

# Importing the Modules

```
In [43]: import pandas as pd
import numpy as nms
import seaborn as sns
import matplotlib.pyplot as plt
```

## Importing the dataset

```
In [7]: covid19 = pd.read_csv("desktop/covid19_dataset.csv")
covid19.head(10)
```

```
Out[7]:
```

	Province/State	Country/Region	Lat	Long	1/22/20	1/23/20	1/24/20	1/25/20	1/26/20	1/27/20	...	4/21/20	4/22/20	4/23/20	4/24/20	4/25/20
0	NaN	Afghanistan	33.0000	65.0000	0	0	0	0	0	0	...	1092	1176	1279	1351	1428
1	NaN	Albania	41.1533	20.1683	0	0	0	0	0	0	...	609	634	663	678	700
2	NaN	Algeria	28.0339	1.6596	0	0	0	0	0	0	...	2811	2910	3007	3127	3200
3	NaN	Andorra	42.5063	1.5218	0	0	0	0	0	0	...	717	723	723	731	740
4	NaN	Angola	-11.2027	17.8739	0	0	0	0	0	0	...	24	25	25	25	25
5	NaN	Antigua and Barbuda	17.0608	-61.7964	0	0	0	0	0	0	...	23	24	24	24	24
6	NaN	Argentina	-38.4161	-63.6167	0	0	0	0	0	0	...	3031	3144	3435	3607	3700
7	NaN	Armenia	40.0691	45.0382	0	0	0	0	0	0	...	1401	1473	1523	1596	1600
8	Australian Capital Territory	Australia	-35.4735	149.0124	0	0	0	0	0	0	...	104	104	104	105	105
9	New South Wales	Australia	-33.8688	151.2093	0	0	0	0	3	4	...	2969	2971	2976	2982	2982

10 rows × 104 columns



# Checking the shape

```
In [8]: covid19.shape
```

```
Out[8]: (266, 104)
```

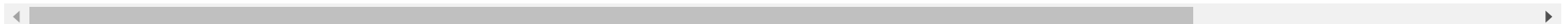
## Deleing the useless columns

```
In [9]: covid19.drop(["Lat","Long"],axis=1,inplace=True)
```

```
In [10]: covid19.head(10)
```

Out[10]:	Province/State	Country/Region	1/22/20	1/23/20	1/24/20	1/25/20	1/26/20	1/27/20	1/28/20	1/29/20	...	4/21/20	4/22/20	4/23/20	4/24/20	4/25/20
0	NaN	Afghanistan	0	0	0	0	0	0	0	0	...	1092	1176	1279	1351	1463
1	NaN	Albania	0	0	0	0	0	0	0	0	...	609	634	663	678	712
2	NaN	Algeria	0	0	0	0	0	0	0	0	...	2811	2910	3007	3127	3256
3	NaN	Andorra	0	0	0	0	0	0	0	0	...	717	723	723	731	738
4	NaN	Angola	0	0	0	0	0	0	0	0	...	24	25	25	25	25
5	NaN	Antigua and Barbuda	0	0	0	0	0	0	0	0	...	23	24	24	24	24
6	NaN	Argentina	0	0	0	0	0	0	0	0	...	3031	3144	3435	3607	3780
7	NaN	Armenia	0	0	0	0	0	0	0	0	...	1401	1473	1523	1596	1677
8	Australian Capital Territory	Australia	0	0	0	0	0	0	0	0	...	104	104	104	105	106
9	New South Wales	Australia	0	0	0	0	3	4	4	4	...	2969	2971	2976	2982	2994

10 rows × 102 columns



```
In [ ]: aggregating the rows by country
```

```
In [11]: covid19_aggregated = covid19.groupby("Country/Region").sum()  
covid19_aggregated.head()
```

```
Out[11]:
```

	1/22/20	1/23/20	1/24/20	1/25/20	1/26/20	1/27/20	1/28/20	1/29/20	1/30/20	1/31/20	...	4/21/20	4/22/20	4/23/20	4/24/20	4/25/20
Country/Region																
Afghanistan	0	0	0	0	0	0	0	0	0	0	...	1092	1176	1279	1351	1463
Albania	0	0	0	0	0	0	0	0	0	0	...	609	634	663	678	712
Algeria	0	0	0	0	0	0	0	0	0	0	...	2811	2910	3007	3127	3256
Andorra	0	0	0	0	0	0	0	0	0	0	...	717	723	723	731	738
Angola	0	0	0	0	0	0	0	0	0	0	...	24	25	25	25	25

