

An update on mortality and morbidity rate to track covid-19 outbreak - A case study

October 24, 2023

1 An update on mortality and morbidity global rates to track covid-19 outbreak - A case study.

1.1 # Background:

- In December, 2019, a local outbreak of pneumonia of initially unknown cause was detected in Wuhan (Hubei, China), and was quickly identified to be caused by a novel coronavirus, namely severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2). The outbreak has since spread to every province of mainland China as well as 27 other countries and regions, with more than 70 000 confirmed cases as of Feb 17, 2020. In response to this ongoing public health emergency, an online interactive dashboard, hosted by the Center for Systems Science and Engineering (CSSE) at Johns Hopkins University, Baltimore, MD, USA, to visualise and track reported cases of coronavirus disease 2019 (COVID-19) in real time. All data collected and displayed are made freely available, initially through Google Sheets and now through a GitHub repository, along with the feature layers of the dashboard, which are now included in the Esri Living Atlas. On March 10, 2023, the Johns Hopkins Coronavirus Resource Center ceased its collecting and reporting of global COVID-19 data * —

1.2 # Objective:

- Understanding and identifying any trends on confirmed (positive), death(mortality), active(recovered) cases * —

1.3 # Method:

- Data was collated from Novel Coronavirus (COVID-19) Cases, provided by JHU CSSE, the data repository for the 2019 Novel Coronavirus Visual Dashboard operated by the Johns Hopkins University Center for Systems Science and Engineering (JHU CSSE). Also, Supported by ESRI Living Atlas Team and the Johns Hopkins University Applied Physics Lab (JHU APL). An opendata source was used to retrieve data [this is a link].(<https://github.com/CSSEGISandData/COVID-19>). * —

[132]: *## Importing libraries for data aquisition and evidence syntheis*

```
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
```

```
import matplotlib.style as style
style.available
import seaborn as sns
%matplotlib inline

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

```
[133]: ## Graphical preparation

import plotly.express as px
import plotly.graph_objs as go
from plotly.subplots import make_subplots
```

```
[134]: ## Using Covid-19 CSSE at John Hopkins University data from J.H github ↵
       ↪ repository

positive_df = pd.read_csv('https://raw.githubusercontent.com/CSSEGISandData/
↪ COVID-19/master/csse_covid_19_data/csse_covid_19_time_series/
↪ time_series_covid19_confirmed_global.csv')
mortality_df = pd.read_csv('https://raw.githubusercontent.com/CSSEGISandData/
↪ COVID-19/master/csse_covid_19_data/csse_covid_19_time_series/
↪ time_series_covid19_deaths_global.csv')
morbidity_df = pd.read_csv('https://raw.githubusercontent.com/CSSEGISandData/
↪ COVID-19/master/csse_covid_19_data/csse_covid_19_time_series/
↪ time_series_covid19_recovered_global.csv')
```

2 An overview of the data sets

```
[136]: ### Shape of the data sets

print(positive_df.shape)
print(mortality_df.shape)
print(morbidity_df.shape)
```

```
(289, 1147)
```

```
(289, 1147)
```

```
(274, 1147)
```

```
[137]: ### Content in the data set

positive_df.head()
```

```
[137]: Province/State Country/Region      Lat      Long  1/22/20  1/23/20  \
0      NaN      Afghanistan  33.93911  67.709953      0      0
1      NaN      Albania    41.15330  20.168300      0      0
2      NaN      Algeria    28.03390   1.659600      0      0
3      NaN      Andorra    42.50630   1.521800      0      0
4      NaN      Angola     -11.20270  17.873900      0      0

      1/24/20  1/25/20  1/26/20  1/27/20  ...  2/28/23  3/1/23  3/2/23  3/3/23  \
0      0      0      0      0      0  ...  209322  209340  209358  209362
1      0      0      0      0      0  ...  334391  334408  334408  334427
2      0      0      0      0      0  ...  271441  271448  271463  271469
3      0      0      0      0      0  ...   47866   47875   47875   47875
4      0      0      0      0      0  ...  105255  105277  105277  105277

      3/4/23  3/5/23  3/6/23  3/7/23  3/8/23  3/9/23
0  209369  209390  209406  209436  209451  209451
1  334427  334427  334427  334427  334443  334457
2  271469  271477  271477  271490  271494  271496
3   47875   47875   47875   47875   47890   47890
4  105277  105277  105277  105277  105288  105288
```

[5 rows x 1147 columns]

```
[138]: mortality_df.head()
```

```
[138]: Province/State Country/Region      Lat      Long  1/22/20  1/23/20  \
0      NaN      Afghanistan  33.93911  67.709953      0      0
1      NaN      Albania    41.15330  20.168300      0      0
2      NaN      Algeria    28.03390   1.659600      0      0
3      NaN      Andorra    42.50630   1.521800      0      0
4      NaN      Angola     -11.20270  17.873900      0      0

      1/24/20  1/25/20  1/26/20  1/27/20  ...  2/28/23  3/1/23  3/2/23  3/3/23  \
0      0      0      0      0      0  ...   7896   7896   7896   7896
1      0      0      0      0      0  ...   3598   3598   3598   3598
2      0      0      0      0      0  ...   6881   6881   6881   6881
3      0      0      0      0      0  ...    165    165    165    165
4      0      0      0      0      0  ...   1933   1933   1933   1933

      3/4/23  3/5/23  3/6/23  3/7/23  3/8/23  3/9/23
0   7896   7896   7896   7896   7896   7896
1   3598   3598   3598   3598   3598   3598
2   6881   6881   6881   6881   6881   6881
3    165    165    165    165    165    165
4   1933   1933   1933   1933   1933   1933
```

[5 rows x 1147 columns]

```
[139]: morbidity_df.head()
```

```
[139]: Province/State Country/Region      Lat      Long  1/22/20  1/23/20  \
0      NaN      Afghanistan  33.93911  67.709953      0      0
1      NaN      Albania    41.15330  20.168300      0      0
2      NaN      Algeria    28.03390   1.659600      0      0
3      NaN      Andorra    42.50630   1.521800      0      0
4      NaN      Angola     -11.20270  17.873900      0      0

      1/24/20  1/25/20  1/26/20  1/27/20  ...  2/28/23  3/1/23  3/2/23  3/3/23  \
0      0      0      0      0      ...      0      0      0      0
1      0      0      0      0      ...      0      0      0      0
2      0      0      0      0      ...      0      0      0      0
3      0      0      0      0      ...      0      0      0      0
4      0      0      0      0      ...      0      0      0      0

      3/4/23  3/5/23  3/6/23  3/7/23  3/8/23  3/9/23
0      0      0      0      0      0      0
1      0      0      0      0      0      0
2      0      0      0      0      0      0
3      0      0      0      0      0      0
4      0      0      0      0      0      0
```

[5 rows x 1147 columns]

```
[140]: ### Identifying missing values within each data sets
```

```
positive_df.isna().sum()
```

```
[140]: Province/State      198
Country/Region          0
Lat                      2
Long                     2
1/22/20                  0
...
3/5/23                   0
3/6/23                   0
3/7/23                   0
3/8/23                   0
3/9/23                   0
Length: 1147, dtype: int64
```

```
[141]: mortality_df.isna().sum()
```

```
[141]: Province/State      198
Country/Region          0
Lat                      2
```

```

Long                2
1/22/20             0
...
3/5/23              0
3/6/23              0
3/7/23              0
3/8/23              0
3/9/23              0
Length: 1147, dtype: int64

```

```
[142]: morbidity_df.isna().sum()
```

```

[142]: Province/State    199
Country/Region         0
Lat                    1
Long                   1
1/22/20                0
...
3/5/23                 0
3/6/23                 0
3/7/23                 0
3/8/23                 0
3/9/23                 0
Length: 1147, dtype: int64

```

```

[143]: ### Identifying unique values within the data sets

positive_df.nunique()

```

```

[143]: Province/State    91
Country/Region        201
Lat                   283
Long                  284
1/22/20                11
...
3/5/23                 287
3/6/23                 287
3/7/23                 287
3/8/23                 287
3/9/23                 287
Length: 1147, dtype: int64

```

```
[144]: mortality_df.nunique()
```

```

[144]: Province/State    91
Country/Region        201
Lat                   283

```

```

Long          284
1/22/20        2
...
3/5/23        244
3/6/23        243
3/7/23        243
3/8/23        243
3/9/23        242
Length: 1147, dtype: int64

```

```
[145]: morbidity_df.nunique()
```

```

[145]: Province/State    75
Country/Region        201
Lat                   272
Long                  272
1/22/20                3
...
3/5/23                1
3/6/23                1
3/7/23                1
3/8/23                1
3/9/23                1
Length: 1147, dtype: int64

```

```
[146]: ### Counting multiple entry by country/region

positive_df['Country/Region'].value_counts()
```

```

[146]: China            34
Canada              16
United Kingdom     15
France             12
Australia          8
..
Guinea             1
Guinea-Bissau      1
Guyana             1
Haiti              1
Zimbabwe           1
Name: Country/Region, Length: 201, dtype: int64

```

```
[147]: mortality_df['Country/Region'].value_counts()
```

```

[147]: China            34
Canada              16
United Kingdom     15

```

```

France          12
Australia       8
..
Guinea          1
Guinea-Bissau   1
Guyana          1
Haiti           1
Zimbabwe        1
Name: Country/Region, Length: 201, dtype: int64

```

```
[148]: morbidity_df['Country/Region'].value_counts()
```

```

[148]: China          34
United Kingdom    15
France           12
Australia         8
Netherlands       5
..
Guinea           1
Guinea-Bissau    1
Guyana           1
Haiti            1
Zimbabwe         1
Name: Country/Region, Length: 201, dtype: int64

```

```

[149]: ## Data cleaning

## Changing the coloumms 'Province/State' & 'Country/Region' & change latest_
↳date to 'Current'.

```

```

[150]: col=positive_df.columns[-1]

positive_df.rename(columns = {'Province/State' : 'Province', 'Country/Region' :
↳'Country', col : 'Current'},inplace = True)
mortality_df.rename(columns = {'Province/State' : 'Province', 'Country/Region' :
↳ 'Country', col : 'Current'},inplace = True)
morbidity_df.rename(columns = {'Province/State' : 'Province', 'Country/Region' :
↳ 'Country', col : 'Current'},inplace = True)
positive_df.head(3)
mortality_df.head(3)
morbidity_df.head(3)

```

```

[150]: Province      Country      Lat      Long  1/22/20  1/23/20  1/24/20  \
0      NaN  Afghanistan  33.93911  67.709953      0      0      0
1      NaN    Albania   41.15330  20.168300      0      0      0
2      NaN    Algeria   28.03390   1.659600      0      0      0

```

	1/25/20	1/26/20	1/27/20	...	2/28/23	3/1/23	3/2/23	3/3/23	3/4/23	\
0	0	0	0	...	0	0	0	0	0	
1	0	0	0	...	0	0	0	0	0	
2	0	0	0	...	0	0	0	0	0	

	3/5/23	3/6/23	3/7/23	3/8/23	Current
0	0	0	0	0	0
1	0	0	0	0	0
2	0	0	0	0	0

[3 rows x 1147 columns]

```
[151]: ## Creating new data frame through combining unique country entries
```

```
[152]: ### Positive cases
```

```
positive = pd.DataFrame(positive_df.groupby('Country').sum())
positive.reset_index(inplace = True)
positive.head(5)
```

/tmp/ipykernel_1139/1101471026.py:3: FutureWarning:

The default value of numeric_only in DataFrameGroupBy.sum is deprecated. In a future version, numeric_only will default to False. Either specify numeric_only or select only columns which should be valid for the function.

```
[152]:
```

	Country	Lat	Long	1/22/20	1/23/20	1/24/20	1/25/20	\
0	Afghanistan	33.93911	67.709953	0	0	0	0	
1	Albania	41.15330	20.168300	0	0	0	0	
2	Algeria	28.03390	1.659600	0	0	0	0	
3	Andorra	42.50630	1.521800	0	0	0	0	
4	Angola	-11.20270	17.873900	0	0	0	0	

	1/26/20	1/27/20	1/28/20	...	2/28/23	3/1/23	3/2/23	3/3/23	3/4/23	\
0	0	0	0	...	209322	209340	209358	209362	209369	
1	0	0	0	...	334391	334408	334408	334427	334427	
2	0	0	0	...	271441	271448	271463	271469	271469	
3	0	0	0	...	47866	47875	47875	47875	47875	
4	0	0	0	...	105255	105277	105277	105277	105277	

	3/5/23	3/6/23	3/7/23	3/8/23	Current
0	209390	209406	209436	209451	209451
1	334427	334427	334427	334443	334457
2	271477	271477	271490	271494	271496
3	47875	47875	47875	47890	47890
4	105277	105277	105277	105288	105288

[5 rows x 1146 columns]

[153]: *### Moratlity cases*

```
mortality = pd.DataFrame(mortality_df.groupby('Country').sum())
mortality.reset_index(inplace = True)
mortality.head(5)
```

/tmp/ipykernel_1139/1933718209.py:3: FutureWarning:

The default value of numeric_only in DataFrameGroupBy.sum is deprecated. In a future version, numeric_only will default to False. Either specify numeric_only or select only columns which should be valid for the function.

[153]:

	Country	Lat	Long	1/22/20	1/23/20	1/24/20	1/25/20	\
0	Afghanistan	33.93911	67.709953	0	0	0	0	
1	Albania	41.15330	20.168300	0	0	0	0	
2	Algeria	28.03390	1.659600	0	0	0	0	
3	Andorra	42.50630	1.521800	0	0	0	0	
4	Angola	-11.20270	17.873900	0	0	0	0	

	1/26/20	1/27/20	1/28/20	...	2/28/23	3/1/23	3/2/23	3/3/23	3/4/23	\
0	0	0	0	...	7896	7896	7896	7896	7896	
1	0	0	0	...	3598	3598	3598	3598	3598	
2	0	0	0	...	6881	6881	6881	6881	6881	
3	0	0	0	...	165	165	165	165	165	
4	0	0	0	...	1933	1933	1933	1933	1933	

	3/5/23	3/6/23	3/7/23	3/8/23	Current
0	7896	7896	7896	7896	7896
1	3598	3598	3598	3598	3598
2	6881	6881	6881	6881	6881
3	165	165	165	165	165
4	1933	1933	1933	1933	1933

[5 rows x 1146 columns]

[154]: *### Morbidity cases*

```
morbidity = pd.DataFrame(morbidity_df.groupby('Country').sum())
morbidity.reset_index(inplace = True)
morbidity.head(5)
```

/tmp/ipykernel_1139/3587905092.py:3: FutureWarning:

The default value of numeric_only in DataFrameGroupBy.sum is deprecated. In a

future version, `numeric_only` will default to `False`. Either specify `numeric_only` or select only columns which should be valid for the function.

```
[154]:
```

	Country	Lat	Long	1/22/20	1/23/20	1/24/20	1/25/20	\
0	Afghanistan	33.93911	67.709953	0	0	0	0	
1	Albania	41.15330	20.168300	0	0	0	0	
2	Algeria	28.03390	1.659600	0	0	0	0	
3	Andorra	42.50630	1.521800	0	0	0	0	
4	Angola	-11.20270	17.873900	0	0	0	0	

	1/26/20	1/27/20	1/28/20	...	2/28/23	3/1/23	3/2/23	3/3/23	3/4/23	\
0	0	0	0	...	0	0	0	0	0	
1	0	0	0	...	0	0	0	0	0	
2	0	0	0	...	0	0	0	0	0	
3	0	0	0	...	0	0	0	0	0	
4	0	0	0	...	0	0	0	0	0	

	3/5/23	3/6/23	3/7/23	3/8/23	Current
0	0	0	0	0	0
1	0	0	0	0	0
2	0	0	0	0	0
3	0	0	0	0	0
4	0	0	0	0	0

[5 rows x 1146 columns]

```
[155]: ### Eliminating cordination data (latitude and Longitude)
```

```
col = positive['Country']
positive.drop(['Lat', 'Long'], axis=1, inplace=True)
positive.head(3)
```

```
[155]:
```

	Country	1/22/20	1/23/20	1/24/20	1/25/20	1/26/20	1/27/20	1/28/20	\
0	Afghanistan	0	0	0	0	0	0	0	
1	Albania	0	0	0	0	0	0	0	
2	Algeria	0	0	0	0	0	0	0	

	1/29/20	1/30/20	...	2/28/23	3/1/23	3/2/23	3/3/23	3/4/23	3/5/23	\
0	0	0	...	209322	209340	209358	209362	209369	209390	
1	0	0	...	334391	334408	334408	334427	334427	334427	
2	0	0	...	271441	271448	271463	271469	271469	271477	

	3/6/23	3/7/23	3/8/23	Current
0	209406	209436	209451	209451
1	334427	334427	334443	334457
2	271477	271490	271494	271496

[3 rows x 1144 columns]

```
[156]: col = mortality['Country']
mortality.drop(['Lat', 'Long'], axis=1, inplace=True)
mortality.head(3)
```

```
[156]:
```

	Country	1/22/20	1/23/20	1/24/20	1/25/20	1/26/20	1/27/20	1/28/20	\
0	Afghanistan	0	0	0	0	0	0	0	
1	Albania	0	0	0	0	0	0	0	
2	Algeria	0	0	0	0	0	0	0	

	1/29/20	1/30/20	...	2/28/23	3/1/23	3/2/23	3/3/23	3/4/23	3/5/23	\
0	0	0	...	7896	7896	7896	7896	7896	7896	
1	0	0	...	3598	3598	3598	3598	3598	3598	
2	0	0	...	6881	6881	6881	6881	6881	6881	

	3/6/23	3/7/23	3/8/23	Current
0	7896	7896	7896	7896
1	3598	3598	3598	3598
2	6881	6881	6881	6881

[3 rows x 1144 columns]

```
[157]: col = morbidity['Country']
morbidity.drop(['Lat', 'Long'], axis=1, inplace=True)
morbidity.head(3)
```

```
[157]:
```

	Country	1/22/20	1/23/20	1/24/20	1/25/20	1/26/20	1/27/20	1/28/20	\
0	Afghanistan	0	0	0	0	0	0	0	
1	Albania	0	0	0	0	0	0	0	
2	Algeria	0	0	0	0	0	0	0	

	1/29/20	1/30/20	...	2/28/23	3/1/23	3/2/23	3/3/23	3/4/23	3/5/23	\
0	0	0	...	0	0	0	0	0	0	
1	0	0	...	0	0	0	0	0	0	
2	0	0	...	0	0	0	0	0	0	

	3/6/23	3/7/23	3/8/23	Current
0	0	0	0	0
1	0	0	0	0
2	0	0	0	0

[3 rows x 1144 columns]

```
[158]: ## Calculating active cases
```

```
[159]: ### Creating new data frame of Active cases
```

```
active= positive.copy()
for i in active.columns[1:]:
    active[i] =active[i] - mortality[i]
active.head()
```

```
[159]:
```

	Country	1/22/20	1/23/20	1/24/20	1/25/20	1/26/20	1/27/20	1/28/20	\
0	Afghanistan	0	0	0	0	0	0	0	
1	Albania	0	0	0	0	0	0	0	
2	Algeria	0	0	0	0	0	0	0	
3	Andorra	0	0	0	0	0	0	0	
4	Angola	0	0	0	0	0	0	0	

	1/29/20	1/30/20	...	2/28/23	3/1/23	3/2/23	3/3/23	3/4/23	3/5/23	\
0	0	0	...	201426	201444	201462	201466	201473	201494	
1	0	0	...	330793	330810	330810	330829	330829	330829	
2	0	0	...	264560	264567	264582	264588	264588	264596	
3	0	0	...	47701	47710	47710	47710	47710	47710	
4	0	0	...	103322	103344	103344	103344	103344	103344	

	3/6/23	3/7/23	3/8/23	Current
0	201510	201540	201555	201555
1	330829	330829	330845	330859
2	264596	264609	264613	264615
3	47710	47710	47725	47725
4	103344	103344	103355	103355

[5 rows x 1144 columns]

```
[160]: ## Data analysis
```

```
[161]: ## Overall cases for each conditions
```

```
[162]: print("Positive Cases :", positive.iloc[:, -1].sum())
print("Mortality Cases :", mortality.iloc[:, -1].sum())
print("Active Cases :", active.iloc[:, -1].sum())
```

```
Positive Cases : 676570149
Mortality Cases : 6881802
Active Cases : 669688347
```

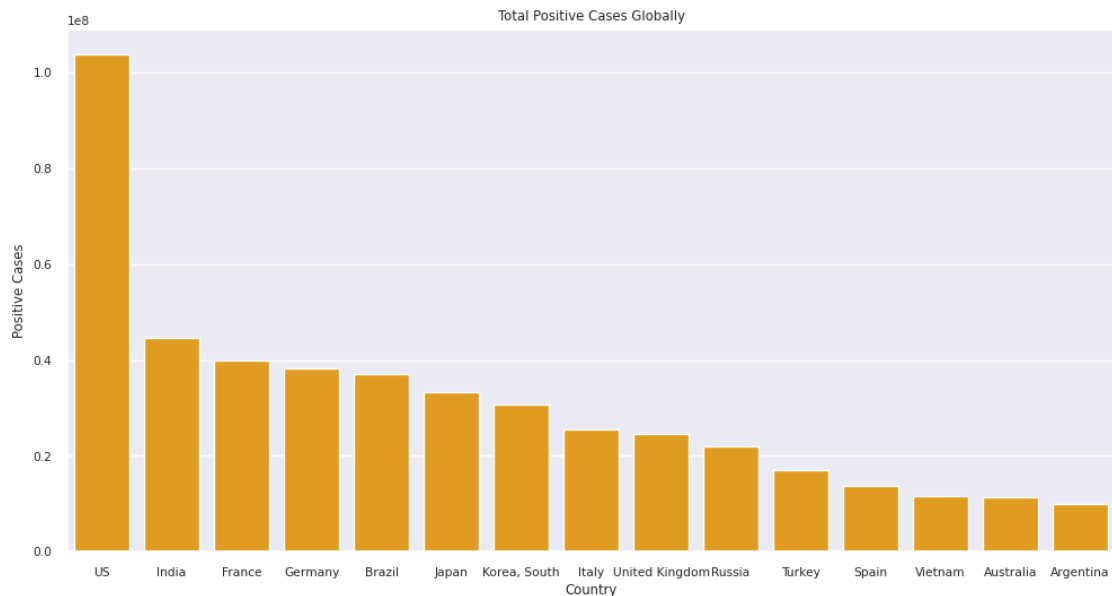
```
[163]: ## Highest records of countries per each conditions.
```

```
[164]: positive_data = positive[['Country', 'Current']].sort_values('Current', ascending_
↪ False)
```

```
mortality_data = mortality[['Country','Current']].
    ↪sort_values('Current',ascending = False)
active_data = active[['Country','Current']].sort_values('Current',ascending =
    ↪False)
```

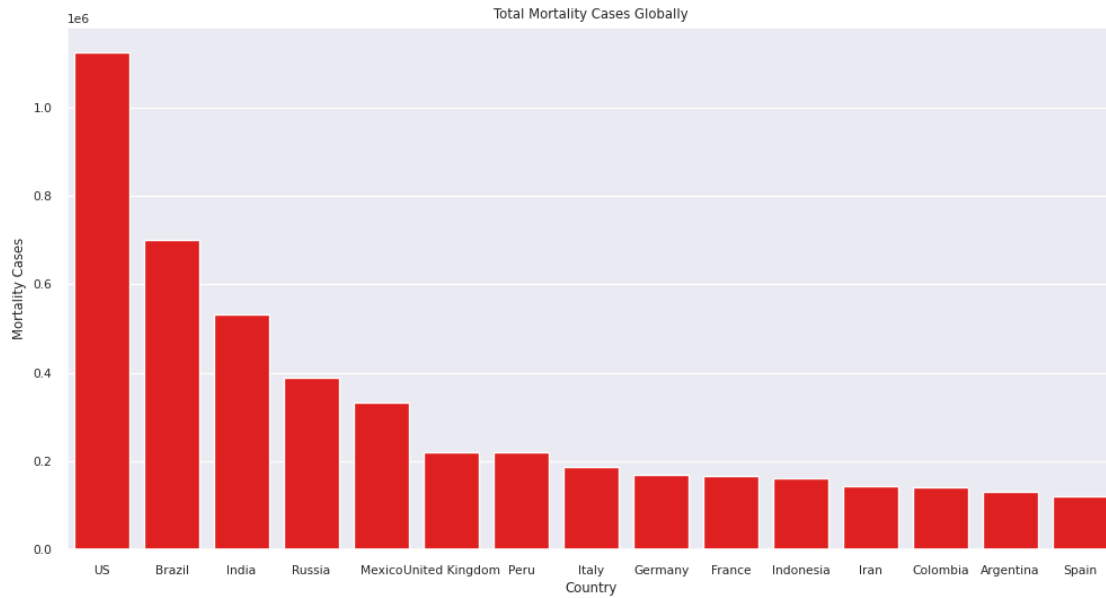
[165]: *### Positive cases*

```
sns.set(font_scale=0.7)
plt.figure(figsize=(12, 6))
fig= sns.barplot(x='Country', y='Current', data=positive_data[:15],
    ↪orient='v',color='Orange')
plt.title('Total Positive Cases Globally')
fig.set(xlabel = 'Country', ylabel = 'Positive Cases')
plt.show()
```



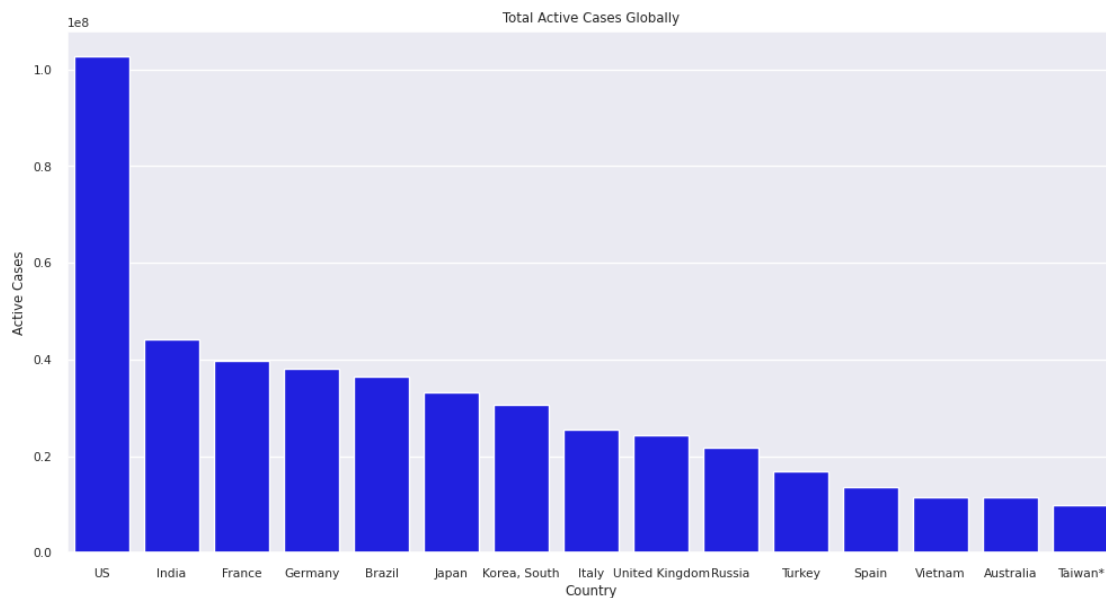
[166]: *### Mortality cases*

```
sns.set(font_scale=0.7)
plt.figure(figsize=(12, 6))
fig= sns.barplot(x='Country', y='Current', data=mortality_data[:15],
    ↪orient='v',color='Red')
plt.title('Total Mortality Cases Globally')
fig.set(xlabel = 'Country', ylabel = 'Mortality Cases')
plt.show()
```



```
[167]: ### Active cases

sns.set(font_scale=0.7)
plt.figure(figsize=(12, 6))
fig= sns.barplot(x='Country', y='Current', data=active_data[:15],
    orient='v',color='blue')
plt.title('Total Active Cases Globally')
fig.set(xlabel = 'Country', ylabel = 'Active Cases')
plt.show()
```



```
[168]: ## Total cases in UK, USA, Australia, India, Brazil, South Africa
```

```
[169]: ### Total positive cases in UK, USA, Australia, India, Brazil, South Africa
```

```
usa_positive = positive[positive.Country == 'US'].iloc[:,1:].sum().values.  
    ↪tolist()  
uk_positive = positive[positive.Country == 'United Kingdom'].iloc[:,1:].sum().  
    ↪values.tolist()  
australia_positive = positive[positive.Country == 'Australria'].iloc[:,1:].sum().  
    ↪values.tolist()  
india_positive = positive[positive.Country == 'India'].iloc[:,1:].sum().values.  
    ↪tolist()  
sa_positive = positive[positive.Country == 'South Africa'].iloc[:,1:].sum().  
    ↪values.tolist()  
brazil_positive = positive[positive.Country == 'Brazil'].iloc[:,1:].sum().  
    ↪values.tolist()
```

```
### Total mortality cases in UK, USA, Australia, India, Brazil, South Africa
```

```
usa_mortality = mortality[mortality.Country == 'US'].iloc[:,1:].sum().values.  
    ↪tolist()  
uk_mortality = mortality[mortality.Country == 'United Kingdom'].iloc[:,1:].  
    ↪sum().values.tolist()  
australia_mortality = mortality[mortality.Country == 'Australria'].iloc[:,1:].  
    ↪sum().values.tolist()  
india_mortality = mortality[mortality.Country == 'India'].iloc[:,1:].sum().  
    ↪values.tolist()  
sa_mortality = mortality[mortality.Country == 'South Africa'].iloc[:,1:].sum().  
    ↪values.tolist()  
brazil_mortality = mortality[mortality.Country == 'Brazil'].iloc[:,1:].sum().  
    ↪values.tolist()
```

```
### Total active cases in UK, USA, Australia, India, Brazil, South Africa
```

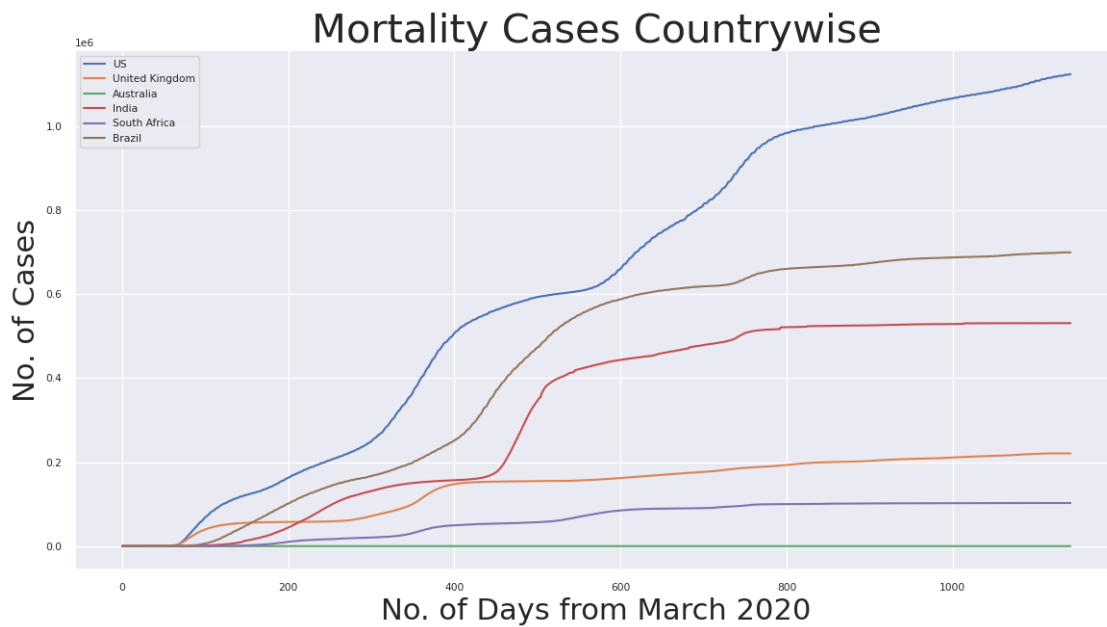
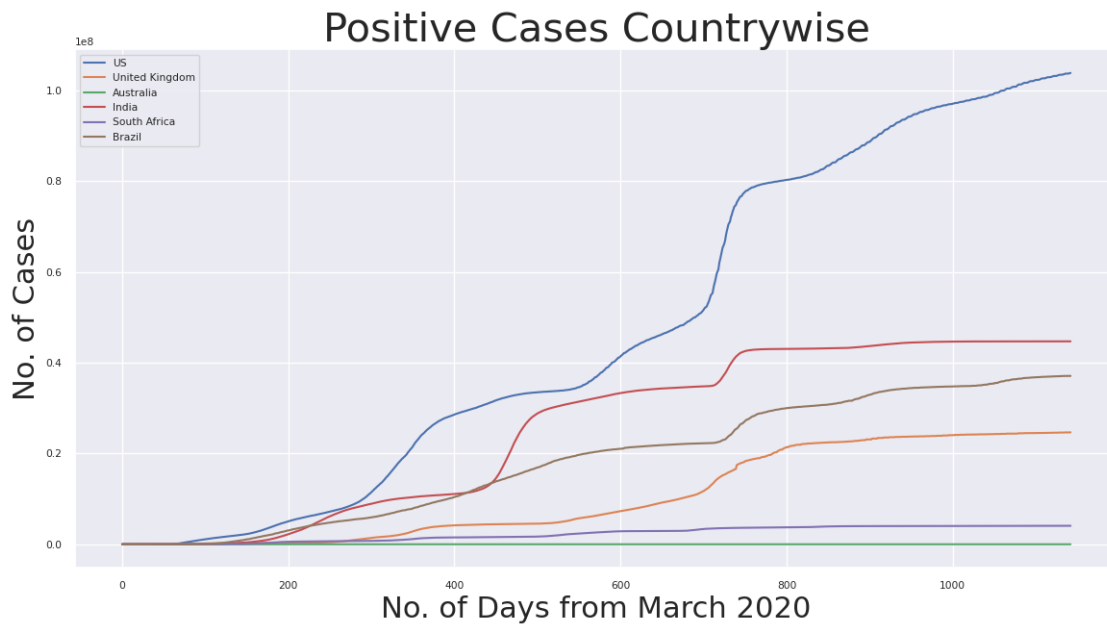
```
usa_active = active[active.Country == 'US'].iloc[:,1:].sum().values.tolist()  
uk_active = active[active.Country == 'United Kingdom'].iloc[:,1:].sum().values.  
    ↪tolist()  
australia_active = active[active.Country == 'Australria'].iloc[:,1:].sum().values.  
    ↪tolist()  
india_active = active[active.Country == 'India'].iloc[:,1:].sum().values.  
    ↪tolist()  
sa_active = active[active.Country == 'South Africa'].iloc[:,1:].sum().values.  
    ↪tolist()
```

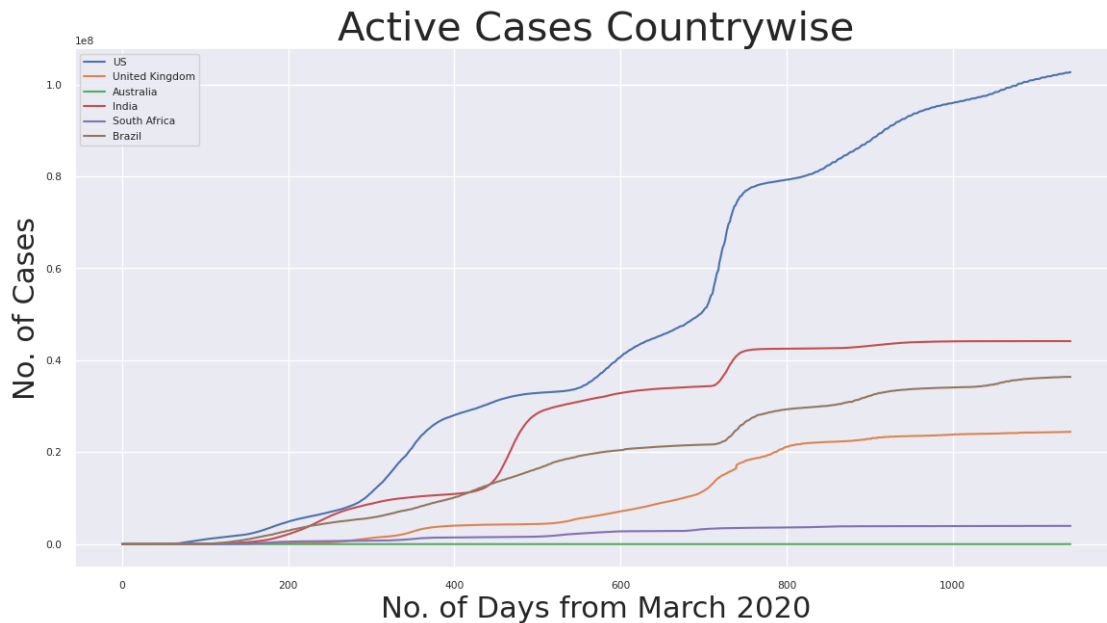
```
brazil_active = active[active.Country == 'Brazil'].iloc[:,1:].sum().values.  
↳tolist()
```

```
[170]: ### Positive cases  
  
plt.figure(figsize=(14,7))  
plt.plot(usa_positive)  
plt.plot(uk_positive)  
plt.plot(australia_positive)  
plt.plot(india_positive)  
plt.plot(sa_positive)  
plt.plot(brazil_positive)  
  
plt.title('Positive Cases Countrywise', size=30)  
plt.xlabel('No. of Days from March 2020', size=22)  
plt.ylabel('No. of Cases', size=22)  
plt.legend(['US', 'United Kingdom', 'Australia', 'India', 'South Africa', 'Brazil'])  
plt.show()  
  
### Mortality cases  
  
plt.figure(figsize=(14,7))  
plt.plot(usa_mortality)  
plt.plot(uk_mortality)  
plt.plot(australia_mortality)  
plt.plot(india_mortality)  
plt.plot(sa_mortality)  
plt.plot(brazil_mortality)  
  
plt.title('Mortality Cases Countrywise', size=30)  
plt.xlabel('No. of Days from March 2020', size=22)  
plt.ylabel('No. of Cases', size=22)  
plt.legend(['US', 'United Kingdom', 'Australia', 'India', 'South Africa', 'Brazil'])  
plt.show()  
  
### Active cases  
  
plt.figure(figsize=(14,7))  
plt.plot(usa_active)  
plt.plot(uk_active)  
plt.plot(australia_active)  
plt.plot(india_active)  
plt.plot(sa_active)  
plt.plot(brazil_active)  
  
plt.title('Active Cases Countrywise', size=30)  
plt.xlabel('No. of Days from March 2020', size=22)
```



```
plt.ylabel('No. of Cases', size=22)
plt.legend(['US', 'United Kingdom', 'Australia', 'India', 'South Africa', 'Brazil'])
plt.show()
```





[171]: # Initial Results:

```
## Initailly the number of positive cases was increasing each month for all the
↳ countries, following the identified number came to a constant rate since two
↳ years from the outbreak, except the United States.
## The mortality and active state of cases follwed a similar patter across the
↳ countries.
## Australia was identified to have an extremly low rates with positive,
↳ mortality and active cases untill now, having US the quite opposite with
↳ hisghest rates.
```

[172]: ### The rate of each conditions since March 2020

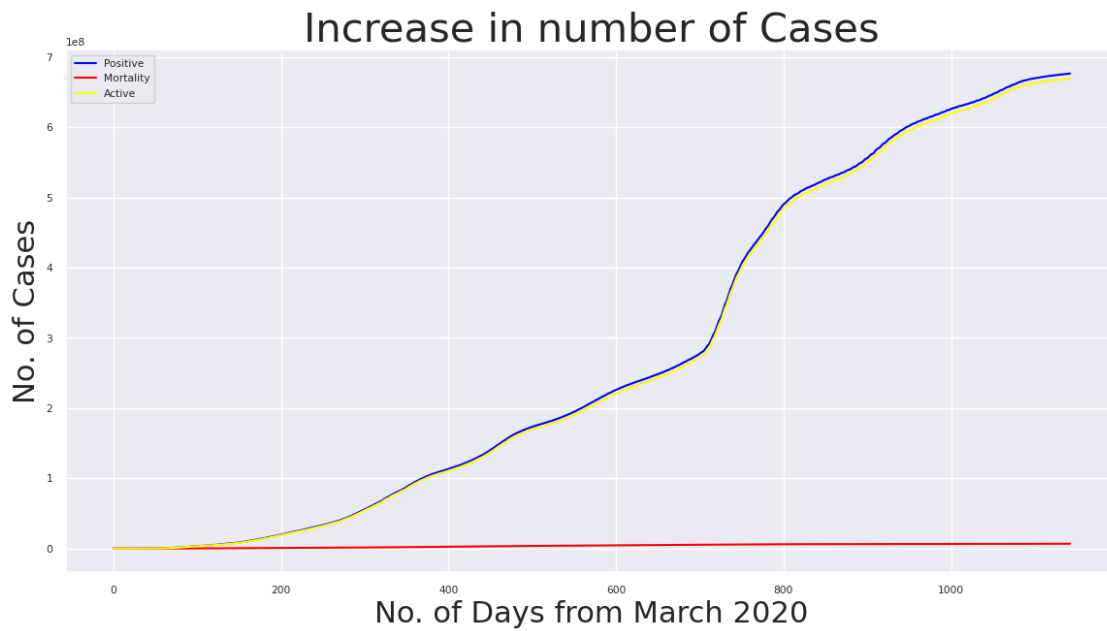
```
positive_date = positive.iloc[:,1:].sum().values.tolist()
mortality_date = mortality.iloc[:,1:].sum().values.tolist()
active_date = active.iloc[:,1:].sum().values.tolist()
```

[173]:

```
plt.figure(figsize=(14,7))
plt.plot(positive_date,color='Blue')
plt.plot(mortality_date,color='Red')
plt.plot(active_date,color='Yellow')

plt.xlabel('No. of Days from March 2020',size=22)
plt.ylabel('No. of Cases',size=22)
plt.title('Increase in number of Cases',size=30)
plt.legend(['Positive','Mortality','Active'])
```

```
plt.show()
```



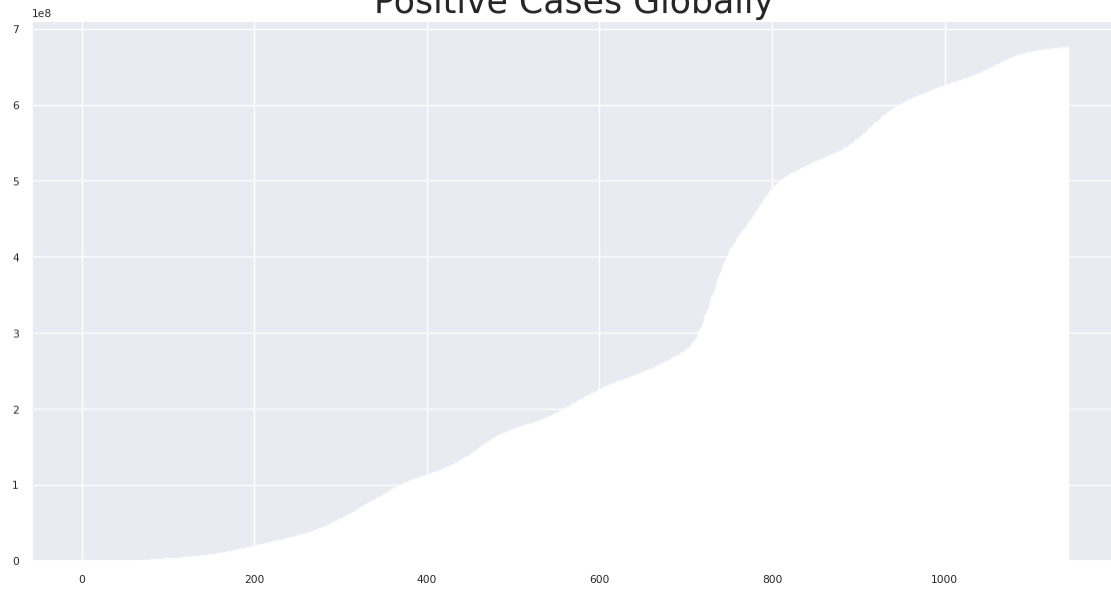
```
[174]: days = [ i for i in range(positive.shape[1] - 1) ]

plt.figure(figsize=(14,7))
plt.bar(days,positive_date,color='Blue')
plt.title('Positive Cases Globally',size=25)
plt.show()

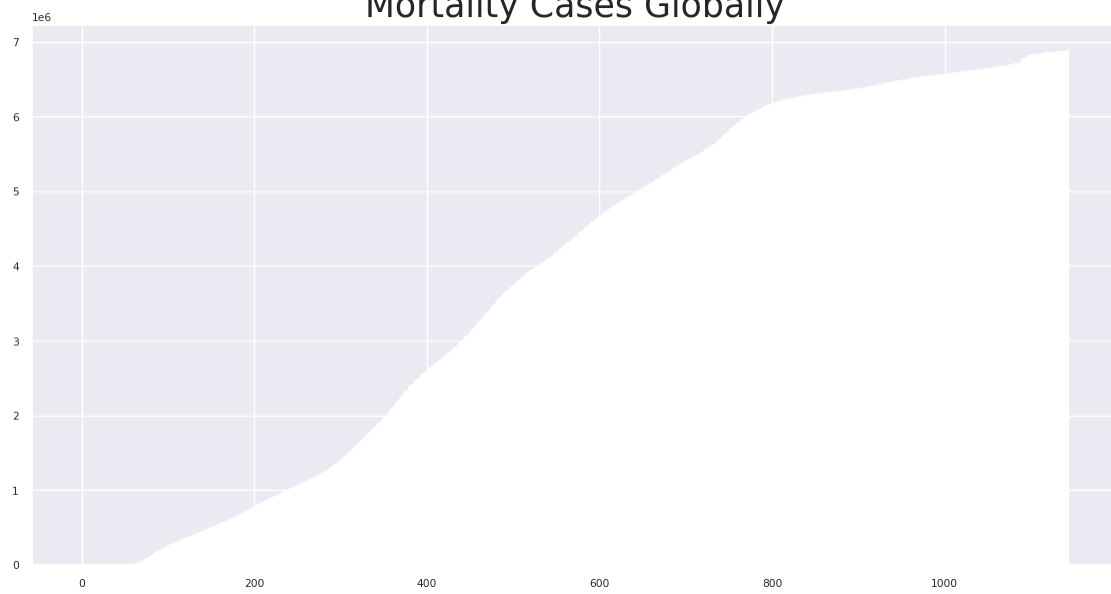
plt.figure(figsize=(14,7))
plt.bar(days,mortality_date,color='Red')
plt.title('Mortality Cases Globally',size=25)
plt.show()

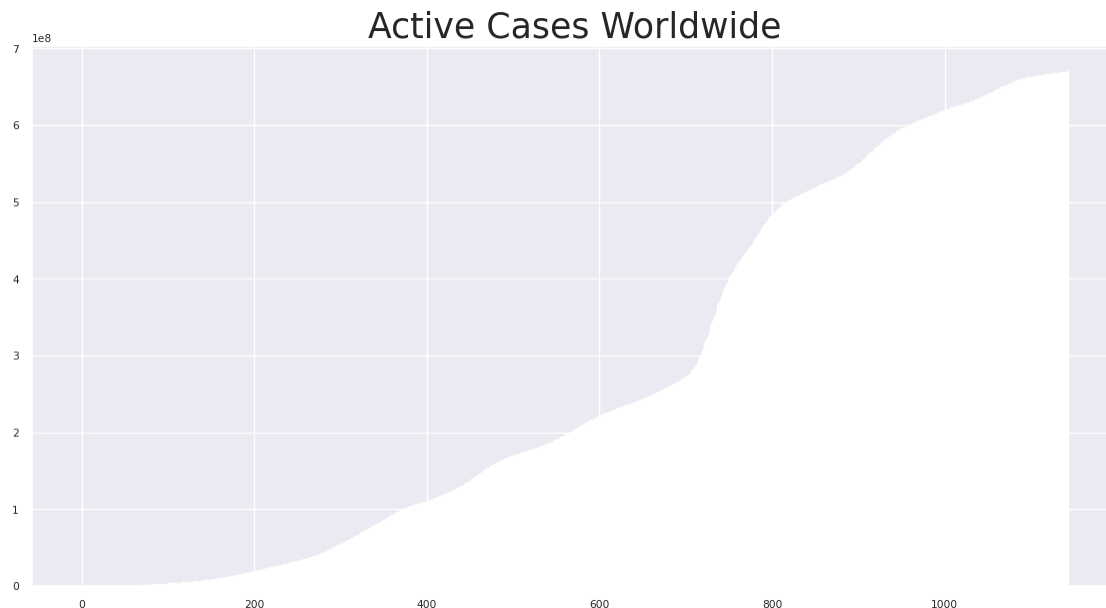
plt.figure(figsize=(14,7))
plt.bar(days,active_date,color='Yellow')
plt.title('Active Cases Worldwide',size=25)
plt.show()
```

Positive Cases Globally



Mortality Cases Globally





```
[175]: # Initial results:

## An exponential growth in positive cases was identified on third month from
↳ the outbreak, having a spike after second year.
## There was a gradual increase in mortality rate from second month onwards.
## A similar path was identified with active cases as covid positive.
```

```
[176]: # The distribution of cases nationwide:
```

```
[177]: ### Positive cases

fig = px.choropleth(positive_data, locations="Country", locationmode='country_
↳ names',
                    color=positive_data['Current'], hover_name="Country",
                    title='Countries with Positive Cases', hover_data=['Current'],
↳ color_continuous_scale="blues")
fig.show()

### Mortality cases

fig = px.choropleth(mortality_data, locations="Country", locationmode='country_
↳ names',
                    color=mortality_data['Current'], hover_name="Country",
                    title='Countries with Mortality_
↳ Cases', hover_data=['Current'], color_continuous_scale="reds")
fig.show()
```

```
### Active cases
```

```
fig = px.choropleth(active_data, locations="Country", locationmode='country_↵  
↵names',  
                    color=active_data['Current'], hover_name="Country",  
                    title='Countries with Active Cases',hover_data=['Current'],↵  
↵color_continuous_scale="ylgn")  
fig.show()
```

Countries with Positive Cases



Countries with Mortality Cases



Countries with Active Cases



```
[178]: # Linear Regression
```

```
[179]: total_positive = np.array(positive_date).reshape(-1,1)
total_mortality = np.array(mortality_date).reshape(-1,1)
total_active = np.array(active_date).reshape(-1,1)
dates = np.array([i for i in range(len(days))]).reshape(-1, 1)
```

```
[180]: ## Linear Graph
```

```
def linear_plot(x,y,reg,title):
    plt.figure(figsize=(14,7))
    plt.scatter(x,y,color='blue')
    plt.plot(x,reg)
    plt.title(title)
```

```
[181]: X_train_positive, X_test_positive, y_train_positive, y_test_positive =
    ↪train_test_split(dates[100:], total_positive[100:], test_size=0.14,
    ↪shuffle=False)
```

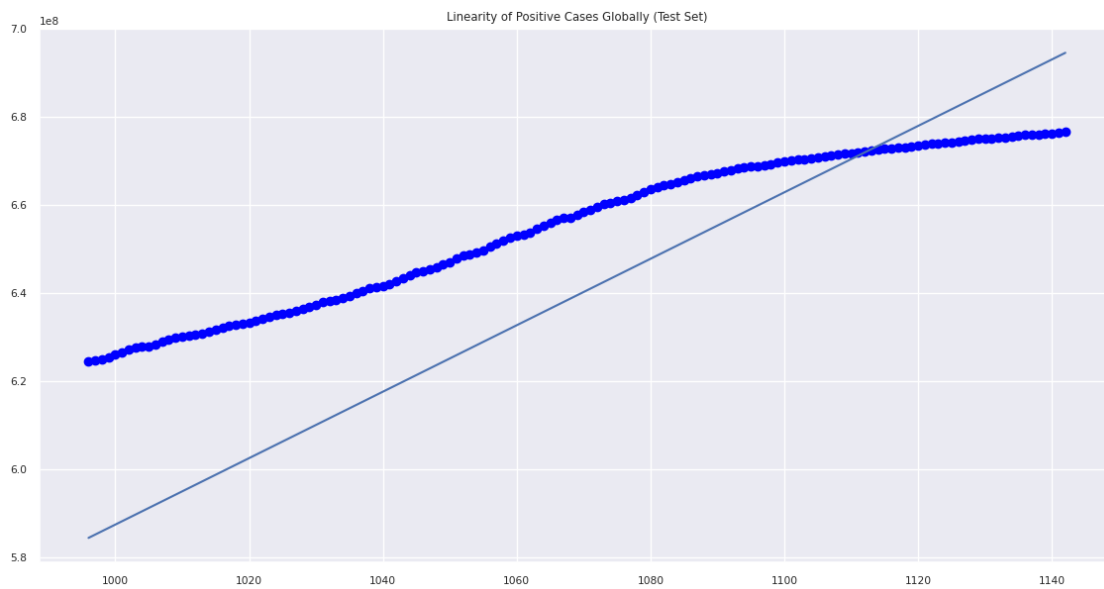
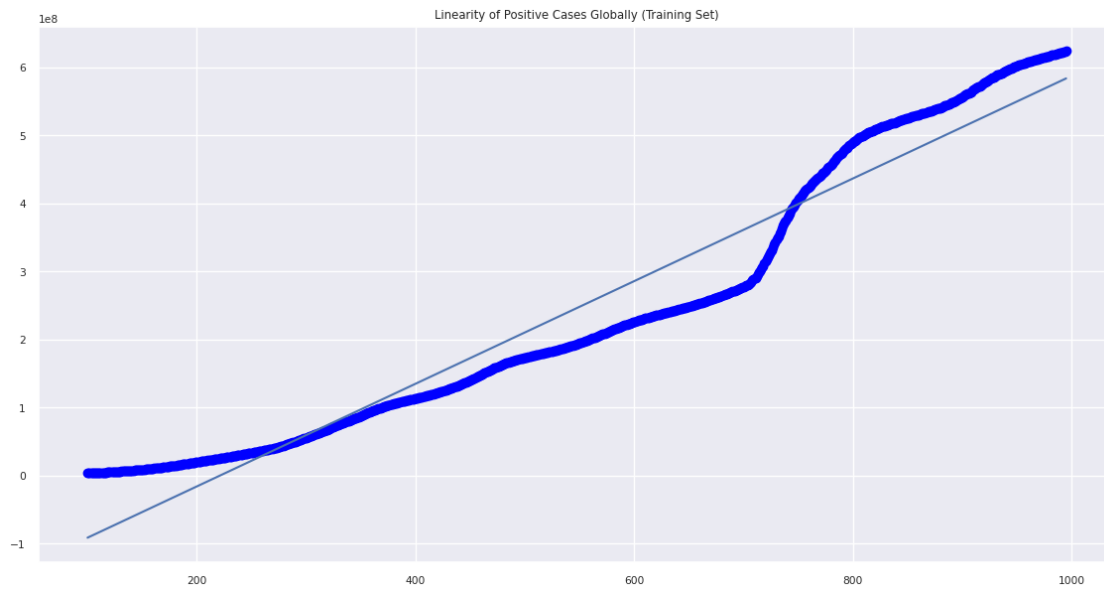
```
[182]: reg = LinearRegression()
reg.fit(X_train_positive, y_train_positive);
```

