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import pandas as pd # Import pandas for data manipulation and analysis
import numpy as np # Import numpy for numerical operations
import seaborn as sns # Import seaborn for statistical data visualization
import matplotlib.pyplot as plt # Import matplotlib for plotting
import plotly.express as px # Import plotly for interactive visualizations
from sklearn.linear_model import LinearRegression # Import LinearRegression for predictive analysis

# Load datasets from online sources
url_cases = "https://covid.ourworldindata.org/data/owid-covid-data.csv"
url_vaccinations = "https://covid.ourworldindata.org/data/vaccinations/vaccinations.csv"
url_population = "https://raw.githubusercontent.com/datasets/population/master/data/population.csv"

cases_data = pd.read_csv(url_cases) # Load COVID-19 case data
vaccinations_data = pd.read_csv(url_vaccinations) # Load vaccination data
population_data = pd.read_csv(url_population) # Load population data

# Function to get the latest population of a given country
def get_population(country):
    pop = population_data[population_data['Country Name'] == country] # Filter population data for the country
    if not pop.empty:
        return pop.iloc[-1]['Value'] # Return the latest population value if found
    return None # Return None if no data found

# Function to analyze COVID-19 data
def analyze_covid_data(countries, start_date_str, end_date_str, metrics):
    combined_data = pd.DataFrame() # Initialize an empty DataFrame to store combined data

    # Data collection and merging for each country
    for country in countries:
        # Extract case and vaccination data for the country
        cases_country = cases_data[cases_data['location'] == country][['date', 'new_cases', 'new_deaths', 'total_cases', 'total_deaths']]
        vaccinations_country = vaccinations_data[vaccinations_data['location'] == country][['date', 'daily_vaccinations', 'total_vaccinations']]

        # Check if data is available for the country
        if cases_country.empty or vaccinations_country.empty:
            print(f"No data found for {country}.")
            continue # Skip to the next country if no data is found

        # Convert date columns to datetime format
        cases_country['date'] = pd.to_datetime(cases_country['date'])
        vaccinations_country['date'] = pd.to_datetime(vaccinations_country['date'])

        # Merge case and vaccination data on the date
        merged_data = pd.merge(cases_country, vaccinations_country, on='date', how='outer')
        merged_data.set_index('date', inplace=True) # Set date as the index

        # Filter merged data by the specified date range
        start_date = pd.to_datetime(start_date_str)
        end_date = pd.to_datetime(end_date_str)
        merged_data = merged_data[(merged_data.index >= start_date) & (merged_data.index <= end_date)]

        # Get the population for the country
        population = get_population(country)
        if population:
            # Calculate metrics per million and vaccination rates
            merged_data['cases_per_million'] = (merged_data['total_cases'] / population) * 1_000_000
            merged_data['deaths_per_million'] = (merged_data['total_deaths'] / population) * 1_000_000
            merged_data['vaccination_rate'] = (merged_data['total_vaccinations'] / population) * 100
            merged_data['recovery_rate'] = (merged_data['total_cases'] - merged_data['total_deaths']) / merged_data['total_cases'] * 100

        # Add a column for the country name
        merged_data['country'] = country
        combined_data = pd.concat([combined_data, merged_data]) # Concatenate data for each country

    # Check if combined data is empty
    if combined_data.empty:
        print("No data available for the specified countries and date range.")
        return

    # Descriptive Analysis
    print("\nDescriptive Analysis:")
    print(combined_data.describe()) # Display descriptive statistics

    # Diagnostic Analysis
    print("\nDiagnostic Analysis:")
    numeric_data = combined_data.select_dtypes(include=[np.number]) # Select numeric columns for analysis

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correlation_matrix = numeric_data.corr() # Calculate correlation matrix
print("Correlation matrix between metrics:")
print(correlation_matrix)

# Visualizations
plt.figure(figsize=(18, 45)) # Set figure size

# Heatmap for Correlation between Metrics
plt.subplot(len(metrics) + 10, 1, 1) # Create a subplot for the heatmap
sns.heatmap(correlation_matrix, annot=True, cmap="YlGnBu", fmt=".2f") # Draw the heatmap
plt.title("Correlation Heatmap of Metrics")
plt.tight_layout()

# Bar Chart for Average Daily New Cases
plt.subplot(len(metrics) + 10, 1, 2) # Create a subplot for bar chart
avg_daily_cases = combined_data.groupby('country')['new_cases'].mean().reset_index() # Calculate average daily cases
sns.barplot(data=avg_daily_cases, x='country', y='new_cases', palette='viridis') # Draw the bar chart
plt.title("Average Daily New Cases by Country")
plt.xlabel("Country")
plt.ylabel("Average Daily New Cases")
plt.xticks(rotation=45) # Rotate x-axis labels

