In [1]:

```
import pandas as pd
import numpy as np
import seaborn as sns #visualisation
import matplotlib.pyplot as plt #visualisation
from sklearn.preprocessing import LabelEncoder #To Encoding
import missingno as msno #Visualisation Matrix NaNs
```

STEPS

Dataframe about Coronavirus Group D

In [2]:

```
from utils.folders_tb import downloader
world = downloader(url='https://covid.ourworldindata.org/data/owid-covid-data.csv')
world
```

Out[2]:

	iso_code	continent	location	date	total_cases	new_cases	new_cases_smoothed	total_deaths	new_deaths	nev
0	AFG	Asia	Afghanistan	2020- 02-24	1.0	1.0	NaN	NaN	NaN	
1	AFG	Asia	Afghanistan	2020- 02-25	1.0	0.0	NaN	NaN	NaN	
2	AFG	Asia	Afghanistan	2020- 02-26	1.0	0.0	NaN	NaN	NaN	
3	AFG	Asia	Afghanistan	2020- 02-27	1.0	0.0	NaN	NaN	NaN	
4	AFG	Asia	Afghanistan	2020- 02-28	1.0	0.0	NaN	NaN	NaN	
62060	ZWE	Africa	Zimbabwe	2021- 01-20	29408.0	733.0	736.000	879.0	54.0	
62061	ZWE	Africa	Zimbabwe	2021- 01-21	30047.0	639.0	668.429	917.0	38.0	
62062	ZWE	Africa	Zimbabwe	2021- 01-22	30523.0	476.0	630.571	962.0	45.0	
62063	ZWE	Africa	Zimbabwe	2021- 01-23	31007.0	484.0	589.429	974.0	12.0	
62064	ZWE	Africa	Zimbabwe	2021- 01-24	31320.0	313.0	588.143	1005.0	31.0	

62065 rows × 55 columns

1

In [5]:

 $\#This\ correlation\ Matrix\ of\ NaNs\ is\ showing\ how\ many\ columns\ have\ a\ lot\ of\ non-values\ msno.matrix\ (world)$

Out[5]:

<AxesSubplot:>

```
msno.heatmap(world)
```

Out[4]:

<AxesSubplot:>

4

.....**)**

In [6]:

world.info()

Ranger Index: 62047 entries, 0 to 62048 Date				
# Column	Rang	eIndex: 62047 entries, 0 to 62046		
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2 10cation 62047 non-null object		_		_
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```
dtypes: float64(50), object(5)
```

memory usage: 26.0+ MB

In [8]:

```
#In this section only it is showing how "date" columns has been changed from object to da
tetime to can work in future functions.
world_date = downloader(url='https://covid.ourworldindata.org/data/owid-covid-data.csv')
world_date["date"] = pd.to_datetime(world_date["date"])
world_date.info()
```

<class 'pandas.core.frame.DataFrame'>
RangeIndex: 62047 entries, 0 to 62046
Data columns (total 55 columns):

#	Column	Non-Null Count	Dtype
0	iso code	61694 non-null	
1	continent	61325 non-null	-
2	location	62047 non-null	_
3	date	62047 non-null	_
4	total cases	61450 non-null	
5	new cases	61443 non-null	
6	new cases smoothed	60484 non-null	
7	total deaths	52913 non-null	
8	new deaths	52912 non-null	
9	new deaths smoothed	60484 non-null	
10		61097 non-null	
11	new cases per million	61090 non-null	
12	new cases smoothed per million	60136 non-null	
13		52573 non-null	
14	new deaths per million	52572 non-null	
15	new deaths smoothed per million	60136 non-null	
16		50035 non-null	
17	-	6856 non-null	float64
18	_	6856 non-null	float64
19	hosp patients	8114 non-null	float64
20	hosp patients per million	8114 non-null	float64
21		672 non-null	float64
22		672 non-null 991 non-null	float64
23	weekly_hosp_admissions	991 non-null	float64
24	weekly_hosp_admissions_per_million	991 non-null	float64
25	total_tests	28702 non-null	float64
26	new_tests	28885 non-null	float64
27	total_tests_per_thousand	28702 non-null	float64
28	new_tests_per_thousand	28885 non-null	float64
29	new_tests_smoothed	32573 non-null	
30	<pre>new_tests_smoothed_per_thousand</pre>	32573 non-null	
31	positive_rate	30771 non-null	
32	——————————————————————————————————————	30282 non-null	
33	_	33772 non-null	_
34	_	922 non-null	
35	new_vaccinations	725 non-null	
	new_vaccinations_smoothed	1196 non-null	
37	total_vaccinations_per_hundred	922 non-null	float64
38	new_vaccinations_smoothed_per_million	1196 non-null	float64
39	stringency_index	55360 non-null	float64
40	population	61694 non-null	float64
41	population_density	60390 non-null	float64
42	median_age	59028 non-null	float64
43	aged_65_older	58344 non-null	float64
44	aged_70_older	58694 non-null	float64
45	gdp_per_capita	59064 non-null	float64
46 47	extreme_poverty	40417 non-null	float64 float64
4 7	<pre>cardiovasc_death_rate diabetes_prevalence</pre>	59665 non-null 60368 non-null	float64
49	female smokers	47084 non-null	float64
50	male smokers	46439 non-null	float64
51	handwashing facilities	29886 non-null	float64
52	hospital beds per thousand	54798 non-null	float64
53	life expectancy	61377 non-null	float64
54	human development index	59329 non-null	float64
	_	27527 HOH HULL	

```
memory usage: 26.0+ MB

In [4]:

from utils.mining_data_tb import countries
gbr = countries(df=world, code= "GBR")
prt = countries(df=world, code= "PRT")
ven = countries(df=world, code= "VEN")
tur = countries(df=world, code= "TUR")
esp = countries(df=world, code= "ESP")
gbr

Out[4]:

iso_code continent location date total_cases new_cases_new_cases_smoothed total_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_new_deaths_ne
```

	iso_code	continent	location	date	total_cases	new_cases	new_cases_smoothed	total_deaths	new_deaths	new_d
58315	GBR	Europe	United Kingdom		2.0	2.0	NaN	NaN	NaN	
58316	GBR	Europe	United Kingdom		2.0	0.0	NaN	NaN	NaN	
58317	GBR	Europe	United Kingdom		2.0	0.0	NaN	NaN	NaN	
58318	GBR	Europe	United Kingdom		8.0	6.0	NaN	NaN	NaN	
58319	GBR	Europe	United Kingdom		8.0	0.0	NaN	NaN	NaN	
					•••			•••		
58670	GBR	Europe	United Kingdom		3515796.0	38992.0	42120.429	93469.0	1826.0	
58671	GBR	Europe	United Kingdom		3553773.0	37977.0	40573.714	94765.0	1296.0	
58672	GBR	Europe	United Kingdom		3594094.0	40321.0	38350.286	96166.0	1401.0	
58673	GBR	Europe	United Kingdom		3627746.0	33652.0	37239.429	97518.0	1352.0	
58674	GBR	Europe	United Kingdom		3657857.0	30111.0	36016.714	98129.0	611.0	

360 rows × 55 columns

In [5]:

```
from utils.mining_data_tb import column_erraser

gbr_dd = column_erraser(df=gbr, col1="date", col2='new_cases', col3="new_deaths")
prt_dd = column_erraser(df=prt, col1="date", col2='new_cases', col3="new_deaths")
ven_dd = column_erraser(df=ven, col1="date", col2='new_cases', col3="new_deaths")
tur_dd = column_erraser(df=tur, col1="date", col2='new_cases', col3="new_deaths")
esp_dd = column_erraser(df=esp, col1="date", col2='new_cases', col3="new_deaths")
```

In [5]:

gbr_dd

Out[5]:

	date	new_cases	new_deaths
58315	2020-01-31	2.0	NaN
58316	2020-02-01	0.0	NaN
58317	2020-02-02	0.0	NaN
58318	2020-02-03	6.0	NaN

58319	2020-0 2afé	new_cases	new_deaths
58670	2021-01-20	38992.0	1826.0
58671	2021-01-21	37977.0	1296.0
58672	2021-01-22	40321.0	1401.0
58673	2021-01-23	33652.0	1352.0
58674	2021-01-24	30111.0	611.0

360 rows × 3 columns

In [6]:

```
gbr_dd.dropna(inplace= True)
prt_dd.dropna( inplace= True)
ven_dd.dropna( inplace= True)
tur_dd.dropna( inplace= True)
esp_dd.dropna( inplace= True)
```

In [7]:

gbr_dd

Out[7]:

	date	new_cases	new_deaths
58350	2020-03-06	79.0	1.0
58351	2020-03-07	55.0	1.0
58352	2020-03-08	54.0	0.0
58353	2020-03-09	147.0	1.0
58354	2020-03-10	259.0	4.0
58670	2021-01-20	38992.0	1826.0
58671	2021-01-21	37977.0	1296.0
58672	2021-01-22	40321.0	1401.0
58673	2021-01-23	33652.0	1352.0
58674	2021-01-24	30111.0	611.0

325 rows × 3 columns

In [18]:

```
#Pregunta C8
gbr.corr()
prt.corr()
ven.corr()
tur.corr()
esp.corr()
```

Out[18]:

	total_cases	new_cases	new_cases_smoothed	total_deaths	new_deaths	new_
total_cases	1.000000e+00	5.306517e-01	8.070279e-01	8.793744e-01	5.678626e-02	
new_cases	5.306517e-01	1.000000e+00	6.480925e-01	3.881492e-01	3.449407e-01	
new_cases_smoothed	8.070279e-01	6.480925e-01	1.000000e+00	6.128911e-01	2.063171e-01	
total_deaths	8.793744e-01	3.881492e-01	6.128911e-01	1.000000e+00	-1.186537e- 01	
		0.44040 7 04	0.000174 04	-1.186537e-	1000000 00	

new_deatns	5.678626e-02 total_cases	3.44940/e-01 new_cases	2.0631/1e-01 new_cases_smoothed	total_deaths	1.000000e+00 new_deaths	new_
new_deaths_smoothed	1.435459e-01	1.676450e-01	2.996779e-01	-1.084045e- 01	7.224396e-01	
total_cases_per_million	1.000000e+00	5.306517e-01	8.070279e-01	8.793744e-01	5.678626e-02	
new_cases_per_million	5.306517e-01	1.000000e+00	6.480924e-01	3.881493e-01	3.449407e-01	
new_cases_smoothed_per_million	8.070280e-01	6.480924e-01	1.000000e+00	6.128912e-01	2.063171e-01	
total_deaths_per_million	8.793744e-01	3.881493e-01	6.128911e-01	1.000000e+00	-1.186538e- 01	
new_deaths_per_million	5.678975e-02	3.449424e-01	2.063185e-01	-1.186484e- 01	1.000000e+00	
new_deaths_smoothed_per_million	1.435458e-01	1.676428e-01	2.996795e-01	-1.084037e- 01	7.224380e-01	
reproduction_rate	-2.704262e- 01	-4.507082e- 02	-1.512950e-01	-5.722246e- 01	-4.038914e- 02	
icu_patients	7.542774e-01	3.425082e-01	6.763201e-01	6.440188e-01	6.224790e-01	
icu_patients_per_million	7.542792e-01	3.425099e-01	6.763184e-01	6.440210e-01	6.224782e-01	
hosp_patients	5.838531e-01	4.455477e-01	8.511182e-01	4.489541e-01	6.179149e-01	
hosp_patients_per_million	5.838531e-01	4.455474e-01	8.511181e-01	4.489542e-01	6.179150e-01	
weekly_icu_admissions	-1.034623e- 02	3.329079e-01	1.568608e-01	-4.810924e- 01	3.140879e-01	
weekly_icu_admissions_per_million	-1.033943e- 02	3.329056e-01	1.568674e-01	-4.810873e- 01	3.140857e-01	
weekly_hosp_admissions	1.560007e-02	4.971079e-01	2.239458e-01	-4.724676e- 01	5.053663e-01	
weekly_hosp_admissions_per_million	1.559985e-02	4.971079e-01	2.239456e-01	-4.724678e- 01	5.053663e-01	
total_tests	9.935334e-01	8.079092e-01	7.706481e-01	9.766989e-01	3.465857e-01	
new_tests	NaN	NaN	NaN	NaN	NaN	
total_tests_per_thousand	9.935333e-01	8.079097e-01	7.706486e-01	9.766988e-01	3.465859e-01	
new_tests_per_thousand	NaN	NaN	NaN	NaN	NaN	
new_tests_smoothed	8.506193e-01	5.695179e-01	9.376576e-01	7.644713e-01	2.675102e-01	
new_tests_smoothed_per_thousand	8.506225e-01	5.694967e-01	9.376329e-01	7.644788e-01	2.675182e-01	
positive_rate	3.259592e-01	3.529368e-01	5.685625e-01	2.010074e-01	2.973048e-01	
tests_per_case	-3.352373e- 01	-2.581484e- 01	-4.299592e-01	-2.933662e- 01	-1.069640e- 01	
total_vaccinations	9.962167e-01	3.741305e-01	9.846909e-01	9.900327e-01	5.333185e-01	
new_vaccinations	-1.315952e- 01	1.886223e-01	7.621172e-02	-1.466593e- 01	-6.665631e- 01	
new_vaccinations_smoothed	6.672829e-01	1.935738e-01	7.559267e-01	6.278346e-01	1.513558e-01	
total_vaccinations_per_hundred	9.961878e-01	3.744837e-01	9.846397e-01	9.898782e-01	5.339127e-01	
new_vaccinations_smoothed_per_million	6.673870e-01	1.940945e-01	7.559833e-01	6.279747e-01	1.518692e-01	
stringency_index	3.689012e-01	2.216712e-01	3.386774e-01	2.378178e-01	3.981147e-01	
population	NaN	NaN	NaN	NaN	NaN	
population_density	1.457344e-15	2.342494e-15	1.116635e-15	-3.433818e- 15	-4.037768e- 16	
median_age	NaN	NaN	NaN	NaN	NaN	
aged_65_older	3.290689e-15	2.735455e-18	1.116635e-15	-3.005899e- 16	5.467811e-16	
aged_70_older	4.853011e-15	-1.243446e- 15	-1.772436e-16	8.803784e-16	3.148134e-16	
ado per capita	3.290689e-15	2.735455e-18	-1.772436e-16	-2.557054e- -	4.343872e-16	

V-r	total_cases	new_cases	new_cases_smoothed	15 total_deaths	new_deaths	new
extreme_poverty	NaN	NaN	NaN	NaN	NaN	
cardiovasc_death_rate	4.271298e-16	1.818611e-15	4.070039e-16	2.007449e-15	6.529491e-16	
diabetes_prevalence	-3.145630e- 15	-1.593129e- 15	-4.741268e-16	1.404248e-15	6.806264e-16	
female_smokers	3.290689e-15	2.735455e-18	6.255658e-17	1.516406e-15	-3.594241e- 16	
male_smokers	3.604038e-15	-1.850639e- 15	-1.909256e-15	-3.425526e- 15	3.386764e-16	
handwashing_facilities	NaN	NaN	NaN	NaN	NaN	
hospital_beds_per_thousand	2.769411e-15	1.616520e-15	-7.877495e-18	-8.803784e- 16	-3.148134e- 16	
life_expectancy	3.700407e-15	7.547041e-16	-5.812549e-16	2.667679e-15	4.374249e-16	
human_development_index	-4.853011e- 15	1.243446e-15	-8.271370e-16	-3.378236e- 15	-2.226180e- 16	

50 rows × 50 columns

1

In [8]:

```
#Pregunta A2
worlda_2 = world.drop(['iso_code','continent','location','date', 'aged_65_older', 'tests
_units'],1)
from utils.mining_data_tb import get_top_abs_correlations
get_top_abs_correlations(df=worlda_2, n=7)
```

Out[8]:

new_cases	new_cases_smoothed	0.990950
total_cases	total_deaths	0.977766
new_cases_smoothed	total_deaths	0.976873
icu_patients	hosp_patients	0.975158
new_tests	new_tests_smoothed	0.975068
new_deaths	new_deaths_smoothed	0.973488
total_cases dtype: float64	new_cases_smoothed	0.970599

In [9]:

worlda_2cor = column_erraser(df= worlda_2, col1='new_cases', col2='new_cases_smoothed',
col3='total_deaths', col4='total_cases', col5='new_tests', col6='new_tests_smoothed', col
7= "icu_patients", col8='hosp_patients', col9='new_deaths', col10='new_deaths_smoothed')
worlda_2cor.corr()

Out[9]:

	total_cases	new_cases	new_cases_smoothed	total_deaths	new_deaths	new_deaths_smoothed	icu_ţ
total_cases	1.000000	0.958567	0.970599	0.977766	0.883569	0.906525	0
new_cases	0.958567	1.000000	0.990950	0.967305	0.933366	0.934160	0.
new_cases_smoothed	0.970599	0.990950	1.000000	0.976873	0.922549	0.944571	0.
total_deaths	0.977766	0.967305	0.976873	1.000000	0.919862	0.944834	0.
new_deaths	0.883569	0.933366	0.922549	0.919862	1.000000	0.973488	0.
new_deaths_smoothed	0.906525	0.934160	0.944571	0.944834	0.973488	1.000000	0.
icu_patients	0.882391	0.890198	0.909069	0.881514	0.915093	0.962369	1.
hosp_patients	0.884858	0.897988	0.915676	0.900303	0.913121	0.956632	0.
new_tests	0.883138	0.857692	0.857943	0.846004	0.701788	0.718675	0.
new_tests_smoothed	0.902680	0.853262	0.870133	0.847494	0.664222	0.713038	0

```
In [7]:
#Pregunta A.4.b - Ejecutado debajo de este código en la sección Visualization Javi.
from utils.mining data tb import A 4 b
gbr1M = A_4_b(df= gbr_dd, column='date', key= "date", freq= '1M')
In [8]:
gbr10D = A 4 b(df= gbr dd, column='date', key= "date", freq= '10D')
In [25]:
prt1M = A 4 b(df= prt dd, column='date', key= "date", freq= '1M')
In [26]:
prt10D = A 4 b(df= prt dd, column='date', key= "date", freq= '10D')
In [27]:
ven1M = A 4 b(df= ven dd, column='date', key= "date", freq= '1M')
In [28]:
ven10D = A 4 b(df= ven dd, column='date', key= "date", freq= '10D')
In [29]:
tur1M = A 4 b(df= tur dd, column='date', key= "date", freq= '1M')
In [30]:
tur10D = A 4 b(df= tur dd, column='date', key= "date", freq= '10D')
In [31]:
esp1M = A 4 b(df= esp dd, column='date', key= "date", freq= '1M')
In [32]:
esp10D = A 4 b(df= esp dd, column='date', key= "date", freq= '10D')
Visualization Javi
In [19]:
from utils.visualization tb import b 2 line
gbrb_2_line = b_2_line(df= gbr_dd, x= gbr_dd['date'], y=gbr_dd.columns, title= 'GBR covi
d deaths')
In [20]:
from utils.visualization tb import b 2 bar
gbrb 2 bar = b 2 bar(df= gbr dd, x= gbr dd['date'], y=gbr dd.columns, title= 'GBR covid
deaths')
In [21]:
from utils.visualization tb import b 2 dots
gbrb 2 dots = b 2 dots(df= gbr dd, x= gbr dd['date'], y=gbr dd.columns, title= 'GBR covi
d deaths')
In [33]:
prtb 2 line = b 2 line(df= prt dd, x= prt dd['date'], y=prt dd.columns, title= 'PRT covi
```

d deaths')

```
In [37]:
prtb 2 bar = b 2 bar(df= prt dd, x= prt dd['date'], y=prt dd.columns, title= 'PRT covid
deaths')
In [38]:
prtb 2 dots = b 2 dots(df= prt dd, x= prt dd['date'], y=prt dd.columns, title= 'PRT covi
d deaths')
In [39]:
venb 2 line = b 2 line(df= ven dd, x= ven dd['date'], y=ven dd.columns, title= 'VEN covi
d deaths')
In [40]:
venb 2 bar = b 2 bar(df= ven dd, x= ven dd['date'], y=ven dd.columns, title= 'VEN covid
deaths')
In [42]:
venb 2 dots = b 2 dots(df= ven dd, x= ven dd['date'], y=ven dd.columns, title= 'VEN covi
d deaths')
In [43]:
turb 2 line = b 2 line(df= tur dd, x= tur dd['date'], y=tur dd.columns, title= 'TUR covi
d deaths')
In [44]:
turb_2_bar = b_2_bar(df= tur_dd, x= tur_dd['date'], y=tur_dd.columns, title= 'TUR covid
deaths')
In [45]:
turb 2 dots = b 2 dots(df= tur dd, x= tur dd['date'], y=tur dd.columns, title= 'TUR covi
d deaths')
In [46]:
espb_2_line = b_2_line(df= esp_dd, x= esp_dd['date'], y=esp_dd.columns, title= 'ESP covi
d deaths')
In [47]:
espb 2 bar = b 2 bar(df= esp dd, x= esp dd['date'], y=esp dd.columns, title= 'ESP covid
deaths')
In [48]:
espb 2 dots = b 2 dots(df= esp dd, x= esp dd['date'], y=esp dd.columns, title= 'ESP covi
d deaths')
In [13]:
sns.heatmap(worlda 2cor.corr())
Out[13]:
<AxesSubplot:>
In [32]:
gbrb1M line = b 2 line(df= gbr1M, x= gbr1M.index, y=gbr1M.columns, title= 'GBR 1M covid
deaths')
   F O O T
```

```
ın [ZZ]:
gbr10D line = b 2 line(df= gbr10D, x= gbr10D.index, y=gbr10D.columns, title= 'GBR 10D co
vid deaths')
In [33]:
prt1M line = b 2 line(df= prt1M, x= prt1M.index, y=prt1M.columns, title= 'PRT 1M covid d
eaths')
In [34]:
prt10D line = b 2 line(df= prt10D, x= prt10D.index, y=prt10D.columns, title= 'PRT 10D co
vid deaths')
In [35]:
ven1M line = b 2 line(df= ven1M, x= ven1M.index, y=ven1M.columns, title= 'VEN 1M covid d
eaths')
In [36]:
ven10D line = b 2 line(df= ven10D, x= ven10D.index, y=ven10D.columns, title= 'VEN 10D co
vid deaths')
In [37]:
tur1M line = b 2 line(df= tur1M, x= tur1M.index, y=tur1M.columns, title= 'TUR 1M covid d
eaths')
In [38]:
tur10D line = b 2 line(df= tur10D, x= tur10D.index, y=tur10D.columns, title= 'TUR 10D co
vid deaths')
In [39]:
esp1M line = b 2 line(df= esp1M, x= esp1M.index, y=esp1M.columns, title= 'ESP 1M covid d
eaths')
In [40]:
esp10D line = b 2 line(df= esp10D, x= esp10D.index, y=esp10D.columns, title= 'ESP 10D co
vid deaths')
Visualization Ariadna
In [3]:
#Is the second way: this plot is showing that the 75% of the global Countries about total
deaths are between 250K and more 500K.
from utils.visualization tb import C10c
C10c (df=world)
<Figure size 432x288 with 0 Axes>
In [3]:
#In this graphic is representing total cases and total deaths, so this data it's the tota
1 since the covid-19 begun. United Kingdom and Spain are in top 10 Ranking of the worst c
ountries that didn't manage very well.
from utils.visualization tb import C10a
C10a(df=world)
```

<Figure size 432x288 with 0 Axes>

In [3]:

"""In this graphic has been thought with new cases and new deaths because they are not ac $\it cumulative$ """

#It is showing the position of our countries: Portugal, Venezuela, Turkey, UK and Spain in general ranking in over the world. Venezuela is the first country with less new cases and new deaths being 116 position in new cases and 106 in new deaths respect 190 countries total. Instead of the last country is United Kingdom in 186 position in new cases and deaths. Spain is the second last country being 182/190 and 180/190 in new cases and deaths respectively.

from utils.visualization tb import position

position(df=world, col1="iso_code", col2="date", col3="new_cases", col4="new_deaths", co 15="continent", col6="location", col7="tests units")

4

<Figure size 432x288 with 0 Axes>

Visualization Group C's Data - Total Cases

Visualization Ariadna

In [4]:

#This graphic from Group C is representing how the cases has gone increased during all Co ronavirus pandemic.

from utils.visualization tb import groupC plot

groupC_plot(df=world)

4

<Figure size 432x288 with 0 Axes>

In []: