

Challenge-11

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

```
— Attaching core tidyverse packages ————— tidyverse 2.0.0 —
✓ dplyr      1.1.0      ✓ readr      2.1.4
✓ forcats    1.0.0      ✓ stringr    1.5.0
✓ ggplot2    3.4.3      ✓ tibble     3.1.8
✓ lubridate  1.9.2      ✓ tidyr      1.3.0
✓ purrr      1.0.1

— Conflicts ————— tidyverse_conflicts() —
✖ dplyr::filter() masks stats::filter()
✖ purrr::flatten() masks jsonlite::flatten()
✖ dplyr::lag() masks stats::lag()
! Use the ]8;;http://conflicted.r-lib.org/ conflicted package ]8;; to force all
conflicts to become errors
```

```
historic_state_data_url <- "https://api.covidactnow.org/v2/states.timeseries.json?&.K
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, 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[...
```

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

# creating new dataframe with relevant data,
time_series_transmission <- tibble(Date=time_series$cdcTransmissionLevelTimeseries[[wh
time_series_transmission$Alaska <- time_series$cdcTransmissionLevelTimeseries[[whi
time_series_transmission$California <- time_series$cdcTransmissionLevelTimeseries[[whi
time_series_transmission$New_Jersey <- time_series$cdcTransmissionLevelTimeseries[[whi
time_series_transmission$Tennessee <- time_series$cdcTransmissionLevelTimeseries[[whic
time_series_transmission$District_of_Columbia <- time_series$cdcTransmissionLevelTimes

print(head(time_series_transmission))
```

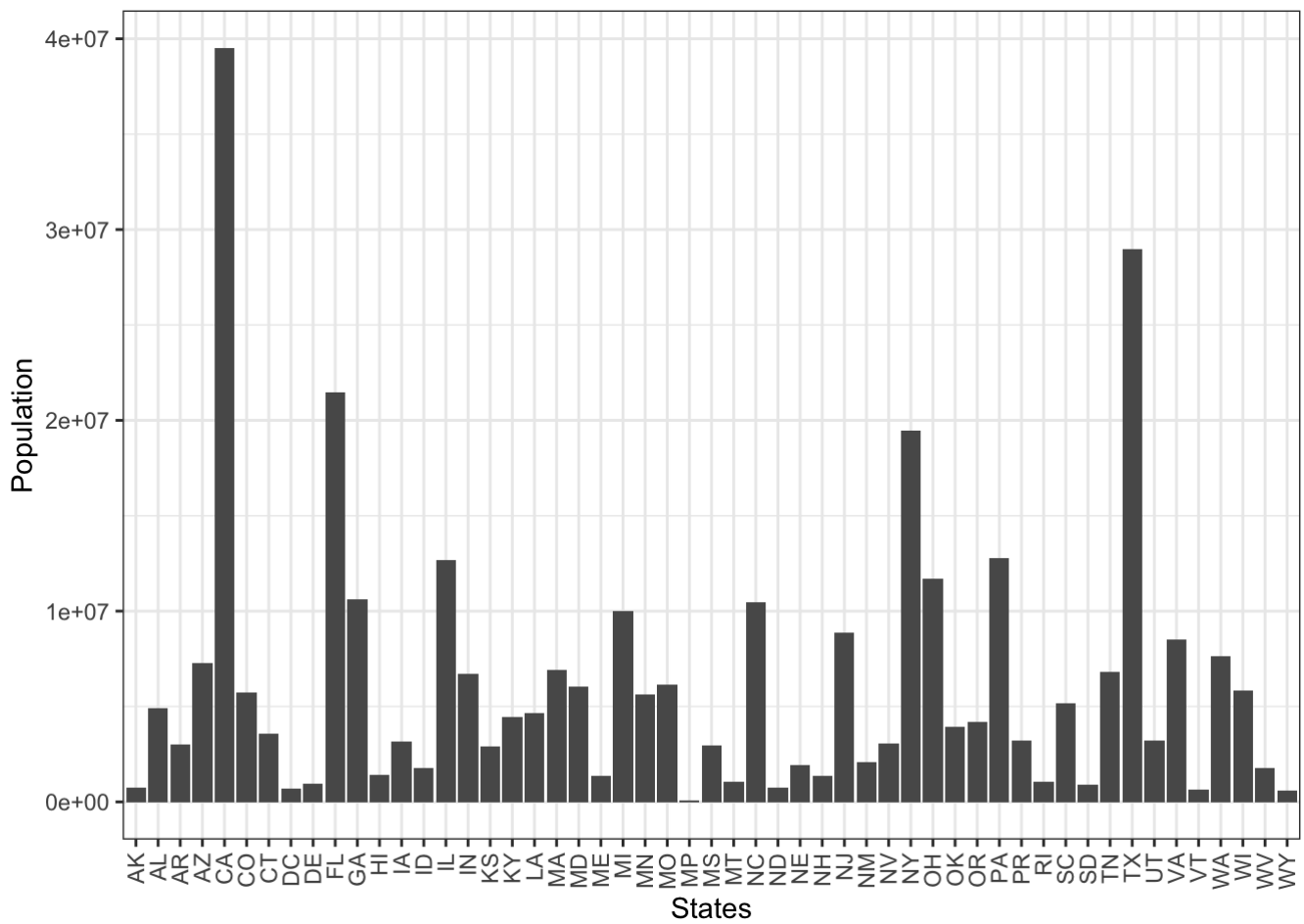
A tibble: 6 × 6

	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

```
# creating new dataframe with dates,
time_series_cases <- list(Alaska = time_series %>% filter(state=="AK") %>% select(date
time_series_cases$California <- time_series %>% filter(state=="CA") %>% select(date,ca
time_series_cases$New_Jersey <- time_series %>% filter(state=="NJ") %>% select(date,ca
time_series_cases$Tennessee <- time_series %>% filter(state=="TN") %>% select(date,cas
time_series_cases$District_of_Columbia <- time_series %>% filter(state=="DC") %>% sele
```

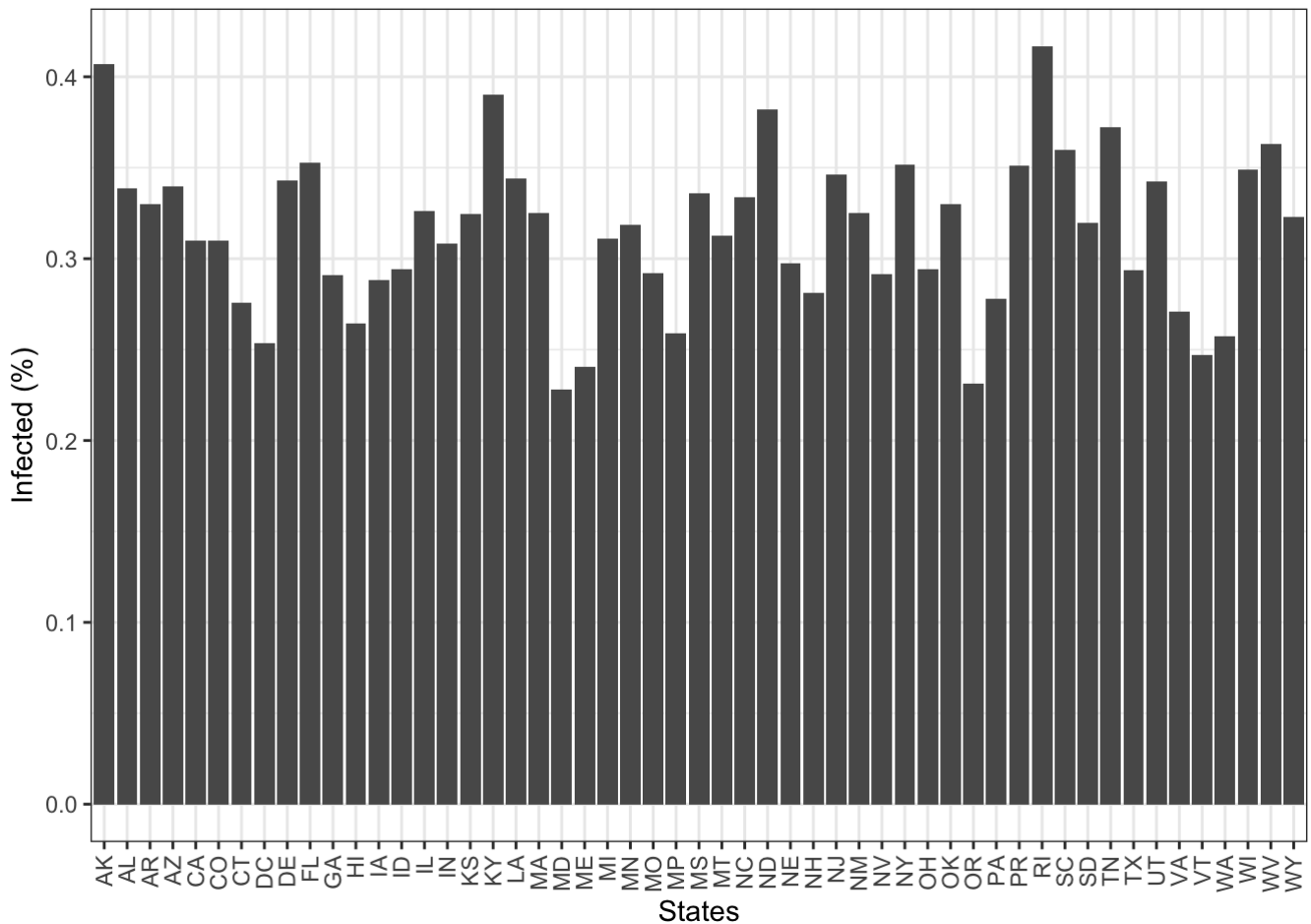
i. What is the population in the various states of U.S.A?

```
library(ggplot2)
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))
```



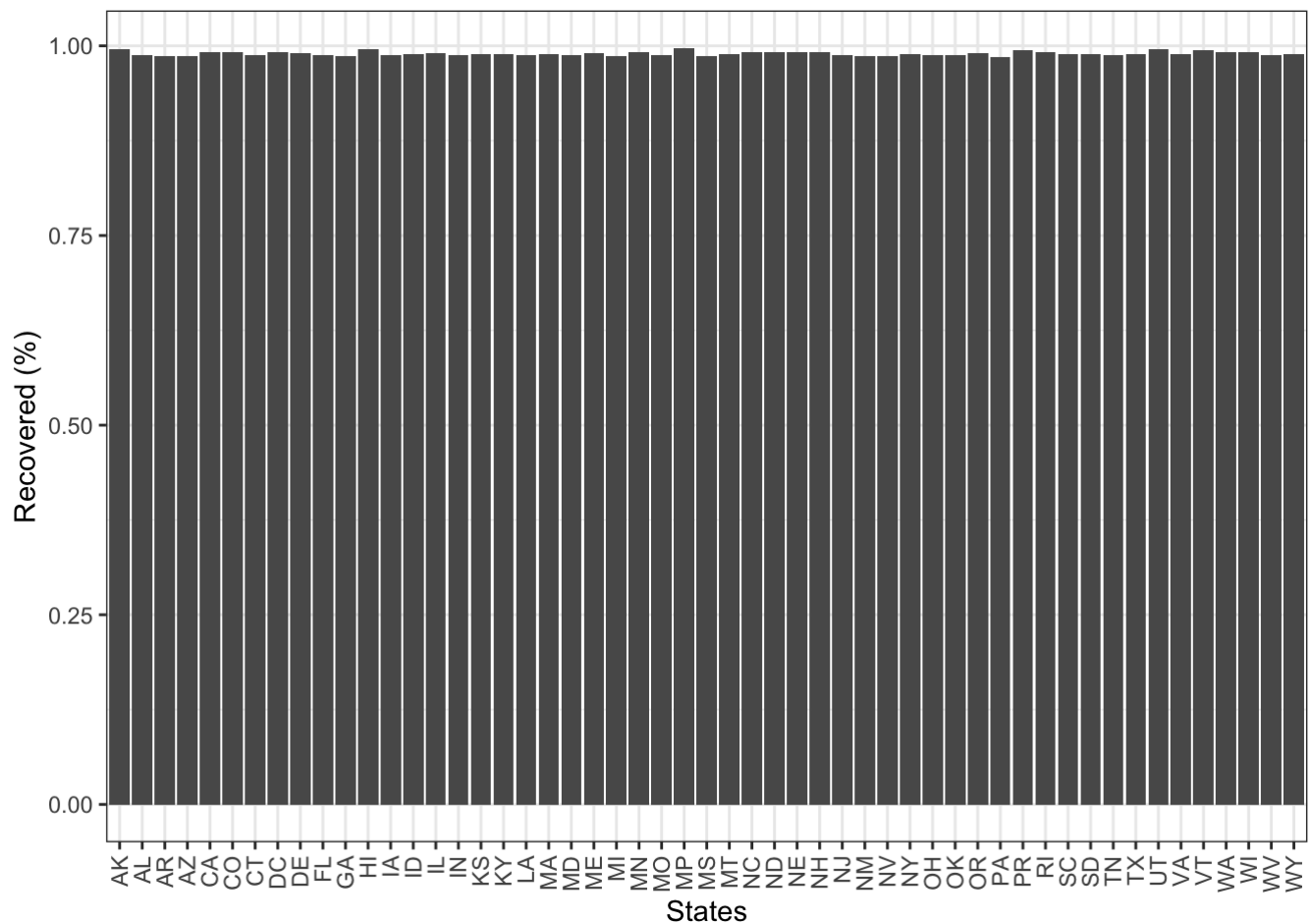
ii. What fraction of the population was infected?

```
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))
```



iii. What fraction of infected persons recovered?

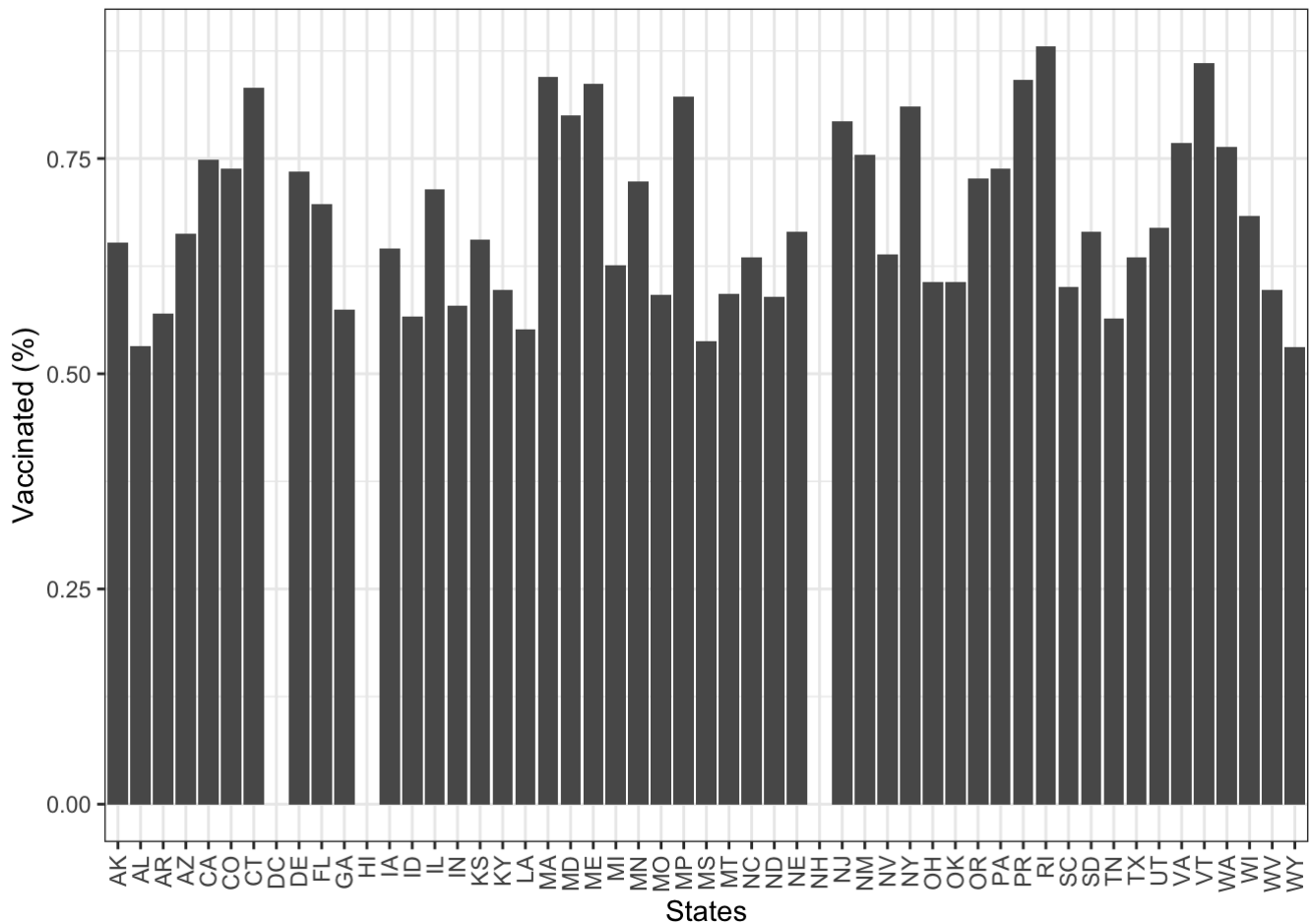
```
ggplot(data, aes(x = state, y = (data$actuals$cases - data$actuals$deaths) / actuals$c
geom_bar(stat="identity") +
labs(x="States",y="Recovered (%)") +
theme_bw() +
scale_x_discrete(guide = guide_axis(angle = 90))
```



iv. What fraction of the population is currently vaccinated?

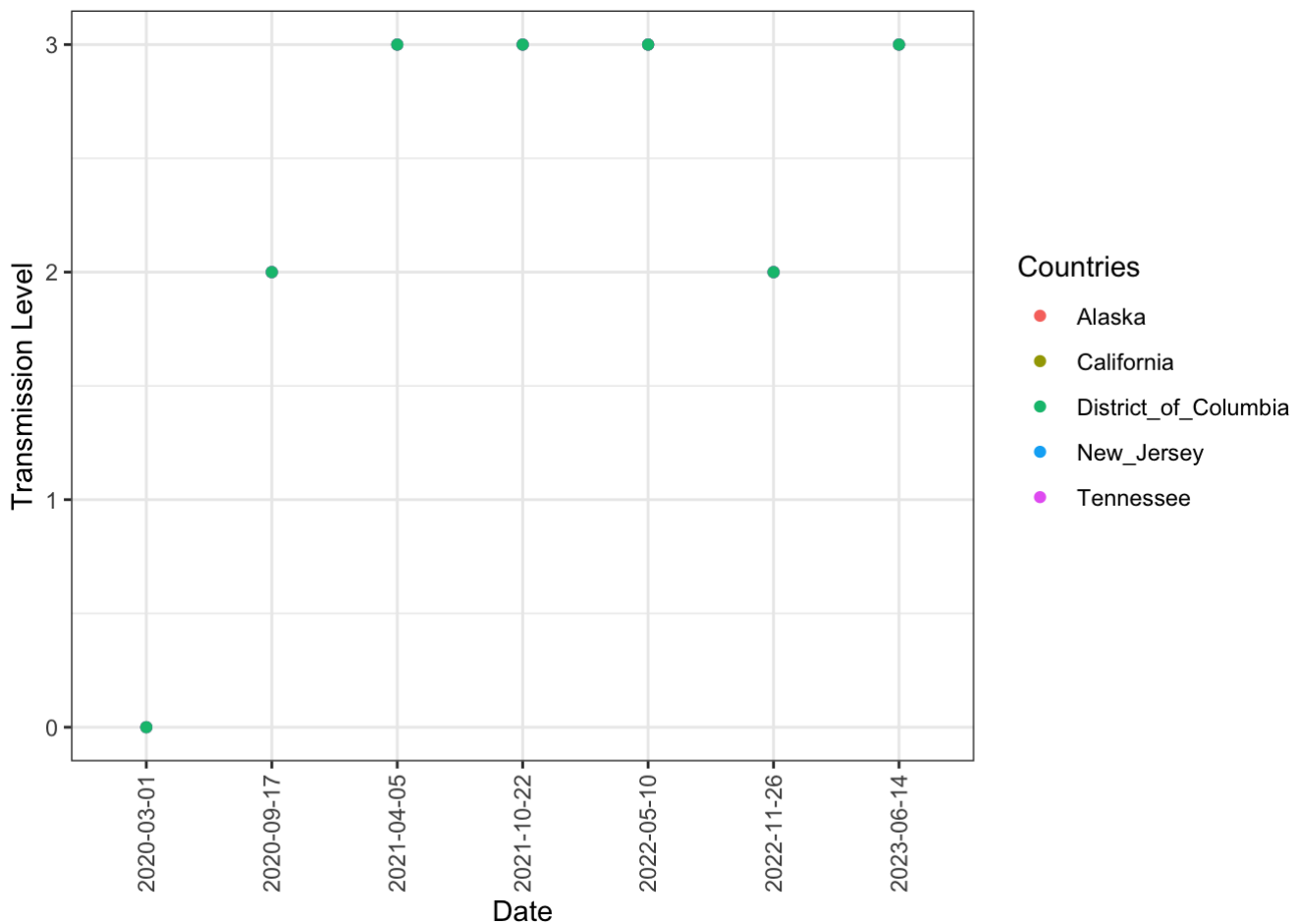
```
ggplot(data, aes(x=state,y=(data$actuals$vaccinationsCompleted/population))) +
  geom_bar(stat="identity") +
  labs(x="States",y="Vaccinated (%)") +
  theme_bw() +
  scale_x_discrete(guide = guide_axis(angle = 90))
```

Warning: Removed 3 rows containing missing values (`position_stack()`).



v. What was the transmission like in the various states?

```
time_series_transmission[seq(1,1334,by=200),]%>%
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_y_continuous(limits=c(0,1))
```



vi. How did the disease progress since it started?

```
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 × 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_discret
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_dis
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_dis
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_disc
fig5<- ggplot(data_to_plot, aes(x=Date_District_of_Columbia,y=Cases_District_of_Columb
geom_point() + labs(x="Date",y="Cases", title="District of Columbia") + theme_bw() + s

plot_grid(
fig1 + theme(legend.justification = c(0,1), axis.text.x = element_text(size = 6)),
fig2 + theme(legend.justification = c(1,0), axis.text.x = element_text(size = 6)),
fig3 + theme(legend.justification = c(0,1), axis.text.x = element_text(size = 6)),
fig4 + theme(legend.justification = c(1,0), axis.text.x = element_text(size = 6)),
fig5 + theme(legend.justification = c(0,1), axis.text.x = element_text(size = 6)),
align = "v", axis = "lr",
nrow = 2, ncol = 1,
labels = LETTERS[1:5],
rel_heights = c(1, 1))

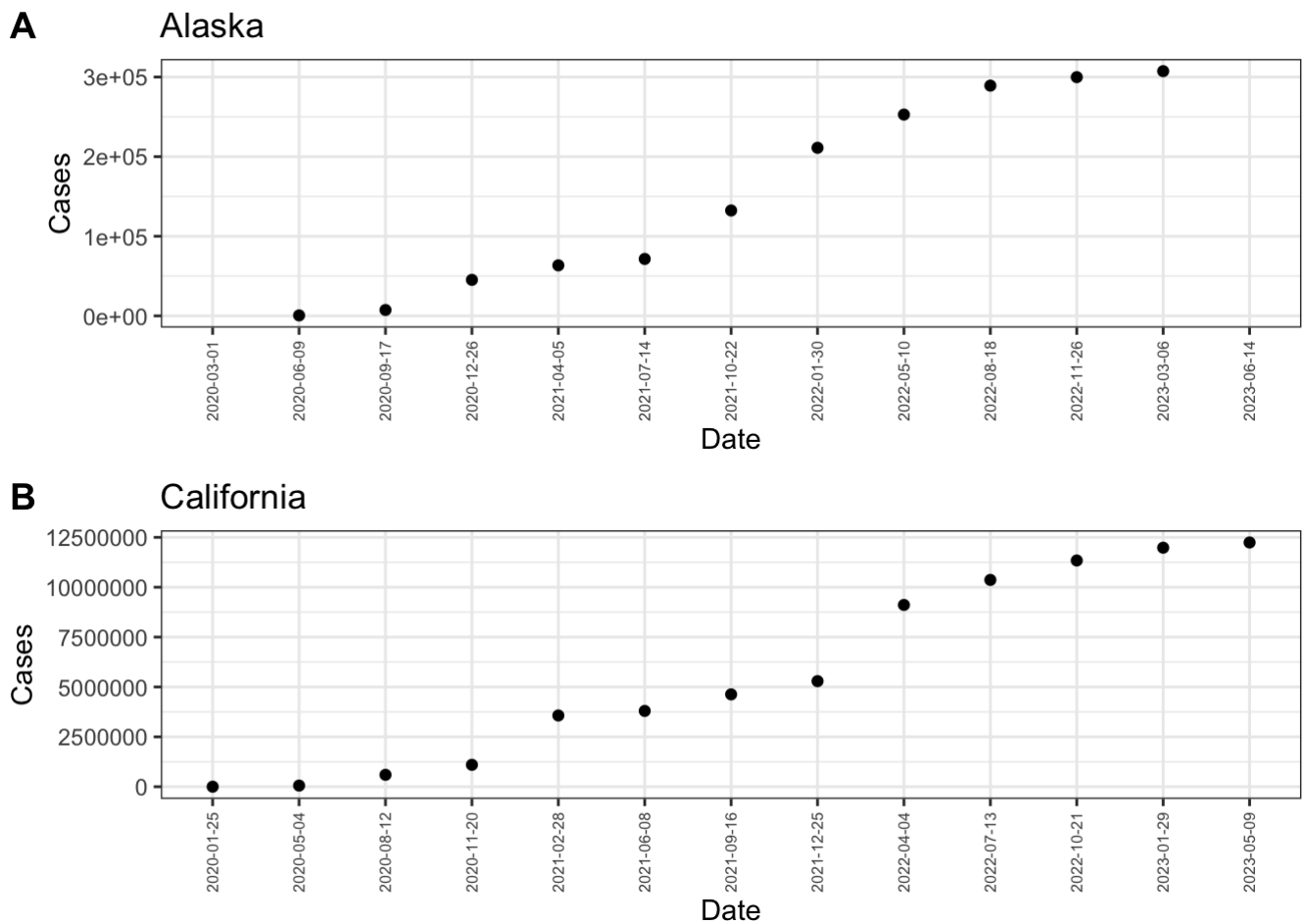
```

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

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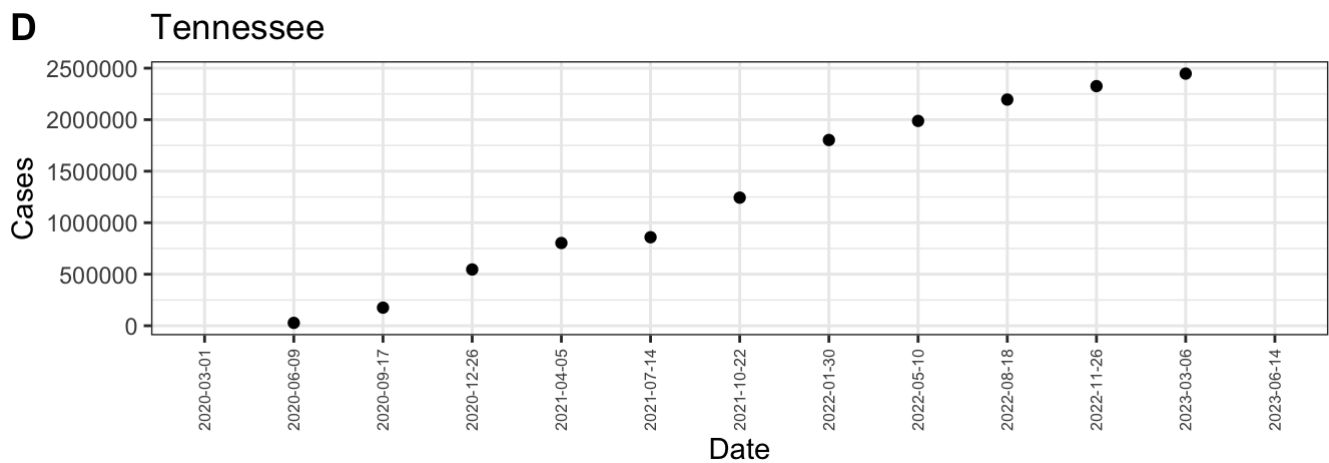
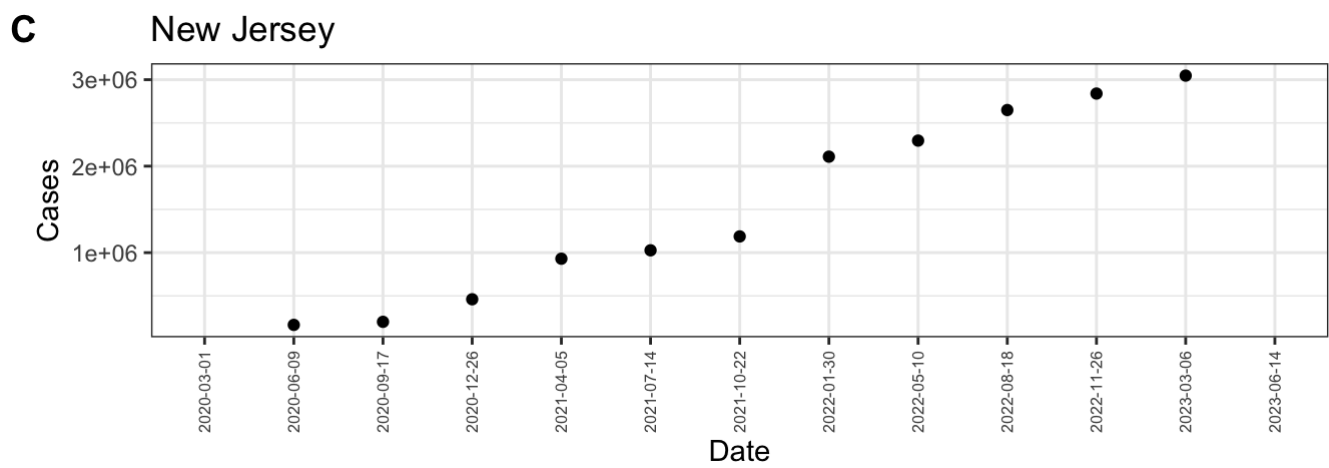
Removed 2 rows containing missing values (`geom_point()`).



```
plot_grid(
  fig3 + theme(legend.justification = c(0,1), axis.text.x = element_text(size = 6)),
  fig4 + theme(legend.justification = c(1,0), axis.text.x = element_text(size = 6)),
  align = "v", axis = "lr",
  nrow = 2, ncol = 1,
  labels = LETTERS[3:4],
  rel_heights = c(1, 1))
```

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

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



```
plot_grid(  
  fig5 + theme(legend.justification = c(0,1), axis.text.x = element_text(size = 6)),  
  align = "v", axis = "lr",  
  nrow = 2, ncol = 1,  
  labels = LETTERS[5],  
  rel_heights = c(1, 1))
```

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

E

District of Columbia