# Pie Chart for Total Cases Distribution
plt.subplot(len(metrics) + 10, 1, 3) # Create a subplot for pie chart
total_cases_per_country = combined_data.groupby('country')['total_cases'].max().reset_index() # Get total cases per country
plt.pie(total_cases_per_country['total_cases'], labels=total_cases_per_country['country'], autopct='%1.1f%%', startangle=140) # Draw the
plt.title("Total Cases Distribution by Country")

# Line Chart for New Cases, New Deaths, and Daily Vaccinations
for i, metric in enumerate(['new_cases', 'new_deaths', 'daily_vaccinations'], 4):
    plt.subplot(len(metrics) + 10, 1, i) # Create a subplot for each metric
    for country in countries:
        country_data = combined_data[combined_data['country'] == country] # Filter data for the country
        if not country_data.empty and country_data[metric].notna().any(): # Check for non-empty data
            sns.lineplot(data=country_data, x=country_data.index, y=metric, label=country) # Draw line plot for the metric

    plt.title(f"{metric.replace('_', ' ').title()} Over Time") # Set title for the plot
    plt.xlabel("Date")
    plt.ylabel(metric.replace("_", " ").title()) # Set ylabel
    plt.legend() # Add legend

# Histogram for New Cases Distribution
plt.subplot(len(metrics) + 10, 1, 8) # Create a subplot for the histogram
sns.histplot(combined_data['new_cases'].dropna(), bins=30, kde=True) # Draw histogram with KDE
plt.title("Distribution of New Cases")
plt.xlabel("New Cases")
plt.ylabel("Frequency")

# Scatter Plot for Cases per Million vs Deaths per Million
plt.subplot(len(metrics) + 10, 1, 9) # Create a subplot for the scatter plot
valid_data = combined_data.dropna(subset=['cases_per_million', 'deaths_per_million']) # Drop rows with NaN values
if not valid_data.empty:
    sns.scatterplot(data=valid_data, x='cases_per_million', y='deaths_per_million', hue='country', palette='deep') # Draw scatter plot
    plt.title("Cases per Million vs Deaths per Million by Country")
    plt.xlabel("Cases per Million")
    plt.ylabel("Deaths per Million")

plt.tight_layout() # Adjust layout
plt.show() # Display plots

# Plotly General Chart for all selected metrics
for country in countries:
    country_data = combined_data[combined_data['country'] == country].reset_index() # Reset index for plotly
    fig = px.line(country_data, x='date', y=metrics,
                  title=f'COVID-19 Metrics Over Time for {country}',
                  labels={'value': 'Count', 'variable': 'Metrics'},
                  template='plotly_dark') # Create a plotly line chart
    fig.show() # Show plotly figure

# Predictive Analysis
print("\nPredictive Analysis:")
for country in countries:
    country_data = combined_data[combined_data['country'] == country] # Filter data for the country
    if not country_data.empty:
        # Preparing data for linear regression
        country_data = country_data[['new_cases']].reset_index() # Reset index for linear regression
        country_data['date_ordinal'] = country_data['date'].map(pd.Timestamp.toordinal) # Convert dates to ordinal format

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# Fit a linear regression model
X = country_data[['date_ordinal']] # Independent variable
y = country_data['new_cases'].fillna(0) # Dependent variable, filling NaNs with 0
model = LinearRegression() # Create linear regression model
model.fit(X, y) # Fit the model

# Making predictions
future_dates = pd.date_range(start=country_data['date'].max() + pd.Timedelta(days=1), periods=30) # Generate future dates
future_dates_ordinal = future_dates.map(pd.Timestamp.toordinal).values.reshape(-1, 1) # Convert future dates to ordinal
predictions = model.predict(future_dates_ordinal) # Make predictions

# Visualizing predictions
plt.figure(figsize=(10, 6)) # Set figure size for predictions
plt.plot(country_data['date'], country_data['new_cases'], label='Historical New Cases', color='blue') # Plot historical data
plt.plot(future_dates, predictions, label='Predicted New Cases', color='orange') # Plot predictions
plt.title(f'Predicted New Cases for {country}') # Set title
plt.xlabel('Date')
plt.ylabel('New Cases')
plt.legend() # Add legend
plt.xticks(rotation=45) # Rotate x-axis labels
plt.show() # Display prediction plot

# Prescriptive Analysis
print("\nPrescriptive Analysis:")
# Simple rules or recommendations based on the latest data
for country in countries:
    country_data = combined_data[combined_data['country'] == country] # Filter data for the country
    if not country_data.empty:
        latest_data = country_data.iloc[-1] # Get the latest data for the country
        print(f'Recommendations for {country}:')
        # Check vaccination and recovery rates to provide recommendations
        if latest_data['vaccination_rate'] < 70:
            print("- Increase vaccination campaigns.")
        if latest_data['recovery_rate'] < 90:
            print("- Focus on improving healthcare infrastructure and recovery protocols.")

