

COVID-19 CASE ANALYSIS – PROJECT

Data Analytics with Cognos (DAC)

Phase 3 – Development Part I

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COVID-19 Case - Data Analysis Report

Executive Summary

This report provides an analysis of the daily death and recovery data for COVID-19 in Germany, France, and Italy. The analysis covers a specific time frame and is accompanied by visual representations.

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Introduction

The COVID-19 pandemic has had a significant impact on countries worldwide. In this report, we focus on Germany, France, and Italy, presenting an analysis of daily death and recovery data to gain insights into the progression of the pandemic.

Data Collection and Sources

The data used in this analysis was collected from reliable sources, such as government health agencies and international health organizations.

Data Preprocessing

Prior to analysis, the data underwent preprocessing, including cleaning, handling missing values, and transforming data formats. This ensured the accuracy and reliability of our analysis.

Data Analysis:

Daily Deaths:

We analyzed the daily death data to:

- Identify peak periods of fatalities.
- Assess the impact of government interventions.
- Determine trends in mortality rates.

Daily Recoveries:

We analyzed the daily recovery data to:

- Understand the rate of recovery.
- Identify patterns in recoveries.
- Assess the effectiveness of healthcare systems.

Key Findings

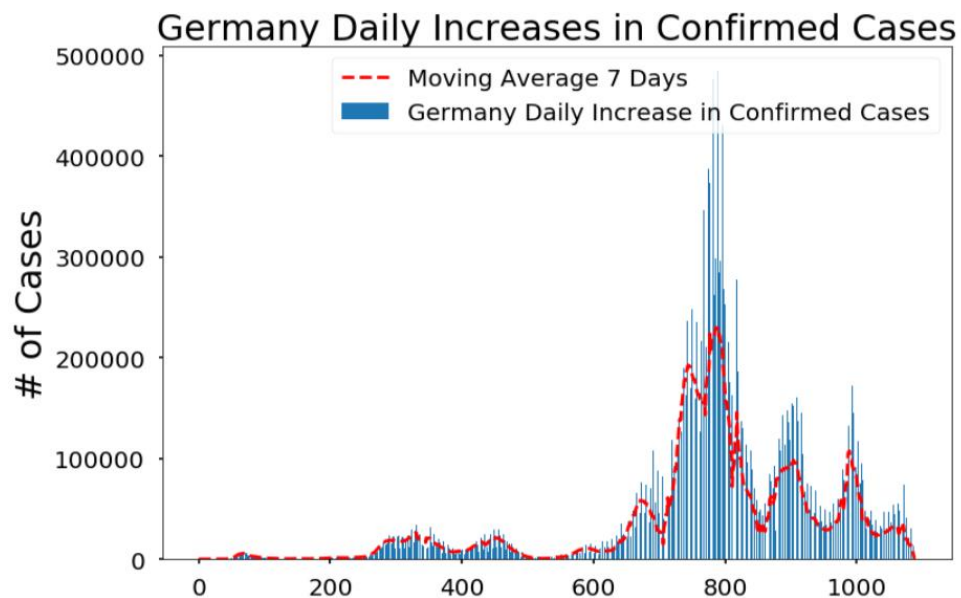
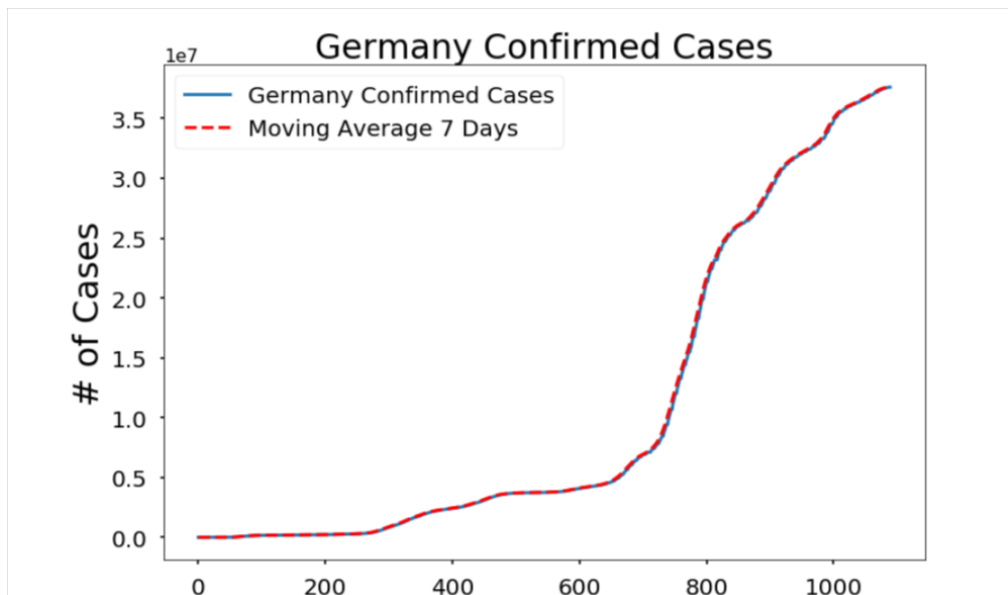
- **Germany:** Key findings for Germany's daily death and recovery data.

