

tl_covid_exercise

Katherine Liu

12/13/2020

```
library(data.table)
```

```
## Warning: package 'data.table' was built under R version 4.0.3
```

```
library(tidyverse)
```

```
## Warning: package 'tidyverse' was built under R version 4.0.3
```

```
## -- Attaching packages ----- tidyverse 1.3.0 --
```

```
## v ggplot2 3.3.2      v purrr  0.3.4
## v tibble  3.0.4      v dplyr  1.0.2
## v tidyr   1.1.2      v stringr 1.4.0
## v readr   1.4.0      v forcats 0.5.0
```

```
## Warning: package 'ggplot2' was built under R version 4.0.3
```

```
## Warning: package 'tibble' was built under R version 4.0.3
```

```
## Warning: package 'tidyr' was built under R version 4.0.3
```

```
## Warning: package 'readr' was built under R version 4.0.3
```

```
## Warning: package 'purrr' was built under R version 4.0.3
```

```
## Warning: package 'dplyr' was built under R version 4.0.3
```

```
## Warning: package 'forcats' was built under R version 4.0.3
```

```
## -- Conflicts ----- tidyverse_conflicts() --
```

```
## x dplyr::between() masks data.table::between()
## x dplyr::filter()  masks stats::filter()
## x dplyr::first()   masks data.table::first()
## x dplyr::lag()      masks stats::lag()
## x dplyr::last()     masks data.table::last()
## x purrr::transpose() masks data.table::transpose()
```

```

covid_data <-
  list.files(pattern = "*.csv") %>%
  map_df(~fread(.))

remove = c("Diamond Princess", "District of Columbia", "Grand Princess", "Guam",
           "Puerto Rico", "American Samoa", "Northern Mariana Islands",
           "Recovered", "Virgin Islands")
state_data = covid_data[!covid_data$Province_State %in% remove]
state_data = select(state_data, -c(Country_Region))

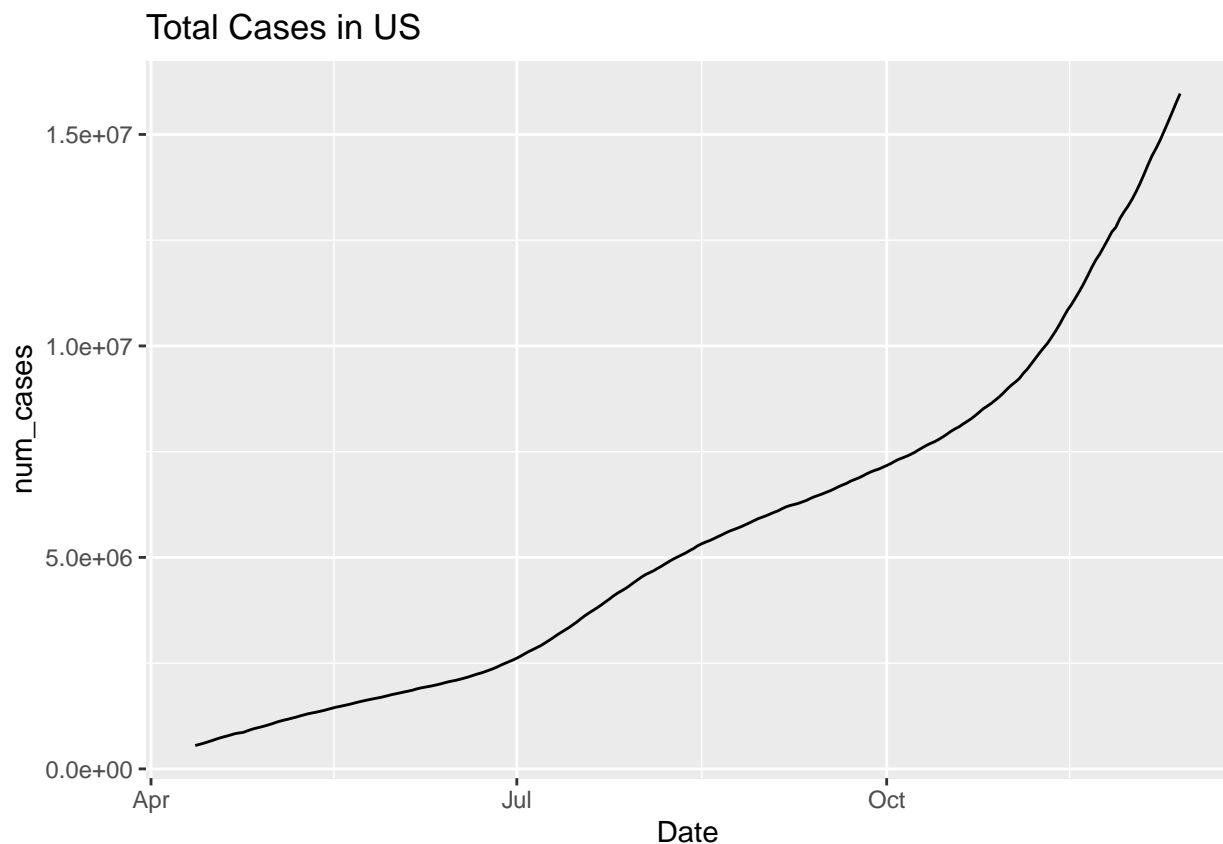
colnames(state_data)[1] = "State"
colnames(state_data)[2] = "Date"
colnames(state_data)[4] = "Long"

state_data$Date = as.Date(state_data$Date, "%Y-%m-%d %H:%M:%S", tz = "GMT")
state_data[59,2] = state_data[58,2]

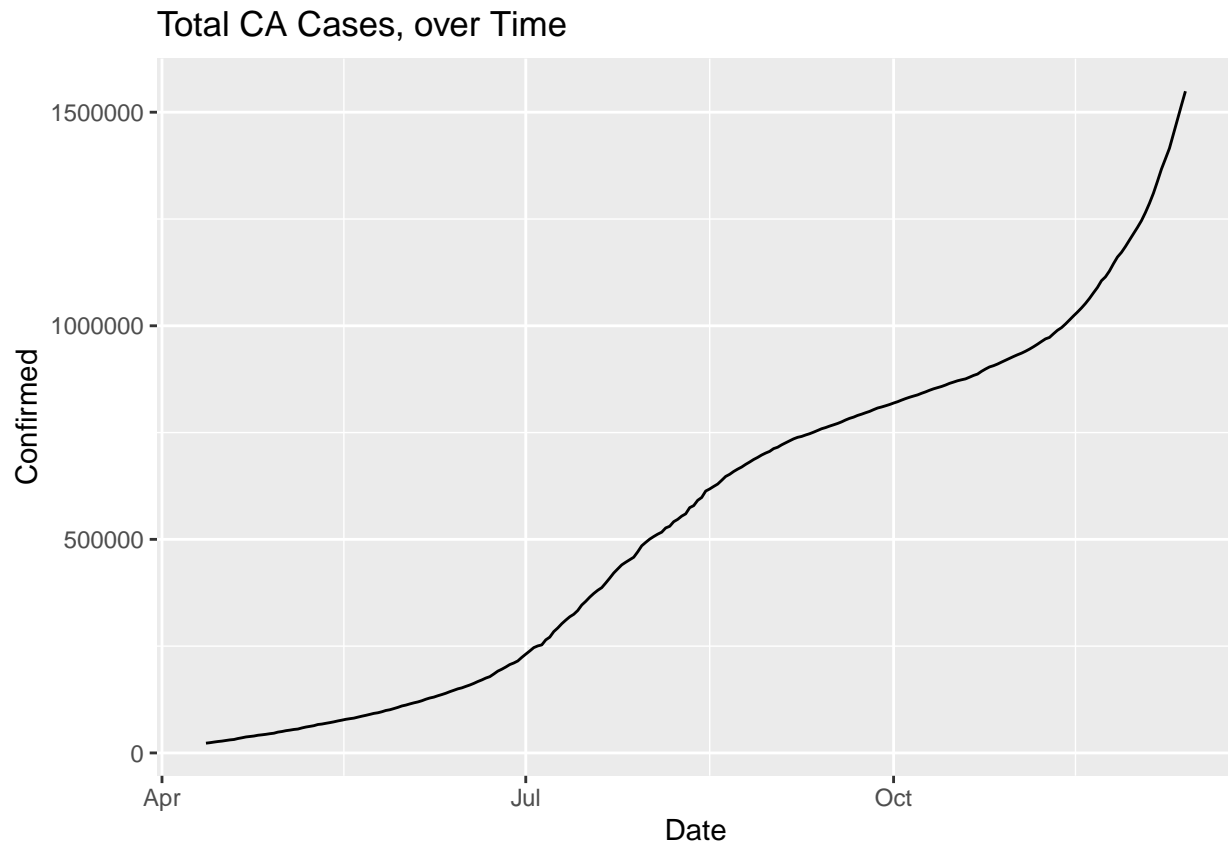
case_totals = state_data %>%
  group_by(Date) %>%
  mutate(num_cases = sum(Confirmed))%>%
  distinct(Date, num_cases)

ggplot(data = case_totals) +
  geom_line(mapping = aes(x = Date, y = num_cases)) +
  ggtitle("Total Cases in US")

```



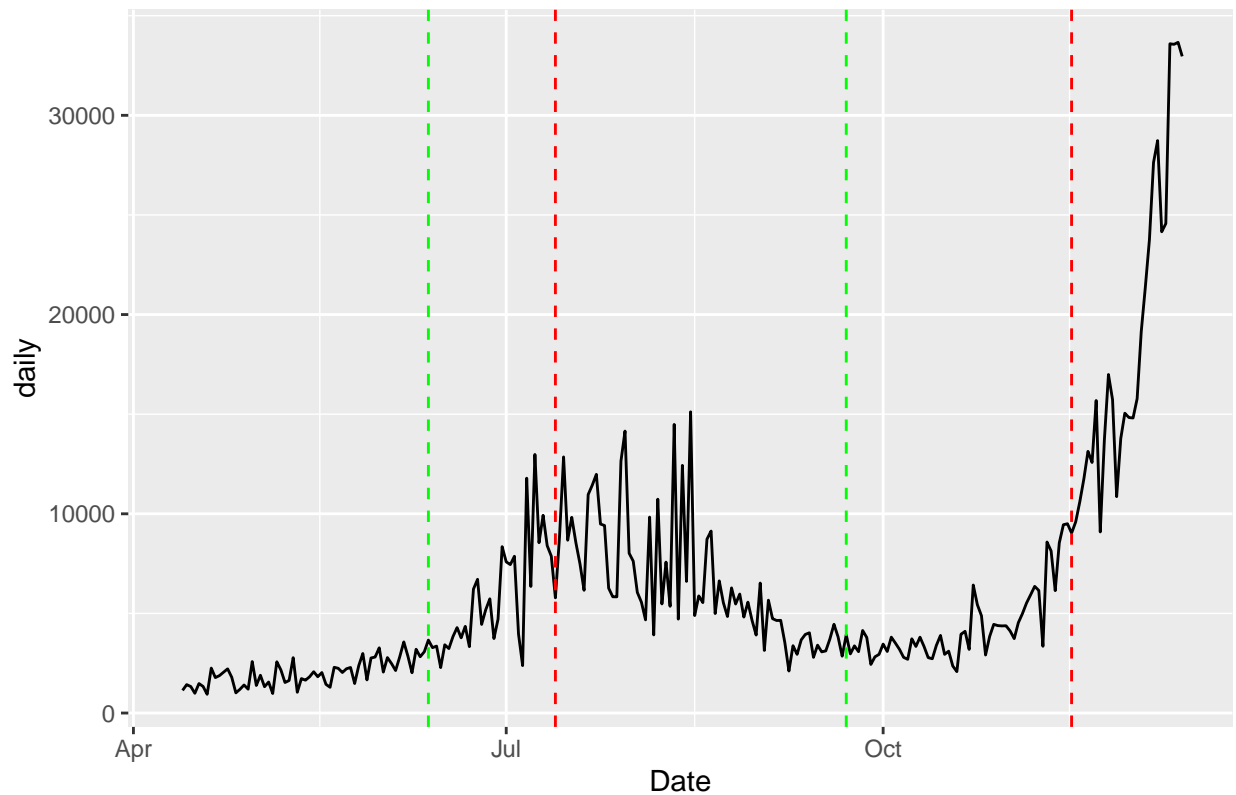
```
state_data %>%
  filter(State == "California") %>%
  ggplot(mapping = aes(x = Date, y = Confirmed)) +
  geom_line() +
  ggtitle("Total CA Cases, over Time")
```



```
state_data %>%
  filter(State == "California") %>%
  mutate(daily = Confirmed - lag(Confirmed)) %>%
  ggplot(mapping = aes(x = Date, y = daily)) +
  geom_line() +
  geom_vline(xintercept = as.numeric(as.Date("2020-06-12")), linetype=2, col = "green") +
  geom_vline(xintercept = as.numeric(as.Date("2020-07-13")), linetype=2, col = "red") +
  geom_vline(xintercept = as.numeric(as.Date("2020-11-16")), linetype=2, col = "red") +
  geom_vline(xintercept = as.numeric(as.Date("2020-09-22")), linetype=2, col = "green") +
  ggtitle("Daily Cases in CA")
```

```
## Warning: Removed 1 row(s) containing missing values (geom_path).
```

Daily Cases in CA



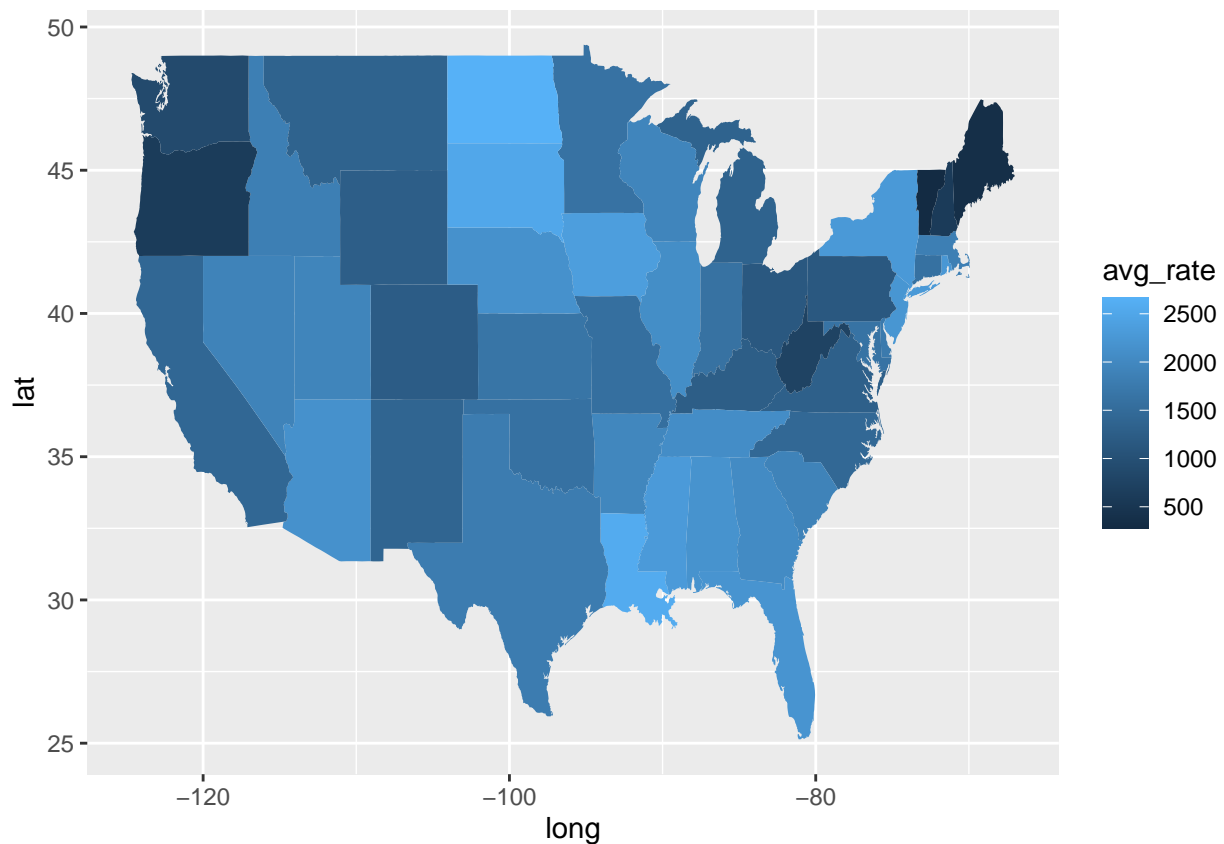
```
(state_avg = state_data %>%
  group_by(State) %>%
  mutate(avg_rate = mean(Incident_Rate)) %>%
  distinct(State, avg_rate) %>%
  arrange(avg_rate)
)
```

```
## # A tibble: 50 x 2
## # Groups:   State [50]
##   State      avg_rate
##   <chr>      <dbl>
## 1 Vermont      281.
## 2 Maine         355.
## 3 Hawaii        500.
## 4 New Hampshire  601.
## 5 Oregon        614.
## 6 West Virginia  745.
## 7 Washington    892.
## 8 Alaska       1079.
## 9 Ohio         1146.
## 10 Pennsylvania 1155.
## # ... with 40 more rows
```

```
state_avg$State = tolower(state_avg$State)
state_df = map_data("state")
```

```
colnames(state_df)[5] = "State"
```

```
merge_state <- inner_join(state_df, state_avg, by = "State")
ggplot() +
  geom_polygon( data=merge_state,
    aes(x=long, y=lat, group=group, fill = avg_rate))
```



```
ci = state_data %>%
  group_by(State) %>%
  dplyr::summarize(mean_rate = mean(Incident_Rate, na.rm = TRUE),
    sd_rate = sd(Incident_Rate, na.rm = TRUE),
    n_rate = n()) %>%
  mutate(se_rate = sd_rate / sqrt(n_rate),
    lower_ci_rate = mean_rate - qt(1 - (0.05 / 1225), n_rate - 1) * se_rate,
    upper_ci_rate = mean_rate + qt(1 - (0.05 / 1225), n_rate - 1) * se_rate)
```

```
## 'summarise()' ungrouping output (override with '.groups' argument)
```

```
ggplot(ci, mapping = aes(x = reorder(State, mean_rate), y = mean_rate)) +
  geom_point() +
  geom_errorbar(data = ci, aes(ymin = mean_rate - 2*sd_rate, ymax = mean_rate + 2 * sd_rate, color = St
  theme(legend.position = "none") +
  coord_flip()
```

