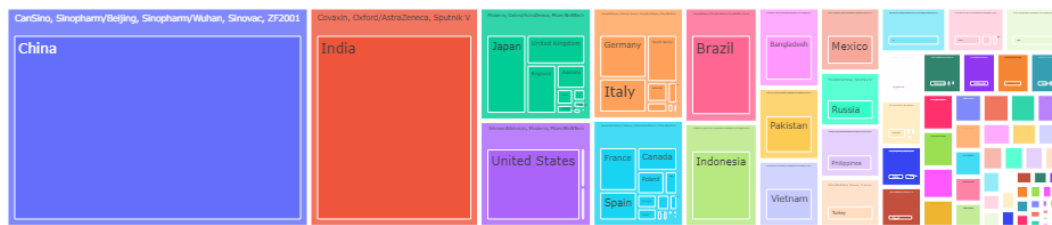
```

Vaccinations'])

fig.update_layout(title = dict(text = 'Total vaccinations per country,
grouped by Vaccine Scheme', x=0.5, y=0.95))
fig.update_traces(hovertemplate = 'Country: %
{customdata[0]}<br>Vaccine: %{customdata[1]}<br>Total Vaccinations: %
{customdata[2]}')
fig.show()

```

Total vaccinations per country, grouped by Vaccine Scheme



- From the above Treemap we can realise that a Bar and Pie Plot may often only show a part of the information that can be observed, whereas a Treemap can accurately show the share of a particular vaccine world-wide, the countries that are using the said vaccine and can even show comparisons between all the countries.
- As the Treemap shows so much information at a time, it can help one understand the data much more accurately.

```

fig = px.choropleth(country_data, locations = 'Country', color =
'Total Vaccinations',
                    locationmode = 'country names',
color_continuous_scale = 'rainbow',
                    hover_name = 'Country', projection = 'natural
earth')

fig.update_layout(title = dict(text = 'Total Vaccinations in every
Country', x=0.5, y=0.95),
                    geo = dict(showocean = True, oceancolor = "#7af8ff",
showland = True,
                    landcolor = "white", showlakes = False,
showframe = False))

fig.show()

```



- In the above visualisation, we can see the countries and the total vaccinations they have completed.

## Statistical Analysis

Which countries started vaccinations first?

```
# Find out which countries started vaccinations earliest
df0['date'] = pd.to_datetime(df0['date'], utc=True)
vacc_start = df0.loc[df0[df0.total_vaccinations >
0].groupby('country')['date'].idxmin()].sort_values('date')
vacc_start.head(5)
```

	country	iso_code	date
total_vaccinations \			
43117	Latvia	LVA	2020-12-04 00:00:00+00:00
1.0			
58523	Norway	NOR	2020-12-08 00:00:00+00:00
5.0			
20826	Denmark	DNK	2020-12-08 00:00:00+00:00
1.0			
82360	United States	USA	2020-12-13 00:00:00+00:00
30288.0			
13403	Canada	CAN	2020-12-14 00:00:00+00:00
5.0			

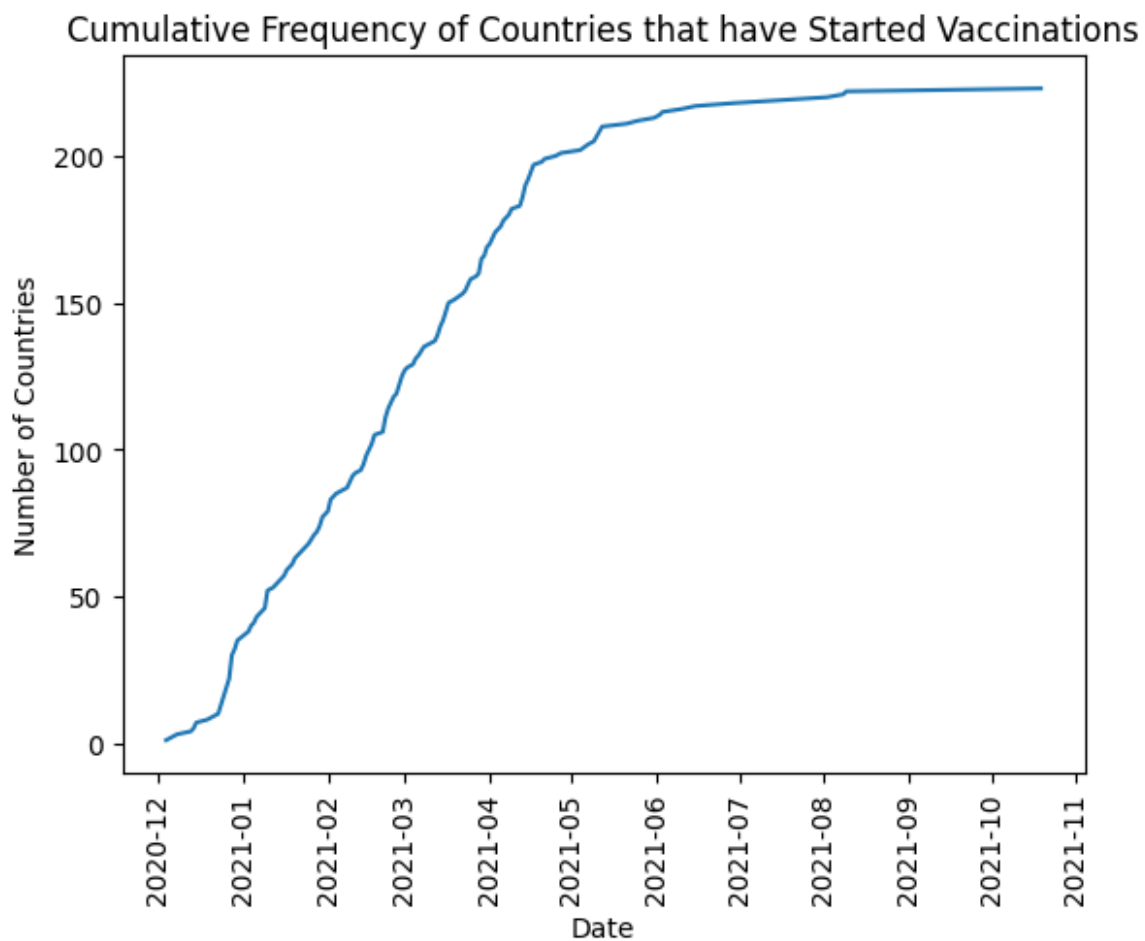
	people_vaccinated	people_fully_vaccinated
daily_vaccinations_raw \		
43117	1.0	NaN
NaN		
58523	5.0	NaN
5.0		
20826	1.0	NaN
NaN		
82360	25125.0	5897.0

NaN		
13403	5.0	NaN
NaN		
	daily_vaccinations	total_vaccinations_percent \
43117	NaN	0.00
58523	1.0	0.00
20826	NaN	0.00
82360	NaN	0.01
13403	NaN	0.00
	people_vaccinated_percent	people_fully_vaccinated_percent \
43117	0.00	NaN
58523	0.00	NaN
20826	0.00	NaN
82360	0.01	0.0
13403	0.00	NaN
	daily_vaccinations_per_million	\
43117	NaN	
58523	0.0	
20826	NaN	
82360	NaN	
13403	NaN	
	vaccines	\
43117	Johnson&Johnson, Moderna, Novavax, Pfizer/BioN...	
58523	Moderna, Pfizer/BioNTech	
20826	Johnson&Johnson, Moderna, Pfizer/BioNTech	
82360	Johnson&Johnson, Moderna, Pfizer/BioNTech	
13403	Johnson&Johnson, Moderna, Oxford/AstraZeneca, ...	
	source_name	\
43117	National Health Service	
58523	Norwegian Institute of Public Health	
20826	Statens Serum Institute	
82360	Centers for Disease Control and Prevention	
13403	Official data from provinces via covid19tracke...	
	source_website	
43117	<a href="https://data.gov.lv/dati/eng/dataset/covid19-v...">https://data.gov.lv/dati/eng/dataset/covid19-v...</a>	
58523	<a href="https://github.com/folkehelseinstituttet/surve...">https://github.com/folkehelseinstituttet/surve...</a>	
20826	<a href="https://covid19.ssi.dk/overvagningsdata/downlo...">https://covid19.ssi.dk/overvagningsdata/downlo...</a>	
82360	<a href="https://data.cdc.gov/Vaccinations/COVID-19-Vac...">https://data.cdc.gov/Vaccinations/COVID-19-Vac...</a>	
13403	<a href="https://covid19tracker.ca/vaccinationtracker.html">https://covid19tracker.ca/vaccinationtracker.html</a>	

How have the cumulative number of countries adopting covid-19 vaccinations evolved over time? How is this trend?

```
# Cumulative distribution of vaccination start dates
events = pd.Series(vacc_start.date.value_counts())
events.index = pd.to_datetime(events.index)
events.sort_index(inplace=True)

plt.plot(events.cumsum())
plt.xticks(rotation=90)
plt.title('Cumulative Frequency of Countries that have Started Vaccinations')
plt.xlabel('Date')
plt.ylabel('Number of Countries')
plt.show()
```



What are the top 20 countries in terms of total number of vaccines administered?

```
vacc_total = df0.loc[df0.groupby('country')
['total_vaccinations'].idxmax()].sort_values('total_vaccinations', ascending=False)
vacc_total = pd.concat((vacc_total, vacc_total["vaccines"].str.split(", ", expand = True)), axis=1)
```

```
# Plot out which countries have performed most vaccinations in
descending order
plt.figure(figsize=(10, 4))
plt.bar(vacc_total.country[0:20], vacc_total.total_vaccinations[0:20])
plt.xticks(rotation=90)
plt.title('Top 20 Countries by Total Vaccinations')
plt.xlabel('Country')
plt.ylabel('Total Vaccinations (x 10Million)')
plt.show()
```