France: Key findings for **France's** daily death and recovery data.

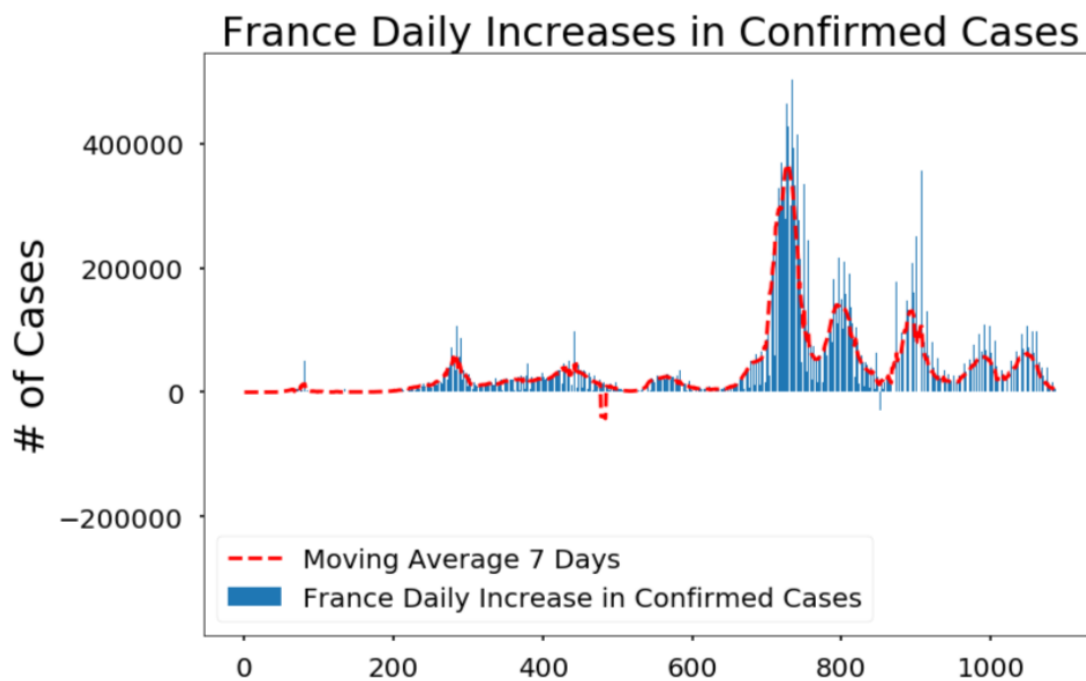
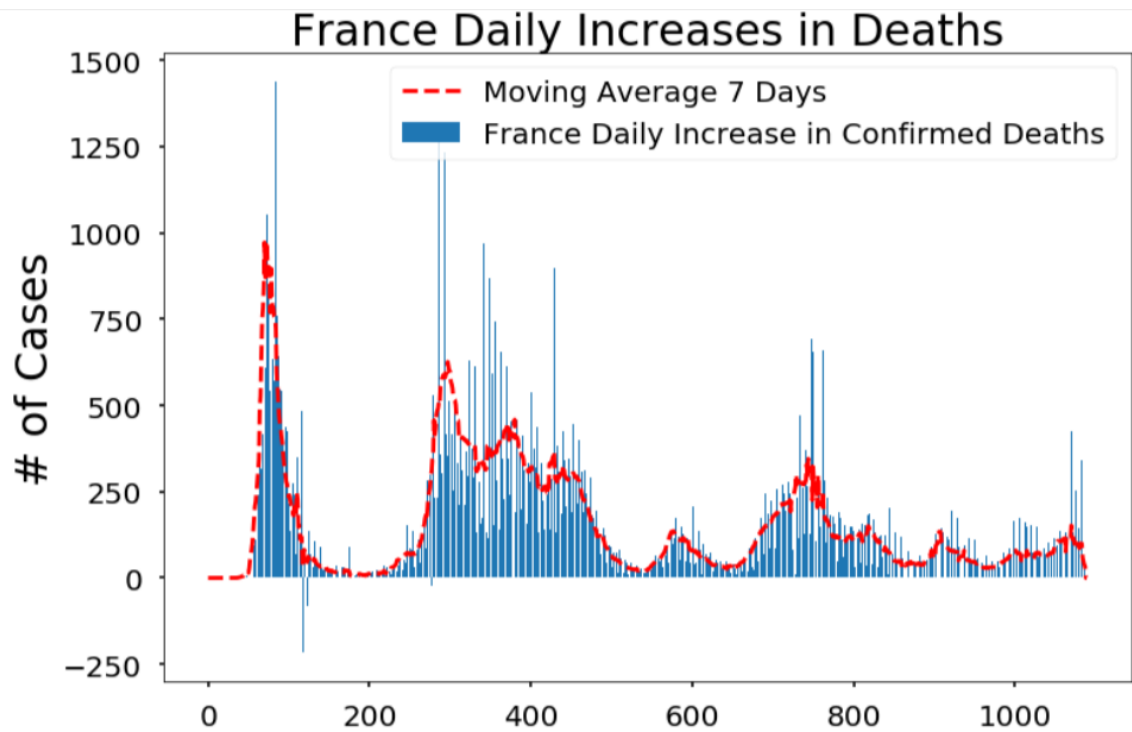
- **Italy:** Key findings for Italy's daily death and recovery data.

