	The Data (10 marks) The data used for this project comes from ECDC, the European Agency for Disease Prevention and Control, an official agent of the European Union. There are two datasets that are used as part of this project, the official report of the ECDC daily covid cases/deaths across the EU/EEA area and the vaccination data from the same region. These datasets can be found on the following links: Link 1 Link 2 The dataset for the cases/deaths consists of 13 columns in total excluding the indexes. They are date, day, month, year, country, new cases for that date, new deaths for that date, country code, country geographical code, country population and continent. It is important to note that this dataset contains the data only from the 1st of March 2021 until the 28th of October 2021. Data before March were reported on a weekly basis and are part of a different dataset. By observing the dataset, it becomes apparent that some piece of data do not offer any new information. For example we can identify the day/month/year by looking at the respective columns and as a result the date string is redundant. The same also applies to countries which have 3 different identifiers, while the continent column offers nothing as obviously this dataset concerns european countries only. Later when analysing this data we can trim some of these columns so the dataset becomes smaller and easier to handle with a computer programme. The dataset for the vaccine data consists of 12 columns in total excluding the indexes. These are the year-week ISO(week of the year identifier), number of first doses administered, number of first doses refused, number of second doses administered, number of unknown doses administered, number of doses supplied to that country, region , country, population, target group, vaccine and denominator. The official and more detailed data dictionary can be found here. At first sight it is clear that the various european countries have a different reporting system. For instance, some countries report by re
	redundant. The same also applies to countries which have 3 different identifiers, while the continent column offers nothing as obviously this dataset concerns european countries only. Later when analysing this data we can trim some of these columns so the dataset becomes smaller and easier to handle with a computer programme. The dataset for the vaccine data consists of 12 columns in total excluding the indexes. These are the year-week ISO(week of the year identifier), number of first doses administered, number of first doses refused, number of second doses administered, number of unknown doses administered, number of doses supplied to that country, region, country, population, target group, vaccine and denominator. The official and more detailed data dictionary can be found here. At first sight it is clear that the various european countries have a different reporting system. For instance, some countries report by region while some countries don't and just use the country identifier in the region
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:	These datasets are constructed from ECDC using the official reports for covid cases/deaths of member states, so as a result they are the most accurate data available for the EU area. However there are a couple caveats that need to be noted. Firstly, the number of covid cases per country heavily depends on how many covid tests that specific country performs per day. Some countries may not invest heavily in detecting covid so as a result they have lower numbers in new cases. Secondly, since all the data mentioned above describes countries that differ in population significantly, we have to keep in mind that raw numbers offer minimal insight on the covid impact, with percentages and ratios being better descriptors of the situation. Project Aim and Objectives (5 marks) The Covid-19 pandemic has had a great impact on everyday life for the past two years. The general aim of this project is to take some very simple daily reported data for each European country, like daily covid cases, deaths and vaccine doses administred, and convert them to more meaningful information that can give a better insight on the situation so far across Europe. In more detail, the following three steps summarise the aim of this project: 1. The fatality rate of the virus is measured for each European country. The fatality rate equals the total number of covid deaths divided.
	 The fatality rate of the virus is measured for each European country. The fatality rate equals the total number of covid deaths divided by the number of covid cases and is used to described in common terms how deadly the virus is. The fatality rate is a very useful indicator for two reasons. Firstly, the average fatality rate is a good indicator to determine the actual death rate of covid, and secondly it is important to see if the death rate varies significantly between different countries and why. The total cases per 100.000 population are calculated. The official report for each country usually contains the new cases for each day as a simple number. However in order to compare the overall situation between countries and see where covid is more pervalent, it is important to convert those numbers to a ratio per 100.000 population, which allows to compare numbers on equal terms. This ratio is calculated using the following formula:(Events occurance)*100.00/Population. This ratio is afterwards used to classify countries according to covid risk level. Although there is no standard framework to assess risk levels and each authority may use a different model, this project uses a framework developed by the Harvard Global Health Institute and can be found online here. The correlation between vaccine doses administered and deaths from covid is found. A negative correlation is a good indicative that the vaccines do indeed offer strong protection againist death from covid. To calculate the correlation the following mathematical ρx; = cov(X, Y)/σxσγ , which is the standard method when using the pandas' built-in correlation function. Specific Objective(s) Objective 1: Measure the fatality rate of each country. Objective 2: Calculate the case per population ratio and classify countries according to covid risk level.
	Objective 3: Find the correlation between vaccination and death rates. System Design (5 marks) Architecture The following flowchart describes the flow of the programme. Each time a module is run we get a new stage which is described in the task section. Meanwhile the boxes represent the dataframes that exist in the memory. Some abbreviations are used. FR stands for fatality rate while DF for dataframe. Stage 1
	Veccination DF Clean Vaccination DF Clean
	In the following code snippet all python packages that are required for this project are imported and the data from the online source is loaded into two pandas' dataframes. The first 5 rows of each dataframe is printed to confirm that the data was successfully read. import matplotlib.pyplot as plt import pandas import numpy as np import geopandas as gpd df_cases = pandas.read_csv('https://opendata.ecdc.europa.eu/covid19/nationalcasedeath_eueea_daily_ei/csv', df_vacc = pandas.read_csv('https://opendata.ecdc.europa.eu/covid19/vaccine_tracker/csv/data.csv', encoding=#Read the csv from the link the ECDC provides into a pandas dataframe. print(df_cases.head(5)) print(df_vacc.head(5)) #Print the first 5 rows of the dataframe to confirm it was successfully read.
	dateRep day month year cases deaths countriesAndTerritories geoId
	3
:	Since this is not an ongoing project and we know exactly what kind of analysis we want to do beforehand, it is safe to delete some columns that either offer information already given by another column(i.e. country name and country code), or delete columns that give information not needed as part of this project. del df_cases['continentExp'] del df_cases['dateRep'] #del df_cases['countriesAndTerritories'] del df_cases['countryterritoryCode'] del df_vacc['FirstDoseRefused'] del df_vacc['NumberDosesReceived'] del df_vacc['Region'] del df_vacc['TargetGroup'] del df_vacc['TargetGroup'] del df_vacc['Vaccine']
	<pre>del df_vacc['Denominator'] del df_vacc['YearWeekISO'] #Delete columns not used at all further down the analysis print(df_cases.head(5)) print(df_vacc.head(5)) #Check the new format of our dataframes day month year cases deaths countriesAndTerritories geoId popData2020 0 29 10 2021 4232 13 Austria AT 8901064 1 28 10 2021 4407 18 Austria AT 8901064 2 27 10 2021 3388 13 Austria AT 8901064 2 27 10 2021 3388 13 Austria AT 8901064 3 26 10 2021 2765 15 Austria AT 8901064 4 25 10 2021 3655 1 Austria AT 8901064 5 FirstDose SecondDose UnknownDose Population ReportingCountry 0 0 0 8901064 AT</pre>
n [4]:	1 0 0 0 8901064 AT 2 0 0 0 8901064 AT 3 5141 0 0 8901064 AT 4 0 0 0 8901064 AT Anote on deleting columns In general it is not advisable to delete information from our data. Although they may not be needed currently, the scope of the analysis may change in the future. However in this instance, where the project will not be an ongoing system, some columns were deleted to make the debugging part easier and faster, plus make the different outputs of the programme more compact. Check for NaNs Now that we have removed unused columns from our dataframes, we should perform one last step before starting the actual analysis, which to check the entire dataframes for any NaN values. df_cases.isnull().values.any() df_vacc.isnull().values.any() False
	The above function returned false, which means that no NaN values were found. We now know that there are no missing/placeholder data. Objective 1: Calculate the fatality rate To get the fatality rate for each country, first the total amount of deaths and cases have to be calculated. Afterwards this info is passed to another dataframe where the fatality rate is calculated as a new column. The new dataframe is going to be much smaller so it is more efficient to calculate the rate at the new dataframe and avoid redundant calculations. df_cases['sumcases'] = df_cases['cases'].groupby(df_cases['geoId']).transform('sum') #Find the sum of column named 'cases' by 'geoId' and add it to a new column name 'sumcases' df_cases['sumdeaths'] = df_cases['deaths'].groupby(df_cases['geoId']).transform('sum') #Similar with above df_total=df_cases.drop_duplicates(subset = ["geoId"]).copy() #Sum is the same for all rows of each country, so drop duplicates
	<pre>df_total['fatalityrate%'] = (100* df_total['sumdeaths']/df_total['sumcases']) #Find fatality rate in percentage form del df_total['day'] del df_total['year'] del df_total['year'] del df_total['cases'] del df_total['deaths'] #Since we dropped duplicates the columns deleted above make no sense in the new dataframe print(df_total.head(5)) # Print 5 rows to check countriesAndTerritories geoId popData2020 sumcases sumdeaths \ 0</pre>
	Croatia HR 4058165 458090 9142 Gyprus CY 888005 124017 571 fatalityrate% 1 1.364550 243 1.921560 486 4.000047 729 1.995678 972 0.460421 Objective 2a: Calculate cases per 100,000 population To calculate the cases per 100,000 population, first the seven-day average for new case has to be found according to the latest date. As a result only the rows that contain the final 7 days of the data are kept and the rest are deleted. Afterwards the sum of cases and deaths is calculated like in objective 1, but now accounting only for 7 days of the reported data. The end result of the objective is inserted into the
	<pre>df_7day=df_cases.copy() df_7day=df_7day.loc[(df_7day['day'] >= 22) & (df_7day['month'] == 10)] #Dataset used is from 28th so last 7 days are 28,27,26,25,24,23,22 df_7day['sumcases'] = df_7day['cases'].groupby(df_7day['geoId']).transform('sum') #sum of cases for last 7 days df_7day['sevenavg'] = df_7day['sumcases']/7 #divide by 7 to get average df_7day=df_7day.drop_duplicates(subset = ["geoId"]) #drop duplicates since all rows contain the same average per geoId #df_7day.drop(df_7day.columns[[0,1,2,3,4,7,8]], axis = 1, inplace = True) #delete columns by position that were copied from df_cases and make no sense with no duplicates df_7day['casespercapita']=df_7day['sevenavg']*100000/df_7day['popData2020'] #calculate case per capita</pre>
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,	Objective 2b: Classify countries according to covid risk level At this step the dataframe is iterated and each row is assigned a value according to the framework used. This value is stored in the new risklevel column. df_7day['risklevel']='' #add new empty column for i, row in df_7day.iterrows(): if df_7day.loc[i,'casespercapita'] >25: df_7day.loc[i,'risklevel'] = 'red' elif df_7day.loc[i,'casespercapita'] >10: df_7day.loc[i,'risklevel'] = 'orange' elif df_7day.loc[i,'casespercapita'] >1:
	<pre>df_7day.loc[i,'risklevel'] = 'yellow' else: df_7day.loc[i,'risklevel'] = 'green' #iterate columns and add colour according to Harvard framework print(df_7day) day month year cases deaths countriesAndTerritories geoId \ 0 29 10 2021 4232 13</pre>
	1944 29 10 2021 673 4 Finland FI 2187 29 10 2021 6461 33 France FR 2430 29 10 2021 24668 121 Germany DE 2673 29 10 2021 2957 31 Greece EL 2916 29 10 2021 4041 37 Hungary HU 3159 29 10 2021 100 0 Iceland IS 3402 29 10 2021 2598 0 Ireland IE 3645 29 10 2021 24863 50 Italy IT 3888 29 10 2021 2919 64 Latvia LV 4131 29 10 2021 2919 64 Latvia LV 4131 29 10 2021 6 0 Liechtenstein LI 4374 29 10 2021 3519 34 Lithuania LT 4617 29 10 2021 138 0 Luxembourg LU 4860 29 10 2021 15 0 Malta MT 5103 29 10 2021 15 0 Malta MT 5103 29 10 2021 168
	0 8901064 29546 11055 4220.857143 47.419692 red 243 11522440 35599 25976 5085.571429 44.136237 red 486 6951482 37866 23718 5409.428571 77.816911 red 729 4058165 24954 9142 3564.857143 87.844066 red 972 888005 1165 571 166.428571 18.741851 orange 1215 10693939 33278 30705 4754.000000 44.455088 red 1458 5822763 11510 2709 1644.285714 28.238926 red 1701 1328976 12914 1502 1844.857143 138.817943 red 1944 5525292 4860 117622 6382.285714 12.565593 orange 2187 67320216 44676 117622 6382.285714 12.565593 orange 2673 10718565 26956 15801 3850.857143 35.926984 </td
	5589 37958138 48275 76773 6896.428571 18.168511 orange 5832 10295909 6277 18149 896.714286 8.709423 yellow 6075 19328838 112605 46430 16086.428571 83.225016 red 6318 5457873 31066 12977 4438.000000 81.313728 red 6561 2095861 17044 5050 2434.857143 116.174553 red 6804 47332614 13711 87322 1958.714286 4.138192 yellow 7047 10327589 4424 15018 632.000000 6.119531 yellow **Objective 3a: Find vaccine doses per 100.000 population per country** Similar to previous scenarios, we have to find the sum of the vaccine dosages administered by each country. The only difference here is that we have to sum 3 different columns, for the 1st,2nd and unknown dosages. target_columns = ['FirstDose', 'SecondDose', 'UnknownDose']
	<pre>#mark columns to be summed df_vacc['AllDose'] = df_vacc[target_columns].sum(axis=1) #sum per row for now df_vacc['SumDoses'] = df_vacc['AllDose'].groupby(df_vacc['ReportingCountry']).transform('sum') #sum new made column AllDose into SumDoses df_vacc=df_vacc.drop_duplicates(subset = ["ReportingCountry"]) #drop duplicates df_vacc['dosepercapita'] = df_vacc['SumDoses']*100000/df_vacc['Population'] df_vacc.drop(df_vacc.columns[[0,2,1,5]], axis = 1, inplace = True) print(df_vacc.head(5)) Population ReportingCountry SumDoses dosepercapita 0 8901064</pre>
	<pre>df_total['deathspercapita']= df_total['sumdeaths']*100000/df_total['popData2020'] #df_vacc.drop(df_vacc.columns[[0,2,1,4,5,3,9,10,11,6,7,12]], axis = 1, inplace = True) print(df_total) countriesAndTerritories geoId</pre>
	2673 Germany DE 83160/11 4559120 95606 2673 Greece EL 10718565 731167 15801 2916 Hungary HU 9769526 863419 30729 3159 Iceland IS 364134 13413 33 3402 Ireland IE 4964440 438124 5436 3645 Italy IT 59641488 4757231 132004 3888 Latvia LV 1907675 211702 3173 4131 Liechtenstein LI 38747 3557 61 4374 Lithuania LT 2794090 403051 5800 4617 Luxembourg LU 626108 81501 843 4860 Malta MT 514564 37612 460 5103 Netherlands NL 17407585 2108549 18371 5346 Norway NO 5367580 203742 900 5589 Poland PL 37958138 2998891 76773 5832 Portugal PT 10295909 1088133 18149 6075 Romania RO 19328838 1616027 46430
	6318 Slovakia SK 5457873 880635 12977 6561 Slovenia SI 2095861 331206 5050 6804 Spain ES 47332614 5008887 87322 7047 Sweden SE 10327589 1170422 15018 fatalityrate* deathspercapita 0 1.364550 124.198635 243 1.921560 225.438362 486 4.000047 341.193432 729 1.995678 225.274231 972 0.460421 64.301440 1215 1.752482 287.125259 1458 0.707688 46.524305
	1701 0.789691 113.019347 1944 0.735224 20.885774 2187 1.645810 174.720176 2430 2.097027 114.957053 2673 2.161066 147.417122 2916 3.558990 314.539313 3159 0.246030 9.062598 3402 1.240745 109.498755 3645 2.774807 221.329153 3888 1.498805 166.328122 4131 1.714928 157.431543 4374 1.439024 207.581001 4617 1.034343 134.641308 4860 1.223014 89.396071
	\$1.223014
[10]:	<pre>df_total['dosepercapita'] = df_total.geoId.map(df_vacc.set_index('ReportingCountry')['dosepercapita']) #map the dosepercapita from df_vacc to df_total according to geoId=ReportingCountry print(df_total) countriesAndTerritories geoId popData2020 sumcases sumdeaths \ 0</pre>
	1944 Finland FI 5525292 156959 1154 2187 France FR 67320216 7146755 117622 2430 Germany DE 83166711 4559120 95606 2673 Greece EL 10718565 731167 15801 2916 Hungary HU 9769526 863419 30729 3159 Iceland IS 364134 13413 33 3402 Ireland IE 4964440 438124 5436 3645 Italy IT 59641488 4757231 132004 3888 Latvia LV 1907675 211702 3173 4131 Liechtenstein LI 38747 3557 61 4374 Lithuania LT 2794090 403051 5800 4617 Luxembourg LU 626108 81501 843 4860 Malta MT 514564 37612 460 5103 Netherlands NL 17407585 2108549 18371 <
	5589 Poland PL 37958138 2998891 76773 5832 Portugal PT 10295909 1088133 18149 6075 Romania RO 19328838 1616027 46430 6318 Slovakia SK 5457873 880635 12977 6561 Slovenia SI 2095861 331206 5050 6804 Spain ES 47332614 5008887 87322 7047 Sweden SE 10327589 1170422 15018 fatalityrate* deathspercapita dosepercapita 0 1.364550 124.198635 250263.811158 243 1.921560 225.438362 291769.017673 486 4.000047 341.193432 80724.369278 729 1.995678 225.274231 177911.124856 972 0.460421 64.301440 268377.205083 1215 1.752482 287.125259 233329.187683
	1458 0.707688 46.524305 304923.968226 1701 0.789691 113.019347 218102.132770 1944 0.735224 20.885774 581989.150981 2187 1.645810 174.720176 548571.700958 2430 2.097027 114.957053 131397.543183 2673 2.161066 147.417122 353843.420271 2916 3.558990 314.539313 233692.770765 3159 0.246030 9.062598 285275.200888 3402 1.240745 109.498755 310421.356689 3645 2.774807 221.329153 432716.747443 3888 1.498805 166.328122 194185.277891 4131 1.714928 157.431543 118798.874752 4374 1.439024 207.581001 353897.691198 4617 1.034343 134.641308 250164.348643 4860 1.223014 89.396071 311723.323046 5103 0.871263 105.534455 130134.587882
	5103
	g=df_total['deathspercapita'].corr(df_total['dosepercapita']) print(g) -0.2855131011220151 Project Outcome (10 + 10 marks) Overview of Results By now we have run all code modules and seen the output of each one. A brief observation of the data is enough to see that there is a negative correlation between death and vaccination rates in EU as a whole. Moreover the data from the last 7 days show that currently there is no single country in the EU where the virus is contained(green colour), with them majority of them being in red state, meaning the regardless of death rates covid is rapidly spreading. Finally there seems to be a big variation between the fatality rates for each country. Let's see however each objective set at the beginning of the project separately and provide some visualisations. Objective 1: Fatality rate of each country Explanation of Results The following bar chart gives a representation of the distribution of fatality rates for each European country. Bulgaria, Romania and Hungary
:	seem to lead the "race" in fatality rates whilst Cyprus, Norway and Iceland come last. It is apparent even at a first glance that there is a significant variance among european countries. For instance Bulgaria, has a circa 4% fatality rate which is nearly 8 times greater than that of Cyprus. The code snippet underneath the bar chart calculates the coefficient variation(CV) for the fatality rates percentage. As a general rule thumb a CV>1 is considered to be high which is also the case here. This variance could be the result of low vaccination rates in some countries or different strategies in treating it that proved less effective. Visualisation df_total.plot(kind='bar', x='geoId', y='fatalityrate%', xlabel='Country', ylabel='Fatality Rate', title='Fatality Rate', titl
[13]:	### 25 15 10 10 15 10 10 10 10 10 1
1	Objective 2: Covid Risk Classification Explanation of Results According to the framework used for covid risk classification, a country contains the virus if it is labeled as green, whilst yellow, orange and red mean that the virus is spreading. The output of the objective 2 code snippet showed that zero countries fell in the green area. This is confirmed by the visualisation shown below. A map was of Europe was drawn using geopandas. The countries are coloured according to their risk level. Countries in gray/black are not part of the data. It is clear that all of Europe is in high covid risk currently. This is a sign that despite efforts made and the vaccine programme rollout the pandemic is still not under control and further preventive measures may be required in the future.
	<pre>Visualisation world = gpd.read_file(gpd.datasets.get_path('naturalearth_lowres')) # or plot Africa continent ax2 = world[world.continent == 'Europe'].plot(figsize=(10,10), edgecolor=u'black', cmap='gray') t_array = df_7day[['countriesAndTerritories','risklevel']].to_numpy() ax2.set_ylim([30, 75]) ax2.set_xlim([-20, 34.4]) for row in t_array: val1 = row[0] val2 = row[1] if val2 =='red': world[world.name == val1].plot(edgecolor=u'gray', color='red', ax=ax2) elif val2=='orange': world[world.name == val1].plot(edgecolor=u'gray', color='orange', ax=ax2) elif val2=='yellow':</pre>
	<pre>world[world.name == val1].plot(edgecolor=u'gray', color='yellow', ax=ax2) else: world[world.name == val1].plot(edgecolor=u'gray', color='green', ax=ax2) # the place to plot additional vector data (points, lines) plt.ylabel('Latitude') plt.xlabel('Longitude') plt.show() C:\Users\Loukas\anaconda3\envs\my_env\lib\site-packages\geopandas\plotting.py:678: UserWarning: The GeoDataF e you are attempting to plot is empty. Nothing has been displayed. warnings.warn(C:\Users\Loukas\anaconda3\envs\my_env\lib\site-packages\geopandas\plotting.py:678: UserWarning: The GeoDataF e you are attempting to plot is empty. Nothing has been displayed. warnings.warn(75 70-</pre>
	65 - 60 - 55 - 50 - 45 - 40 -
	Some error messages appear. They seem to happen because the border/map for countries is downloaded from external source due to geopandas. Objective 3: Correlation between death and vaccination rates Explanation of Results The correlation matrix underneath shows that there is a negative correlation between the values of deathspercapita and dosepercapita. It expected that vaccines decrease death rate and this is proof that they work. However from this scope we examine the correlation only between vaccine and death rate. In a more realistic scenario it is possible that more variables affect the final outcome.
	Visualisation new = df_total[['deathspercapita', 'dosepercapita']].copy() f = plt.figure(figsize=(15, 15)) plt.matshow(new.corr(), fignum=3) cb = plt.colorbar(fraction=0.1) cb.ax.tick_params(labelsize=14) plt.title('Correlation Matrix', fontsize=16); <figure 0="" 1080x1080="" axes="" size="" with=""> Correlation Matrix 0 1.0</figure>
	0 - 0.8 - 0.6 - 0.4