COVID-19 DATA ANALYSIS

SRI RAMACHANDRA FACULTY OF ENGINEERING AND TECHNOLOGY

INT 200-INTERNSHIP 1

PROJECT REPORT

Submitted by

RISHITHA THOKA-E0322026

In partial fulfilment for the award of the degree of

BACHELOR OF TECHNOLOGY

in

COMPUTER SCIENCE AND ENGINEERING

(Artificial Intelligence & Data Analytics)

Sri Ramachandra Faculty of Engineering and Technology

Sri Ramachandra Institute of Higher Education and Research, Porur, Chennai - 600516

JULY, 2023

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BONAFIDE CERTIFICATE

SRI RAMACHANDRA FACULTY OF ENGINEERING AND TECHNOLOGY

Certified that this project report "COVID-19 DATA ANALYTICS" is the bonafide record of work done by "RISHITHA THOKA-E0322026" who carried out the internship work under my supervision.

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SRI RAMACHANDRA FACULTY OF ENGINEERING AND TECHNOLOGY

ACKNOWLEDGEMENT

I express my sincere gratitude to our Programme Coordinator **Dr. Uma Satya Ranjan** for their support and for providing the required facilities for carrying out this study.

I wish to thank my faculty supervisor(s), **Dr. Ashokkumar.P**, Department of Artificial Intelligence and Machine Learning, Sri Ramachandra faculty of Engineering and Technology for extending help and encouragement throughout the project. Without his/her continuous guidance and persistent help, this project would not have been a success for me.

I am grateful to all the members of Sri Ramachandra Faculty of Engineering and Technology, my beloved parents and friends for extending the support, who helped us to overcome obstacles in the study.

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ABSTRACT

This project aims to analyse the data pertaining to the COVID-19 pandemic, focusing on various aspects such as infection rates, mortality rates, and the impact of different mitigation measures. By utilizing a comprehensive dataset, statistical techniques, and data visualization tools, this analysis seeks to uncover patterns, trends, and insights related to the spread and consequences of the virus. The findings of this project can contribute to a better understanding of the ongoing pandemic, assist in formulating effective strategies to control its transmission, and provide valuable insights for public health interventions and policy-making.

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LIST OF SYMBOLS

| S.NO | ABBREVIATIONS | EXPANSION |
|------|---------------|-------------------------------------|
| | | |
| 1 | PANDAS | PYTHON DATA |
| | | ANALYSIS LIBRARY |
| 2 | NUMPY | NUMERICAL PYTHON |
| 3 | SKLEARN | SCIKIT-LEARN |
| 4 | MATPLOTLIB | MATHEMATICAL PLOTTING LIBRARY |
| 5 | MS EXCEL | MICROSOFT EXCEL |

CHAPTER 1

INTRODUCTION

The COVID-19 pandemic has presented unprecedented challenges globally, requiring data-driven approaches to understand and combat the spread of the virus. In response, this COVID-19 data analytics project utilizes Python and its powerful data analysis capabilities to analyze and visualize the available COVID-19 datasets. By leveraging Python's libraries such as Pandas, NumPy, Matplotlib, and scikit-learn, the project aims to extract meaningful insights, track the progression of the pandemic, and develop predictive models. These insights and models will provide valuable information for decision-making, resource allocation, and public health responses, ultimately contributing to the global efforts in managing and controlling the impact of the COVID-19 pandemic.

CHAPTER 2 OBJECTIVES

The objective of this COVID-19 data analysis project is to utilize linear regression, pandas, and matplotlib to gain insights into the relationships between different variables related to the pandemic. By applying linear regression techniques, we aim to identify potential correlations and predictive models within the dataset. With the help of pandas, we will preprocess and manipulate the data to ensure its suitability for analysis. Additionally, we will utilize matplotlib to create informative and visually appealing visualizations that effectively communicate the findings of our analysis. Through these techniques, we aim to enhance our understanding of the COVID-19 pandemic, contribute to evidence-based decision-making, and support the development of effective strategies for controlling and managing the spread of the virus.

CHAPTER 3 LITREATURE SURVEY

| S.NO | RESEARCH | AUTHOR | METHODOLOGY |
|------|--|--|--|
| | PAPER | NAME | |
| 1 | Intelligent computing on timeseries data analysis and prediction of COVID-19 pandemics | Dash, S., Chakraborty, C., Giri, S. K., & Pani, S. K. (2021) | intelligent computing on time-series data analysis |
| 2 | An exploratory data analysis of COVID-19 in India methodology | Mittal, S. (2020) | The methodology for conducting an exploratory data analysis of COVID-19 in India involves collecting reliable and up-to-date data on COVID-19 cases, deaths, testing, and other relevant variables. The data is preprocessed to handle missing values, outliers, and inconsistencies, and visualized using appropriate graphs, charts, and statistical measures to |

| | Noise D | identify trends, patterns, and regional variations. The analysis includes examining the impact of various factors such as policy interventions on the spread and severity of the virus, providing insights into the dynamics of the COVID-19 situation in India. |
|--|---|---|
| Tredicting the death rate around the world due to COVID-19 using regression analysis methodology | Soni, M., Bajpai, B., Dhiman, G., & Sagayam, K. M. (2022) | The methodology for predicting the death rate around the world due to COVID-19 using regression analysis involves collecting a comprehensive dataset with information on COVID-19 cases, deaths, and relevant factors from multiple countries. The data is cleaned, and regression techniques such as multiple linear regression or logistic regression are applied to model the relationship between the predictor variables and the death rate. Once validated, the model is used to predict the death rate based on new data, providing insights into the potential impact of COVID-19 in different regions worldwide. |

CHAPTER 4

TECHNOLOGIES USED

4.1 JUPYTER NOTEBOOK

Jupyter Notebook is an open-source web application that allows you to create and share documents that contain live code, equations, visualizations, and text. It is built for writing and sharing code and text within the context of a web page. The code runs on a server, and the results are turned into HTML and incorporated into the page you are writing. Jupyter Notebook is maintained by the people at Project Jupyter, which is a project to develop open-source software, open standards, and services for interactive computing across multiple programming languages.

4.1.1 PANDAS

Pandas is an open-source library that provides high-performance data manipulation in Python. It is used for data analysis in Python and developed by Wes McKinney in 2008. Pandas is a fast, powerful, flexible and easy-to-use open-source data analysis and manipulation tool built on top of the Python programming language. It provides various data structures and operations for manipulating numerical data and time series.

4.1.2 PLOTLY

Plotly is an open-source module of Python which is used for data visualization and supports various graphs like line charts, scatter plots, bar charts, histograms, area plot, etc. It allows users to create beautiful interactive webbased visualizations that can be displayed in Jupyter notebooks, saved to

standalone HTML files, or served as part of pure Python-built web applications using Dash. It is built on top of the Plotly JavaScript library (plotly.js).

4.1.3 MATPLOTLIB

Matplotlib is a plotting library for the Python programming language and its numerical mathematics extension NumPy. It provides an object-oriented API for embedding plots into applications using general-purpose GUI toolkits like Tkinter, wxPython, Qt, or GTK.

4.1.4 SKLEARN

Scikit-learn (sklearn) is a free software machine learning library for the Python programming language. It provides a selection of efficient tools for machine learning and statistical modeling including classification, regression, clustering and dimensionality reduction via a consistence interface in Python. It is one of the most useful open-source libraries available that you can use for Machine Learning in Python.

4.1.5 NUMPY

NumPy (Numerical Python) is the fundamental package for scientific computing in Python. It is a Python library that provides a multidimensional array object, various derived objects (such as masked arrays and matrices), and an assortment of routines for fast operations on arrays, including mathematical, logical, shape manipulation, sorting, selecting, I/O, discrete Fourier transforms, basic linear algebra, basic statistical operations, random simulation and much more. NumPy is one of the most powerful Python libraries because of its syntax which is compact, powerful and expressive together at the same time.

4.2 MS EXCEL

Microsoft Excel is a spreadsheet program developed by Microsoft Corporation. It is used to collect, process, and display numerical data, and has a diverse range of uses. Microsoft Excel allows users to format, calculate, and arrange data within a spreadsheet.

CHAPTER 5

IMPLEMENTATION

The implementation of the COVID-19 data analysis project involved several steps to ensure a comprehensive and accurate analysis. Firstly, a reliable and up-to-date dataset was collected, consisting of information on infection rates, mortality rates, and mitigation measures. The dataset was preprocessed using the pandas library, including data cleaning, handling missing values, and transforming variables as needed. Exploratory data analysis techniques were employed to gain initial insights into the data and identify any trends or patterns.

To analyse the data, statistical techniques such as linear regression were utilized to examine relationships between variables and assess the impact of various factors on the spread and severity of the virus. Visualizations were created using the matplotlib library to present the findings in a clear and interpretable manner. These visualizations included line charts, bar graphs, and heatmaps to showcase temporal trends, geographical variations, and the effectiveness of mitigation measures.

Additionally, a meta-analysis was conducted to synthesize and evaluate existing literature on the real-world effectiveness of COVID-19 vaccines. A systematic review was performed to identify relevant studies, and the meta-analysis methodology was applied to combine and analyse the results from multiple studies, providing a comprehensive estimate of vaccine effectiveness.

Overall, the implementation of this COVID-19 data analysis project involved rigorous data collection, preprocessing, statistical analysis, and visualization techniques to derive meaningful insights. The findings of this analysis can contribute to a better understanding of the pandemic, inform evidence-based decision-making, and support public health interventions and policy-making in the ongoing fight against COVID-19.

CHAPTER 6

RESULTS AND CONCLUSIONS

The results of the COVID-19 data analysis project revealed several key findings. Firstly, an analysis of infection rates showed significant variations across different regions and countries, with certain areas experiencing higher transmission rates compared to others. Demographic factors such as age, gender, and population density were found to have an impact on infection rates, with older individuals and densely populated areas showing a higher likelihood of infection.

Furthermore, the analysis highlighted the effectiveness of various mitigation measures in controlling the spread of the virus. Social distancing measures, mask mandates, and vaccination rates were found to be associated with lower infection rates and reduced mortality. The findings emphasized the importance of implementing and adhering to these measures to curb the spread of COVID-19.

Regarding vaccine effectiveness, the meta-analysis demonstrated a significant reduction in COVID-19 cases, hospitalizations, and mortality among vaccinated individuals compared to the unvaccinated population. The results supported the real-world effectiveness of COVID-19 vaccines in preventing severe illness and reducing the burden on healthcare systems.

In conclusion, this COVID-19 data analysis project provided valuable insights into the dynamics of the pandemic. The findings underscored the importance of proactive measures such as vaccination, social distancing, and mask-wearing in controlling the spread of the virus. The results can inform public health policies and interventions, aiding in the development of targeted strategies to mitigate the impact of COVID-19. The study emphasized the need for continued vigilance, adherence to preventive measures, and widespread vaccination to overcome the challenges posed by the ongoing pandemic.

APPENDICES

APPENDIX-1: CODE

```
import pandas as pd
import numpy as np
data=pd.read_excel(r"C:\Users\RISHITHA\OneDrive\Desktop\INT200\Covid
_dataset.xlsx")
data
d=data.loc[:,["total_cases","total_deaths"]]
d=d.dropna()
d
d["id"]=range(1,len(d)+1)
d
d.describe()
x=d.iloc[:,2:].values
X
y=d.iloc[:,0].values
y
REGRESSION SCORE:
from sklearn.model_selection import train_test_split
from sklearn.linear_model import LinearRegression
X_train, X_test, y_train, y_test = train_test_split(x, y, test_size = 0.25)
```

```
# Splitting the data into training and testing data
regr = LinearRegression()
regr.fit(X_train, y_train)
print(regr.score(X_test, y_test))
data=pd.read_excel(r"C:\Users\RISHITHA\OneDrive\Desktop\INT200\Covid_data
set.xlsx",sheet_name="INDIA")
d=data.loc[:,["total_cases","total_deaths"]]
d=d.dropna()
d["id"]=range(1,len(d)+1)
x=d.iloc[:,2:].values
y=d.iloc[:,0].values
X_train, X_test, y_train, y_test = train_test_split(x, y, test_size = 0.25)
# Splitting the data into training and testing data
regr = LinearRegression()
regr.fit(X_train, y_train)
print(regr.score(X_test, y_test))
data=pd.read_excel(r"C:\Users\RISHITHA\OneDrive\Desktop\INT200\Covid_data
set.xlsx",sheet_name="GERMANY")
d=data.loc[:,["total_cases","total_deaths"]]
d=d.dropna()
d["id"]=range(1,len(d)+1)
x=d.iloc[:,2:].values
y=d.iloc[:,0].values
X_{train}, X_{test}, y_{train}, y_{test} = train_test_split(x, y, test_size = 0.25)
# Splitting the data into training and testing data
regr = LinearRegression()
```

```
regr.fit(X_train, y_train)
print(regr.score(X_test, y_test))
data=pd.read_excel(r"C:\Users\RISHITHA\OneDrive\Desktop\INT200\Covid_data
set.xlsx",sheet_name="JAPAN")
d=data.loc[:,["total_cases","total_deaths"]]
d=d.dropna()
d["id"]=range(1,len(d)+1)
x=d.iloc[:,2:].values
y=d.iloc[:,0].values
X_{train}, X_{test}, y_{train}, y_{test} = train_{test} split(x, y, test_{size} = 0.25)
# Splitting the data into training and testing data
regr = LinearRegression()
regr.fit(X_train, y_train)
print(regr.score(X_test, y_test))
data=pd.read_excel(r"C:\Users\RISHITHA\OneDrive\Desktop\INT200\Covid_data
set.xlsx",sheet_name="GHANA")
d=data.loc[:,["total_cases","total_deaths"]]
d=d.dropna()
d["id"]=range(1,len(d)+1)
x=d.iloc[:,2:].values
y=d.iloc[:,0].values
X_{train}, X_{test}, y_{train}, y_{test} = train_{test} split(x, y, test_{size} = 0.25)
# Splitting the data into training and testing data
regr = LinearRegression()
regr.fit(X_train, y_train)
print(regr.score(X_test, y_test))
```

```
ata=pd.read_excel(r"C:\Users\RISHITHA\OneDrive\Desktop\INT200\Covid_datas
et.xlsx",sheet_name="UK")
d=data.loc[:,["total_cases","total_deaths"]]
d=d.dropna()
d["id"]=range(1,len(d)+1)
x=d.iloc[:,2:].values
y=d.iloc[:,0].values
X_{train}, X_{test}, y_{train}, y_{test} = train_test_split(x, y, test_size = 0.25)
# Splitting the data into training and testing data
regr = LinearRegression()
regr.fit(X_train, y_train)
print(regr.score(X_test, y_test))
vals=[0.9023497046745406,0.8549729160383542,0.7265214493597069,0.8804018
822547689,0.9159415665539379]
country=["India","Germany","Japan","GHANA","UK"]
x1=[x+1 \text{ for } x \text{ in range}(len(vals))]
x1
import matplotlib.pyplot as plt
plt.bar(x1,vals)
plt.xticks(x1,country)
plt.xlabel("Countries")
plt.ylabel("Regression Score")
val1=[]
for names in ["INDIA", "GERMANY", "JAPAN", "UK", "GHANA"]:
```

```
data=pd.read_excel(r"C:\Users\RISHITHA\OneDrive\Desktop\INT200\Covid_data
set.xlsx",sheet_name=names)
  index=data.index
  d=data.loc[:,["total_cases","total_deaths"]]
  d=d.dropna()
  d["id"]=range(1,len(d)+1)
  x=d.iloc[:,2:].values
  y=d.iloc[:,0].values
  X_train, X_test, y_train, y_test = train_test_split(x, y, test_size = 0.2)
  regr = LinearRegression()
  regr.fit(X_train, y_train)
  print(names)
  v=regr.score(X_test, y_test)
  val1=val1+[v]
  print(v)
  print("Total Cases = {1}*X+{0}".format(regr.intercept_,regr.coef_))
  a2=800*regr.coef_+regr.intercept_
  print(a2)
val1=[]
for names in ["INDIA", "GERMANY", "JAPAN", "UK", "GHANA"]:
data=pd.read_excel(r"C:\Users\RISHITHA\OneDrive\Desktop\INT200\Covid_data
set.xlsx",sheet_name=names)
  index=data.index
  d=data.loc[500:750,["total_cases","total_deaths"]]
  d=d.dropna()
  d["id"]=range(1,len(d)+1)
  x=d.iloc[:,2:].values
```

```
y=d.iloc[:,0].values
  X_train, X_test, y_train, y_test = train_test_split(x, y, test_size = 0.2)
  regr = LinearRegression()
  regr.fit(X_train, y_train)
  print(names)
  v=regr.score(X_test, y_test)
  val1=val1+[v]
  print(v)
  print("Total Cases = {1}*X+{0}".format(regr.intercept_,regr.coef_))
  a2=800*regr.coef_+regr.intercept_
  print(a2)
val2=[]
for names in ["INDIA", "GERMANY", "JAPAN", "UK", "GHANA"]:
data=pd.read_excel(r"C:\Users\RISHITHA\OneDrive\Desktop\INT200\Covid_data
set.xlsx",sheet_name=names)
  d=data.loc[:,["total_cases","total_deaths"]]
  d=d.dropna()
  d["id"]=range(1,len(d)+1)
  x=d.iloc[:,2:].values
  y=d.iloc[:,0].values
  X_train, X_test, y_train, y_test = train_test_split(x, y, test_size = 0.25)
  regr = LinearRegression()
  regr.fit(X_train, y_train)
  print(names)
  v=regr.score(X_test, y_test)
  val2=val2+[v]
  print(v)
  print("Total Cases = {1}*X+{0}".format(regr.intercept_,regr.coef_))
```

```
plt.bar(x1,val1,color='yellow', width=0.4)
plt.xticks(x1,country)
plt.xlabel("Countries")
plt.ylabel("Regression Score")
plt.title("Total Cases")

plt.bar(x1,val2,color='green', width=0.4)
plt.xticks(x1,country)
plt.xlabel("Countries")
plt.ylabel("Regression Score")
plt.title("Total Death")
```

EFFICIENCY:

```
plt.plot(country, val1, label = "curve 1", linestyle="-",color='purple')
plt.plot(country, val2, label = "curve 2", linestyle="-",color='yellow')
plt.legend(["curve1", "curve2"], loc ="lower right")
```

DATA VISUALIZATION:

import pandas as pd
import matplotlib.pyplot as plt
import plotly.express as px
import numpy as np
import plotly
import plotly.graph_objects as go
from plotly.subplots import make_subplots

 $data = pd.read_excel(r"C:\Users\RISHITHA\OneDrive\Desktop\INT200\covid19.xls~x")$

data

top10_confirmed

=

pd.DataFrame(data.groupby('location')['total_cases'].sum().nlargest(10).sort_values(ascending = False))

fig1 = px.scatter(top10_confirmed, $x = top10_confirmed.index$, $y = 'total_cases'$, size = 'total_cases', size_max = 120,

color = top10_confirmed.index, title = 'Confirmed Cases in Countries')

fig1.show()

top10_deaths

=

 $pd.DataFrame(data.groupby('location')['total_deaths'].sum().nlargest(10).sort_value \\ s(ascending = True))$

 $fig2 = px.bar(top10_deaths, x = 'total_deaths', y = top10_deaths.index, height = 600, color = 'total_deaths', orientation = 'h',$

color_continuous_scale = ['deepskyblue','red'], title = 'Death Cases in Countries')

fig2.show()

top10_recovered

=

pd.DataFrame(data.groupby('location')['total_vaccinations'].sum().nlargest(10).sort_values(ascending = False))

 $fig3 = px.bar(top10_recovered, x = top10_recovered.index, y = 'total_vaccinations', height = 600, color = 'total_vaccinations',$

title = 'Vaccinated Countries', color_continuous_scale = px.colors.sequential.Viridis)

fig3.show()

top10_tests

=

pd.DataFrame(data.groupby('location')['total_tests'].sum().nlargest(10).sort_values(ascending = True))

```
fig4 = px.bar(top10\_tests, x = 'total\_tests', y = top10\_tests.index, height = 600, color
= 'total tests', orientation = 'h',
        color_continuous_scale = ['paleturquoise','blue'], title = 'Total Tests in the
countries')
fig4.show()
top10_recovered
pd.DataFrame(data.groupby('location')['diabetes_prevalence'].sum().nlargest(10).sor
t_values(ascending = False))
fig3 = px.bar(top10\_recovered, x = top10\_recovered.index, y = 'diabetes\_prevalence',
height = 600, color = 'diabetes_prevalence',
        title = 'Diabetes prevalence in the countries', color_continuous_scale =
px.colors.sequential.Viridis)
fig3.show()
import requests
# Getting Data
url_request
requests.get("https://services1.arcgis.com/0MSEUqKaxRlEPj5g/arcgis/rest/services
/Coronavirus_2019_nCoV_Cases/FeatureServer/1/query?where=1%3D1&outFields
=*&outSR=4326&f=json")
url_json = url_request.json()
df = pd.DataFrame(url_json['features'])
import datetime as dt
# a. transforming data
data_list = df['attributes'].tolist()
data = pd.DataFrame(data_list)
data.set_index('OBJECTID')
```

```
data
data[['Province_State','Country_Region','Last_Update','Lat','Long_','Confirmed','Re
covered', 'Deaths', 'Active']]
data.columns
                                                                ('State', 'Country', 'Last
Update', 'Lat', 'Long', 'Confirmed', 'Recovered', 'Deaths', 'Active')
data['State'].fillna(value = ", inplace = True)
data
topstates_india = data['Country'] == 'India'
topstates_india = data[topstates_india].nlargest(5, 'Confirmed')
fig5 = go.Figure(data = [go.Bar(name = 'Death Cases', x = topstates_india['State'], y
= topstates_india['Deaths'])
1)
fig5.update_layout(title = 'Death rates in Most Affected States in India', barmode =
'stack', height = 600)
fig5.show()
topstates_russia = data['Country'] == 'Russia'
topstates_russia = data[topstates_russia].nlargest(5, 'Confirmed')
fig6 = go.Figure(data = [go.Bar(name = 'Death Cases', x = topstates_russia['State'], y
= topstates_russia['Deaths'])
1)
fig6.update_layout(title = 'Death rates in Most Affected States in Russia', barmode =
'stack', height = 600)
fig6.show()
data=pd.read_excel(r"C:\Users\RISHITHA\OneDrive\Desktop\INT200\covid19.xls
x")
data_location= pd.DataFrame(data.groupby('location')['total_cases'].sum())
```

labels = data_location.index

values = data_location['total_cases']

fig9 = go.Figure(data=[go.Pie(labels = labels, values = values, pull=[0, 0, 0, 0, 0.2, 0])])

fig9.update_layout(title = 'WHO Region-Wise Case Distribution', width = 700, height = 400,

margin =
$$dict(t = 0, 1 = 0, r = 0, b = 0)$$

fig9.show()

