

# Covid-19 variants in California

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Importing libraries

```
library(ggplot2)
```

```
## Warning: package 'ggplot2' was built under R version 4.1.3
```

```
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(scales)
```

```
## Warning: package 'scales' was built under R version 4.1.3
```

Loading dataset

```
dataset = read.csv('covid19_variants.csv')
```

```
dataset$date = as.Date(dataset$date)
```

```
head(dataset,20)
```

```
##           date      area area_type variant_name specimens percentage
## 1 2021-01-01 California      State      Epsilon         28         47.46
## 2 2021-01-01 California      State      Other         29         49.15
## 3 2021-01-01 California      State      Alpha          1          1.69
## 4 2021-01-01 California      State      Beta           0          0.00
## 5 2021-01-01 California      State      Total         59        100.00
## 6 2021-01-01 California      State      Gamma          0          0.00
```

```
## 7 2021-01-01 California State Mu 0 0.00
## 8 2021-01-01 California State Delta 0 0.00
## 9 2021-01-01 California State Omicron 1 1.69
## 10 2021-01-01 California State Lambda 0 0.00
## 11 2021-01-02 California State Delta 1 0.93
## 12 2021-01-02 California State Omicron 1 0.93
## 13 2021-01-02 California State Epsilon 46 42.99
## 14 2021-01-02 California State Beta 0 0.00
## 15 2021-01-02 California State Mu 0 0.00
## 16 2021-01-02 California State Gamma 0 0.00
## 17 2021-01-02 California State Other 53 49.53
## 18 2021-01-02 California State Lambda 0 0.00
## 19 2021-01-02 California State Total 107 100.00
## 20 2021-01-02 California State Alpha 6 5.61
## specimens_7d_avg percentage_7d_avg
## 1 NA NA
## 2 NA NA
## 3 NA NA
## 4 NA NA
## 5 NA NA
## 6 NA NA
## 7 NA NA
## 8 NA NA
## 9 NA NA
## 10 NA NA
## 11 NA NA
## 12 NA NA
## 13 NA NA
## 14 NA NA
## 15 NA NA
## 16 NA NA
## 17 NA NA
## 18 NA NA
## 19 NA NA
## 20 NA NA
```

Selecting for variants of interest

```
dataset = dataset%>%
  filter(dataset$variant_name != 'Other' & dataset$variant_name != 'Total')
```

Plotting the data

```
ggplot(dataset, aes(x=date, y=percentage, color=variant_name)) +
  geom_line() + ggtitle("COVID-19 Variants in California") +
  xlab("Date") + ylab("Percentage of Sequenced Specimens") +
  scale_x_date(labels = date_format("%Y-%m"))
```

COVID-19 Variants in California

