In Depth Analysis of COVID 19 Data for 2 Small European Nations

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Library Used

```
library(tidyverse)
library(lubridate)
```

Import Data

Import COVID 19 Data by the given URL as a .csv file.

```
url_in <- 'https://raw.githubusercontent.com/CSSEGISandData/COVID-19/master/csse_covid_19_data/csse_cov
file names <-
 c('time_series_covid19_confirmed_global.csv',
    'time_series_covid19_deaths_global.csv',
   'time_series_covid19_recovered_global.csv')
urls <- str_c(url_in, file_names)</pre>
global_cases <- read_csv(urls[1])</pre>
## Rows: 289 Columns: 1147
## Delimiter: ","
         (2): Province/State, Country/Region
## dbl (1145): Lat, Long, 1/22/20, 1/23/20, 1/24/20, 1/25/20, 1/26/20, 1/27/20,...
## 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.
global_deaths <- read_csv(urls[2])</pre>
## Rows: 289 Columns: 1147
## -- Column specification ----
## Delimiter: ","
        (2): Province/State, Country/Region
## dbl (1145): Lat, Long, 1/22/20, 1/23/20, 1/24/20, 1/25/20, 1/26/20, 1/27/20,...
## 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: 274 Columns: 1147

## -- Column specification ------

## Delimiter: ","

## chr (2): Province/State, Country/Region

## dbl (1145): Lat, Long, 1/22/20, 1/23/20, 1/24/20, 1/25/20, 1/26/20, 1/27/20,...

##

## 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.

Data Clean Up

Cleaning the raw data so that it's more usable. This involves omitting some column values especially those that are marked NA and name editing.

```
#For Global cases
global_cases <- global_cases %>%
  pivot_longer(cols = -c('Province/State', 'Country/Region', Lat, Long),
              names_to = 'date',
              values_to = 'cases') %>%
  select(-c(Lat, Long)) %>%
  rename(province_state = 'Province/State', country_region = 'Country/Region')
#For Global deaths
global_deaths <- global_deaths %>%
  pivot_longer(cols = -c('Province/State', 'Country/Region', Lat, Long),
              names_to = 'date',
              values to = 'deaths') %>%
  select(-c(Lat, Long)) %>%
  rename(province_state = 'Province/State', country_region = 'Country/Region')
#For Combining global cases and global deaths
global <- global_cases %>%
  full_join(global_deaths) %>%
  mutate(date = mdy(date)) %>%
 filter(cases > 0)
```

Joining with 'by = join_by(province_state, country_region, date)'

```
global <- global %>% filter(cases > 0)
summary(global)
```

```
country_region
                                         date
## province_state
                                                            cases
## Length:306827
                    Length: 306827
                                     Min. :2020-01-22 Min. :
                                                                      1
## Class :character Class :character
                                     1st Qu.:2020-12-12 1st Qu.:
                                                                   1316
## Mode :character Mode :character
                                     Median :2021-09-16 Median :
                                                                   20365
##
                                     Mean :2021-09-11 Mean : 1032863
##
                                     3rd Qu.:2022-06-15 3rd Qu.: 271281
                                     Max. :2023-03-09 Max. :103802702
##
```

```
##
       deaths
##
  Min.
         :
                 0
   1st Qu.:
                 7
## Median:
               214
## Mean
         : 14405
## 3rd Qu.:
              3665
  Max.
        :1123836
#Global by country
global_by_country <- global %>%
 group_by(country_region, date) %>%
 summarize(cases = sum(cases), deaths = sum(deaths))
## 'summarise()' has grouped output by 'country_region'. You can override using
## the '.groups' argument.
#Retrieving Iceland data
Iceland <- global_by_country %>%
 filter(country_region == 'Iceland') %>%
 mutate(new_cases = cases - lag(cases),
        new_deaths = deaths - lag(deaths)) %>%
 select(everything()) %>% drop_na()
summary(Iceland)
## country_region
                          date
                                             cases
                                                              deaths
## Length:1105
                            :2020-02-29
                                                          Min. : 0.00
                     Min.
                                         Min.
                                               :
                                                      1
                                                          1st Qu.: 27.00
## Class:character 1st Qu.:2020-12-01
                                         1st Qu.: 5413
## Mode :character Median :2021-09-03
                                         Median : 10956
                                                          Median : 33.00
##
                     Mean
                            :2021-09-03
                                         Mean : 74533
                                                               : 77.27
                                                          Mean
                    3rd Qu.:2022-06-06
##
                                         3rd Qu.:188924
                                                          3rd Qu.:153.00
                    Max.
##
                            :2023-03-09
                                         Max. :209137
                                                          Max.
                                                                :263.00
##
     new cases
                    new deaths
  Min. :-273.0 Min. :-39.000
##
  1st Qu.: 0.0 1st Qu.: 0.000
## Median :
              2.0
                   Median : 0.000
## Mean : 189.3
                   Mean : 0.238
   3rd Qu.: 47.0
                    3rd Qu.: 0.000
  Max. :7408.0
                   Max. : 52.000
#Retrieving Malta data
Malta <- global_by_country %>%
 filter(country_region == 'Malta') %>%
 mutate(new_cases = cases - lag(cases),
        new_deaths = deaths - lag(deaths)) %>%
 select(everything()) %>% drop_na()
summary(Malta)
## country_region
                          date
                                             cases
                                                              deaths
                            :2020-03-08 Min. :
                                                          Min. : 0.0
## Length:1097
                     Min.
                                                      3
```

Class:character 1st Qu.:2020-12-07

1st Qu.: 10659

1st Qu.:155.0

```
##
   Mode :character
                    Median :2021-09-07
                                         Median : 36553
                                                         Median :445.0
                           :2021-09-07
##
                                         Mean : 51021
                                                               :435.1
                     Mean
                                                         Mean
##
                     3rd Qu.:2022-06-08
                                         3rd Qu.: 95467
                                                         3rd Qu.:723.0
##
                            :2023-03-09
                                                                :828.0
                     Max.
                                         Max.
                                                :117610
                                                         Max.
##
     new_cases
                     new_deaths
         : -42.0
                 Min. :-2.0000
##
  Min.
   1st Qu.: 13.0
                   1st Qu.: 0.0000
   Median: 43.0
                   Median : 0.0000
##
                   Mean : 0.7548
##
   Mean : 107.2
##
   3rd Qu.: 123.0
                   3rd Qu.: 1.0000
  Max.
        :1677.0
                   Max. : 7.0000
```

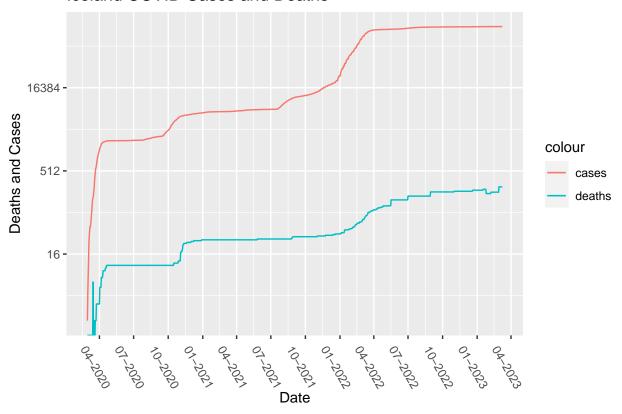
Visualizations

Now that all the data is organized, we can make some visualizations to answer some questions especially how changes in Canadian cases and deaths relate to the world as whole.

1. How many cases and deaths were there to date in Iceland?

Warning: Transformation introduced infinite values in continuous y-axis

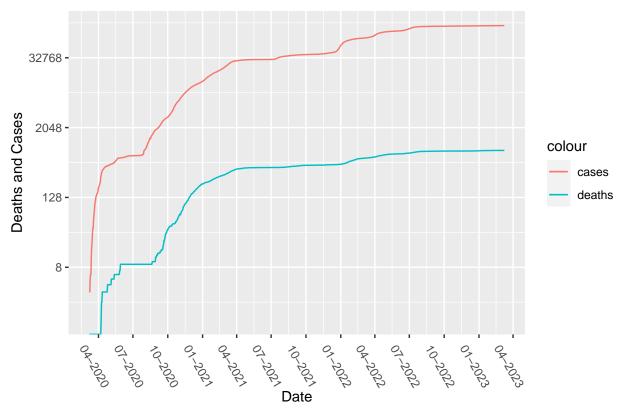
Iceland COVID Cases and Deaths



2. How many cases and deaths were there to date in Malta?

Warning: Transformation introduced infinite values in continuous y-axis

Malta COVID Cases and Deaths

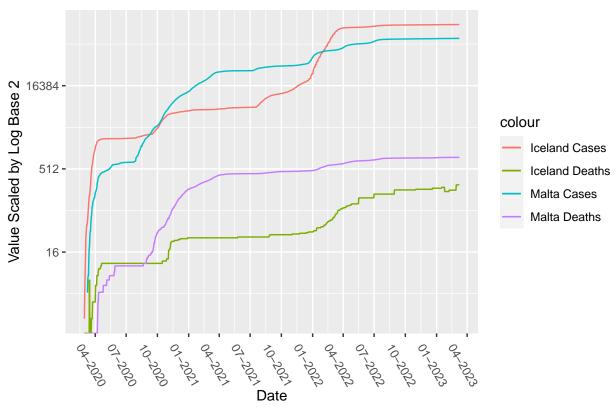


3. How does Iceland and Malta compare to each other?

