Lab_15

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G	<pre># Import vaccination data vax <- read.csv('covid19vacchead(vax)</pre>	cinesbyzipo	ode_tes	t.csv')		
	as_of_date zip_code_tabulati	on_area lo	cal_hea	lth_ju	risdiction		county
1	2021-01-05	92240			Riverside	Riv	erside
2	2021-01-05	91302			os Angeles		ngeles
3	2021-01-05	93420		San Lı	ıis Obispo	San Luis	Obispo
	2021-01-05	91901			San Diego	San	Diego
5	2021-01-05	94110		San	Francisco	San Fra	ncisco
6	2021-01-05	91902			San Diego	San	Diego
	${\tt vaccine_equity_metric_quarti}$	le		vem_s	source		
1		1 Healthy	Places	Index	Score		
2		4 Healthy					
3		3 Healthy					
4		3 Healthy					
5		4 Healthy	Places	Index	Score		

 ${\tt age12_plus_population~age5_plus_population~tot_population}$

29270.5

6

4 Healthy Places Index Score

35278

33093

```
2
                                                          26712
                 23163.9
                                          25899
3
                 26694.9
                                          29253
                                                          30740
4
                 15549.8
                                          16905
                                                          18162
5
                 64350.7
                                          68320
                                                          72380
6
                 16620.7
                                                          18896
                                          18026
 persons_fully_vaccinated persons_partially_vaccinated
                          NA
2
                          15
                                                        614
3
                         NA
                                                         NA
4
                                                         NA
                          NA
5
                          17
                                                       1268
6
                          15
                                                        397
 percent_of_population_fully_vaccinated
                                         NA
1
2
                                  0.000562
3
                                         NA
4
                                         NA
5
                                  0.000235
6
                                  0.000794
 percent_of_population_partially_vaccinated
1
2
                                       0.022986
3
                                             NA
4
                                             NA
5
                                       0.017519
6
                                       0.021010
 percent_of_population_with_1_plus_dose booster_recip_count
1
                                         NA
                                                              NA
2
                                  0.023548
                                                              NA
3
                                         NA
                                                              NA
4
                                         NA
                                                              NΑ
5
                                  0.017754
                                                              NA
6
                                  0.021804
                                                              NA
 bivalent_dose_recip_count eligible_recipient_count
                                                       2
1
                           NA
2
                           NA
                                                      15
3
                           NA
                                                       4
4
                           NA
                                                       8
5
                           NA
                                                      17
6
                           NA
                                                      15
```

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

head(vax\$as_of_date)

- [1] "2021-01-05" "2021-01-05" "2021-01-05" "2021-01-05" "2021-01-05"
- [6] "2021-01-05"

tail(vax\$as_of_date)

- [1] "2022-11-22" "2022-11-22" "2022-11-22" "2022-11-22" "2022-11-22"
- [6] "2022-11-22"

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

A: persons_fully_vaccinated

Q2. What column details the Zip code tabulation area?

A: zip_code_tabulation_area

Q3. What is the earliest date in this dataset?

A: 2021-01-05

Q4. What is the latest date in this dataset?

A: 2022-11-22

skimr::skim(vax)

Table 1: Data summary

Name vax

Table 1: Data summary

174636
18
5
13
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
local_health_jurisdiction	0		1	0	15	495	62	0
county	0		1	0	15	495	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_missim	mplete	nna ben	sd	p0	p25	p50	p75	p100	hist
zip_code_tabulation_a	rea 0	1.00	93665.	11817.3	399000	192257	.7933658	.5905380	.5997635	.0
vaccine_equity_metric_	_&64 8tile	0.95	2.44	1.11	1	1.00	2.00	3.00	4.0	
age12_plus_population	. 0	1.00	18895.	.0148993	.880	1346.9	9513685	. 1301756	.128556	.7
$age5_plus_population$	0	1.00	20875.	.2241105	.980	1460.5	5015364	.0304877	.0100190	2.0
$tot_population$	8514	0.95	23372.	72/2628	.5112	2126.0	0018714	.038168	.001116	5.0
persons_fully_vaccinate	e d 14921	0.91	13466.	3144722	.461	883.00	8024.0	022529	.007186	.0
persons_partially_vacc	in 4921	0.91	1707.5	01998.8	30 11	167.00	1194.0	02547.0	039204	.0
percent_of_population	_ 1f8666 5_vac	c On&9 ec	10.55	0.25	0	0.39	0.59	0.73	1.0	
percent_of_population	_1p8:6:6:5 ally_	_ 0 a& 9 in	1a 0e01 8	0.09	0	0.05	0.06	0.08	1.0	
percent_of_population	_1.956 2_1_	p 0.18 9_d	o © e61	0.25	0	0.46	0.65	0.79	1.0	
booster_recip_count	70421	0.60	5655.1	76867.4	4911	280.00	2575.0	009421.0	0058304	.0
bivalent_dose_recip_co	o d:6 6958	0.10	1646.0	22161.8	34 11	109.00	719.00	2443.0	0018109	.0
eligible_recipient_coun	t 0	1.00	12309.	.1194555	.83 0	466.00	5810.0	0021140	.0806696	.0

sum (is.na(vax\$persons_fully_vaccinated))

```
[1] 14921
```

```
sum (is.na(vax$persons_fully_vaccinated))/nrow(vax)
```

[1] 0.08544057

Q5. How many numeric columns are in this dataset?

A: 13

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

A: 14921

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

A: 0.085

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

Working with dates

```
library(lubridate)

Warning: package 'lubridate' was built under R version 4.2.2

Loading required package: timechange

Warning: package 'timechange' was built under R version 4.2.2

Attaching package: 'lubridate'

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

date, intersect, setdiff, union
```

```
today()

[1] "2022-11-27"

# 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 691 days

# Using the last and the first date value we can now determine how many days the dataset s vax$as_of_date[nrow(vax)] - vax$as_of_date[1]

Time difference of 686 days

# Finding how many unique dates are in the dataset sum( !is.na( unique(vax$as_of_date) ))

[1] 99
```

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

A: 691

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

A: 99

Working with ZIP codes

```
library(terra)
```

library(zipcodeR)

```
Warning: package 'zipcodeR' was built under R version 4.2.2
  geocode_zip('92037')
# A tibble: 1 x 3
 zipcode
          lat lng
        <dbl> <dbl>
 <chr>
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
     92037
               92109
                         2.33
  # More usefully, we can pull census data about ZIP code areas (including median household
  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>
                                        <blob> <chr> <chr> <dbl> <dbl> <chr>
 <chr>
         Standard La Jol~ La Jol~ <raw 20 B> San D~ CA
                                                              32.8 -117. Pacific
1 92037
         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, ...
```

Pull data for all ZIP codes in the dataset

#zipdata <- reverse_zipcode(vax\$zip_code_tabulation_area)</pre>

Focus on the San Diego area

```
library(dplyr)
Attaching package: 'dplyr'
The following objects are masked from 'package:terra':
    intersect, union
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")</pre>
  nrow(sd)
[1] 10593
  sum( !is.na( unique(sd$zip_code_tabulation_area) ))
[1] 107
  sd.10 <- filter(vax, county == "San Diego" &</pre>
                   age5_plus_population > 10000)
  sd$zip_code_tabulation_area[which.max(sd$age12_plus_population)]
[1] 92154
```

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

A: 107

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

A: 92154

Using dplyr select all San Diego "county" entries on "as_of_date" "2022-11-15" and use this for the following questions.

[1] 0.752562

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

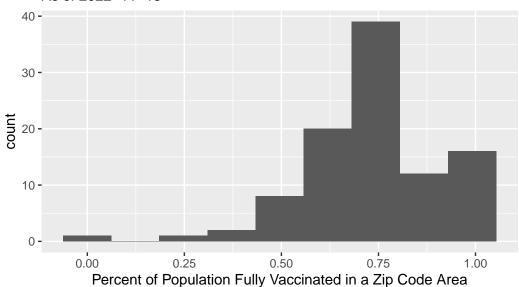
A: persons_fully_vaccinated / tot_population = 0.752562

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"?

A: see plot below

Warning: Removed 8 rows containing non-finite values (stat_bin).

Histogram of Vaccination Rates Accross San Diego County As of 2022–11–15



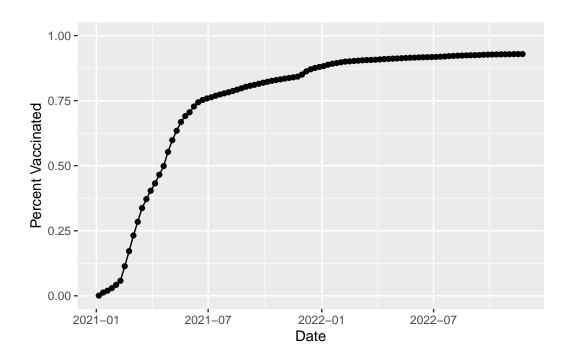
Focuse on UCSD/La Jolla

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

[1] 36144

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

```
labs(x="Date", y="Percent Vaccinated")
p2
```

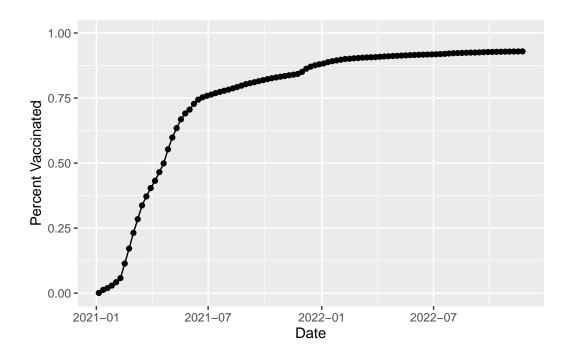


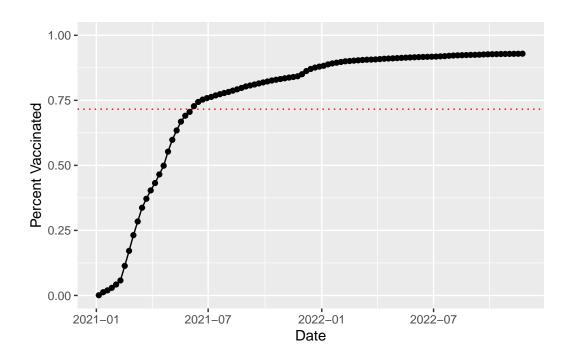
Comparing to similar sized areas

[1] 0.7159711

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?

p2





skimr::skim(vax.36)

Table 4: Data summary

Name	vax.36
Number of rows	411
Number of columns	18
Column type frequency:	
character	4
Date	1
numeric	13
Group variables	None

Variable type: character

skim_variable	n_missing	complete_	_rate	min	max	empty	n_unique	whitespace
local_health_jurisdiction	0		1	4	15	0	37	0
county	0		1	4	15	0	36	0
vem source	0		1	26	26	0	1	0

skim_variable	n_missing	$complete_{-}$	_rate	min	max	empty	n_unique	whitespace
redacted	0		1	2	2	0	1	0

Variable type: Date

skim_variable	n_missing	complete_rate	min	max	median	n_unique
as_of_date	0	1	2022-11-15	2022-11-15	2022-11-15	1

Variable type: numeric

skim_variable n_missingmpl	ete <u>m</u> reate	sd	p0	p25	p50	p75	p100	hist
zip_code_tabulation_area 0	1 92862	.110716.	6090001	.0901761	.5902646	.0904517	.0906003	.00
vaccine_equity_metric_quartile	1 2.35	1.11	1.00	1.00	2.00	3.00	4.00	
age12_plus_population 0	1 46847.	.402057	7.3321650	.937693	.5453985	.4503931	.5838556	.70
age5_plus_population 0	1 52012.	.3123620	0.1336181	.001612	.548573	.0009167	.5100190	2.00
tot_population 0	1 55640.	.9114745	5. 13 8007	.0201393	.002212	.0622910	.0101116	5.00
persons_fully_vaccinated 0	1 39837.	.281739	0.8107422	.031926	.5307064	.0405033	.5807151	.00
persons_partially_vaccinated	1 4077.7	2 02620.	741733.0	002813.0	03542.0	04666.0	039160	.00
percent_of_population_fully_vaccin	ated).72	0.11	0.38	0.64	0.72	0.79	1.00	
percent_of_population_partially_va	tcin@t@7d	0.05	0.04	0.06	0.06	0.08	0.98	
percent_of_population_with_1_plus	\$ _d 0 s ₹ 9	0.11	0.44	0.71	0.79	0.86	1.00	
booster_recip_count 0	1 22817.	.37812.	128603.0	017134	.5201640	.0207265	.556744	.00
bivalent_dose_recip_count0	1 5618.6	52952.	701375.0	03418.5	504941.0	07269.5	5016829	.00
eligible_recipient_count 0	1 39609.	.3111653	3.387321	.031819	.5336758	.004903	.586696	.00

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"?

A: Mean: 7.172851e-01

Min: 3.78501e-01 1st Q: 6.396185e-01

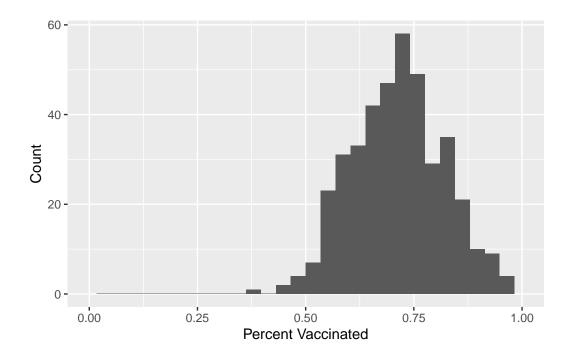
Median: 7.15524e-01 3rd Q: 7.879820e-01 Max: 1.00000e+00

Q18. Using ggplot generate a histogram of this data.

A: see plot below

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

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



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

```
vax %>% filter(as_of_date == "2022-11-15") %>%
    filter(zip_code_tabulation_area=="92109") %>%
    select(percent_of_population_fully_vaccinated)

percent_of_population_fully_vaccinated
1 0.693299
```

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

A: both are below the average value previously calculated

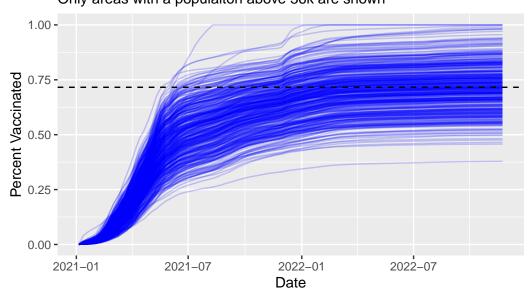
Q20. Finally make a time course plot of vaccination progress for all areas in the full dataset with a $age5_plus_population > 36144$.

```
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 a populaiton above 36k are shown') +
   geom_hline(yintercept=0.7159711 , linetype='dashed')
```

Warning: Removed 184 row(s) containing missing values (geom_path).

Vaccination rate across California Only areas with a populaiton above 36k are shown



Q21. How do you feel about traveling for Thanksgiving Break and meeting for in-person class afterwards? A: I prefer Online class for next week so I can be more flexible with my schedule