



COVID -19 VACCINATION

Lorem ipsum dolor sit amet, consectetuer adipiscing elit, sed diam nonummy nibh euismod tincidunt ut laoreet

shutterstock®

IMAGE ID: 1894960762
www.shutterstock.com

- *Creating a COVID-19 vaccine analysis program would typically involve data analysis and visualization. Below are examples of Python code snippets using popular libraries like NumPy, Pandas, and Matplotlib to get you started.*

1Data Retrieval:

You can fetch COVID-19 vaccine data from sources like the CDC or WHO, or use publicly available datasets. Here's an example using Pandas to read a CSV file:

- *import pandas as pd# Read vaccine data from a CSV filevaccine_data = pd.read_csv('vaccine_data.csv')*

2.Data Exploration:

You might want to explore the dataset by checking for missing values, data types, and basic statistics:

```
# Check for missing values  
missing_values =  
vaccine_data.isnull().sum()  
  
# Get basic statistics  
summary_stats = vaccine_data.describe()
```

- **3.Data Visualization:**
- *Visualize vaccine distribution or coverage using Matplotlib or another plotting library:*

```
import matplotlib.pyplot as plt
```

```
# Plot vaccine coverage over time
```

```
plt.figure(figsize=(10, 6))
```

```
plt.plot(vaccine_data['Date'], vaccine_data['Coverage'])
```

```
plt.xlabel('Date')plt.ylabel('Vaccine Coverage (%)')plt.title('COVID-19 Vaccine Coverage Over Time')
```

- **4.Data Analysis:**
- You can perform specific analyses, like calculating the average vaccine coverage or identifying regions with lower coverage:

Calculate average vaccine coverage

```
avg_coverage =  
vaccine_data['Coverage'].mean()
```

Find regions with lower coverage

```
low_coverage_regions =  
vaccine_data[vaccine_data['Coverage'] < 50]
```

- *5.Machine Learning (Optional):*
- *If you have enough data, you can build predictive models. Here's a simple linear regression example using scikit-learn:*

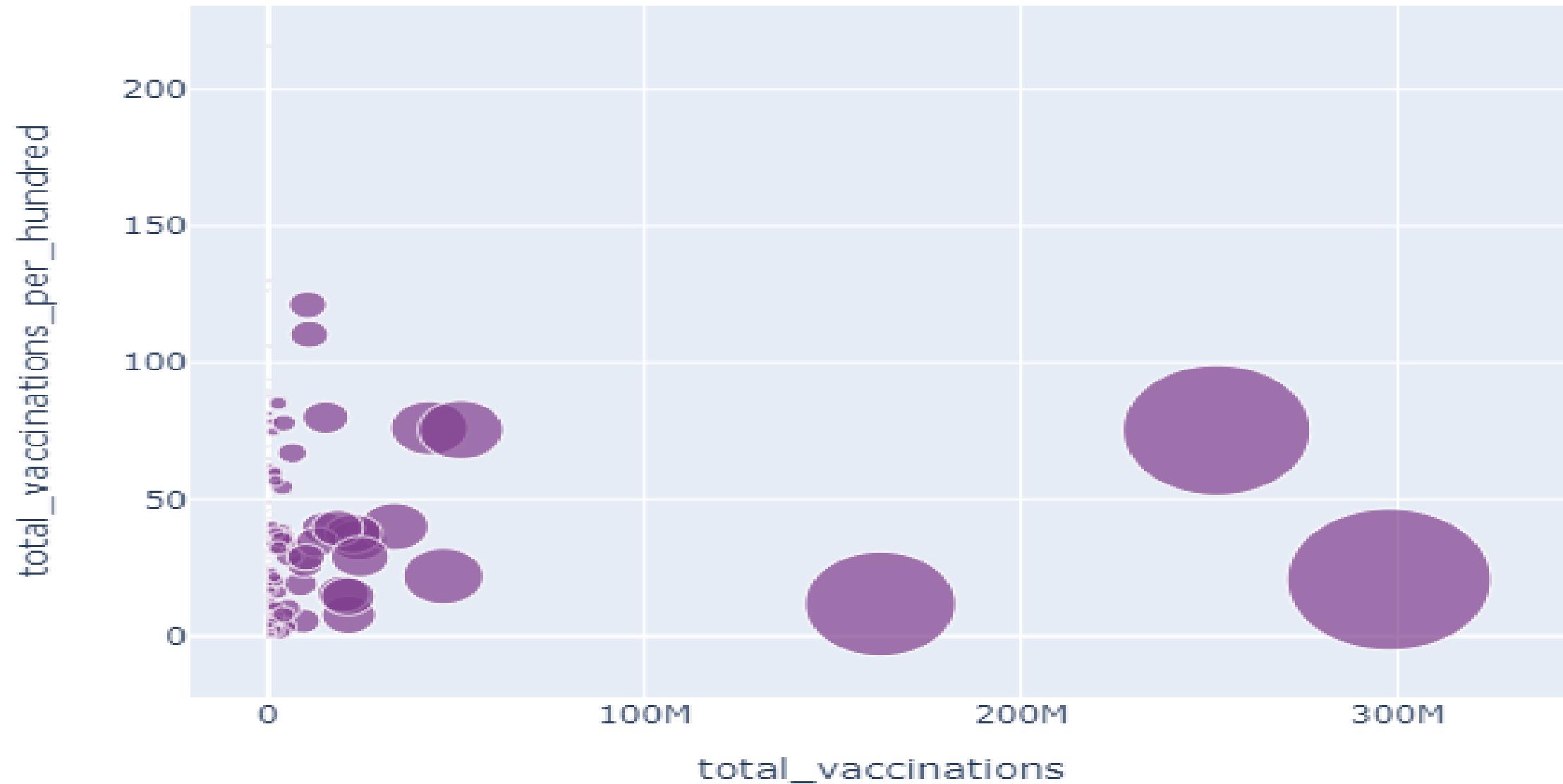
```
from sklearn.model_selection import  
train_test_split  
from sklearn.linear_model import  
LinearRegression
```

- *# Split the data into training and testing sets*
- *X = vaccine_data[['Population']]*
- *y = vaccine_data['Coverage']*
- *X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)*
- *# Create and train a linear regression model*
- *model = LinearRegression()model.fit(X_train, y_train)*

- *Relation between Total Vaccinations and Total Vaccinations per Hundred:*

```
fig = px.scatter(new_df,x =  
'total_vaccinations',y='total_vaccinations_per_hundred',  
size='total_vaccinations', hover_name = 'country',size_max =  
50, title="Total vs Total vaccinations per hundred grouped  
by Vaccines", color_discrete_sequence =  
px.colors.qualitative.Bold) fig.show()
```

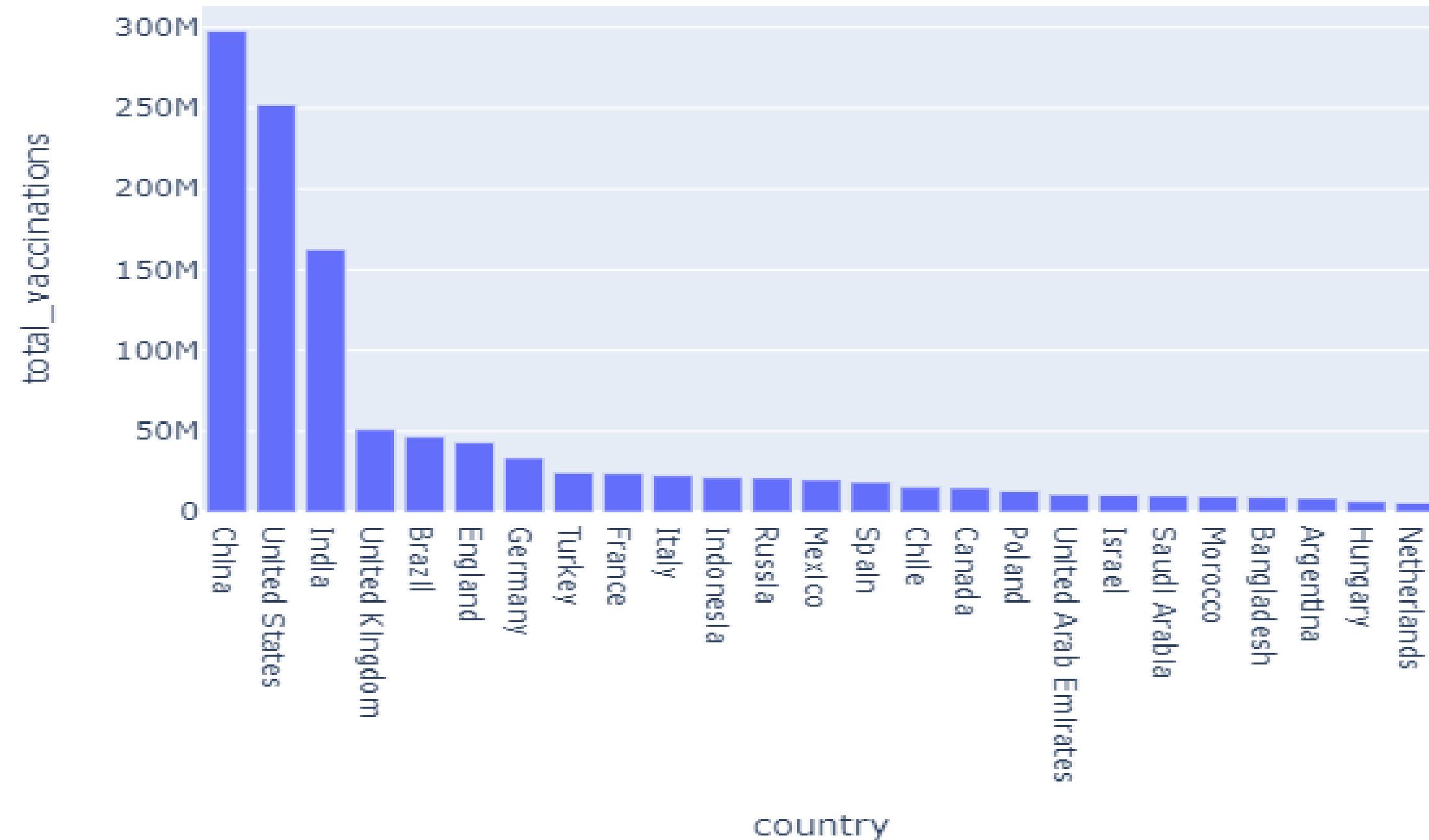
Total vs Total vaccinations per hundred grouped by Vaccines



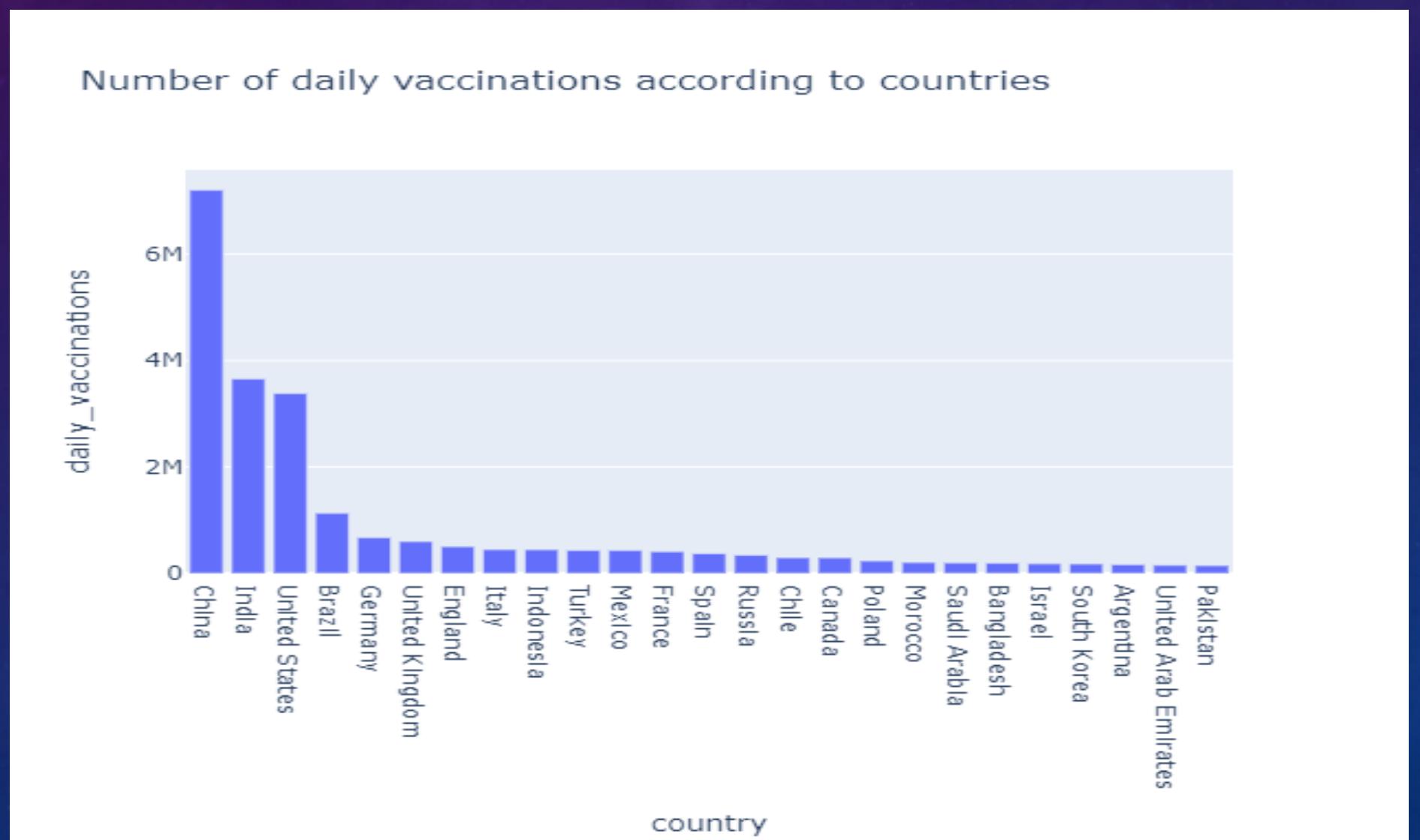
- *What is the number of total vaccinations & daily vaccinations according to countries?*

```
data =  
new_df[['country','total_vaccinations']].nlargest(2  
5,'total_vaccinations') fig = px.bar(data, x =  
'country',y = 'total_vaccinations',title="Number of  
total vaccinations according to countries",)  
fig.show()
```