# User input for analysis
print("COVID-19 Data Analysis")
user_start_date = input("Enter the start date in YYYY-MM-DD format (e.g., '2021-01-01'): ")
user_end_date = input("Enter the end date in YYYY-MM-DD format (e.g., '2021-12-31'): ")

user_countries = input("Enter the country names (comma-separated, e.g., 'India, United States, Brazil'): ")
countries_list = [country.strip() for country in user_countries.split(',')] # Create a list of countries

# Define available metrics for visualization
available_metrics = ['new_cases', 'new_deaths', 'total_cases', 'total_deaths', 'daily_vaccinations', 'total_vaccinations', 'recovery_rate', '']
print("\nAvailable metrics to visualize:", available_metrics)

user_metrics = [metric.strip() for metric in input("Enter metrics to visualize (comma-separated): ").split(',')] # Get user-defined metrics
user_metrics = [metric for metric in user_metrics if metric in available_metrics] # Filter metrics based on availability

# Call the analysis function with validated input
analyze_covid_data(countries_list, user_start_date, user_end_date, user_metrics)

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COVID-19 Data Analysis

Enter the start date in YYYY-MM-DD format (e.g., '2021-01-01'): 2019-02-02

Enter the end date in YYYY-MM-DD format (e.g., '2021-12-31'): 2021-02-02

Enter the country names (comma-separated, e.g., 'India, United States, Brazil'): India, United States, Brazil

Available metrics to visualize: ['new_cases', 'new_deaths', 'total_cases', 'total_deaths', 'daily_vaccinations', 'total_vaccinations', 'cases_per_million', 'deaths_per_million', 'vaccination_rate', 'recovery_rate']
 Enter metrics to visualize (comma-separated): new_cases, total_cases

Descriptive Analysis:

	new_cases	new_deaths	total_cases	total_deaths
count	1.185000e+03	1185.000000	1.185000e+03	1185.000000
mean	3.858880e+04	699.631224	4.145087e+06	96044.806751
std	1.595090e+05	2533.500104	5.106481e+06	98454.431616
min	0.000000e+00	0.000000	0.000000e+00	0.000000
25%	0.000000e+00	0.000000	2.649600e+04	824.000000
50%	0.000000e+00	0.000000	2.153010e+06	79827.000000
75%	0.000000e+00	0.000000	6.796322e+06	153214.000000
max	1.667151e+06	23312.000000	2.586303e+07	452123.000000

	daily_vaccinations	total_vaccinations	cases_per_million
count	8.500000e+01	8.800000e+01	1185.000000
mean	5.098592e+05	8.973659e+06	11360.212604
std	4.568875e+05	1.159965e+07	15836.276891
min	9.970000e+02	0.000000e+00	0.000000
25%	1.822300e+05	1.045193e+06	42.253346
50%	2.968930e+05	2.972606e+06	4819.349614
75%	9.376920e+05	1.338541e+07	18890.756739
max	1.493026e+06	4.119782e+07	77925.641533

	deaths_per_million	vaccination_rate	recovery_rate
count	1185.000000	88.000000	1080.000000
mean	293.418975	2.641500	97.202970
std	347.493128	3.534550	1.640581
min	0.000000	0.000000	93.074818
25%	1.126867	0.165739	96.650020
50%	99.591930	0.881718	97.257887
75%	583.512051	4.033041	98.283198
max	1362.252247	12.412954	100.000000

Diagnostic Analysis:

Correlation matrix between metrics:

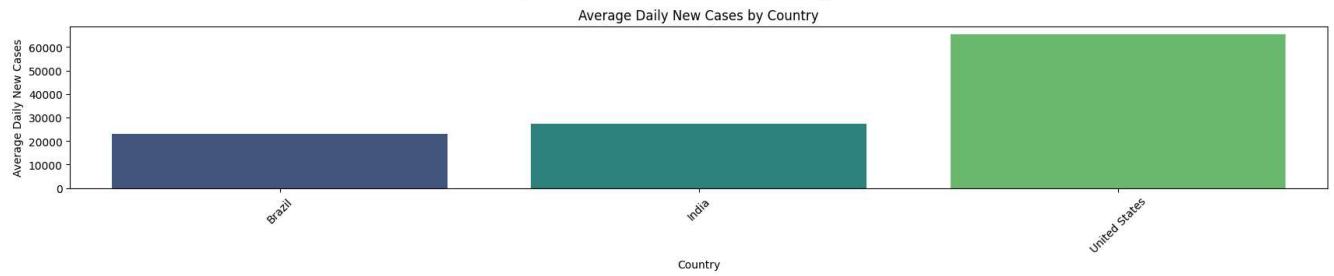
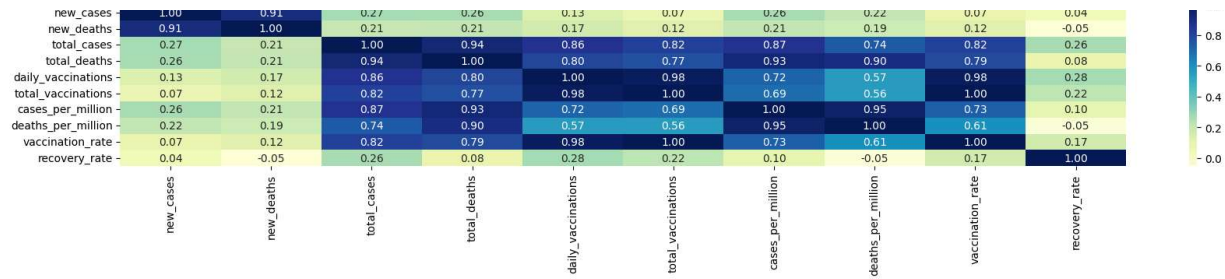
	new_cases	new_deaths	total_cases	total_deaths
new_cases	1.000000	0.907235	0.269764	0.260520
new_deaths	0.907235	1.000000	0.207278	0.214541
total_cases	0.269764	0.207278	1.000000	0.936397
total_deaths	0.260520	0.214541	0.936397	1.000000
daily_vaccinations	0.131788	0.169131	0.863561	0.804891
total_vaccinations	0.068279	0.117451	0.816996	0.770008
cases_per_million	0.256876	0.213810	0.867446	0.927979
deaths_per_million	0.217127	0.193110	0.744785	0.901732
vaccination_rate	0.074072	0.123565	0.821496	0.791870
recovery_rate	0.040855	-0.051620	0.257356	0.082083

	daily_vaccinations	total_vaccinations	cases_per_million
new_cases	0.131788	0.068279	0.256876
new_deaths	0.169131	0.117451	0.213810
total_cases	0.863561	0.816996	0.867446
total_deaths	0.804891	0.770008	0.927979
daily_vaccinations	1.000000	0.979255	0.715656
total_vaccinations	0.979255	1.000000	0.690603
cases_per_million	0.715656	0.690603	1.000000
deaths_per_million	0.568331	0.559017	0.953064
vaccination_rate	0.976248	0.997241	0.726154
recovery_rate	0.275938	0.222542	0.100733

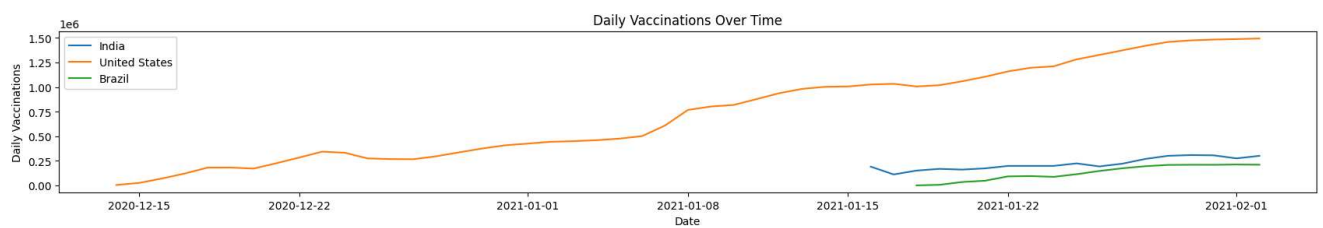
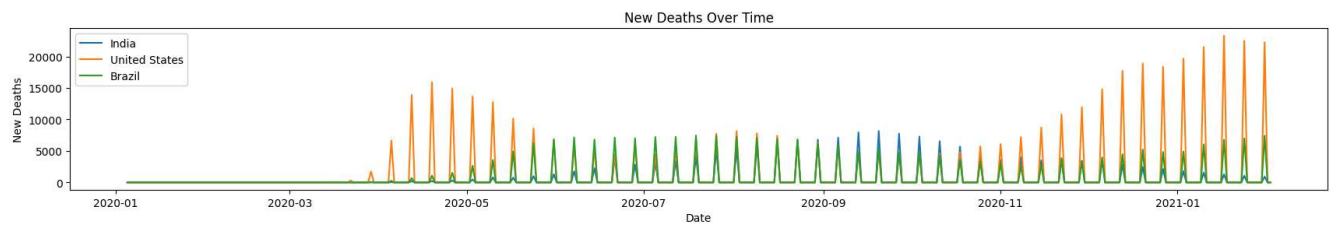
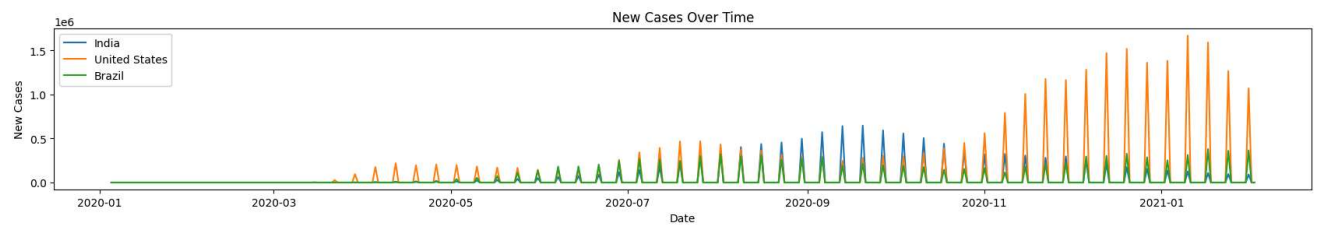
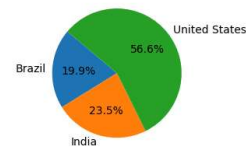
	deaths_per_million	vaccination_rate	recovery_rate
new_cases	0.217127	0.074072	0.040855
new_deaths	0.193110	0.123565	-0.051620
total_cases	0.744785	0.821496	0.257356
total_deaths	0.901732	0.791870	0.082083
daily_vaccinations	0.568331	0.976248	0.275938
total_vaccinations	0.559017	0.997241	0.222542
cases_per_million	0.953064	0.726154	0.100733
deaths_per_million	1.000000	0.605576	-0.047489
vaccination_rate	0.605576	1.000000	0.170209
recovery_rate	-0.047489	0.170209	1.000000

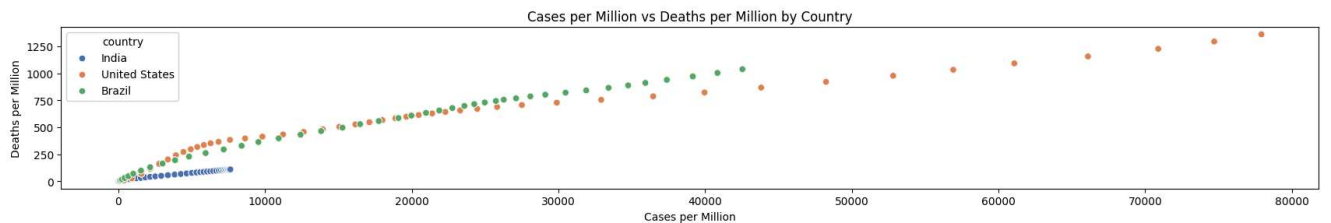
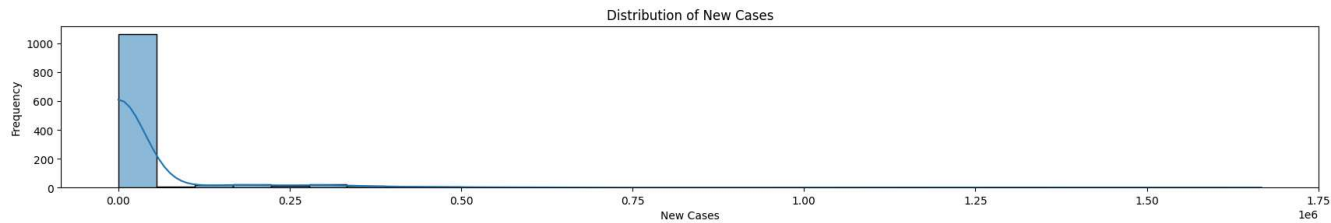
<ipython-input-3-bd08e9fe0737>:93: FutureWarning:

Passing `palette` without assigning `hue` is deprecated and will be removed in v0.14.0. Assign the `x` variable to `hue` and set `legend`

