

## Importing required Libraries

```
import pandas as pd
import numpy as np
import seaborn as sns
import matplotlib.pyplot as plt
import plotly.express as px
import plotly.graph_objs as go
from plotly.subplots import make_subplots
from plotly.offline import plot, iplot, init_notebook_mode
init_notebook_mode(connected=True)

corona_dataset_csv = pd.read_csv('covid19_Confirmed_dataset.csv')
corona_dataset_csv.head(10)
```

Province/State	Country/Region	Lat
NaN	Afghanistan	33.0000
NaN	Albania	41.1533
NaN	Algeria	28.0339
NaN	Andorra	42.5063
NaN	Angola	-11.2027
NaN	Antigua and Barbuda	17.0608
NaN	Argentina	-38.4161
NaN	Armenia	40.0691
Australian Capital Territory	Australia	-35.4735
New South Wales	Australia	-33.8688

4	0	0	0	0	0	0	...	24
5	0	0	0	0	0	0	...	23
6	0	0	0	0	0	0	...	3031
7	0	0	0	0	0	0	...	1401
8	0	0	0	0	0	0	...	104
9	0	0	0	0	3	4	...	2969

	4/22/20	4/23/20	4/24/20	4/25/20	4/26/20	4/27/20	4/28/20
4/29/20 \							
0	1176	1279	1351	1463	1531	1703	1828
1939							
1	634	663	678	712	726	736	750
766							
2	2910	3007	3127	3256	3382	3517	3649
3848							
3	723	723	731	738	738	743	743
743							
4	25	25	25	25	26	27	27
27							
5	24	24	24	24	24	24	24
24							
6	3144	3435	3607	3780	3892	4003	4127
4285							
7	1473	1523	1596	1677	1746	1808	1867
1932							
8	104	104	105	106	106	106	106
106							
9	2971	2976	2982	2994	3002	3004	3016
3016							

	4/30/20
0	2171
1	773
2	4006
3	745
4	27
5	24
6	4428
7	2066
8	106
9	3025



7	0	0	0	0	0	0	...	1401
8	0	0	0	0	0	0	...	104
9	0	0	3	4	4	4	...	2969

	4/22/20	4/23/20	4/24/20	4/25/20	4/26/20	4/27/20	4/28/20
0	1176	1279	1351	1463	1531	1703	1828
1	634	663	678	712	726	736	750
2	2910	3007	3127	3256	3382	3517	3649
3	723	723	731	738	738	743	743
4	25	25	25	25	26	27	27
5	24	24	24	24	24	24	24
6	3144	3435	3607	3780	3892	4003	4127
7	1473	1523	1596	1677	1746	1808	1867
8	104	104	105	106	106	106	106
9	2971	2976	2982	2994	3002	3004	3016

	4/30/20
0	2171
1	773
2	4006
3	745
4	27
5	24
6	4428
7	2066
8	106
9	3025

[10 rows x 102 columns]

# Aggregating the rows with country

```
corona_dataset_aggregated =
corona_dataset_csv.groupby("Country/Region").sum()
```

```
corona_dataset_aggregated.head(10).sum(axis=1)
```

```
Country/Region
Afghanistan      28462
Albania           17864
Algeria           74325
Andorra           21893
Angola            649
Antigua and Barbuda 678
Argentina         84105
Armenia           40610
Australia         224354
Austria           502063
dtype: int64
```

```
corona_dataset_aggregated.shape
```

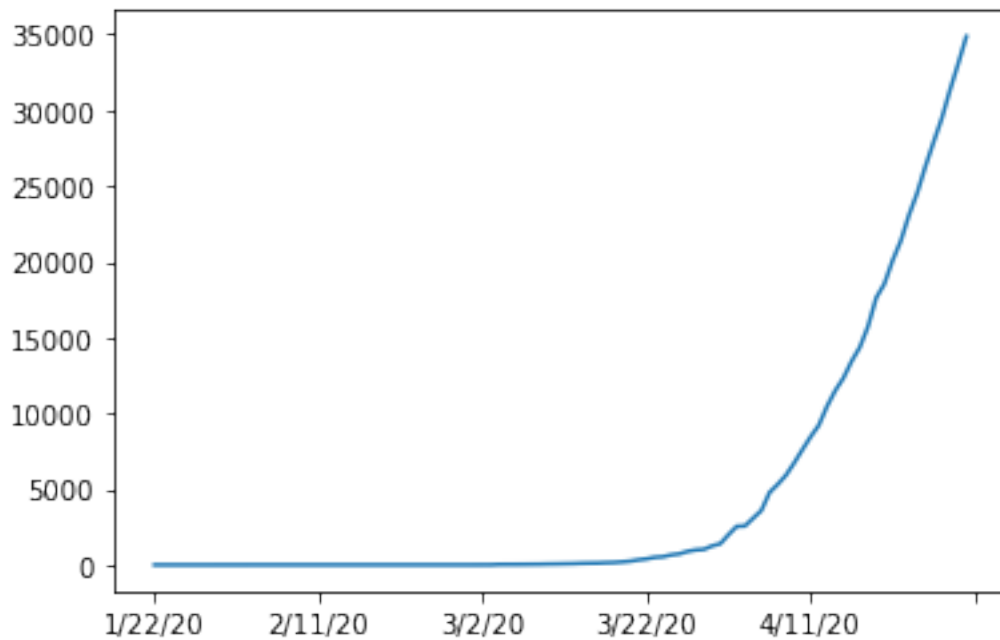
```
(187, 100)
```

```
#from matplotlib.pyplot import figure
#countries=list(corona_dataset_aggregated.index)
#total_cases=corona_dataset_aggregated.sum(axis=1)
#plt.figure(figsize=(100,40),dpi=380)
#plt.xticks(rotation=90,fontsize=25)
#plt.bar(countries,total_cases)
#plt.show()
#figure(num=None, figsize=(8, 6), dpi=80, facecolor='w',
edgecolor='k')
```

## Visualizing data related to a country for example India,China,Spain,Italy

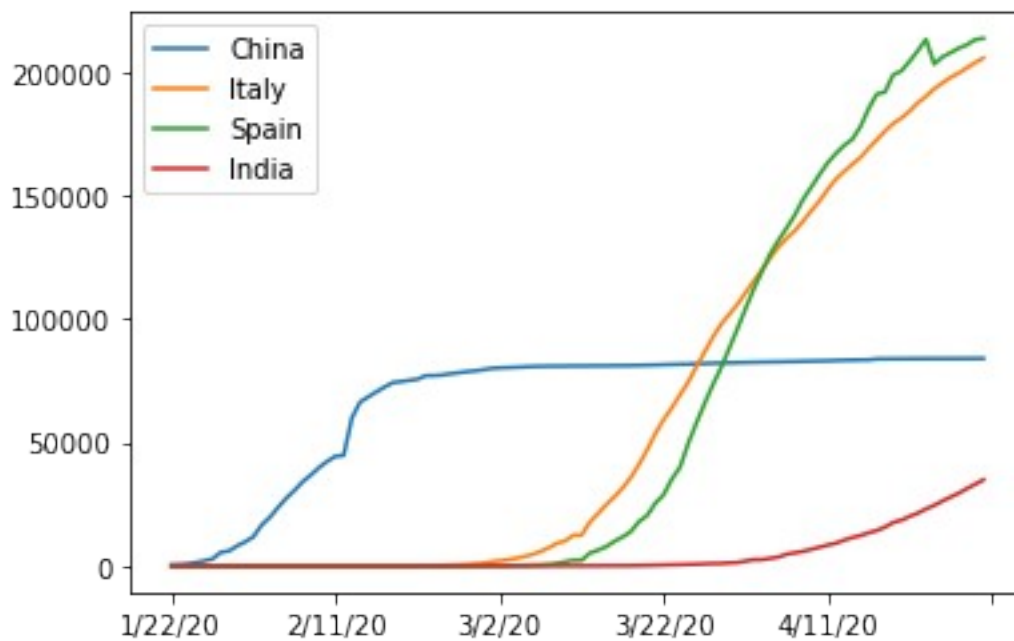
```
corona_dataset_aggregated.loc['India'].plot()
```

```
<AxesSubplot:>
```



```
corona_dataset_aggregated.loc['China'].plot()
corona_dataset_aggregated.loc['Italy'].plot()
corona_dataset_aggregated.loc['Spain'].plot()
corona_dataset_aggregated.loc['India'].plot()
plt.legend()
```

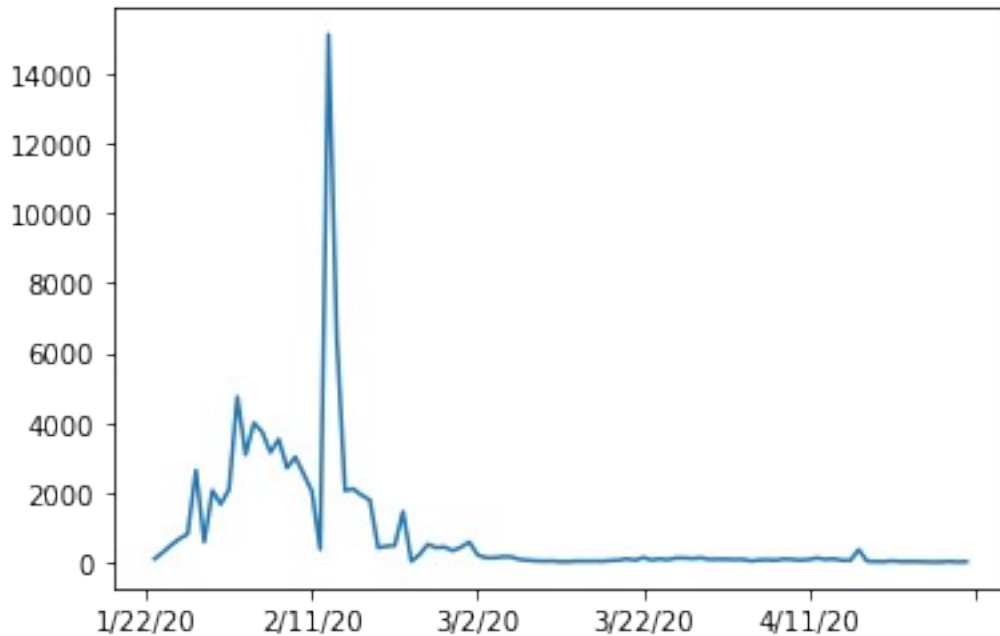
<matplotlib.legend.Legend at 0x1cfd9066310>



TO GET THE RATE AT WHICH CASES ARE GROWING(DIFFERENTIATING THE CURVE)

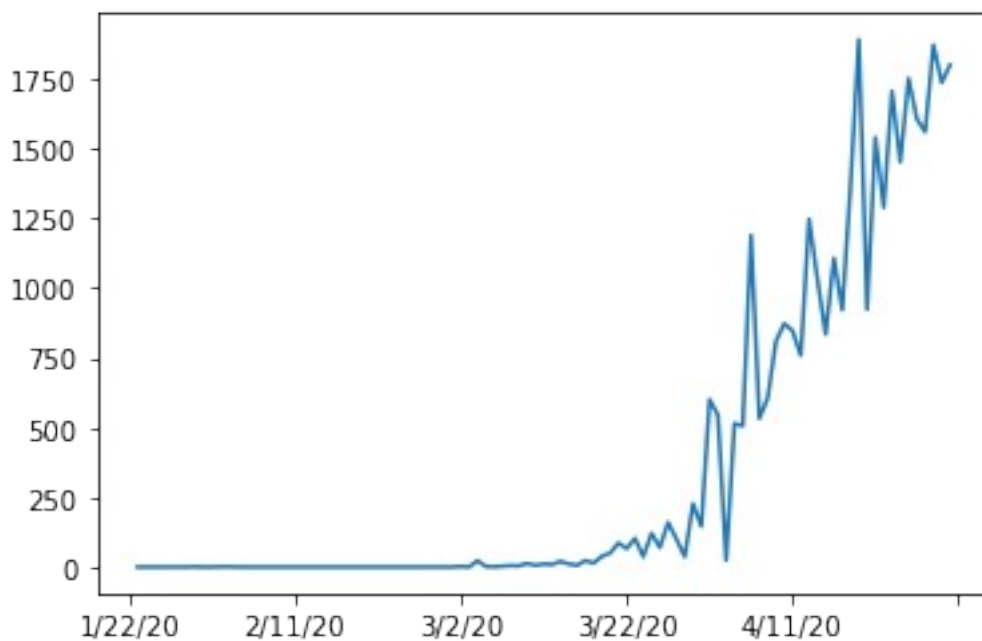
```
corona_dataset_aggregated.loc['China'].diff().plot()
```