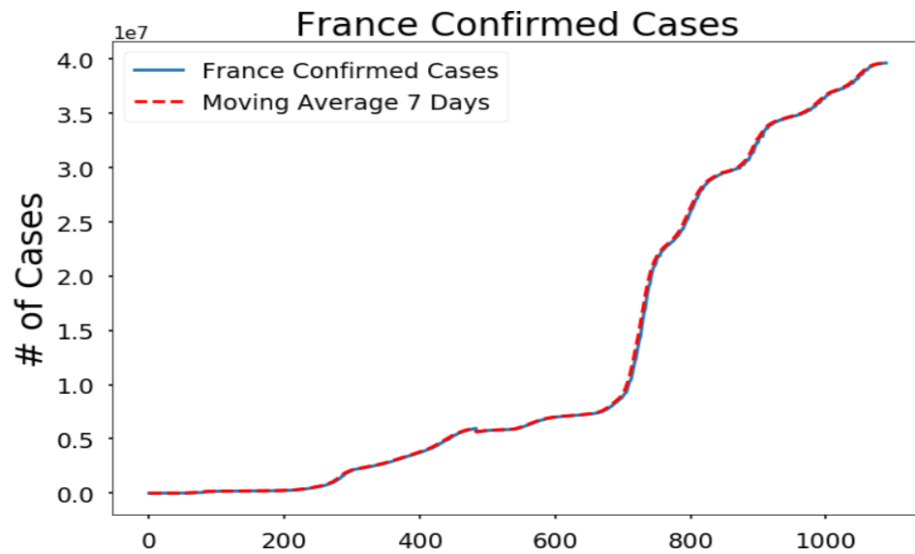
VISUALIZATIONS

Germany:

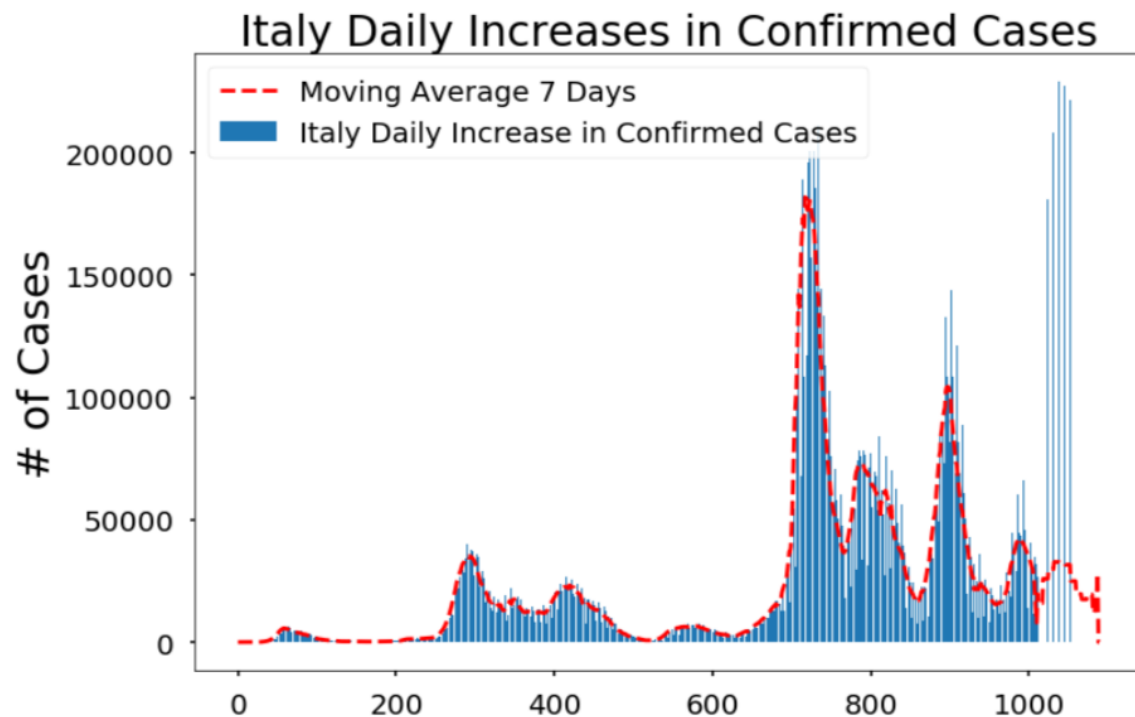


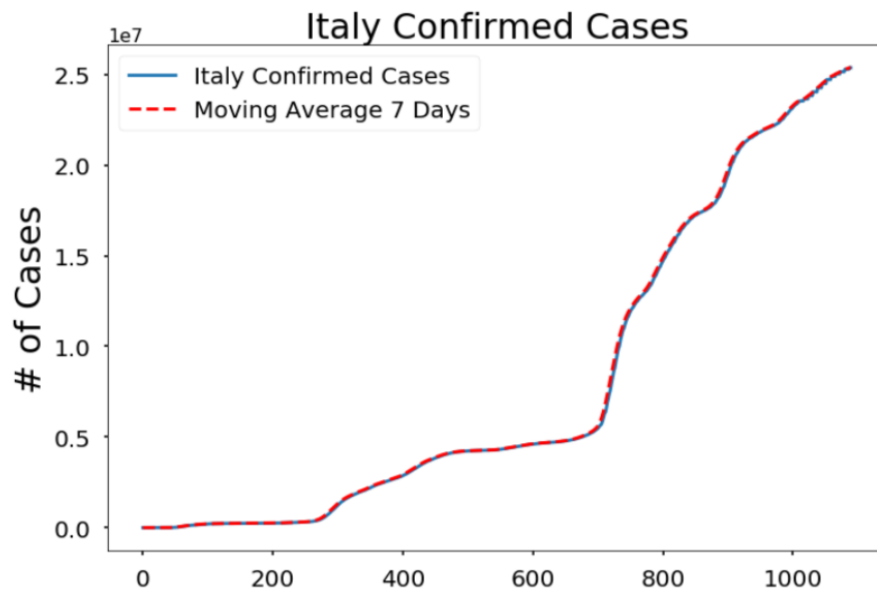
France:



