

EPI Info CDC

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Libraries Used

```
library(tidyverse)
```

```
## -- Attaching packages -----
## v ggplot2 3.3.0      v purrr  0.3.3
## v tibble  2.1.3      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 -----
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()     masks stats::lag()
```

```
library(lubridate)
```

```
##
## Attaching package: 'lubridate'

## The following object is masked from 'package:base':
##
##     date
```

Load CDC data

Read CSV File

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

## Parsed with column specification:
## cols(
##   Date = col_character(),
##   cases = col_double()
## )
```

Clean data and calculate cumulative number of cases

```
names(cdc)[1:2] <- c("Date",
                    "Number of new cases")
cdc$cum <- cumsum(cdc$`Number of new cases`)
cdc$Date <- as.Date(cdc$Date,
                  format = "%d-%b-%y")
```

Data

```
cdc %>%
  data.frame
```

##	Date	Number.of.new.cases	cum
## 1	2020-01-12	0	0
## 2	2020-01-13	0	0
## 3	2020-01-14	2	2
## 4	2020-01-15	0	2
## 5	2020-01-16	1	3
## 6	2020-01-17	0	3
## 7	2020-01-18	0	3
## 8	2020-01-19	0	3
## 9	2020-01-20	1	4
## 10	2020-01-21	1	5
## 11	2020-01-22	2	7
## 12	2020-01-23	0	7
## 13	2020-01-24	1	8
## 14	2020-01-25	3	11
## 15	2020-01-26	0	11
## 16	2020-01-27	0	11
## 17	2020-01-28	3	14
## 18	2020-01-29	1	15
## 19	2020-01-30	1	16
## 20	2020-01-31	0	16
## 21	2020-02-01	1	17
## 22	2020-02-02	4	21
## 23	2020-02-03	0	21
## 24	2020-02-04	0	21
## 25	2020-02-05	0	21
## 26	2020-02-06	2	23
## 27	2020-02-07	1	24

## 28	2020-02-08	1	25
## 29	2020-02-09	0	25
## 30	2020-02-10	3	28
## 31	2020-02-11	6	34
## 32	2020-02-12	0	34
## 33	2020-02-13	4	38
## 34	2020-02-14	3	41
## 35	2020-02-15	7	48
## 36	2020-02-16	3	51
## 37	2020-02-17	13	64
## 38	2020-02-18	14	78
## 39	2020-02-19	8	86
## 40	2020-02-20	12	98
## 41	2020-02-21	23	121
## 42	2020-02-22	23	144
## 43	2020-02-23	23	167
## 44	2020-02-24	50	217
## 45	2020-02-25	39	256
## 46	2020-02-26	73	329
## 47	2020-02-27	68	397
## 48	2020-02-28	98	495
## 49	2020-02-29	91	586
## 50	2020-03-01	165	751
## 51	2020-03-02	180	931
## 52	2020-03-03	213	1144
## 53	2020-03-04	235	1379
## 54	2020-03-05	257	1636
## 55	2020-03-06	331	1967
## 56	2020-03-07	349	2316
## 57	2020-03-08	496	2812
## 58	2020-03-09	692	3504
## 59	2020-03-10	659	4163
## 60	2020-03-11	791	4954
## 61	2020-03-12	1048	6002
## 62	2020-03-13	1105	7107
## 63	2020-03-14	1051	8158
## 64	2020-03-15	991	9149
## 65	2020-03-16	916	10065
## 66	2020-03-17	455	10520
## 67	2020-03-18	281	10801
## 68	2020-03-19	180	10981
## 69	2020-03-20	52	11033
## 70	2020-03-21	13	11046
## 71	2020-03-22	5	11051
## 72	2020-03-23	4	11055
## 73	2020-03-24	0	11055

Visualize all data

```
epi <- ggplot(data = cdc)

epi + geom_point(aes(x = Date,
                     y = `Number of new cases`))+
```

```
# geom_line(aes(x = Date,
#               y = `Number of new cases`),
#           linetype = 2) +
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'
```

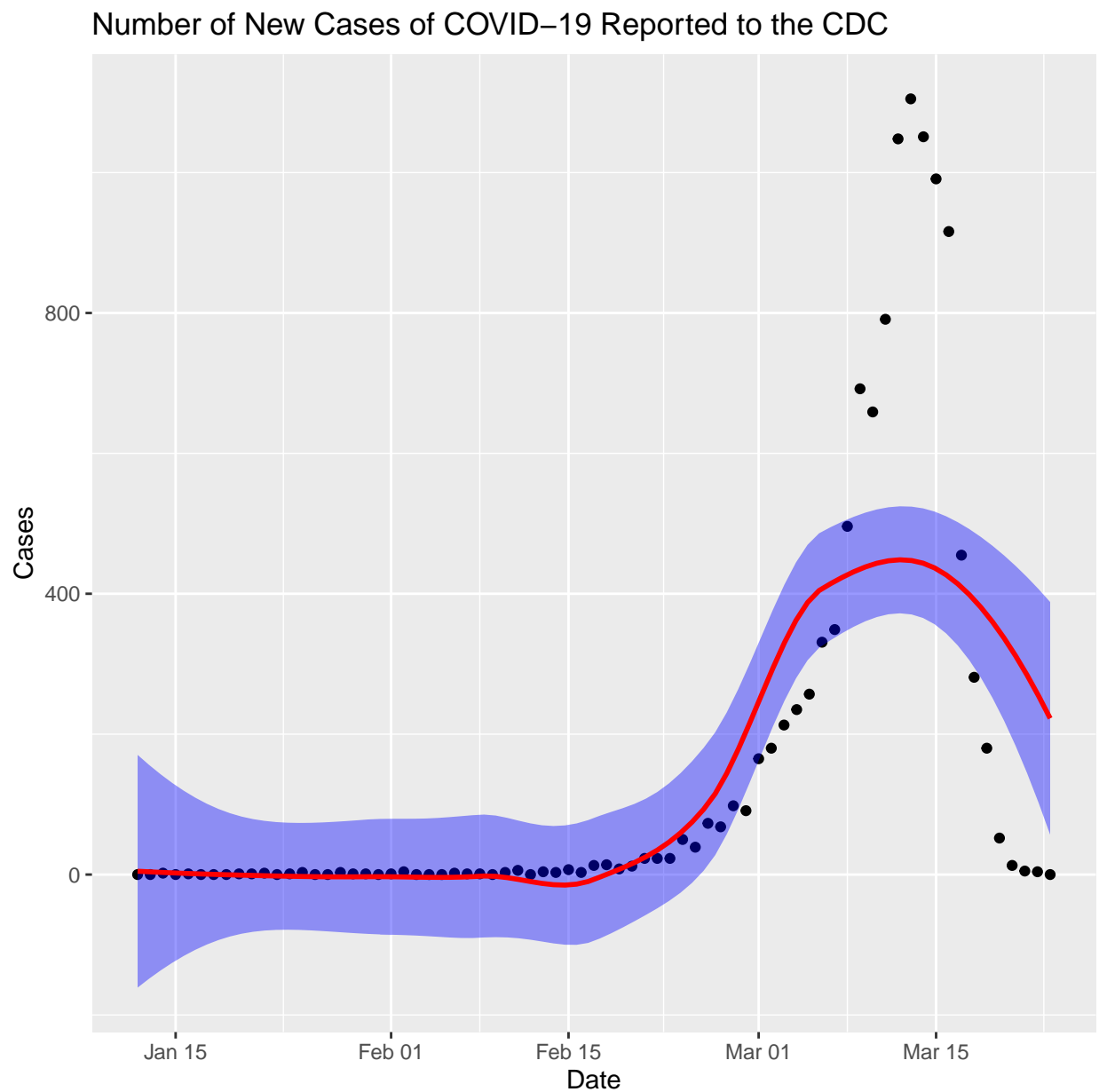


Figure 1: Epi curve 1

```

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'
```

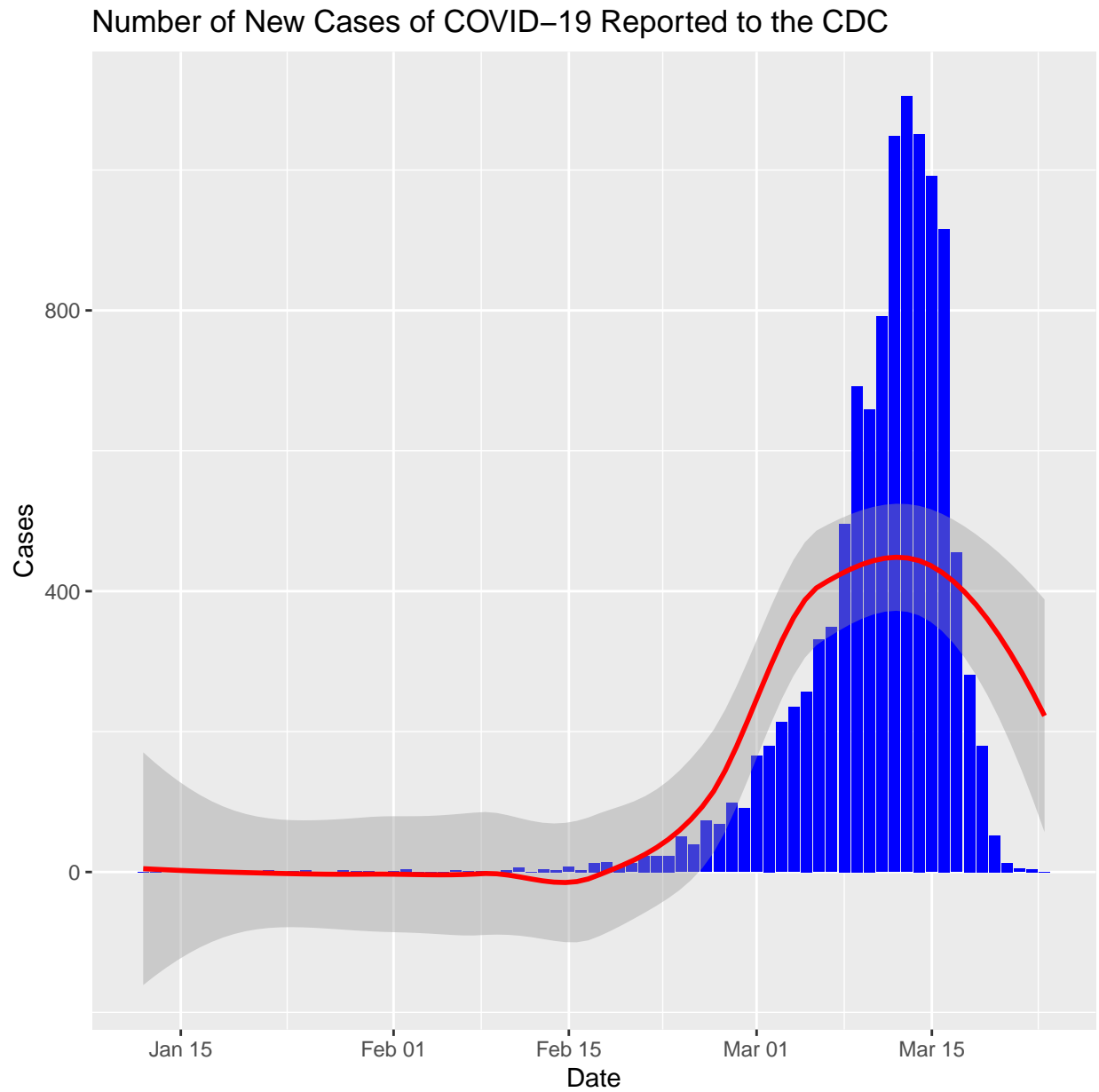


Figure 2: Epi curve 2, traditional

```

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",
    title = "Cumulative Number of Cases of COVID-19 Reported to the CDC") +
  geom_hline(yintercept = mean(cdc$cum)) +
  geom_hline(yintercept = median(cdc$cum),
    lty = 2)

```

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

Filter to remove incomplete reporting

remove dates on or after 5 March as this data may not be completely reported

```

cdc <- cdc %>%
  filter(Date < as.Date("2020-03-14"))

```

Visualize

```
epi <- ggplot(data = cdc)
```

```

epi + geom_point(aes(x = Date,
  y = `Number of new cases`))+
  # geom_line(aes(x = Date,
  # y = `Number of new cases`,
  # linetype = 2) +
  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,

```

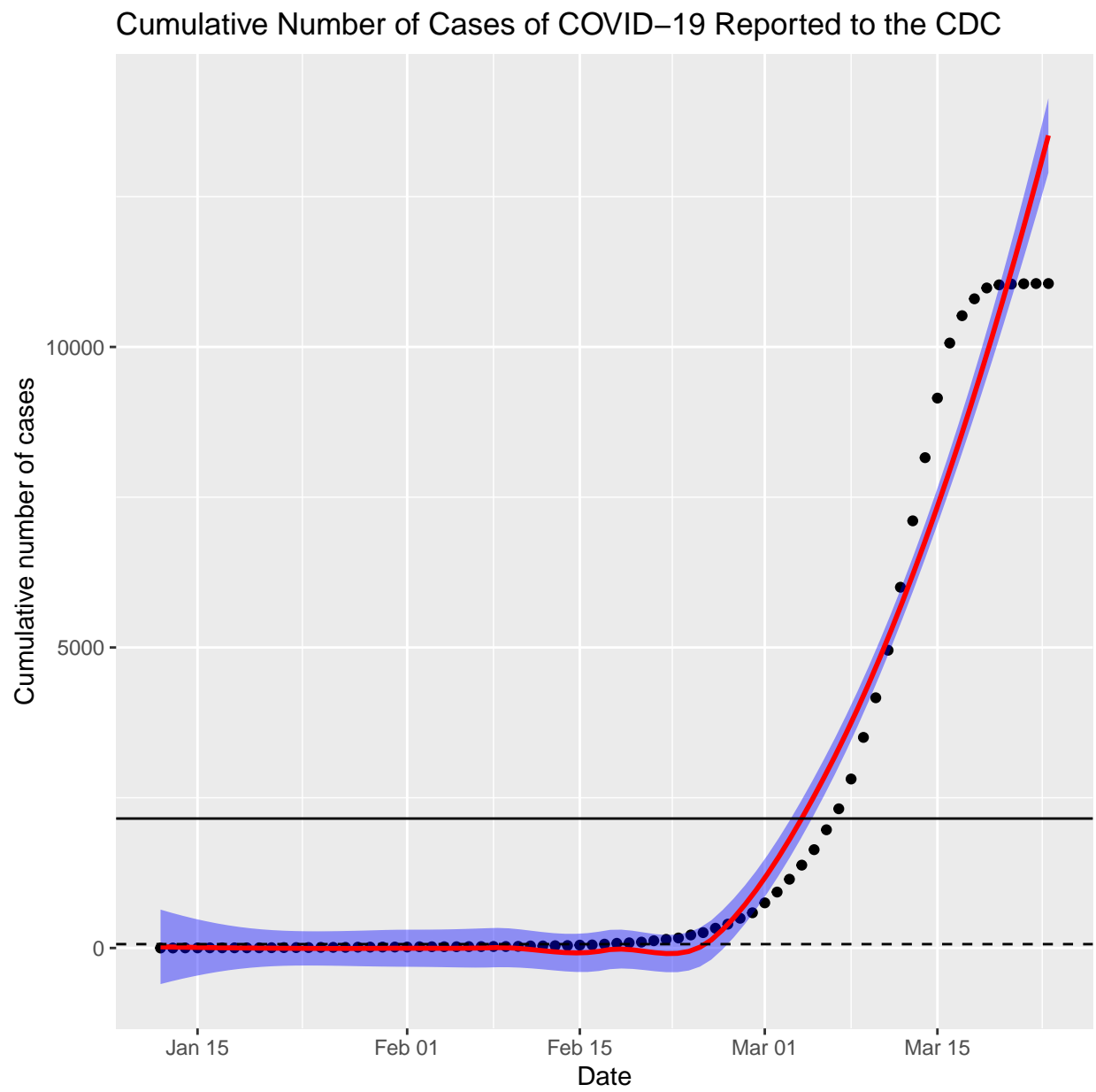


