Covid

Ben Lewis

2025-01-05

```
library(stringr)
library(tidyverse)
## -- Attaching core tidyverse packages ----- tidyverse 2.0.0 --
## v dplyr 1.1.4 v purrr 1.0.2
## v forcats 1.0.0 v readr
                                   2.1.5
## v ggplot2 3.5.1
                      v tibble
                                   3.2.1
## v lubridate 1.9.4
                       v tidyr
                                   1.3.1
## -- Conflicts ------ tidyverse_conflicts() --
## x dplyr::filter() masks stats::filter()
## x dplyr::lag() masks stats::lag()
## i Use the conflicted package (<a href="http://conflicted.r-lib.org/">http://conflicted.r-lib.org/</a>) to force all conflicts to become error
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", "t
urls <- str_c(url_in, file_names)</pre>
global_cases <- read_csv(urls[1])</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.
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.

```
us_cases <- read_csv(urls[3])</pre>
## Rows: 3342 Columns: 1154
## -- Column specification -----
## Delimiter: ","
          (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.
us_deaths <- read_csv(urls[4])</pre>
## Rows: 3342 Columns: 1155
## -- Column specification -----
## Delimiter: ","
          (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.
global_cases <- global_cases %>% pivot_longer(cols = -c(`Province/State`, `Country/Region`, `Lat`, `Lon,
global_deaths <- global_deaths %>% pivot_longer(cols = -c(`Province/State`, `Country/Region`, `Lat`, `L
global <- global_cases %>% full_join(global_deaths) %>% rename(Country_Region = `Country/Region`, Provi
## Joining with 'by = join_by('Province/State', 'Country/Region', date)'
global <- global %>% filter(cases > 0)
us_cases <- us_cases %>% pivot_longer(cols = -(UID:Combined_Key),
                                     names_to = "date",
                                     values_to = "cases") %>%
  select(Admin2:cases) %>%
  mutate(date = mdy(date)) %>%
  select(-c(Lat, Long_))
us_deaths <- us_deaths %>%
  pivot_longer(cols = -(UID:Population),
               names_to = "date",
               values_to = "deaths") %>%
  select(Admin2:deaths) %>%
  mutate(date = mdy(date)) %>%
  select(-c(Lat, Long_))
us <- us_cases %>%
 full_join(us_deaths)
## Joining with 'by = join_by(Admin2, Province_State, Country_Region,
## Combined_Key, date) '
```

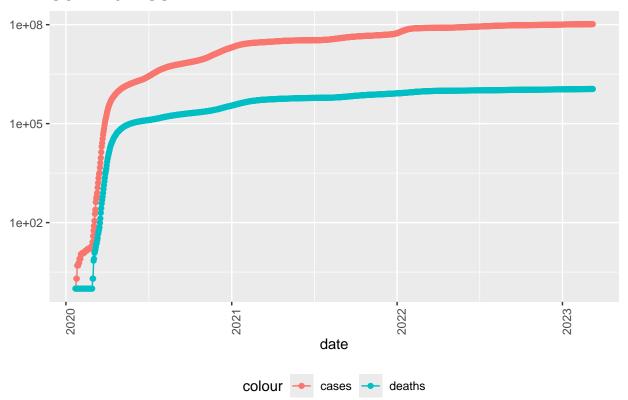
```
global <- global %>%
  unite("Combined_Key",
       c(Province_State, Country_Region),
        sep = ", ",
       na.rm = TRUE,
       remove = FALSE)
uid lookup url <- "https://raw.githubusercontent.com/CSSEGISandData/COVID-19/master/csse covid 19 data/
uid <- read_csv(uid_lookup_url) %>%
  select(-c(Lat, Long_, Combined_Key, code3, iso2, iso3, Admin2))
## Rows: 4321 Columns: 12
## -- Column specification -----
## Delimiter: ","
## chr (7): iso2, iso3, FIPS, Admin2, Province_State, Country_Region, Combined_Key
## dbl (5): UID, code3, Lat, Long_, Population
## 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 <- global %>%
 left_join(uid, by = c("Province_State", "Country_Region")) %>%
  select(-c(UID, FIPS)) %>%
  select(Province_State, Country_Region, date,
         cases, deaths, Population,
         Combined_Key)
us_by_state <- us %>%
  group_by(Province_State, Country_Region, date) %>%
  summarize(cases = sum(cases), deaths = sum(deaths),
            Population = sum(Population)) %>%
 mutate(deaths_per_mill = deaths *1000000 / Population) %>%
  select(Province_State, Country_Region, date,
         cases, deaths, deaths_per_mill, Population) %>%
  ungroup()
## 'summarise()' has grouped output by 'Province_State', 'Country_Region'. You can
## override using the '.groups' argument.
us_totals <- us_by_state %>%
  group_by(Country_Region, date) %>%
  summarize(cases = sum(cases), deaths = sum(deaths),
            Population = sum(Population)) %>%
  mutate(deaths_per_mill = deaths *1000000 / Population) %>%
  select(Country_Region, date,
         cases, deaths, deaths_per_mill, Population) %>%
  ungroup()
## 'summarise()' has grouped output by 'Country Region'. You can override using
```

the '.groups' argument.

tail(us_totals)