5 rows × 100 columns



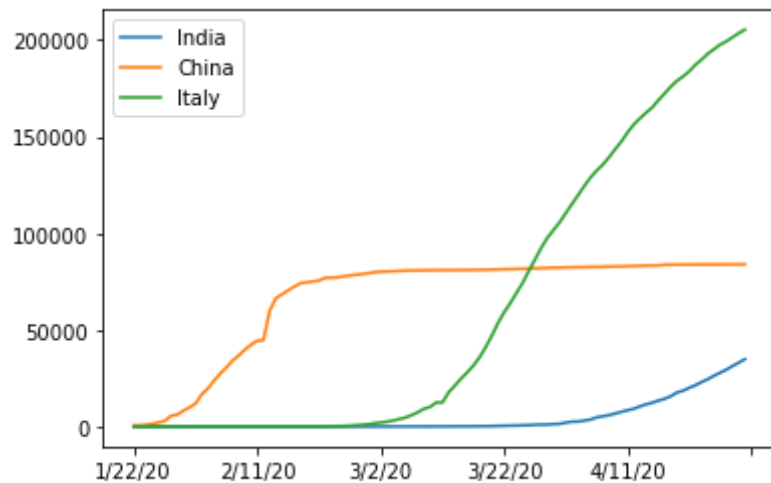
```
In [12]: covid19_aggregated.shape
```

```
Out[12]: (187, 100)
```

## Visualizing data related to countries

```
In [13]: covid19_aggregated.loc["India"].plot()  
covid19_aggregated.loc["China"].plot()  
covid19_aggregated.loc["Italy"].plot()  
plt.legend()
```

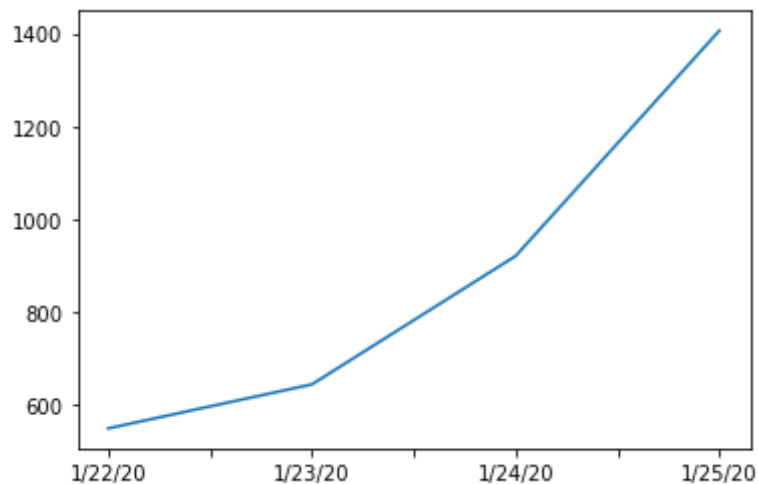
```
Out[13]: <matplotlib.legend.Legend at 0x1f10bd94e80>
```



To calculate a good measure

```
In [14]: covid19_aggregated.loc["China"][:4].plot()
```

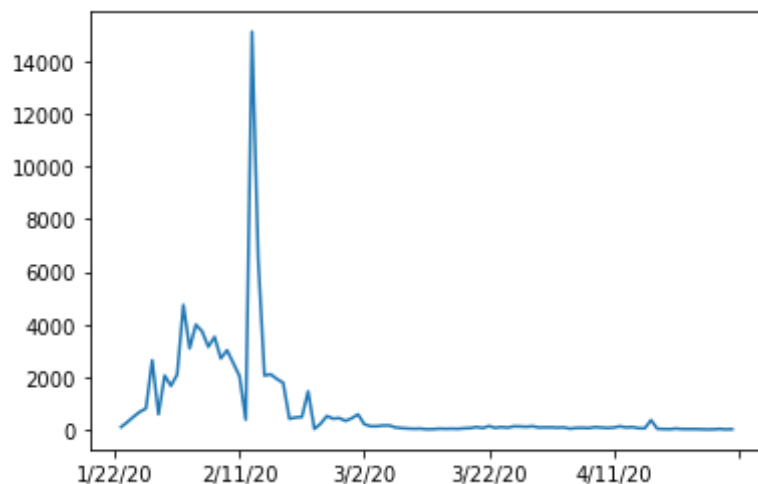
Out[14]: <AxesSubplot:>



# Calculating and plotting the first derivative

```
In [15]: covid19_aggregated.loc["China"].diff().plot()
```

```
Out[15]: <AxesSubplot:>
```



## Finding maximum infection rate

```
In [16]: covid19_aggregated.loc["China"].diff().max()
```

```
Out[16]: 15136.0
```

```
In [17]: covid19_aggregated.loc["India"].diff().max()
```

```
Out[17]: 1893.0
```

```
In [18]: covid19_aggregated.loc["Italy"].diff().max()
```

```
Out[18]: 6557.0
```

# Finding maximum infection rate for all the countries

```
In [19]: countries = list(covid19_aggregated.index)
max_infection_rates = []
for c in countries :
    max_infection_rates.append(covid19_aggregated.loc[c].diff().max())
covid19_aggregated["max_infection_rate"] = max_infection_rates
```

```
In [20]: covid19_aggregated.head()
```

```
Out[20]:
```

	1/22/20	1/23/20	1/24/20	1/25/20	1/26/20	1/27/20	1/28/20	1/29/20	1/30/20	1/31/20	...	4/22/20	4/23/20	4/24/20	4/25/20	4/26/20
Country/Region																
Afghanistan	0	0	0	0	0	0	0	0	0	0	...	1176	1279	1351	1463	1531
Albania	0	0	0	0	0	0	0	0	0	0	...	634	663	678	712	726
Algeria	0	0	0	0	0	0	0	0	0	0	...	2910	3007	3127	3256	3382
Andorra	0	0	0	0	0	0	0	0	0	0	...	723	723	731	738	738
Angola	0	0	0	0	0	0	0	0	0	0	...	25	25	25	25	26

5 rows × 101 columns



# Creating a new dataframe with only needed columns

```
In [21]: covid19 = pd.DataFrame(covid19_aggregated["max_infection_rate"])
```

```
In [22]: covid19.head()
```

```
Out[22]:
```

	max_infection_rate
Country/Region	
Afghanistan	232.0
Albania	34.0

max_infection_rate	
Country/Region	
Algeria	199.0
Andorra	43.0
Angola	5.0

## Importing a new dataset

```
In [23]: happyreport = pd.read_csv("Desktop/worldwide_happiness_report.csv")
```

```
In [24]: happyreport.head()
```

```
Out[24]:
```

	Overall rank	Country or region	Score	GDP per capita	Social support	Healthy life expectancy	Freedom to make life choices	Generosity	Perceptions of corruption
0	1	Finland	7.769	1.340	1.587	0.986	0.596	0.153	0.393
1	2	Denmark	7.600	1.383	1.573	0.996	0.592	0.252	0.410
2	3	Norway	7.554	1.488	1.582	1.028	0.603	0.271	0.341
3	4	Iceland	7.494	1.380	1.624	1.026	0.591	0.354	0.118
4	5	Netherlands	7.488	1.396	1.522	0.999	0.557	0.322	0.298

## Removing useless columns

```
In [25]: useless_cols = ["Overall rank", "Score", "Generosity", "Perceptions of corruption"]
```

```
In [26]: happyreport.drop(useless_cols, axis=1, inplace=True)
happyreport.head()
```

```
Out[26]:
```

	Country or region	GDP per capita	Social support	Healthy life expectancy	Freedom to make life choices
0	Finland	1.340	1.587	0.986	0.596

	Country or region	GDP per capita	Social support	Healthy life expectancy	Freedom to make life choices
1	Denmark	1.383	1.573	0.996	0.592
2	Norway	1.488	1.582	1.028	0.603
3	Iceland	1.380	1.624	1.026	0.591
4	Netherlands	1.396	1.522	0.999	0.557

## Changing the indices of the dataframe

```
In [27]: happyreport.set_index("Country or region", inplace=True)
```

```
In [28]: happyreport.head()
```

```
Out[28]:
```

	GDP per capita	Social support	Healthy life expectancy	Freedom to make life choices
Country or region				
Finland	1.340	1.587	0.986	0.596
Denmark	1.383	1.573	0.996	0.592
Norway	1.488	1.582	1.028	0.603
Iceland	1.380	1.624	1.026	0.591
Netherlands	1.396	1.522	0.999	0.557

## Joining the two datasets

## COVID19 DATASET

```
In [29]: covid19.head()
```

```
Out[29]:
```

	max_infection_rate
--	--------------------



Country/Region	max_infection_rate
----------------	--------------------

Country/Region
----------------

Afghanistan	232.0
Albania	34.0
Algeria	199.0
Andorra	43.0
Angola	5.0

```
In [30]: covid19.shape
```

```
Out[30]: (187, 1)
```

## WORLD HAPPINESS DATASET

```
In [31]: happyreport.head()
```

```
Out[31]:
```

	GDP per capita	Social support	Healthy life expectancy	Freedom to make life choices
--	----------------	----------------	-------------------------	------------------------------

Country or region
-------------------

Finland	1.340	1.587	0.986	0.596
Denmark	1.383	1.573	0.996	0.592
Norway	1.488	1.582	1.028	0.603
Iceland	1.380	1.624	1.026	0.591
Netherlands	1.396	1.522	0.999	0.557

```
In [32]: happyreport.shape
```

```
Out[32]: (156, 4)
```

# Using Join method

```
In [33]: newdata = covid19.join(happyreport, how="inner")
newdata.head()
```

```
Out[33]:
```

	max_infection_rate	GDP per capita	Social support	Healthy life expectancy	Freedom to make life choices
<b>Afghanistan</b>	232.0	0.350	0.517	0.361	0.000
<b>Albania</b>	34.0	0.947	0.848	0.874	0.383
<b>Algeria</b>	199.0	1.002	1.160	0.785	0.086
<b>Argentina</b>	291.0	1.092	1.432	0.881	0.471
<b>Armenia</b>	134.0	0.850	1.055	0.815	0.283

## Correlation matrix

```
In [34]: newdata.corr()
```

```
Out[34]:
```

	max_infection_rate	GDP per capita	Social support	Healthy life expectancy	Freedom to make life choices
<b>max_infection_rate</b>	1.000000	0.250118	0.191958	0.289263	0.078196
<b>GDP per capita</b>	0.250118	1.000000	0.759468	0.863062	0.394603
<b>Social support</b>	0.191958	0.759468	1.000000	0.765286	0.456246
<b>Healthy life expectancy</b>	0.289263	0.863062	0.765286	1.000000	0.427892
<b>Freedom to make life choices</b>	0.078196	0.394603	0.456246	0.427892	1.000000

## Visualisation of the results

```
In [42]: newdata.head()
```

```
Out[42]:
```

	max_infection_rate	GDP per capita	Social support	Healthy life expectancy	Freedom to make life choices
--	--------------------	----------------	----------------	-------------------------	------------------------------

	max_infection_rate	GDP per capita	Social support	Healthy life expectancy	Freedom to make life choices
<b>Afghanistan</b>	232.0	0.350	0.517	0.361	0.000
<b>Albania</b>	34.0	0.947	0.848	0.874	0.383
<b>Algeria</b>	199.0	1.002	1.160	0.785	0.086
<b>Argentina</b>	291.0	1.092	1.432	0.881	0.471
<b>Armenia</b>	134.0	0.850	1.055	0.815	0.283

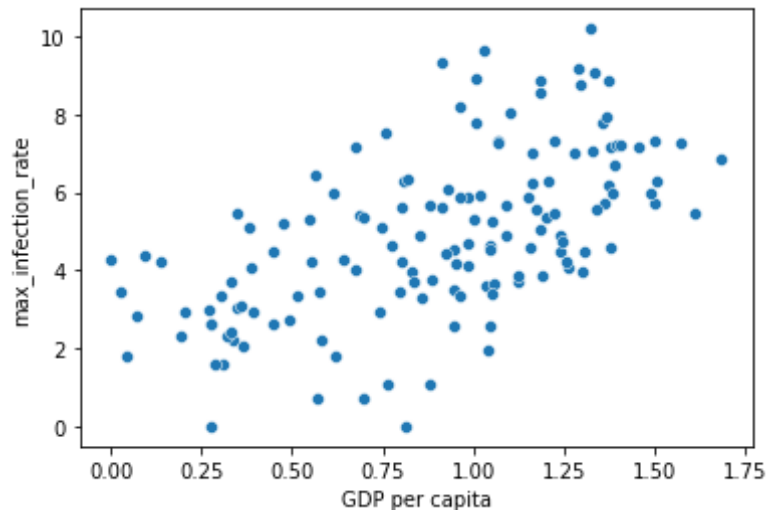
## Plotting GDP vs Maximum infection rate

```
In [35]: x = newdata["GDP per capita"]
y = newdata["max_infection_rate"]
sns.scatterplot(x, nms.log(y))
```

