

# COVID19 Vaccination Rate Mini Project

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Here we downloaded the data from file called "Statewide COVID-19 Vaccines Administered by ZIP Code" using website <https://data.ca.gov/dataset/covid-19-vaccine-progress-dashboard-data-by-zip-code>

#First step is to import/read vaccination data

```
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?

persons\_fully\_vaccinated

Q2. What column details the Zip code tabulation area?

zip\_code\_tabulation\_area

Q3. What is the earliest date in this dataset?

2021-01-05

Q4. What is the latest date in this dataset?

2022-03-01

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

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

```
skimr::skim(vax)
```

Table 1: 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	93665.111817.39	90001	92257.7593658.5095380.5097635.0					
vaccine_equity_metric_quarter	5307	0.95	2.44	1.11	1	1.00	2.00	3.00	4.0	
age12_plus_population	0	1.00	18895.0418993.91	0	1346.95	13685.1031756.1288556.7				
age5_plus_population	0	1.00	20875.2421106.02	0	1460.50	15364.0034877.00101902.0				
persons_fully_vaccinated	18338	0.83	12155.6113063.88	11	1066.25	7374.50	20005.0077744.0			
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_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?

9

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)?

```
round((18338/107604)*100, 2)
```

```
## [1] 17.04
```

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

Not everyone gets them from the CDC, such as people who get them somewhere else such as the VA or elsewhere.

#Working with Dates

```
library(lubridate)
```

```
##
```

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

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

```
today()
```

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

```
# Specify that we are using the year-month-day format  
vax$as_of_date <- ymd(vax$as_of_date)
```

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

```
## Time difference of 422 days
```

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

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

```
age <- today() - ymd("2000-01-20")  
age
```

```
## Time difference of 8078 days
```

```
time_length(age, "year")
```

```
## [1] 22.11636
```

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

```
time difference of 2 days
```

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

```
## Time difference of 2 days
```

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

```
61 unique dates in the dataset
```

```
udates <- unique(vax$as_of_date)  
length(udates)
```

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

```
#Working with Zipcodes
```

```
library(zipcodeR)
```

```
geocode_zip('92037')
```

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

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

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

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

```
## # A tibble: 2 x 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>
```

```
# Pull data for all ZIP codes in the dataset
zipdata <- reverse_zipcode(vax$zip_code_tabulation_area )
head(zipdata)
```

```
## # A tibble: 6 x 24
##   zipcode zipcode_type major_city post_office_city common_city_list county state
##   <chr>   <chr>         <chr>         <chr>         <blob> <chr> <chr>
## 1 90001   Standard      Los Angel~ Los Angeles, CA      <raw 44 B> Los A~ CA
## 2 90002   Standard      Los Angel~ Los Angeles, CA      <raw 47 B> Los A~ CA
## 3 90003   Standard      Los Angel~ Los Angeles, CA      <raw 23 B> Los A~ CA
## 4 90004   Standard      Los Angel~ Los Angeles, CA      <raw 34 B> Los A~ CA
## 5 90005   Standard      Los Angel~ Los Angeles, CA      <raw 34 B> Los A~ CA
## 6 90006   Standard      Los Angel~ Los Angeles, CA      <raw 23 B> Los A~ 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. We have two main choices on how to do this. The first using base R the second using the dplyr package:

```
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
```

```
sd.10 <- filter(vax, county == "San Diego" &
  age5_plus_population > 10000)
```

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

107 distinct zip codes listed for SD county

```
uzip <- unique(sd$zip_code_tabulation_area)
length(uzip)
```

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

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

92154

```
sdmax12 <- which.max(sd$age12_plus_population)
print(sdmax12)
```

```
## [1] 91
```

```
sd$zip_code_tabulation_area[sdmax12]
```

```
## [1] 92154
```

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

```
sd$as_of_date[nrow(sd)]
```

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

Let's do this with the most recent date in the data-set (2022-03-01).

Q13. What is the overall average “Percent of Population Fully Vaccinated” value for all San Diego “County” as of “2022-03-01”?

```
#filter to the day  
sd.latest <- filter(sd, as_of_date == "2022-03-01")  
round(mean(sd.latest$percent_of_population_fully_vaccinated, na.rm = TRUE), 2)
```

```
## [1] 0.71
```

```
summary(sd.latest$percent_of_population_fully_vaccinated)
```

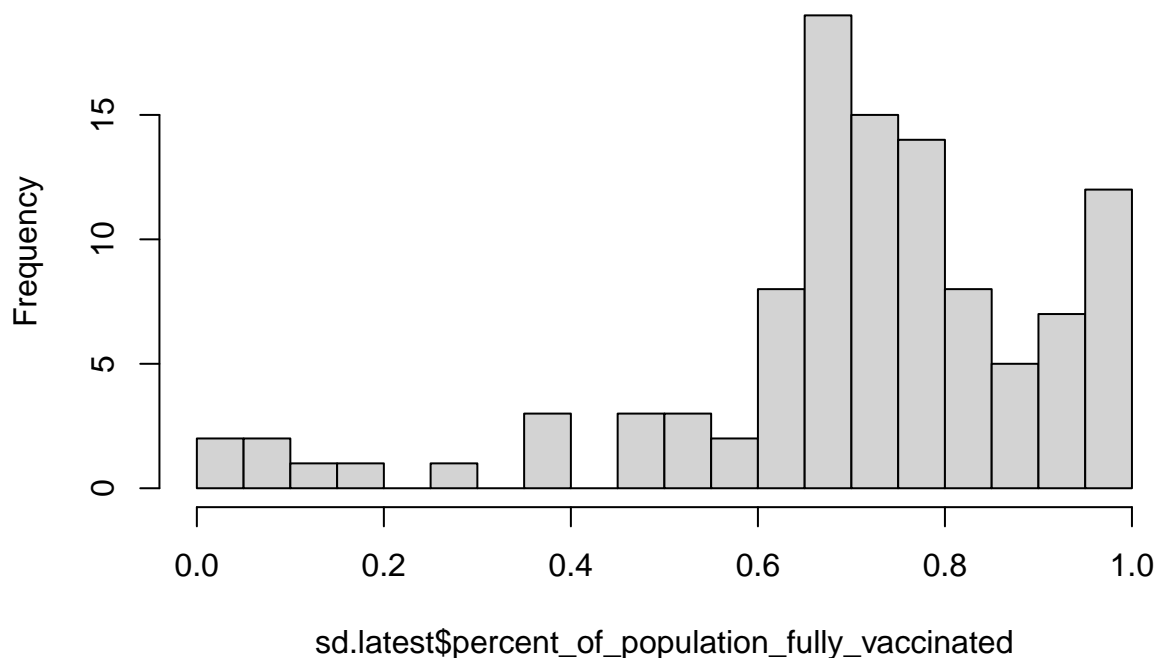
```
##      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.     NA's  
## 0.01017 0.65132 0.72452 0.70529 0.82567 1.00000         1
```

71% overall average

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 “2022-03-01”?

```
hist(sd.latest$percent_of_population_fully_vaccinated, breaks = 30)
```

## Histogram of sd.latest\$percent\_of\_population\_fully\_vaccinated

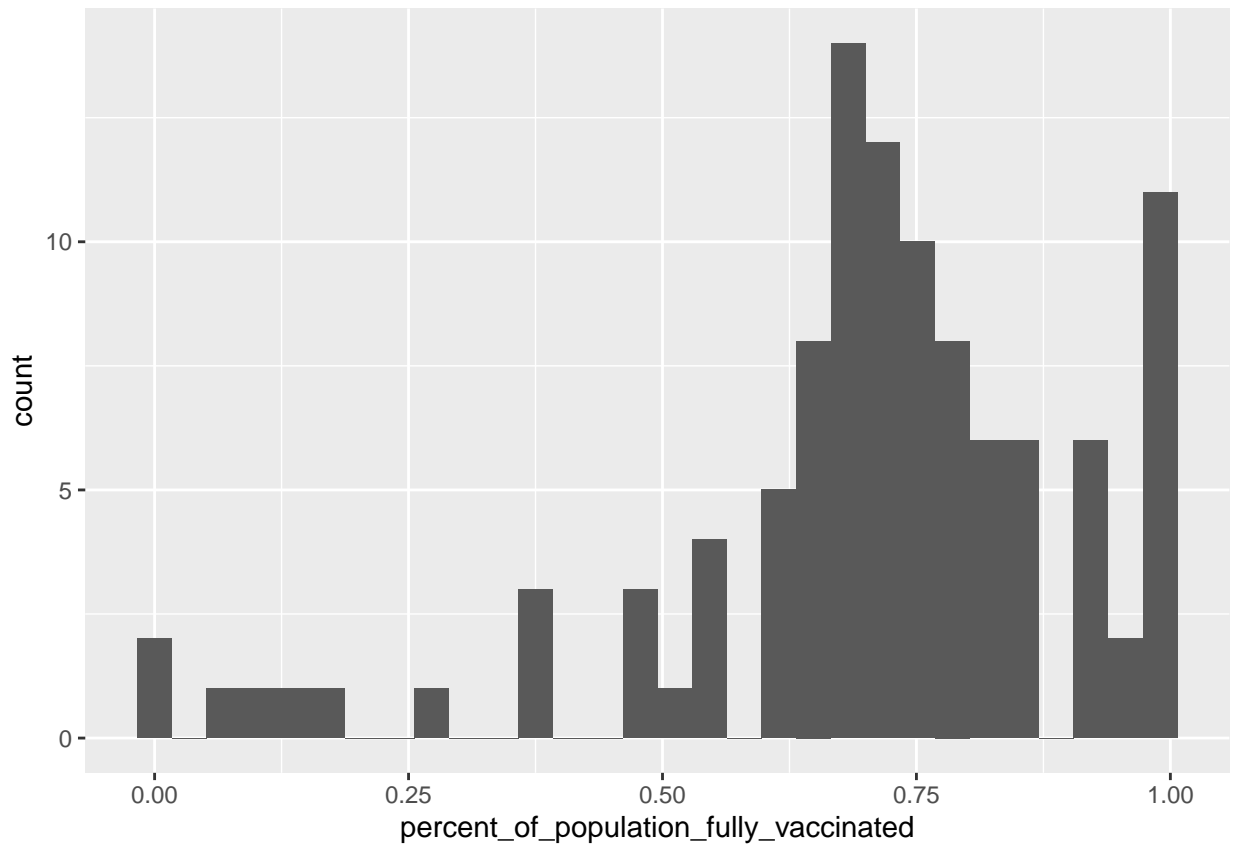


```
library(ggplot2)

ggplot(sd.latest) +aes(percent_of_population_fully_vaccinated)+ geom_histogram()

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

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



### Focus on UCSD/La Jolla

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

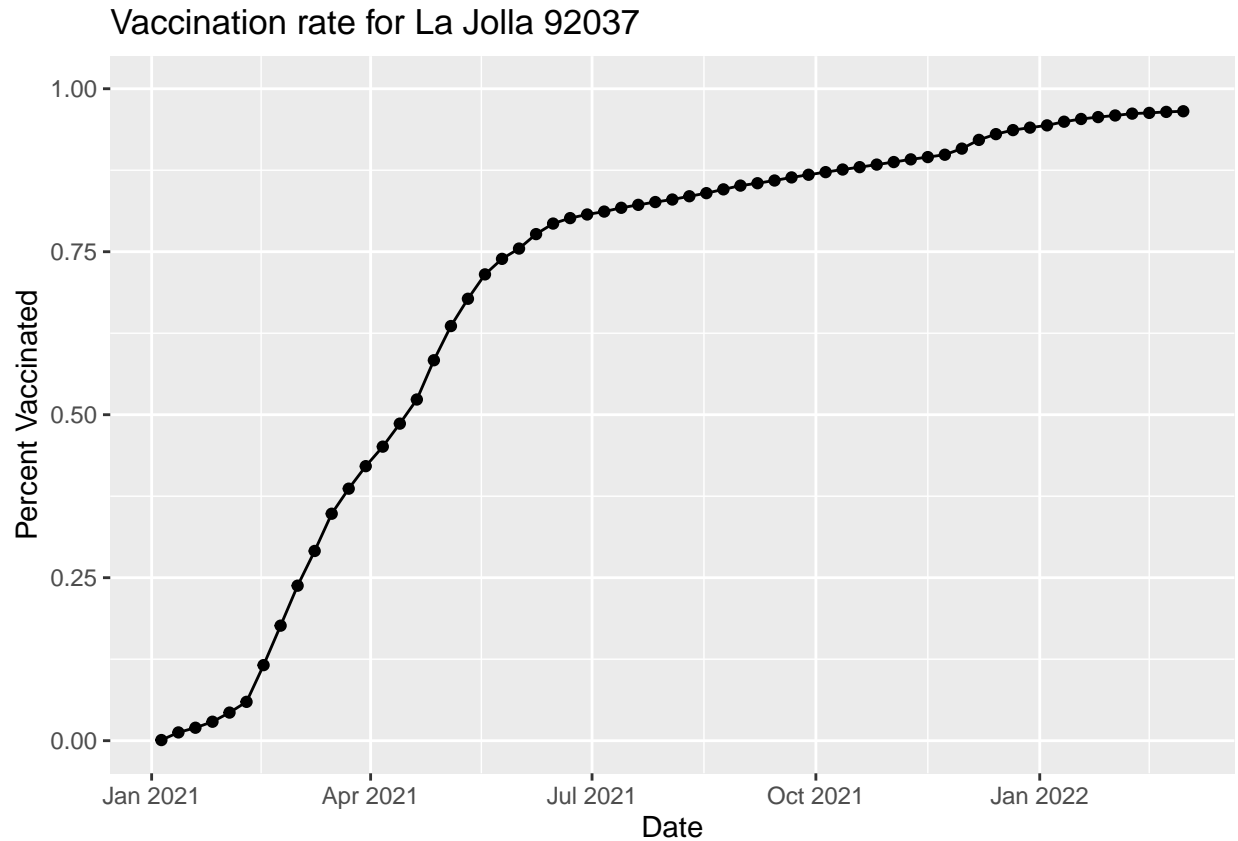
```
## [1] 36144
```

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

```
baseplot <- ggplot(ucsd) +
  aes(as_of_date,
      percent_of_population_fully_vaccinated) +
  geom_point() +
```



```
geom_line(group=1) +
ylim(c(0,1)) +
labs(x= "Date", y="Percent Vaccinated", title= "Vaccination rate for La Jolla 92037" )
baseplot
```



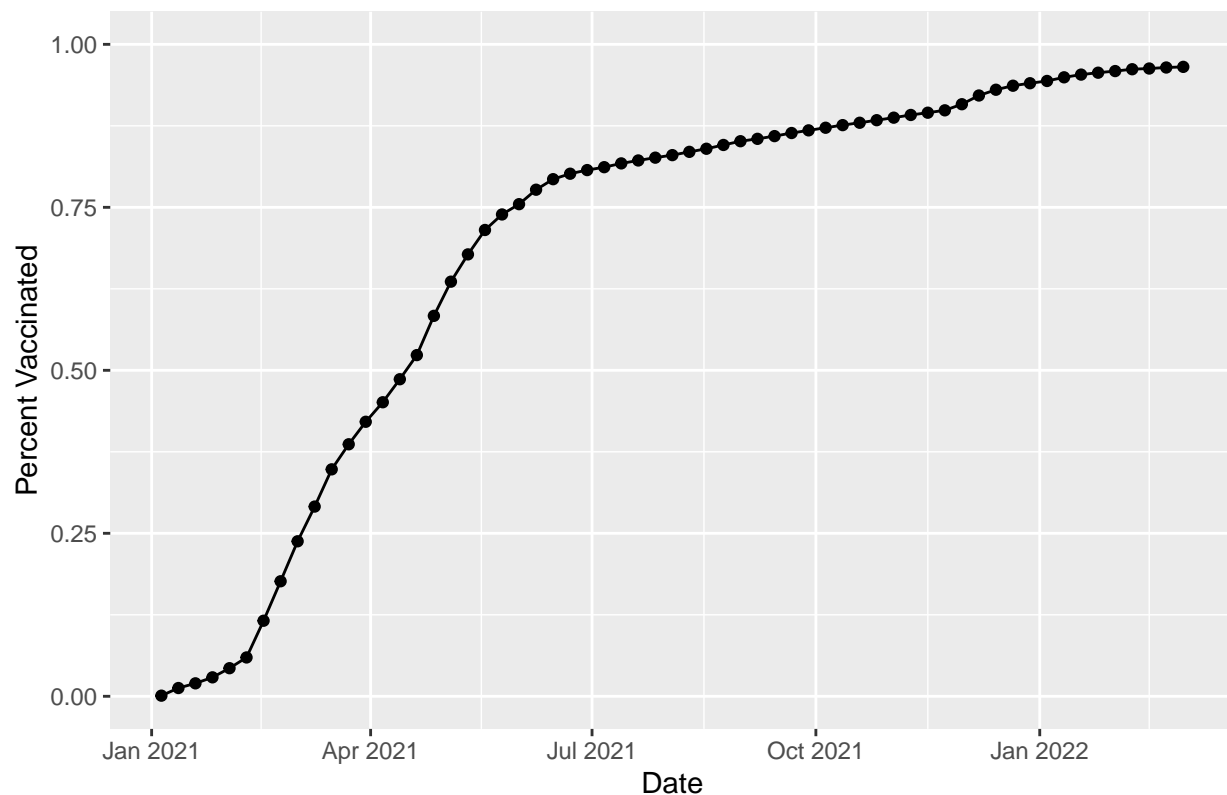
#Comparing to similar sized areas

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?

Add a line showing the average vaccination rate for all zip code areas with a population jyst as large as 92037

```
baseplot + labs(title= "Vaccination rate for CA 92037 (UCSD)")
```

Vaccination rate for CA 92037 (UCSD)



Add a line showing the average vaccination rate for all zip code areas with a population just as large as 92037

```
# Subset to all CA areas with a population as large as 92037
vax.36 <- filter(vax, age5_plus_population > 36144 &
  as_of_date == "2022-03-01")
```

```
head(vax.36)
```

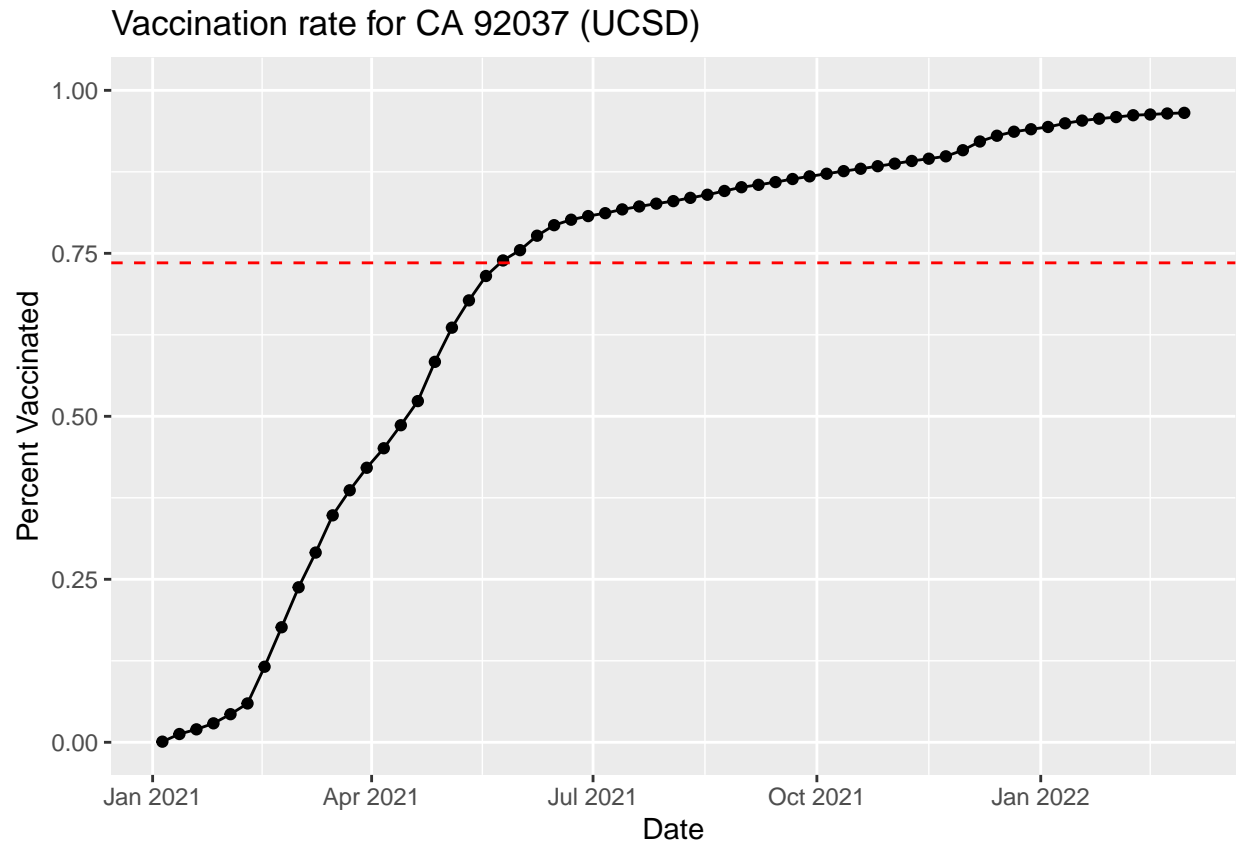
```
##   as_of_date zip_code_tabulation_area local_health_jurisdiction    county
## 1 2022-03-01          95628      Sacramento Sacramento
## 2 2022-03-01          90808      Long Beach Los Angeles
## 3 2022-03-01          92507      Riverside Riverside
## 4 2022-03-01          92626      Orange Orange
## 5 2022-03-01          93257      Tulare Tulare
## 6 2022-03-01          90011      Los Angeles Los Angeles
##   vaccine_equity_metric_quartile      vem_source
## 1                3 Healthy Places Index Score
## 2                4 Healthy Places Index Score
## 3                1 Healthy Places Index Score
## 4                3 Healthy Places Index Score
## 5                1 Healthy Places Index Score
## 6                1 Healthy Places Index Score
##   age12_plus_population age5_plus_population persons_fully_vaccinated
## 1              35579.0              38694              28842
## 2              33952.3              37179              29383
## 3              51432.5              55253              34455
```

```
## 4          44238.8          47883          33767
## 5          61519.8          70784          42919
## 6          87902.8          101902          65342
##  persons_partially_vaccinated percent_of_population_fully_vaccinated
## 1              1990              0.745387
## 2              2112              0.790312
## 3              3947              0.623586
## 4              2937              0.705198
## 5              5868              0.606338
## 6             15255              0.641224
##  percent_of_population_partially_vaccinated
## 1              0.051429
## 2              0.056806
## 3              0.071435
## 4              0.061337
## 5              0.082900
## 6              0.149703
##  percent_of_population_with_1_plus_dose booster_recip_count redacted
## 1              0.796816              16913          No
## 2              0.847118              17253          No
## 3              0.695021              15073          No
## 4              0.766535              17595          No
## 5              0.689238              17740          No
## 6              0.790927              19928          No
```

```
ave.36 <- mean(vax.36$percent_of_population_fully_vaccinated, na.rm = TRUE)
ave.36
```

```
## [1] 0.7353974
```

```
baseplot + labs(title= "Vaccination rate for CA 92037 (UCSD)") + geom_hline(yintercept=ave.36, linetype=
```



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”?

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

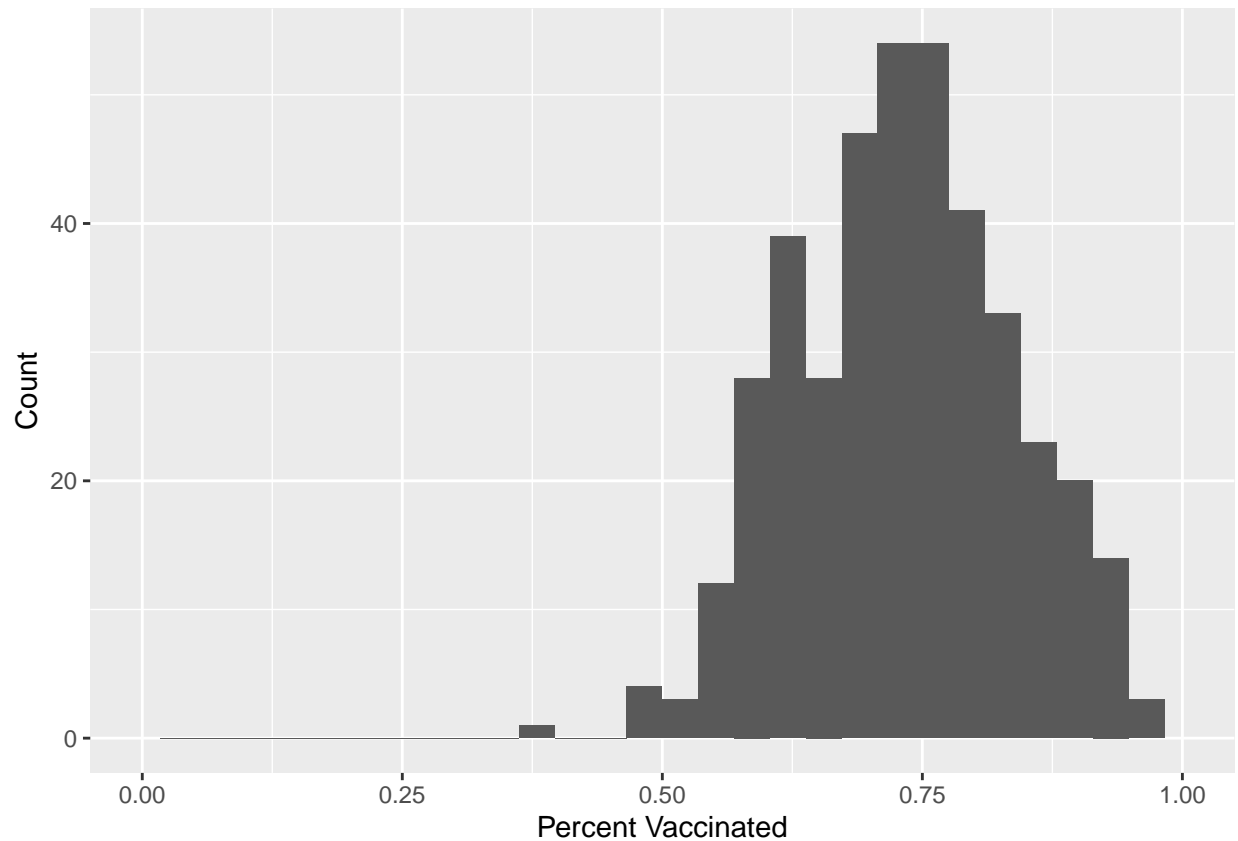
```
##      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
## 0.3890  0.6554  0.7350  0.7354  0.8044  1.0000
```

Q18. Using ggplot generate a histogram of this data.

```
library(ggplot2)
ggplot(vax.36)+ aes(percent_of_population_fully_vaccinated) + geom_histogram() +xlim(c(0,1)) + labs(x=

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

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



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 == "2022-03-01") %>%
  filter(zip_code_tabulation_area=="92040") %>%
  select(percent_of_population_fully_vaccinated)
```

```
## percent_of_population_fully_vaccinated
## 1 0.551981
```

```
vax %>% filter(as_of_date == "2022-03-01") %>%
  filter(zip_code_tabulation_area=="92109") %>%
  select(percent_of_population_fully_vaccinated)
```

```
## percent_of_population_fully_vaccinated
## 1 0.723778
```

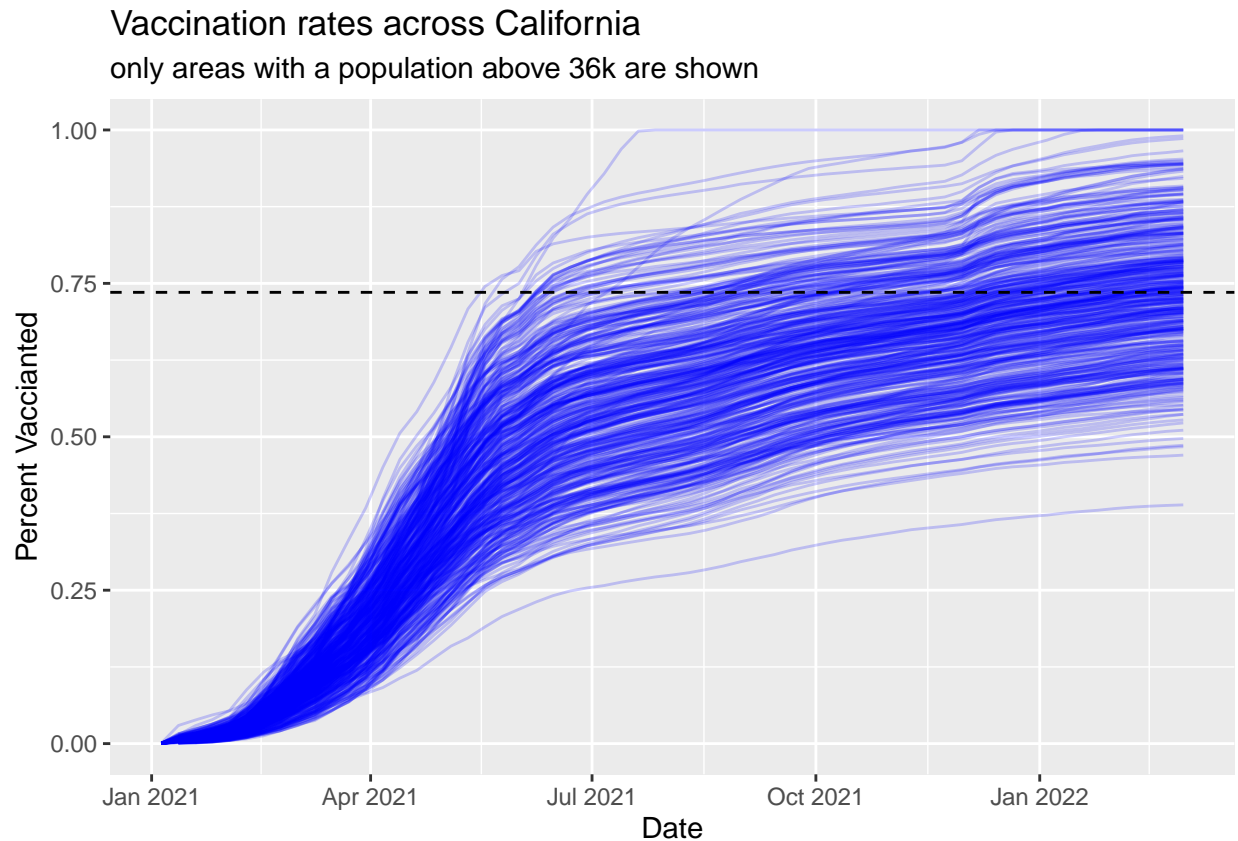
Both values are below the mean we found above.  $0.754 > 0.724$   $0.754 > 0.552$

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(c(0,1)) +
  labs(x="Date", y="Percent Vaccianted",
       title="Vaccination rates across California",
       subtitle="only areas with a population above 36k are shown") +
  geom_hline(yintercept = ave.36, linetype="dashed")
```

## 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?

I graduate this quarter so while I would be down to travel, I won't be a student anymore next quarter (sad)