```
## # A tibble: 6 x 6
##
    Country_Region date
                                   cases deaths deaths_per_mill Population
     <chr>
                                                           <dbl>
##
                                   <dbl>
## 1 US
                    2023-03-04 103650837 1122172
                                                           3371. 332875137
## 2 US
                                                           3371.
                    2023-03-05 103646975 1122134
                                                                  332875137
## 3 US
                    2023-03-06 103655539 1122181
                                                           3371. 332875137
## 4 US
                   2023-03-07 103690910 1122516
                                                           3372. 332875137
## 5 US
                    2023-03-08 103755771 1123246
                                                           3374. 332875137
                                                           3376. 332875137
## 6 US
                    2023-03-09 103802702 1123836
```

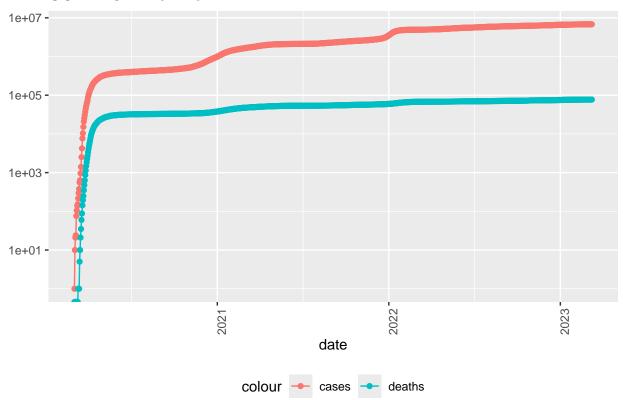
COVID19 in US



```
state <- "New York"
us_by_state %>%
filter(Province_State == state) %>%
filter(cases > 0) %>%
ggplot(aes(x = date, y = cases)) +
geom_line(aes(color = "cases")) +
geom_point(aes(color = "cases")) +
geom_line(aes(y = deaths, color = "deaths")) +
geom_point(aes(y = deaths, color = "deaths")) +
scale_y_log10() +
theme(legend.position = "bottom", axis.text.x = element_text(angle = 90)) +
labs(title = str_c("COVID19 in ", state), y = NULL)
```

Warning in scale_y_log10(): log-10 transformation introduced infinite values.
log-10 transformation introduced infinite values.

COVID19 in New York



```
max(us_totals$date)
```

[1] "2023-03-09"

max(us_totals\$deaths)

[1] 1123836

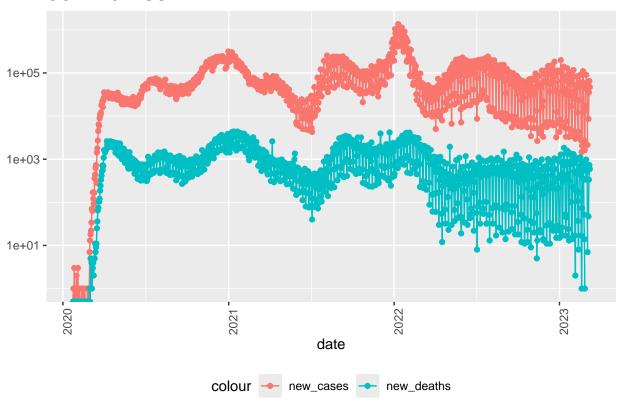
```
us_by_state <- us_by_state %>%
 mutate(new_cases = cases - lag(cases),
        new_deaths = deaths - lag(deaths))
us_totals <- us_totals %>%
 mutate(new_cases = cases - lag(cases),
        new_deaths = deaths - lag(deaths))
tail(us_totals %>% select(new_cases, new_deaths, everything()))
## # A tibble: 6 x 8
## new_cases new_deaths Country_Region date
                                                    cases deaths deaths_per_mill
                                      <date>
##
        <dbl>
                 <dbl> <chr>
                                                    <dbl> <dbl>
                                                                            <dbl>
                                      2023-03-04 1.04e8 1.12e6
                      7 US
                                                                            3371.
## 1
         2147
                                      2023-03-05 1.04e8 1.12e6
                     -38 US
## 2
        -3862
                                                                            3371.
                     47 US
                                      2023-03-06 1.04e8 1.12e6
## 3
        8564
                                                                           3371.
                                      2023-03-07 1.04e8 1.12e6
## 4
       35371
                    335 US
                                                                            3372.
                                       2023-03-08 1.04e8 1.12e6
## 5
       64861
                     730 US
                                                                            3374.
        46931
## 6
                     590 US
                                        2023-03-09 1.04e8 1.12e6
                                                                            3376.
## # i 1 more variable: Population <dbl>
us_totals %>%
 ggplot(aes(x = date, y = new_cases)) +
 geom_line(aes(color = "new_cases")) +
 geom_point(aes(color = "new_cases")) +
 geom_line(aes(y = new_deaths, color = "new_deaths")) +
 geom_point(aes(y = new_deaths, color = "new_deaths")) +
 scale_y_log10() +
 theme(legend.position = "bottom",
       axis.text.x = element_text(angle = 90)) +
 labs(title = "COVID19 in US", y = NULL)
## Warning in transformation$transform(x): NaNs produced
## Warning in transformation $\text{transform}(x): log-10 transformation introduced
## infinite values.
## Warning in transformation$transform(x): NaNs produced
## Warning in scale_y_log10(): log-10 transformation introduced infinite values.
## Warning in transformation$transform(x): NaNs produced
## Warning in scale_y_log10(): log-10 transformation introduced infinite values.
## Warning in transformation$transform(x): NaNs produced
## Warning in scale_y_log10(): log-10 transformation introduced infinite values.
## Warning: Removed 1 row containing missing values or values outside the scale range
## ('geom_line()').
```

Warning: Removed 2 rows containing missing values or values outside the scale range
('geom_point()').

Warning: Removed 1 row containing missing values or values outside the scale range
('geom_line()').

Warning: Removed 4 rows containing missing values or values outside the scale range
('geom_point()').

COVID19 in US



R Markdown

This is an R Markdown document. Markdown is a simple formatting syntax for authoring HTML, PDF, and MS Word documents. For more details on using R Markdown see http://rmarkdown.rstudio.com.

When you click the \mathbf{Knit} button a document will be generated that includes both content as well as the output of any embedded R code chunks within the document. You can embed an R code chunk like this:

summary(cars)

```
##
                         dist
        speed
                           : 2.00
##
          : 4.0
    1st Qu.:12.0
                   1st Qu.: 26.00
##
    Median:15.0
                   Median: 36.00
                           : 42.98
           :15.4
                   Mean
##
    Mean
##
    3rd Qu.:19.0
                   3rd Qu.: 56.00
##
    Max.
           :25.0
                   Max.
                           :120.00
```

Including Plots

You can also embed plots, for example:



Note that the echo = FALSE parameter was added to the code chunk to prevent printing of the R code that generated the plot.