<AxesSubplot:>



```
corona_dataset_aggregated.loc['India'].diff().plot()
```

<AxesSubplot:>



### Find max infection rates for China and India

```
corona_dataset_aggregated.loc['China'].diff().max()
```

15136.0

```
corona_dataset_aggregated.loc['India'].diff().max()
```

1893.0

```
corona_dataset_aggregated.loc['India',:]
```

```
1/22/20      0
1/23/20      0
1/24/20      0
1/25/20      0
1/26/20      0
```

```
...
4/26/20    27890
4/27/20    29451
4/28/20    31324
4/29/20    33062
4/30/20    34863
```

Name: India, Length: 100, dtype: int64

### Finding average infection rates

```
corona_dataset_aggregated.loc['China'].diff().mean()
```

842.5050505050505

```
corona_dataset_aggregated.loc['India'].diff().mean()
```

352.1515151515151

### Find maximum infection rate for all of the countries.

```
countries = list(corona_dataset_aggregated.index)
```

```
ave_infection_rates = []
```

```
for country in countries :
```

```
ave_infection_rates.append(corona_dataset_aggregated.loc[country].diff
().mean())
```

```
corona_dataset_aggregated['mean infection rate'] = ave_infection_rates
```

```
corona_dataset_aggregated.head()
```

```
          1/22/20  1/23/20  1/24/20  1/25/20  1/26/20  1/27/20
1/28/20  \
Country/Region
```

```
Afghanistan      0      0      0      0      0      0
0
```



Albania	0	0	0	0	0	0
0						
Algeria	0	0	0	0	0	0
0						
Andorra	0	0	0	0	0	0
0						
Angola	0	0	0	0	0	0
0						

	1/29/20	1/30/20	1/31/20	...	4/22/20	4/23/20
4/24/20 \						
Country/Region				...		

Afghanistan	0	0	0	...	1176	1279
1351						
Albania	0	0	0	...	634	663
678						
Algeria	0	0	0	...	2910	3007
3127						
Andorra	0	0	0	...	723	723
731						
Angola	0	0	0	...	25	25
25						

	4/25/20	4/26/20	4/27/20	4/28/20	4/29/20	
4/30/20 \						
Country/Region						
Afghanistan	1463	1531	1703	1828	1939	2171
Albania	712	726	736	750	766	773
Algeria	3256	3382	3517	3649	3848	4006
Andorra	738	738	743	743	743	745
Angola	25	26	27	27	27	27

	mean infection rate
Country/Region	
Afghanistan	21.929293
Albania	7.808081
Algeria	40.464646
Andorra	7.525253
Angola	0.272727

[5 rows x 101 columns]

## Create a new dataframe with only needed column

```
corona_data = pd.DataFrame(corona_dataset_aggregated['mean infection rate'])
```

```
corona_data.head(10)
```

Country/Region	mean infection rate
Afghanistan	21.929293
Albania	7.808081
Algeria	40.464646
Andorra	7.525253
Angola	0.272727
Antigua and Barbuda	0.242424
Argentina	44.727273
Armenia	20.868687
Australia	68.343434
Austria	156.080808

## Importing the world happiness report

```
world_happiness_report = pd.read_csv("worldwide_happiness_report.csv")  
world_happiness_report.head()
```

	Overall rank	Country or region	Score	GDP per capita	Social support \
0	1	Finland	7.769	1.340	1.587
1	2	Denmark	7.600	1.383	1.573
2	3	Norway	7.554	1.488	1.582
3	4	Iceland	7.494	1.380	1.624
4	5	Netherlands	7.488	1.396	1.522

	Healthy life expectancy	Freedom to make life choices	Generosity \
0	0.986	0.596	0.153
1	0.996	0.592	0.252
2	1.028	0.603	0.271
3	1.026	0.591	0.354
4	0.999	0.557	0.322

```

    Perceptions of corruption
0          0.393
1          0.410
2          0.341
3          0.118
4          0.298

```

```
world_happiness_report.dtypes
```

```

Overall rank          int64
Country or region     object
Score                 float64
GDP per capita         float64
Social support         float64
Healthy life expectancy float64
Freedom to make life choices float64
Generosity             float64
Perceptions of corruption float64
dtype: object

```

```
world_happiness_report.shape
```

```
(156, 9)
```

### Dropping useless Columns

```
columns_to_dropped = ['Overall rank', 'Score', 'Generosity', 'Perceptions
of corruption']
```

```
world_happiness_report.drop(columns_to_dropped,axis=1 , inplace=True)
```

```
world_happiness_report.head()
```

```

    Country or region  GDP per capita  Social support  Healthy life
expectancy \
0          Finland          1.340          1.587
0.986
1          Denmark          1.383          1.573
0.996
2          Norway          1.488          1.582
1.028
3          Iceland          1.380          1.624
1.026
4          Netherlands          1.396          1.522
0.999

```

```

    Freedom to make life choices
0          0.596
1          0.592
2          0.603
3          0.591
4          0.557

```

## Changing the indices of the dataframe

```
world_happiness_report.set_index(['Country or region'],inplace=True)
world_happiness_report.head()
```

Country or region	GDP per capita	Social support	Healthy life expectancy \
Finland	1.340	1.587	0.986
Denmark	1.383	1.573	0.996
Norway	1.488	1.582	1.028
Iceland	1.380	1.624	1.026
Netherlands	1.396	1.522	0.999

Country or region	Freedom to make life choices
Finland	0.596
Denmark	0.592
Norway	0.603
Iceland	0.591
Netherlands	0.557

## Joining the two datasets

```
data = world_happiness_report.join(corona_data,how="inner")
data.head()
```

Country or region	GDP per capita	Social support	Healthy life expectancy \
Finland	1.340	1.587	0.986
Denmark	1.383	1.573	0.996
Norway	1.488	1.582	1.028
Iceland	1.380	1.624	1.026
Netherlands	1.396	1.522	0.999

Country or region	Freedom to make life choices	mean infection rate
Finland	0.596	50.454545
Denmark	0.592	94.505051
Norway	0.603	78.161616

Iceland	0.591	18.151515
Netherlands	0.557	399.111111

### Correlation Matrix

```
data.corr()
```

*# it is representing the correlation between every two columns of our dataset*

	GDP per capita	Social support \
GDP per capita	1.000000	0.759468
Social support	0.759468	1.000000
Healthy life expectancy	0.863062	0.765286
Freedom to make life choices	0.394603	0.456246
mean infection rate	0.329073	0.241618

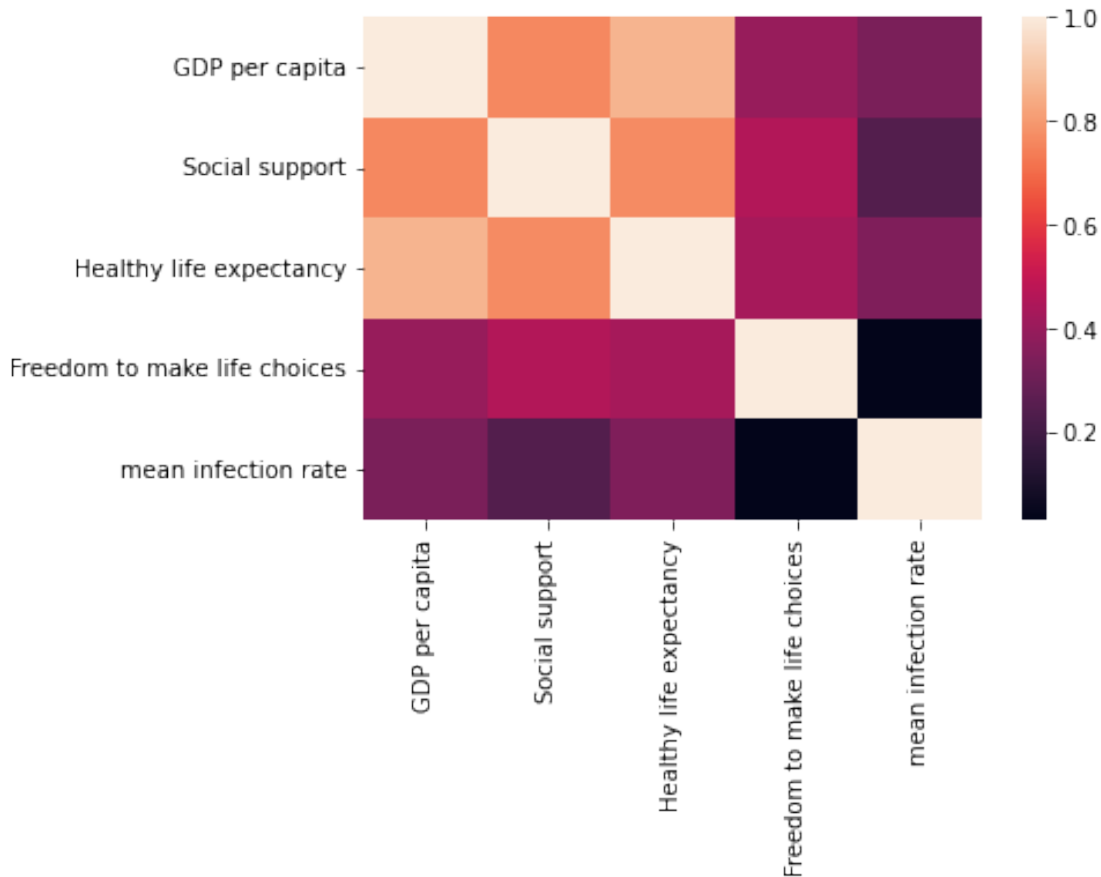
	Healthy life expectancy \
GDP per capita	0.863062
Social support	0.765286
Healthy life expectancy	1.000000
Freedom to make life choices	0.427892
mean infection rate	0.341890

	Freedom to make life choices \
GDP per capita	0.394603
Social support	0.456246
Healthy life expectancy	0.427892
Freedom to make life choices	1.000000
mean infection rate	0.029595

	mean infection rate
GDP per capita	0.329073
Social support	0.241618
Healthy life expectancy	0.341890
Freedom to make life choices	0.029595
mean infection rate	1.000000

```
sns.heatmap(data.corr())
```