Figure 3: Cumulative cases

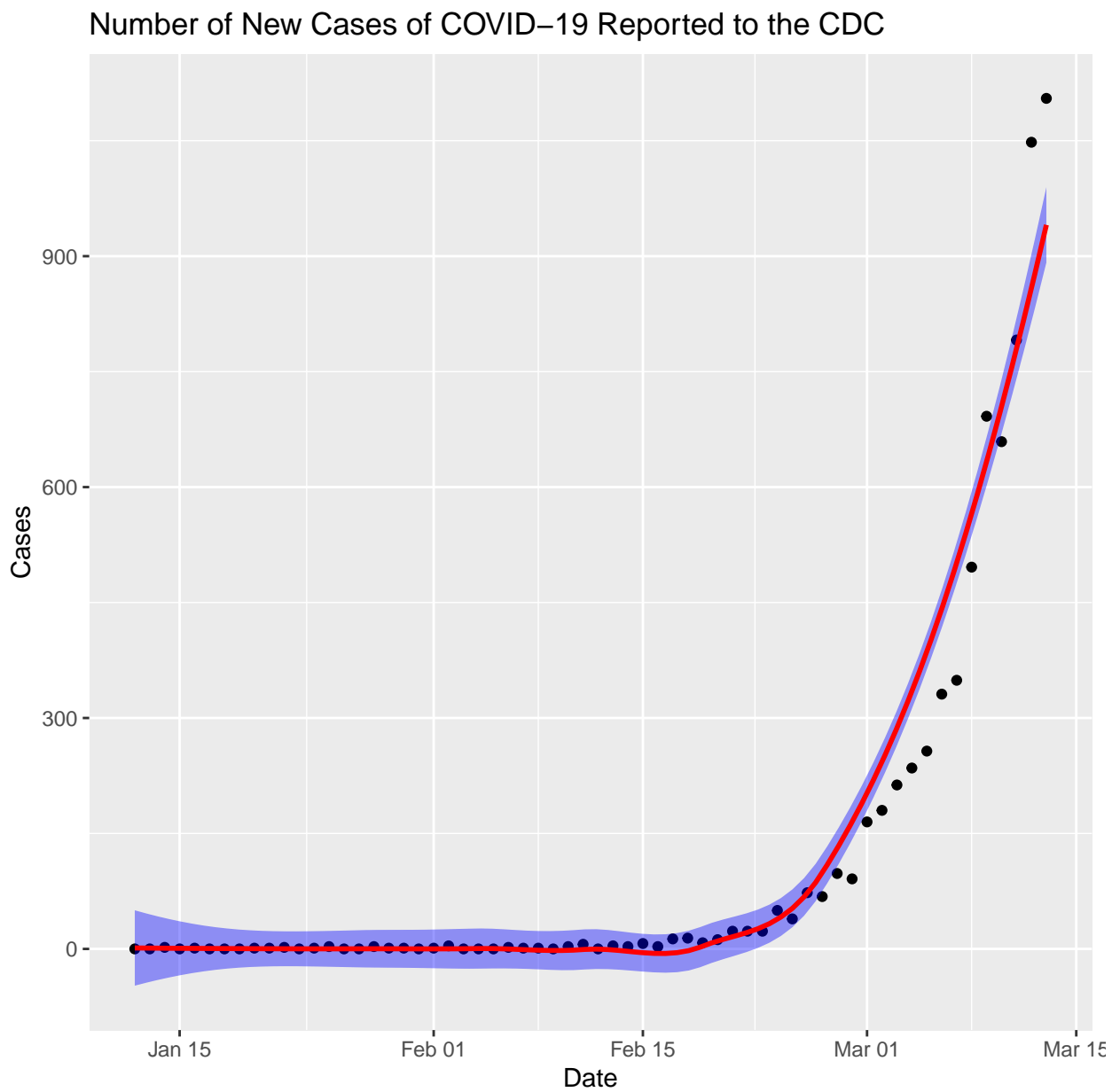


Figure 4: Epi curve 1

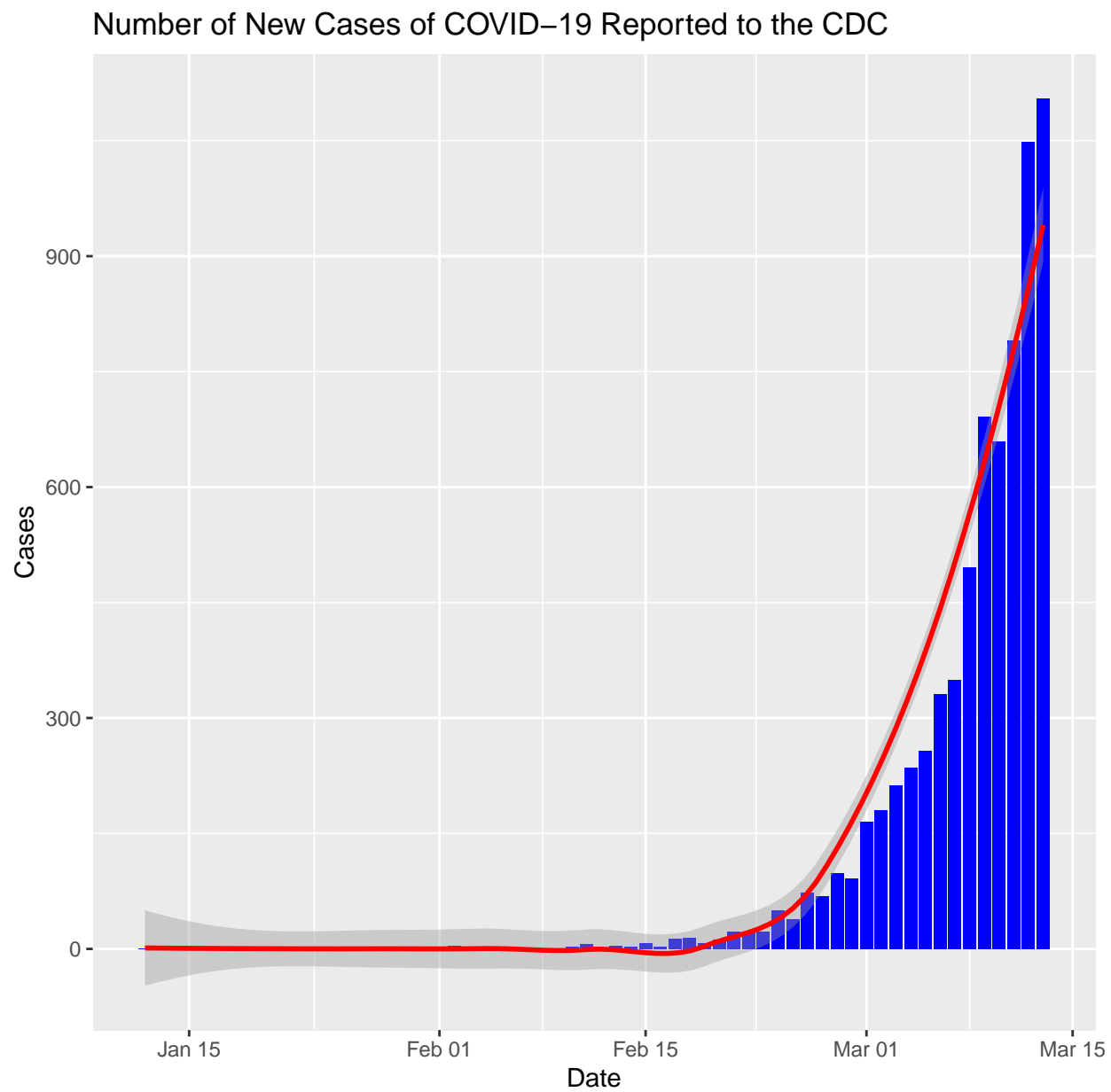


Figure 5: Epi curve 2, traditional