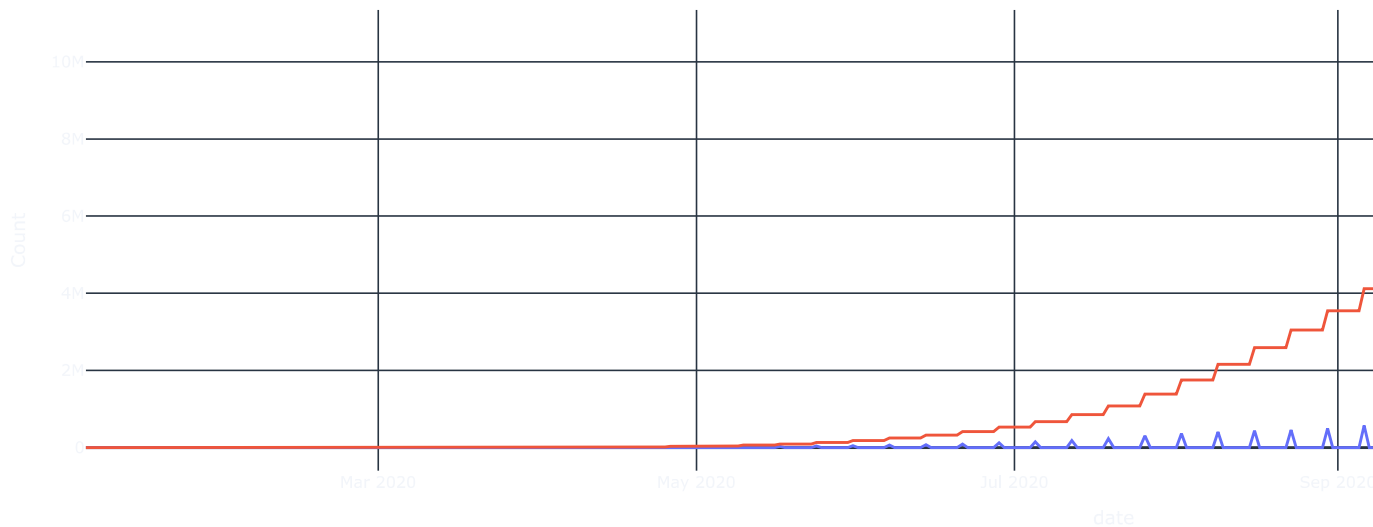


Total Cases Distribution by Country

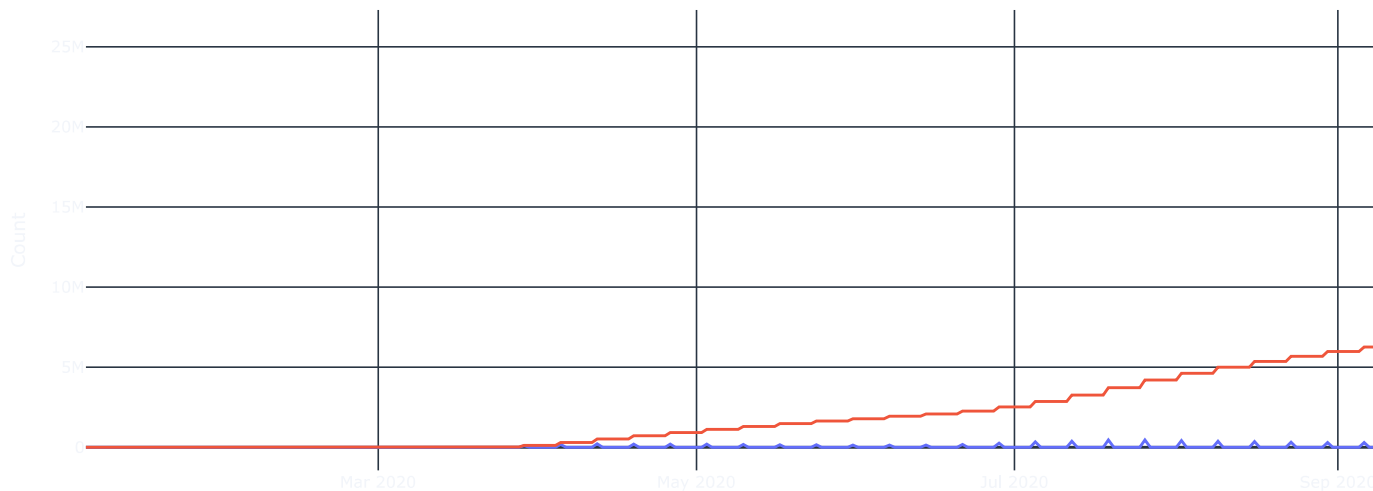




COVID-19 Metrics Over Time for India

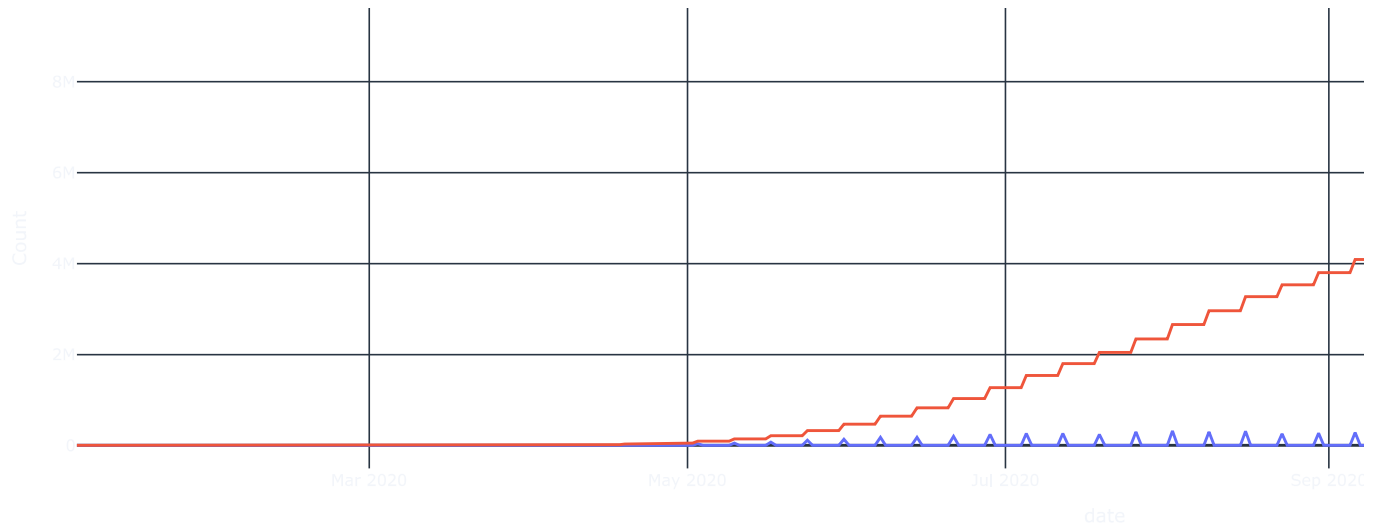


COVID-19 Metrics Over Time for United States



date

COVID-19 Metrics Over Time for Brazil

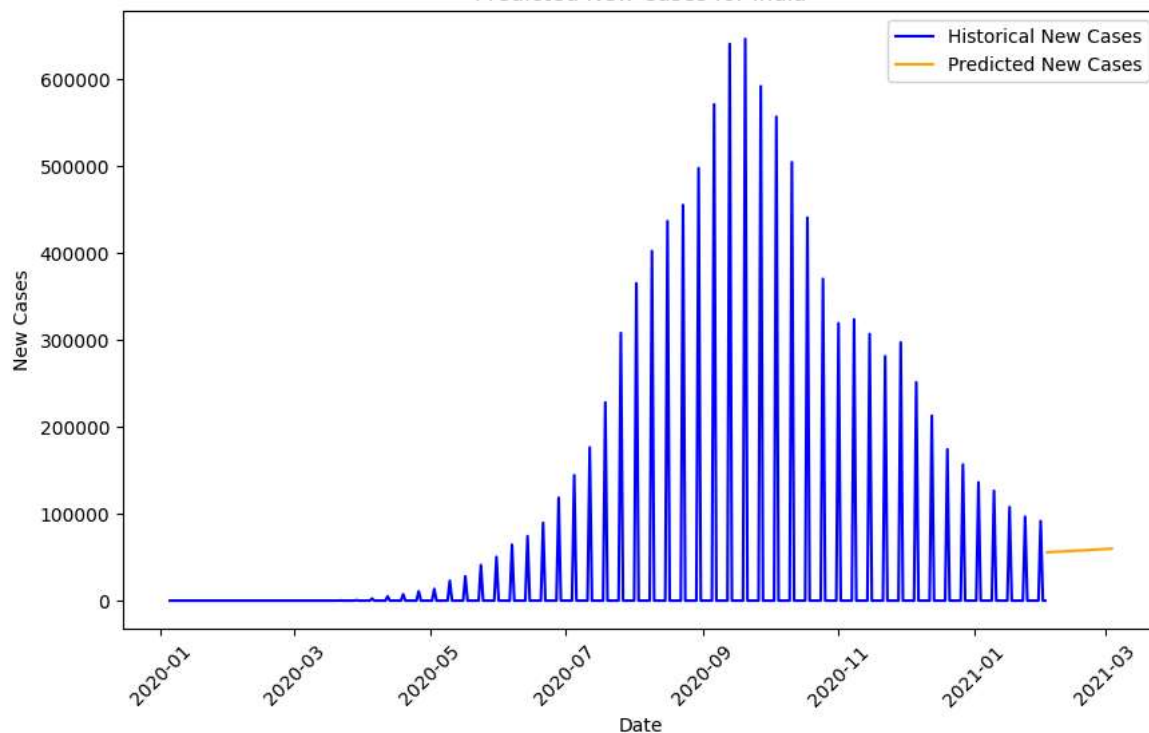


Predictive Analysis:

/usr/local/lib/python3.10/dist-packages/sklearn/base.py:493: UserWarning:

X does not have valid feature names, but LinearRegression was fitted with feature names

