

Challenge-11

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2023-10-30

Step 1: Procuring Data

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

```
## -- Attaching core tidyverse packages ----- tidyverse 2.0.0 --
## v dplyr      1.1.3      v readr      2.1.4
## v forcats    1.0.0      v stringr   1.5.0
## v ggplot2    3.4.3      v tibble    3.2.1
## v lubridate  1.9.3      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=9b75214f4f3041"
raw_data <- GET(historic_state_data_url)
```

Extracting data

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

Exploring the data

```
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 <lg1> NA, NA, NA, NA, NA, NA, NA, NA, NA, NA, NA,~
## $ hsaName <lg1> NA, NA, NA, NA, NA, NA, NA, NA, NA, NA, NA,~
## $ level <chr> "state", "state", "state", "state", "st~
## $ lat <lg1> NA, NA, NA, NA, NA, NA, NA, NA, NA, NA, NA,~
## $ locationId <chr> "iso1:us#iso2:us-ak", "iso1:us#iso2:us--
## $ long <lg1> NA, 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, 3,~
## $ 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> "2023-10-30", "2023-10-30", "2023-10~
## $ url <chr> "https://covidactnow.org/us/alaska-ak",~
## $ metricsTimeseries <list> [<data.frame[1334 x 14]>], [<data.fr~
## $ actualsTimeseries <list> [<data.frame[1334 x 20]>], [<data.f~
## $ riskLevelsTimeseries <list> [<data.frame[1334 x 3]>], [<data.fr~
## $ cdcTransmissionLevelTimeseries <list> [<data.frame[1334 x 2]>], [<data.frame[~
## $ communityLevelsTimeseries <list> [<data.frame[1334 x 3]>], [<data.frame[~
```

Mapping Variables to Questions

```
data$state
```

```
## [1] "AK" "AL" "AR" "AZ" "CA" "CO" "CT" "DC" "DE" "FL" "GA" "HI" "IA" "ID" "IL"
## [16] "IN" "KS" "KY" "LA" "MA" "MD" "ME" "MI" "MN" "MO" "MP" "MS" "MT" "NC" "ND"
## [31] "NE" "NH" "NJ" "NM" "NV" "NY" "OH" "OK" "OR" "PA" "PR" "RI" "SC" "SD" "TN"
## [46] "TX" "UT" "VA" "VT" "WA" "WI" "WV" "WY"
```

```
data$population
```

```
## [1] 731545 4903185 3017804 7278717 39512223 5758736 3565287 705749
## [9] 973764 21477737 10617423 1415872 3155070 1787065 12671821 6732219
## [17] 2913314 4467673 4648794 6892503 6045680 1344212 9986857 5639632
## [25] 6137428 53605 2976149 1068778 10488084 762062 1934408 1359711
## [33] 8882190 2096829 3080156 19453561 11689100 3956971 4217737 12801989
## [41] 3193694 1059361 5148714 884659 6829174 28995881 3205958 8535519
## [49] 623989 7614893 5822434 1792147 578759
```

```
data$actuals$cases
```

```
## [1] 297588 1659936 995043 2474154 12251820 1783663 982335 178904
## [9] 333772 7572282 3087729 374264 908936 526118 4136659 2076326
## [17] 945923 1743117 1600610 2242176 1379385 322982 3106362 1795771
## [25] 1790763 13886 1000415 333959 3501404 291093 575134 382013
## [33] 3075271 681242 898511 6838769 3441458 1306350 975856 3559331
## [41] 1122076 441466 1852019 282895 2542163 8508204 1097475 2310846
## [49] 154243 1957759 2030717 650556 187034
```

```
data$state
```

```
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## [46] "TX" "UT" "VA" "VT" "WA" "WI" "WV" "WY"
```

```
data$actuals$cases - data$actuals$deaths
```

```
## [1] 296120 1638798 981928 2440703 12149934 1769278 969998 177469
## [9] 330378 7484034 3044947 372408 898139 520639 4094654 2049880
## [17] 935709 1724494 1581639 2217532 1362645 319944 3063489 1781001
## [25] 1767832 13842 986941 330244 3472345 288580 570045 378961
## [33] 3039153 672006 886503 6761247 3399251 1290193 966306 3508284
## [41] 1116168 437551 1832096 279664 2512614 8414424 1092113 2287103
## [49] 153276 1941573 2014117 642431 185003
```

```
data$state
```

```
## [1] "AK" "AL" "AR" "AZ" "CA" "CO" "CT" "DC" "DE" "FL" "GA" "HI" "IA" "ID" "IL"
## [16] "IN" "KS" "KY" "LA" "MA" "MD" "ME" "MI" "MN" "MO" "MP" "MS" "MT" "NC" "ND"
## [31] "NE" "NH" "NJ" "NM" "NV" "NY" "OH" "OK" "OR" "PA" "PR" "RI" "SC" "SD" "TN"
## [46] "TX" "UT" "VA" "VT" "WA" "WI" "WV" "WY"
```

```
data$actuals$vaccinationsCompleted
```

```
## [1] 477592 2611593 1720209 4821350 29588939 4248431 2967081 NA
## [9] 715787 14971549 6103647 NA 2036465 1012257 9054864 3896474
## [17] 1909898 2669260 2561641 5825021 4835390 1124112 6254485 4082263
## [25] 3634453 44044 1600364 633531 6659428 448757 1286860 NA
## [33] 7043583 1582545 1966594 15763340 7089473 2399774 3065513 9444480
## [41] 2688410 932541 3091956 588257 3852924 18406327 2146291 6556391
## [49] 537193 5814062 3981606 1071540 307619
```

```
data$state
```

```
## [1] "AK" "AL" "AR" "AZ" "CA" "CO" "CT" "DC" "DE" "FL" "GA" "HI" "IA" "ID" "IL"
## [16] "IN" "KS" "KY" "LA" "MA" "MD" "ME" "MI" "MN" "MO" "MP" "MS" "MT" "NC" "ND"
## [31] "NE" "NH" "NJ" "NM" "NV" "NY" "OH" "OK" "OR" "PA" "PR" "RI" "SC" "SD" "TN"
## [46] "TX" "UT" "VA" "VT" "WA" "WI" "WV" "WY"
```

```
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=="
```

```
# 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$state=="CA")]]$cdcTransmissionLevel

time_series_transmission$New_Jersey <-
  time_series$cdcTransmissionLevelTimeseries[[which(data$state=="NJ")]]$cdcTransmissionLevel

time_series_transmission$Tennessee <-
  time_series$cdcTransmissionLevelTimeseries[[which(data$state=="TN")]]$cdcTransmissionLevel

time_series_transmission$District_of_Columbia <-
  time_series$cdcTransmissionLevelTimeseries[[which(data$state=="DC")]]$cdcTransmissionLevel

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)

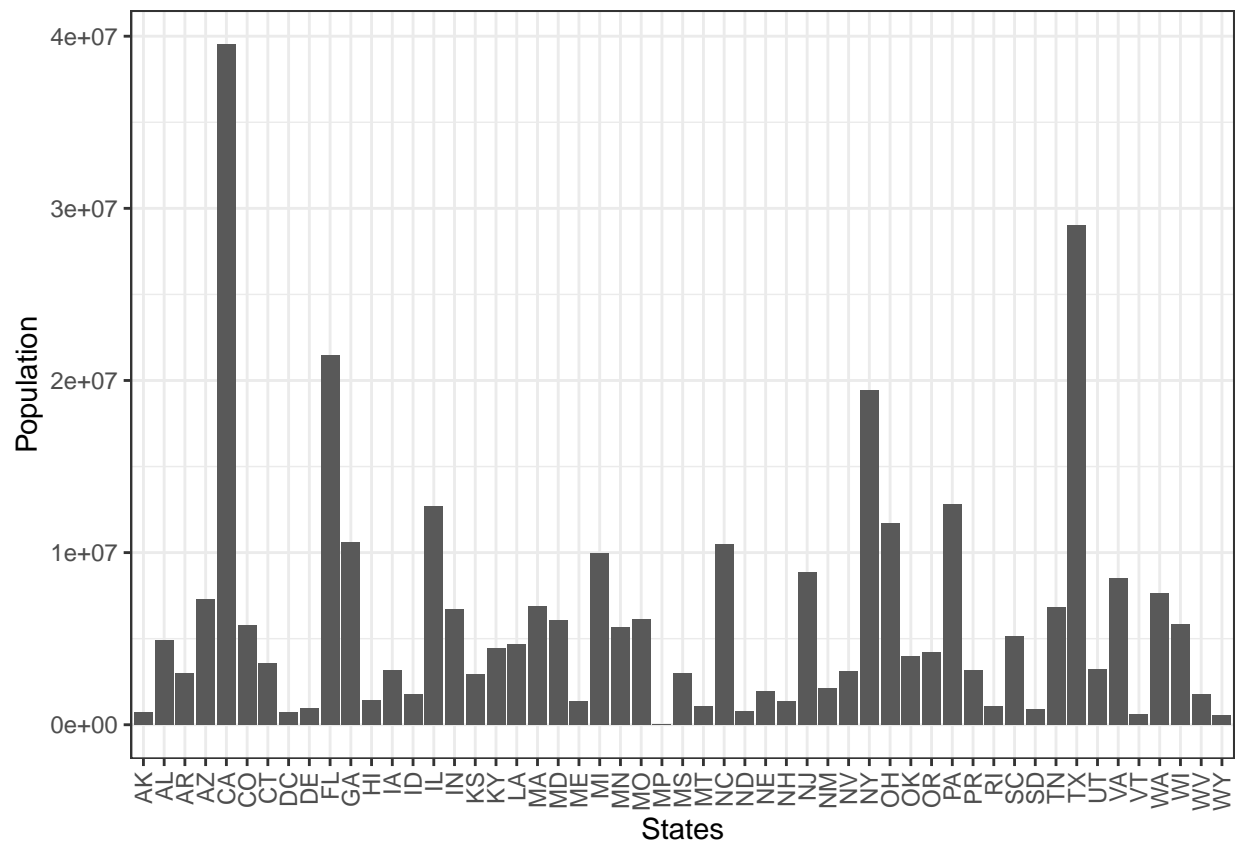
```

Analysing Data

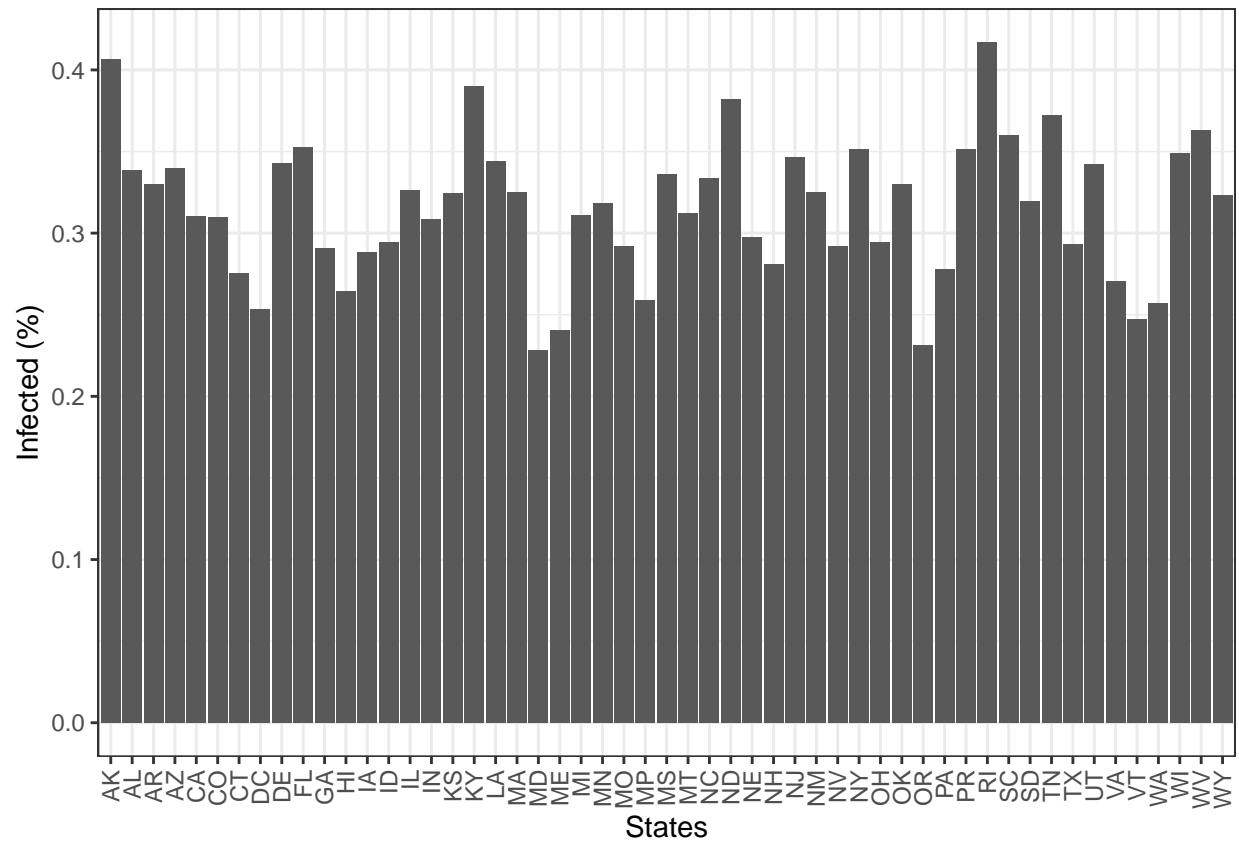
```

ggplot(data, aes(x=state,y=population)) +
  geom_bar(stat="identity") +
  labs(x="States",y="Population") +
  theme_bw() + scale_x_discrete(guide = guide_axis(angle=90))

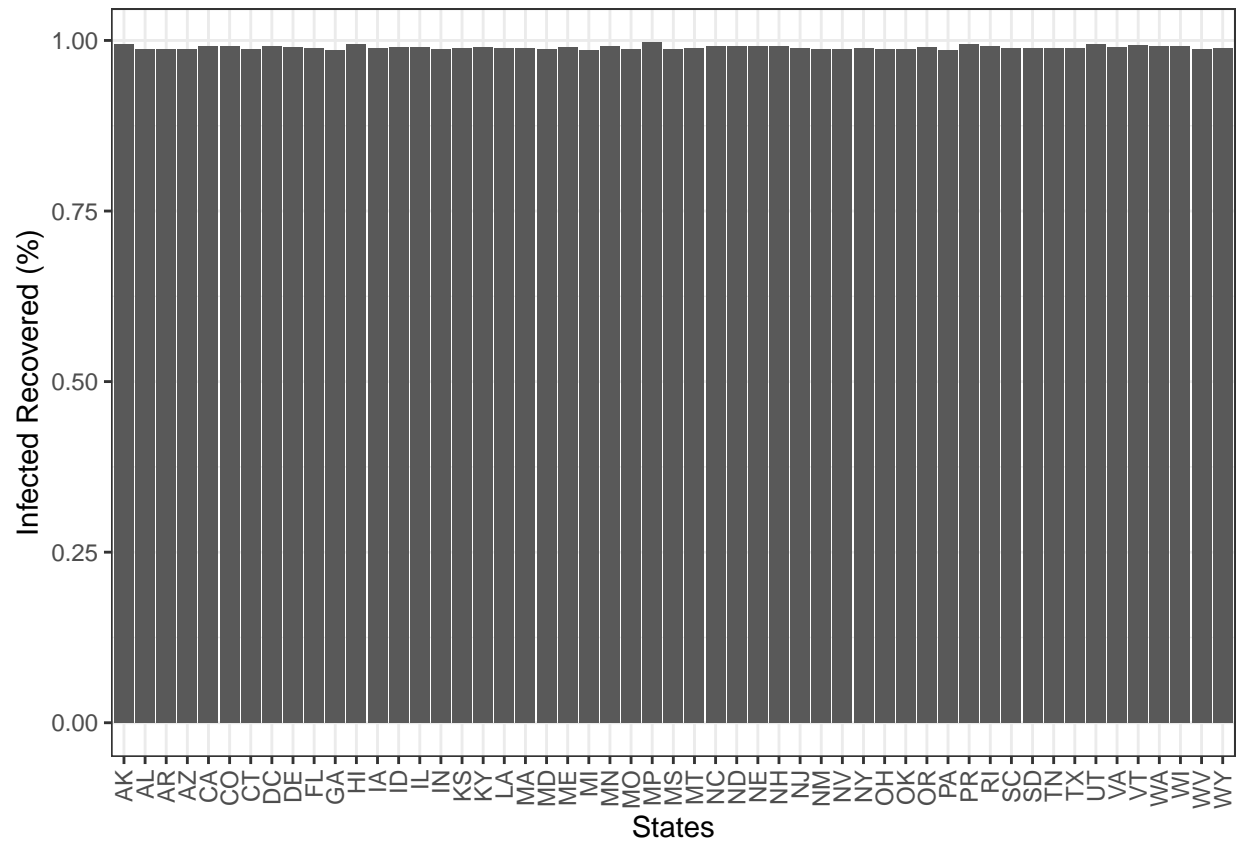
```



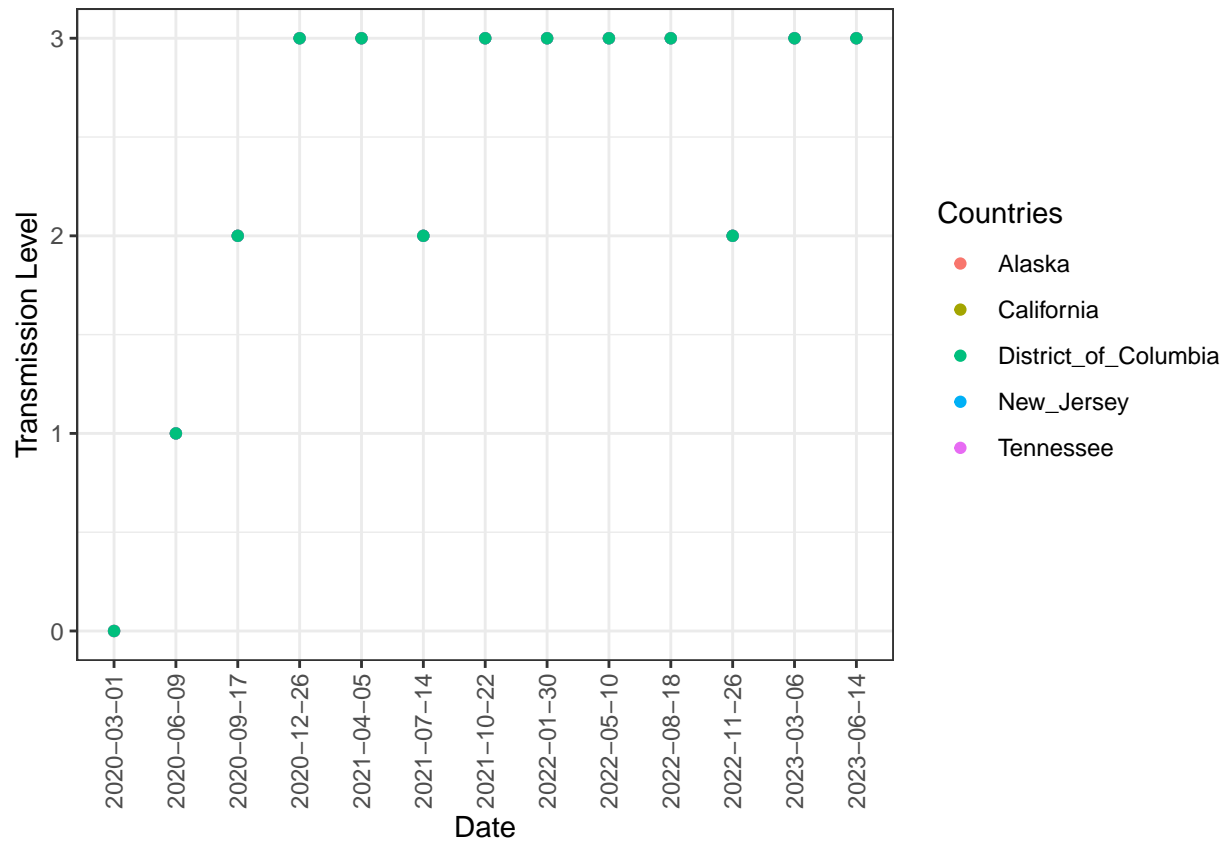
```
ggplot(data, aes(x=state,y=(data$actuals$cases/population))) +
  geom_bar(stat="identity") +
  labs(x="States",y="Infected (%)") +
  theme_bw() + scale_x_discrete(guide = guide_axis(angle=90))
```



```
ggplot(data, aes(x=state,y=((data$actuals$cases - data$actuals$deaths)/data$actuals$cases))) +
  geom_bar(stat="identity") +
  labs(x="States",y="Infected Recovered (%)") +
  theme_bw() + scale_x_discrete(guide = guide_axis(angle=90))
```



```
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() + scale_x_discrete(guide = guide_axis(angle=90))
```



```
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() + scale_x_discrete(guide = guide_

fig2<- ggplot(data_to_plot, aes(x=Date_California,y=Cases_California)) +
  geom_point() + labs(x="Date",y="Cases", title="California") + theme_bw() + scale_x_discrete(guide = gu

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() + scale_x_discrete(guide = gu

fig4<- ggplot(data_to_plot, aes(x=Date_Tennessee,y=Cases_Tennessee)) +
  geom_point() + labs(x="Date",y="Cases", title="Tennessee") + theme_bw() + scale_x_discrete(guide = gui

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() + scale_x_discrete(guide = guide_axis(angle=90))

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= 2,
  ncol = 3,labels = LETTERS[1:5],
  rel_heights = c(1,1))
```

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

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

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

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