```
ggplot() +
  geom_line(data = Iceland, aes(x = date, y = deaths, color = 'Iceland Deaths')) +
  geom_line(data = Iceland, aes(x = date, y = cases, color = 'Iceland Cases')) +
  geom_line(data = Malta, aes(x = date, y = deaths, color = 'Malta Deaths')) +
  geom_line(data = Malta, aes(x = date, y = cases, color = 'Malta Cases')) +
  scale_y_continuous(trans = 'log2') +
  labs(title = 'Iceland and Malta Cases and Deaths', x = 'Date', y = 'Value Scaled by Log Base 2') +
  scale_x_date(date_labels = '%m-%Y', date_breaks = '3 month') +
  theme(axis.text.x = element_text(angle = 300))
```

- ## Warning: Transformation introduced infinite values in continuous y-axis
- ## Transformation introduced infinite values in continuous y-axis





Modeling

Now that we have a basic picture of what cases look like over time in Iceland and Malta, let's dive into modeling for predicted death based on number of cases for both countries.

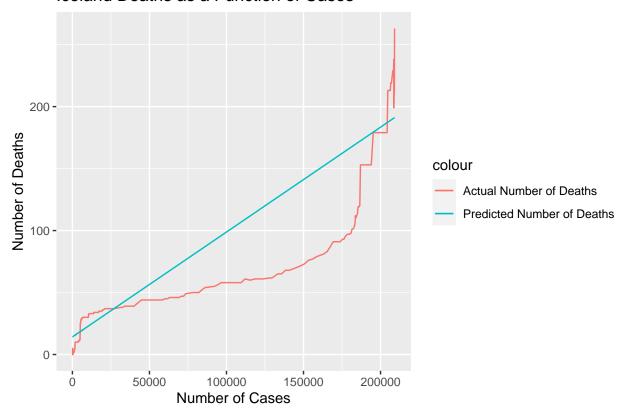
```
Iceland_death_model <- lm(deaths ~ cases, data = Iceland)
summary(Iceland_death_model)</pre>
```

