

lab17

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Import

```
url = "covid19vaccinesbyzipcode_test.csv" # data fulled from 2023-02-28

vax <- read.csv(url)
tail(vax[order(vax$as_of_date),])
```

	as_of_date	zip_code	tabulation_area	local_health_jurisdiction	
197563	2023-02-21		95076	Santa Cruz	
197564	2023-02-21		93280	Kern	
197565	2023-02-21		93301	Kern	
197566	2023-02-21		95046	Santa Clara	
197567	2023-02-21		93437	Santa Barbara	
197568	2023-02-21		95010	Santa Cruz	
	county	vaccine_equity_metric_quartile		vem_source	
197563	Santa Cruz		2	Healthy Places Index Score	
197564	Kern		1	Healthy Places Index Score	
197565	Kern		1	Healthy Places Index Score	
197566	Santa Clara		4	Healthy Places Index Score	
197567	Santa Barbara		NA	No VEM Assigned	
197568	Santa Cruz		3	Healthy Places Index Score	
	age12_plus_population	age5_plus_population	tot_population		
197563	70940.9	80312	86905		
197564	23052.1	26032	28138		
197565	9731.2	11151	12325		
197566	5087.5	5696	5918		
197567	2494.5	2871	3387		
197568	7602.0	8436	8856		
	persons_fully_vaccinated	persons_partially_vaccinated			
197563	68622	6516			

197564	13439	1829	
197565	7578	1148	
197566	5291	336	
197567	457	988	
197568	7752	695	
percent_of_population_fully_vaccinated			
197563	0.789621		
197564	0.477610		
197565	0.614848		
197566	0.894052		
197567	0.134928		
197568	0.875339		
percent_of_population_partially_vaccinated			
197563	0.074978		
197564	0.065001		
197565	0.093144		
197566	0.056776		
197567	0.291704		
197568	0.078478		
percent_of_population_with_1_plus_dose booster_recip_count			
197563	0.864599	40293	
197564	0.542611	5768	
197565	0.707992	3714	
197566	0.950828	3391	
197567	0.426632	103	
197568	0.953817	5600	
bivalent_dose_recip_count eligible_recipient_count redacted			
197563	15013	68362	No
197564	1132	13418	No
197565	1168	7563	No
197566	1267	5276	No
197567	31	456	No
197568	2831	7741	No

Q1 persons_fully_vaccinated

Q2 zip_code_tabulation_area

Q3 2021-01-05

Q4 2023-02-21 (this report is done on 2023-02-28, the data will update throughout time)

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

Table 1: Data summary

Name	vax
Number of rows	197568
Number of columns	18
Column type frequency:	
character	5
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	112	0
local_health_jurisdiction	0	1	0	15	560	62	0
county	0	1	0	15	560	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	1817.38	9000	192257.75	3658.50	5380.50	7635.0	
vaccine_equity_metric_quantile	0	0.95	2.44	1.11	1	1.00	2.00	3.00	4.0	
age12_plus_population	0	1.00	18895.04	8993.87	0	1346.95	13685.13	1756.18	8556.7	
age5_plus_population	0	1.00	20875.21	1105.97	0	1460.50	15364.03	14877.00	1902.0	
tot_population	9632	0.95	23372.72	2628.51	12	2126.00	18714.08	168.00	11165.0	
persons_fully_vaccinated	16424	0.92	13933.28	5034.25	11	927.75	8548.00	23255.08	7533.0	
persons_partially_vaccinated	16424	0.92	1701.03	2026.98	11	165.00	1196.00	2534.00	39832.0	
percent_of_population_2008_vaccinated	20688	0.90	0.57	0.25	0	0.42	0.60	0.74	1.0	
percent_of_population_2008_partially_vaccinated	20688	0.90	0.08	0.09	0	0.05	0.06	0.08	1.0	
percent_of_population_2012_1_plus_dose	21712	0.89	0.63	0.24	0	0.48	0.67	0.81	1.0	
booster_recip_count	72754	0.63	5791.60	7111.11	11	295.00	2724.00	372.75	9493.0	
bivalent_dose_recip_count	158541	0.20	2867.36	3529.71	11	186.00	1374.00	4513.50	27175.0	
eligible_recipient_count	0	1.00	12772.28	4887.79	0	501.00	6309.50	21926.58	7215.0	

Q5 13

Q6 16424

```
1 - 0.9168691
```

```
[1] 0.0831309
```

Q7 8.31%

Q8 Some population might not be fully registered in the medical system.

Working with dates

```
library(lubridate)
```

Attaching package: 'lubridate'

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

date, intersect, setdiff, union

```
today()
```

```
[1] "2023-03-07"
```

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

How many days have passed since the first vaccination reported in this dataset?

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

Time difference of 791 days

How many days the dataset span?

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

Time difference of 777 days

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

```
today() - tail(vax$as_of_date,1)
```

Time difference of 14 days

Q9 7

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

[1] 112

Q10 112

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.
```

```
# distance between two zip code
zip_distance('92037','92109')
```

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

```
# get information from zip code
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, ...
```

```
sd <- vax[vax$county=="San Diego", ]
length(unique(sd$zip_code_tabulation_area))
```

```
[1] 107
```

```
Q11. 107
```

```
sd[which.max(sd$age12_plus_population),"zip_code_tabulation_area"]
```

```
[1] 92154
```

```
Q12 92154
```

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

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

```
[1] 0.7380452
```

```
Q13 73.8%
```

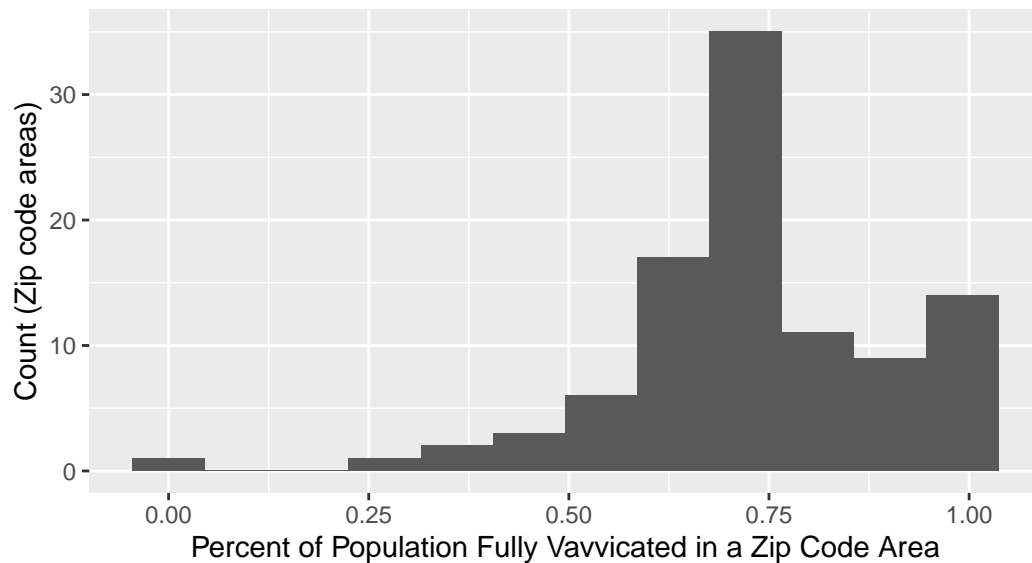
```
Q14
```

```
library(ggplot2)

sd.tmp <- filter(vax, county == "San Diego" &
                 as_of_date == "2022-11-15")
ggplot(sd.tmp, aes(percent_of_population_fully_vaccinated)) +
  geom_histogram(bins=12) +
  labs(x="Percent of Population Fully Vavvicated in a Zip Code Area",
       y="Count (Zip code areas)",
       title="Histogram of Vavvination Rates Accross San Diego County",
       subtitle = "As of 2022-11-15")
```

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

Histogram of Vavvination Rates Accross San Diego County
As of 2022-11-15



focus on UCSD/La Jolla

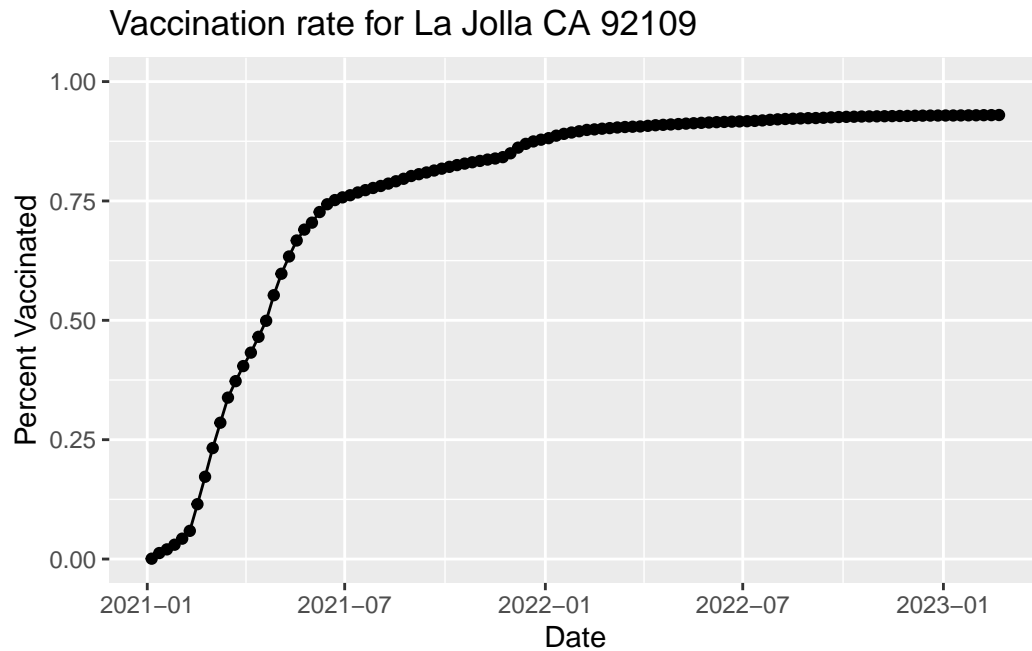
```
ucsd <- filter(sd, zip_code_tabulation_area=="92037")
ucsd[1,]$age5_plus_population
```

[1] 36144

Q15.

```
ucsd.graph <- 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")

ucsd.graph
```

Similar size area

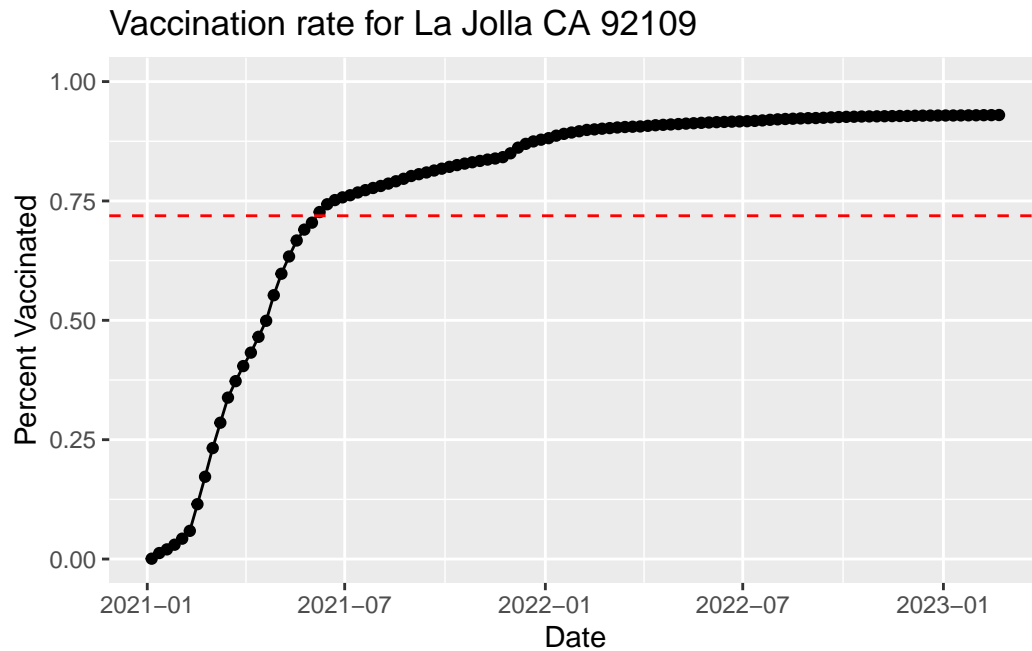
```
# Subset to all CA areas with a population as large as 92037
vax.36 <- filter(vax, age5_plus_population > 36144 &
  as_of_date == "2022-11-15")
```

Q16.

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

```
[1] 0.7190084
```

```
ucsd.graph +
  geom_hline(yintercept=vax.mean, color="red", linetype=2)
```



Q17

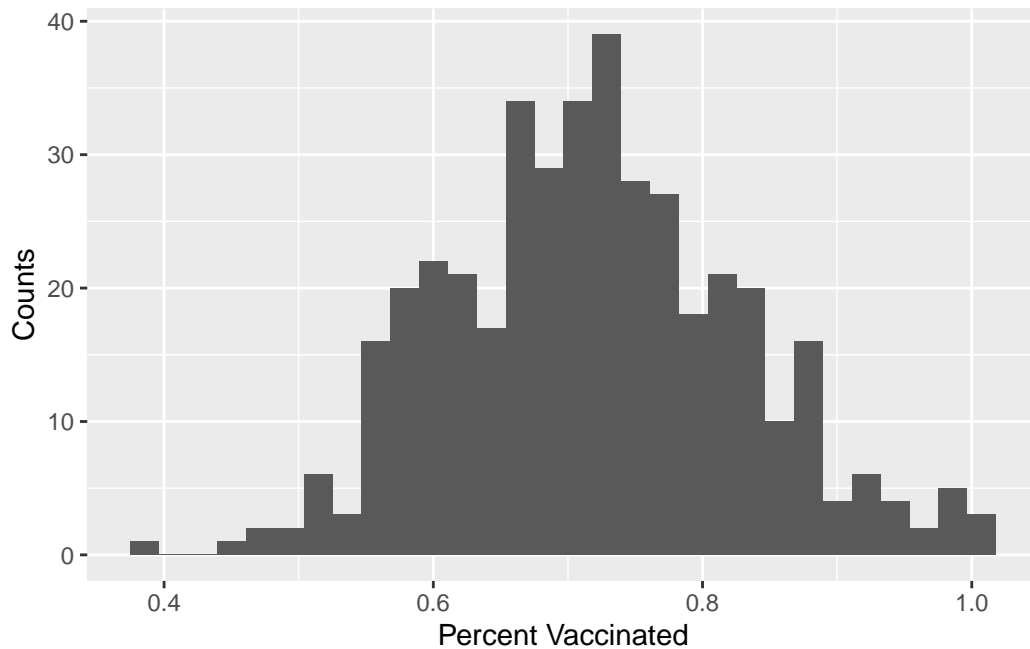
```
summary(vax.36$percent_of_population_fully_vaccinated)
```

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
0.3784	0.6444	0.7163	0.7190	0.7883	1.0000

Q18

```
ggplot(vax.36)+
  geom_histogram(aes(percent_of_population_fully_vaccinated)) +
  labs(x="Percent Vaccinated", y="Counts")
```

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



Q19

```
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                                0.548979
```

0.549 < 0.719

The area 92040 is below the average.

```
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                                0.692683
```

0.693 < 0.719

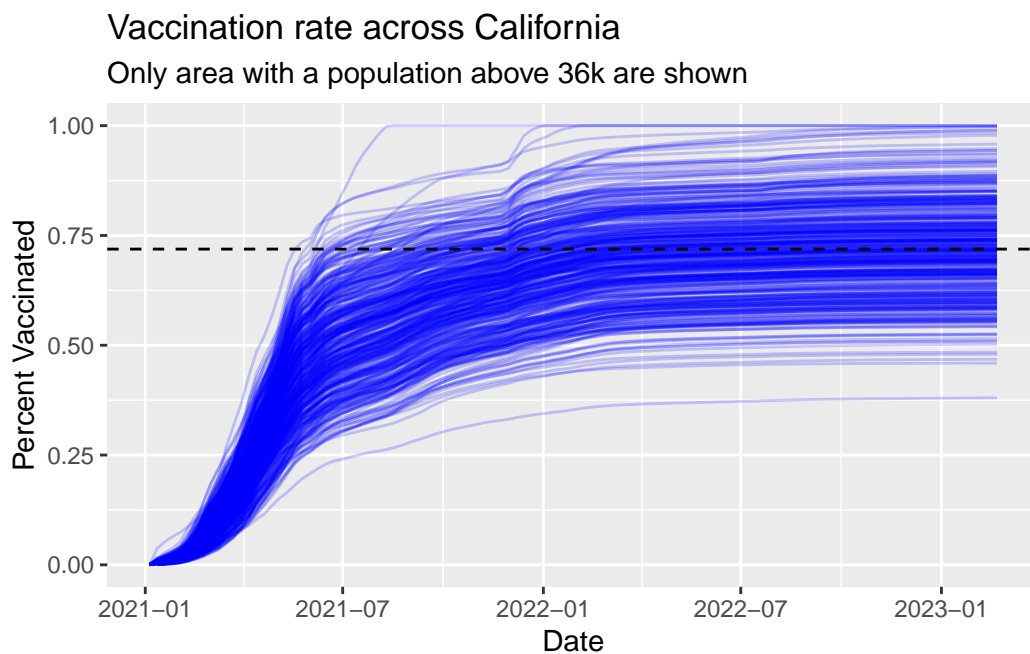
The area 92109 is below the average.

Q20

```
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(0.00, 1.00) +
  labs(x="Date", y="Percent Vaccinated",
       title="Vaccination rate across California",
       subtitle="Only area with a population above 36k are shown") +
  geom_hline(yintercept = vax.mean, linetype=2)
```

Warning: Removed 183 rows containing missing values (`geom_line()`).



About this document

```
sessionInfo()
```

```
R version 4.2.2 (2022-10-31)
Platform: aarch64-apple-darwin20 (64-bit)
Running under: macOS Monterey 12.5
```

```
Matrix products: default
```

```
BLAS:   /Library/Frameworks/R.framework/Versions/4.2-arm64/Resources/lib/libRblas.0.dylib
```

```
LAPACK: /Library/Frameworks/R.framework/Versions/4.2-arm64/Resources/lib/libRlapack.dylib
```

```
locale:
```

```
[1] en_US.UTF-8/en_US.UTF-8/en_US.UTF-8/C/en_US.UTF-8/en_US.UTF-8
```

```
attached base packages:
```

```
[1] stats      graphics  grDevices  utils      datasets  methods   base
```

```
other attached packages:
```

```
[1] ggplot2_3.4.1  dplyr_1.1.0    zipcodeR_0.3.5  lubridate_1.9.2
```

```
loaded via a namespace (and not attached):
```

```
[1] Rcpp_1.0.10      lattice_0.20-45  tidyr_1.3.0      class_7.3-21
[5] digest_0.6.31    utf8_1.2.3       R6_2.5.1         repr_1.1.6
[9] RSQLite_2.3.0    evaluate_0.20    e1071_1.7-13     httr_1.4.4
[13] pillar_1.8.1     rlang_1.0.6      curl_5.0.0       uuid_1.1-0
[17] rstudioapi_0.14  raster_3.6-14    blob_1.2.3       rmarkdown_2.20
[21] labeling_0.4.2   readr_2.1.4      stringr_1.5.0    munsell_0.5.0
[25] bit_4.0.5        proxy_0.4-27     compiler_4.2.2   xfun_0.37
[29] pkgconfig_2.0.3  tigris_2.0.1     base64enc_0.1-3  htmltools_0.5.4
[33] tidyselect_1.2.0 tibble_3.1.8     codetools_0.2-19 fansi_1.0.4
[37] crayon_1.5.2     tzdb_0.3.0       withr_2.5.0      sf_1.0-9
[41] tidycensus_1.3.2 rappdirs_0.3.3   grid_4.2.2       gtable_0.3.1
[45] jsonlite_1.8.4   lifecycle_1.0.3 DBI_1.1.3         magrittr_2.0.3
[49] scales_1.2.1     units_0.8-1      KernSmooth_2.23-20 cli_3.6.0
[53] stringi_1.7.12   cachem_1.0.6     farver_2.1.1     sp_1.6-0
[57] skimr_2.1.5      xml2_1.3.3       ellipsis_0.3.2   generics_0.1.3
[61] vctrs_0.5.2      tools_4.2.2      bit64_4.0.5      glue_1.6.2
[65] purrr_1.0.1      hms_1.1.2        fastmap_1.1.0    yaml_2.3.7
[69] colorspace_2.1-0 timechange_0.2.0 terra_1.7-3      classInt_0.4-9
[73] rvest_1.0.3      memoise_2.0.1    knitr_1.42
```