C:\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.

warnings.warn(

```
Out[35]: <AxesSubplot:xlabel='GDP per capita', ylabel='max_infection_rate'>
```

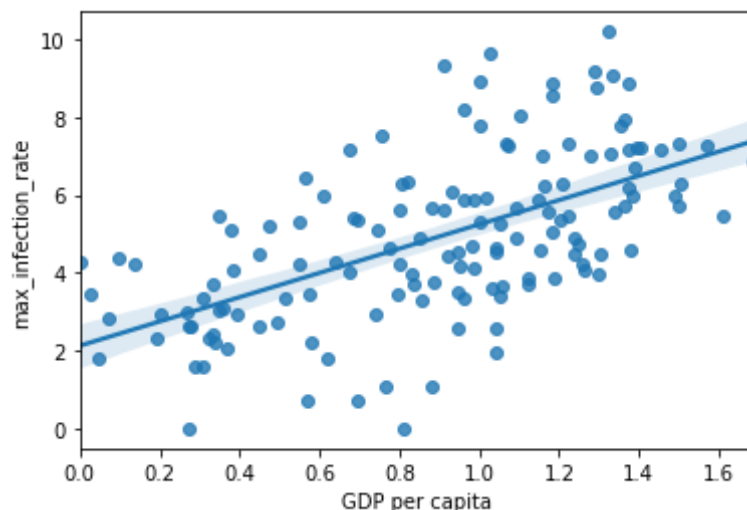


# Using regression plot

```
In [36]: sns.regplot(x,nms.log(y))
```

C:\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.  
warnings.warn()

```
Out[36]: <AxesSubplot:xlabel='GDP per capita', ylabel='max_infection_rate'>
```

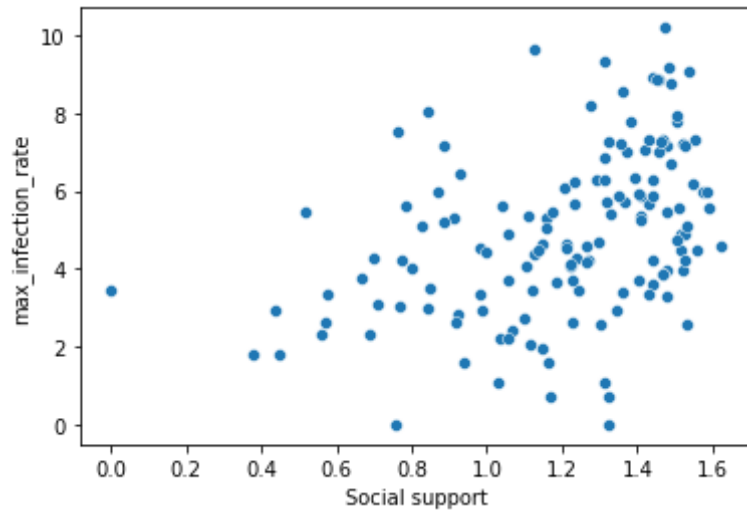


## Plotting social support vs maximum infection rate

```
In [37]: x = newdata["Social support"]  
y = newdata["max_infection_rate"]  
sns.scatterplot(x,nms.log(y))
```

C:\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.  
warnings.warn()

```
Out[37]: <AxesSubplot:xlabel='Social support', ylabel='max_infection_rate'>
```



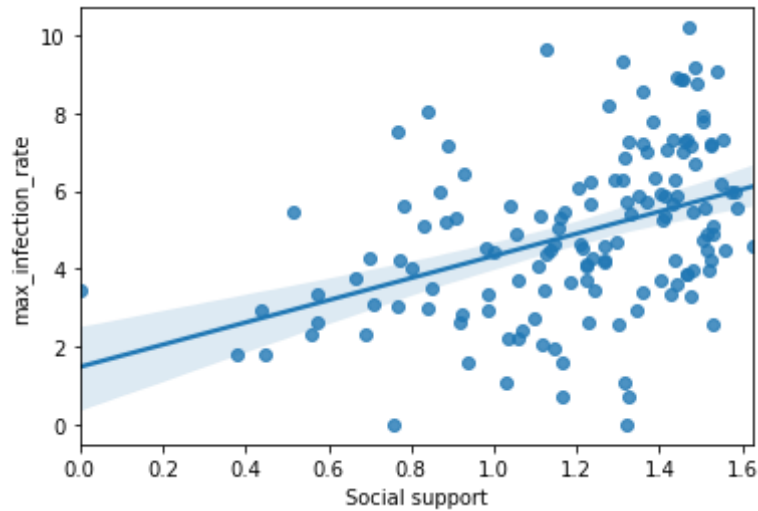
## Regression plot

```
In [38]: sns.regplot(x,nms.log(y))
```

C:\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.

```
warnings.warn(
```

```
Out[38]: <AxesSubplot:xlabel='Social support', ylabel='max_infection_rate'>
```

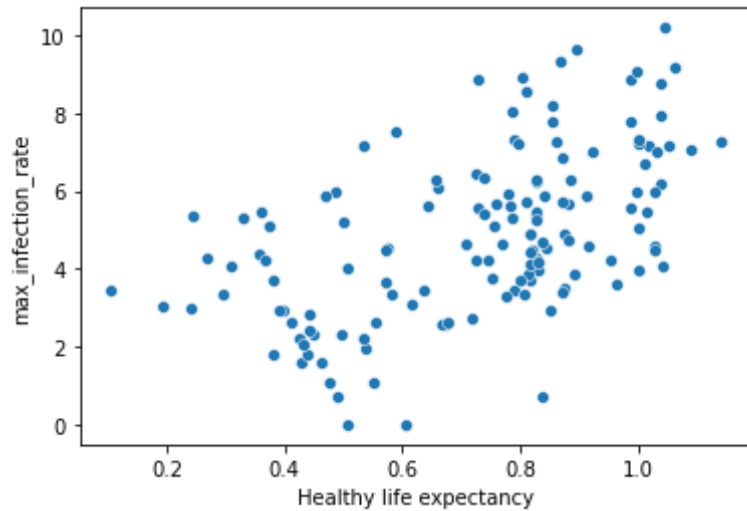


## Plotting healthy life expectancy vs maximum infection rate

```
In [39]: x = newdata["Healthy life expectancy"]  
y = newdata["max_infection_rate"]  
sns.scatterplot(x,nms.log(y))
```

C:\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.  
warnings.warn(

```
Out[39]: <AxesSubplot:xlabel='Healthy life expectancy', ylabel='max_infection_rate'>
```



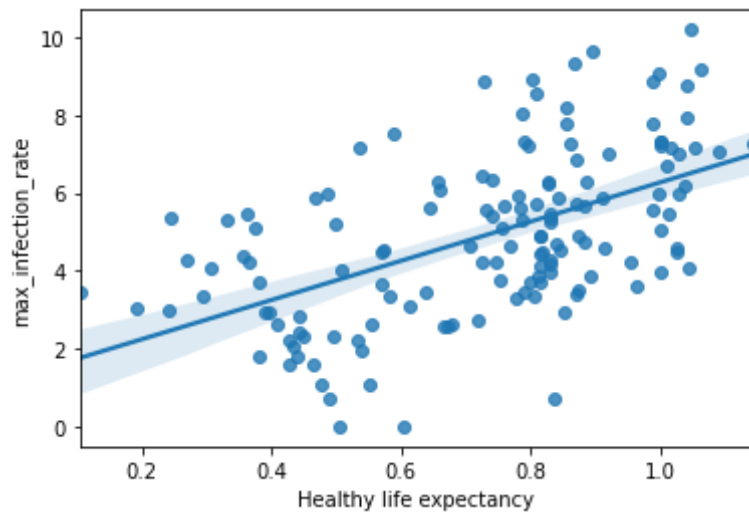
## Regression plot

```
In [40]: sns.regplot(x,nms.log(y))
```

C:\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.

```
warnings.warn(
```

```
Out[40]: <AxesSubplot:xlabel='Healthy life expectancy', ylabel='max_infection_rate'>
```



In [ ]:

In [ ]:

In [ ]:

In [ ]:

In [ ]: