

class14

Ethan Lai

3/4/2022

```
vax <- read.csv( "covid19vaccinesbyzipcode_test.csv" )
head(vax)
```

```
## as_of_date zip_code_tabulation_area local_health_jurisdiction county
## 1 2021-01-05 92549 Riverside Riverside
## 2 2021-01-05 92130 San Diego San Diego
## 3 2021-01-05 92397 San Bernardino San Bernardino
## 4 2021-01-05 94563 Contra Costa Contra Costa
## 5 2021-01-05 94519 Contra Costa Contra Costa
## 6 2021-01-05 91042 Los Angeles Los Angeles
## vaccine_equity_metric_quartile vem_source
## 1 3 Healthy Places Index Score
## 2 4 Healthy Places Index Score
## 3 3 Healthy Places Index Score
## 4 4 Healthy Places Index Score
## 5 3 Healthy Places Index Score
## 6 2 Healthy Places Index Score
## age12_plus_population age5_plus_population persons_fully_vaccinated
## 1 2348.4 2461 NA
## 2 46300.3 53102 61
## 3 3695.6 4225 NA
## 4 17216.1 18896 NA
## 5 16861.2 18678 NA
## 6 23962.2 25741 NA
## persons_partially_vaccinated percent_of_population_fully_vaccinated
## 1 NA NA
## 2 27 0.001149
## 3 NA NA
## 4 NA NA
## 5 NA NA
## 6 NA NA
## percent_of_population_partially_vaccinated
## 1 NA
## 2 0.000508
## 3 NA
## 4 NA
## 5 NA
## 6 NA
## percent_of_population_with_1_plus_dose booster_recip_count
## 1 NA NA
## 2 0.001657 NA
## 3 NA NA
## 4 NA NA
## 5 NA NA
## 6 NA NA
## redacted
## 1 Information redacted in accordance with CA state privacy requirements
## 2 Information redacted in accordance with CA state privacy requirements
## 3 Information redacted in accordance with CA state privacy requirements
## 4 Information redacted in accordance with CA state privacy requirements
## 5 Information redacted in accordance with CA state privacy requirements
## 6 Information redacted in accordance with CA state privacy requirements
```

Q1. What column details the total number of people fully vaccinated?

Column 9, persons_fully_vaccinated

Q2. What column details the Zip code tabulation area?

Column 2, zip_code_tabulation_area

Q3. What is the earliest date in this dataset?

```
min(vax$as_of_date)
```

```
## [1] "2021-01-05"
```

2021-01-05

Q4. What is the latest date in this dataset

```
max(vax$as_of_date)
```

```
## [1] "2022-03-01"
```

2022-03-01

As we have done previously, let's call the skim() function from the skimr package to get a quick overview of this dataset:

```
library("skimr")
skimr::skim(vax)
```

Data summary







Name	vax
Number of rows	107604
Number of columns	15
Column type frequency:	
character	5
numeric	10
Group variables	
None	

Variable type: character

skim_variable	n_missing	complete_rate	min	max	empty	n_unique	whitespace
as_of_date	0	1	10	10	0	61	0
local_health_jurisdiction	0	1	0	15	305	62	0
county	0	1	0	15	305	59	0
vem_source	0	1	15	26	0	3	0
redacted	0	1	2	69	0	2	0

Variable type: numeric

skim_variable	n_missing	complete_rate	mean	sd	p0	p25	p50	p75	p100	hist
zip_code_tabulation_area	0	1.00	93665.11	1817.39	90001	92257.75	93658.50	95380.50	97635.0	
vaccine_equity_metric_quartile	5307	0.95	2.44	1.11	1	1.00	2.00	3.00	4.0	
age12_plus_population	0	1.00	18895.04	18993.91	0	1346.95	13685.10	31756.12	88556.7	
age5_plus_population	0	1.00	20875.24	21106.02	0	1460.50	15364.00	34877.00	101902.0	

