

avs-nexus

July 25, 2024

```
[1]: import json
import folium
import requests
import mimetypes
import http.client
import numpy as np
import pandas as pd
import plotly.express as px
import matplotlib.pyplot as plt
import plotly.graph_objects as go
from folium.plugins import HeatMap
from pandas.io.json import json_normalize
import matplotlib.pyplot as plt
import seaborn as sns

[2]: # conn= http.client.HTTPSConnection("api.covid19api.com")
# payload=''
# headers={}
# conn.request("GET", "/summary", payload, headers)
# res=conn.getresponse()
# data=res.read().decode('UTF-8')

[3]: # pd.set_option("display.max_rows", None, "display.max_columns", None)

[4]: # covid1=json.loads(data)

[5]: # pd.json_normalize(covid1['Countries'], sep=",")
# df=pd.DataFrame(covid1['Countries'])
# df

[6]: # covid2=df.drop(columns=['ID', 'CountryCode', 'Slug', 'Date', 'Premium'], axis=1)
# covid2=covid2.sort_values(by=['TotalConfirmed'], ascending=False)
# covid2['Country'] = covid2['Country'].replace('Russian Federation', 'Russia')
# covid2['Country'] = covid2['Country'].replace('Iran, Islamic Republic of', 'Iran')
# covid2['Country'] = covid2['Country'].replace('Venezuela (Bolivarian Republic)', 'Venezuela')
```

```
# covid3=covid2
# covid2
```

```
[7]: # SARS
sars_03 = pd.read_csv(r"C:
↳\Users\Asus\Desktop\Project\DV\DV_Project\sars_2003_complete_dataset_clean.
↳csv",parse_dates=['Date'])
sars_03 = sars_03[['Date', 'Country', 'Cumulative number of case(s)', 'Number of_
↳deaths', 'Number recovered']]
sars_03.columns = ['Date', 'Country', 'Cases', 'Deaths', 'Recovered']
sars_03 = sars_03.groupby(['Date', 'Country'])
sars_03 = sars_03.sum().reset_index()
sars_03['Cases'] = sars_03['Cases'].fillna(0)
sars_03['Deaths'] = sars_03['Deaths'].fillna(0)
sars_03['Recovered'] = sars_03['Recovered'].fillna(0)
sars_latest = sars_03[sars_03['Date'] == max(sars_03['Date'])].reset_index()
sars_latest=sars_latest.sort_values(by=['Cases'],ascending=False)

# H1N1
h1n1_09 = pd.read_csv(r"C:
↳\Users\Asus\Desktop\Project\DV\DV_Project\Pandemic_H1N1_2009.
↳csv",parse_dates=['Update Time'])
h1n1_09 = h1n1_09[['Update Time', 'Country', 'Cases','Deaths']]
h1n1_09 = h1n1_09.groupby(['Update Time', 'Country'])
h1n1_09 = h1n1_09.sum().reset_index()
h1n1_09['Cases'] = h1n1_09['Cases'].fillna(0)
h1n1_09['Deaths'] = h1n1_09['Deaths'].fillna(0)
h1n1_09['Deaths'] = h1n1_09['Deaths'].astype('int')
h1n1_latest=h1n1_09[h1n1_09['Update Time'] == max(h1n1_09['Update Time'])].
↳reset_index()
h1n1_latest=h1n1_latest.sort_values(by=['Cases'],ascending=False)
h1n1_latest=h1n1_latest.drop([43])

# EBOLA
ebola_14 = pd.read_csv(r"C:
↳\Users\Asus\Desktop\Project\DV\DV_Project\ebola_2014_2016_clean.
↳csv",parse_dates=['Date'])
ebola_14 = ebola_14[['Date', 'Country',
'Cumulative no. of confirmed, probable and suspected cases', 'Cumulative no. of_
↳confirmed, probable and suspected deaths']]
ebola_14.columns = ['Date', 'Country', 'Cases', 'Deaths']
ebola_14 = ebola_14.groupby(['Date', 'Country'])
ebola_14 = ebola_14.sum().reset_index()
ebola_14['Cases'] = ebola_14['Cases'].fillna(0)
ebola_14['Deaths'] = ebola_14['Deaths'].fillna(0)
ebola_14['Cases'] = ebola_14['Cases'].astype('int')
```

```

ebola_14['Deaths'] = ebola_14['Deaths'].astype('int')
ebola_latest = ebola_14[ebola_14['Date'] == max(ebola_14['Date'])].reset_index()
ebola_latest=ebola_latest.sort_values(by=['Cases'],ascending=False)

#MERS
mers_latest = pd.read_csv(r"C:
↪\Users\Asus\Desktop\Project\DV\DV_Project\country_count_latest.csv")
mers_latest=mers_latest.sort_values(by=['Confirmed'],ascending=False)

```

```
[8]: sars_latest
```

```

[8]:      index      Date      Country  Cases  Deaths  Recovered
2      2508 2003-07-11      China    5327      348      4941
7      2513 2003-07-11  Hong Kong SAR, China  1755      298      1433
26     2532 2003-07-11      Taiwan, China    671       84       507
1      2507 2003-07-11      Canada     250       38       194
21     2527 2003-07-11      Singapore    206       32       172
29     2535 2003-07-11      United States    75        0        67
30     2536 2003-07-11      Viet Nam     63        5        58
16     2522 2003-07-11      Philippines    14        2        12
6      2512 2003-07-11      Germany     10        0         9
14     2520 2003-07-11      Mongolia     9         0         9
27     2533 2003-07-11      Thailand     9         2         7
5      2511 2003-07-11      France        7         1         6
13     2519 2003-07-11      Malaysia     5         2         3
10     2516 2003-07-11      Italy         4         0         4
28     2534 2003-07-11      United Kingdom  4         0         4
8      2514 2003-07-11      India         3         0         3
18     2524 2003-07-11      Republic of Korea  3         0         3
24     2530 2003-07-11      Sweden         3         0         3
9      2515 2003-07-11      Indonesia     2         0         2
22     2528 2003-07-11      South Africa     1         1         0
25     2531 2003-07-11      Switzerland     1         0         1
23     2529 2003-07-11      Spain          1         0         1
0      2506 2003-07-11      Brazil          1         0         1
20     2526 2003-07-11      Russian Federation  1         0         0
19     2525 2003-07-11      Romania         1         0         1
17     2523 2003-07-11      Republic of Ireland  1         0         1
12     2518 2003-07-11      Macao SAR, China    1         0         1
11     2517 2003-07-11      Kuwait          1         0         1
4      2510 2003-07-11      Finland          1         0         1
3      2509 2003-07-11      Colombia          1         0         1
15     2521 2003-07-11      New Zealand     1         0         1

```

```
[9]: h1n1_latest
```

```
[9]:      index      Update Time      Country  Cases  Deaths
116  1802 2009-07-06 09:00:00 United States of America 33902    170
72   1758 2009-07-06 09:00:00 Mexico 10262    119
18   1704 2009-07-06 09:00:00 Canada 7983    25
115  1801 2009-07-06 09:00:00 United Kingdom 7447    3
21   1707 2009-07-06 09:00:00 Chile 7376    14
..   ...   ...   ...   ...   ...
106  1792 2009-07-06 09:00:00 Syria 1    0
75   1761 2009-07-06 09:00:00 Myanmar 1    0
82   1768 2009-07-06 09:00:00 Palau 1    0
84   1770 2009-07-06 09:00:00 Papua New Guinea 1    0
135  1821 2009-07-06 09:00:00 Virgin Islands 1    0
```

[135 rows x 5 columns]

```
[10]: ebola_latest
```

```
[10]:      index      Date      Country  Cases  Deaths
6     2375 2016-03-23 Sierra Leone 14122    3955
2     2371 2016-03-23 Liberia 10680    4813
0     2369 2016-03-23 Guinea 3804    2536
4     2373 2016-03-23 Nigeria 20    8
3     2372 2016-03-23 Mali 8    6
9     2378 2016-03-23 United States of America 4    1
1     2370 2016-03-23 Italy 1    0
5     2374 2016-03-23 Senegal 1    0
7     2376 2016-03-23 Spain 1    0
8     2377 2016-03-23 United Kingdom 1    0
```

```
[11]: mers_latest
```

```
[11]:      Country  Confirmed
19      Saudi Arabia 2058
18  Republic of Korea 186
24  United Arab Emirates 87
10      Jordan 28
15      Oman 24
17      Qatar 19
8      Iran 6
23  United Kingdom 5
11      Kuwait 4
21      Tunisia 3
6      Germany 3
20      Thailand 3
0      Algeria 2
25  United States of America 2
16      Philippines 2
```

13	Malaysia	2
14	Netherlands	2
1	Austria	2
12	Lebanon	2
5	France	2
9	Italy	1
7	Greece	1
22	Turkey	1
4	Egypt	1
3	China	1
2	Bahrain	1
26	Yemen	1

```
[12]: # fig = px.treemap(covid3, names='Country', parents=[""]*len(covid3),
      ↪values='TotalConfirmed')
      # fig.add_trace(go.Treemap(
      #     labels = list(covid3.Country),
      #     parents = [""]*len(covid3),
      #     values = list(covid3.TotalConfirmed),
      #     textinfo = "label+value+percent parent",
      #     ),row = 1, col = 1)
      # fig.show()
```

```
[13]: # m=folium.Map(tiles="Stamen Terrain",min_zoom=1.5)
```

```
[14]: #ERROR
      url='https://raw.githubusercontent.com/python-visualization/folium/master/
      ↪examples/data'
      country_shapes=f'{url}/world-countries.json'
```

```
[15]: # folium.Choropleth(
      #     geo_data=country_shapes,
      #     min_zoom=2,
      #     name='Covid-19',
      #     data=covid3,
      #     columns=['Country','TotalConfirmed'],
      #     key_on='feature.properties.name',
      #     fill_color='OrRd',
      #     nan_fill_color='white',
      #     legend_name='Total Confirmed Covid Cases',
      # ).add_to(m)
```

```
[16]: # covid3.update(covid3['TotalConfirmed'].map('Total Confirmed:{}'.format))
      # covid3.update(covid3['TotalRecovered'].map('Total Recovered:{}'.format))
```

```
[14]: coordinates=pd.read_csv(r"C:
      ↪\Users\Asus\Desktop\Project\DV\DV_Project\world_country_and_usa_states_latitude_and_longitude
      ↪.csv")
coordinates = coordinates.rename(columns={'country': 'Country'})
coordinates= coordinates.drop(columns=['country_code',
      ↪'usa_state_code', 'usa_state_latitude', 'usa_state_longitude', 'usa_state'])
coordinates
```

```
[14]:
```

	latitude	longitude	Country
0	42.546245	1.601554	Andorra
1	23.424076	53.847818	United Arab Emirates
2	33.939110	67.709953	Afghanistan
3	17.060816	-61.796428	Antigua and Barbuda
4	18.220554	-63.068615	Anguilla
..
240	15.552727	48.516388	Yemen
241	-12.827500	45.166244	Mayotte
242	-30.559482	22.937506	South Africa
243	-13.133897	27.849332	Zambia
244	-19.015438	29.154857	Zimbabwe

