covid19-notebook

December 30, 2020

1 Visualisation of COVID-19 datas

The goal of this notebook is to display on a Choropleth map informations related to COVID-19 at different moment in time. We will use the open source dataset maintained by *Our World in Data*.

1.0.1 Table of content

1.1 1. Fetching the dataset

First of all we need to import different libraries in order to get the datas, work with them and visualize them.

[1]:
```
import numpy as np # Library to handle data in a vectorized manner
import pandas as pd # Library for data analsysis
import branca.colormap as cmp
pd.set_option('display.max_columns', None)
pd.set_option('display.max_rows', None)
import json # Library to handle JSON files

!pip3 install folium # Install the folium library
import folium # Map rendering library
```

```
Requirement already satisfied: folium in /usr/local/lib/python3.7/site-packages
(0.11.0)
Requirement already satisfied: jinja2>=2.9 in /usr/local/lib/python3.7/site-
packages (from folium) (2.10.3)
Requirement already satisfied: requests in /usr/local/lib/python3.7/site-
packages (from folium) (2.22.0)
Requirement already satisfied: branca>=0.3.0 in /usr/local/lib/python3.7/site-
packages (from folium) (0.4.1)
Requirement already satisfied: numpy in /usr/local/lib/python3.7/site-packages
(from folium) (1.18.1)
Requirement already satisfied: MarkupSafe>=0.23 in
/usr/local/lib/python3.7/site-packages (from jinja2>=2.9->folium) (1.1.1)
Requirement already satisfied: idna<2.9,>=2.5 in /usr/local/lib/python3.7/site-
```

```
packages (from requests->folium) (2.8)
Requirement already satisfied: certifi>=2017.4.17 in
/usr/local/lib/python3.7/site-packages (from requests->folium) (2019.11.28)
Requirement already satisfied: urllib3!=1.25.0,!=1.25.1,<1.26,>=1.21.1 in
/usr/local/lib/python3.7/site-packages (from requests->folium) (1.25.8)
Requirement already satisfied: chardet<3.1.0,>=3.0.2 in
/usr/local/lib/python3.7/site-packages (from requests->folium) (3.0.4)
```

Now that all necessary libraries have been imported, we need to put the dataset into a pandas dataframe. To do so we will use the read_csv function offered by pandas to get datas directly from the link of the repository.

[2]:
```
covid_df = pd.read_csv('https://covid.ourworldindata.org/data/owid-covid-data.
↪csv')
print('Dataset successfully downloaded !')
```

```
Dataset successfully downloaded !
```

Now that the dataset have been downloaded and transformed into a dataframe, we analyse the content of the dataframe.

[3]:
```
covid_df.head()
```

[3]:

	iso_code	continent	location	date	total_cases	new_cases
0	AFG	Asia	Afghanistan	2020-02-24	1.0	1.0
1	AFG	Asia	Afghanistan	2020-02-25	1.0	0.0
2	AFG	Asia	Afghanistan	2020-02-26	1.0	0.0
3	AFG	Asia	Afghanistan	2020-02-27	1.0	0.0
4	AFG	Asia	Afghanistan	2020-02-28	1.0	0.0

	new_cases_smoothed	total_deaths	new_deaths	new_deaths_smoothed
0	NaN	NaN	NaN	NaN
1	NaN	NaN	NaN	NaN
2	NaN	NaN	NaN	NaN
3	NaN	NaN	NaN	NaN
4	NaN	NaN	NaN	NaN

	total_cases_per_million	new_cases_per_million
0	0.026	0.026
1	0.026	0.000
2	0.026	0.000
3	0.026	0.000
4	0.026	0.000

	new_cases_smoothed_per_million	total_deaths_per_million
0	NaN	NaN
1	NaN	NaN
2	NaN	NaN
3	NaN	NaN

```
4                              NaN                          NaN

   new_deaths_per_million  new_deaths_smoothed_per_million  reproduction_rate  \
0                     NaN                              NaN                NaN
1                     NaN                              NaN                NaN
2                     NaN                              NaN                NaN
3                     NaN                              NaN                NaN
4                     NaN                              NaN                NaN

   icu_patients  icu_patients_per_million  hosp_patients  \
0           NaN                       NaN            NaN
1           NaN                       NaN            NaN
2           NaN                       NaN            NaN
3           NaN                       NaN            NaN
4           NaN                       NaN            NaN

   hosp_patients_per_million  weekly_icu_admissions  \
0                        NaN                    NaN
1                        NaN                    NaN
2                        NaN                    NaN
3                        NaN                    NaN
4                        NaN                    NaN

   weekly_icu_admissions_per_million  weekly_hosp_admissions  \
0                                NaN                     NaN
1                                NaN                     NaN
2                                NaN                     NaN
3                                NaN                     NaN
4                                NaN                     NaN

   weekly_hosp_admissions_per_million  new_tests  total_tests  \
0                                 NaN        NaN          NaN
1                                 NaN        NaN          NaN
2                                 NaN        NaN          NaN
3                                 NaN        NaN          NaN
4                                 NaN        NaN          NaN

   total_tests_per_thousand  new_tests_per_thousand  new_tests_smoothed  \
0                       NaN                     NaN                 NaN
1                       NaN                     NaN                 NaN
2                       NaN                     NaN                 NaN
3                       NaN                     NaN                 NaN
4                       NaN                     NaN                 NaN

   new_tests_smoothed_per_thousand  positive_rate  tests_per_case tests_units  \
0                              NaN            NaN             NaN         NaN
1                              NaN            NaN             NaN         NaN
```

```
2                            NaN            NaN             NaN         NaN
3                            NaN            NaN             NaN         NaN
4                            NaN            NaN             NaN         NaN

   total_vaccinations  total_vaccinations_per_hundred  stringency_index  \
0                 NaN                             NaN              8.33
1                 NaN                             NaN              8.33
2                 NaN                             NaN              8.33
3                 NaN                             NaN              8.33
4                 NaN                             NaN              8.33

   population  population_density  median_age  aged_65_older  aged_70_older  \
0  38928341.0              54.422        18.6          2.581          1.337
1  38928341.0              54.422        18.6          2.581          1.337
2  38928341.0              54.422        18.6          2.581          1.337
3  38928341.0              54.422        18.6          2.581          1.337
4  38928341.0              54.422        18.6          2.581          1.337

   gdp_per_capita  extreme_poverty  cardiovasc_death_rate  \
0        1803.987              NaN                597.029
1        1803.987              NaN                597.029
2        1803.987              NaN                597.029
3        1803.987              NaN                597.029
4        1803.987              NaN                597.029

   diabetes_prevalence  female_smokers  male_smokers  handwashing_facilities  \
0                 9.59             NaN           NaN                  37.746
1                 9.59             NaN           NaN                  37.746
2                 9.59             NaN           NaN                  37.746
3                 9.59             NaN           NaN                  37.746
4                 9.59             NaN           NaN                  37.746

   hospital_beds_per_thousand  life_expectancy  human_development_index
0                         0.5            64.83                    0.498
1                         0.5            64.83                    0.498
2                         0.5            64.83                    0.498
3                         0.5            64.83                    0.498
4                         0.5            64.83                    0.498
```

1.2 2. Understanding and preparation of the data

Now that we have correctly save the dataset we need to keep only few columns for this representation. We only want to display the total cases in each country over time. To do so we extract 5 columns: - iso_code - continent - location - date - total_cases

The others columns could be very useful in a more advanced context were the cause of contamination could be searched.

[4]:
```
simp_covid_df = covid_df[['iso_code', 'continent', 'location', 'date',␣
↪'total_cases']]
simp_covid_df['location'].unique()

# We don't want the general number of case so we remove the 'World' entry
simp_covid_df = simp_covid_df[simp_covid_df.location != 'World']
```

Since we only want to check a specific date we need to create a new DataFrame with only the values of the desired day. Here we choose the christmas day.

[5]:
```
xmas_df = simp_covid_df[simp_covid_df['date'] == '2020-12-25']
xmas_df.head(50)
```

[5]:

	iso_code	continent	location	date	\
305	AFG	Asia	Afghanistan	2020-12-25	
601	ALB	Europe	Albania	2020-12-25	
910	DZA	Africa	Algeria	2020-12-25	
1213	AND	Europe	Andorra	2020-12-25	
1498	AGO	Africa	Angola	2020-12-25	
1790	ATG	North America	Antigua and Barbuda	2020-12-25	
2122	ARG	South America	Argentina	2020-12-25	
2426	ARM	Asia	Armenia	2020-12-25	
2765	AUS	Oceania	Australia	2020-12-25	
3074	AUT	Europe	Austria	2020-12-25	
3379	AZE	Asia	Azerbaijan	2020-12-25	
3668	BHS	North America	Bahamas	2020-12-25	
3978	BHR	Asia	Bahrain	2020-12-25	
4280	BGD	Asia	Bangladesh	2020-12-25	
4568	BRB	North America	Barbados	2020-12-25	
4874	BLR	Europe	Belarus	2020-12-25	
5204	BEL	Europe	Belgium	2020-12-25	
5486	BLZ	North America	Belize	2020-12-25	
5775	BEN	Africa	Benin	2020-12-25	
6074	BTN	Asia	Bhutan	2020-12-25	
6368	BOL	South America	Bolivia	2020-12-25	
6668	BIH	Europe	Bosnia and Herzegovina	2020-12-25	
6943	BWA	Africa	Botswana	2020-12-25	
7251	BRA	South America	Brazil	2020-12-25	
7547	BRN	Asia	Brunei	2020-12-25	
7844	BGR	Europe	Bulgaria	2020-12-25	
8139	BFA	Africa	Burkina Faso	2020-12-25	
8413	BDI	Africa	Burundi	2020-12-25	
8751	KHM	Asia	Cambodia	2020-12-25	
9050	CMR	Africa	Cameroon	2020-12-25	
9389	CAN	North America	Canada	2020-12-25	
9674	CPV	Africa	Cape Verde	2020-12-25	
9964	CAF	Africa	Central African Republic	2020-12-25	

10250	TCD	Africa		Chad	2020-12-25
10561	CHL	South America		Chile	2020-12-25
10904	CHN	Asia		China	2020-12-25
11203	COL	South America		Colombia	2020-12-25
11447	COM	Africa		Comoros	2020-12-25
11737	COG	Africa		Congo	2020-12-25
12036	CRI	North America		Costa Rica	2020-12-25
12330	CIV	Africa		Cote d'Ivoire	2020-12-25
12639	HRV	Europe		Croatia	2020-12-25
12932	CUB	North America		Cuba	2020-12-25
13228	CYP	Europe		Cyprus	2020-12-25
13566	CZE	Europe		Czechia	2020-12-25
13860	COD	Africa	Democratic Republic of Congo		2020-12-25
14198	DNK	Europe		Denmark	2020-12-25
14485	DJI	Africa		Djibouti	2020-12-25
14768	DMA	North America		Dominica	2020-12-25
15072	DOM	North America		Dominican Republic	2020-12-25

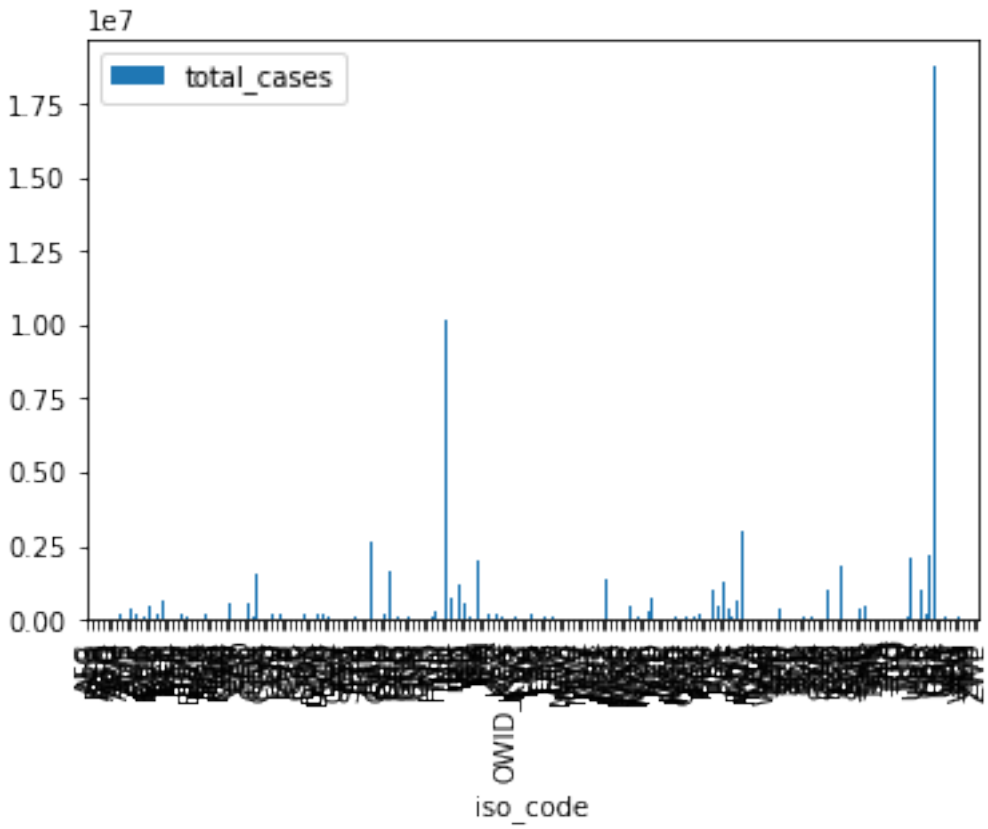
	total_cases
305	50810.0
601	55380.0
910	97441.0
1213	7756.0
1498	17099.0
1790	155.0
2122	1574554.0
2426	156763.0
2765	28297.0
3074	349055.0
3379	211764.0
3668	7788.0
3978	91304.0
4280	507265.0
4568	347.0
4874	183006.0
5204	637246.0
5486	10490.0
5775	3205.0
6074	576.0
6368	153590.0
6668	108891.0
6943	14025.0
7251	7448560.0
7547	152.0
7844	196915.0
8139	6134.0
8413	786.0

```
8751         363.0
9050       26277.0
9389      540939.0
9674       11696.0
9964        4948.0
10250       1971.0
10561     595831.0
10904      95460.0
11203    1574707.0
11447        715.0
11737       6571.0
12036     162990.0
12330      22081.0
12639     203962.0
12932      10900.0
13228      19366.0
13566     664863.0
13860      16472.0
14198     149926.0
14485       5804.0
14768         88.0
15072     165035.0
```

It seems to have large gaps between total cases. This will be an issue in the choropleth map because there will not be enough color shade and thus, the map will not have much interest. Let's check using a plot to make sure it's not only on the 50 first rows.

[6]:
```
xmas_df.plot.bar(x='iso_code', y='total_cases')
```

[6]:
```
<matplotlib.axes._subplots.AxesSubplot at 0x126b9ea50>
```



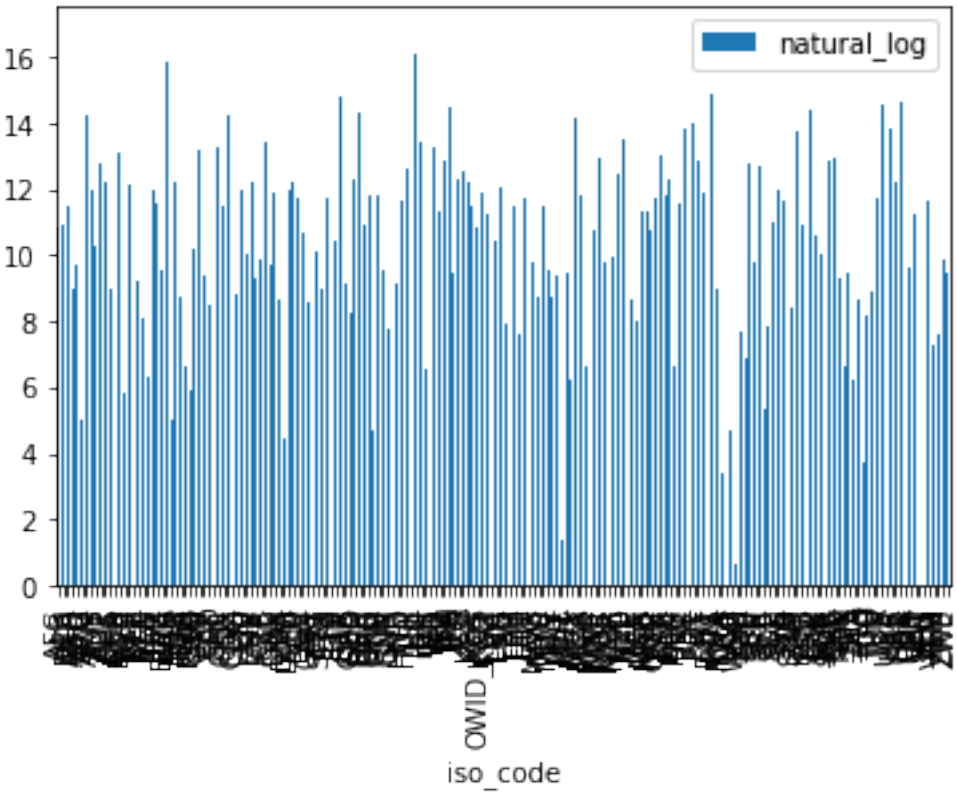
The legend on the plot is not really understandable but what we can see is that the total number of cases have large gaps. In order to solve the problem for the representation we will create a new column with the natural logarithm value of the total_cases in order to normalise the data and see the variations on the map.

[7]:
```
xmas_df['natural_log'] = np.log(xmas_df['total_cases'])
xmas_df.plot.bar(x='iso_code', y='natural_log')
```

```
/usr/local/lib/python3.7/site-packages/ipykernel_launcher.py:1:
SettingWithCopyWarning:
A value is trying to be set on a copy of a slice from a DataFrame.
Try using .loc[row_indexer,col_indexer] = value instead

See the caveats in the documentation: https://pandas.pydata.org/pandas-
docs/stable/user_guide/indexing.html#returning-a-view-versus-a-copy
  """Entry point for launching an IPython kernel.
```

[7]:
```
<matplotlib.axes._subplots.AxesSubplot at 0x127719d10>
```



Now that the data have been normalised and that the values seems more close to each other, let's create the choropleth map in order to visualise the evolution of COVID-19 accross the world.

1.3 3. Building the Choropleth map

We isolated the wanted columns but now we need to display it on a Choropleth map. First we will do a fixed map with the datas on the day of christmas and then we will do an interactive map with all the available datas from February 24th 2020 to today.

To do so we need to get a GeoJSON file in order to link data to countries on the map. This open source file is perfect for what we are looking for. An other solution would have to generate the map using this website.

[8]:
```
!wget --quiet 'https://raw.githubusercontent.com/johan/world.geo.json/master/
↪countries.geo.json'

print('Geo JSON file downloaded !')
```

```
Geo JSON file downloaded !
```

Now that we have the GeoJSON file, let's create a world map centered on the coordinates [0,0].

[9]:
```
world_geo = r'countries.geo.json'
```

And now to create the `Choropleth` map, we will use the *choropleth* method with the following main parameters:

1. world_geo, which is the GeoJSON file.
2. xmas_df, which is the dataframe containing the data.
3. columns, which represents the columns in the dataframe that will be used to create the `Choropleth` map.
4. key_on, which is the key or variable in the GeoJSON file that contains the name of the variable of interest.

[10]:
```
static_map = folium.Map(location=[0, 0], zoom_start=2)

folium.Choropleth(
    geo_data=world_geo,
    data=xmas_df,
    columns=['iso_code', 'natural_log'],
    key_on='feature.id',
    fill_color='OrRd',
    fill_opacity=0.7,
    line_opacity=0.2,
    legend_name='Number of COVID Cases in the world on Christmas 2020'
).add_to(static_map)
```

[10]:
```
<folium.features.Choropleth at 0x127f15a90>
```

[12]:
```
static_map
```

[12]:
```
<folium.folium.Map at 0x127f15050>
```

As we can see from the map, it's mostly the countries in the north of the emispherere that are affected by the virus. It can be explained by different factors. Either it's because of the temperature because she is lower in the north but we can see that in south america there is a large amount of case so it's not only this. Since the virus is more effective in the cold it's a tangible explanation. An other possibility is that the number of test performed and the population amount of the countries are inequal so the number are biased.

Either way we can see that the virus is more present in richer country where the health cares are better. It shows that the life lived by those richer population are not optimal to defeat the virus. It also demonstrate that poorer countries can't be tested enough.

[ ]: