# First Year Exam: Bioinformatics Portion

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## **Packages**

Load all of the necessary packages

```
library(dplyr)
## Warning: package 'dplyr' was built under R version 4.1.3
## 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
##
library(ggplot2)
## Warning: package 'ggplot2' was built under R version 4.1.3
library(lubridate)
## Warning: package 'lubridate' was built under R version 4.1.3
## Attaching package: 'lubridate'
## The following objects are masked from 'package:base':
##
       date, intersect, setdiff, union
##
```

### Data import

Read in the data from the .csv as a tibble

Look at the structure of the data

head(data)

```
## # A tibble: 6 x 8
## date area area_type variant_name specimens percentage specimens_7d_avg
## <chr> <chr> <chr> <chr> <dbl> <dbl> <dbl>
```

```
## 1 2021-01-01 Calif~ State
                                 Alpha
                                                               1.69
                                                                                   NA
                                                       1
## 2 2021-01-01 Calif~ State
                                 Beta
                                                       0
                                                                                   NΑ
                                                               0
## 3 2021-01-01 Calif~ State
                                 Mu
                                                       0
                                                               0
                                                                                   NA
## 4 2021-01-01 Calif~ State
                                                       0
                                                               0
                                                                                   NΑ
                                 Gamma
## 5 2021-01-01 Calif~ State
                                 Total
                                                      59
                                                             100
                                                                                   NA
## 6 2021-01-01 Calif~ State
                                 Omicron
                                                                                   NA
                                                       1
                                                               1.69
## # ... with 1 more variable: percentage 7d avg <dbl>
```

## Data processing

Make dates more workable

```
data <- data %>%
  mutate(date = ymd(date)) # Change the date column from a <chr> to a <date>
```

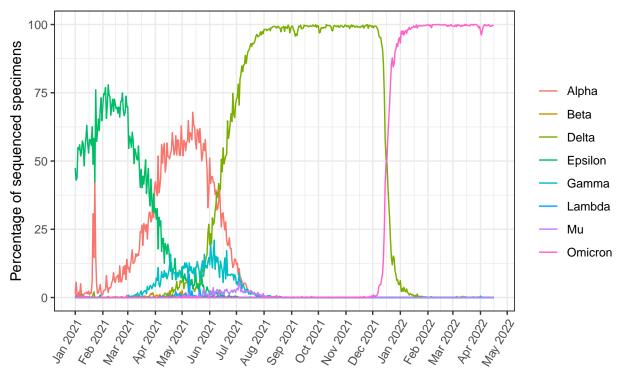
Subset the data to match the example

```
data <- data %>%
  filter(date < as.Date("2022-04-17")) %>% # drop late dates
  filter(variant_name != "Other", variant_name != "Total") # drop "other" & "total"
```

#### Data visulization

Plot the data

## Covid-19 Variants in California



Data Source: <a href="https://www.cdph.ca.gov">https://www.cdph.ca.gov</a>

#### Session information

#### sessionInfo()

```
## R version 4.1.2 (2021-11-01)
## Platform: x86_64-w64-mingw32/x64 (64-bit)
## Running under: Windows 10 x64 (build 19044)
##
## Matrix products: default
##
## locale:
## [1] LC_COLLATE=English_United States.1252
## [2] LC_CTYPE=English_United States.1252
## [3] LC_MONETARY=English_United States.1252
## [4] LC_NUMERIC=C
## [5] LC_TIME=English_United States.1252
##
## attached base packages:
## [1] stats
                 graphics grDevices utils
                                               datasets methods
                                                                   base
## other attached packages:
## [1] lubridate_1.8.0 ggplot2_3.3.6
                                       dplyr_1.0.9
##
## loaded via a namespace (and not attached):
  [1] highr_0.9
                                          compiler_4.1.2
                        pillar_1.7.0
                                                           tools_4.1.2
   [5] digest_0.6.29
                         evaluate_0.15
                                          lifecycle_1.0.1 tibble_3.1.7
```

```
## [9] gtable_0.3.0
                        pkgconfig_2.0.3 rlang_1.0.2
                                                          cli_3.3.0
## [13] rstudioapi_0.13
                        yaml_2.3.5
                                         xfun_0.31
                                                          fastmap_1.1.0
## [17] withr_2.5.0
                                                          generics_0.1.2
                                         knitr_1.39
                        stringr_1.4.0
## [21] vctrs_0.4.1
                        grid_4.1.2
                                         tidyselect_1.1.2 glue_1.6.2
## [25] R6_2.5.1
                        fansi_1.0.3
                                         rmarkdown_2.14
                                                          purrr_0.3.4
## [29] farver_2.1.0
                        magrittr_2.0.3
                                         scales_1.2.0
                                                          ellipsis_0.3.2
## [33] htmltools_0.5.2
                        colorspace_2.0-3 labeling_0.4.2
                                                          utf8_1.2.2
## [37] stringi_1.7.6
                        munsell_0.5.0
                                         crayon_1.5.1
```