

COVID-19 Project: Mass. Spread of COVID-19

This project uses the packages tidyverse, which now contains lubridate. Please make sure it is installed before trying to knit this project.

COVID-19 was tracked on a global scale likely with more detail than any pandemics of the past. We are going to look at the different counties of Massachusetts and plot both the number of cases and deaths over time. Additionally, we will try to model and predict the number of cases and deaths in the state overall.

We will discover which counties have the:

- highest growth in cases
- most cases per population
- highest growth in deaths
- most deaths per population

This project is broken down into 4 steps.

1. Get the data
2. Convert the data into something useful
3. Create a model and present the data
4. Identify possible biases

Step 1: Import data in a way that's reproducible

- Install all the packages used in this project and load the corresponding libraries
- Use the tidyverse package to read the csv directly from the source
- The data source is provided by Johns Hopkins University on GitHub at: <https://github.com/CSSEGISandData/COVID-19>
- The exact directory for the download is currently: <https://raw.githubusercontent.com/CSSEGISandData/COVID-19/master/data/us> and the individual files are: time_series_covid19_confirmed_US.csv, and time_series_covid19_deaths_US.csv.

A. Install and load the libraries

```
##
## The downloaded binary packages are in
## /var/folders/h4/0yj9g1kx5bvvgbvm3b1209tr0000gn/T//Rtmp7bSx0H/downloaded_packages

## -- Attaching core tidyverse packages ----- tidyverse 2.0.0 --
## v dplyr      1.1.3      v readr      2.1.4
## v forcats    1.0.0      v stringr    1.5.0
## v ggplot2    3.4.3      v tibble     3.2.1
## v lubridate  1.9.3      v tidyr      1.3.0
```

```
## v purrr      1.0.2
## -- Conflicts ----- tidyverse_conflicts() --
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()   masks stats::lag()
## i Use the conflicted package (<http://conflicted.r-lib.org/>) to force all conflicts to become errors
```

B. - D. Use the Tidyverse package to Read the CSV directly from the data sources.

https://raw.githubusercontent.com/CSSEGISandData/COVID-19/master/csse_covid_19_data/csse_covid_19_time_series/csse_covid_19_time_series.csv

https://raw.githubusercontent.com/CSSEGISandData/COVID-19/master/csse_covid_19_data/csse_covid_19_time_series/csse_covid_19_time_series.csv

```
## Rows: 3342 Columns: 1154
## -- Column specification -----
## Delimiter: ","
## chr      (6): iso2, iso3, Admin2, Province_State, Country_Region, Combined_Key
## dbl (1148): UID, code3, FIPS, Lat, Long_, 1/22/20, 1/23/20, 1/24/20, 1/25/20...
##
## i Use 'spec()' to retrieve the full column specification for this data.
## i Specify the column types or set 'show_col_types = FALSE' to quiet this message.
## Rows: 3342 Columns: 1155
## -- Column specification -----
## Delimiter: ","
## chr      (6): iso2, iso3, Admin2, Province_State, Country_Region, Combined_Key
## dbl (1149): UID, code3, FIPS, Lat, Long_, Population, 1/22/20, 1/23/20, 1/24...
##
## i Use 'spec()' to retrieve the full column specification for this data.
## i Specify the column types or set 'show_col_types = FALSE' to quiet this message.
```

Step 2. Convert the Data into something usefull.