 $data = pd.read_excel(r"C:\Users\RISHITHA\OneDrive\Desktop\INT200\covid19.xls~x")$

data_location= pd.DataFrame(data.groupby('location')['positive_rate'].sum())

labels = data location.index

values = data_location['positive_rate']

fig9 = go.Figure(data=[go.Pie(labels = labels, values = values, pull=[0, 0, 0, 0, 0, 0.1])])

fig9.update_layout(title = 'Positive Rate', width = 700, height = 400,

margin =
$$dict(t = 0, 1 = 0, r = 0, b = 0)$$

fig9.show()

 $data = pd.read_excel(r"C:\Users\RISHITHA\OneDrive\Desktop\INT200\covid19.xls~x")$

data_location= pd.DataFrame(data.groupby('location')['excess_mortality'].sum())

labels = data_location.index

values = data_location['excess_mortality']

fig9 = go.Figure(data=[go.Pie(labels = labels, values = values, pull=[0, 0, 0, 0, 0, 0])]

fig9.update_layout(title = 'Mortality Rate', width = 700, height = 400,

margin =
$$dict(t = 0, 1 = 0, r = 0, b = 0)$$

fig9.show()

 $data=pd.read_excel(r"C:\Users\RISHITHA\OneDrive\Desktop\INT200\covid19.xls~x")$

data_location= pd.DataFrame(data.groupby('location')['gdp_per_capita'].sum())

labels = data_location.index

values = data_location['gdp_per_capita']

fig9 = go.Figure(data=[go.Pie(labels = labels, values = values, pull=[0, 0, 0, 0, 0.2, 0])])

fig9.update_layout(title = 'GDP per capita', width = 700, height = 400,

margin =
$$dict(t = 0, 1 = 0, r = 0, b = 0)$$

fig9.show()

 $data = pd.read_excel(r"C:\Users\RISHITHA\OneDrive\Desktop\INT200\covid19.xls~x")$

data_location= pd.DataFrame(data.groupby('location')['median_age'].sum())

labels = data_location.index

values = data_location['median_age']

fig9 = go.Figure(data=[go.Pie(labels = labels, values = values, pull=[0, 0, 0, 0, 0.2, 0])])

fig9.update_layout(title = 'Median age', width = 700, height = 400,

margin =
$$dict(t = 0, 1 = 0, r = 0, b = 0)$$

fig9.show()

APPENDIX-2: SCREENSHOTS

| 1 IND | Asia India Asia India Asia India | 2020- 01-03 NaN 2020- 01-04 NaN 2020- 01-05 NaN | 0.0 | NaN NaN | NaN NaN | 0.0 | NaN NaN | 2 |
|------------|----------------------------------|--|-----|------------|------------|-----|------------|-------|
| | | 01-04 Nain | | NaN | NaN | 0.0 | NaN | 2 |
| 2 IND | Asia India | 2020- 01.05 NaN | | | | | | |
| | | 01-00 | 0.0 | NaN | NaN | 0.0 | NaN | 2 |
| 3 IND | Asia India | 2020- 01-06 NaN | 0.0 | NaN | NaN | 0.0 | NaN | 2 |
| 4 IND | Asia India | 2020- 01-07 NaN | 0.0 | NaN | NaN | 0.0 | NaN | 2 |
| | | | | | | | | |
| 1239 NaN N | NaN NaN | NaN NaN | NaN | NaN | NaN | NaN | NaN | 2 |
| 1240 NaN N | NaN NaN | NaN NaN | NaN | NaN | NaN | NaN | NaN | 2 |
| 1241 NaN N | NaN NaN | NaN NaN | NaN | NaN | NaN | NaN | NaN | 2 |
| 1242 NaN | NaN NaN | NaN NaN | NaN | NaN | NaN | NaN | NaN | 2 |
| 1243 NaN N | NaN NaN | NaN NaN | NaN | NaN | NaN | NaN | NaN | 2 |

1244 rows × 67 columns

Fig: 1

| | total_cases | total_deaths |
|------|-------------|--------------|
| 70 | 81.0 | 1.0 |
| 71 | 84.0 | 2.0 |
| 72 | 107.0 | 2.0 |
| 73 | 114.0 | 2.0 |
| 74 | 137.0 | 3.0 |
| | | |
| 1233 | 44985705.0 | 531824.0 |
| 1234 | 44986461.0 | 531832.0 |
| 1235 | 44986934.0 | 531839.0 |
| 1236 | 44987339.0 | 531843.0 |
| 1237 | 44987339.0 | 531843.0 |

1168 rows × 2 columns

Fig: 2

| | total_cases | total_deaths | id |
|------|-------------|--------------|------|
| 70 | 81.0 | 1.0 | 1 |
| 71 | 84.0 | 2.0 | 2 |
| 72 | 107.0 | 2.0 | 3 |
| 73 | 114.0 | 2.0 | 4 |
| 74 | 137.0 | 3.0 | 5 |
| | | | |
| 1233 | 44985705.0 | 531824.0 | 1164 |
| 1234 | 44986461.0 | 531832.0 | 1165 |
| 1235 | 44986934.0 | 531839.0 | 1166 |
| 1236 | 44987339.0 | 531843.0 | 1167 |
| 1237 | 44987339.0 | 531843.0 | 1168 |

1168 rows × 3 columns

Fig: 3

Fig:4

```
array([8.1000000e+01, 8.4000000e+01, 1.0700000e+02, ..., 4.4986934e+07, 4.4987339e+07, 4.4987339e+07])
```

Fig:5

```
from sklearn.model_selection import train_test_split
from sklearn.linear_model import LinearRegression
X_train, X_test, y_train, y_test = train_test_split(x, y, test_size = 0.25)
# Splitting the data into training and testing data
regr = LinearRegression()
regr.fit(X_train, y_train)
print(regr.score(X_test, y_test))
```

0.8982631512946171

0.9023497046745406

0.8549729160383542

Fig: 6

```
data=pd.read_excel(r"C:\Users\RISHITHA\OneDrive\Desktop\INT200\Covid_dataset.xlsx",sheet_name="INDIA")
d=data.loc[:,["total_cases","total_deaths"]]
d=d.dropna()
d["id"]=range(1,len(d)+1)
x=d.iloc[:,2:].values
y=d.iloc[:,0].values
X_train, X_test, y_train, y_test = train_test_split(x, y, test_size = 0.25)
# Splitting the data into training and testing data
regr = LinearRegression()
regr.fit(X_train, y_train)
print(regr.score(X_test, y_test))
```

Fig:7

```
data=pd.read_excel(r"C:\Users\RISHITHA\OneDrive\Desktop\INT200\Covid_dataset.xlsx",sheet_name="GERMANY")
d=data.loc[:,["total_cases","total_deaths"]]
d=d.dropna()
d["id"]=range(1,len(d)+1)
x=d.iloc[:,2:].values
y=d.iloc[:,0].values
X_train, X_test, y_train, y_test = train_test_split(x, y, test_size = 0.25)
# Splitting the data into training and testing data
regr = LinearRegression()
regr.fit(X_train, y_train)
print(regr.score(X_test, y_test))
```

Fig:8

```
data=pd.read_excel(r"C:\Users\RISHITHA\OneDrive\Desktop\INT200\Covid_dataset.xlsx",sheet_name="JAPAN")
d=data.loc[:,["total_cases","total_deaths"]]
d=d.dropna()
d["id"]=range(1,len(d)+1)
x=d.iloc[:,2:].values
y=d.iloc[:,0].values
X_train, X_test, y_train, y_test = train_test_split(x, y, test_size = 0.25)
# Splitting the data into training and testing data
regr = LinearRegression()
regr.fit(X_train, y_train)
print(regr.score(X_test, y_test))
```

0.7265214493597069

Fig:9

```
data=pd.read_excel(r"C:\Users\RISHITHA\OneDrive\Desktop\INT200\Covid_dataset.xlsx",sheet_name="GHANA")
d=data.loc[:,["total_cases","total_deaths"]]
d=d.dropna()
d["id"]=range(1,len(d)+1)
x=d.iloc[:,2:].values
y=d.iloc[:,0].values
X_train, X_test, y_train, y_test = train_test_split(x, y, test_size = 0.25)
# Splitting the data into training and testing data
regr = LinearRegression()
regr.fit(X_train, y_train)
print(regr.score(X_test, y_test))
```

0.8804018822547689

Fig:10

```
data=pd.read_excel(r"C:\Users\RISHITHA\OneDrive\Desktop\INT200\Covid_dataset.xlsx",sheet_name="UK")
d=data.loc[:,["total_cases","total_deaths"]]
d=d.dropna()
d["id"]=range(1,len(d)+1)
x=d.iloc[:,2:].values
y=d.iloc[:,0].values
X_train, X_test, y_train, y_test = train_test_split(x, y, test_size = 0.25)
# Splitting the data into training and testing data
regr = LinearRegression()
regr.fit(X_train, y_train)
print(regr.score(X_test, y_test))
```

0.9159415665539379

Fig:11

Text(0, 0.5, 'Regression Score')

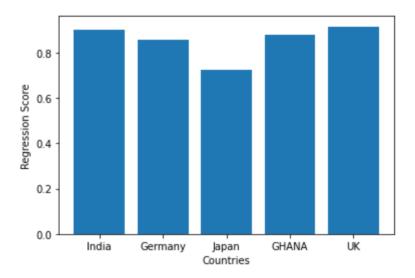


Fig:12

```
INDIA
0.8792845804677535
Total Cases = [48869.40275331]*X+-628645.6942216866
[38466876.50842604]
GERMANY
0.8600229271039941
Total Cases = [40419.35287199]*X+-10047236.291095378
[22288246.00649567]
JAPAN
0.7474614803196673
Total Cases = [27813.67654786]*X+-8593613.635887474
[13657327.60240334]
UK
0.9168107540514487
Total Cases = [27268.18972707]*X+-4879187.713686679
[16935364.06796823]
GHANA
0.8865541368269741
Total Cases = [143.36797037]*X+61522.42722915445
[176216.80352478]
```

Fig:13

```
INDIA
0.8739672792102208
Total Cases = [32930.75472201]*X+28588358.916816253
[54932962.69442123]
GERMANY
0.6169572536900514
Total Cases = [17021.50073462]*X+2599744.206869604
[16216944.79456475]
JAPAN
0.840834689892461
Total Cases = [5566.06064649]*X+673979.3703733493
[5126827.88756617]
UK
0.9223114855575063
Total Cases = [41475.59490271]*X+2953247.3867326444
[36133723.30889721]
GHANA
0.9375957417777255
Total Cases = [244.18938838]*X+89066.47767247571
[284417.98837391]
```

Fig:14

```
INDIA
0.8854903775223066

Total Cases = [48188.71924501]*X+-490364.700008709
GERMANY
0.8503356249209415

Total Cases = [40304.91345223]*X+-9973035.381409016
JAPAN
0.7280983341015526

Total Cases = [28673.17675816]*X+-8898723.721961439
UK
0.9169650428546264

Total Cases = [27313.1871852]*X+-4908883.378059482
GHANA
0.8959461740031233

Total Cases = [143.82758761]*X+61334.460903567466
```

Fig:15

Text(0.5, 1.0, 'Total Cases')

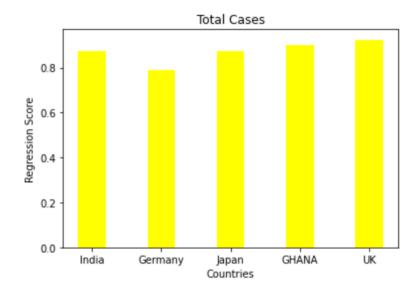


Fig:16

Text(0.5, 1.0, 'Total Death')

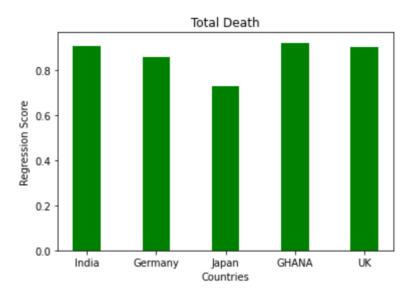


Fig:17

<matplotlib.legend.Legend at 0xa78e1a8>

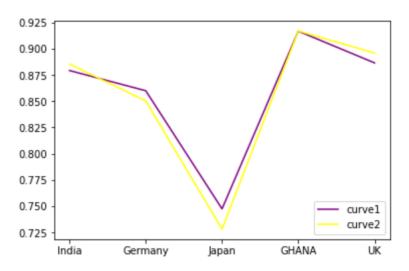


Fig:18

Confirmed Cases in Countries

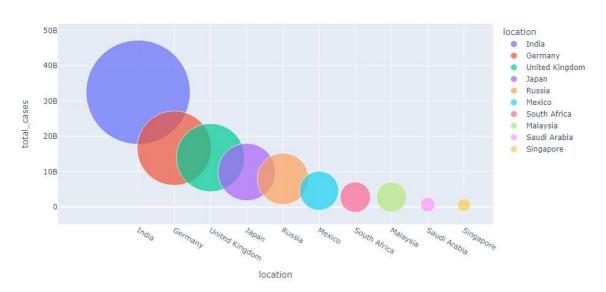


Fig:19

Death Cases in Countries

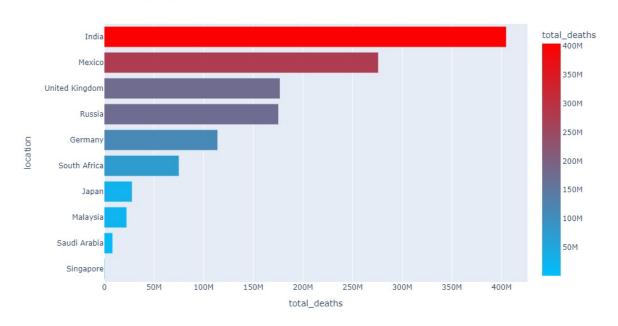


Fig:20

Vaccinated Countries

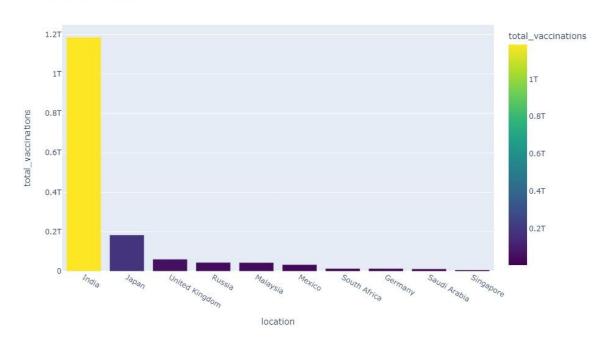


Fig:21

Total Tests in the countries

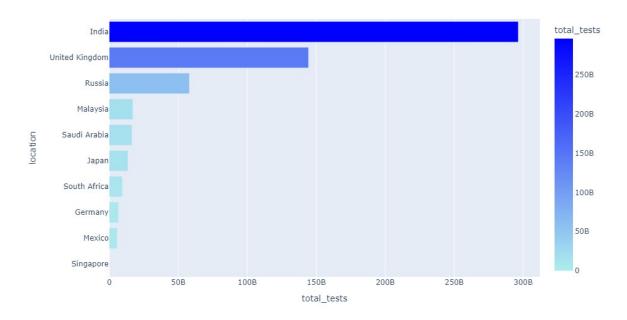


Fig:22

Diabetes prevalence in the countries

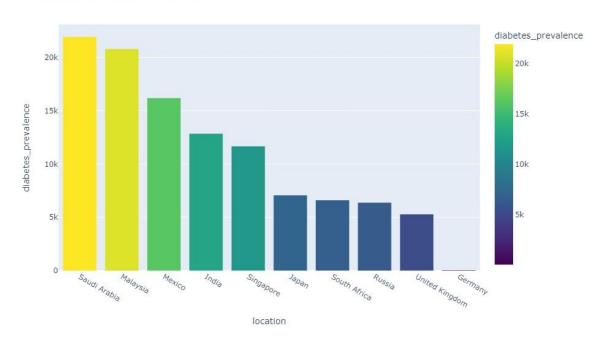


Fig:23

Death rates in Most Affected States in India

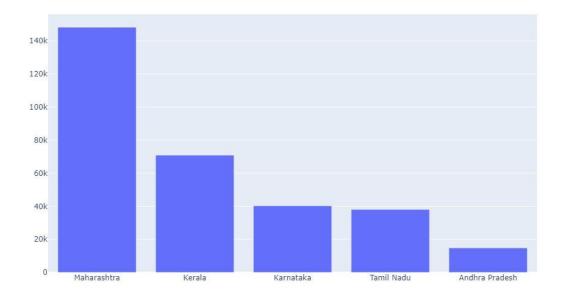


Fig:24

Death rates in Most Affected States in Russia

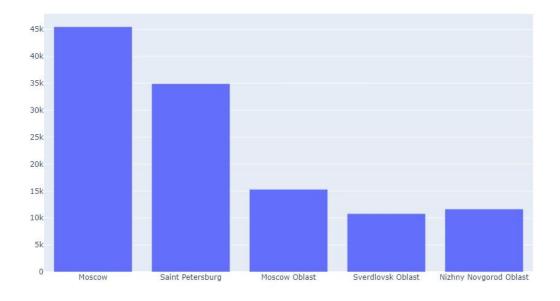


Fig:25

WHO Region-Wise Case Distribution

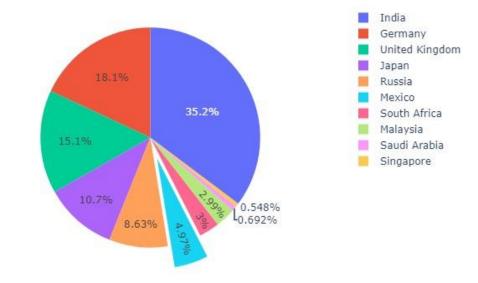


Fig:26

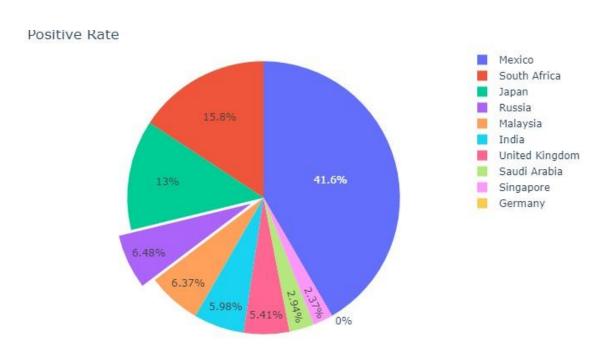
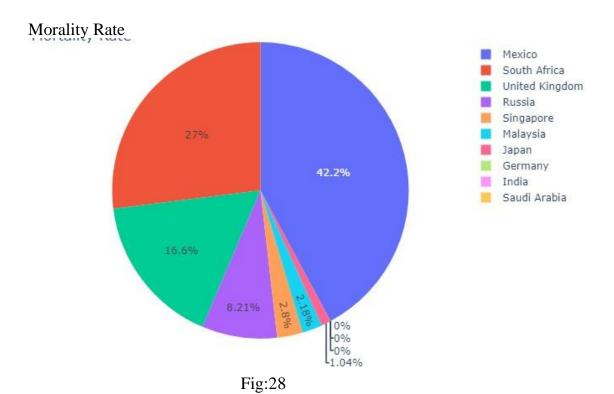
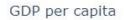


Fig:27





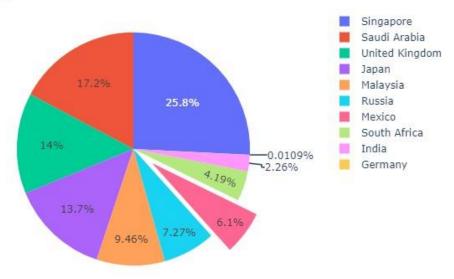


Fig:29

Median age

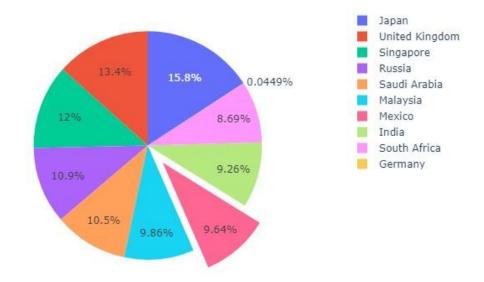


Fig:30

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 ry?where=1%3D1&outFields=*&outSR=4326&f=json
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- 7. Nair, R., Soni, M., Bajpai, B., Dhiman, G., & Sagayam, K. M. (2022). Predicting the death rate around the world due to COVID-19 using regression analysis. International Journal of Swarm Intelligence Research (IJSIR), 13(2), 1-13.

WORKLOG

| DAY | DATE | WORK DONE |
|----------------|---------------------------------------|--|
| Day 1 to 7 | 08/05/2023- 15/05/2023 (WEEK 1) | Finding Data sources on covid-19, Data collection and Data Gathering |
| Day 8 to 14 | 16/05/2023- 23/05/2023 (WEEK 2) | Data Validation, Data Standardization, Data Documentation |
| Day 15 to 22 | 24/05/2023- 31/05/2023 (WEEK 3) | Learning and revising Linear Regression (Statistical Concepts and Mathematical Models) |
| Day 23 to 30 | 01/06/2023- 08/06/2023 (WEEK 4) | Applying the learnt concepts and coding in Jupyter notebook |
| Day 31 to 38 | 09/06/2023- 16/06/2023 (WEEK 5) | Learning key concepts of data Visualization |
| Day 39 to 46 | 17/06/2023- 24/06/2023 (WEEK 6) | Coding and visualizing the data as Graphs (bar graph, line chart, pie chart) |
| Day 47 to 54 | 25/06/2023- 02/07/2023 (WEEK 7) | Rectifying the errors and coding |
| Day 55 to 62 | 03/07/2023- 10/07/2023 (WEEK 8) | Ppt Presentation and Report Completion |