```
[183]: ## Plot training set
```

```
linear_plot(X_train_positive,y_train_positive,reg,
    ↪predict(X_train_positive),'Linearity of Positive Cases Globally (Training_
    ↪Set)')

## Plot test set
```

```
linear_plot(X_test_positive,y_test_positive,reg.  
↪predict(X_test_positive),'Linearity of Positive Cases Globally (Test Set)')
```



[184]: *### The test set predictions are not similar as the training set predictions. ↪
↪As the total positive cases irregular curves, polynomial regression model was ↪
↪approached.*

[185]: *# Polynomial Regression*


```
[186]: ## Positive Cases

#### Data transformation

poly = PolynomialFeatures(degree=5)
poly_X_train_positive = poly.fit_transform(X_train_positive)
poly_X_test_positive = poly.fit_transform(X_test_positive)
```

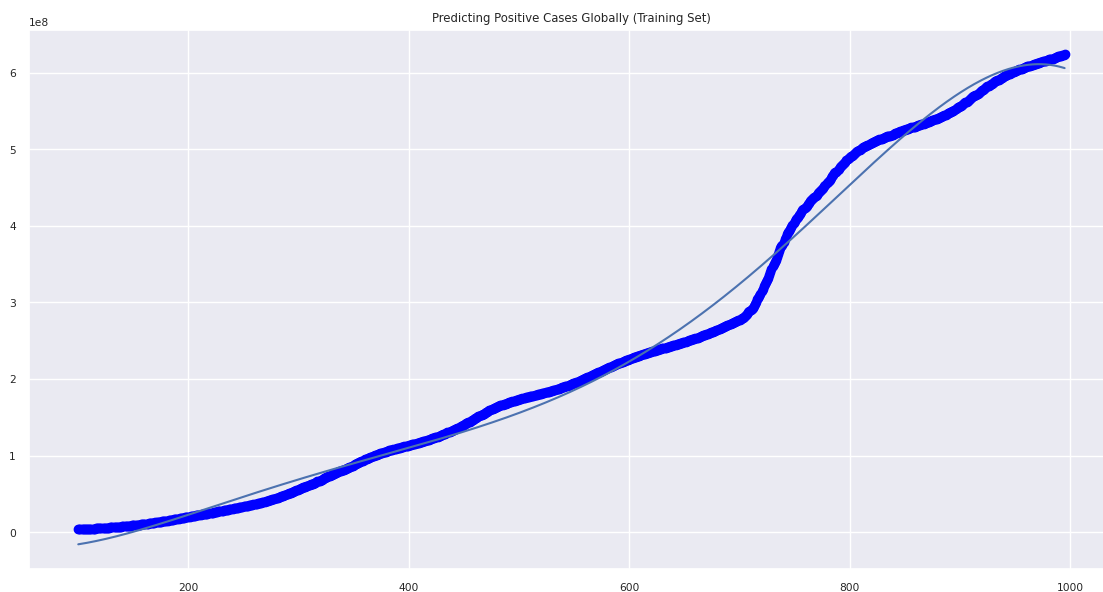
```
[187]: ### polynomial regression for positive cases

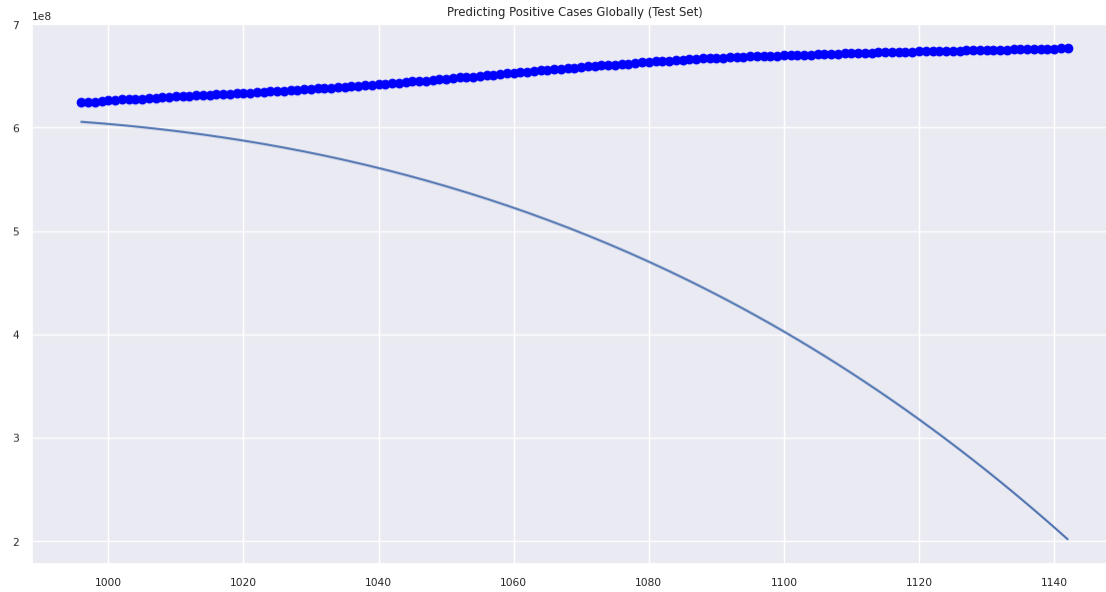
poly_reg = LinearRegression(fit_intercept=False)
poly_reg.fit(poly_X_train_positive, y_train_positive)
```

```
[187]: LinearRegression(fit_intercept=False)
```

```
[188]: ### Plot training set
linear_plot(X_train_positive,y_train_positive,poly_reg.
↳predict(poly_X_train_positive),'Predicting Positive Cases Globally (Training_
↳Set)')

#### Plot test set
linear_plot(X_test_positive,y_test_positive,poly_reg.
↳predict(poly_X_test_positive),'Predicting Positive Cases Globally (Test_
↳Set)')
```

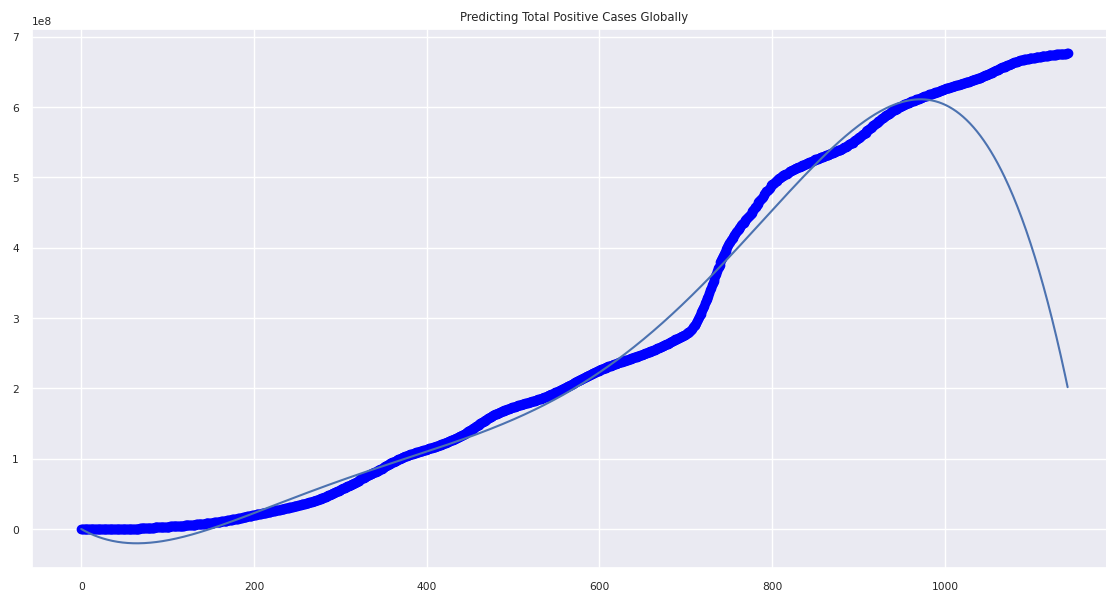




[189]: *### Training set curve showed similar prediction*

[190]: *### Plot of total cases*

```
linear_plot(dates,total_positive,poly_reg.predict(poly.
↪fit_transform(dates)), 'Predicting Total Positive Cases Globally')
```



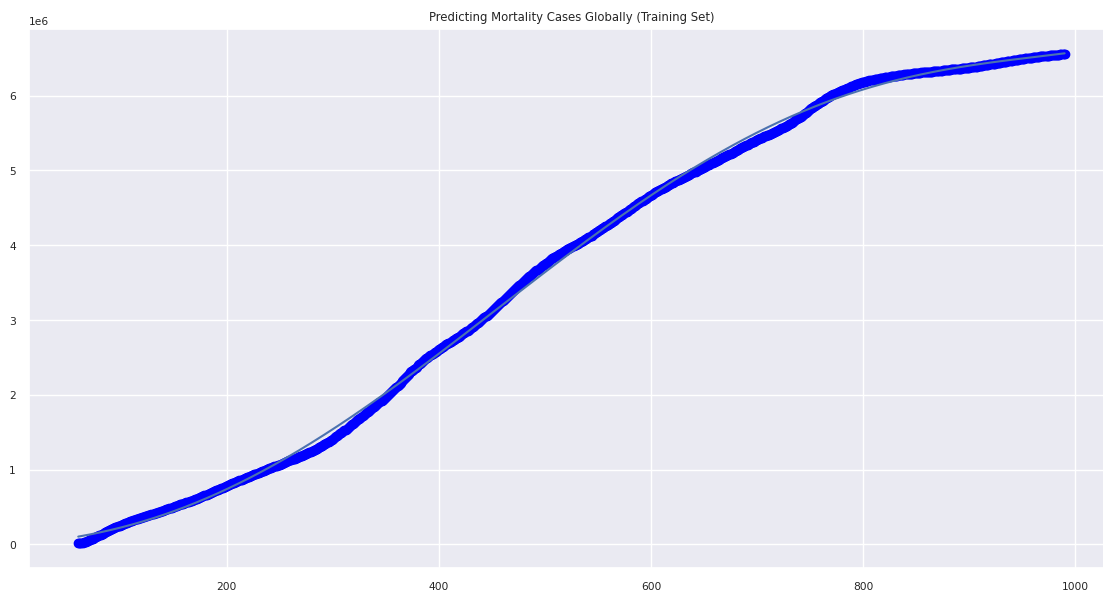
[191]: *## Mortality cases*

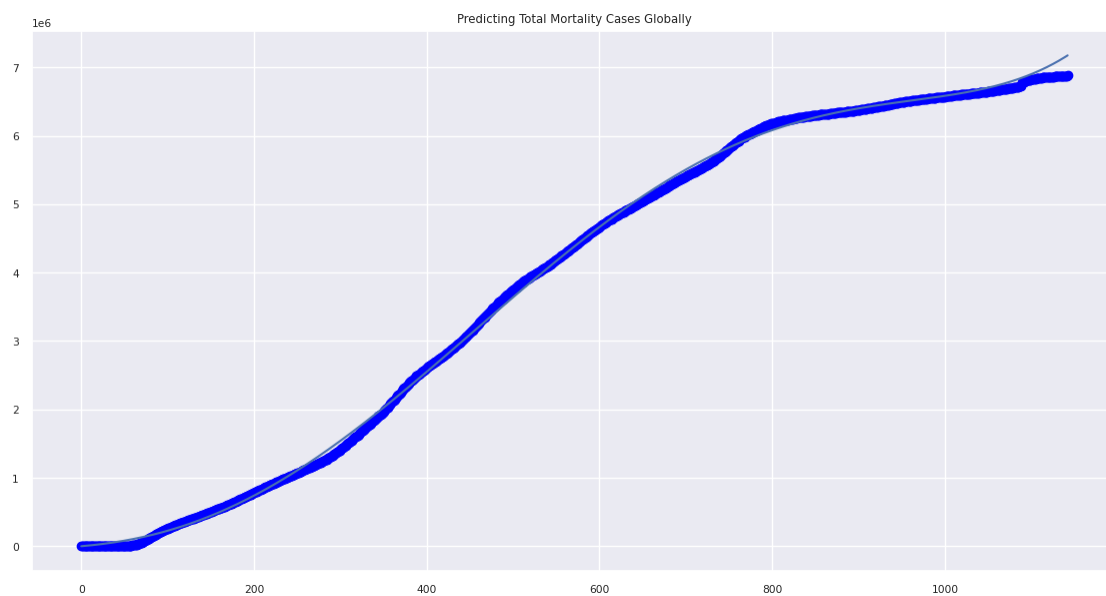
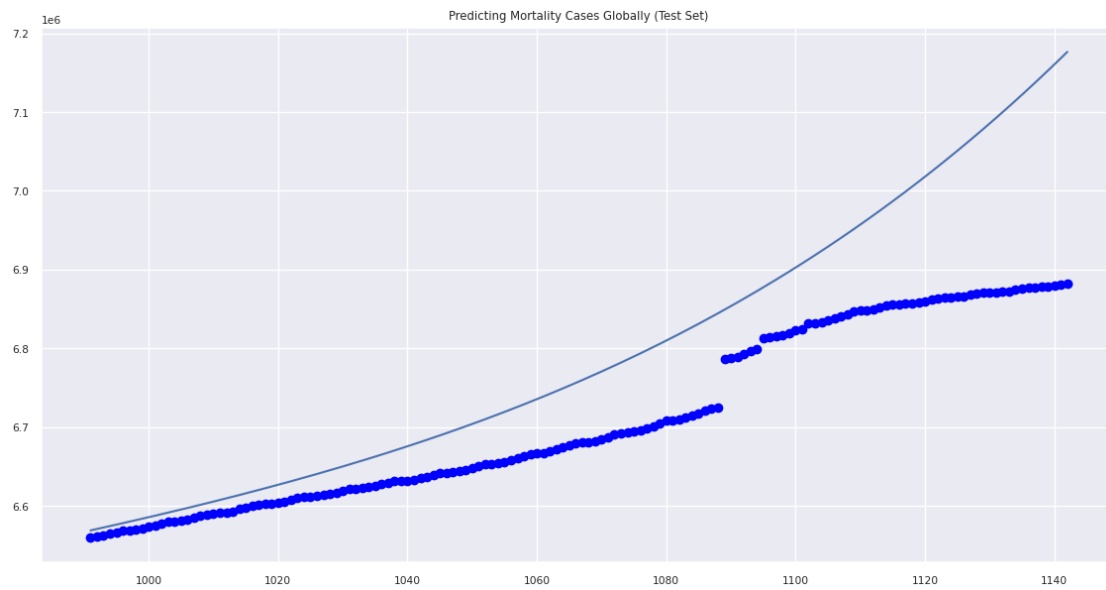
```
[192]: X_train_mortality, X_test_mortality, y_train_mortality, y_test_mortality =  
↳train_test_split(dates[60:], total_mortality[60:], test_size=0.14,  
↳shuffle=False)
```

```
[193]: ### transform our data for polynomial regression  
poly = PolynomialFeatures(degree=5)  
poly_X_train_mortality = poly.fit_transform(X_train_mortality)  
poly_X_test_mortality = poly.fit_transform(X_test_mortality)  
  
### polynomial regression  
poly_reg = LinearRegression(fit_intercept=False)  
poly_reg.fit(poly_X_train_mortality, y_train_mortality)
```

```
[193]: LinearRegression(fit_intercept=False)
```

```
[194]: # Plot training set  
linear_plot(X_train_mortality,y_train_mortality,poly_reg.  
↳predict(poly_X_train_mortality),'Predicting Mortality Cases Globally'  
↳(Training Set)')  
  
# Plot test set  
linear_plot(X_test_mortality,y_test_mortality,poly_reg.  
↳predict(poly_X_test_mortality),'Predicting Mortality Cases Globally (Test'  
↳Set)')  
  
# Plot total cases  
linear_plot(dates,total_mortality,poly_reg.predict(poly.  
↳fit_transform(dates)),'Predicting Total Mortality Cases Globally')
```





```
[195]: ## The mortality rate predictions showed a similar trend throughout the duration.
```

```
[ ]: # Result:
    ---
```

```

* The data sets of confirmed, death, and recovered cases comprised 298, 289 and
  ↳274 entries of countries with 1147 values describing the nation, region and
  ↳cases from 22nd January 2020 to 9th September 2023.
* The total cases identified were 676570149 confirmed, 6881802 deaths and
  ↳669688347 recovered.
* The most confirmed cases were observed in US, India, France, Germany, and
  ↳Brazil; death cases in US, Brazil, India, Russia and Mexico; recovered cases
  ↳in US, India, France, Germany and Brazil.
* Initially, confirmed cases increased monthly for all the countries (US,
  ↳India, Australia, UK, Brazil and South Africa).
* A constant rate was identified two years after the first case, which can
  ↳result from global regulations under quarantine, safety protocols and
  ↳vaccination, except for the United States.
* A similar fashion of death and recovery rates was observed across the five
  ↳nations; uniquely, Australia showed a meagre rate with all conditions,
  ↳whereas the US had the highest rates.
* An exponential growth in confirmed cases was identified after the third
  ↳month, having a spike after the second year, leading to a gradual increase
  ↳in death rates globally from the second month onwards.
---

```

```
[ ]: # Referece
```

```

---
1.      This data set is licensed under the Creative Commons Attribution 4.0
  ↳International (CC BY 4.0) by the Johns Hopkins University on behalf of its
  ↳Center for Systems Science in Engineering. Copyright Johns Hopkins
  ↳University 2020.

2.      Attribute the data as the "COVID-19 Data Repository by the Center for
  ↳Systems Science and Engineering (CSSE) at Johns Hopkins University" or "JHU
  ↳CSSE COVID-19 Data" for short, and the url: https://github.com/
  ↳CSSEGISandData/COVID-19.

3.      For publications that use the data, please cite the following
  ↳publication: "Dong E, Du H, Gardner L. An interactive web-based dashboard to
  ↳track COVID-19 in real time. Lancet Inf Dis. 20(5):533-534. doi: 10.1016/
  ↳S1473-3099(20)30120-1"
-

```