Preview the data.

```
## # A tibble: 6 x 1,154
##       UID iso2 iso3 code3 FIPS Admin2 Province_State Country_Region Lat
##       <dbl> <chr> <chr> <dbl> <dbl> <chr>    <chr>          <chr>    <dbl>
## 1 84001001 US   USA   840  1001 Autauga Alabama      US          32.5
## 2 84001003 US   USA   840  1003 Baldwin Alabama      US          30.7
## 3 84001005 US   USA   840  1005 Barbour Alabama      US          31.9
## 4 84001007 US   USA   840  1007 Bibb Alabama      US          33.0
## 5 84001009 US   USA   840  1009 Blount Alabama      US          34.0
## 6 84001011 US   USA   840  1011 Bullock Alabama      US          32.1
## # i 1,145 more variables: Long_ <dbl>, Combined_Key <chr>, '1/22/20' <dbl>,
## # '1/23/20' <dbl>, '1/24/20' <dbl>, '1/25/20' <dbl>, '1/26/20' <dbl>,
## # '1/27/20' <dbl>, '1/28/20' <dbl>, '1/29/20' <dbl>, '1/30/20' <dbl>,
## # '1/31/20' <dbl>, '2/1/20' <dbl>, '2/2/20' <dbl>, '2/3/20' <dbl>,
## # '2/4/20' <dbl>, '2/5/20' <dbl>, '2/6/20' <dbl>, '2/7/20' <dbl>,
## # '2/8/20' <dbl>, '2/9/20' <dbl>, '2/10/20' <dbl>, '2/11/20' <dbl>,
## # '2/12/20' <dbl>, '2/13/20' <dbl>, '2/14/20' <dbl>, '2/15/20' <dbl>, ...

## # A tibble: 6 x 1,155
##       UID iso2 iso3 code3 FIPS Admin2 Province_State Country_Region Lat
##       <dbl> <chr> <chr> <dbl> <dbl> <chr>    <chr>          <chr>    <dbl>
## 1 84001001 US   USA   840  1001 Autauga Alabama      US          32.5
```

```
## 2 84001003 US      USA      840  1003 Baldwin Alabama      US      30.7
## 3 84001005 US      USA      840  1005 Barbour Alabama      US      31.9
## 4 84001007 US      USA      840  1007 Bibb    Alabama      US      33.0
## 5 84001009 US      USA      840  1009 Blount   Alabama      US      34.0
## 6 84001011 US      USA      840  1011 Bullock  Alabama      US      32.1
## # i 1,146 more variables: Long_ <dbl>, Combined_Key <chr>, Population <dbl>,
## #   '1/22/20' <dbl>, '1/23/20' <dbl>, '1/24/20' <dbl>, '1/25/20' <dbl>,
## #   '1/26/20' <dbl>, '1/27/20' <dbl>, '1/28/20' <dbl>, '1/29/20' <dbl>,
## #   '1/30/20' <dbl>, '1/31/20' <dbl>, '2/1/20' <dbl>, '2/2/20' <dbl>,
## #   '2/3/20' <dbl>, '2/4/20' <dbl>, '2/5/20' <dbl>, '2/6/20' <dbl>,
## #   '2/7/20' <dbl>, '2/8/20' <dbl>, '2/9/20' <dbl>, '2/10/20' <dbl>,
## #   '2/11/20' <dbl>, '2/12/20' <dbl>, '2/13/20' <dbl>, '2/14/20' <dbl>, ...
```

```
cases <- confirmed_us %>%
  pivot_longer(cols = -c(UID:Combined_Key), names_to = "date", values_to = "Cases")%>%
  select(-c(iso2, iso3, code3, FIPS, UID, Country_Region))%>%
  mutate(date = mdy(date))

summary(cases)
```

Pivot the data so each day isn't a separate column.

```
##      Admin2      Province_State      Lat      Long_
## Length:3819906 Length:3819906 Min.   :-14.27 Min.   :-174.16
## Class :character Class :character 1st Qu.: 33.90 1st Qu.: -97.81
## Mode  :character Mode  :character Median : 38.01 Median : -89.49
##                                     Mean  : 36.72 Mean  : -88.64
##                                     3rd Qu.: 41.58 3rd Qu.: -82.31
##                                     Max.   : 69.31 Max.   : 145.67
##      Combined_Key      date      Cases
## Length:3819906 Min.   :2020-01-22 Min.   : -3073
## Class :character 1st Qu.:2020-11-02 1st Qu.:   330
## Mode  :character Median :2021-08-15 Median :   2272
##                                     Mean  :2021-08-15 Mean  :  14088
##                                     3rd Qu.:2022-05-28 3rd Qu.:   8159
##                                     Max.   :2023-03-09 Max.   :3710586
```

```
deaths <- deaths_us %>%
  pivot_longer(cols = -c(UID:Population), names_to = "date", values_to = "deaths")%>%
  select(-c(iso2, iso3, code3, FIPS, UID, Country_Region))%>%
  mutate(date = mdy(date))

summary(deaths)
```

```
##      Admin2      Province_State      Lat      Long_
## Length:3819906 Length:3819906 Min.   :-14.27 Min.   :-174.16
## Class :character Class :character 1st Qu.: 33.90 1st Qu.: -97.81
## Mode  :character Mode  :character Median : 38.01 Median : -89.49
##                                     Mean  : 36.72 Mean  : -88.64
##                                     3rd Qu.: 41.58 3rd Qu.: -82.31
```

```
##                               Max.   : 69.31   Max.   : 145.67
## Combined_Key      Population      date      deaths
## Length:3819906   Min.     :      0   Min.   :2020-01-22   Min.   : -82.0
## Class :character 1st Qu.:   9917   1st Qu.:2020-11-02   1st Qu.:   4.0
## Mode  :character Median :  24892   Median :2021-08-15   Median :   37.0
##                               Mean  :  99604   Mean  :2021-08-15   Mean   :  186.9
##                               3rd Qu.:  64979   3rd Qu.:2022-05-28   3rd Qu.:  122.0
##                               Max.   :10039107   Max.   :2023-03-09   Max.   :35545.0
```

```
#summary(mass_density)
```