persons_fully_vaccinated	18338	0.83	12155.61	13063.88	11	1066.25	7374.50	20005.00	77744.0	
persons_partially_vaccinated	18338	0.83	831.74	1348.68	11	76.00	372.00	1076.00	34219.0	
percent_of_population_fully_vaccinated	18338	0.83	0.51	0.26	0	0.33	0.54	0.70	1.0	
percent_of_population_partially_vaccinated	18338	0.83	0.05	0.09	0	0.01	0.03	0.05	1.0	
percent_of_population_with_1_plus_dose	18338	0.83	0.54	0.28	0	0.36	0.58	0.75	1.0	
booster_recip_count	64317	0.40	4100.55	5900.21	11	176.00	1136.00	6154.50	50602.0	

Q5. How many numeric columns are in this dataset?

10

Q6. Note that there are “missing values” in the dataset. How many NA values there in the persons_fully_vaccinated column?

18338

Q7. What percent of persons_fully_vaccinated values are missing (to 2 significant figures)?

0.17

Q8. [Optional]: Why might this data be missing?

Data is redacted for legal reasons

```
library(lubridate)
```

```
##
## Attaching package: 'lubridate'
```

```
## The following objects are masked from 'package:base':
##
##   date, intersect, setdiff, union
```

```
today()
```

```
## [1] "2022-03-05"
```

```
vax$as_of_date <- ymd(vax$as_of_date)
```

Now we can do math with dates. For example: How many days have passed since the first vaccination reported in this dataset?

```
today() - vax$as_of_date[1]
```

```
## Time difference of 424 days
```

Using the last and the first date value we can now determine how many days the dataset span?

```
vax$as_of_date[nrow(vax)] - vax$as_of_date[1]
```

```
## Time difference of 420 days
```

Q9. How many days have passed since the last update of the dataset?

```
today() - vax$as_of_date[nrow(vax)]
```

```
## Time difference of 4 days
```

Q10. How many unique dates are in the dataset (i.e. how many different dates are detailed)?

```
length(unique(vax$as_of_date))
```

```
## [1] 61
```

#Working with ZIP codes

One of the numeric columns in the dataset (namely `vax$zip_code_tabulation_area`) are actually ZIP codes - a postal code used by the United States Postal Service (USPS). In R we can use the `zipcodeR` package to make working with these codes easier. For example, let's install and then load up this package and to find the centroid of the La Jolla 92037 (i.e. UC San Diego) ZIP code

```
library(zipcodeR)
geocode_zip('92037')
```

```
## # A tibble: 1 × 3
##   zipcode lat lng
##   <chr>   <dbl> <dbl>
## 1 92037   32.8 -117.
```

Calculate the distance between the centroids of any two ZIP codes in miles, e.g.

```
zip_distance('92037','92109')
```

```
##   zipcode_a zipcode_b distance
## 1      92037      92109      2.33
```

More usefully, we can pull census data about ZIP code areas (including median household income etc.). For example:

```
reverse_zipcode(c('92037', "92109"))
```

```
## # A tibble: 2 × 24
##   zipcode zipcode_type major_city post_office_city common_city_list county state
##   <chr>   <chr>         <chr>      <chr>                <blob> <chr>  <chr>
## 1 92037   Standard      La Jolla   La Jolla, CA          <raw 20 B> San D... CA
## 2 92109   Standard      San Diego  San Diego, CA          <raw 21 B> San D... CA
## # ... with 17 more variables: lat <dbl>, lng <dbl>, timezone <chr>,
## #   radius_in_miles <dbl>, area_code_list <blob>, population <int>,
## #   population_density <dbl>, land_area_in_sqmi <dbl>,
## #   water_area_in_sqmi <dbl>, housing_units <int>,
## #   occupied_housing_units <int>, median_home_value <int>,
## #   median_household_income <int>, bounds_west <dbl>, bounds_east <dbl>,
## #   bounds_north <dbl>, bounds_south <dbl>
```

#Focus on the San Diego area

Let's now focus in on the San Diego County area by restricting ourselves first to `vax$county == "San Diego"` entries. Using `dplyr` the code would look like this:

```
library(dplyr)
```

```
##
## Attaching package: 'dplyr'
```

```
## The following objects are masked from 'package:stats':
##
##   filter, lag
```

```
## The following objects are masked from 'package:base':
##
##   intersect, setdiff, setequal, union
```

```
sd <- filter(vax, county == "San Diego")  
  
nrow(sd)
```

```
## [1] 6527
```

Q11. How many distinct zip codes are listed for San Diego County?

```
length(unique(sd$zip_code_tabulation_area))
```

```
## [1] 107
```

Q12. What San Diego County Zip code area has the largest 12 + Population in this dataset?

```
byZip <- rowsum(sd$age12_plus_population, sd$zip_code_tabulation_area)  
byZip[which.max(byZip), ]
```

```
## 92154  
## 4658277
```

Zip code 92154, with population 4658277

Using dplyr select all San Diego “county” entries on “as_of_date” “2022-02-22” and use this for the following questions.

```
on22 <- filter(sd, as_of_date == "2022-02-22")
```

Q13. What is the overall average “Percent of Population Fully Vaccinated” value for all San Diego “County” as of “2022-02-22”?

```
mean (on22$percent_of_population_fully_vaccinated, na.rm=TRUE)
```

```
## [1] 0.7041551
```

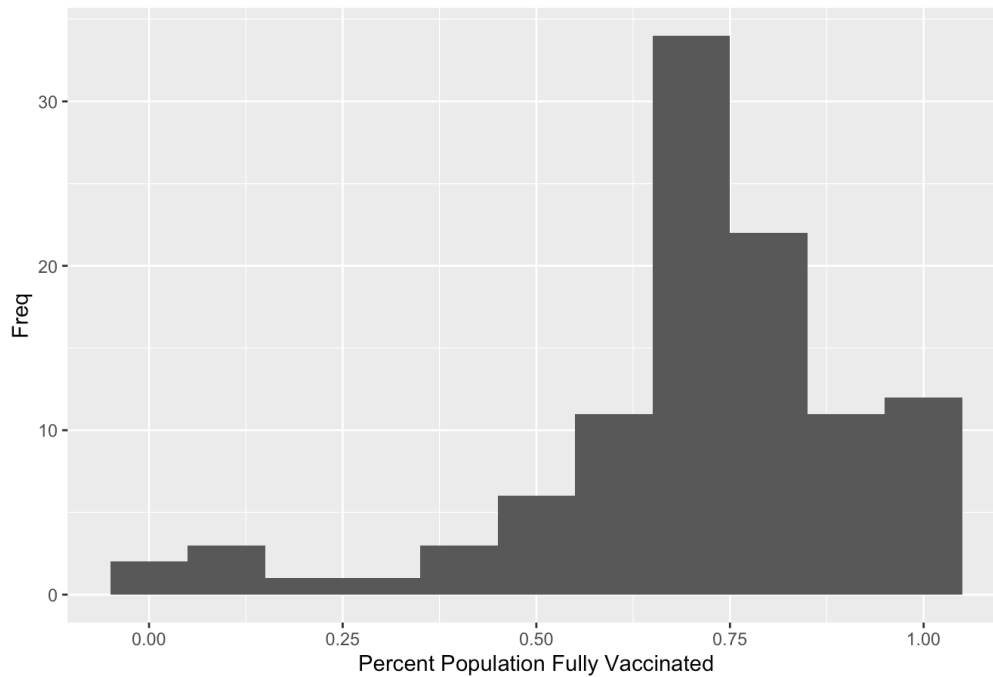
70.41%

Q14. Using either ggplot or base R graphics make a summary figure that shows the distribution of Percent of Population Fully Vaccinated values as of “2022-02-22

```
library (ggplot2)  
  
ggplot(on22, aes(x=percent_of_population_fully_vaccinated)) + geom_histogram(binwidth=0.1) + labs(title="Histogram of Vaccination Rates Across San Diego Counties", x="Percent Population Fully Vaccinated", y="Freq")
```

```
## Warning: Removed 1 rows containing non-finite values (stat_bin).
```