```
##
## Call:
## lm(formula = deaths ~ cases, data = Iceland)
##
## Residuals:
##
                                3Q
       Min
                1Q
                    Median
                                       Max
                                   71.881
  -71.067
           -6.543
                     6.480
                             9.713
##
## Coefficients:
##
                Estimate Std. Error t value Pr(>|t|)
## (Intercept) 1.420e+01 9.288e-01
                                      15.29
                                              <2e-16 ***
               8.461e-04 7.953e-06
                                     106.39
##
  cases
                                              <2e-16 ***
##
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
##
```

```
## Residual standard error: 23.77 on 1103 degrees of freedom
## Multiple R-squared: 0.9112, Adjusted R-squared: 0.9111
## F-statistic: 1.132e+04 on 1 and 1103 DF, p-value: < 2.2e-16</pre>
```

```
Iceland <- Iceland %>%
  mutate(pred_deaths = predict(Iceland_death_model))
Iceland %>%
  ggplot() +
  geom_line(aes(x = cases, y = deaths, color = 'Actual Number of Deaths')) +
  geom_line(aes(x = cases, y = pred_deaths, color = 'Predicted Number of Deaths')) +
  labs(title = 'Iceland Deaths as a Function of Cases', x = 'Number of Cases', y = 'Number of Deaths')
```

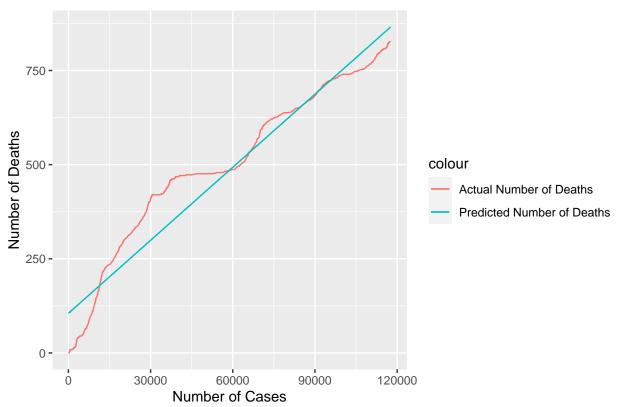
Iceland Deaths as a Function of Cases



```
Malta_death_model <- lm(deaths ~ cases, data = Malta)
summary(Malta_death_model)</pre>
```

```
##
                Estimate Std. Error t value Pr(>|t|)
  (Intercept) 1.052e+02 3.610e+00
                                      29.14
##
                                              <2e-16 ***
               6.465e-03
                         5.384e-05
                                    120.06
##
## Signif. codes:
                   0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
##
## Residual standard error: 77.58 on 1095 degrees of freedom
## Multiple R-squared: 0.9294, Adjusted R-squared: 0.9293
## F-statistic: 1.442e+04 on 1 and 1095 DF, p-value: < 2.2e-16
Malta <- Malta %>%
  mutate(pred_deaths = predict(Malta_death_model))
Malta %>%
  ggplot() +
  geom_line(aes(x = cases, y = deaths, color = 'Actual Number of Deaths')) +
  geom_line(aes(x = cases, y = pred_deaths, color = 'Predicted Number of Deaths')) +
  labs(title = 'Malta Deaths as a Function of Cases', x = 'Number of Cases', y = 'Number of Deaths')
```

Malta Deaths as a Function of Cases



Based on the Linear Regression Models, we can see that Malta's trend follows very closely with the predicted outcomes based on the number of cases present. Iceland on the other hand managed to reduce deaths falling under the predicted outcome.

Bias

The possible biases present in this data include data collection, how cases were tested and deemed positive, how local climate might impact transmission rates, how the healthcare systems differ in both countries in

how they manage patients with COVID 19. Responding to these biases may present a fuller picture as to how these two small European nations varied differently in how they handeld COVID 19 Deaths relative to the number of cases.