Initial Filtering and Joins Filter the other states' data out, then join the data for Cases and Deaths in Massachusetts.

```
## Joining with 'by = join_by(Admin2, Province_State, Lat, Long_, Combined_Key,
## date)'
```

More Mutation and Filtering Because of the population correlation, mutate the Mass rows to add deaths per 1000 and cases per 1000. While we're at it, create a couple other date formats and remove the ... from Long. Additionally, we will filter for 0 Population as this will cause divide by zero errors.

```
## Admin2      Province_State      Lat      Long_
## Length:16002 Length:16002   Min.   :41.29   Min.   : -73.21
## Class :character Class :character 1st Qu.:41.79   1st Qu.: -72.59
## Mode  :character Mode  :character Median :42.24   Median : -71.16
##                               Mean  :42.11   Mean   : -71.47
##                               3rd Qu.:42.37   3rd Qu.: -70.81
##                               Max.   :42.67   Max.   : -70.09
## Combined_Key      date      Cases      Population
## Length:16002   Min.   :2020-01-22   Min.   :      0   Min.   : 11399
## Class :character 1st Qu.:2020-11-02   1st Qu.:  1475   1st Qu.: 124944
## Mode  :character Median :2021-08-15   Median : 23196   Median : 493787
##                               Mean  :2021-08-15   Mean   : 65074   Mean   : 492322
##                               3rd Qu.:2022-05-28   3rd Qu.:104131   3rd Qu.: 789034
##                               Max.   :2023-03-09   Max.   :437431   Max.   :1611699
## deaths      deaths_per_k      cases_per_k      month_year
## Min.   :      0   Min.   :0.0000   Min.   : 0.000   Length:16002
## 1st Qu.:   77   1st Qu.:0.7267   1st Qu.: 7.139   Class :character
## Median :  683   Median :1.8719   Median : 75.010   Mode  :character
## Mean   : 1028   Mean   :1.7397   Mean   :106.850
## 3rd Qu.: 1794   3rd Qu.:2.6503   3rd Qu.:203.338
## Max.   : 4822   Max.   :4.5843   Max.   :368.944
## Lng      month
## Min.   : -73.21   Min.   : 1.000
## 1st Qu.: -72.59   1st Qu.: 3.000
## Median : -71.16   Median : 6.000
## Mean   : -71.47   Mean   : 6.335
## 3rd Qu.: -70.81   3rd Qu.: 9.000
## Max.   : -70.09   Max.   :12.000
```

```
## # A tibble: 6 x 14
```

```
##   Admin2      Province_State   Lat Long_ Combined_Key date      Cases Population
##   <chr>      <chr>            <dbl> <dbl> <chr>      <date>    <dbl>      <dbl>
## 1 Barnstable Massachusetts    41.7 -70.3 Barnstable,~ 2020-01-22    0      212990
## 2 Barnstable Massachusetts    41.7 -70.3 Barnstable,~ 2020-01-23    0      212990
## 3 Barnstable Massachusetts    41.7 -70.3 Barnstable,~ 2020-01-24    0      212990
## 4 Barnstable Massachusetts    41.7 -70.3 Barnstable,~ 2020-01-25    0      212990
## 5 Barnstable Massachusetts    41.7 -70.3 Barnstable,~ 2020-01-26    0      212990
## 6 Barnstable Massachusetts    41.7 -70.3 Barnstable,~ 2020-01-27    0      212990
## # i 6 more variables: deaths <dbl>, deaths_per_k <dbl>, cases_per_k <dbl>,
## #   month_year <chr>, Lng <dbl>, month <dbl>
```

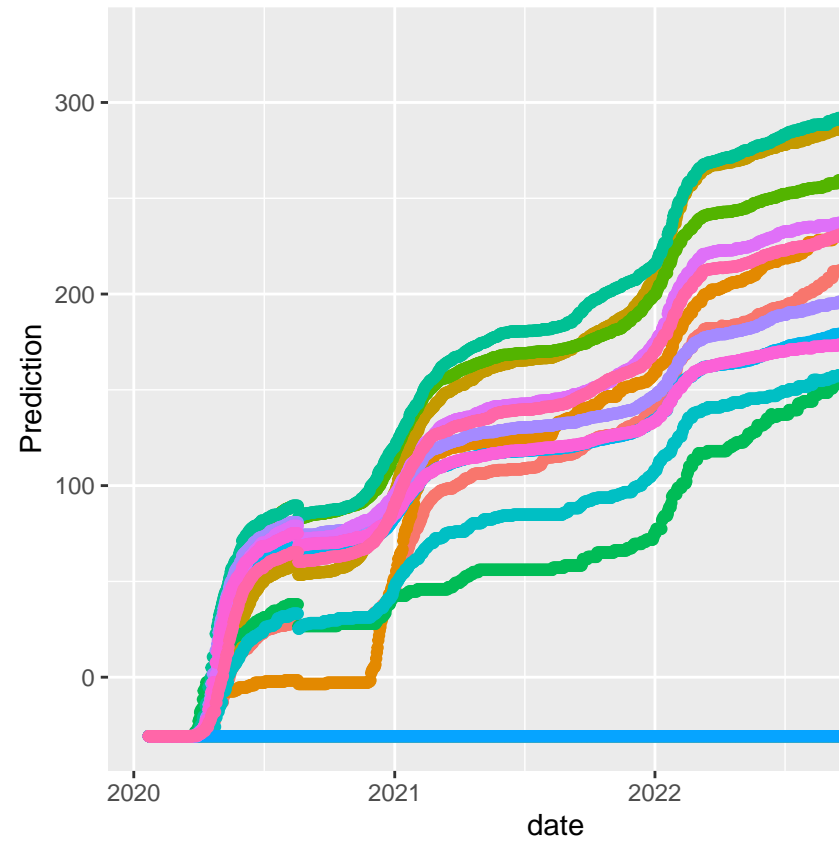
Step 3. Analyze the data, create a model and present everything

Quick Correlation Check To begin with, we do a quick correlation analysis to try to get a better sense of the relationship between the columns of data. I'm looking for correlations between the deaths, cases, and the population.

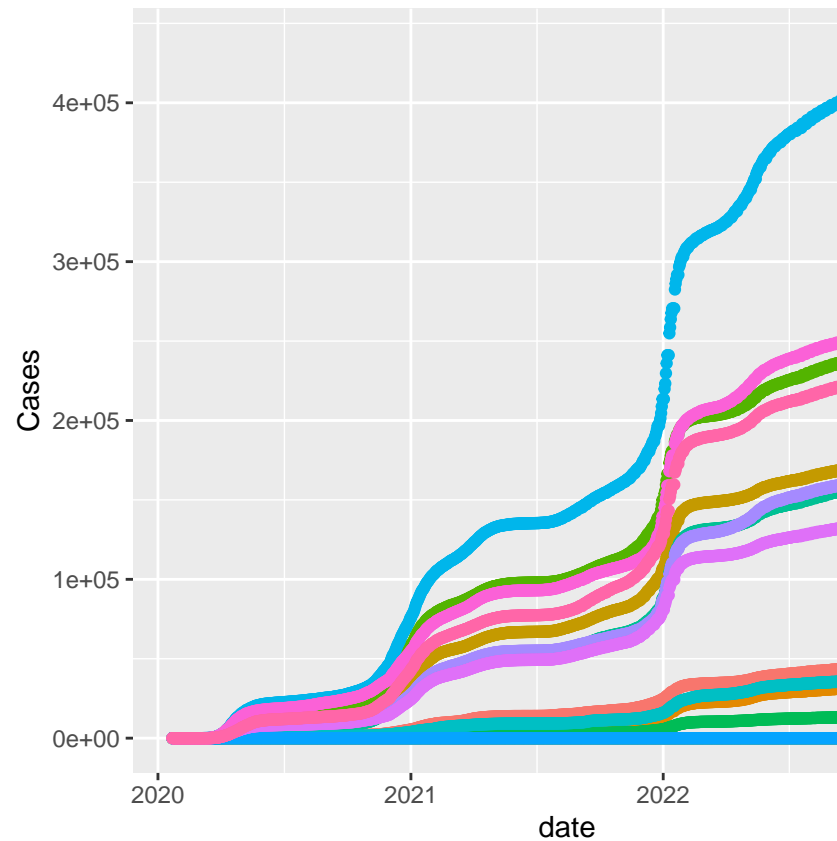
```
## [1] "Deaths & Population: "
## [1] 0.7976191
## [1] "Cases & Population: "
## [1] 0.6327786
## [1] "Cases & Deaths: "
## [1] 0.924681
## [1] "Cases/1000 & Deaths/1000: "
## [1] 0.91824
```

Build a model. We will build a model based on cases per 1000 and deaths per 1000, output the summary, then add the predictions to the Mass. county data.

```
##
## Call:
## lm(formula = cases_per_k ~ deaths_per_k, data = Mass)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -79.130 -38.676   6.675  30.785 146.494
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)  -30.7845     0.5764  -53.41  <2e-16 ***
## deaths_per_k   79.1149     0.2698  293.29  <2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 42.33 on 16000 degrees of freedom
## Multiple R-squared:  0.8432, Adjusted R-squared:  0.8432
## F-statistic: 8.602e+04 on 1 and 16000 DF, p-value: < 2.2e-16
```



Plot Predictions for the individual Counties.



Plot the Actual County Data for comparison

Group the Mass. data by county.

```
##      Admin2      Max_Deaths  Total_Deaths  Max_Cases
## Length:14      Min.   : 0.0      Min.   : 0      Min.   : 0
## Class :character 1st Qu.: 459.8    1st Qu.: 276102    1st Qu.: 36522
## Mode  :character Median : 2050.0    Median : 1382960   Median : 157826
##                      Mean  : 1735.9    Mean  : 1175195   Mean  : 143892
##                      3rd Qu.: 2498.8    3rd Qu.: 1665867   3rd Qu.: 226117
##                      Max.   : 4822.0    Max.   : 3378924   Max.   : 437431
## Total_Cases      Population
## Min.   : 0      Min.   : 11399
## 1st Qu.: 15948862 1st Qu.: 133916
## Median : 81161396 Median : 493787
## Mean   : 74379791 Mean   : 492322
## 3rd Qu.: 118428435 3rd Qu.: 768469
## Max.   : 220834357 Max.   : 1611699

## # A tibble: 6 x 6
##   Admin2      Max_Deaths Total_Deaths Max_Cases Total_Cases Population
##   <chr>      <dbl>      <dbl>      <dbl>      <dbl>      <dbl>
## 1 Barnstable      785      447242      49617      23514236      212990
## 2 Berkshire       480      276606      35456      15223406      124944
## 3 Bristol       2555     1619263     182344     98339486      565217
## 4 Dukes           0           0           0           0          17332
## 5 Essex        3272     2235421     256987     140031284     789034
```

## 6 Franklin	198	108143	14736	6453660	70180
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Step 4: Add Bias Identification

Data Bias

Massachusetts is one state in the United States, out of the entire planet may not be an accurate sample. It is difficult to say how accurate the data itself is or how consistent it is from county to county. Furthermore, as more was known about the COVID-19 virus, methods of accurately identifying cases and deaths are likely to improve. This may skew the data.

Personal Bias

On a personal note, I chose to examine only Massachusetts. I did so based on the basis of my perception of Massachusetts as a place with cutting edge medicine and unbiased data, with both rural and very urban areas. This may be have be completely wrong.