

Texas Covid Data Project

Nathen Byford, Connor Bryson, Miguel Iglesias

11/11/2021

Part 1: Data manipulation

```
tb <- read_xlsx("./data/data.xlsx", skip = 2)

cases <- tb |>
  rename(county = "County Name") |>
  gather(-1, key = "date", value = "cases") |>
  mutate(date = str_extract(date, "[0-9]{2}-[0-9]{2}-[0-9]{4}"),
         date = mdy(date))

cases
```

```
## # A tibble: 160,420 x 3
##   county    date    cases
##   <chr>    <date>    <dbl>
## 1 Anderson 2020-03-04     0
## 2 Andrews 2020-03-04     0
## 3 Angelina 2020-03-04     0
## 4 Aransas 2020-03-04     0
## 5 Archer   2020-03-04     0
## 6 Armstrong 2020-03-04     0
## 7 Atascosa 2020-03-04     0
## 8 Austin   2020-03-04     0
## 9 Bailey   2020-03-04     0
## 10 Bandera 2020-03-04     0
## # ... with 160,410 more rows
```

Part 2: Data merging

```
pop_dat <- read_csv("./data/county-populations.csv")
```

```
## Rows: 255 Columns: 2
```

```
## -- Column specification -----
## Delimiter: ","
## chr (1): county
## dbl (1): 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.
```

```
cases <- left_join(cases, pop_dat)
```

```
## Joining, by = "county"
```

```
cases
```

```
## # A tibble: 160,420 x 4
##   county   date      cases population
##   <chr>   <date>   <dbl>      <dbl>
## 1 Anderson 2020-03-04     0      58199
## 2 Andrews  2020-03-04     0      22269
## 3 Angelina 2020-03-04     0      90437
## 4 Aransas  2020-03-04     0      27699
## 5 Archer   2020-03-04     0       8344
## 6 Armstrong 2020-03-04     0       1948
## 7 Atascosa 2020-03-04     0      51831
## 8 Austin   2020-03-04     0      30402
## 9 Bailey   2020-03-04     0       7692
## 10 Bandera 2020-03-04     0      21246
## # ... with 160,410 more rows
```

part 3: Data visualization

```
# Find top 16 counties, and total.
top_count <- tail(pop_dat[order(pop_dat$population), ], n = 17) |> pull(county)

cases |> filter(county %in% top_count, county != "Total") |>
  group_by(county) |>
  mutate(new_cases = c(cases[1], diff(cases)),
         new_cases = slide_dbl(new_cases, mean, .before = 6)) |>
  ggplot(aes(x = date, y = new_cases, group = county)) +
  geom_line(aes(color = county), alpha = .6) +
  scale_y_continuous(labels = scales::comma, limits = c(0, NA)) +
  scale_color_viridis_d() +
  labs(title = "Incident Cases",
       subtitle = "For 16 Largest Counties in Texas",
       y = "Cases",
       x = "Date")
```