Histogram of Vaccination Rates Across San Diego Counties



#Focus on UCSD/La Jolla

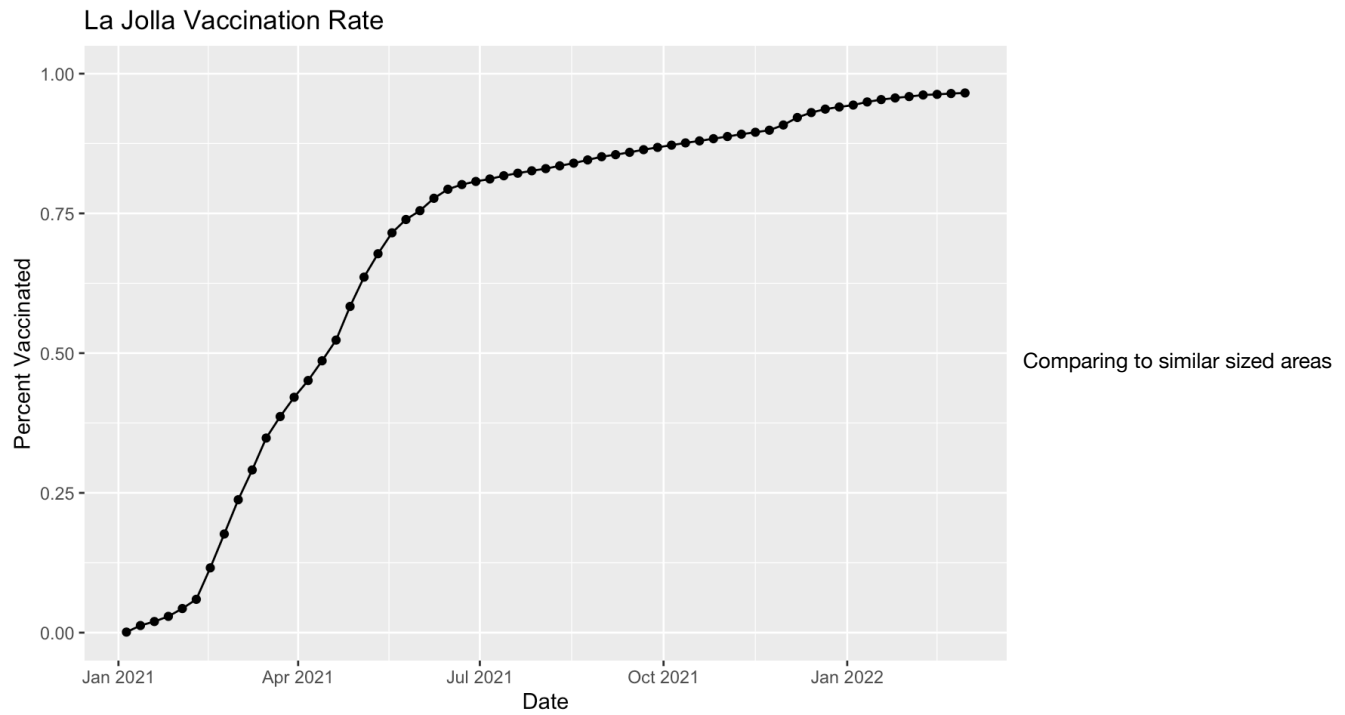
UC San Diego resides in the 92037 ZIP code area and is listed with an age 5+ population size of 36,144.

```
ucsd <- filter(sd, zip_code_tabulation_area=="92037")
ucsd[1,]$age5_plus_population
```

```
## [1] 36144
```

Q15. Using ggplot make a graph of the vaccination rate time course for the 92037 ZIP code area:

```
ucsdplot<- ggplot(ucsd) +
  aes(as_of_date,
      percent_of_population_fully_vaccinated) +
  geom_point() +
  geom_line(group=1) +
  ylim(c(0,1)) +
  labs(title = "La Jolla Vaccination Rate", x="Date", y="Percent Vaccinated")
ucsdplot
```



Let's return to the full dataset and look across every zip code area with a population at least as large as that of 92037 on as_of_date "2022-02-22".

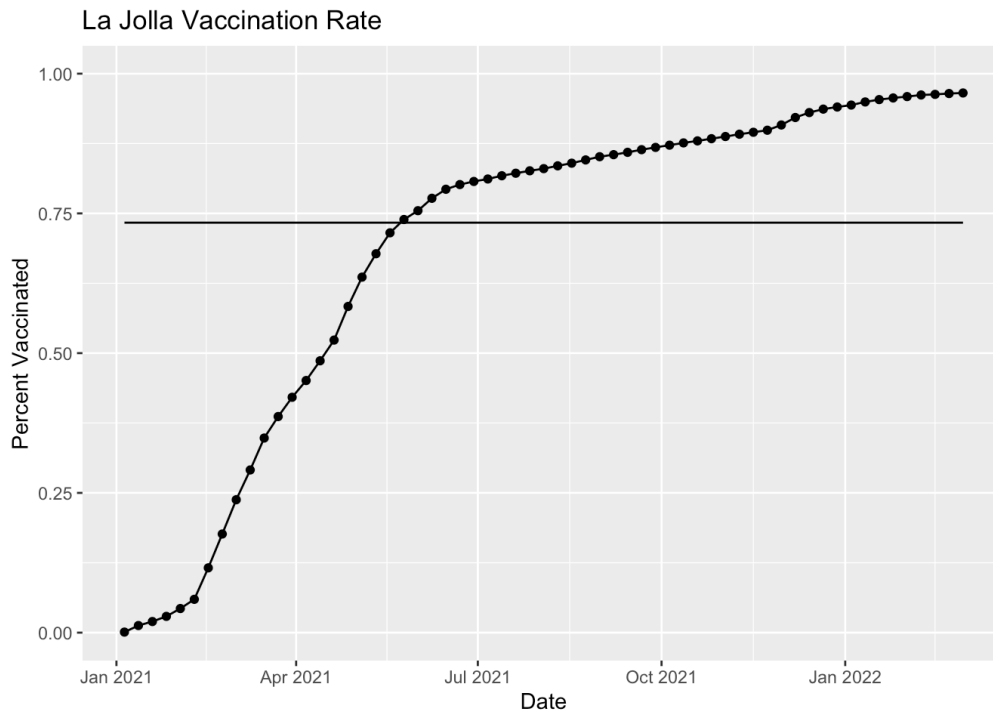
```
# Subset to all CA areas with a population as large as 92037
vax.36 <- filter(vax, age5_plus_population > 36144 &
  as_of_date == "2022-02-22")

head(vax.36)
```

```
## as_of_date zip_code_tabulation_area local_health_jurisdiction county
## 1 2022-02-22 92840 Orange Orange
## 2 2022-02-22 92064 San Diego San Diego
## 3 2022-02-22 92508 Riverside Riverside
## 4 2022-02-22 95403 Sonoma Sonoma
## 5 2022-02-22 90001 Los Angeles Los Angeles
## 6 2022-02-22 92802 Orange Orange
## vaccine_equity_metric_quartile vem_source
## 1 2 Healthy Places Index Score
## 2 4 Healthy Places Index Score
## 3 3 Healthy Places Index Score
## 4 3 Healthy Places Index Score
## 5 1 Healthy Places Index Score
## 6 2 Healthy Places Index Score
## age12_plus_population age5_plus_population persons_fully_vaccinated
## 1 47302.5 51902 40725
## 2 42177.1 46855 34266
## 3 32415.3 36303 21925
## 4 38545.9 42294 33158
## 5 47175.7 54805 43075
## 6 35113.6 39393 29268
## persons_partially_vaccinated percent_of_population_fully_vaccinated
## 1 4324 0.784652
## 2 6861 0.731320
## 3 1714 0.603945
## 4 2833 0.783988
## 5 13917 0.785968
## 6 6138 0.742975
## percent_of_population_partially_vaccinated
## 1 0.083311
## 2 0.146430
## 3 0.047214
## 4 0.066983
## 5 0.253937
## 6 0.155814
## percent_of_population_with_1_plus_dose booster_recip_count redacted
## 1 0.867963 20654 No
## 2 0.877750 15499 No
## 3 0.651159 10753 No
## 4 0.850971 18659 No
## 5 1.000000 13408 No
## 6 0.898789 12816 No
```

Q16. Calculate the mean “Percent of Population Fully Vaccinated” for ZIP code areas with a population as large as 92037 (La Jolla) as_of_date “2022-02-22”. Add this as a straight horizontal line to your plot from above with the `geom_hline()` function?

```
meanpop<- mean(vax.36$percent_of_population_fully_vaccinated)
ucsdplot + geom_line(y=meanpop)
```

Q17. What is the 6 number summary

(Min, 1st Qu., Median, Mean, 3rd Qu., and Max) of the “Percent of Population Fully Vaccinated” values for ZIP code areas with a population as large as 92037 (La Jolla) as_of_date “2022-02-22”?

```
fivenum(vax.36$percent_of_population_fully_vaccinated)
```

```
## [1] 0.3881090 0.6539015 0.7332750 0.8027110 1.0000000
```

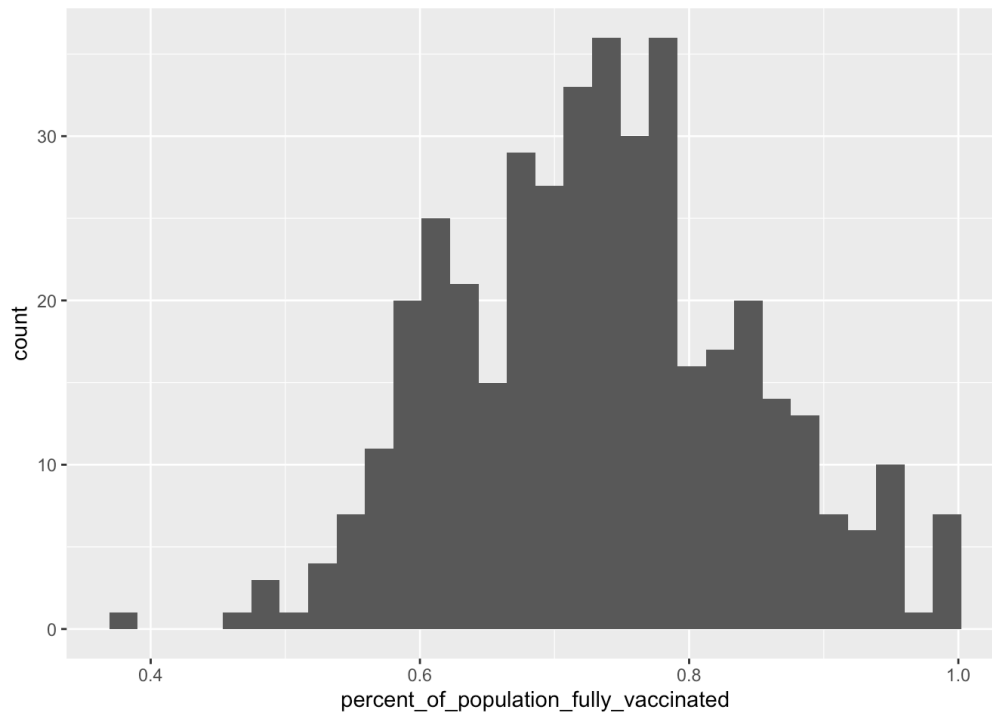
```
mean (vax.36$percent_of_population_fully_vaccinated)
```

```
## [1] 0.733385
```

