

# Week 11 Challenge

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2023-11-21

```
library(httr)
library(jsonlite)
library(tidyverse)
```

```
## -- Attaching core tidyverse packages ----- tidyverse 2.0.0 --
## v dplyr      1.1.2      v readr      2.1.4
## v forcats    1.0.0      v stringr   1.5.0
## v ggplot2    3.4.4      v tibble    3.2.1
## v lubridate  1.9.2      v tidyr     1.3.0
## v purrr      1.0.2
```

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

```
historic_state_data_url <- "https://api.covidactnow.org/v2/states.timeseries.json?apiKey=0204bfb158084b"
raw_data <- GET(historic_state_data_url)
```

```
data <- fromJSON(rawToChar(raw_data$content))
glimpse(data)
```

```
## Rows: 53
## Columns: 25
## $ fips      <chr> "02", "01", "05", "04", "06", "08", "09~
## $ country   <chr> "US", "US", "US", "US", "US", "US", "US~
## $ state     <chr> "AK", "AL", "AR", "AZ", "CA", "CO", "CT~
## $ county    <lgl> NA, NA, NA, NA, NA, NA, NA, NA, NA, NA,~
## $ hsa       <lgl> NA, NA, NA, NA, NA, NA, NA, NA, NA, NA,~
## $ hsaName    <lgl> NA, NA, NA, NA, NA, NA, NA, NA, NA, NA,~
## $ level     <chr> "state", "state", "state", "state", "st~
## $ lat       <lgl> NA, NA, NA, NA, NA, NA, NA, NA, NA, NA,~
## $ locationId <chr> "iso1:us#iso2:us-ak", "iso1:us#iso2:us~
## $ long      <lgl> NA, NA, NA, NA, NA, NA, NA, NA, NA, NA,~
## $ population <int> 731545, 4903185, 3017804, 7278717, 3951~
## $ hsaPopulation <int> NA, NA, NA, NA, NA, NA, NA, NA, NA, NA,~
## $ metrics    <df[,14]> <data.frame[26 x 14]>
## $ riskLevels <df[,6]> <data.frame[26 x 6]>
## $ cdcTransmissionLevel <int> 2, 4, 3, 3, 1, 4, 4, 1, 4, 4, 2, 2,~
## $ communityLevels <df[,2]> <data.frame[26 x 2]>
## $ actuals     <df[,19]> <data.frame[26 x 19]>
```

```
## $ annotations <df[,30]> <data.frame[26 x 30]>
## $ lastUpdatedDate <chr> "2023-11-18", "2023-11-18", "2023-11-~
## $ url <chr> "https://covidactnow.org/us/alaska-ak", ~
## $ metricsTimeseries <list> [<data.frame[1351 x 14]>], [<data.fr~
## $ actualsTimeseries <list> [<data.frame[1351 x 20]>], [<data.f~
## $ riskLevelsTimeseries <list> [<data.frame[1351 x 3]>], [<data.fr~
## $ cdcTransmissionLevelTimeseries <list> [<data.frame[1351 x 2]>], [<data.frame[~
## $ communityLevelsTimeseries <list> [<data.frame[1351 x 3]>], [<data.frame[~
```

```
time_series <- data %>%
  unnest(actualsTimeseries)
```

```
# Creating a new dataframe with needed data
# Save date
```

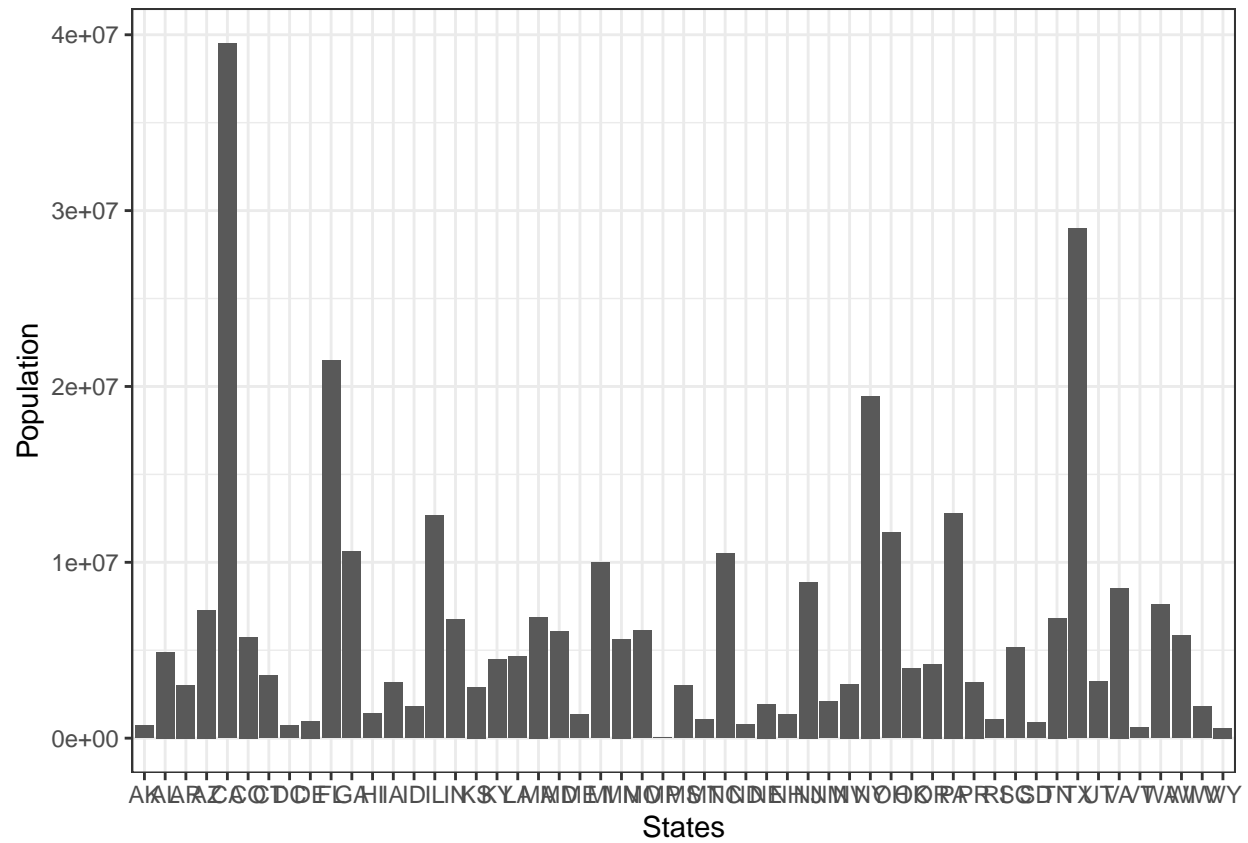
```
time_series_transmission <- tibble(Date=time_series$cdcTransmissionLevelTimeseries[[which(data$state=="C~
# Transmission levels in each state
time_series_transmission$Alaska <- time_series$cdcTransmissionLevelTimeseries[[which(data$state=="AK")]]
cdcTransmissionLevel
time_series_transmission$California <- time_series$cdcTransmissionLevelTimeseries[[which(data$state=="C~
time_series_transmission$New_Jersey <- time_series$cdcTransmissionLevelTimeseries[[which(data$state=="N~
time_series_transmission$Tennessee <- time_series$cdcTransmissionLevelTimeseries[[which(data$state=="TN~
time_series_transmission$District_of_Columbia <- time_series$cdcTransmissionLevelTimeseries[[which(data~
print(head(time_series_transmission))
```

```
## # A tibble: 6 x 6
##   Date      Alaska California New_Jersey Tennessee District_of_Columbia
##   <chr>      <int>      <int>      <int>      <int>      <int>
## 1 2020-03-01      0          0          0          0          0
## 2 2020-03-02      0          0          0          0          0
## 3 2020-03-03      0          0          0          0          0
## 4 2020-03-04      0          0          0          0          0
## 5 2020-03-05      0          0          0          0          0
## 6 2020-03-06      0          0          0          0          0
```

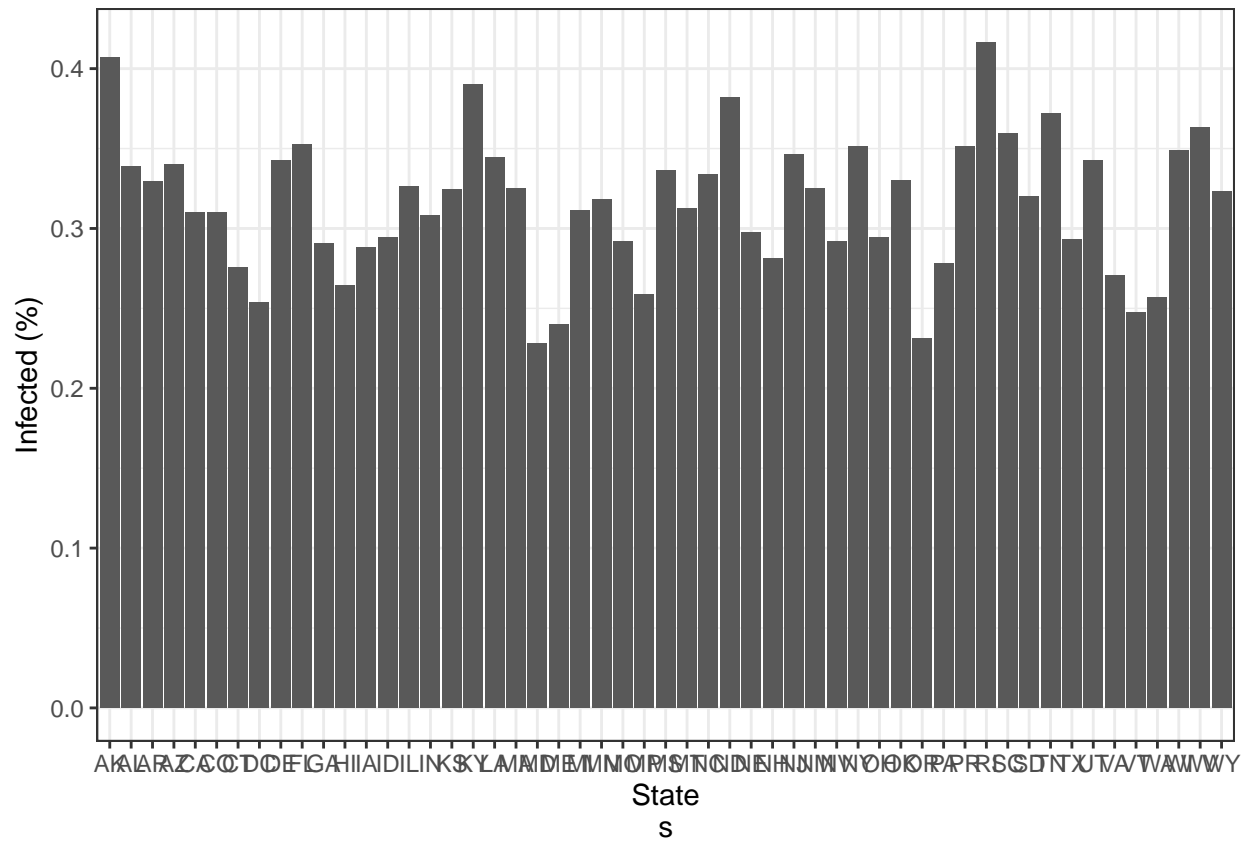
```
# New data-frame with dates
```

```
time_series_cases <- list(Alaska = time_series %>% filter(state=="AK") %>% select(date,cases))
# Cases of each state
time_series_cases$California <- time_series %>% filter(state=="CA") %>% select(date,cases)
time_series_cases$New_Jersey <- time_series %>% filter(state=="NJ") %>% select(date,cases)
time_series_cases$Tennessee <- time_series %>% filter(state=="TN") %>% select(date,cases)
time_series_cases$District_of_Columbia <- time_series %>% filter(state=="DC") %>% select(date,cases)
```

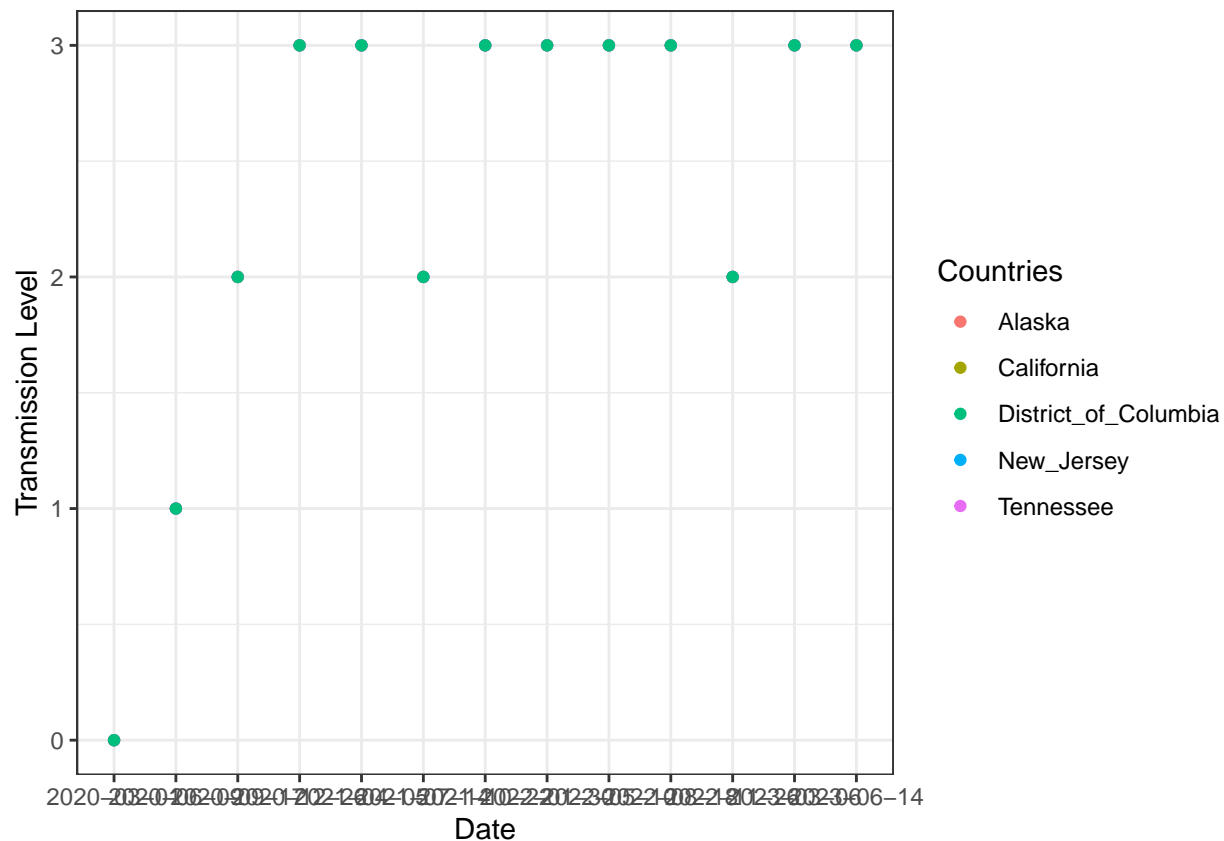
```
ggplot(data, aes(x=state,y=population)) + geom_bar(stat="identity") +labs(x="States",y="Population") +
```



```
ggplot(data, aes(x=state,y=(data$actuals$cases/population))) + geom_bar(stat="identity") + labs(x="States",y="Infected (%)") + theme_bw()
```



```
time_series_transmission[seq(1,1300,by=100),]%>%
pivot_longer(cols=Alaska:District_of_Columbia,names_to="Countries",values_to="Transmission") %>%
ggplot(aes(x=Date,y=Transmission,colour=Countries,group=Countries)) +
geom_point(show.legend=TRUE) + labs(x="Date",y="Transmission Level")+theme_bw()
```



```
data_to_plot <- tibble(Date_Alaska = time_series_cases$Alaska$date[seq(1,1300,by=100)],
Cases_Alaska = time_series_cases$Alaska$cases[seq(1,1300,by=100)],
Date_California = time_series_cases$California$date[seq(1,1300,by=100)],
Cases_California = time_series_cases$California$cases[seq(1,1300,by=100)],
Date_New_Jersey = time_series_cases$New_Jersey$date[seq(1,1300,by=100)],
Cases_New_Jersey = time_series_cases$New_Jersey$cases[seq(1,1300,by=100)],
Date_Tennessee = time_series_cases$Tennessee$date[seq(1,1300,by=100)],
Cases_Tennessee = time_series_cases$Tennessee$cases[seq(1,1300,by=100)], Date_District_of_Columbia = time_series_cases$District_of_Columbia$date[seq(1,1300,by=100)],
Cases_District_of_Columbia = time_series_cases$District_of_Columbia$cases[seq(1,1300,by=100)])
data_to_plot
```

```
## # A tibble: 13 x 10
##   Date_Alaska Cases_Alaska Date_California Cases_California Date_New_Jersey
##   <chr>          <int> <chr>          <int> <chr>
## 1 2020-03-01      NA 2020-01-25          1 2020-03-01
## 2 2020-06-09      620 2020-05-04       56333 2020-06-09
## 3 2020-09-17     7413 2020-08-12     595097 2020-09-17
## 4 2020-12-26    45247 2020-11-20    1096427 2020-12-26
## 5 2021-04-05    63486 2021-02-28    3569578 2021-04-05
## 6 2021-07-14    71539 2021-06-08    3798225 2021-07-14
## 7 2021-10-22   132393 2021-09-16    4629146 2021-10-22
## 8 2022-01-30   211117 2021-12-25    5291605 2022-01-30
## 9 2022-05-10   252847 2022-04-04    9110544 2022-05-10
## 10 2022-08-18   289203 2022-07-13   10365785 2022-08-18
## 11 2022-11-26   299841 2022-10-21   11338846 2022-11-26
## 12 2023-03-06   307377 2023-01-29   11980312 2023-03-06
```

```
## 13 2023-06-14          NA 2023-05-09          12242634 2023-06-14
## # i 5 more variables: Cases_New_Jersey <int>, Date_Tennessee <chr>,
## #   Cases_Tennessee <int>, Date_District_of_Columbia <chr>,
## #   Cases_District_of_Columbia <int>
```

```
library(cowplot)
```

```
##
```

```
## Attaching package: 'cowplot'
```

```
## The following object is masked from 'package:lubridate':
```

```
##
```

```
##      stamp
```

```
fig1<- ggplot(data_to_plot, aes(x=Date_Alaska,y=Cases_Alaska)) +
geom_point() + labs(x="Date",y="Cases", title="Alaska") + theme_bw()
fig2<- ggplot(data_to_plot, aes(x=Date_California,y=Cases_California)) +
geom_point() + labs(x="Date",y="Cases", title="California") + theme_bw()
fig3<- ggplot(data_to_plot, aes(x=Date_New_Jersey,y=Cases_New_Jersey)) +
geom_point() + labs(x="Date",y="Cases", title="New Jersey") + theme_bw()
fig4<- ggplot(data_to_plot, aes(x=Date_Tennessee,y=Cases_Tennessee)) +
geom_point() + labs(x="Date",y="Cases", title="Tennessee") + theme_bw()
fig5<- ggplot(data_to_plot, aes(x=Date_District_of_Columbia,y=Cases_District_of_Columbia)) +
geom_point() + labs(x="Date",y="Cases", title="District of Columbia") + theme_bw()
```

```
plot_grid(fig1 + theme(legend.justification = c(0,1)),
          fig2 + theme(legend.justification = c(1,0)),
          fig3 + theme(legend.justification = c(0,1)),
          fig4 + theme(legend.justification = c(1,0)),
          fig5 + theme(legend.justification = c(0,1)),
          align = "v", axis = "lr", nrow=3,
          ncol = 2,labels = LETTERS[1:5],
          rel_heights = c(1,2))
```

```
## Warning: Removed 2 rows containing missing values ('geom_point()').
```

```
## Removed 2 rows containing missing values ('geom_point()').
```

```
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```

```
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```