<AxesSubplot:>



## Visualization of the results

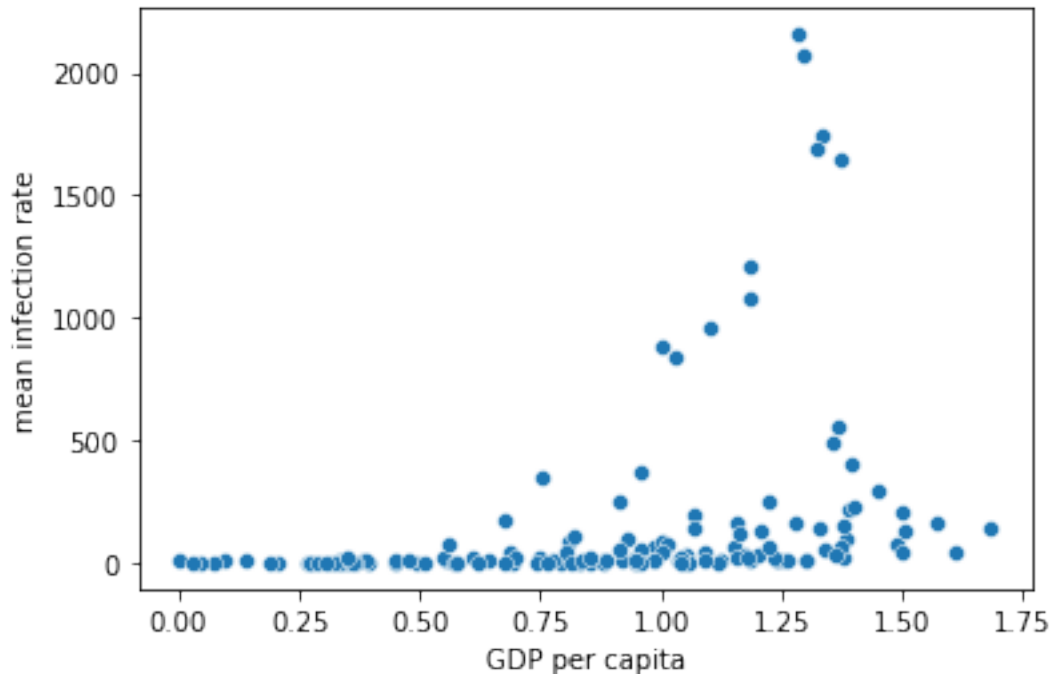
### Plotting GDP vs maximum Infection rate

```
x = data['GDP per capita']
y = data['mean infection rate']
sns.scatterplot(x,y) #VISUALISATION IS DIFFICULT BECAUSE OF
DIFFERENCE Y SCALE AND X-SCALE
```

D:\Users\dell\anaconda3\lib\site-packages\seaborn\\_decorators.py:36:  
FutureWarning:

Pass the following variables as keyword args: x, y. From version 0.12, the only valid positional argument will be `data`, and passing other arguments without an explicit keyword will result in an error or misinterpretation.

```
<AxesSubplot:xlabel='GDP per capita', ylabel='mean infection rate'>
```

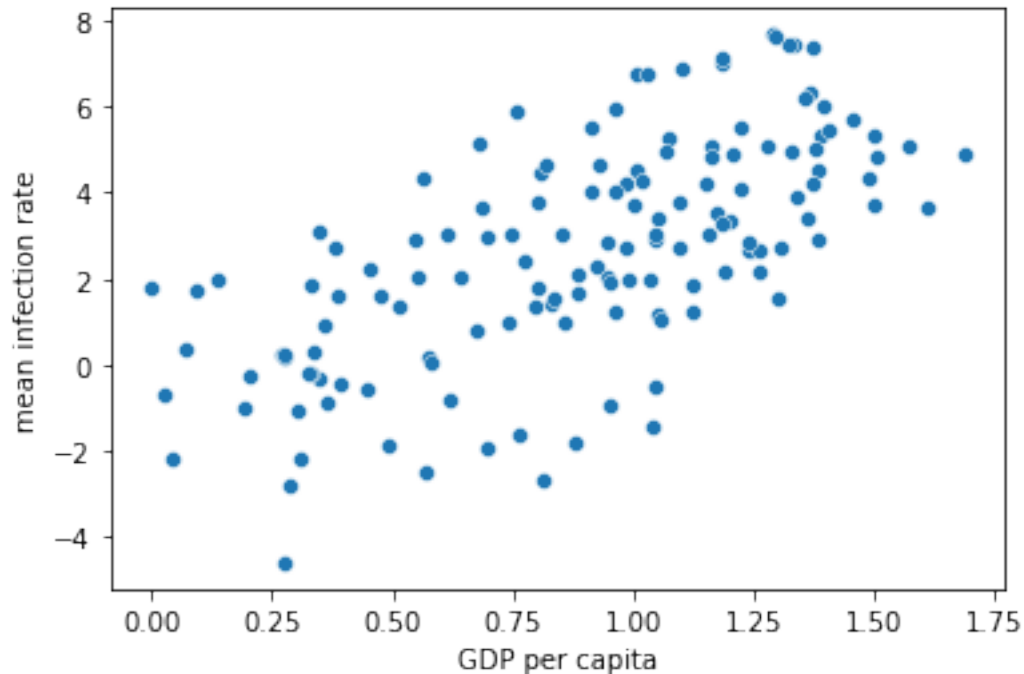


```
x = data['GDP per capita']  
y = data['mean infection rate']  
sns.scatterplot(x,np.log(y))
```

D:\Users\dell\anaconda3\lib\site-packages\seaborn\\_decorators.py:36:  
FutureWarning:

Pass the following variables as keyword args: x, y. From version 0.12, the only valid positional argument will be `data`, and passing other arguments without an explicit keyword will result in an error or misinterpretation.

<AxesSubplot:xlabel='GDP per capita', ylabel='mean infection rate'>



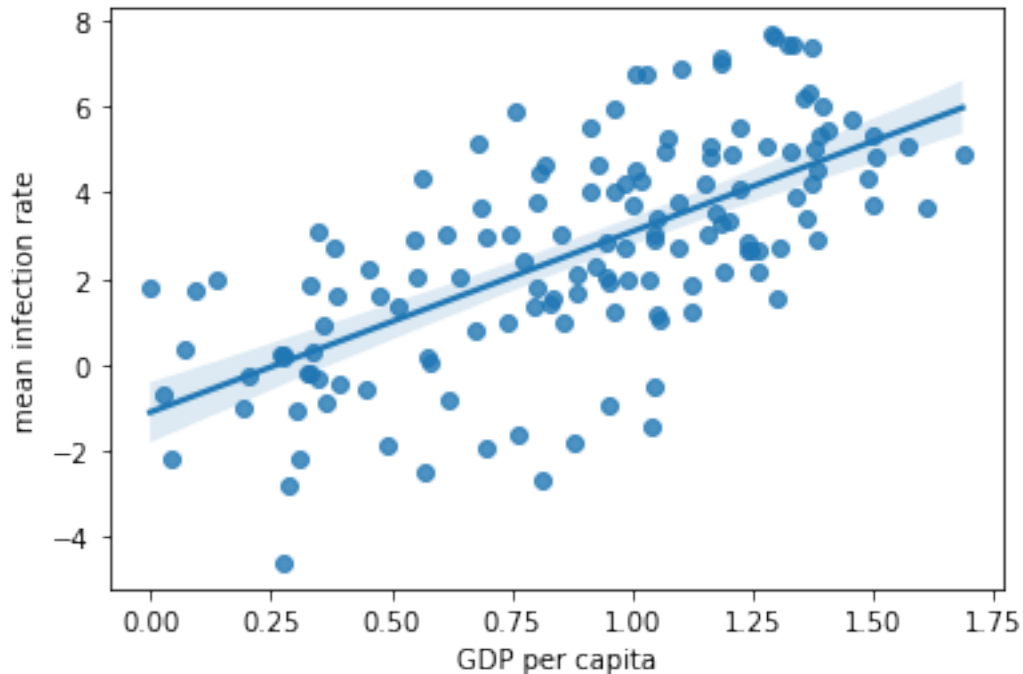
```
sns.regplot(x,np.log(y))
```

D:\Users\dell\anaconda3\lib\site-packages\seaborn\\_decorators.py:36:  
FutureWarning:

Pass the following variables as keyword args: x, y. From version 0.12, the only valid positional argument will be `data`, and passing other arguments without an explicit keyword will result in an error or misinterpretation.

```
<AxesSubplot:xlabel='GDP per capita', ylabel='mean infection rate'>
```





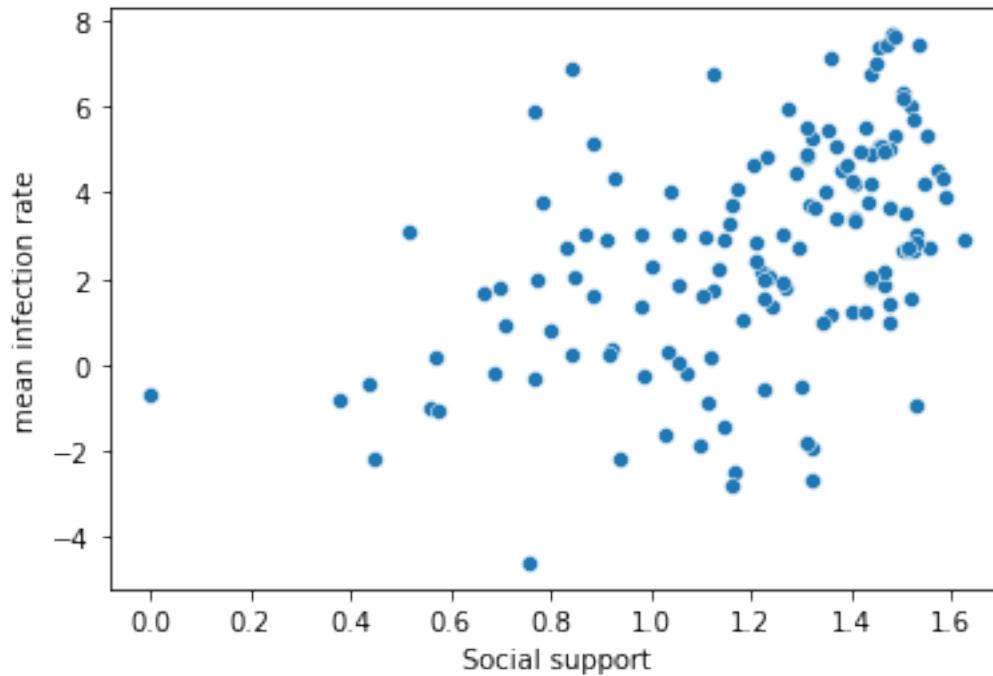
### Plotting Social support vs maximum Infection rate

```
x = data['Social support']
y = data['mean infection rate']
sns.scatterplot(x,np.log(y))
```

D:\Users\dell\anaconda3\lib\site-packages\seaborn\\_decorators.py:36:  
FutureWarning:

Pass the following variables as keyword args: x, y. From version 0.12, the only valid positional argument will be `data`, and passing other arguments without an explicit keyword will result in an error or misinterpretation.

```
<AxesSubplot:xlabel='Social support', ylabel='mean infection rate'>
```