Number of total vaccinations according to countries



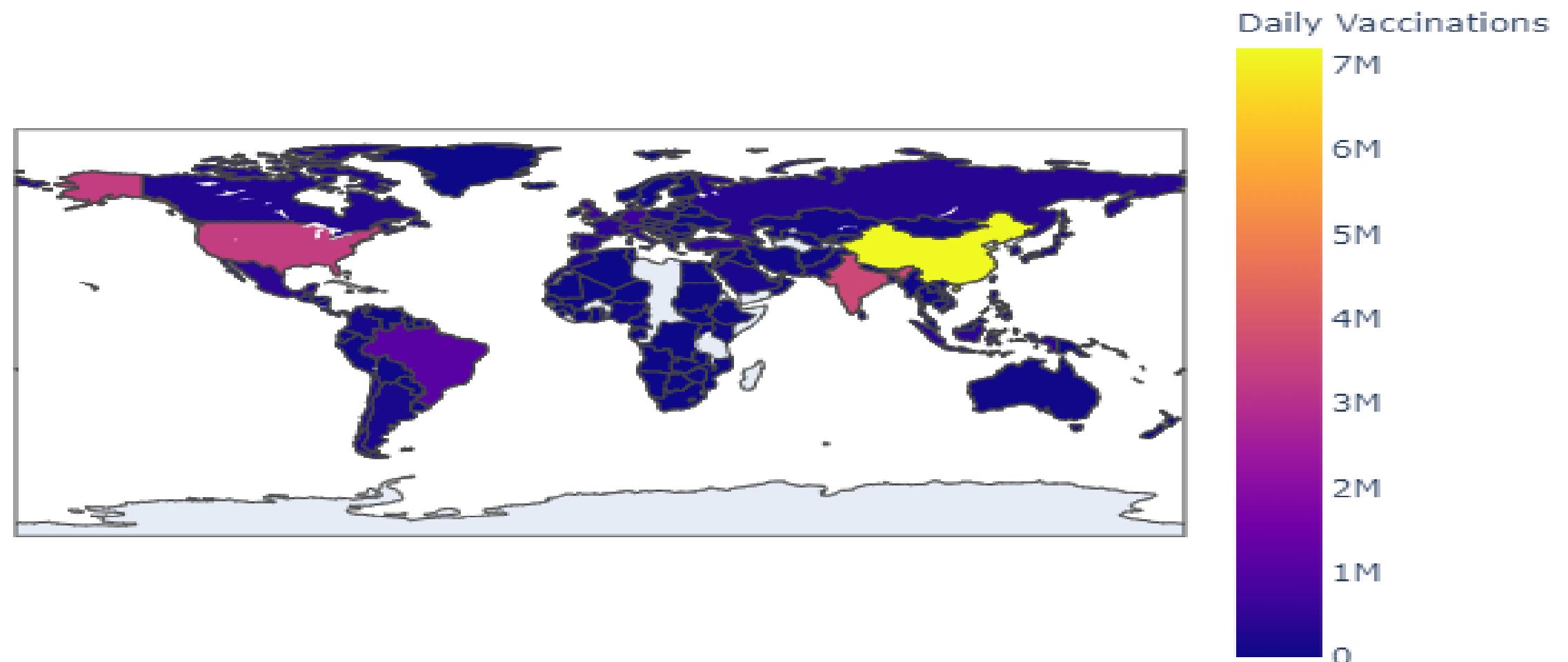
```
• data =  
new_df[['country','daily_vaccinations']].nlargest( 25,'daily_vacci  
nations') fig = px.bar(data, x = 'country',y =  
'daily_vaccinations',title="Number of daily vaccinations  
according to countries",) fig.show()
```



- *Daily Vaccinations per Countries:*

```
fig = go.Choropleth(locations =  
new_df["country"],locationmode = 'country names', z =  
new_df['daily_vaccinations'], text =  
new_df['country'],colorbar = dict(title= "Daily  
Vaccinations")) data = [fig] layout = go.Layout(title =  
'Daily Vaccinations per Countries') fig = dict(data =  
data,layout = layout) iplot(fig)
```

Daily Vaccinations per Countries



- *The conclusions from the available data up to that point indicated that COVID-19 vaccines:*

- 1*Are effective in preventing COVID-19 infection.*

- 2*Significantly reduce the risk of severe illness, hospitalization, and death.*

- 3. Contribute to herd immunity, slowing the spread of the virus.
- 2. Have demonstrated safety profiles, with side effects typically being mild and temporary.
- 3. May require booster shots for ongoing protection due to the emergence of new variants.