[245 rows x 3 columns]

```
[15]: # covid_final=pd.merge(covid3,coordinates,on='Country')
```

```
[16]: def plotDot(point):
      folium.CircleMarker(location=[point.latitude,point.longitude],
                           radius=5,
                           weight=2,
                           popup=[point.Country,point.TotalConfirmed,point.
      ↪TotalRecovered],
                           fill_color='#000000').add_to(m)
```

```
[20]: # covid_final.apply(plotDot,axis=1)
      # m.fit_bounds(m.get_bounds())
      # m
```

```
[21]: # m1=folium.Map(tiles="StamenToner",min_zoom=2)
```

```
[22]: # deaths=covid_final['TotalDeaths'].astype(float)
      # lat=covid_final['latitude'].astype(float)
      # lon=covid_final['longitude'].astype(float)
      # m1=m1.add_child(HeatMap(zip(lat,lon,deaths),radius=0))
```

```
[23]: # def plotDot(point):
      #     folium.CircleMarker(location=[point.latitude,point.longitude],
      #                           radius=5,
```

```
# weight=2,
# popup=[point.Country,point.TotalDeaths],
# fill_color='#000000').add_to(m1)
```

```
[24]: # covid_final.apply(plotDot,axis=1)
      # m1.fit_bounds(m1.get_bounds())
      # m1
```

```
[17]: h_df=h1n1_latest
fig = px.treemap(h_df, names='Country', parents=[""]*len(h_df), values='Cases')
fig.add_trace(go.Treemap(
    labels = list(h_df.Country),
    parents = [""]*len(h_df),
    values = list(h_df.Cases),
    textinfo = "label+value+percent parent",
    ),row = 1, col = 1)
fig.show()
```

```
[18]: #ERROR IN URL
m6=folium.Map(tiles="Stamen Terrain",min_zoom=1.5)
url='https://raw.githubusercontent.com/python-visualization/folium/master/
examples/data'
country_shapes=f'{url}/world-countries.json'
folium.Choropleth(
    geo_data=country_shapes,
    min_zoom=2,
    name='H1N1',
    data=h1n1_latest,
    columns=['Country','Cases'],
    key_on='feature.properties.name',
    fill_color='OrRd',
    nan_fill_color='white',
    legend_name='Total Confirmed H1N1 Cases',
).add_to(m6)
h1n1_final=pd.merge(h1n1_latest,coordinates,on='Country')
h1n1_final
```

```
[18]:
```

	index	Update Time	Country	Cases	Deaths	latitude	\
0	1758	2009-07-06 09:00:00	Mexico	10262	119	23.634501	
1	1704	2009-07-06 09:00:00	Canada	7983	25	56.130366	
2	1801	2009-07-06 09:00:00	United Kingdom	7447	3	55.378051	
3	1707	2009-07-06 09:00:00	Chile	7376	14	-35.675147	
4	1689	2009-07-06 09:00:00	Australia	5298	10	-25.274398	
..	
101	1752	2009-07-06 09:00:00	Libya	1	0	26.335100	
102	1757	2009-07-06 09:00:00	Mauritius	1	0	-20.348404	
103	1792	2009-07-06 09:00:00	Syria	1	0	34.802075	

```

104  1768 2009-07-06 09:00:00          Palau      1      0   7.514980
105  1770 2009-07-06 09:00:00 Papua New Guinea  1      0  -6.314993

```

```

    longitude
0   -102.552784
1   -106.346771
2    -3.435973
3   -71.542969
4   133.775136
..      ...
101   17.228331
102   57.552152
103   38.996815
104  134.582520
105  143.955550

```

[106 rows x 7 columns]

```

[27]: def plotDot(point):
        folium.CircleMarker(location=[point.latitude,point.longitude],
                             radius=5,
                             weight=2,
                             popup=[point.Country,point.Cases,point.Deaths],
                             fill_color='#000000').add_to(m6)

```

```

[28]: h1n1_final.apply(plotDot,axis=1)
m6.fit_bounds(m6.get_bounds())
m6

```

[28]: <folium.folium.Map at 0x1ef57b5b0a0>

```

[29]: s_df=sars_latest
fig = px.treemap(s_df, names='Country', parents=[""]*len(s_df), values='Cases')
fig.add_trace(go.Treemap(
    labels = list(s_df.Country),
    parents = [""]*len(s_df),
    values = list(s_df.Cases),
    textinfo = "label+value+percent parent",
    ),row = 1, col = 1)
fig.show()

```

```

[30]: m4=folium.Map(tiles="Stamen Terrain",min_zoom=1.5)
url='https://raw.githubusercontent.com/python-visualization/folium/master/
    ↪examples/data'
country_shapes=f'{url}/world-countries.json'
folium.Choropleth(
    geo_data=country_shapes,

```



```

min_zoom=2,
name='SARS',
data=sars_latest,
columns=['Country','Cases'],
key_on='feature.properties.name',
fill_color='OrRd',
nan_fill_color='white',
legend_name='Total Confirmed SARS Cases',
).add_to(m4)
sars_final=pd.merge(sars_latest,coordinates,on='Country')
sars_final

```

```

[30]:
index      Date      Country  Cases  Deaths  Recovered  latitude  \
0      2508 2003-07-11      China   5327     348      4941  35.861660
1      2507 2003-07-11      Canada    250      38       194  56.130366
2      2527 2003-07-11    Singapore    206      32       172   1.352083
3      2535 2003-07-11  United States     75       0        67  37.090240
4      2522 2003-07-11    Philippines    14       2        12  12.879721
5      2512 2003-07-11      Germany    10       0         9  51.165691
6      2520 2003-07-11     Mongolia     9       0         9  46.862496
7      2533 2003-07-11     Thailand     9       2         7  15.870032
8      2511 2003-07-11      France     7       1         6  46.227638
9      2519 2003-07-11     Malaysia     5       2         3   4.210484
10     2516 2003-07-11        Italy     4       0         4  41.871940
11     2534 2003-07-11  United Kingdom     4       0         4  55.378051
12     2514 2003-07-11        India     3       0         3  20.593684
13     2530 2003-07-11        Sweden     3       0         3  60.128161
14     2515 2003-07-11     Indonesia     2       0         2  -0.789275
15     2528 2003-07-11  South Africa     1       1         0 -30.559482
16     2531 2003-07-11    Switzerland     1       0         1  46.818188
17     2529 2003-07-11        Spain     1       0         1  40.463667
18     2506 2003-07-11        Brazil     1       0         1 -14.235004
19     2525 2003-07-11      Romania     1       0         1  45.943161
20     2517 2003-07-11      Kuwait     1       0         1  29.311660
21     2510 2003-07-11      Finland     1       0         1  61.924110
22     2509 2003-07-11    Colombia     1       0         1   4.570868
23     2521 2003-07-11   New Zealand     1       0         1 -40.900557

```

```

longitude
0  104.195397
1 -106.346771
2  103.819836
3  -95.712891
4  121.774017
5   10.451526
6  103.846656
7  100.992541

```

```

8      2.213749
9    101.975766
10    12.567380
11    -3.435973
12    78.962880
13    18.643501
14   113.921327
15    22.937506
16     8.227512
17    -3.749220
18   -51.925280
19    24.966760
20    47.481766
21    25.748151
22   -74.297333
23   174.885971

```

```

[31]: def plotDot(point):
        folium.CircleMarker(location=[point.latitude,point.longitude],
                             radius=5,
                             weight=2,
                             popup=[point.Country,point.Cases,point.Deaths],
                             fill_color='#000000').add_to(m4)

```

```

[32]: sars_final.apply(plotDot,axis=1)
      m4.fit_bounds(m4.get_bounds())
      m4

```

```

[32]: <folium.folium.Map at 0x1ef57e7fa00>

```

```

[33]: m_df=mers_latest
      fig = px.treemap(m_df, names='Country', parents=[""]*len(m_df),
                      ↪values='Confirmed')
      fig.add_trace(go.Treemap(
          labels = list(m_df.Country),
          parents = [""]*len(m_df),
          values = list(m_df.Confirmed),
          textinfo = "label+value+percent parent",
          ),row = 1, col = 1)
      fig.show()

```

```

[34]: m5=folium.Map(tiles="Stamen Terrain",min_zoom=1.5)
      url='https://raw.githubusercontent.com/python-visualization/folium/master/
      ↪examples/data'
      country_shapes=f'{url}/world-countries.json'
      folium.Choropleth(
          geo_data=country_shapes,

```

```

min_zoom=2,
name='MERS',
data=mers_latest,
columns=['Country','Confirmed'],
key_on='feature.properties.name',
fill_color='OrRd',
nan_fill_color='white',
legend_name='Total Confirmed MERS Cases',
).add_to(m5)

```

[34]: <folium.features.Choropleth at 0x1ef58152700>

```

[35]: mers_final=pd.merge(mers_latest,coordinates,on='Country')
mers_final

```

```

[35]:

```

	Country	Confirmed	latitude	longitude
0	Saudi Arabia	2058	23.885942	45.079162
1	United Arab Emirates	87	23.424076	53.847818
2	Jordan	28	30.585164	36.238414
3	Oman	24	21.512583	55.923255
4	Qatar	19	25.354826	51.183884
5	Iran	6	32.427908	53.688046
6	United Kingdom	5	55.378051	-3.435973
7	Kuwait	4	29.311660	47.481766
8	Tunisia	3	33.886917	9.537499
9	Germany	3	51.165691	10.451526
10	Thailand	3	15.870032	100.992541
11	Algeria	2	28.033886	1.659626
12	Philippines	2	12.879721	121.774017
13	Malaysia	2	4.210484	101.975766
14	Netherlands	2	52.132633	5.291266
15	Austria	2	47.516231	14.550072
16	Lebanon	2	33.854721	35.862285
17	France	2	46.227638	2.213749
18	Italy	1	41.871940	12.567380
19	Greece	1	39.074208	21.824312
20	Turkey	1	38.963745	35.243322
21	Egypt	1	26.820553	30.802498
22	China	1	35.861660	104.195397
23	Bahrain	1	25.930414	50.637772
24	Yemen	1	15.552727	48.516388

```

[36]: def plotDot(point):
        folium.CircleMarker(location=[point.latitude,point.longitude],
                             radius=5,
                             weight=2,
                             popup=[point.Country,point.Confirmed],

```

```
fill_color='#000000').add_to(m5)
```

```
[37]: mers_final.apply(plotDot,axis=1)
m5.fit_bounds(m5.get_bounds())
m5
```

```
[37]: <folium.folium.Map at 0x1ef581525b0>
```

```
[38]: e_df=ebola_latest
fig = px.treemap(e_df, names='Country', parents=[""]*len(e_df), values='Cases')
fig.add_trace(go.Treemap(
    labels = list(e_df.Country),
    parents = [""]*len(e_df),
    values = list(e_df.Cases),
    textinfo = "label+value+percent parent",
    ),row = 1, col = 1)
fig
```

```
[39]: m3=folium.Map(tiles="Stamen Terrain",min_zoom=1.5)
url='https://raw.githubusercontent.com/python-visualization/folium/master/
↳examples/data'
country_shapes=f'{url}/world-countries.json'
folium.Choropleth(
    geo_data=country_shapes,
    min_zoom=2,
    name='EBOLA',
    data=ebola_latest,
    columns=['Country','Cases'],
    key_on='feature.properties.name',
    fill_color='OrRd',
    nan_fill_color='white',
    legend_name='Total Confirmed EBOLA Cases',
).add_to(m3)
```

```
[39]: <folium.features.Choropleth at 0x1ef58152c40>
```

```
[40]: ebola_final=pd.merge(ebola_latest,coordinates,on='Country')
ebola_final
```

```
[40]:
```