```

#y = `cum`)) +
geom_smooth(aes(x = Date,
                 y = `cum`,
                 color = "red",
                 fill = "blue") +
labs(y = "Cumulative number of cases",
      title = "Cumulative Number of Cases of COVID-19 Reported to the CDC") +
geom_hline(yintercept = mean(cdc$cum)) +
geom_hline(yintercept = median(cdc$cum),
           lty = 2)

```

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

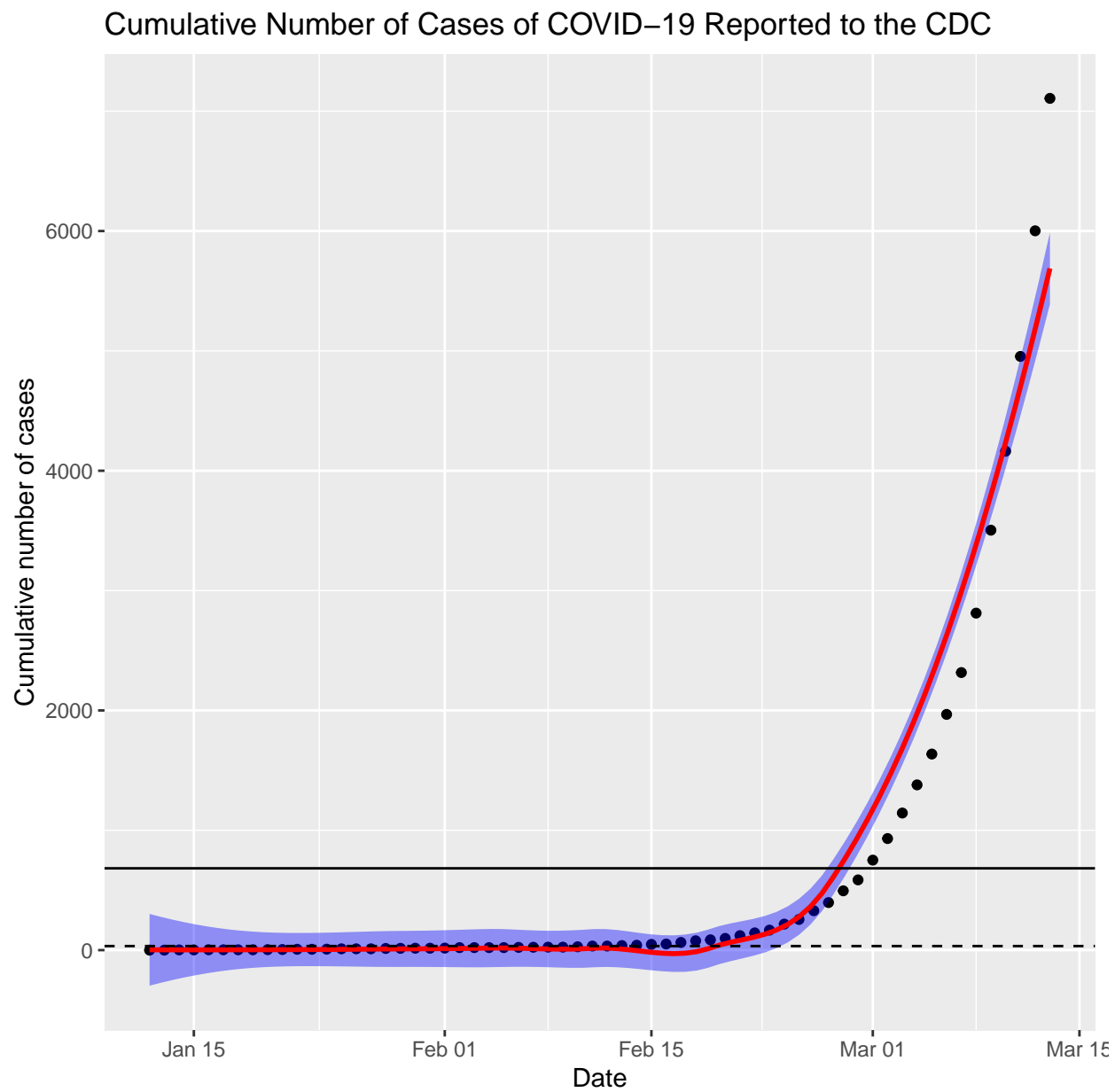


Figure 6: Cumulative cases