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Covid-19 Variants in California

Libraries

Read file

```
# Calling the libraries that will be used to generate the plot
library(ggplot2) # For plots
library(lubridate) # For date-time data
##
## Attaching package: 'lubridate'
## The following objects are masked from 'package:base':
##
##
       date, intersect, setdiff, union
library(dplyr) # For data manipulation
## Warning: package 'dplyr' was built under R version 4.1.2
## 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
```

```
# To read the .csv file downloaded from the California Health and Human Services (CHHS) open data site data <- read.csv("covid19_variants.csv")
```

Filter data

I'll use 'dplyr' to filter out unnecessary rows (containing the values 'Other' and 'Total' in the 'variant_name' column) and change the format of the date values in the 'date' column to date instead of character.

```
# To remove the rows that contain "variant_name" = "Other", "Total" (as they're not included in the plo
# To change the date values from "character" to "date" format
clean.data <- data %>% filter(!variant_name %in% c('Other', 'Total')) %>% mutate(date = as.Date(date))
```

Assigning variables

We use this to simplify things when writing code, substituting longer names to simpler and shorter words.

```
# Assigning variables
Percentage <- clean.data$percentage
Date <- clean.data$date
Variant <- clean.data$variant_name</pre>
```

Plotting the data

I'll use 'ggplot2' and I'll first specify the data frame and the variables that we're interested in, coloring each Covid-19 variant with a different color (each variant will be a line in the graph). Then, I'll select the type of graph I would like to represent the data in, in this case I'll use a line chart. I'll modify the theme of the plot so that it resembles the example and I'll modify the dates with 'lubricate'. In addition, I'll add labels to my plot so that it's easily understandable and a caption at the bottom.

```
# Making the plot

plot <- ggplot(data = clean.data, aes(x=Date, y=Percentage, color=Variant)) +

# Select the data frame, x and y axis variables, and the variable by which the data will be colored b

geom_line(aes(group=Variant)) +

# To make a line chart, where every line will be a category of the "Variant" column, meaning a differ

theme_bw() +

# Specifies the theme used for the plot: in this case a white background and thin grey grid lines

scale_x_date(date_breaks = "1 month", date_labels = "%b %Y") +

# Specifies how the dates in the x-axis will be represented, separated by 1 month gaps, and with the

theme(axis.text.x=element_text(angle=60, hjust=1)) +

# Specifies that the label for the dates in the x-axis is angled

labs(title = "Covid-19 Variants in California", x = NULL, color = NULL, caption = "Data Source: <http

# To label the title of the plot, remove the x-axis label to only keep the dates, remove the label of

ylab("Percentage of sequenced specimens") # To add a y-axis label

plot
```