	index	Date	Country	Cases	Deaths	latitude	longitude
0	2375	2016-03-23	Sierra Leone	14122	3955	8.460555	-11.779889
1	2371	2016-03-23	Liberia	10680	4813	6.428055	-9.429499
2	2369	2016-03-23	Guinea	3804	2536	9.945587	-9.696645
3	2373	2016-03-23	Nigeria	20	8	9.081999	8.675277
4	2372	2016-03-23	Mali	8	6	17.570692	-3.996166
5	2370	2016-03-23	Italy	1	0	41.871940	12.567380
6	2374	2016-03-23	Senegal	1	0	14.497401	-14.452362

```

7  2376 2016-03-23      Spain      1      0  40.463667 -3.749220
8  2377 2016-03-23 United Kingdom  1      0  55.378051 -3.435973

```

```

[41]: def plotDot(point):
        folium.CircleMarker(location=[point.latitude,point.longitude],
                             radius=5,
                             weight=2,
                             popup=[point.Country,point.Cases,point.Deaths],
                             fill_color='#000000').add_to(m3)

```

```

[42]: ebola_final.apply(plotDot,axis=1)
m3.fit_bounds(m3.get_bounds())
m3

```

```

[42]: <folium.folium.Map at 0x1ef583d4640>

```

```

[43]: df = pd.read_csv(r'C:\Users\Asus\Desktop\Project\DV\DV_Project\PANDEMIC.csv')
df.sort_values('CONFIRMED', ascending =False)

```

```

[43]:
    EPIDEMIC  CONFIRMED  DEATHS  RECOVERED  FATALITY RATE  \
4  COVID-19  261442954  5215001  236167794      1.994699
0    H1N1      491382   18499    472883      3.764688
3    EBOLA    28646    11323    17323      39.527334
1    SARS     8437     813     7452      9.636127
2    MERS     2562     881     1681      34.387197

```

```

    NUMBER OF COUNTRIES FIRST REPORTED COUNTRY  YEAR
4              224           Wuhan, China  2019
0              213      Veracruz,Mexico  2009
3               11              Guinea  2013
1               29              China  2002
2               27  Jeddah, Saudi Arabia  2012

```

```

[44]: color = ['aqua', 'lightcoral', 'yellowgreen', 'mediumpurple', 'mediumturquoise']
rcolor = ['mediumturquoise', 'mediumpurple', 'yellowgreen', 'lightcoral',
          'aqua']
df2 = df.sort_values('DEATHS')
fig = px.bar(df2, x='EPIDEMIC', y='CONFIRMED',color = "EPIDEMIC")
fig.show()

```

```

[45]: df2 = df.sort_values('CONFIRMED')
fig = px.bar(df2, x='EPIDEMIC', y='CONFIRMED',color = "EPIDEMIC", log_y=True)
fig.show()

```

```

[46]: df2 = df.sort_values('DEATHS')
fig = px.bar(df2, x='EPIDEMIC', y='DEATHS',color = "EPIDEMIC", log_y=True)
fig.show()

```

```
[47]: df2 = df.sort_values('FATILITY RATE')
fig = px.bar(df2, x='EPIDEMIC', y='FATILITY RATE',color = "EPIDEMIC")
fig.show()
```

```
[48]: color = ['aqua', 'lightcoral', 'yellowgreen', 'mediumpurple', 'mediumturquoise']
rcolor = ['mediumturquoise', 'mediumpurple', 'yellowgreen', 'lightcoral', 'aqua']
df2 = df.sort_values('DEATHS')
fig = px.bar(df2, x='EPIDEMIC', y='DEATHS',color = "EPIDEMIC")
fig.show()
```

```
[49]: df2 = df.sort_values('RECOVERED')
fig = px.bar(df2, x='EPIDEMIC', y='RECOVERED',color = "EPIDEMIC")
fig.show()
```

```
[50]: df2 = df.sort_values('RECOVERED')
fig = px.bar(df2, x='EPIDEMIC', y='RECOVERED',color = "EPIDEMIC",log_y=True)
fig.show()
```

```
[51]: df2 = df.sort_values('CONFIRMED')
fig = px.bar(df2, x='EPIDEMIC', y='CONFIRMED',color = "EPIDEMIC")
fig.show()
```

```
[52]: df2 = df.sort_values('NUMBER OF COUNTRIES',ascending=False)
fig = px.bar(df2, x='NUMBER OF COUNTRIES', y='EPIDEMIC',color = "EPIDEMIC",
orientation='h')
fig.show()
```

```
[53]: fig = px.scatter(df2, x="NUMBER OF COUNTRIES", y="FATILITY RATE",size="FATILITY RATE",
color="EPIDEMIC", hover_name="EPIDEMIC")
fig
```

```
[54]: df2 = df.sort_values('FATILITY RATE')
fig = px.bar(df2, x='EPIDEMIC', y='FATILITY RATE',color = "EPIDEMIC")
fig.show()
```

```
[55]: fig = px.scatter(df, x="CONFIRMED", y="DEATHS",size="FATILITY RATE",
color="EPIDEMIC", hover_name="EPIDEMIC", log_x=True, log_y=True)
fig
```

```
[56]: fig = px.scatter(df, x="CONFIRMED", y="RECOVERED",size="FATILITY RATE",
color="EPIDEMIC", hover_name="EPIDEMIC", log_x=True, log_y=True)
fig
```

```
[57]: df2 = df.sort_values('NUMBER OF COUNTRIES',ascending=False)
fig = px.scatter(df2, x='NUMBER OF COUNTRIES', y='EPIDEMIC')
```

```
fig.show()
```

```
[58]: df2 = df.sort_values('DEATHS')
fig = px.area(df2, x='EPIDEMIC', y='CONFIRMED', color = "EPIDEMIC")
fig.show()
```

```
[59]: df2 = df.sort_values('DEATHS')
fig = px.area(df2, x='EPIDEMIC', y='CONFIRMED', color = "EPIDEMIC", log_y=True)
fig.show()
```

```
[60]: df2 = df.sort_values('FATILITY RATE')
fig = px.line(df2, x='EPIDEMIC', y='FATILITY RATE')
fig.show()
```

```
[61]: df2 = df.sort_values('NUMBER OF COUNTRIES', ascending=False)
fig = px.area(df2, x='EPIDEMIC', y='NUMBER OF COUNTRIES')
fig.show()
```

```
[62]: covid = pd.read_csv(r"C:\Users\Asus\Desktop\Project\DV\DV_Project\owid-covid-data1.
↳ csv", parse_dates=['date'])
covid = covid[['continent', 'location', 'date', 'total_cases', '
↳ new_cases', 'new_cases_smoothed', 'total_deaths', 'new_deaths', 'total_vaccinations']]
# covid.columns = ['Date', 'Country', 'Cases', 'Deaths', 'Recovered']
covid = covid.groupby(['date', 'location'])
covid = covid.sum().reset_index()
covid['total_cases'] = covid['total_cases'].fillna(0)
covid['total_deaths'] = covid['total_deaths'].fillna(0)
covid['total_vaccinations'] = covid['total_vaccinations'].fillna(0)
covid_latest = covid[covid['date'] == max(covid['date'])].reset_index()
covid_latest = covid_latest.sort_values(by=['total_cases'], ascending=False)
```

