Q15 - First Year Exam

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Load the data and packages

```
df <- read.csv("covid19_variants.csv")</pre>
library(ggplot2)
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
library(lubridate)
##
## Attaching package: 'lubridate'
## The following objects are masked from 'package:base':
##
       date, intersect, setdiff, union
Look at data
head(df)
##
         date
                    area area_type variant_name specimens percentage
## 1 1/1/2021 California
                              State
                                            Alpha
                                                          1
                                                                   1.69
## 2 1/1/2021 California
                                            Beta
                                                          0
                                                                   0.00
                              State
## 3 1/1/2021 California
                              State
                                               Mu
                                                          0
                                                                   0.00
## 4 1/1/2021 California
                                                          0
                                                                  0.00
                              State
                                           Gamma
## 5 1/1/2021 California
                              State
                                            Total
                                                         59
                                                                 100.00
## 6 1/1/2021 California
                                                                  1.69
                              State
                                         Omicron
                                                          1
     specimens_7d_avg percentage_7d_avg
## 1
                   NA
                                      NA
```

Remove 'Total' and 'Other' variants since these aren't in the plot given to us

```
df <- df[!df$variant_name %in% c("Other", "Total"), ]
head(df)</pre>
```

```
area area_type variant_name specimens percentage
         date
## 1 1/1/2021 California
                              State
                                            Alpha
                                                           1
                                                                   1.69
## 2 1/1/2021 California
                              State
                                             Beta
                                                           0
                                                                   0.00
## 3 1/1/2021 California
                              State
                                               Mu
                                                           0
                                                                   0.00
## 4 1/1/2021 California
                              State
                                            Gamma
                                                           0
                                                                   0.00
## 6 1/1/2021 California
                              State
                                          Omicron
                                                          1
                                                                   1.69
## 7 1/1/2021 California
                              State
                                          Epsilon
                                                          28
                                                                  47.46
     specimens_7d_avg percentage_7d_avg
## 1
                    NA
                                       NA
## 2
                    NA
                                       NA
## 3
                    NA
                                       NA
## 4
                    NA
                                       NA
## 6
                    NA
                                       NA
## 7
                    NA
                                       NA
```

Change the date to a 'date' variable rather than a 'character' variable using the 'lubridate' package

```
# observe column variable in wrong format class(df$date)
```

[1] "character"

```
df$date[1]
```

[1] "1/1/2021"

```
# fix above, make new column and add to df
# note some will fail to parse as csv doesn't have dates for all rows

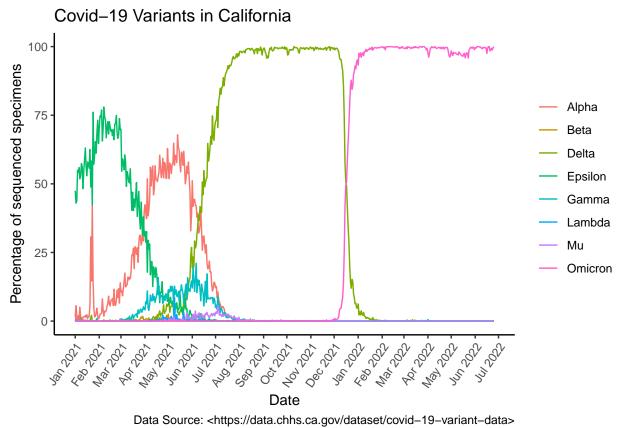
df$new_date <- mdy(df$date)
head(df)</pre>
```

```
##
         date
                    area area_type variant_name specimens percentage
## 1 1/1/2021 California
                             State
                                           Alpha
                                                                  1.69
                                                         1
## 2 1/1/2021 California
                             State
                                            Beta
                                                         0
                                                                  0.00
                                                                  0.00
## 3 1/1/2021 California
                             State
                                              Mu
                                                         0
## 4 1/1/2021 California
                             State
                                           Gamma
                                                         0
                                                                  0.00
## 6 1/1/2021 California
                             State
                                         Omicron
                                                         1
                                                                  1.69
## 7 1/1/2021 California
                                         Epsilon
                                                        28
                                                                 47.46
                             State
     specimens_7d_avg percentage_7d_avg new_date
```

```
## 1
                                     NA 2021-01-01
                   NA
## 2
                                     NA 2021-01-01
                   NA
## 3
                   NA
                                     NA 2021-01-01
## 4
                   NA
                                     NA 2021-01-01
                                     NA 2021-01-01
## 6
                   NA
## 7
                   NA
                                     NA 2021-01-01
```

Plot:

```
ggplot(df, aes(new_date,percentage)) +
# group by variant (hence excluding total and other)
 geom_line(aes(color = variant_name)) +
# set theme
 theme_classic() +
# now, format axes
 scale_x_date(date_labels = "%b %Y", date_breaks = "1 month") +
# add axes titles
 ylab("Percentage of sequenced specimens") +
 xlab("Date") +
# make x axis readable (rotate), remove legend title
 theme(axis.text.x = element_text(angle = 55,
                                   hjust = 1),
        legend.position = "right",
       legend.title = element_blank()) +
# title plot
 ggtitle("Covid-19 Variants in California") +
# add caption to include website where data was found
 labs(caption = "Data Source: <https://data.chhs.ca.gov/dataset/covid-19-variant-data>")
```



Data Source: https://data.chhs.ca.gov/dataset/covid-19-variant-data