

1st_year_final_CBrennan

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Importing data and saving as a variable:

```
covid_data <- read.csv("covid-19-variant-data.csv")
head(covid_data)
```

```
##      date      area area_type variant_name specimens percentage
## 1 2021-01-01 California      State      Delta          0         0.00
## 2 2021-01-01 California      State      Alpha          1         1.69
## 3 2021-01-01 California      State      Other         29        49.15
## 4 2021-01-01 California      State      Total         59       100.00
## 5 2021-01-01 California      State      Beta          0         0.00
## 6 2021-01-01 California      State      Omicron         1         1.69
##  specimens_7d_avg percentage_7d_avg
## 1                NA                NA
## 2                NA                NA
## 3                NA                NA
## 4                NA                NA
## 5                NA                NA
## 6                NA                NA
```

Loading packages required to make plot:

```
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
```

```
library(scales)  
library(gridExtra)
```

```
##  
## Attaching package: 'gridExtra'
```

```
## The following object is masked from 'package:dplyr':  
##  
##    combine
```

```
library(ggthemes)
```

Filtering out unwanted data:

```
covid_data4 <- filter(covid_data, variant_name!="Total" & variant_name!="Other")
```

Converting “date” column to date format:

```
covid_data4$date <- as.Date(covid_data4$date)
```

Making plot which uses one tick per month on x-axis. X-axis format is in “month-year”.

```
ggplot(covid_data4, aes(x = date, y=percentage, group=variant_name)) +  
  geom_line(aes(color=variant_name)) +  
  theme(axis.text.x = element_text(angle = 60, vjust = 0.5)) +  
  scale_x_date(labels=date_format("%h-%y"), breaks=date_breaks("1 month")) +  
  xlab("Data Source:<https://www.cdph.ca.gov/>") +  
  ylab("Percentage of Sequenced Specimens") +  
  ggtitle("Covid-19 Variants in California") +  
  theme(axis.title.x=element_text(size=8, hjust = 1))
```

Covid-19 Variants in California