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

```
[64]: # # plot_map(covid, 'total_cases', [(0,"blue"), (1,"red")])
# plot_map(covid, 'total_cases', 'matter')
```

```
[65]: # plot_map(covid, 'total_deaths', 'matter')
```

```
[66]: # plot_map(covid, 'total_vaccinations', 'matter')
```

```
[67]: # def plot_daywise(col, hue):
#     fig = px.bar(covid_latest, x="date", y=col, width=700,
# ↪color_discrete_sequence=[hue])
#     fig.update_layout(title=col, xaxis_title="", yaxis_title="")
#     fig.show()
```

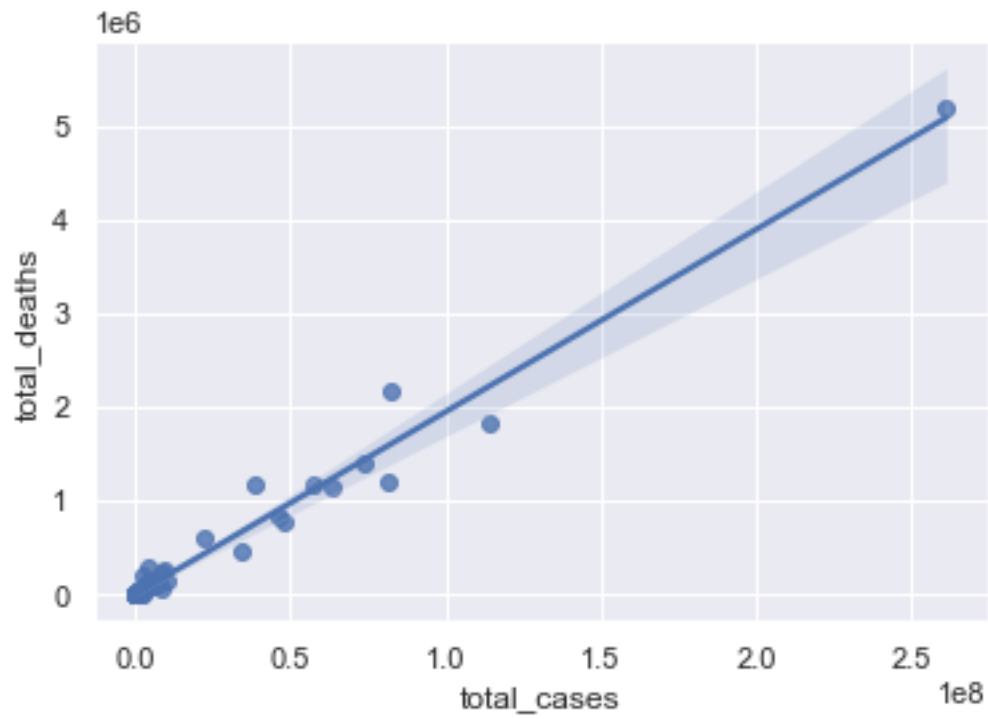
```
[68]: # def plot_daywise_line(col, hue):
#     fig = px.line(covid_latest, x="date", y=col, width=700,
# ↪color_discrete_sequence=[hue])
#     fig.update_layout(title=col, xaxis_title="", yaxis_title="")
#     fig.show()
```

```
[69]: # temp = covid_latest['date'].groupby('date')['total_cases', 'total_deaths'].
# ↪sum().reset_index()
# temp = temp.melt(id_vars="date", value_vars=['total_cases',
# ↪'total_deaths'], var_name='Case', value_name='Count')
# temp.head()

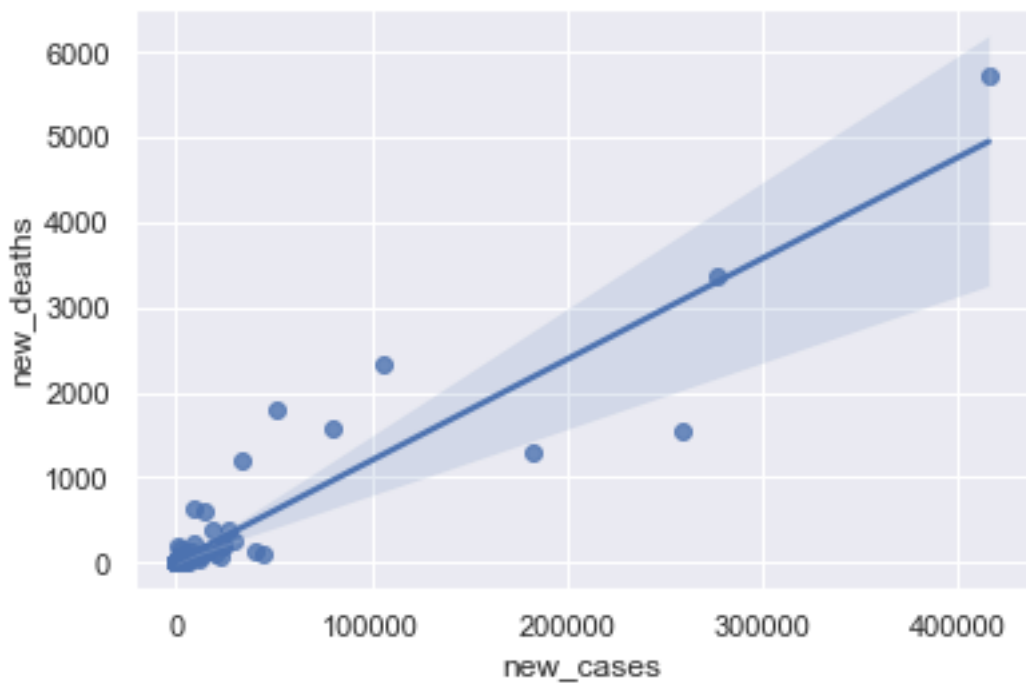
# fig = px.area(temp, x="date", y="Count", color='Case', height=600, width=700,
# ↪title='Cases over time', color_discrete_sequence = [rec, dth,
# ↪act])
# fig.update_layout(xaxis_rangeslider_visible=True)
# fig.show()
```

```
[70]: sns.set_theme(color_codes=True)
```

```
[71]: # covid = sns.load_dataset(r"C:
# ↪\Users\Asus\Desktop\Project\DV\DV_Project\owid-covid-data.csv")
sns.regplot(x="total_cases", y="total_deaths", data=covid_latest);
```

```
[72]: sns.regplot(x="new_cases", y="new_deaths", data=covid_latest);
```



```
[73]: temp = covid[['date', 'total_cases', 'total_deaths']]
temp = temp.melt(id_vars='date', value_vars=[ 'total_cases','total_deaths'],
                var_name='Variable', value_name='Count')
px.line(temp, x='date', y='Count', color='Variable')
```

```
[74]: temp = covid[['date', 'total_cases', 'total_deaths']]
temp = temp.melt(id_vars='date', value_vars=[ 'total_cases','total_deaths'],
                var_name='Variable', value_name='Count')
px.line(temp, x='date', y='Count', color='Variable',log_y=True)
```

```
[75]: fig = px.scatter(covid_latest, x='total_cases', y='total_deaths',
    ↪color='total_cases',
height=700, hover_name='location',
title='Confirmed vs Deaths',
color_discrete_sequence=px.colors.qualitative.Vivid)
fig.update_traces(textposition='top center')
# fig.update_layout(showlegend=False)
# fig.update_layout(xaxis_rangeflider_visible=True)
fig.show()
```

```
[76]: fig = px.scatter(covid_latest, x='total_cases', y='total_deaths',
    ↪color='total_cases',
height=700, hover_name='location', log_x=True, log_y=True,
title='Confirmed vs Deaths',
color_discrete_sequence=px.colors.qualitative.Vivid)
fig.update_traces(textposition='top center')
# fig.update_layout(showlegend=False)
# fig.update_layout(xaxis_rangeflider_visible=True)
fig.show()
```

```
[77]: def plot_bubble(col, pal):
    temp = covid_latest[covid_latest[col]>0].sort_values('location',
    ↪ascending=False)
    fig = px.scatter(temp, x='date', y='location', size=col, color=col,
    ↪height=3000,
                        color_continuous_scale=pal)
    fig.update_layout(yaxis = dict(dtick = 1))
    fig.update(layout_coloraxis_showscale=False)
    fig.show()
```

```
[78]: plot_bubble('new_cases', 'Viridis')
```

```
[79]: def plot_treemap(col):
    fig = px.treemap(covid_latest, path=["location"], values=col, height=700,
                    title=col, color_discrete_sequence = px.colors.qualitative.
    ↪Dark2)
```

```
fig.data[0].textinfo = 'label+text+value'
fig.show()
```

```
[80]: plot_treemap('total_cases')
```

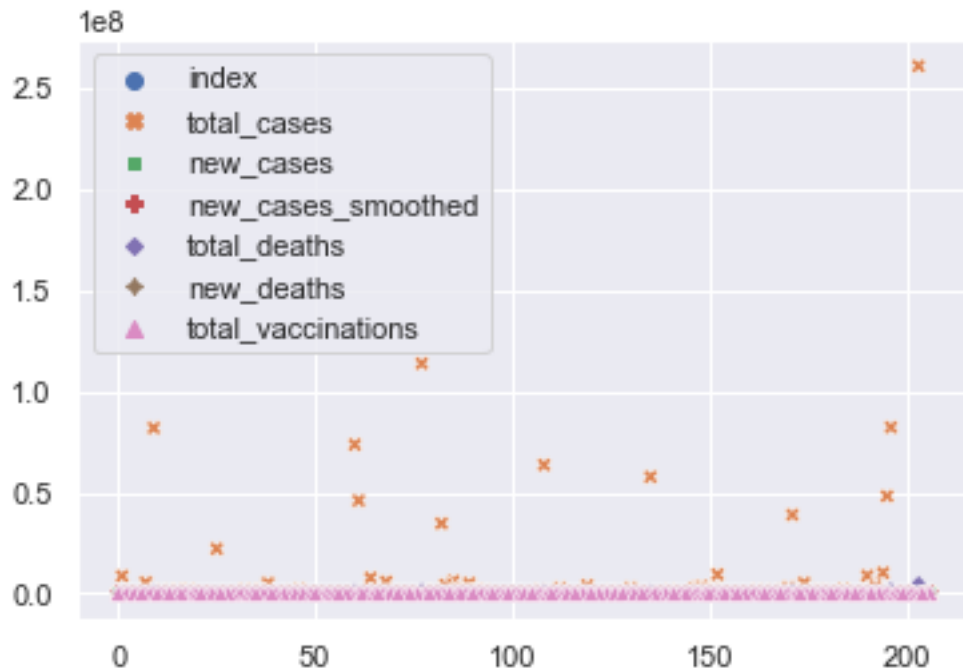
```
[81]: plot_treemap('total_deaths')
```

```
[82]: # plot_treemap('total_vaccinations')
```

```
[83]: fig = px.scatter(covid_latest.sort_values('total_deaths', ascending=False).
    ↪iloc[:20, :],
    x='total_cases', y='total_deaths', color='location',
    ↪size='total_cases',
    height=700, text='location', log_x=True, log_y=True,
    title='Deaths vs Confirmed (Scale is in log10)')
fig.update_traces(textposition='top center')
fig.update_layout(showlegend=False)
fig.update_layout(xaxis_rangeslider_visible=True)
fig.show()
```

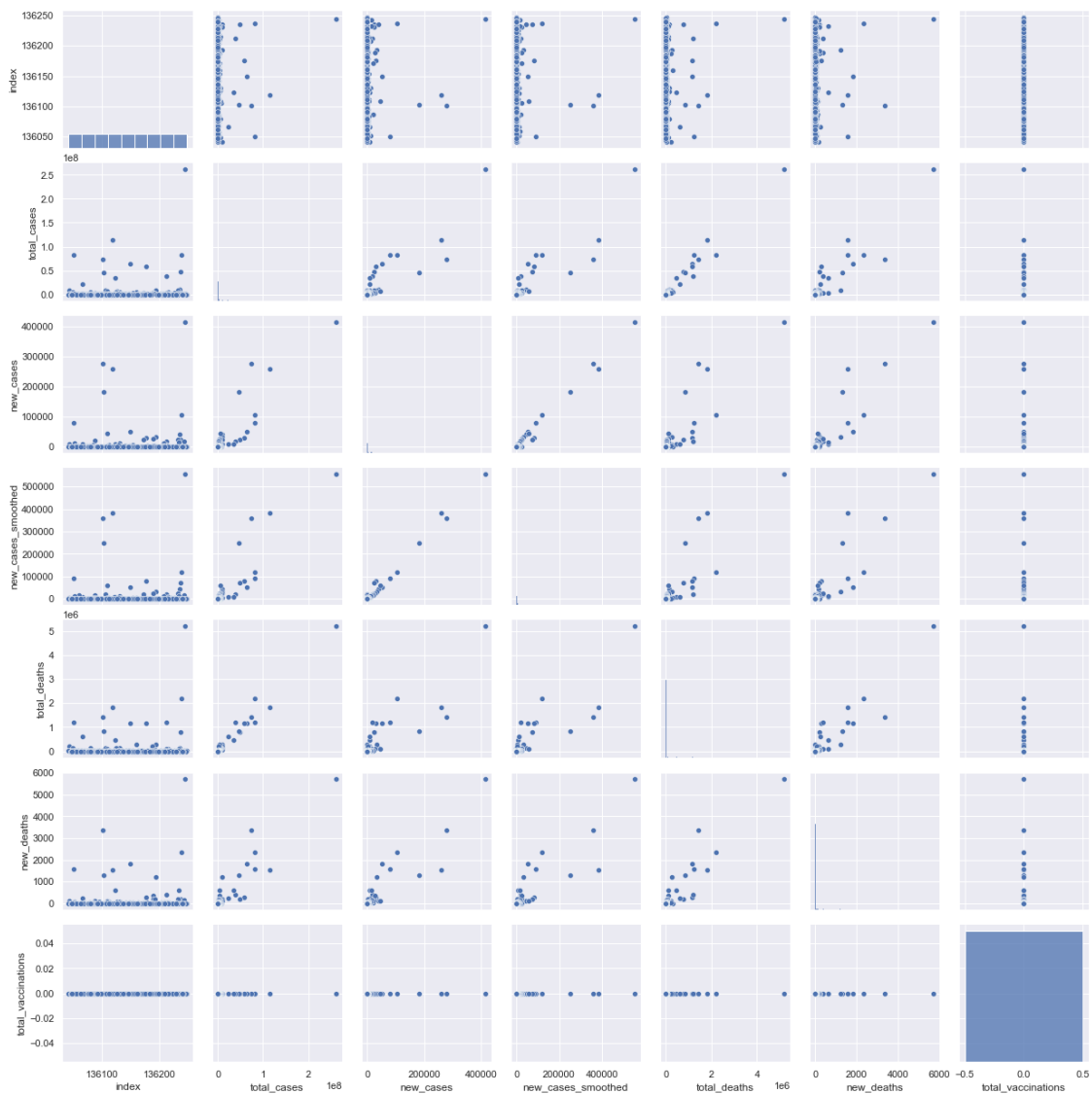
```
[84]: sns.scatterplot(data=covid_latest)
```

```
[84]: <AxesSubplot:>
```



```
[85]: sns.pairplot(data=covid_latest)
```

```
[85]: <seaborn.axisgrid.PairGrid at 0x1ef56423a30>
```



```
[ ]:
```