

class 17 Vaccination Mini Project

Getting Started

```
vax <- read.csv("statewide-covid-19-vaccines-administered-by-zip-code.csv")  
head(vax)  
tail(vax)
```

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?

2022-11-22

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

Table 1: Data summary

Name	vax
Number of rows	174636
Number of columns	18
Column type frequency:	
character	5

Table 1: Data summary

numeric	13
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	99	0	0
local_health_jurisdiction	0	1	0	15	495	62	0	0
county	0	1	0	15	495	59	0	0
vem_source	0	1	15	26	0	3	0	0
redacted	0	1	2	69	0	2	0	0

Variable type: numeric

skim_variable	n_missing	complete	mean	sd	p0	p25	p50	p75	p100	hist
zip_code_tabulation_area	0	1.00	93665.11	1817.39	0000	192257.75	3658.50	5380.50	7635.0	
vaccine_equity_metric_618tile	0	0.95	2.44	1.11	1	1.00	2.00	3.00	4.0	
age12_plus_population	0	1.00	18895.04	8993.88	0	1346.95	13685.13	1756.18	8556.7	
age5_plus_population	0	1.00	20875.24	1105.98	0	1460.50	15364.03	1877.00	1902.0	
tot_population	8514	0.95	23372.72	2628.51	2	2126.00	18714.08	168.00	1165.0	
persons_fully_vaccinated	14921	0.91	13466.34	722.46	1	883.00	8024.00	2529.08	7186.0	
persons_partially_vaccinated	14921	0.91	1707.50	1998.80	11	167.00	1194.00	2547.00	39204.0	
percent_of_population_18065_vaccinated	18665	0.89	0.55	0.25	0	0.39	0.59	0.73	1.0	
percent_of_population_18065_fully_vaccinated	18665	0.89	0.08	0.09	0	0.05	0.06	0.08	1.0	
percent_of_population_18065_1_plus_dose	19762	0.89	0.61	0.25	0	0.46	0.65	0.79	1.0	
booster_recip_count	70421	0.60	5655.17	6867.49	11	280.00	2575.00	9421.00	58304.0	
bivalent_dose_recip_count	156958	0.10	1646.02	2161.84	11	109.00	719.00	2443.00	18109.0	
eligible_recipient_count	0	1.00	12309.10	4555.83	0	466.00	5810.00	21140.00	86696.0	

```
# To find all the NA values in the persons_fully_vaccinated column
sum( is.na(vax$persons_fully_vaccinated) )
```

```
[1] 14921
```

Q5. How many numeric columns are in this dataset?

13

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

14921

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

9.2%

Working with Dates

```
library(lubridate)
```

Loading required package: timechange

Attaching package: 'lubridate'

The following objects are masked from 'package:base':

`date`, `intersect`, `setdiff`, `union`

```
today()
```

```
[1] "2022-11-28"
```

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

Using this format, we can determine the span of the datasets.

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

Time difference of 692 days

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

Time difference of 686 days

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

6 Days

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

99 unique dates

Working with Zip-Codes

```
library(zipcodeR)
```

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

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

```
# Calculate distance of zipcode centers in miles
```

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

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

You can pull useful data from zipcodes with this function.

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

```
# A tibble: 2 x 24
  zipcode zipcode_~1 major~2 post_~3 common_c~4 county state lat lng timez~5
  <chr>   <chr>         <chr>   <chr>         <blob> <chr> <chr> <dbl> <dbl> <chr>
1 92037   Standard    La Jol~ La Jol~ <raw 20 B> San D~ CA   32.8 -117. Pacific
2 92109   Standard    San Di~ San Di~ <raw 21 B> San D~ CA   32.8 -117. Pacific
```

```
# ... with 14 more variables: 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>, and abbreviated variable names
#   1: zipcode_type, 2: major_city, 3: post_office_city, ...
```

Focus on the San Diego Area

```
# Subset to San Diego county only areas
sd <- vax[ vax$county == "San Diego" , ]
```

or, use 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] 10593
```

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

```
filter (vax, county == "San Diego" &
        age12_plus_population > 70000 &
        as_of_date == "2022-11-15")
```

	as_of_date	zip_code	tabulation_area	local_health_jurisdiction	county
1	2022-11-15		92126	San Diego	San Diego
2	2022-11-15		91911	San Diego	San Diego
3	2022-11-15		92154	San Diego	San Diego
		vaccine_equity_metric_quartile		vem_source	
1		4	Healthy Places Index Score		
2		2	Healthy Places Index Score		
3		2	Healthy Places Index Score		
		age12_plus_population	age5_plus_population	tot_population	
1		71820.2	77775	82658	
2		71642.8	79225	84026	
3		76365.2	82971	88979	
		persons_fully_vaccinated	persons_partially_vaccinated		
1		60484	5255		
2		83188	16550		
3		87151	17243		
		percent_of_population_fully_vaccinated			
1		0.731738			
2		0.990027			
3		0.979456			
		percent_of_population_partially_vaccinated			
1		0.063575			
2		0.196963			
3		0.193787			
		percent_of_population_with_1_plus_dose	booster_recip_count		
1		0.795313	39544		
2		1.000000	44281		
3		1.000000	45961		
		bivalent_dose_recip_count	eligible_recipient_count	redacted	
1		10069	59905	No	
2		6992	82731	No	
3		7033	86696	No	

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

107

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

92154

```
sd.11 <- filter(vax, county == "San Diego" &
  as_of_date == "2022-11-15")

fullyVaccPercent <- sd.11$percent_of_population_fully_vaccinated

mean(fullyVaccPercent[!is.na(fullyVaccPercent)])
```

[1] 0.7369099

Q13. What is the overall average “Percent of Population Fully Vaccinated” value for all San Diego “County” as of “2022-11-15”?

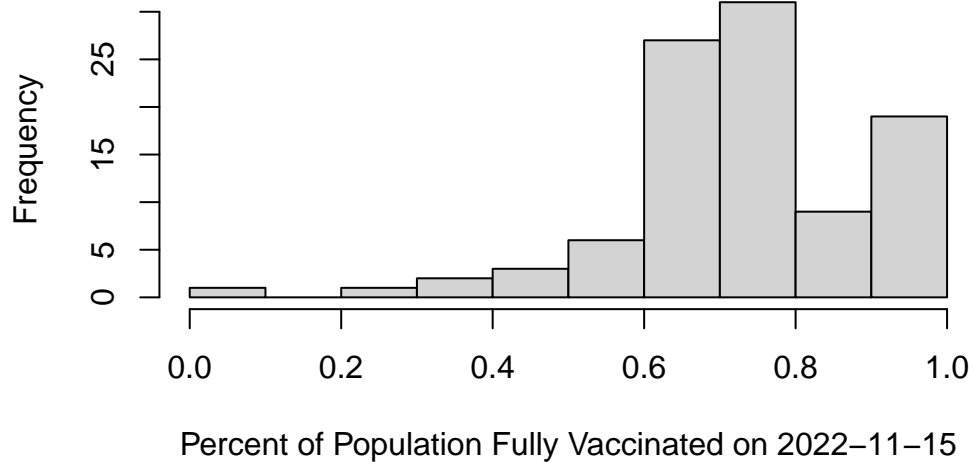
73.69%

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 “2022-11-15”?

```
library(ggplot2)

hist(fullyVaccPercent,
  main = "Histogram of Vaccination Rates Across San Diego County",
  xlab = "Percent of Population Fully Vaccinated on 2022-11-15")
```

Histogram of Vaccination Rates Across San Diego Count



Focus on UCSD/La Jolla

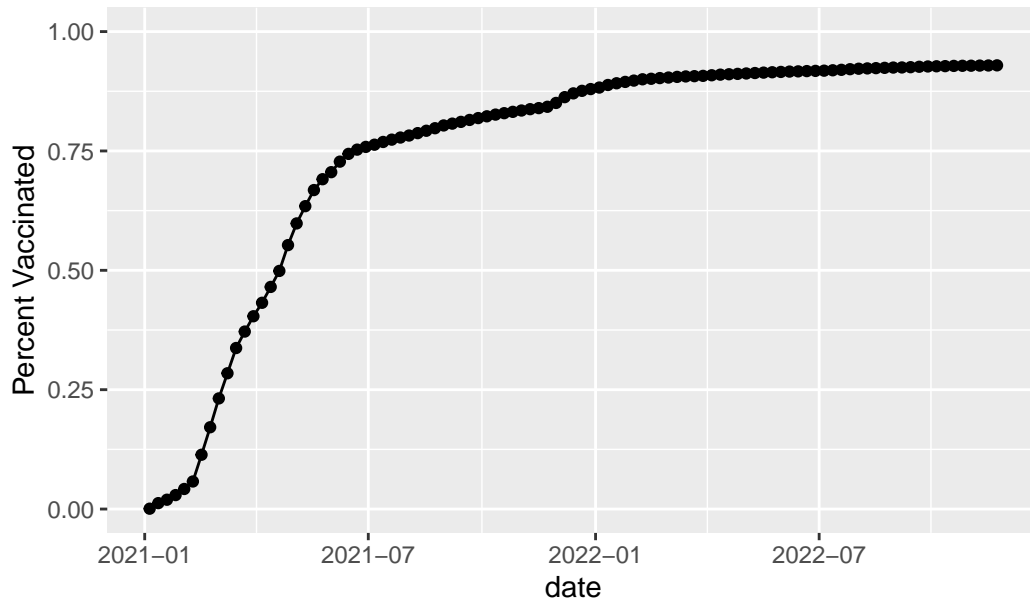
```
ucsd <- filter(sd, zip_code_tabulation_area=="92037")  
  
ucsd[1,]$age5_plus_population
```

[1] 36144

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

```
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 CA 92109")
```


Vaccination rate for La Jolla CA 92109



Comparing to Similar Sized Area

```
# Subset to all CA areas with a population as large as 92037
vax.36 <- filter(vax, age5_plus_population > 36144 &
  as_of_date == "2022-11-15")

head(vax.36)
```

	as_of_date	zip_code_tabulation_area	local_health_jurisdiction	county
1	2022-11-15	92236	Riverside	Riverside
2	2022-11-15	92130	San Diego	San Diego
3	2022-11-15	94121	San Francisco	San Francisco
4	2022-11-15	94551	Alameda	Alameda
5	2022-11-15	94112	San Francisco	San Francisco
6	2022-11-15	94303	Santa Clara	Santa Clara

	vaccine_equity_metric_quartile	vem_source
1	1	Healthy Places Index Score
2	4	Healthy Places Index Score
3	4	Healthy Places Index Score
4	4	Healthy Places Index Score
5	3	Healthy Places Index Score
6	3	Healthy Places Index Score

	age12_plus_population	age5_plus_population	tot_population
1	38505.3	42923	45477
2	46300.3	53102	56134
3	39105.0	41363	43616
4	38947.9	43399	47227
5	75681.8	81107	84707
6	40033.3	44989	48244

	persons_fully_vaccinated	persons_partially_vaccinated
1	30465	3858
2	52380	5751
3	36566	2373
4	32557	2333
5	78358	4646
6	41275	4175

	percent_of_population_fully_vaccinated
1	0.669899
2	0.933124
3	0.838362
4	0.689373
5	0.925048
6	0.855547

	percent_of_population_partially_vaccinated
1	0.084834
2	0.102451
3	0.054407
4	0.049400
5	0.054848
6	0.086539

	percent_of_population_with_1_plus_dose	booster_recip_count
1	0.754733	12943
2	1.000000	34821
3	0.892769	28345
4	0.738773	20223
5	0.979896	56744
6	0.942086	26288

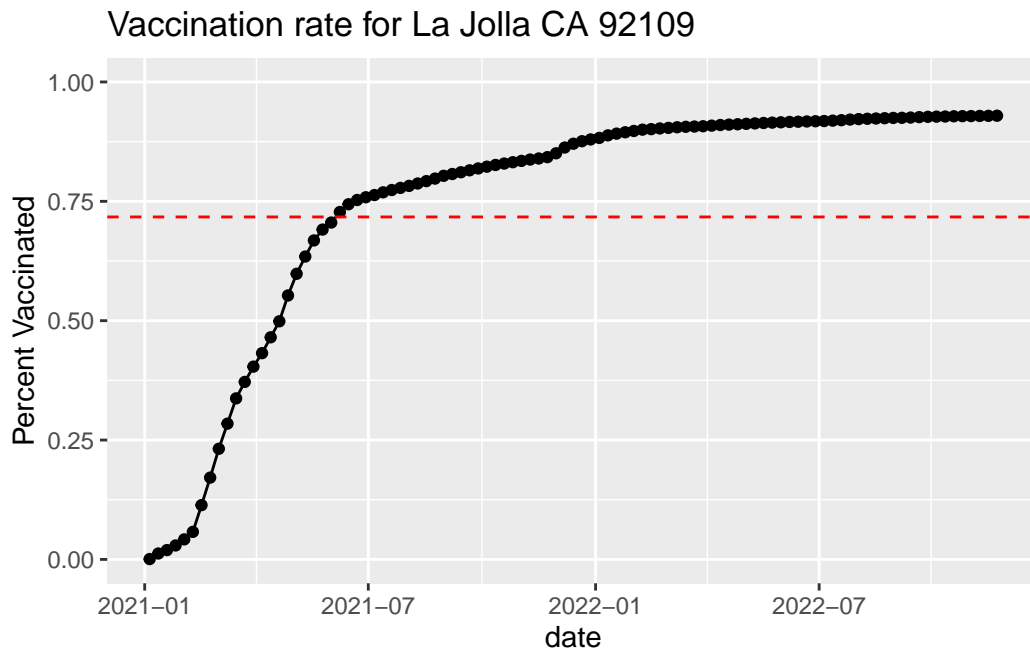
	bivalent_dose_recip_count	eligible_recipient_count	redacted
1	1395	30375	No
2	11203	51780	No
3	10994	36013	No
4	5568	32234	No
5	16019	77580	No
6	8573	40853	No

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 “2022-11-15”. Add this as a straight horizontal line to your plot from above with the `geom_hline()` function?

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

```
[1] 0.7172851
```

```
ggplot(ucsd) +  
  aes(as_of_date,  
       percent_of_population_fully_vaccinated) +  
  geom_point() +  
  geom_line(group=1) +  
  ylim(c(0,1)) +  
  geom_hline(yintercept = 0.7172851, linetype = "dashed", col = "red") +  
  labs(x = "date", y="Percent Vaccinated",  
       title = "Vaccination rate for La Jolla CA 92109")
```



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 “2022-11-15”?

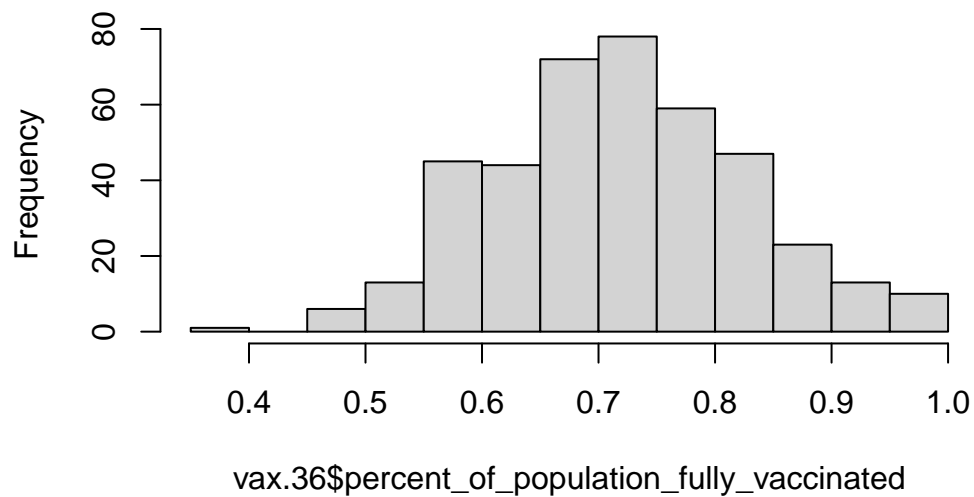
```
summary(vax.36$percent_of_population_fully_vaccinated)
```

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
0.3785	0.6396	0.7155	0.7173	0.7880	1.0000

Q18. Using ggplot generate a histogram of this data.

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

Histogram of vax.36\$percent_of_population_fully_vaccinated



Q19. Is the 92109 and 92040 ZIP code areas above or below the average value you calculated for all these above?

```
filter(vax, zip_code_tabulation_area == "92109", as_of_date == "2022-11-15")$percent_of_po
```

```
[1] 0.693299
```

```
filter(vax, zip_code_tabulation_area == "92040", as_of_date == "2022-11-15")$percent_of_po
```

```
[1] 0.546646
```

Both of these area codes are below the average.

```
vax %>% filter(as_of_date == "2022-11-15") %>%  
  filter(zip_code_tabulation_area=="92040") %>%  
  select(percent_of_population_fully_vaccinated)
```

```
percent_of_population_fully_vaccinated  
1                                0.546646
```

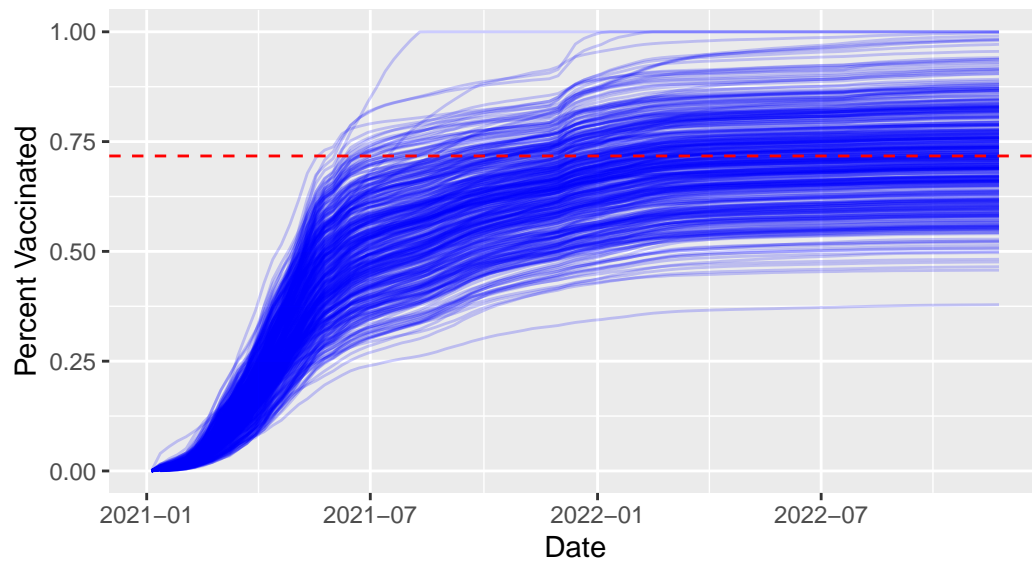
Q20. Finally make a time course plot of vaccination progress for all areas in the full dataset with a age5_plus_population > 36144.

```
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 Vaccinated",  
       title = "Vaccination Rate Across California",  
       subtitle = "Only areas with a population above 36k are shown") +  
  geom_hline(yintercept = 0.7172851, linetype="dashed", col = "red")
```

Warning: Removed 184 rows containing missing values (`geom_line()`).

Vaccination Rate Across California

Only areas with a population above 36k are shown



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

Great.