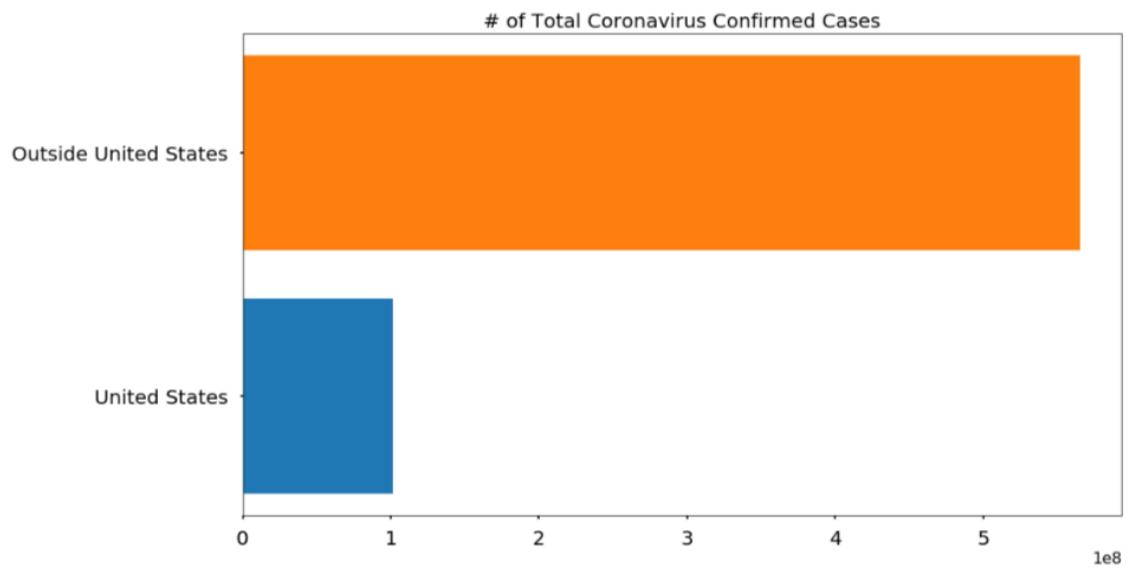


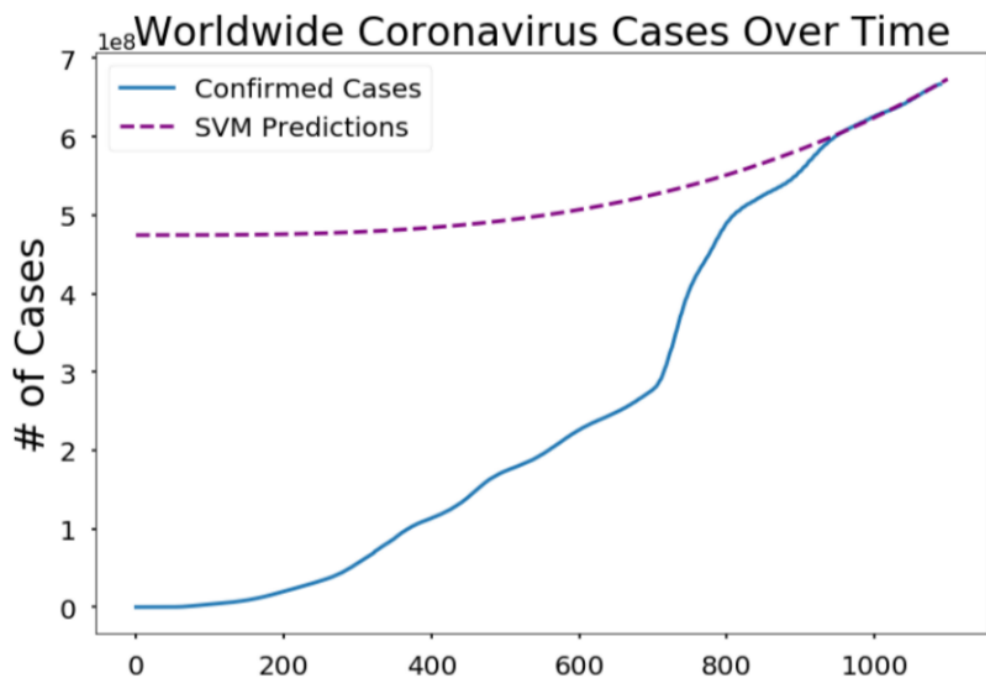
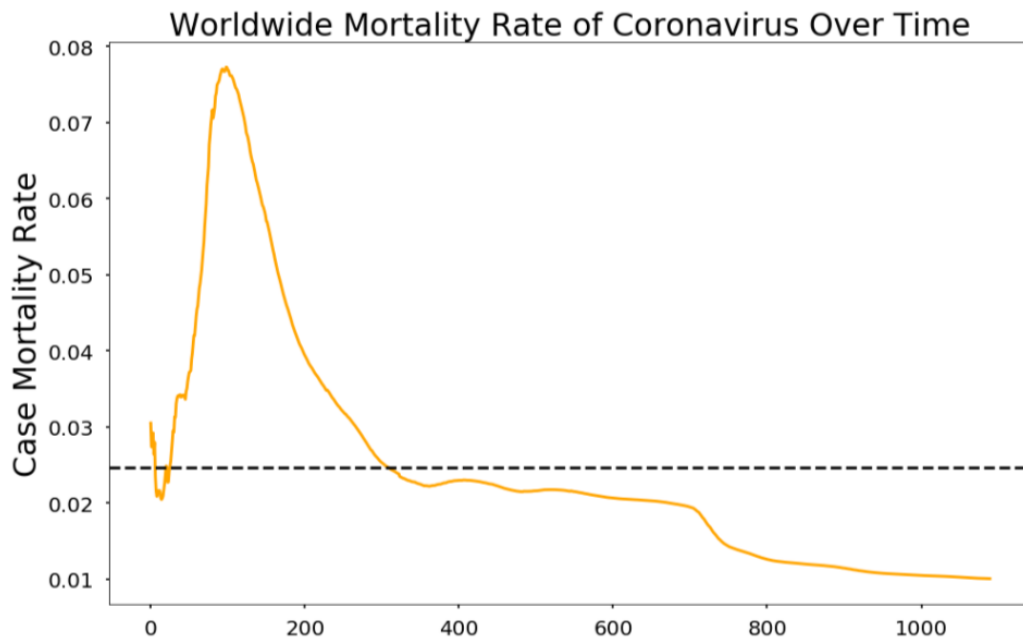
Italy

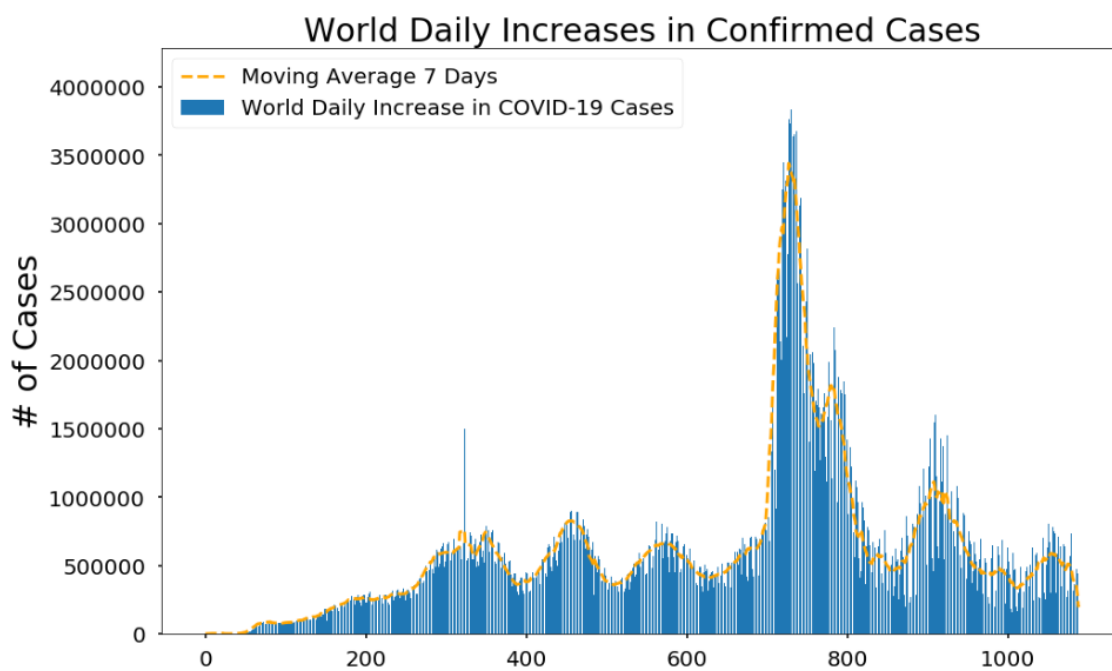
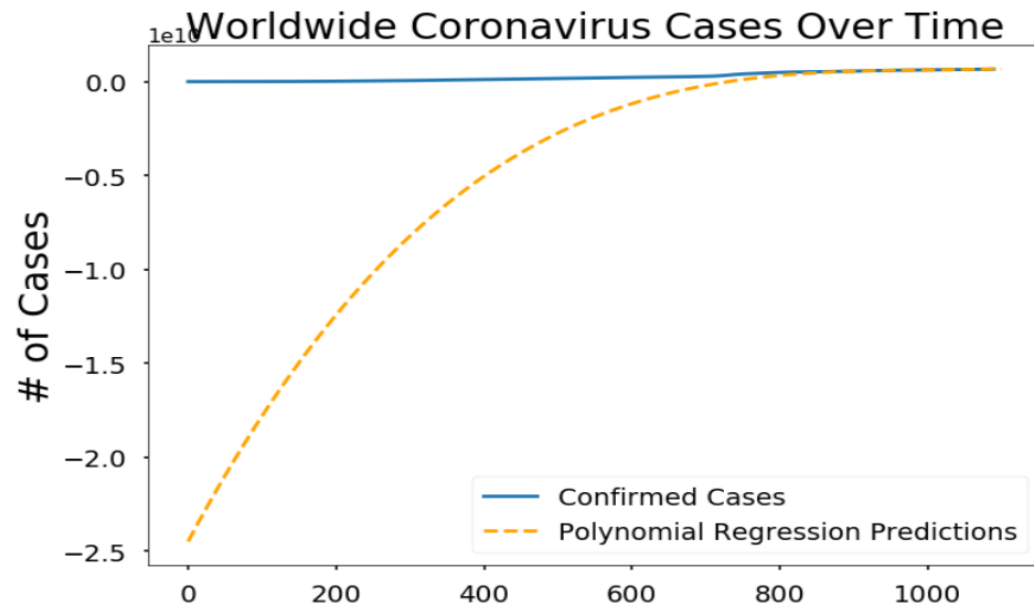


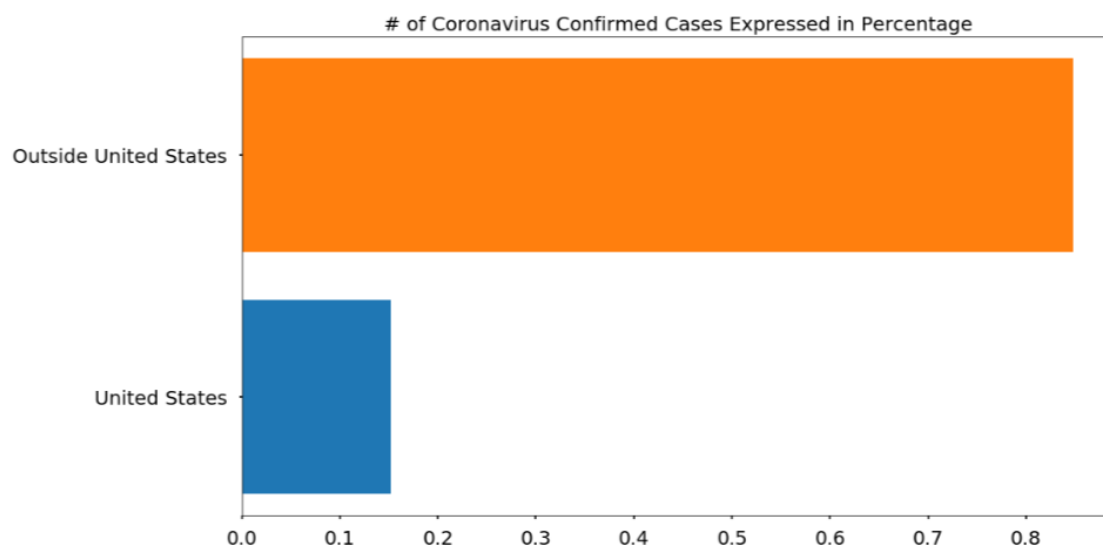
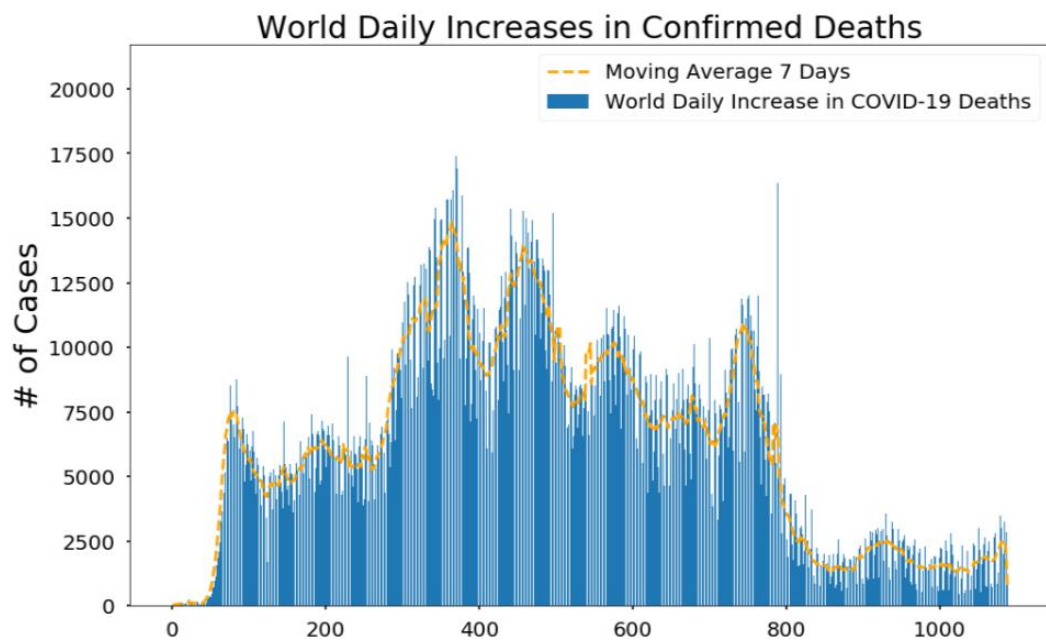


World trend:









PROGRAM:

```
import numpy as np

import matplotlib.pyplot as plt

import pandas as pd

from sklearn.linear_model import LinearRegression, BayesianRidge
from sklearn.model_selection import RandomizedSearchCV, train_test_split
from sklearn.preprocessing import PolynomialFeatures
from sklearn.svm import SVR
from sklearn.metrics import mean_squared_error, mean_absolute_error
import datetime


# Import and preprocess COVID-19 data
confirmed_df = pd.read_csv('confirmed_data_url')
deaths_df = pd.read_csv('deaths_data_url')
latest_data = pd.read_csv('latest_data_url')
us_medical_data = pd.read_csv('us_medical_data_url')


# Extract relevant columns
confirmed_cols = confirmed_df.keys()
deaths_cols = deaths_df.keys()
confirmed = confirmed_df.loc[:, confirmed_cols[4]:]
deaths = deaths_df.loc[:, deaths_cols[4]:]


# Analyze global COVID-19 data
```

```

num_dates = len(confirmed.keys())
ck = confirmed.keys()
dk = deaths.keys()
world_cases = []
total_deaths = []
mortality_rate = []

# Calculate total cases, deaths, and mortality rate
for i in range(num_dates):
    confirmed_sum = confirmed[ck[i]].sum()
    death_sum = deaths[dk[i]].sum()
    world_cases.append(confirmed_sum)
    total_deaths.append(death_sum)
    mortality_rate.append(death_sum / confirmed_sum)

# Define functions for data analysis and visualization
def daily_increase(data):
    # Calculate daily increase in data
    d = []
    for i in range(len(data)):
        if i == 0:
            d.append(data[0])
        else:
            d.append(data[i] - data[i-1])
    return d

def moving_average(data, window_size):

```

```
# Calculate moving average of data
moving_average = []
for i in range(len(data)):
    if i + window_size < len(data):
        moving_average.append(np.mean(data[i:i+window_size]))
    else:
        moving_average.append(np.mean(data[i:len(data)]))
return moving_average
```

```
# Specify window size for moving averages
window = 7
```

```
# Analyze and visualize COVID-19 cases and deaths
world_daily_increase = daily_increase(world_cases)
world_confirmed_avg = moving_average(world_cases, window)
world_daily_increase_avg = moving_average(world_daily_increase, window)
world_daily_death = daily_increase(total_deaths)
world_death_avg = moving_average(total_deaths, window)
world_daily_death_avg = moving_average(world_daily_death, window)
```

```
# Prepare data for regression modeling
days_since_1_22 = np.array([i for i in range(len(ck))]).reshape(-1, 1)
world_cases = np.array(world_cases).reshape(-1, 1)
total_deaths = np.array(total_deaths).reshape(-1, 1)
days_in_future = 10
future_forecast = np.array([i for i in range(len(ck) + days_in_future)]).reshape(-1, 1)
adjusted_dates = future_forecast[:-10]
```

```
start = '1/22/2020'

start_date = datetime.datetime.strptime(start, '%m/%d/%Y')

future_forecast_dates = []


# Generate future dates for forecasting
for i in range(len(future_forecast)):

    future_forecast_dates.append((start_date +
datetime.timedelta(days=i)).strftime('%m/%d/%Y'))


# Train and test data for regression models

days_to_skip = 830

X_train_confirmed, X_test_confirmed, y_train_confirmed, y_test_confirmed =
train_test_split(days_since_1_22[days_to_skip:], world_cases[days_to_skip:], test_size=0.10,
shuffle=False)


# Support Vector Regression (SVM) model for confirmed cases

svm_confirmed = SVR(shrinking=True, kernel='poly', gamma=0.01, epsilon=1, degree=3, C=0.1)

svm_confirmed.fit(X_train_confirmed, y_train_confirmed)

svm_pred = svm_confirmed.predict(future_forecast)


# Polynomial regression model for confirmed cases

poly = PolynomialFeatures(degree=3)

poly_X_train_confirmed = poly.fit_transform(X_train_confirmed)

poly_X_test_confirmed = poly.fit_transform(X_test_confirmed)

poly_future_forecast = poly.fit_transform(future_forecast)

linear_model = LinearRegression(normalize=True, fit_intercept=False)

linear_model.fit(poly_X_train_confirmed, y_train_confirmed)

test_linear_pred = linear_model.predict(poly_X_test_confirmed)
```

```
linear_pred = linear_model.predict(poly_future_forecast)
```

```
# Bayesian Ridge Polynomial Regression model for confirmed cases
```

```
tol = [1e-6, 1e-5, 1e-4, 1e-3, 1e-2]
```

```
alpha_1 = [1e-7, 1e-6, 1e-5, 1e-4, 1e-3]
```

```
alpha_2 = [1e-7, 1e-6, 1e-5, 1e-4, 1e-3]
```

```
lambda_1 = [1e-7, 1e-6, 1e-5, 1e-4, 1e-3]
```

```
lambda_2 = [1e-7, 1e-6, 1e-5, 1e-4, 1e-3]
```

```
normalize = [True, False]
```

```
bayesian_grid = {
```

```
    'tol': tol,
```

```
    'alpha_1': alpha_1,
```

```
    'alpha_2': alpha_2,
```

```
    'lambda_1': lambda_1,
```

```
    'lambda_2': lambda_2,
```

```
    'normalize': normalize
```

```
}
```

```
bayesian = BayesianRidge(fit_intercept=False)
```

```
bayesian_search = RandomizedSearchCV(bayesian, bayesian_grid,  
scoring='neg_mean_squared_error', cv=3, return_train_score=True, n_jobs=-1, n_iter=40,  
verbose=1)
```

```
bayesian_search.fit(bayesian_poly_X_train_confirmed, y_train_confirmed)
```

```
bayesian_confirmed = bayesian_search.best_estimator_
```

```
test_bayesian_pred = bayesian_confirmed.predict(bayesian_poly_X_test_confirmed)
```

```
bayesian_pred = bayesian_confirmed.predict(bayesian_poly_future_forecast)
```

```
# Visualize top 10 total COVID-19 cases for specific countries

countries = ['US', 'India', 'Brazil', 'France', 'Germany', 'United Kingdom', 'Italy', 'Korea, South',
'Russia', 'Turkey']

for country in countries:
    country_visualizations(country)

# Compare COVID-19 cases and deaths in selected countries

compare_countries = ['India', 'US', 'Brazil', 'Russia', 'United Kingdom', 'France']
graph_name = ['Coronavirus Confirmed Cases', 'Coronavirus Confirmed Deaths']

for num in range(2):
    plt.figure(figsize=(12, 8))
    for country in compare_countries:
        plt.plot(get_country_info(country)[num])
    plt.legend(compare_countries, prop={'size': 20})
    plt.xlabel('Days since 1/22/2020', size=30)
    plt.ylabel('# of Cases', size=30)
    plt.title(graph_name[num], size=30)
    plt.xticks(size=20)
    plt.yticks(size=20)
    plt.show()
```


Conclusion

The analysis of daily death and recovery data for Germany, France, and Italy provides valuable insights into the COVID-19 pandemic's impact in these countries. Our findings offer information that can guide policy decisions, healthcare resource allocation, and public health measures.