Q18. Using ggplot generate a histogram of this data.

```
ggplot(vax.36, aes(x=percent_of_population_fully_vaccinated)) + geom_histogram()
```

```
## `stat_bin()` using `bins = 30`. Pick better value with `binwidth`.
```



Q19. Is the 92109 and 92040 ZIP

code areas above or below the average value you calculated for all these above?

```
vax %>% filter(as_of_date == "2022-02-22") %>%
  filter(zip_code_tabulation_area=="92040" | zip_code_tabulation_area=="92109") %>%
  select(percent_of_population_fully_vaccinated)
```

```
## percent_of_population_fully_vaccinated
## 1 0.551304
```

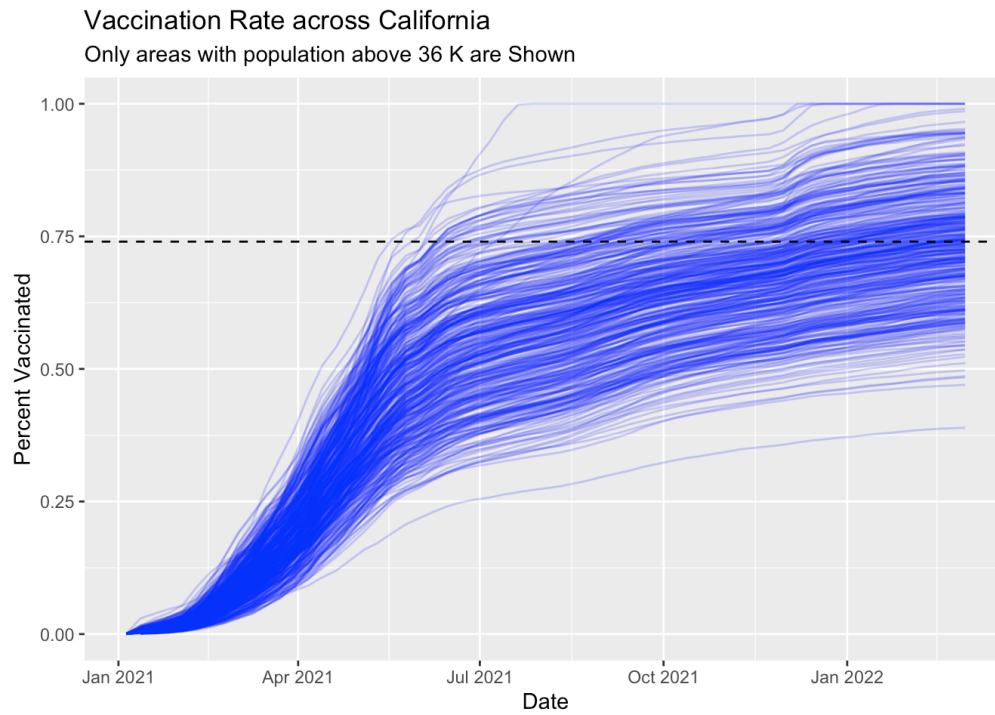
Below.

Q20. Finally make a time course plot of vaccination progress for all areas in the full dataset with a age5_plus_population > 36144.

```
vax.36.all <- filter(vax, age5_plus_population > 36144)

ggplot(vax.36.all) +
  aes(as_of_date,
    percent_of_population_fully_vaccinated,
    group=zip_code_tabulation_area) +
  geom_line(alpha=0.2, color="blue") +
  ylim(0, 1) +
  labs(x="Date", y="Percent Vaccinated",
    title="Vaccination Rate across California",
    subtitle="Only areas with population above 36 K are Shown") +
  geom_hline(yintercept = 0.74, linetype=2)
```

```
## Warning: Removed 311 row(s) containing missing values (geom_path).
```



Q21. How do you feel about traveling for Spring Break and meeting for in-person class afterwards?

Pretty much okay!