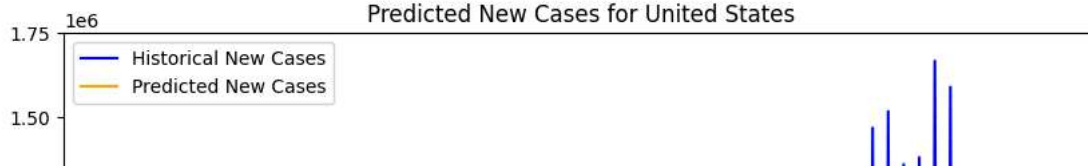
Predicted New Cases for India

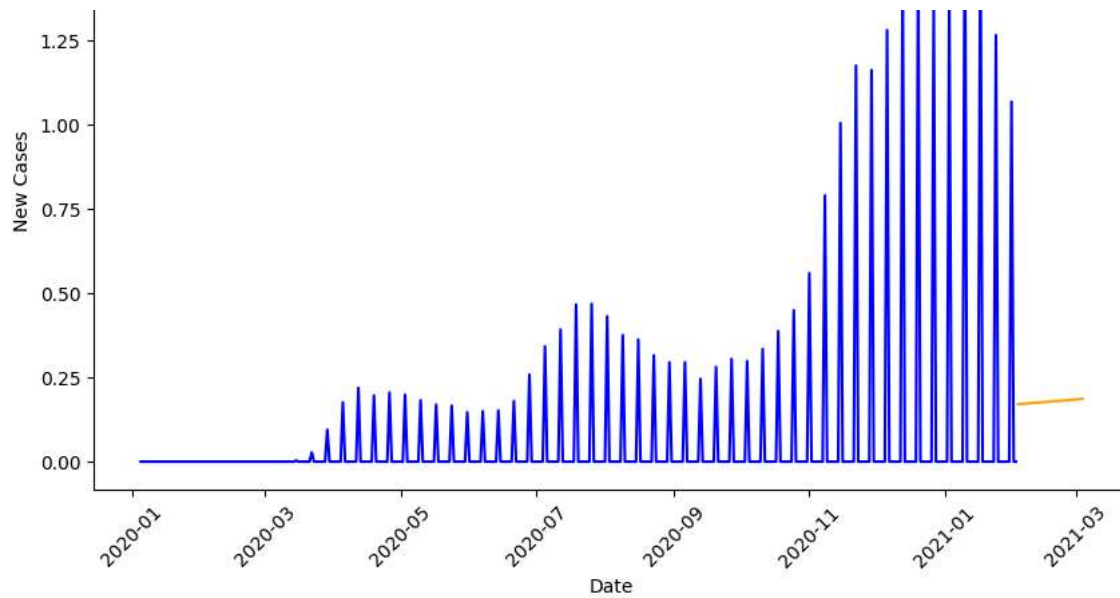


/usr/local/lib/python3.10/dist-packages/sklearn/base.py:493: UserWarning:

X does not have valid feature names, but LinearRegression was fitted with feature names

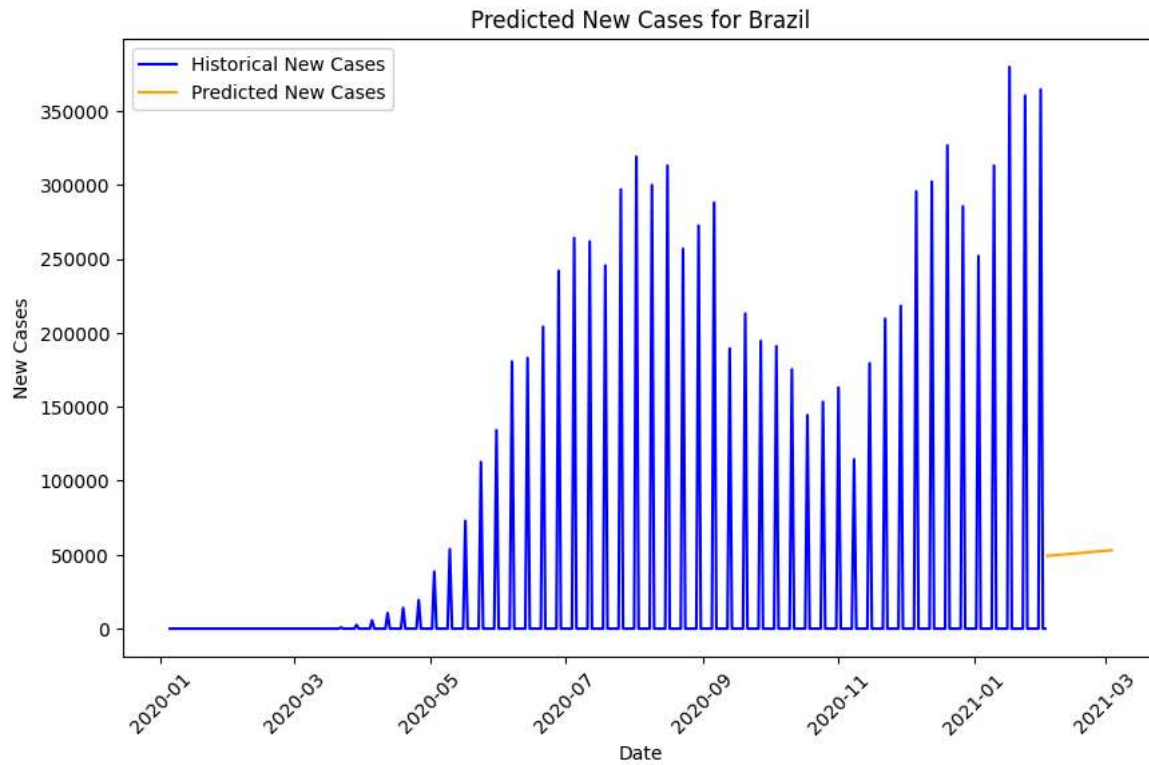
Predicted New Cases for United States





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/usr/local/lib/python3.10/dist-packages/sklearn/base.py:493: UserWarning:
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```
X does not have valid feature names, but LinearRegression was fitted with feature names
```



Prescriptive Analysis:

Recommendations for India:

- Increase vaccination campaigns.

Recommendations for United States:

- Increase vaccination campaigns.

Recommendations for Brazil:

- Increase vaccination campaigns.