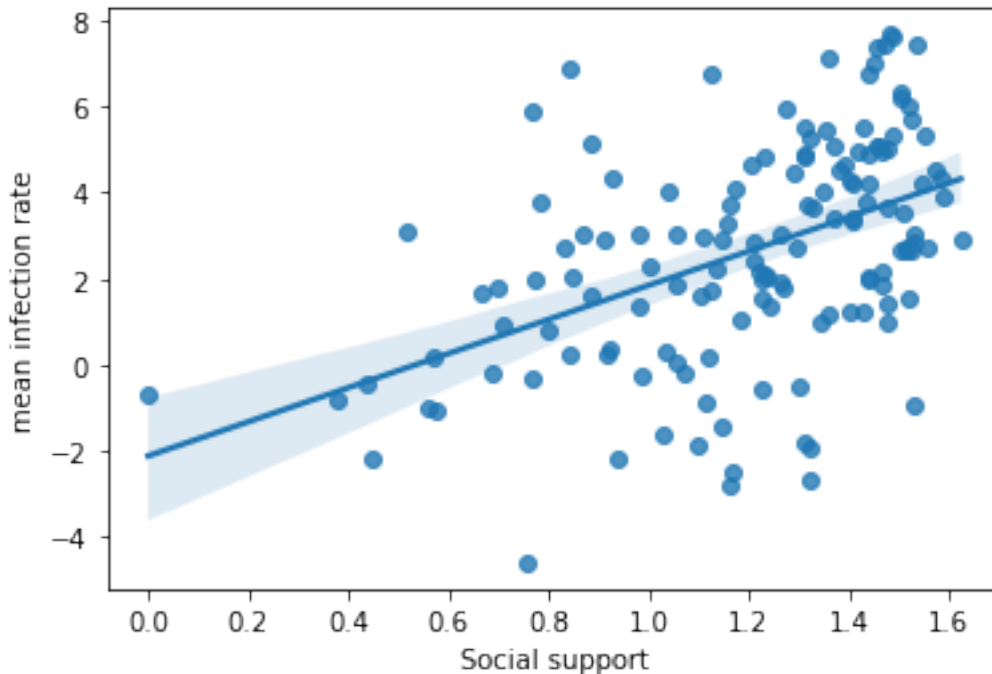


```
sns.regplot(x,np.log(y))
```

D:\Users\dell\anaconda3\lib\site-packages\seaborn\\_decorators.py:36:  
FutureWarning:

Pass the following variables as keyword args: x, y. From version 0.12,  
the only valid positional argument will be `data`, and passing other  
arguments without an explicit keyword will result in an error or  
misinterpretation.

```
<AxesSubplot:xlabel='Social support', ylabel='mean infection rate'>
```



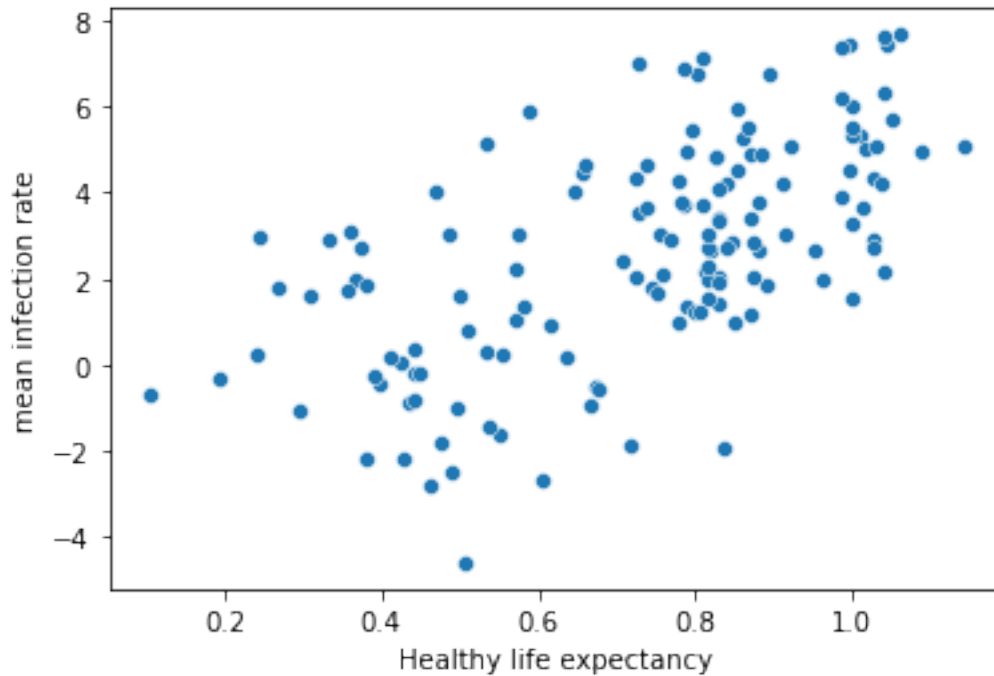
### Plotting Healthy life expectancy vs maximum Infection rate

```
x = data['Healthy life expectancy']  
y = data['mean infection rate']  
sns.scatterplot(x,np.log(y))
```

D:\Users\dell\anaconda3\lib\site-packages\seaborn\\_decorators.py:36:  
FutureWarning:

Pass the following variables as keyword args: x, y. From version 0.12, the only valid positional argument will be `data`, and passing other arguments without an explicit keyword will result in an error or misinterpretation.

```
<AxesSubplot:xlabel='Healthy life expectancy', ylabel='mean infection rate'>
```

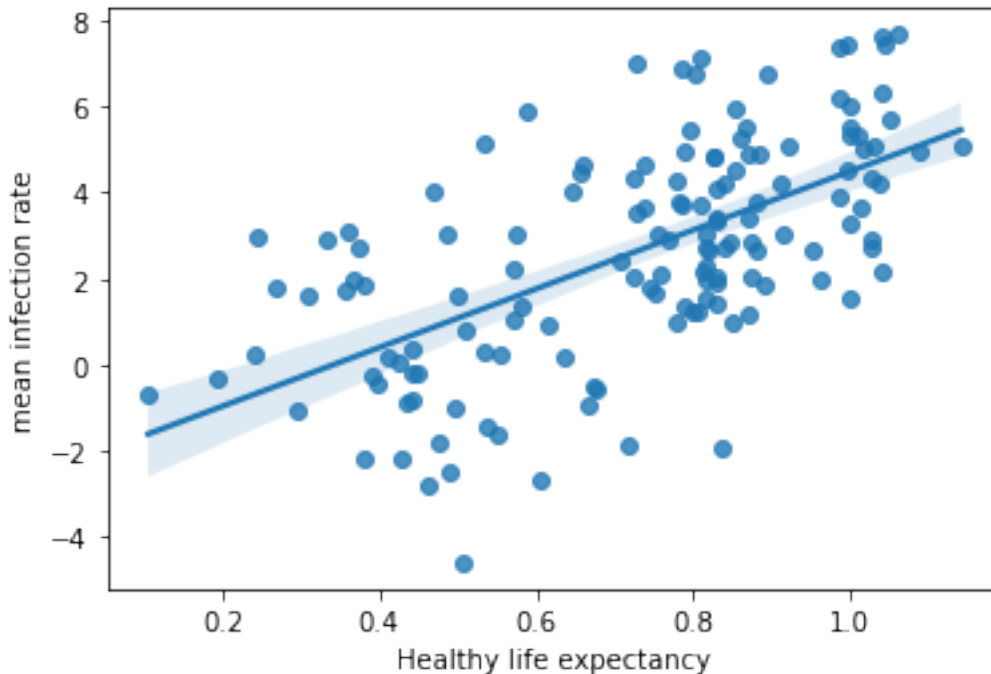


```
sns.regplot(x,np.log(y))
```

D:\Users\dell\anaconda3\lib\site-packages\seaborn\\_decorators.py:36:  
FutureWarning:

Pass the following variables as keyword args: x, y. From version 0.12,  
the only valid positional argument will be `data`, and passing other  
arguments without an explicit keyword will result in an error or  
misinterpretation.

```
<AxesSubplot:xlabel='Healthy life expectancy', ylabel='mean infection  
rate'>
```



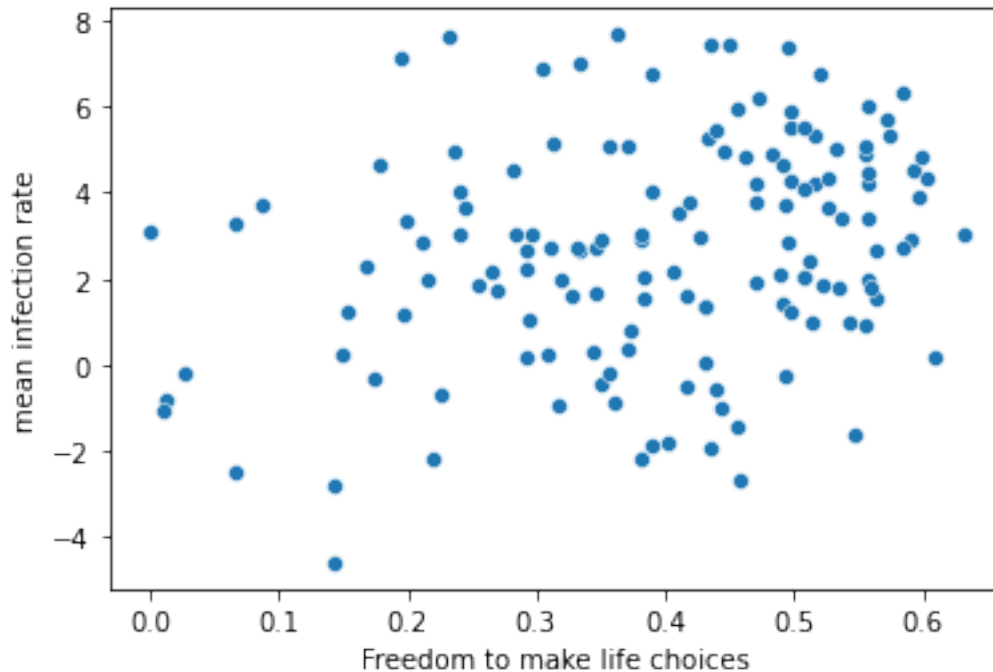
### Plotting Freedom to make life choices vs maximum Infection rate

```
x = data['Freedom to make life choices']
y = data['mean infection rate']
sns.scatterplot(x,np.log(y))
```

D:\Users\dell\anaconda3\lib\site-packages\seaborn\\_decorators.py:36:  
FutureWarning:

Pass the following variables as keyword args: x, y. From version 0.12, the only valid positional argument will be `data`, and passing other arguments without an explicit keyword will result in an error or misinterpretation.

```
<AxesSubplot:xlabel='Freedom to make life choices', ylabel='mean  
infection rate'>
```

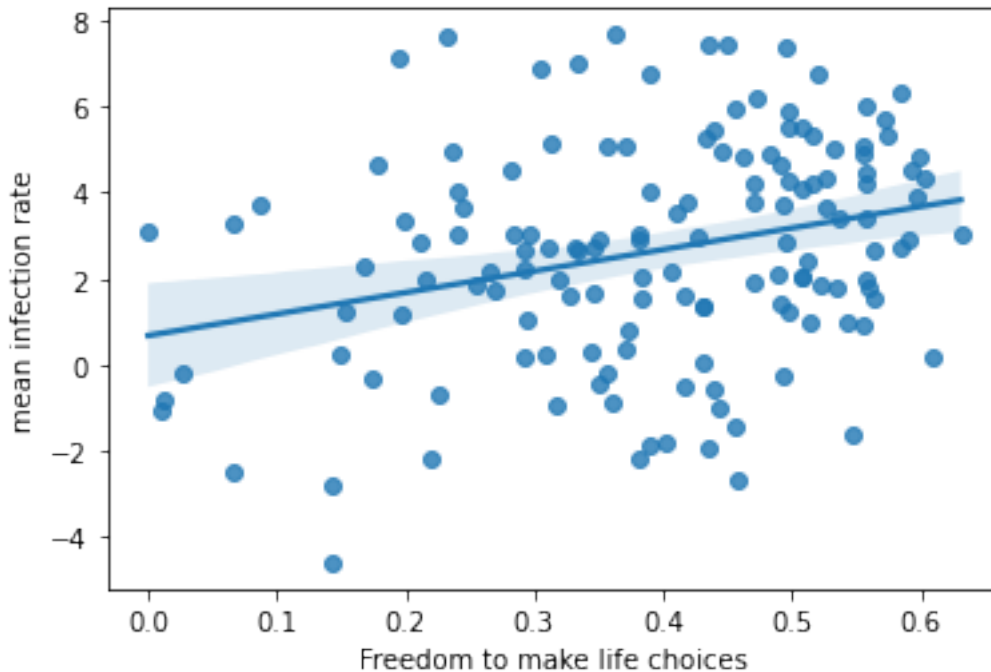


```
sns.regplot(x,np.log(y))
```

D:\Users\dell\anaconda3\lib\site-packages\seaborn\\_decorators.py:36:  
FutureWarning:

Pass the following variables as keyword args: x, y. From version 0.12, the only valid positional argument will be `data`, and passing other arguments without an explicit keyword will result in an error or misinterpretation.

```
<AxesSubplot:xlabel='Freedom to make life choices', ylabel='mean  
infection rate'>
```



```
from covid import Covid
```

```
covid = Covid()
```

```
covid.get_data()
```

```
len(covid.get_data())
```

```
196
```

```
countries = covid.list_countries()
```

```
italy_cases = covid.get_status_by_country_id(115)
```

```
india_cases = covid.get_status_by_country_name("india")
india_cases
```

```
{'id': '80',
 'country': 'India',
 'confirmed': 34674643,
 'active': None,
 'deaths': 474479,
 'recovered': None,
 'latitude': 20.593684,
 'longitude': 78.96288,
 'last_update': 1639146106000}
```

```
df=pd.DataFrame(data=covid.get_data())
```

```
df.head()
```

```
\
    id          country  confirmed  active  deaths  recovered  latitude
```

0	183	US	49666654	None	794656	None	40.000000
1	80	India	34674643	None	474479	None	20.593684
2	24	Brazil	22177059	None	616457	None	-14.235000
3	187	United Kingdom	10722083	None	146592	None	55.000000
4	145	Russia	9782723	None	281571	None	61.524000

	longitude	last_update
0	-100.00000	1639146106000
1	78.96288	1639146106000
2	-51.92530	1639146106000
3	-3.00000	1639146106000
4	105.31880	1639146106000

```
def plot_map(df, col, pal):
    df = df[df[col]>0]
    fig = px.choropleth(df, locations="country", locationmode='country
names',
                        color=col, hover_name="country",
                        title=col, hover_data=[col],
                        color_continuous_scale=pal)

    fig.show()

plot_map(df, 'confirmed', 'matter')
```

```
{"data":[{"hovertemplate":"<b>{hovertext}</b><br><br>country=%
{location}<br>confirmed={z}<extra></extra>","locationmode":"country
names","coloraxis":"coloraxis","customdata":[[49666654],[34674643],
[22177059],[10722083],[9782723],[8986377],[8209911],[6427417],
[6150843],[5350867],[5273178],[5164780],[5088008],[4258560],[3911714],
[3785036],[3717640],[3093452],[2870459],[2835996],[2678465],[2312736],
[2249671],[2160780],[2086702],[1914527],[1831341],[1790571],[1778370],
[1727654],[1578550],[1382272],[1349134],[1288366],[1270407],[1265056],
[1226055],[1221095],[1182922],[1181294],[1111923],[1061645],[998492],
[989814],[963428],[951092],[882822],[823786],[742641],[719006],
[711364],[685162],[671432],[644206],[621328],[610855],[601028],
[570672],[567906],[550136],[547892],[542181],[530126],[525750],
[503606],[487953],[480090],[463580],[463277],[437689],[436602],
[413608],[409490],[401897],[384912],[378423],[377450],[372711],
[368497],[366634],[342115],[305368],[304679],[280469],[277964],
[271979],[260658],[255796],[244867],[227446],[225640],[218362],
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[183863],[161176],[159214],[157601],[155817],[152586],[139368],
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```



```
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```

```
df.drop(['latitude', 'longitude', 'id'], axis=1, inplace=True)
df.head()
```

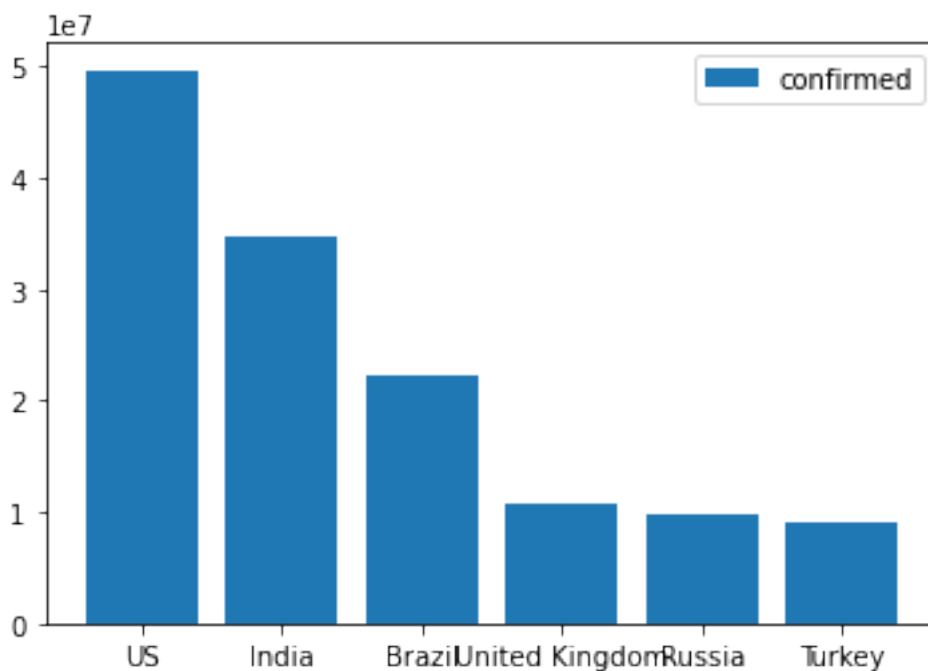
	country	confirmed	active	deaths	recovered	last_update
0	US	49666654	None	794656	None	1639146106000
1	India	34674643	None	474479	None	1639146106000
2	Brazil	22177059	None	616457	None	1639146106000
3	United Kingdom	10722083	None	146592	None	1639146106000
4	Russia	9782723	None	281571	None	1639146106000

```
df.set_index('country', inplace=True)
df.head()
```

country	confirmed	active	deaths	recovered	last_update
US	49666654	None	794656	None	1639146106000
India	34674643	None	474479	None	1639146106000
Brazil	22177059	None	616457	None	1639146106000
United Kingdom	10722083	None	146592	None	1639146106000
Russia	9782723	None	281571	None	1639146106000

```
countries=df.index[:6]
confirm=df['confirmed'][:6].to_numpy()
plt.bar(countries,confirm,label='confirmed')
plt.legend()
```

<matplotlib.legend.Legend at 0x1cfdba9dee0>



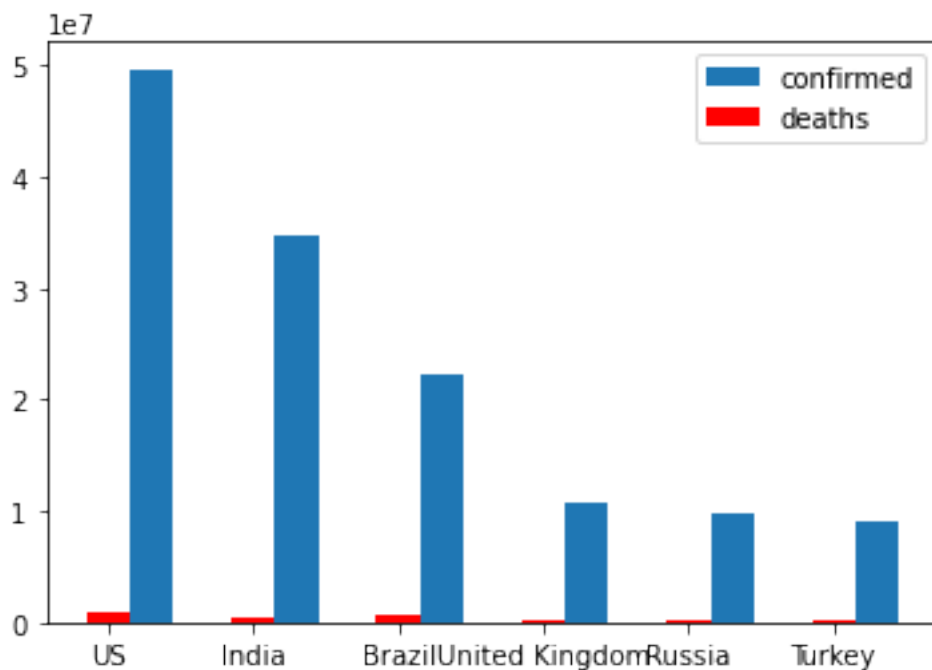
```
countries=df.index[:6]
confirm=df['confirmed'][:6].to_numpy()
total_deaths=df['deaths'][:6].to_numpy()
```

```
xpos=np.arange(6)
```

```
plt.bar(xpos+0.3,confirm,label='confirmed',width=0.3)
plt.bar(xpos,total_deaths,label='deaths',width=0.3,color='red')
plt.xticks(xpos,countries)
plt.legend()
```

<matplotlib.legend.Legend at 0x1cfdbb19df0>





```
final=df.join(corona_data,how='inner')
len(final)
```

```
186
```

```
final.head()
```

	confirmed	active	deaths	recovered	last_update \
US	49666654	None	794656	None	1639146106000
India	34674643	None	474479	None	1639146106000
Brazil	22177059	None	616457	None	1639146106000
United Kingdom	10722083	None	146592	None	1639146106000
Russia	9782723	None	281571	None	1639146106000

	mean infection rate
US	10802.252525
India	352.151515
Brazil	880.676768
United Kingdom	1742.232323
Russia	1075.737374

## Joining Data from covid library and World Happiness Index

```
data1 = world_happiness_report.join(df,how="inner")
```

```
data1.head(10)
```

	GDP per capita	Social support	Healthy life expectancy \
--	----------------	----------------	---------------------------

Finland	1.340	1.587	0.986
Denmark	1.383	1.573	0.996
Norway	1.488	1.582	1.028
Iceland	1.380	1.624	1.026
Netherlands	1.396	1.522	0.999
Switzerland	1.452	1.526	1.052
Sweden	1.387	1.487	1.009
New Zealand	1.303	1.557	1.026
Canada	1.365	1.505	1.039
Austria	1.376	1.475	1.016

recovered \	Freedom to make life choices	confirmed	active	deaths
Finland	0.596	201051	None	1421
None				
Denmark	0.592	542181	None	2996
None				
Norway	0.603	305368	None	1135
None				
Iceland	0.591	19159	None	35
None				
Netherlands	0.557	2870459	None	20372
None				
Switzerland	0.572	1111923	None	11739
None				
Sweden	0.574	1226055	None	15185
None				
New Zealand	0.585	12623	None	44
None				
Canada	0.584	1831341	None	29930
None				
Austria	0.532	1221095	None	13076
None				

	last_update
Finland	1639146106000
Denmark	1639146106000
Norway	1639146106000
Iceland	1639146106000

```
Netherlands 1639146106000
Switzerland 1639146106000
Sweden       1639146106000
New Zealand  1639146106000
Canada       1639146106000
Austria      1639146106000
```

### Scaling data

```
data1.drop(['active', 'recovered', 'last_update'], axis=1, inplace =
True)
data1.head()
```

	GDP per capita	Social support	Healthy life
expectancy \			
Finland	1.340	1.587	0.986
Denmark	1.383	1.573	0.996
Norway	1.488	1.582	1.028
Iceland	1.380	1.624	1.026
Netherlands	1.396	1.522	0.999

	Freedom to make life choices	confirmed	deaths
Finland	0.596	201051	1421
Denmark	0.592	542181	2996
Norway	0.603	305368	1135
Iceland	0.591	19159	35
Netherlands	0.557	2870459	20372

```
from sklearn.preprocessing import MinMaxScaler
scaler = MinMaxScaler()
scaled = scaler.fit_transform(data1)
```

```
from pandas import DataFrame
datax = DataFrame(scaled)
datax.head()
```

	0	1	2	3	4	5
0	0.795724	0.977217	0.850386	0.944532	0.005722	0.002300
1	0.821259	0.968596	0.860039	0.938193	0.015561	0.004855
2	0.883610	0.974138	0.890927	0.955626	0.008731	0.001836
3	0.819477	1.000000	0.888996	0.936609	0.000476	0.000052
4	0.828979	0.937192	0.862934	0.882726	0.082713	0.033042

## Performing linear regression

```
from sklearn.linear_model import LinearRegression
regression_model = LinearRegression()

x = datax.iloc[:, -6].values #GDP
y = datax.iloc[:, -1].values #deaths
x= x.reshape(-1,1)
regression_model.fit(x,y) #GDP vs Deaths

LinearRegression()

y_predicted = regression_model.predict(x)

from sklearn.metrics import mean_squared_error ,r2_score
mse = mean_squared_error(y,y_predicted)

mse

0.015631230121152265

rmse = np.sqrt(mean_squared_error(y,y_predicted))
r2 = r2_score(y,y_predicted)

rmse

0.12502491800098198

r2

0.021593705847284927
```

## GDP alone cannot explain variance of deaths

## Gradient descent approach

```
# Gradient descent_Build the model
m = 0
c = 0
L = 0.00001
epochs = 10000 #the number of iterations to perform gradation
X = datax.iloc[:, -6].values #attributes to determine dependant
variable/ Class
Y = datax.iloc[:, -1].values #dependant variable/ Class
n = float(len(X))
n

144.0

for i in range(epochs):
    Y_pred = m*X + c #The current predicted value of Y
    D_m = (-2/n) * sum(X* (Y - Y_pred)) # Derivative wrt m
    D_c = (-2/n) * sum(Y - Y_pred) # Derivative of c
```

```

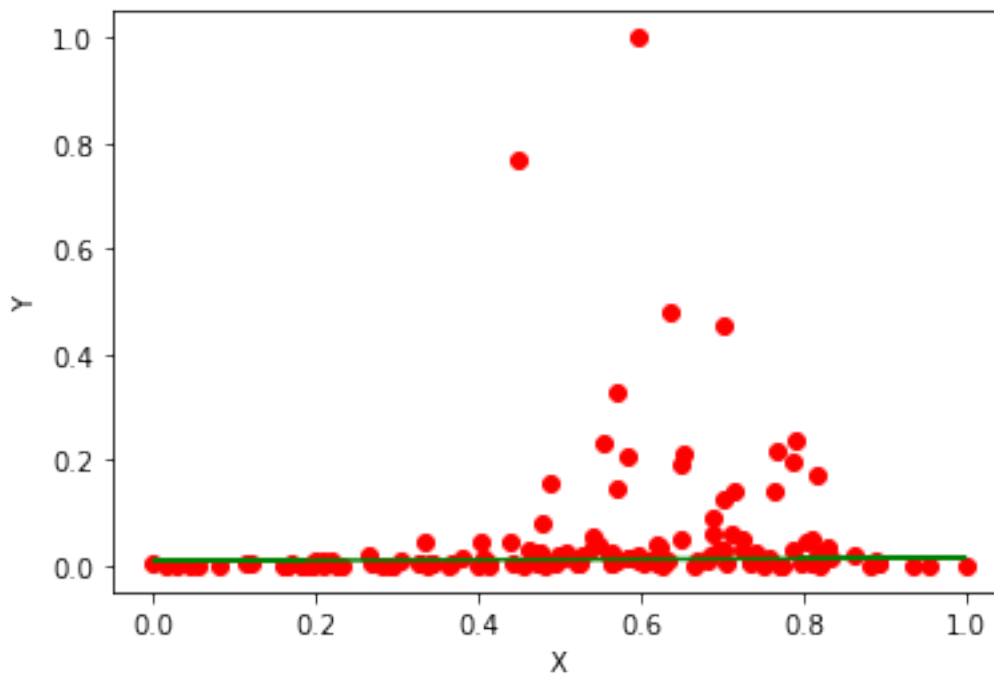
m = m - L* D_m # Update m
c = c - L*D_c # Update c
print (m,c)

0.005452208077273036 0.008711837822156551

Y_pred = m*X + c
plt.scatter(X,Y,color = "red")
plt.plot(X,Y_pred,color = "green")
plt.xlabel("X")
plt.ylabel("Y")

Text(0, 0.5, 'Y')

```



## Outliers of the Data Frame

### Social support index vs Deaths due to covid

```

x = datax.iloc[:, -5].values #attributes to determine dependant
variable/ Class
y = datax.iloc[:, -1].values #dependant variable/ Class
x= x.reshape(-1,1)
regression_model.fit(x,y)

LinearRegression()

y_predicted = regression_model.predict(x)

mse = mean_squared_error(y,y_predicted)
mse

```

0.015785897837850917

```
rmse = np.sqrt(mean_squared_error(y,y_predicted))
r2 = r2_score(y,y_predicted)
r2
```

0.011912582458575072

## Healthy Life Expectancy Vs Deaths due to covid

```
x = datax.iloc[:, -4].values #attributes to determine dependant variable/ Class
y = datax.iloc[:, -1].values #dependant variable/ Class
x= x.reshape(-1,1)
regression_model.fit(x,y)
y_predicted = regression_model.predict(x)
mse = mean_squared_error(y,y_predicted)
mse
```

0.0156979693574256

```
rmse = np.sqrt(mean_squared_error(y,y_predicted))
r2 = r2_score(y,y_predicted)
r2
```

0.017416293811848615

## Vaccinations

```
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import plotly.express as px
import plotly.graph_objects as go
import matplotlib.patches as mpatches
from plotly.subplots import make_subplots
from wordcloud import WordCloud
import seaborn as sns
sns.set(color_codes = True)
sns.set(style="whitegrid")
import plotly.figure_factory as ff
from plotly.colors import n_colors
```

```
df = pd.read_csv('country_vaccinations.csv')
df.head()
```

	country	iso_code	date	total_vaccinations
people_vaccinated \				
0	Afghanistan	AFG	2021-02-22	0.0
0.0				
1	Afghanistan	AFG	2021-02-23	NaN

NaN				
2	Afghanistan	AFG	2021-02-24	NaN
NaN				
3	Afghanistan	AFG	2021-02-25	NaN
NaN				
4	Afghanistan	AFG	2021-02-26	NaN
NaN				

	people_fully_vaccinated	daily_vaccinations_raw	daily_vaccinations
\			
0	NaN	NaN	NaN
1	NaN	NaN	1367.0
2	NaN	NaN	1367.0
3	NaN	NaN	1367.0
4	NaN	NaN	1367.0

	total_vaccinations_per_hundred	people_vaccinated_per_hundred	\
0	0.0	0.0	
1	NaN	NaN	
2	NaN	NaN	
3	NaN	NaN	
4	NaN	NaN	

	people_fully_vaccinated_per_hundred	daily_vaccinations_per_million
\		
0	NaN	NaN
1	NaN	34.0
2	NaN	34.0
3	NaN	34.0
4	NaN	34.0

	vaccines	\
0	Johnson&Johnson, Oxford/AstraZeneca, Pfizer/Bi...	
1	Johnson&Johnson, Oxford/AstraZeneca, Pfizer/Bi...	
2	Johnson&Johnson, Oxford/AstraZeneca, Pfizer/Bi...	
3	Johnson&Johnson, Oxford/AstraZeneca, Pfizer/Bi...	
4	Johnson&Johnson, Oxford/AstraZeneca, Pfizer/Bi...	

source\_name \

```

0 World Health Organization
1 World Health Organization
2 World Health Organization
3 World Health Organization
4 World Health Organization

```

```

                                source_website
0 https://reliefweb.int/sites/reliefweb.int/file...
1 https://reliefweb.int/sites/reliefweb.int/file...
2 https://reliefweb.int/sites/reliefweb.int/file...
3 https://reliefweb.int/sites/reliefweb.int/file...
4 https://reliefweb.int/sites/reliefweb.int/file...

```

```
df.isnull().sum()
```

```

country                0
iso_code                0
date                   0
total_vaccinations     25685
people_vaccinated       27190
people_fully_vaccinated 30037
daily_vaccinations_raw  31124
daily_vaccinations      253
total_vaccinations_per_hundred 25685
people_vaccinated_per_hundred 27190
people_fully_vaccinated_per_hundred 30037
daily_vaccinations_per_million 253
vaccines                0
source_name             0
source_website          0
dtype: int64

```

```

df.fillna(value = 0, inplace = True)
df.total_vaccinations = df.total_vaccinations.astype(int)
df.people_vaccinated = df.people_vaccinated.astype(int)
df.people_fully_vaccinated = df.people_fully_vaccinated.astype(int)
df.daily_vaccinations_raw = df.daily_vaccinations_raw.astype(int)
df.daily_vaccinations = df.daily_vaccinations.astype(int)
df.total_vaccinations_per_hundred =
df.total_vaccinations_per_hundred.astype(int)
df.people_fully_vaccinated_per_hundred =
df.people_fully_vaccinated_per_hundred.astype(int)
df.daily_vaccinations_per_million =
df.daily_vaccinations_per_million.astype(int)
df.people_vaccinated_per_hundred =
df.people_vaccinated_per_hundred.astype(int)
date = df.date.str.split('-', expand = True)
date

```

```

      0    1    2
0  2021  02  22

```



```

1      2021  02  23
2      2021  02  24
3      2021  02  25
4      2021  02  26
...
55819  2021  11  01
55820  2021  11  02
55821  2021  11  03
55822  2021  11  04
55823  2021  11  05

```

[55824 rows x 3 columns]

```

df['year'] = date[0]
df['month'] = date[1]
df['day'] = date[2]
df.year = pd.to_numeric(df.year)
df.month = pd.to_numeric(df.month)
df.day = pd.to_numeric(df.day)
df.date = pd.to_datetime(df.date)
df.head()

```

	country	iso_code	date	total_vaccinations
0	Afghanistan	AFG	2021-02-22	0
1	Afghanistan	AFG	2021-02-23	0
2	Afghanistan	AFG	2021-02-24	0
3	Afghanistan	AFG	2021-02-25	0
4	Afghanistan	AFG	2021-02-26	0

	people_fully_vaccinated	daily_vaccinations_raw	daily_vaccinations
0	0	0	0
1	0	0	1367
2	0	0	1367
3	0	0	1367
4	0	0	1367

	total_vaccinations_per_hundred	people_vaccinated_per_hundred
0	0	0

1	0	0
2	0	0
3	0	0
4	0	0

	people_fully_vaccinated_per_hundred	daily_vaccinations_per_million
\		
0	0	0
1	0	34
2	0	34
3	0	34
4	0	34

	vaccines	\
0	Johnson&Johnson, Oxford/AstraZeneca, Pfizer/Bi...	
1	Johnson&Johnson, Oxford/AstraZeneca, Pfizer/Bi...	
2	Johnson&Johnson, Oxford/AstraZeneca, Pfizer/Bi...	
3	Johnson&Johnson, Oxford/AstraZeneca, Pfizer/Bi...	
4	Johnson&Johnson, Oxford/AstraZeneca, Pfizer/Bi...	

	source_name	\
0	World Health Organization	
1	World Health Organization	
2	World Health Organization	
3	World Health Organization	
4	World Health Organization	

	source_website	year	month	day
0	https://reliefweb.int/sites/reliefweb.int/file...	2021	2	22
1	https://reliefweb.int/sites/reliefweb.int/file...	2021	2	23
2	https://reliefweb.int/sites/reliefweb.int/file...	2021	2	24
3	https://reliefweb.int/sites/reliefweb.int/file...	2021	2	25
4	https://reliefweb.int/sites/reliefweb.int/file...	2021	2	26

df.describe()

	total_vaccinations	people_vaccinated	people_fully_vaccinated
\			
count	5.582400e+04	5.582400e+04	5.582400e+04

mean	9.954941e+06	5.624392e+06	3.438866e+06
std	1.057154e+08	3.158674e+07	1.941963e+07
min	-2.147484e+09	0.000000e+00	0.000000e+00
25%	0.000000e+00	0.000000e+00	0.000000e+00
50%	3.108000e+04	4.969000e+03	0.000000e+00
75%	2.477739e+06	1.369812e+06	6.764480e+05
max	2.142580e+09	1.100842e+09	1.070386e+09

	daily_vaccinations_raw	daily_vaccinations \
count	5.582400e+04	5.582400e+04
mean	1.134868e+05	1.297107e+05
std	8.412834e+05	8.228933e+05
min	0.000000e+00	0.000000e+00
25%	0.000000e+00	9.220000e+02
50%	0.000000e+00	7.007000e+03
75%	1.709025e+04	4.154275e+04
max	2.474100e+07	2.242429e+07

	total_vaccinations_per_hundred people_vaccinated_per_hundred \	
count	55824.000000	55824.000000
mean	28.357785	15.636644
std	45.498547	24.255363
min	0.000000	0.000000
25%	0.000000	0.000000
50%	0.000000	0.000000
75%	42.000000	26.000000
max	269.000000	120.000000

	people_fully_vaccinated_per_hundred daily_vaccinations_per_million \
count	55824.000000

```

55824.000000
mean          11.008222
3480.269669
std           20.232838
4241.817117
min           0.000000
0.000000
25%           0.000000
586.000000
50%           0.000000
2205.000000
75%           12.000000
5095.000000
max           118.000000
117497.000000

```

```

              year      month      day
count  55824.000000  55824.000000  55824.000000
mean    2020.993730    6.266946    15.846106
std       0.078934     2.637807     8.869602
min    2020.000000     1.000000     1.000000
25%    2021.000000     4.000000     8.000000
50%    2021.000000     6.000000    16.000000
75%    2021.000000     8.000000    24.000000
max    2021.000000    12.000000    31.000000

```

```
df.country.unique()
```

```

array(['Afghanistan', 'Albania', 'Algeria', 'Andorra', 'Angola',
      'Anguilla', 'Antigua and Barbuda', 'Argentina', 'Armenia',
      'Aruba',
      'Australia', 'Austria', 'Azerbaijan', 'Bahamas', 'Bahrain',
      'Bangladesh', 'Barbados', 'Belarus', 'Belgium', 'Belize',
      'Benin',
      'Bermuda', 'Bhutan', 'Bolivia', 'Bonaire Sint Eustatius and
      Saba',
      'Bosnia and Herzegovina', 'Botswana', 'Brazil',
      'British Virgin Islands', 'Brunei', 'Bulgaria', 'Burkina Faso',
      'Burundi', 'Cambodia', 'Cameroon', 'Canada', 'Cape Verde',
      'Cayman Islands', 'Central African Republic', 'Chad', 'Chile',
      'China', 'Colombia', 'Comoros', 'Congo', 'Cook Islands',
      'Costa Rica', 'Cote d'Ivoire', 'Croatia', 'Cuba', 'Curacao',
      'Cyprus', 'Czechia', 'Democratic Republic of Congo', 'Denmark',
      'Djibouti', 'Dominica', 'Dominican Republic', 'Ecuador',
      'Egypt',
      'El Salvador', 'England', 'Equatorial Guinea', 'Estonia',
      'Eswatini', 'Ethiopia', 'Faeroe Islands', 'Falkland Islands',
      'Fiji', 'Finland', 'France', 'French Polynesia', 'Gabon',
      'Gambia',
      'Georgia', 'Germany', 'Ghana', 'Gibraltar', 'Greece',

```

```

'Greenland',
    'Grenada', 'Guatemala', 'Guernsey', 'Guinea', 'Guinea-Bissau',
    'Guyana', 'Haiti', 'Honduras', 'Hong Kong', 'Hungary',
'Iceland',
    'India', 'Indonesia', 'Iran', 'Iraq', 'Ireland', 'Isle of Man',
    'Israel', 'Italy', 'Jamaica', 'Japan', 'Jersey', 'Jordan',
    'Kazakhstan', 'Kenya', 'Kiribati', 'Kosovo', 'Kuwait',
    'Kyrgyzstan', 'Laos', 'Latvia', 'Lebanon', 'Lesotho',
'Liberia',
    'Libya', 'Liechtenstein', 'Lithuania', 'Luxembourg', 'Macao',
    'Madagascar', 'Malawi', 'Malaysia', 'Maldives', 'Mali',
'Malta',
    'Mauritania', 'Mauritius', 'Mexico', 'Moldova', 'Monaco',
    'Mongolia', 'Montenegro', 'Montserrat', 'Morocco',
'Mozambique',
    'Myanmar', 'Namibia', 'Nauru', 'Nepal', 'Netherlands',
    'New Caledonia', 'New Zealand', 'Nicaragua', 'Niger',
'Nigeria',
    'Niue', 'North Macedonia', 'Northern Cyprus', 'Northern
Ireland',
    'Norway', 'Oman', 'Pakistan', 'Palestine', 'Panama',
    'Papua New Guinea', 'Paraguay', 'Peru', 'Philippines',
'Pitcairn',
    'Poland', 'Portugal', 'Qatar', 'Romania', 'Russia', 'Rwanda',
    'Saint Helena', 'Saint Kitts and Nevis', 'Saint Lucia',
    'Saint Vincent and the Grenadines', 'Samoa', 'San Marino',
    'Sao Tome and Principe', 'Saudi Arabia', 'Scotland', 'Senegal',
    'Serbia', 'Seychelles', 'Sierra Leone', 'Singapore',
    'Sint Maarten (Dutch part)', 'Slovakia', 'Slovenia',
    'Solomon Islands', 'Somalia', 'South Africa', 'South Korea',
    'South Sudan', 'Spain', 'Sri Lanka', 'Sudan', 'Suriname',
'Sweden',
    'Switzerland', 'Syria', 'Taiwan', 'Tajikistan', 'Tanzania',
    'Thailand', 'Timor', 'Togo', 'Tokelau', 'Tonga',
    'Trinidad and Tobago', 'Tunisia', 'Turkey', 'Turkmenistan',
    'Turks and Caicos Islands', 'Tuvalu', 'Uganda', 'Ukraine',
    'United Arab Emirates', 'United Kingdom', 'United States',
    'Uruguay', 'Uzbekistan', 'Vanuatu', 'Venezuela', 'Vietnam',
    'Wales', 'Wallis and Futuna', 'Yemen', 'Zambia', 'Zimbabwe'],
dtype=object)

```

```

def size(m,n):
    fig = plt.gcf();
    fig.set_size_inches(m,n);

wordCloud = WordCloud(
    background_color='white',
    max_font_size = 50).generate(' '.join(df.country))
plt.figure(figsize=(15,7))
plt.axis('off')

```

```
plt.imshow(wordCloud)
plt.show()
```



```
country_wise_total_vaccinated = {}
for country in df.country.unique() :
    vaccinated = 0
    for i in range(len(df)) :
        if df.country[i] == country :
            vaccinated += df.daily_vaccinations[i]
    country_wise_total_vaccinated[country] = vaccinated
# made a seperate dict from the df
country_wise_total_vaccinated_df =
pd.DataFrame.from_dict(country_wise_total_vaccinated,
orient='index',
columns =
['total_vaccinted_till_date'])
# converted dict to df
country_wise_total_vaccinated_df.sort_values(by =
'total_vaccinted_till_date', ascending = False, inplace = True)
country_wise_total_vaccinated_df
```

C:\Users\dell\AppData\Local\Temp\ipykernel\_8784\469794026.py:6:  
RuntimeWarning:

overflow encountered in long\_scalars

	total_vaccinted_till_date
India	1070252854
United States	426856166
Brazil	277092585
Indonesia	196518393

Japan	189787850
...	...
Niue	2484
Tokelau	1966
Burundi	610
Pitcairn	43
China	-2007914084

[223 rows x 1 columns]

```
datar = country_wise_total_vaccinated_df.join(data1,how="inner")
```

```
datar.head(10)
```

support \	total_vaccinted_till_date	GDP per capita	Social
India	1070252854	0.755	
0.765			
Brazil	277092585	1.004	
1.439			
Indonesia	196518393	0.931	
1.203			
Japan	189787850	1.327	
1.419			
Mexico	125372969	1.070	
1.323			
Turkey	117406602	1.183	
1.360			
Germany	112306162	1.373	
1.454			
Pakistan	107371250	0.677	
0.886			
Russia	105282120	1.183	
1.452			
United Kingdom	102153121	1.333	
1.538			

\	Healthy life expectancy	Freedom to make life choices
India	0.588	0.498
Brazil	0.802	0.390
Indonesia	0.660	0.491
Japan	1.088	0.445

Mexico	0.861	0.433
Turkey	0.808	0.195
Germany	0.987	0.495
Pakistan	0.535	0.313
Russia	0.726	0.334
United Kingdom	0.996	0.450

	confirmed	deaths
India	34674643	474479
Brazil	22177059	616457
Indonesia	4258560	143918
Japan	1727654	18367
Mexico	3911714	296188
Turkey	8986377	78602
Germany	6427417	105000
Pakistan	1288366	28812
Russia	9782723	281571
United Kingdom	10722083	146592

```
from sklearn.preprocessing import MinMaxScaler
scaler = MinMaxScaler()
scaled = scaler.fit_transform(datar)
```

```
from pandas import DataFrame
data_scaled = DataFrame(scaled)
data_scaled.head()
```

	0	1	2	3	4	5
6						
0	1.000000	0.448337	0.471059	0.466216	0.789223	1.000000
	0.769686					
1	0.742327	0.596200	0.886084	0.672780	0.618067	0.639548
	1.000000					
2	0.716151	0.552850	0.740764	0.535714	0.778130	0.122748
	0.233456					
3	0.713965	0.788005	0.873768	0.948842	0.705230	0.049752
	0.029790					
4	0.693038	0.635392	0.814655	0.729730	0.686212	0.112744
	0.480466					

```
from sklearn.linear_model import LinearRegression
regression_model = LinearRegression()
from sklearn.metrics import mean_squared_error ,r2_score
```



```

x = data_scaled.iloc[:, -7].values #attributes to determine dependant
variable/ Class
y = data_scaled.iloc[:, -1].values #dependant variable/ Class
x=x.reshape(-1,1)
regression_model.fit(x,y)
y_predicted = regression_model.predict(x)
mse = mean_squared_error(y,y_predicted)
mse

```

0.013823870773761234

```

rmse = np.sqrt(mean_squared_error(y,y_predicted))
r2 = r2_score(y,y_predicted)
r2

```

0.1449725320537083

```

x = data_scaled.iloc[:, -7].values #attributes to determine dependant
variable/ Class
y = data_scaled.iloc[:, -2].values #dependant variable/ Class
x=x.reshape(-1,1)
regression_model.fit(x,y)
y_predicted = regression_model.predict(x)
mse = mean_squared_error(y,y_predicted)
mse

```

0.009501470187026878

```

rmse = np.sqrt(mean_squared_error(y,y_predicted))
r2 = r2_score(y,y_predicted)
r2

```

0.22022315976696827

## Multinomial regression

Performing multinomial regreesion on (Vaccination,GDP) vs Deaths due to Covid

	0	1	2	3	4	5
6						
0	1.000000	0.448337	0.471059	0.466216	0.789223	1.000000
0.769686						
1	0.742327	0.596200	0.886084	0.672780	0.618067	0.639548
1.000000						
2	0.716151	0.552850	0.740764	0.535714	0.778130	0.122748
0.233456						
3	0.713965	0.788005	0.873768	0.948842	0.705230	0.049752

```

0.029790
4      0.693038  0.635392  0.814655  0.729730  0.686212  0.112744
0.480466
..      ...      ...      ...      ...      ...      ...
...
137  0.652380  0.207838  0.471675  0.083977  0.275753  0.000088
0.000289
138  0.652353  0.181710  0.354064  0.183398  0.015848  0.000294
0.000211
139  0.652349  0.191805  0.423645  0.332046  0.041204  0.000666
0.001225
140  0.652309  0.027316  0.275246  0.265444  0.348653  0.000518
0.000057
141  0.000000  0.611045  0.692734  0.760618  0.825674  0.003156
0.007861

```

```
[142 rows x 7 columns]
```

```

from sklearn import linear_model
import statsmodels.api as sm
from sklearn.metrics import r2_score
X = data_scaled[[0,1]] # here we have 2 variables for multiple
regression. If you just want to use one variable for simple linear
regression, then use X = df['Interest_Rate'] for
example. Alternatively, you may add additional variables within the
brackets
Y = data_scaled[6]

```

```

# with sklearn
regr = linear_model.LinearRegression()
regr.fit(X, Y)

```

```

print('Intercept: \n', regr.intercept_)
print('Coefficients: \n', regr.coef_)
print('R sqaure: \n')
regr.score(X, Y)

```

```

Intercept:
-0.4866183114101143
Coefficients:
[0.75670296 0.07414227]
R sqaure:

```

```
0.16382754007909928
```

## Performing multinomial regreesion on (Vaccination,GDP) vs Deaths due to Covid using stastical models

```

# with statsmodels
X = sm.add_constant(X) # adding a constant

```

```
model = sm.OLS(Y, X).fit()
predictions = model.predict(X)
```

```
print_model = model.summary()
print(print_model)
```

```
D:\Users\dell\anaconda3\lib\site-packages\statsmodels\tsa\
tsatools.py:142: FutureWarning:
```

In a future version of pandas all arguments of concat except for the argument 'objs' will be keyword-only

### OLS Regression Results

```
=====
=====
Dep. Variable:                6    R-squared:
0.164
Model:                        OLS    Adj. R-squared:
0.152
Method:                Least Squares    F-statistic:
13.62
Date:                Fri, 10 Dec 2021    Prob (F-statistic):
3.98e-06
Time:                20:38:27    Log-Likelihood:
104.07
No. Observations:                142    AIC:
-202.1
Df Residuals:                139    BIC:
-193.3
Df Model:                2

Covariance Type:                nonrobust
```

```
=====
=====
coef    std err          t    P>|t|    [0.025
0.975]
-----
-----
const    -0.4866    0.104    -4.660    0.000    -0.693
-0.280
0         0.7567    0.155     4.885    0.000     0.450
1.063
1         0.0741    0.042     1.770    0.079    -0.009
0.157
=====
=====
```

Omnibus:	168.792	Durbin-Watson:
1.099		
Prob(Omnibus):	0.000	Jarque-Bera (JB):
4361.505		
Skew:	4.557	Prob(JB):
0.00		
Kurtosis:	28.575	Cond. No.
24.7		

=====

=====

Notes:

[1] Standard Errors assume that the covariance matrix of the errors is correctly specified.

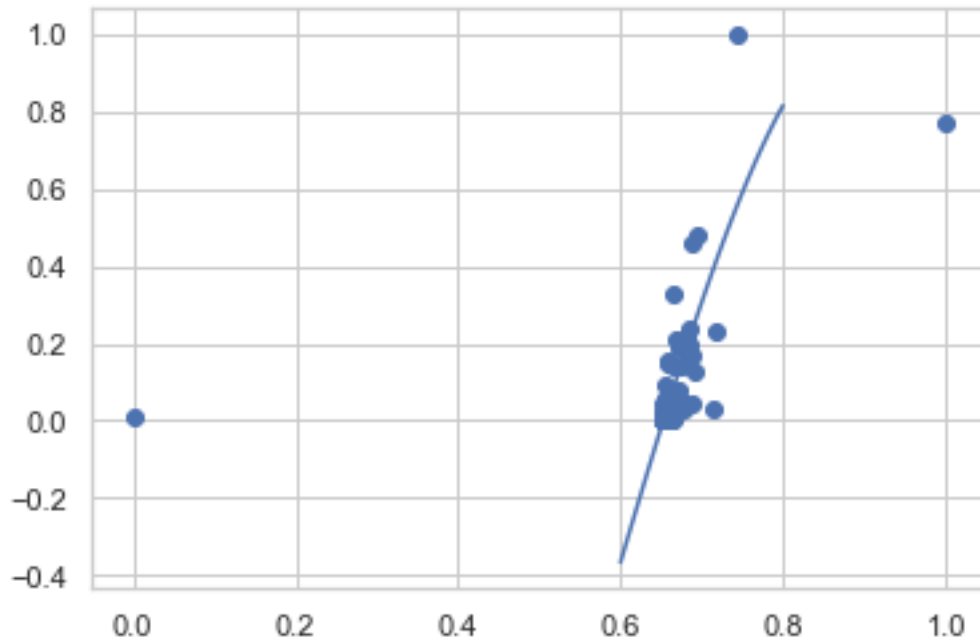
## Polynomial Regression

```
X = data_scaled[0] # Vaccinated list of different countries
Y = data_scaled[6] # Covid Confirmed cases
```

### Polynomial regression (Vaccinations vs Confirmed cases)

```
import numpy as np
mymodel = np.poly1d(np.polyfit(X, Y, 3))
myline = np.linspace(0.6, 0.8, 100)

plt.scatter(X, Y)
plt.plot(myline, mymodel(myline))
plt.show()
```



### Rsquare value of Vaccinations vs Confirmed cases

```
print(r2_score(Y, mymodel(X)))
```

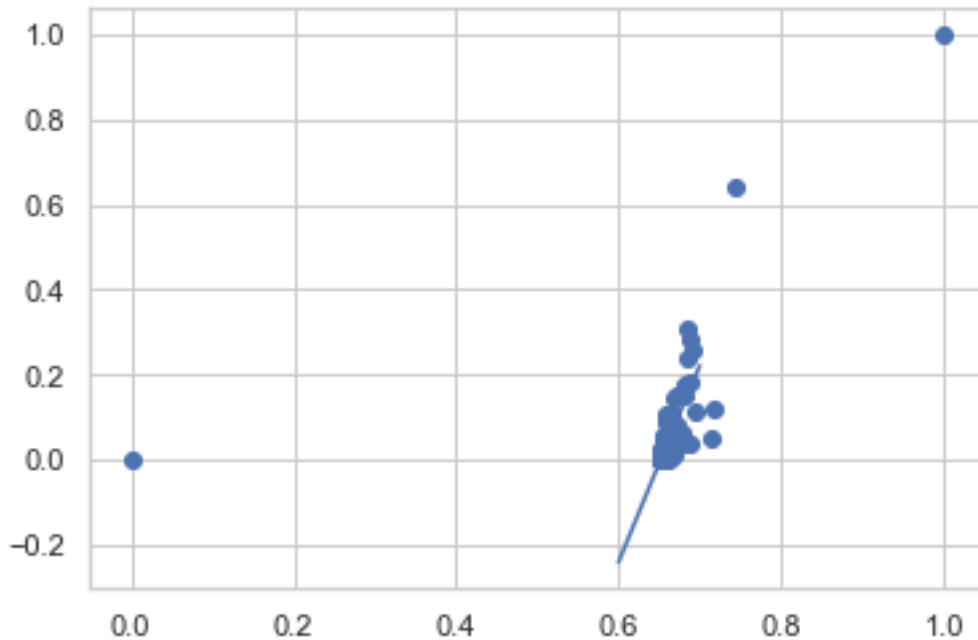
0.6847344781455831

### Polynomial regression (Vaccinations vs Deaths)

```
X = data_scaled[0] #Vaccinated list
Y =data_scaled[5] # Deaths due to covid
```

```
import numpy as np
mymodel = np.poly1d(np.polyfit(X, Y, 3))
myline = np.linspace(0.6, 0.7, 100)
```

```
plt.scatter(X, Y)
plt.plot(myline, mymodel(myline))
plt.show()
print("R2 value of Vaccinations vs Deaths:")
print(r2_score(Y, mymodel(X)))
```



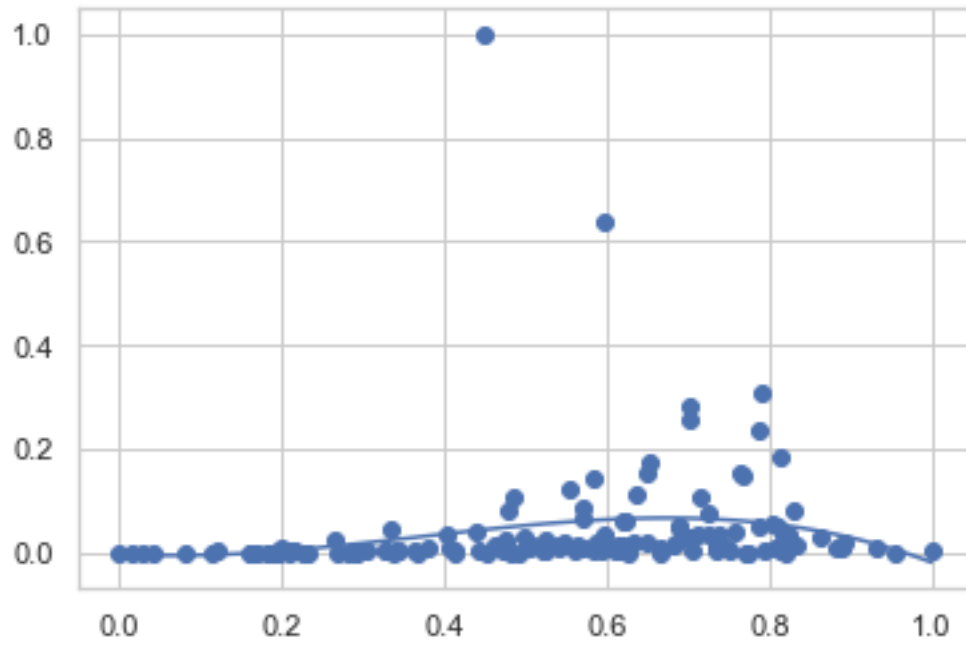
R2 value of Vaccinations vs Deaths:  
0.8397867607203168

### Polynomial Regression (GDP vs Deaths)

```
X = data_scaled[1]
Y = data_scaled[5]

import numpy as np
mymodel = np.poly1d(np.polyfit(X, Y, 3))
myline = np.linspace(0, 1, 100)

plt.scatter(X, Y)
plt.plot(myline, mymodel(myline))
plt.show()
print("R2 value of Gdp vs Deaths:")
print(r2_score(Y, mymodel(X)))
```



R2 value of Gdp vs Deaths:  
0.049696032274289315