# EPI Info CDC

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## Contents

Libraries Used	1
Load CDC data         Read CSV File          Clean data and calculate cumulative number of cases	2 2 2
Visualize all data	2
Filter to remove incomplete reporting Visualize	<b>7</b> 7
Libraries Used	
library(tidyverse)	
## Attaching packages	
## v ggplot2 3.3.0 v purrr 0.3.3 ## v tibble 3.0.0 v dplyr 0.8.5 ## v tidyr 1.0.2 v stringr 1.4.0 ## v readr 1.3.1 v forcats 0.5.0	
## Conflicts	
<pre>## x dplyr::filter() masks stats::filter() ## x dplyr::lag() masks stats::lag()</pre>	
library(lubridate)	
## ## Attaching package: 'lubridate'	
<pre>## The following objects are masked from 'package:dplyr': ##  intersect, setdiff, union</pre>	
<pre>## The following objects are masked from 'package:base': ##</pre>	
## date, intersect, setdiff, union	

### Load CDC data

### Read CSV File

```
cdc <- read_csv(file = "./data/CDC_data.csv")

## Warning: Missing column names filled in: 'X3' [3]

## Warning: Duplicated column names deduplicated: '25 mar data' => '25 mar

## data_1' [12]
```

### Clean data and calculate cumulative number of cases

### Visualize all data

```
epi <- ggplot(data = cdc)
epi + geom_point(aes(x = Date,
              y = `Number of new cases`))+
     geom_smooth(aes(x = Date,
              y = `Number of new cases`),
              color = "red",
              fill = "blue") +
    labs(y = "Cases",
         title = "Number of New Cases of COVID-19 Reported to the CDC")
## `geom_smooth()` using method = 'loess' and formula 'y ~ x'
epi + geom_col(aes(x=Date,
                   y= Number of new cases ),
               fill= "blue") +
     geom smooth(aes(x=Date,
                   y=`Number of new cases`),
                 color = "red") +
     labs(y = "Cases",
          title = "Number of New Cases of COVID-19 Reported to the CDC")
## `geom_smooth()` using method = 'loess' and formula 'y ~ x'
epi + geom_point(aes(x = Date,
              y = cum)+
    \# geom\_line(aes(x = Date,
              #y = `cum`)) +
     geom_smooth(aes(x = Date,
              y = cum),
              color = "red",
              fill = "blue") +
    labs(y = "Cumulative number of cases",
```

# Number of New Cases of COVID-19 Reported to the CDC

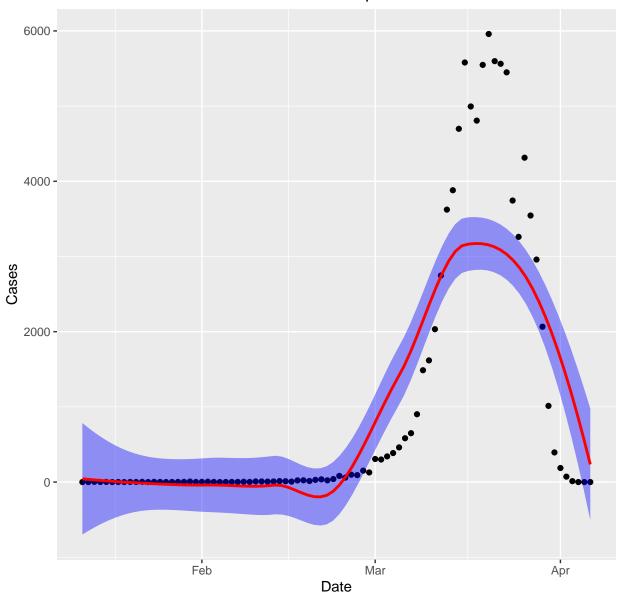


Figure 1: Epi curve 1

# Number of New Cases of COVID-19 Reported to the CDC 6000 -4000 -Cases 2000 -0 -

Figure 2: Epi curve 2, traditional

Date

Mar

Apr

Feb

##  $geom_smooth()$  using method = 'loess' and formula 'y ~ x'

## Cumulative Number of Cases of COVID-19 Reported to the CDC

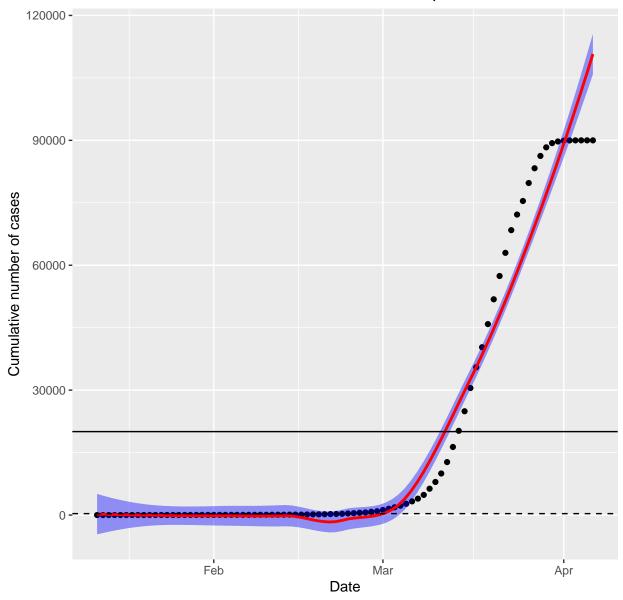


Figure 3: Cumulative cases

```
gr <- cdc %>%
  filter(`Number of new cases` != 0)

ggplot(data = gr) +
  geom_line(aes(y = `Number of new cases`,
```

##  $geom_smooth()$  using method = 'loess' and formula 'y ~ x'

## Daily Growth Rate

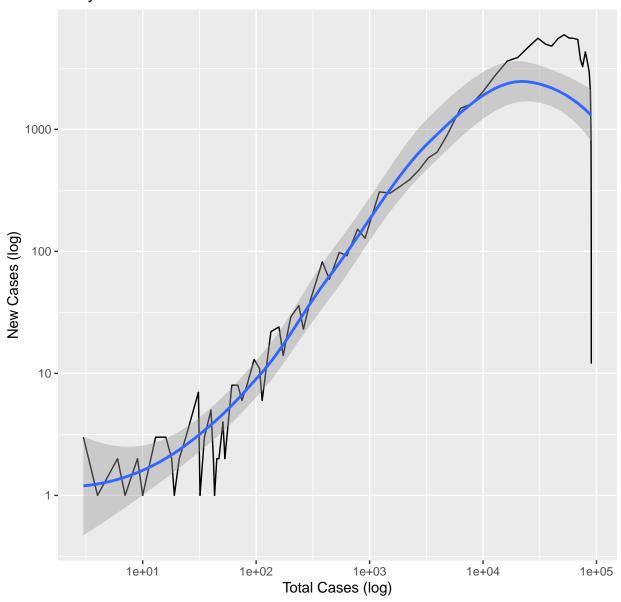


Figure 4: Growth Rate

```
gr <- cdc %>%
     group_by(`Week Number`) %>%
     summarise(cases = sum(`Number of new cases`),
               cs = sum(cum)
ggplot(data = gr) +
     geom_line(aes(y = `cases`,
                  x = cs)) +
     geom_point(aes(y = `cases`,
                    x = cs) +
     scale_x_log10() +
     scale_y_log10() +
     geom_smooth(aes(y = `cases`,
                    x = cs)) +
 labs(title = "Weekly Growth Rate",
      x = "Total Cases (log)",
      y = "New Cases (log)")
```

## `geom\_smooth()` using method = 'loess' and formula 'y ~ x'

### Filter to remove incomplete reporting

remove dates on or after 25 March as this data may not be completely reported

#### Visualize

```
epi <- ggplot(data = cdc)</pre>
epi + geom_point(aes(x = Date,
              y = `Number of new cases`))+
     geom_smooth(aes(x = Date,
              y = `Number of new cases`),
              color = "red",
              fill = "blue") +
     labs(y = "Cases",
          title = "Number of New Cases of COVID-19 Reported to the CDC")
## `geom_smooth()` using method = 'loess' and formula 'y ~ x'
epi + geom_col(aes(x=Date,
                   y=`Number of new cases`),
               fill= "blue") +
     geom_smooth(aes(x=Date,
                   y=`Number of new cases`),
                 color = "red") +
     labs(y = "Cases",
          title = "Number of New Cases of COVID-19 Reported to the CDC")
## `geom_smooth()` using method = 'loess' and formula 'y ~ x'
epi + geom_point(aes(x = Date,
                     y = cum)+
```

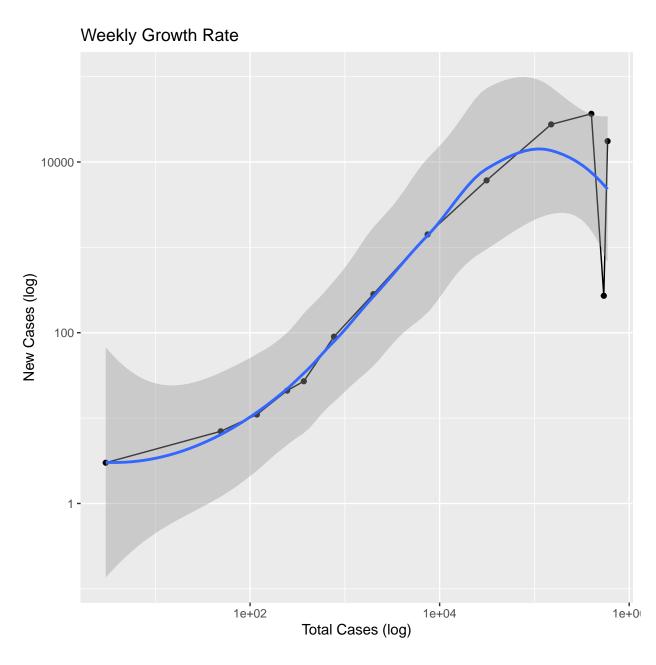


Figure 5: weekly growth rate

# Number of New Cases of COVID-19 Reported to the CDC

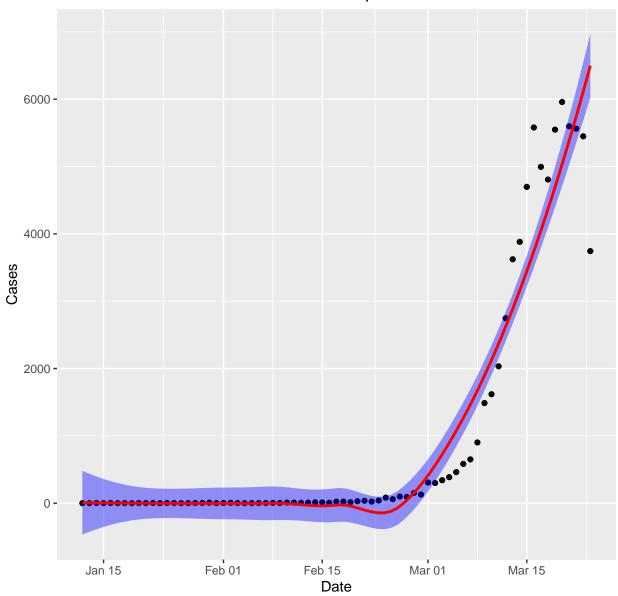


Figure 6: Epi curve 1

# Number of New Cases of COVID-19 Reported to the CDC

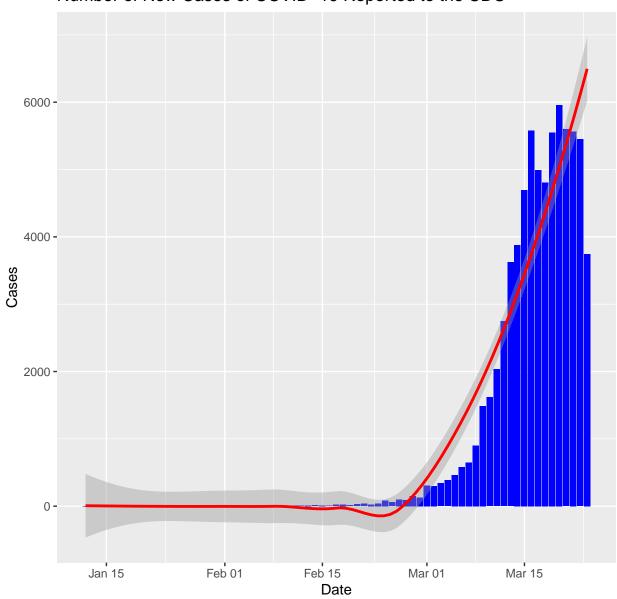


Figure 7: Epi curve 2, traditional

##  $geom_smooth()$  using method = 'loess' and formula 'y ~ x'

## Cumulative Number of Cases of COVID-19 Reported to the CDC

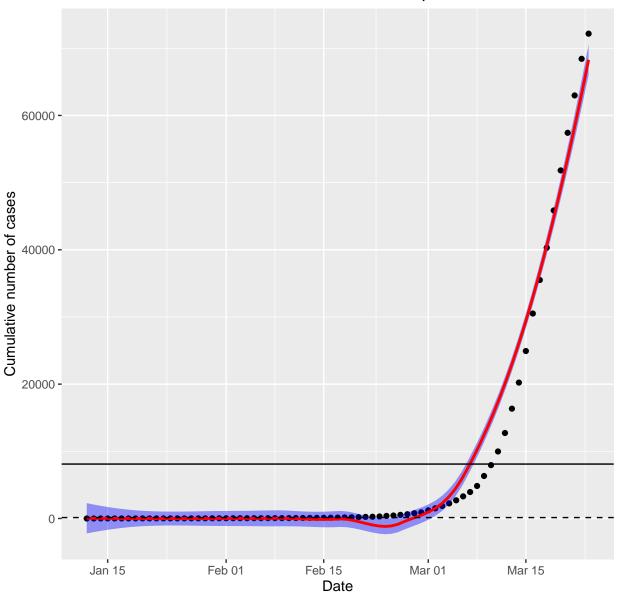


Figure 8: Cumulative cases

```
gr <- cdc %>%
 filter(`Number of new cases` != 0)
ggplot(data = gr) +
 geom_line(aes(y = `Number of new cases`,
                   x = cum) +
 scale_x_log10() +
 scale_y_log10() +
 geom_smooth(aes(y = `Number of new cases`,
             x = cum) +
 labs(title = "Daily Growth Rate",
      x = "Total Cases (log)",
      y = "New Cases (log)")
## `geom_smooth()` using method = 'loess' and formula 'y ~ x'
gr <- cdc %>%
    group_by(`Week Number`) %>%
    summarise(cases = sum(`Number of new cases`),
              cs = sum(cum)
ggplot(data = gr) +
    geom_line(aes(y = `cases`,
                  x = cs)) +
    geom_point(aes(y = `cases`,
                   x = cs)) +
    scale_x_log10() +
    scale_y_log10() +
    geom_smooth(aes(y = `cases`,
                   x = cs)
 labs(title = "Weekly Growth Rate",
      x = "Total Cases (log)",
      y = "New Cases (log)")
```

## `geom\_smooth()` using method = 'loess' and formula 'y ~ x'

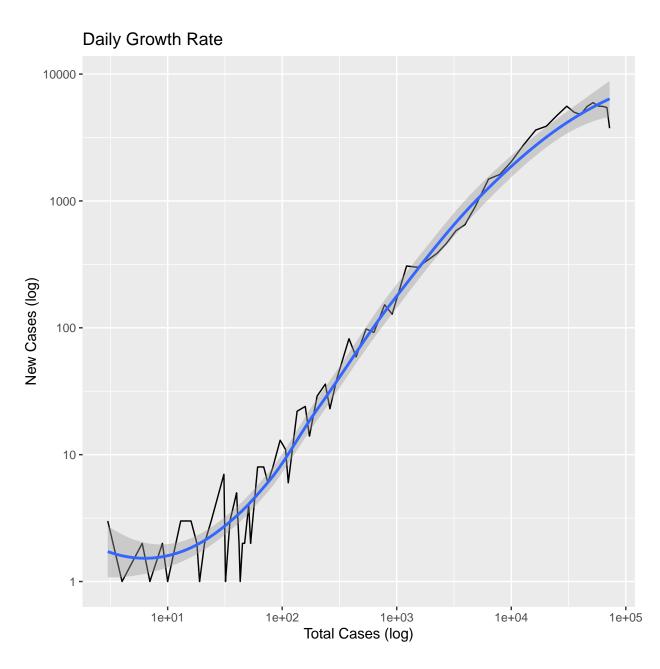


Figure 9: Growth Rate

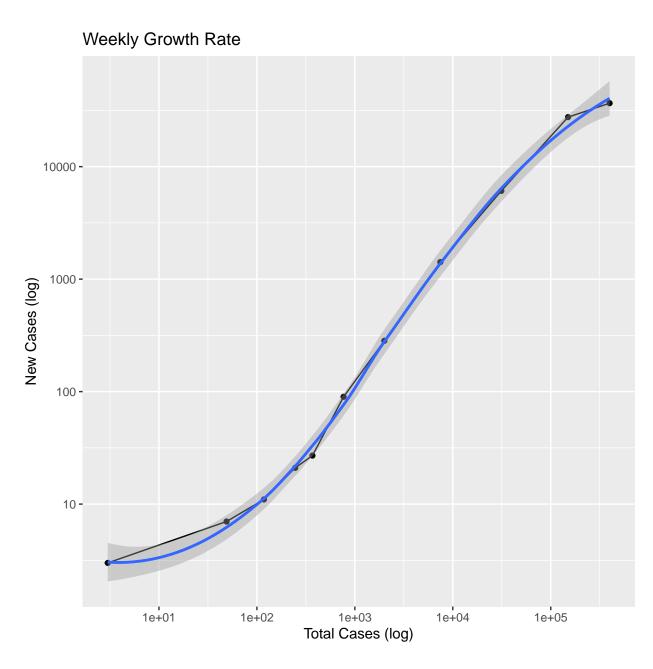


Figure 10: weekly growth rate