

Group 25 Case Study

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```
Case <- fread('Case.csv')
PatientInfo <- fread('PatientInfo.csv')
Policy <- fread('Policy.csv')
Region <- fread('Region.csv')
SearchTrend <- fread('SearchTrend.csv')
SeoulFloating <- fread('SeoulFloating.csv')
Time <- fread('Time.csv')
TimeAge <- fread('TimeAge.csv')
TimeGender <- fread('TimeGender.csv')
TimeProvince <- fread('TimeProvince.csv')
Weather <- fread('Weather.csv')
```

Work on Case file

```
case_dt1 <- Case %>% select(province, confirmed) #change to df function
case_dt1 <- as.data.table(case_dt1) [, lapply(.SD, sum),
                                     by = .(province = tolower(province))]
case_dt1 <- case_dt1[order(case_dt1$confirmed), ]
case_dt1 <- case_dt1[, proportion := confirmed / sum(confirmed) * 100]
```

ANALYSIS ON PROVINCE

Work on TimeProvince file

```
timeprovince_dt <- TimeProvince %>% group_by(province)

outliers_dt <- TimeProvince[province == "Gyeongsangbuk-do" | province == "Daegu"]
outliers_dt <- outliers_dt %>% select(date, confirmed)
outliers_dt <- outliers_dt %>%
  group_by(date) %>%
  summarize(confirmed = sum(confirmed)) %>%
  ungroup

## `summarise()` ungrouping output (override with `.groups` argument)
names(outliers_dt)[1:2] <- c("date", "confirmed_outliers")
time_dt <- merge(Time, outliers_dt, by = "date", all = FALSE)
time_dt <- time_dt[, no_outliers := confirmed - confirmed_outliers]
```

Plot Nr.1: Population* in provinces

(*Data is obtained from [here](#))

```

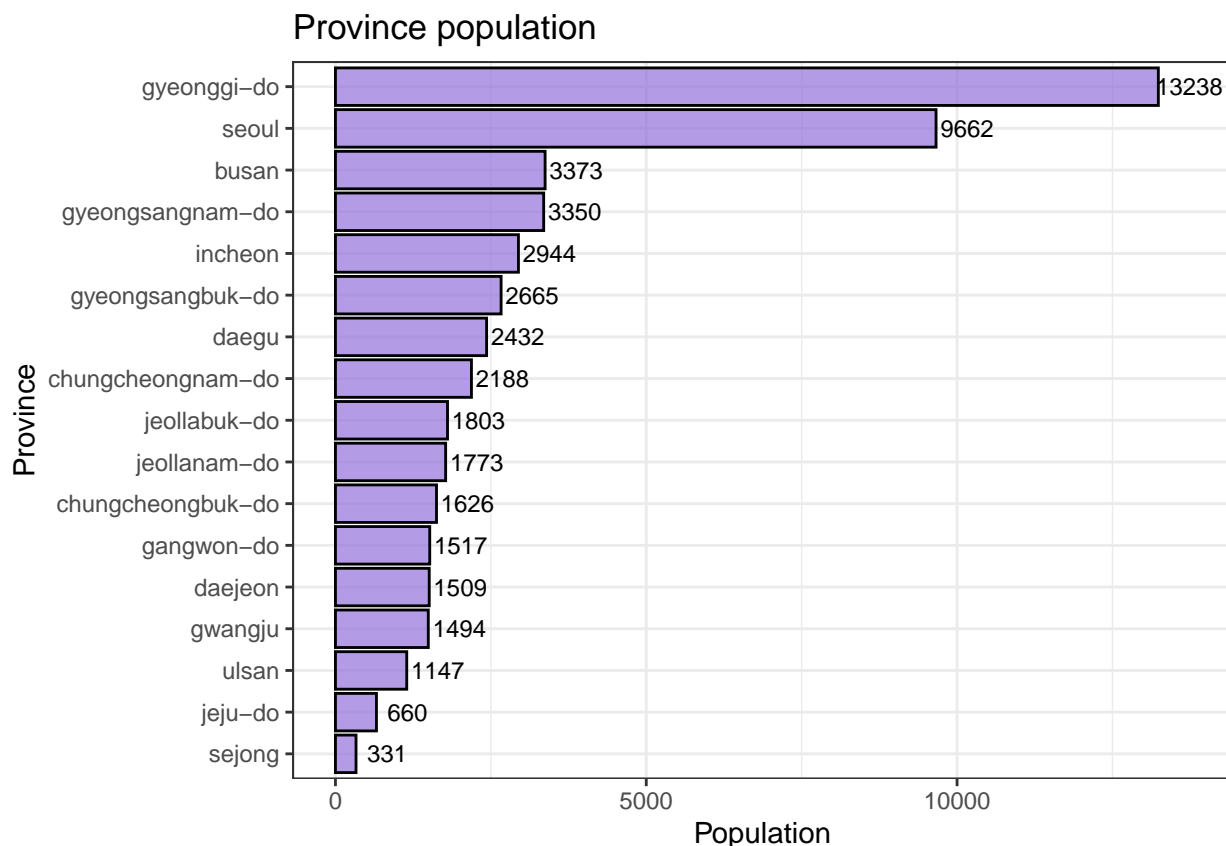
case_dt1 <- case_dt1[order(case_dt1$province), ] #sort in alph. order
case_dt1 <- case_dt1[, population := as.integer(c("3373", "1626", "2188", "2432", "1509",
"1517", "1494", "13238", "2665", "3350",
"2944", "660", "1803", "1773", "331",
"9662", "1147"))]

case_dt1 <- case_dt1[order(case_dt1$population), ]

plot1 <- ggplot(case_dt1, aes(x = reorder(province, population), y = population)) +
  geom_bar(stat = "identity",
    color = "black",
    fill = "mediumpurple",
    alpha = 0.7) +
  coord_flip() +
  geom_text(aes(label = population, y = population + 500), size = 3) +
  labs(x = 'Province', y = 'Population') +
  ggtitle("Province population") +
  theme_bw()

```

plot1



Plot Nr.2: Cases in provinces

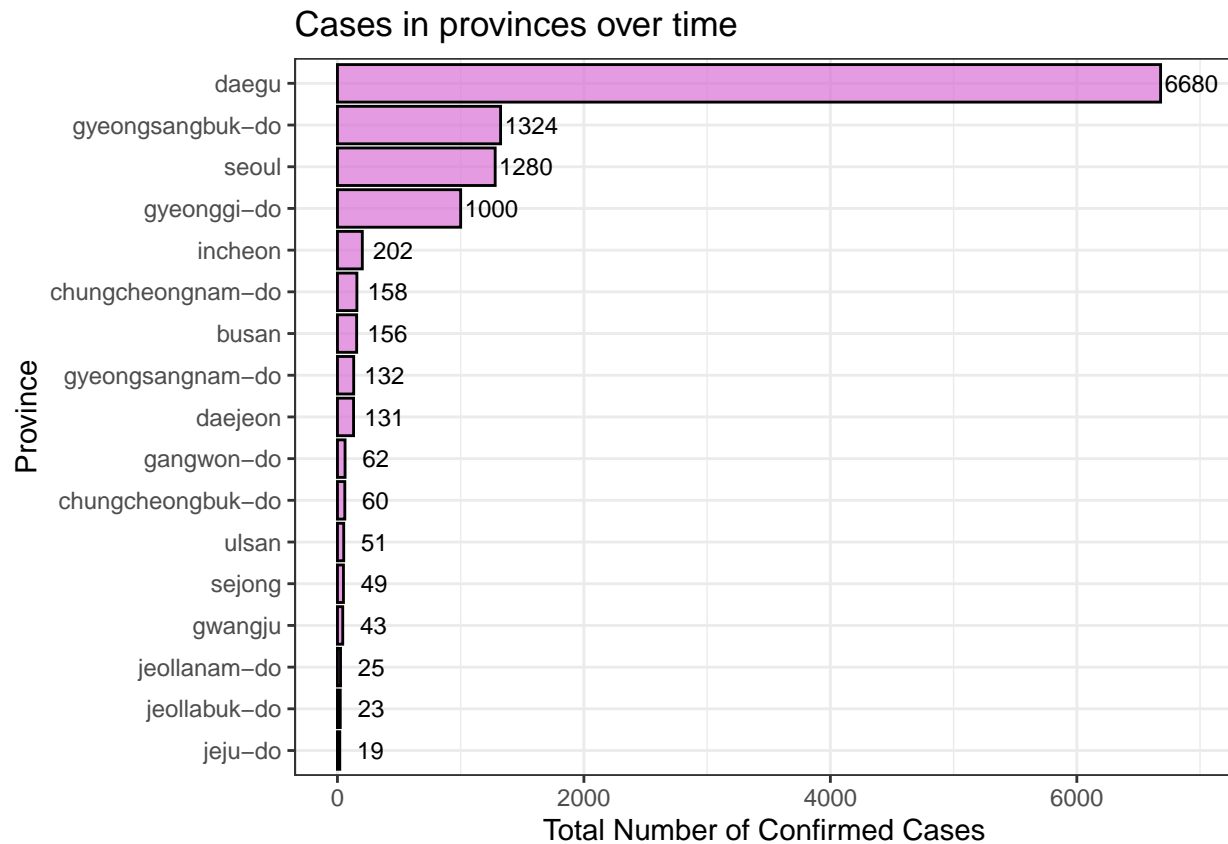
```

plot2 <- ggplot(case_dt1, aes(x=reorder(province, confirmed), y=confirmed)) +
  geom_bar(stat="identity", color="black", fill= "orchid", alpha = 0.7) +
  coord_flip() +

```

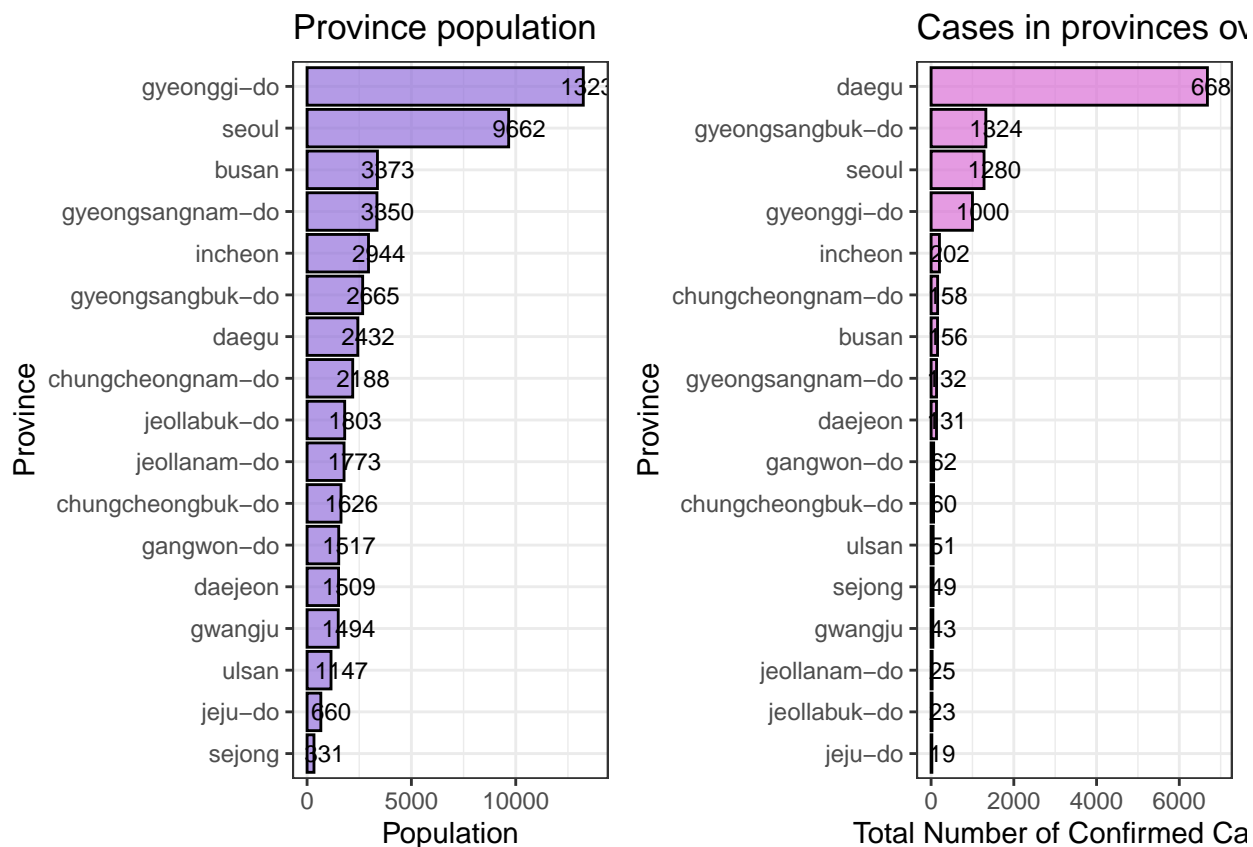
```
geom_text(aes(label=confirmed, y = confirmed+250), size=3) +
labs(x='Province', y='Total Number of Confirmed Cases') +
ggtitle("Cases in provinces over time") +
theme_bw()
```

plot2



Plot Nr.3: Comparison of population and number of cases

```
require(gridExtra)
grid.arrange(plot1, plot2, ncol=2)
```

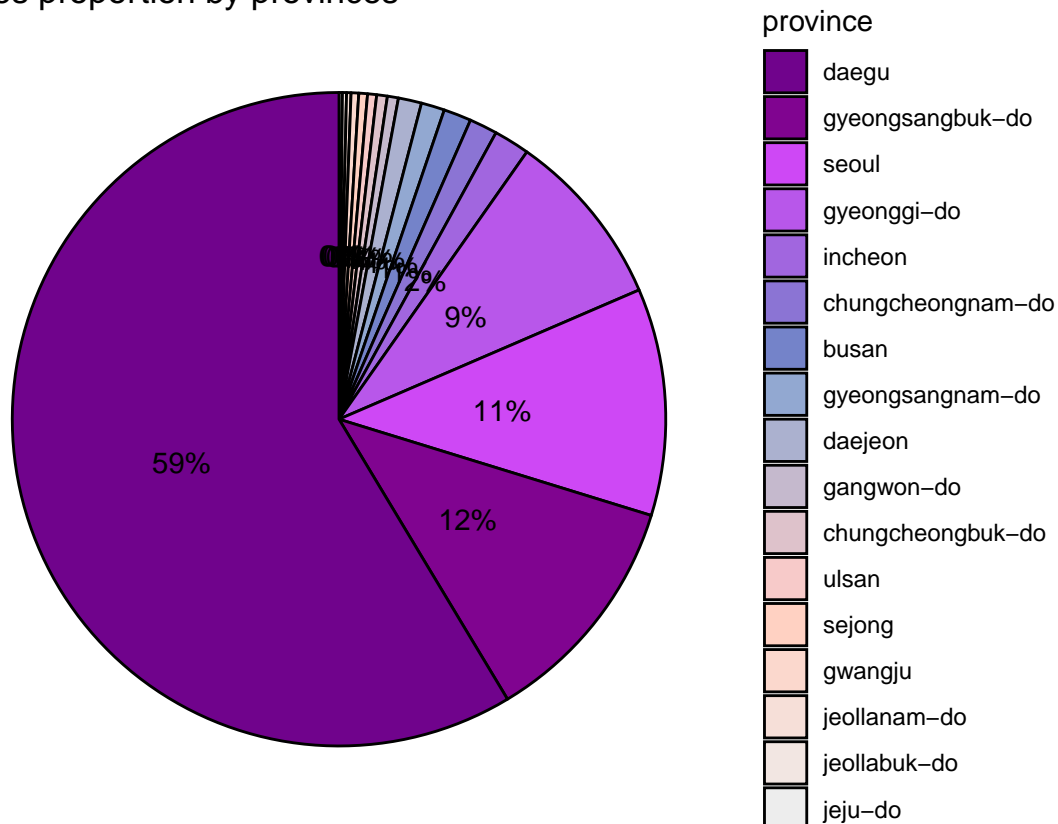


Plot Nr.4: Proportion of total cases by province

```
case_dt1 <- case_dt1[order(case_dt1$proportion),]
case_dt1$province <- factor(case_dt1$province,
                             levels = rev(as.character(case_dt1$province)))
case_dt1_x <- case_dt1 %>%
  arrange(desc(province)) %>%
  mutate(y = proportion / sum(proportion), y.stack = cumsum(y)) %>%
  ungroup()

ggplot(case_dt1_x, aes(x='', label=paste0(round(proportion), "%"), y=y, fill=province))+
  geom_bar(stat="identity", width=1, color="black") +
  coord_polar("y") +
  geom_text(aes(label=paste0(round(proportion), "%")), position=position_stack(vjust=0.5))+
  ggtitle("Cases proportion by provinces")+
  scale_fill_manual(values = rev(c("#ededed", "#f2e6e2", "#f6dfd8", "#fbd8cd", "#ffd1c2",
                                   "#f7cac9", "#dec2cb", "#c5b9cd", "#abb1cf", "#92a8d1",
                                   "#7483c9", "#8b74d4", "#a166df", "#b857ea", "#ce48f5",
                                   "#800490", "#70038c")))) +
  theme_void()
```

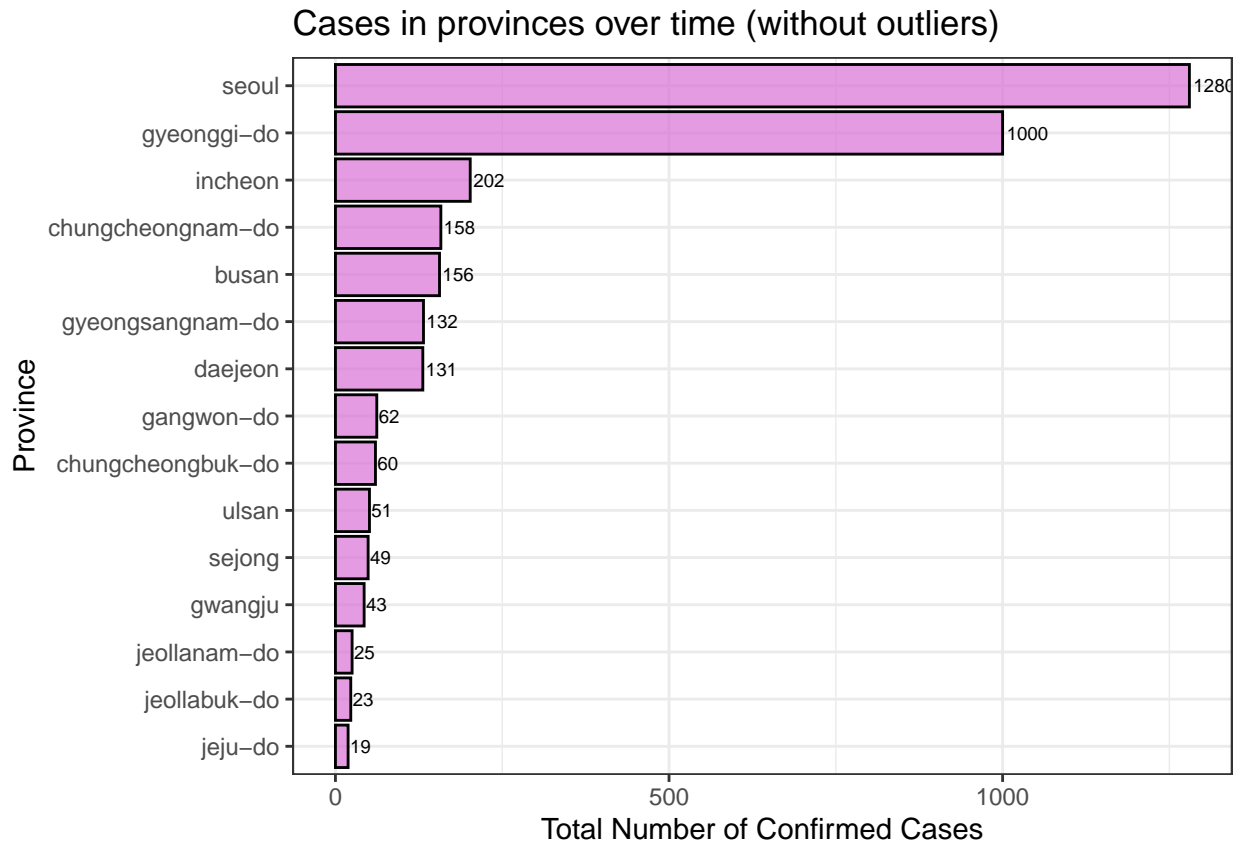
Cases proportion by provinces



Plot Nr.5: Cases in provinces without 2 outliers

```
case_dt2<-case_dt1[!(case_dt1$province=="daegu" | case_dt1$province=="gyeongsangbuk-do"),]
plot5<-ggplot (case_dt2, aes(x=reorder(province, confirmed),y=confirmed))+
  geom_bar(stat="identity", color="black",fill= "orchid", alpha = 0.7)+
  coord_flip()+
  geom_text(aes(label=confirmed), hjust= - 0.1, size=2.5)+
  labs( x='Province', y='Total Number of Confirmed Cases') +
  ggtitle("Cases in provinces over time (without outliers)")+
  theme_bw()
```

plot5

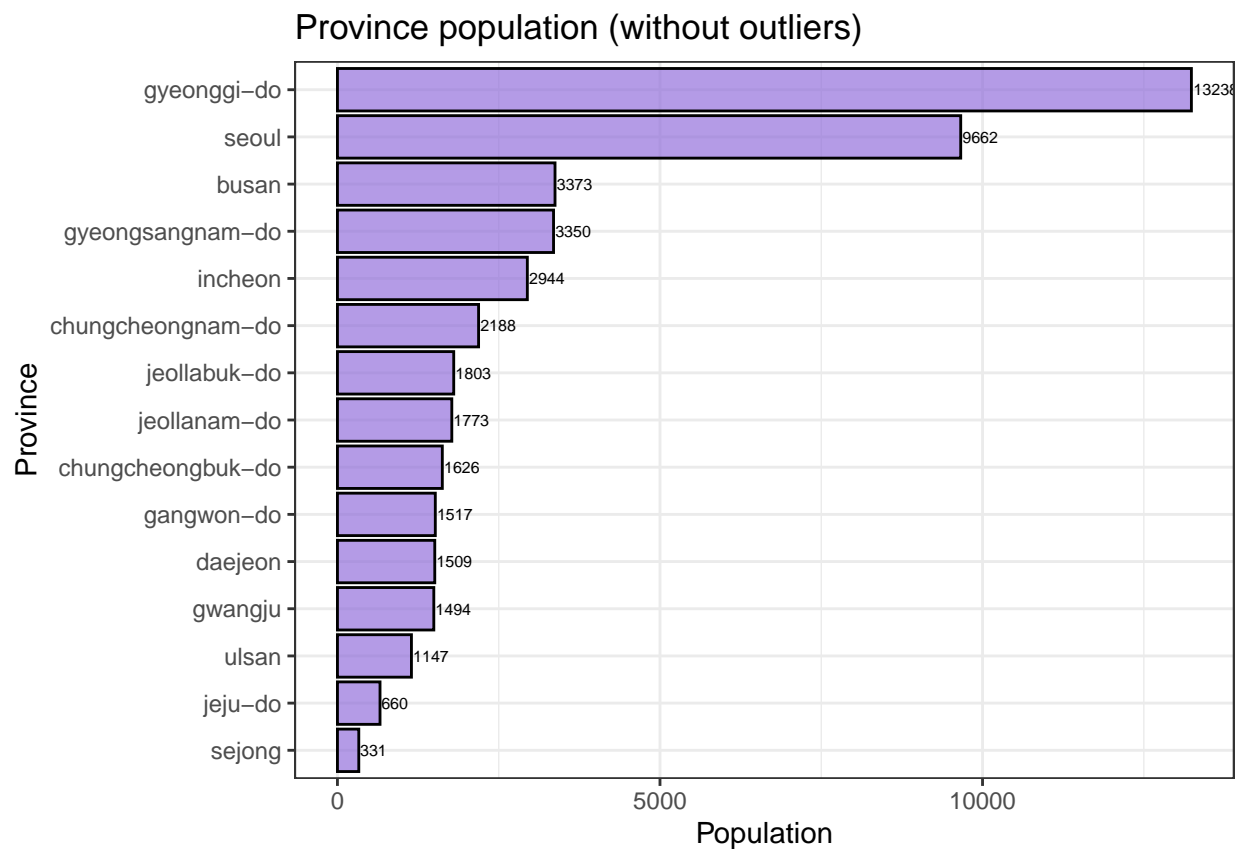


Plot Nr.6: Population in provinces without 2 outliers

```
case_dt1 <- case_dt1[order(case_dt1$province),] #sort in alph. order
case_dt1 <- case_dt1[, population:= as.integer(c("3373","1626","2188",
"2432","1509","1517",
"1494","13238","2665",
"3350","2944","660",
"1803","1773","331",
"9662","1147")))]

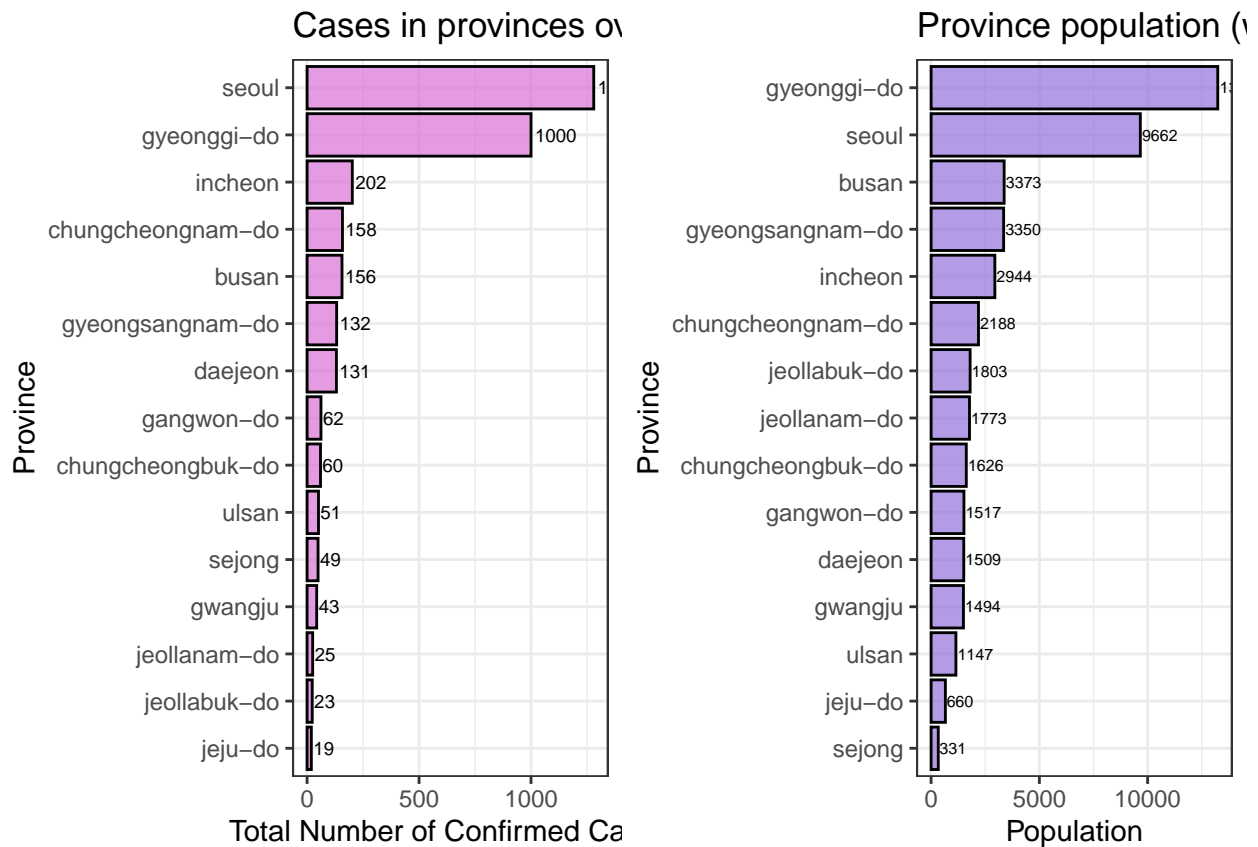
plot6 <- ggplot (case_dt2, aes(x=reorder(province, population), y=population))+
  geom_bar(stat="identity", color="black",fill= "mediumpurple", alpha = 0.7)+
  coord_flip()+
  geom_text(aes(label=population), hjust=-0.05, size=2.2)+
  labs( x='Province', y='Population') +
  ggtitle("Province population (without outliers)")+
  theme_bw()

plot6
```



Plot Nr.7: Comparison of Plot Nr.1 and Plot Nr.4

```
require(gridExtra)
grid.arrange(plot5, plot6, ncol=2)
```



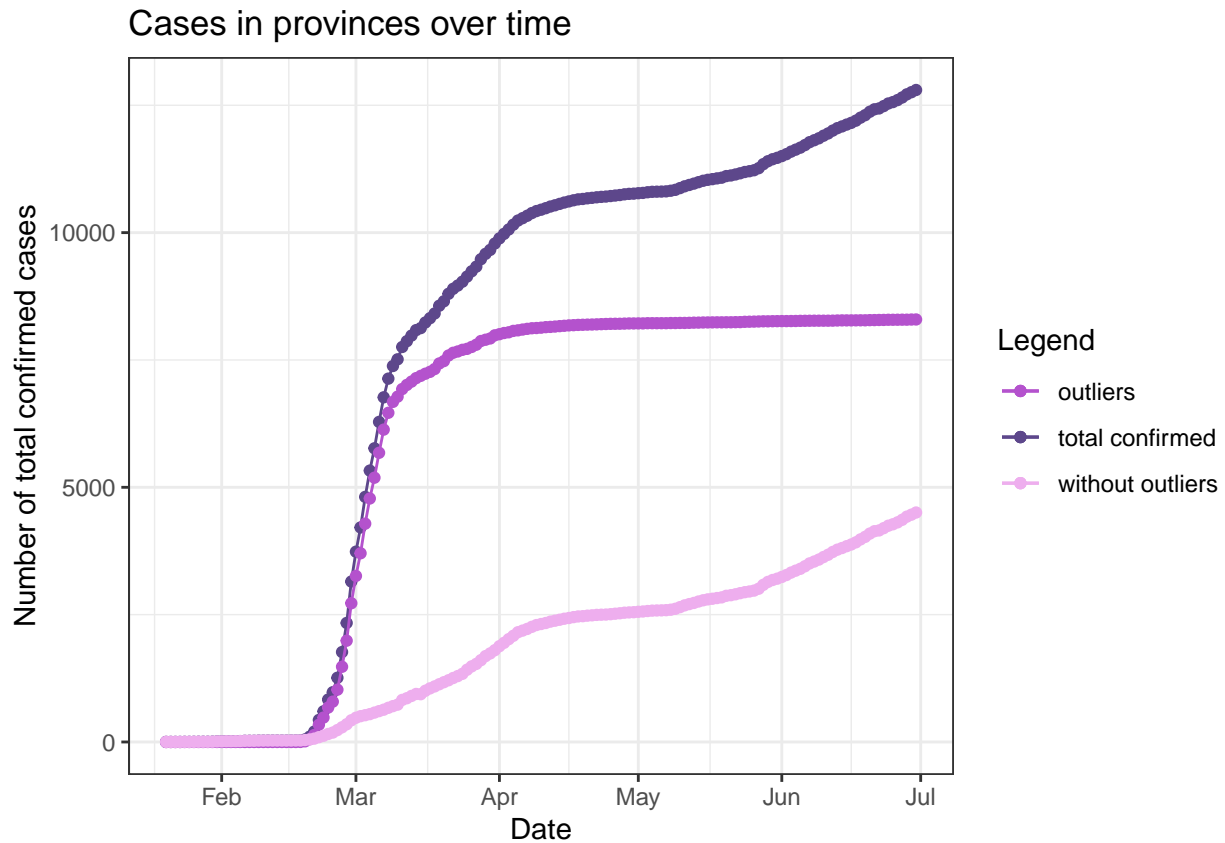
Plot Nr.8: General overview

```

colors <- c("total confirmed" = "mediumpurple4",
            "outliers" = "mediumorchid3",
            "without outliers" = "plum2")

ggplot(time_dt, aes(x=date)) +
  geom_line(aes(y=confirmed, color="total confirmed")) +
  geom_point(aes(y=confirmed, color="total confirmed")) +
  geom_line(aes(y=confirmed_outliers, color="outliers")) +
  geom_point(aes(y=confirmed_outliers, color="outliers")) +
  geom_line(aes(y=no_outliers, color="without outliers")) +
  geom_point(aes(y=no_outliers, color="without outliers")) +
  theme(axis.text.x = element_text(angle = 45, vjust = 0.5, hjust=1)) +
  labs(x='Date', y='Number of total confirmed cases', color="Legend") +
  ggtitle("Cases in provinces over time")+
  scale_color_manual(values = colors)+
  theme_bw()

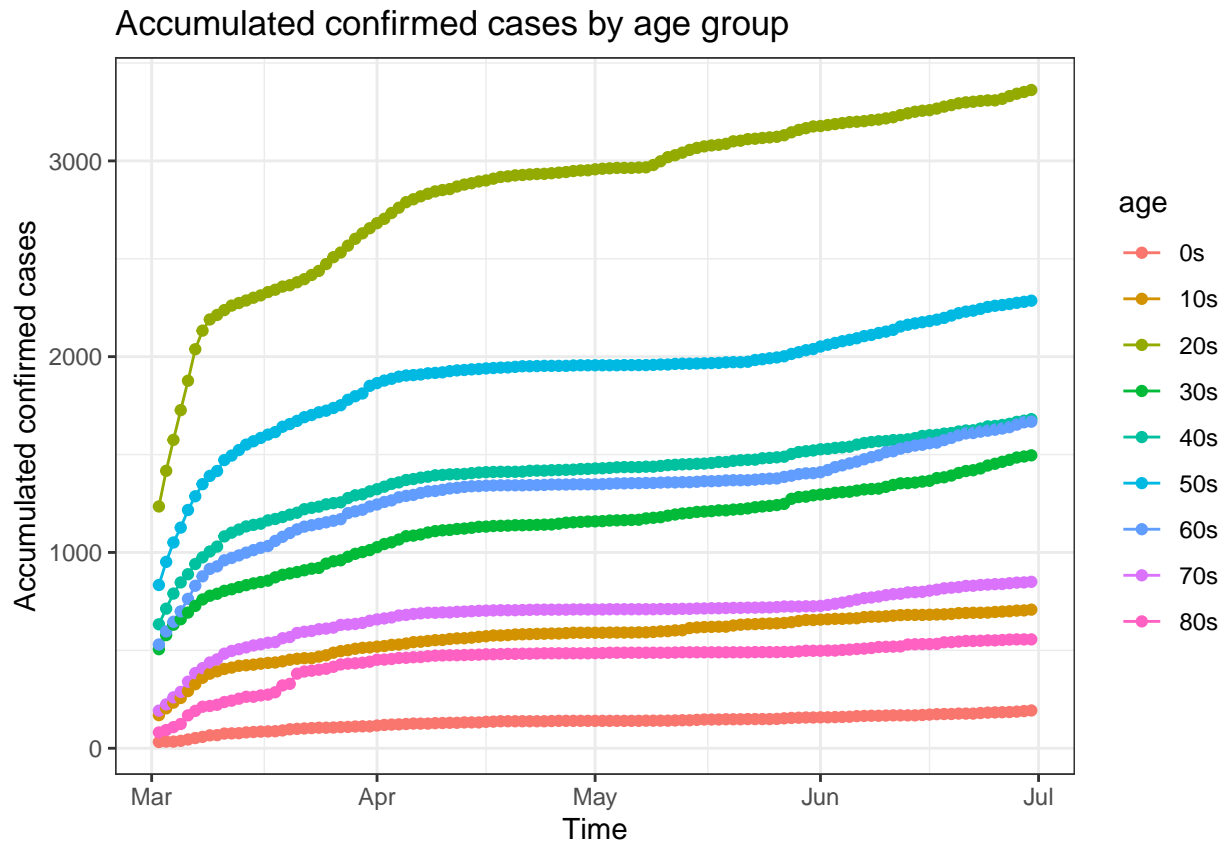
```

ANALYSIS ON GENDER AND AGE

Plot Nr.9: Accumulated confirmed cases by age group

```
TimeAge[, date := as.Date(date, "%Y-%m-%d")]
ggplot(TimeAge, aes(
  x = date,
  y = confirmed,
  color = age,
  group = age
)) +
  geom_line() + geom_point() +
  ggtitle("Accumulated confirmed cases by age group") +
  theme(axis.text.x = element_text(angle = 30, hjust = 1)) +
  labs(x = "Time", y = "Accumulated confirmed cases") + theme_bw()
```

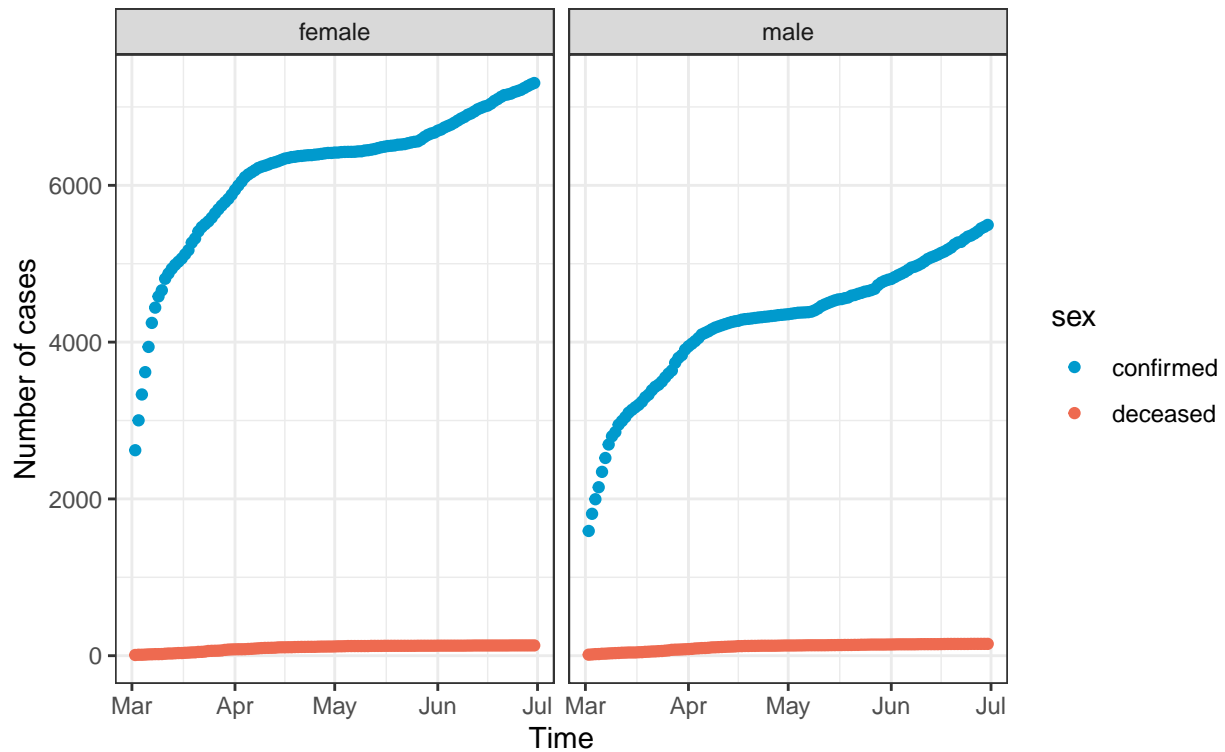


```
TimeGender[, date := as.Date(date, "%Y-%m-%d")]
gender_reshape <- melt(TimeGender,
  id.vars = c("date", "sex", "time"),
  measure.vars = c("confirmed", "deceased"))
female_time <- gender_reshape[sex == "female"]
male_time <- gender_reshape[sex == "male"]
```

Plot Nr.10: Accumulated confirmed and deceased cases over time

```
ggplot(gender_reshape, aes(x = date, y = value, col = variable)) +
  geom_point() +
  labs(x = "Time", y = "Number of cases",
    title = "Accumulated confirmed and deceased cases over time",
    subtitle = "male and female") +
  scale_color_manual("sex", values = c("deceased" = "coral2",
    "confirmed" = "deepskyblue3")) +
  facet_wrap(~ sex) +
  theme_bw()
```

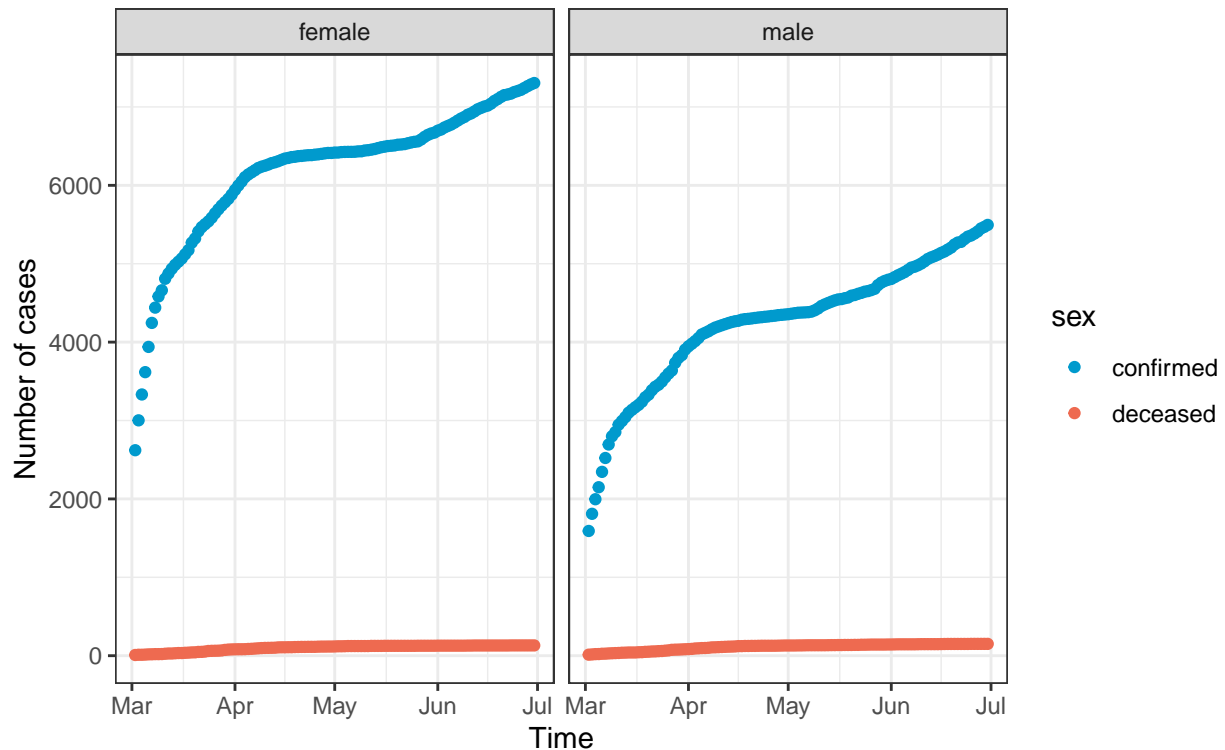
Accumulated confirmed and deceased cases over time
male and female



Plot Nr.11: Accumulated confirmed and deceased cases over time
By gender

```
ggplot(gender_reshape, aes(x = date, y = value, col = variable)) + geom_point() +
  labs(
    x = "Time",
    y = "Number of cases",
    title = "Accumulated confirmed and deceased cases over time",
    subtitle = "By Gender"
  ) +
  scale_color_manual("sex",
    values = c("deceased" = "coral2", "confirmed" = "deepskyblue3")) +
  facet_wrap(~ sex) + theme_bw()
```

Accumulated confirmed and deceased cases over time
By Gender



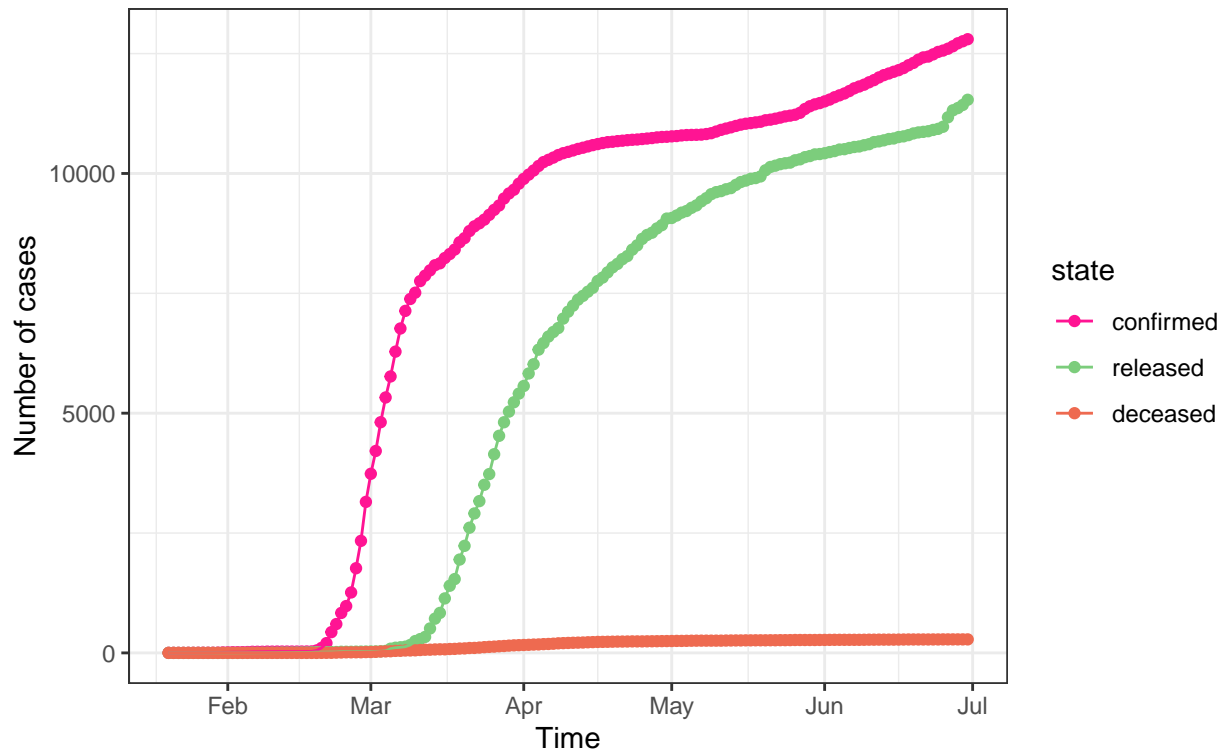
Plot Nr.12: Accumulated number of cases over time

By state of patients

```
# reshape time data
time_reshape <- melt(Time,
  id.vars = "date",
  measure.vars = c("confirmed", "released", "deceased"))

ggplot(time_reshape, aes(x = date, y = value, col = variable)) +
  geom_point() + geom_line() +
  labs(x = "Time",
    y = "Number of cases",
    title = "Accumulated number of cases over time",
    subtitle = "By state of patients") +
  scale_color_manual("state",
    values = c("deceased" = "coral2",
      "confirmed" = "deeppink",
      "released" = "palegreen3")) +
  theme_bw()
```

Accumulated number of cases over time
By state of patients



Working on PatientInfo

```
PatientInfo[sex == "female", .N]
```

```
## [1] 2218
```

```
PatientInfo[sex == "male", .N]
```

```
## [1] 1825
```

```
PatientInfo$age <- factor(PatientInfo$age,
  levels = c("0s", "10s", "20s", "30s",
    "40s", "50s", "60s", "70s",
    "80s", "90s", "100s"))
```

```
patient_state_sex <- PatientInfo[sex != "", .(sex, age, confirmed_date, state)]
patient_state_sex[, state := as.factor(state)]
patient_state_sex[, sex := as.factor(sex)]
sapply(patient_state_sex, class)
```

```
## $sex
## [1] "factor"
##
## $age
## [1] "factor"
##
## $confirmed_date
## [1] "IDate" "Date"
```

```
##
## $state
## [1] "factor"

dead_patients_bysex <- patient_state_sex[state == "deceased"]
```

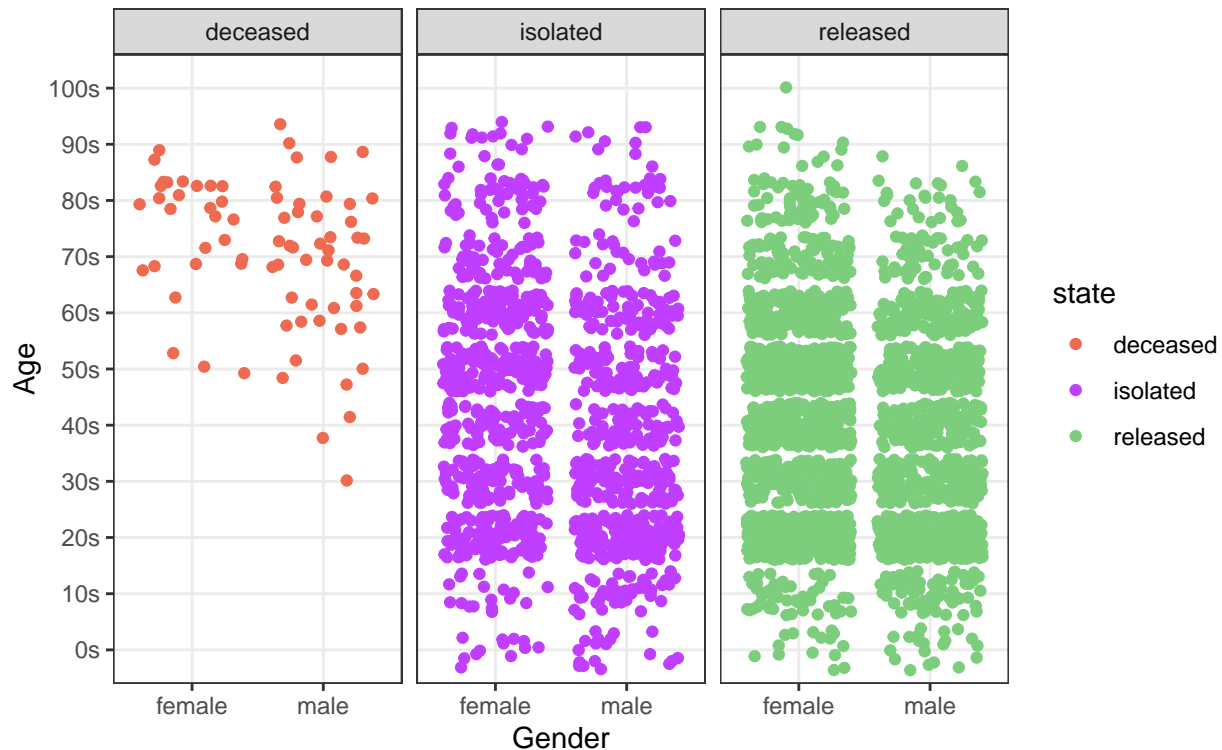
Plot Nr.13: Confirmed case distribution by state

Gender vs Age

```
patient_state_sex %>%
  filter(is.na(confirmed_date) == F) %>%
  filter(is.na(age) == F) %>%
  ggplot(., aes(x = sex, y = age, col = state)) +
  geom_jitter() +
  labs(
    title = "Confirmed case distribution by state",
    subtitle = "Gender vs Age",
    y = "Age",
    x = "Gender"
  ) + facet_wrap( ~ state) +
  scale_color_manual(
    "state",
    values = c(
      "deceased" = "coral2",
      "isolated" = "darkorchid1",
      "released" = "palegreen3"
    )
  ) +
  theme_bw()
```

Confirmed case distribution by state

Gender vs Age

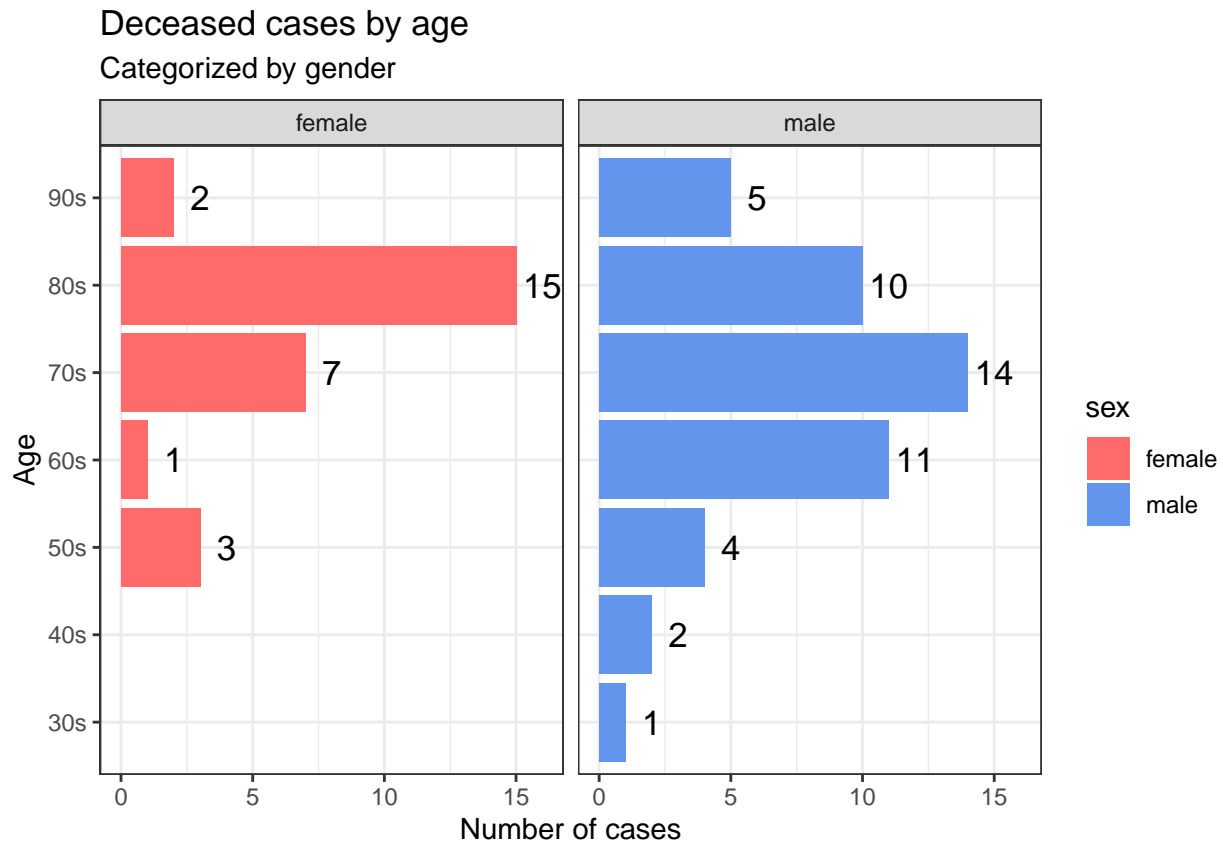


Plot Nr.14: Deceased cases by age

Categorized by gender

```
dead_patients_bysex %>%
  group_by(sex, age) %>%
  summarise(count = n()) %>%
  ggplot(., aes(x = age, y = count, fill = sex)) + geom_bar(stat = "identity") +
  geom_text(aes(label = count, y = count + 1), size = 4.5) +
  labs(
    x = "Age",
    y = "Number of cases",
    title = "Deceased cases by age",
    subtitle = "Categorized by gender"
  ) +
  scale_fill_manual("sex",
    values = c("female" = "indianred1", "male" = "cornflowerblue")) +
  facet_wrap(~ sex) + coord_flip() + theme_bw()
```

`summarise()` regrouping output by 'sex' (override with `.groups` argument)



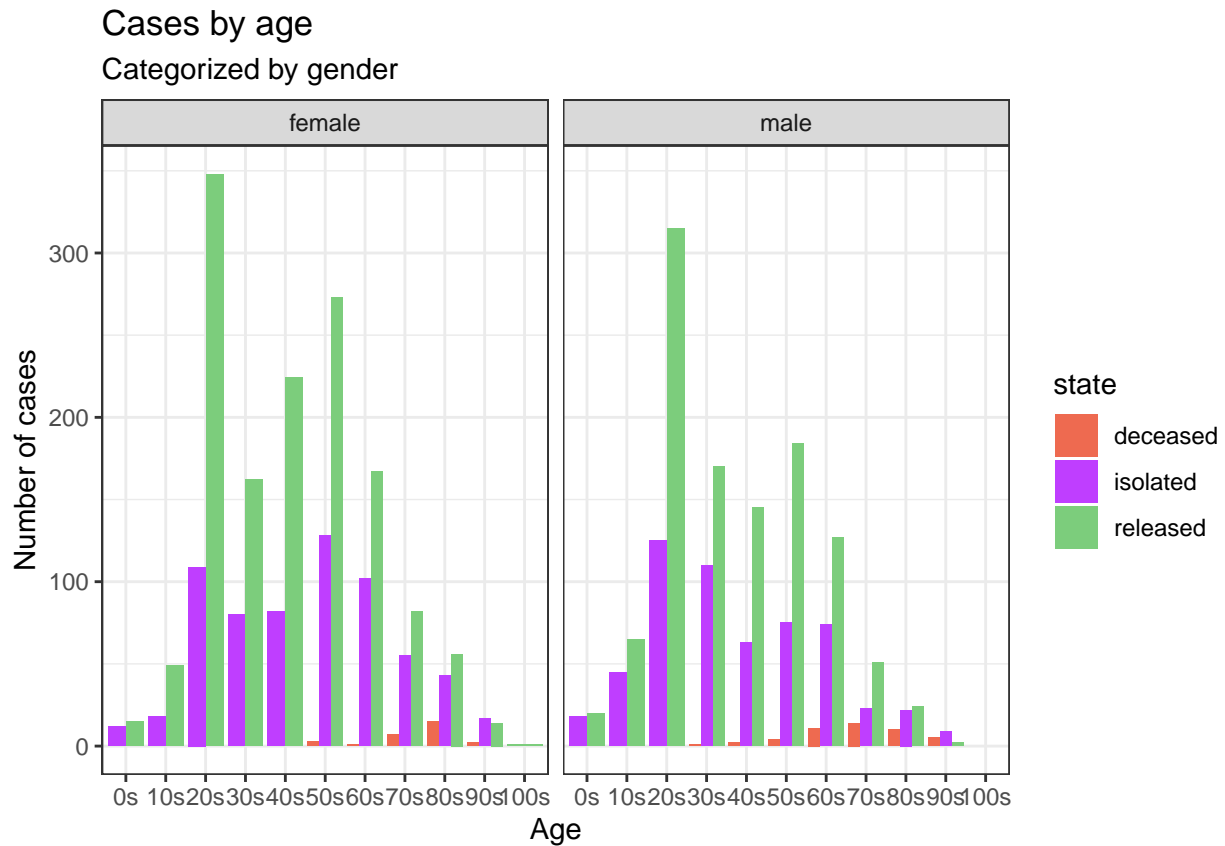
Plot Nr.15: Cases by age

Categorized by gender

```
patient_state_sex %>%
  filter(is.na(confirmed_date) == F) %>%
  filter(is.na(age) == F) %>%
  group_by(sex, age, state) %>%
  summarise(count = n()) %>%
  ggplot(., aes(x = age, y = count, fill = state)) +
  geom_bar(stat = "identity", position = "dodge") +
  labs(
    x = "Age",
    y = "Number of cases",
    title = "Cases by age",
    subtitle = "Categorized by gender"
  ) +
  scale_fill_manual(
    "state",
    values = c(
      "deceased" = "coral2",
      "isolated" = "darkorchid1",
      "released" = "palegreen3"
    )
  ) +
  facet_wrap(~ sex) + theme_bw()
```



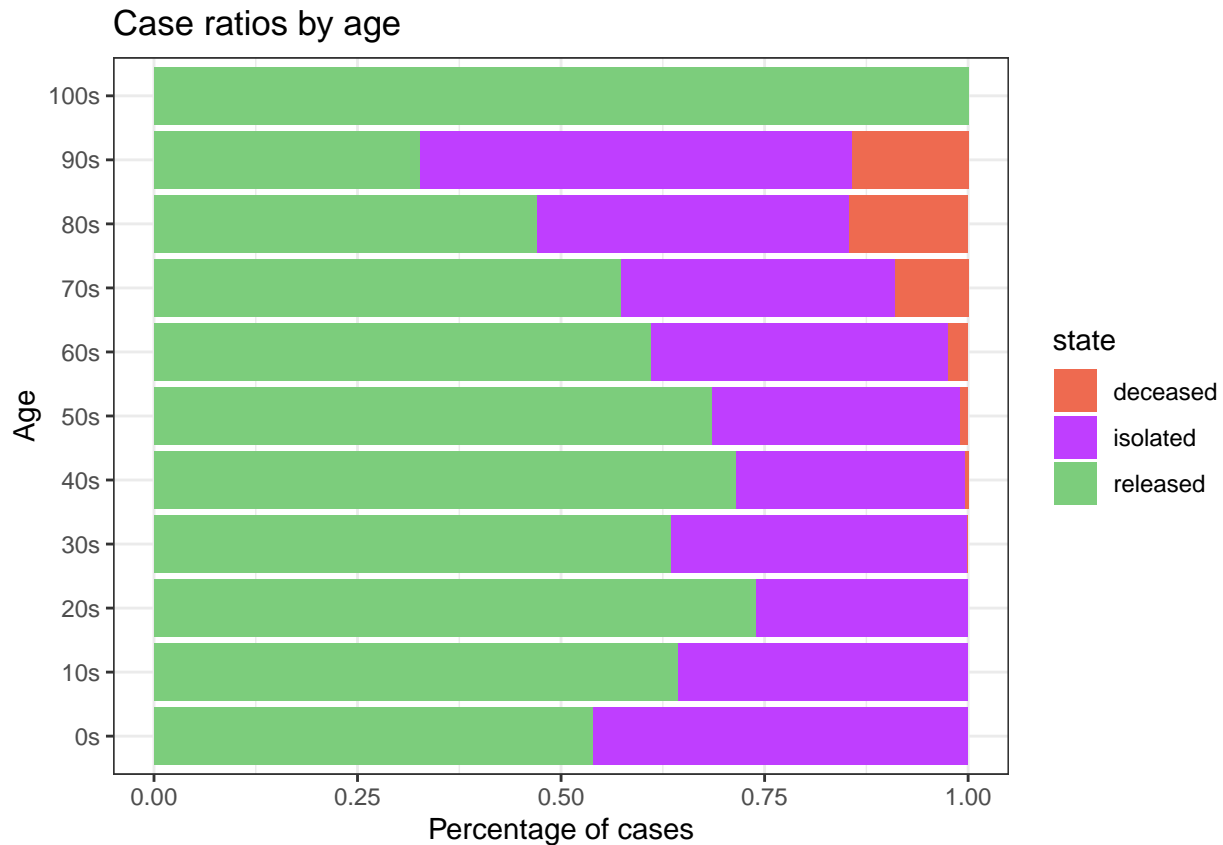
```
## `summarise()` regrouping output by 'sex', 'age' (override with `.groups` argument)
```



Plot Nr.16: Case ratios by age

```
patient_state_sex %>%
  filter(is.na(confirmed_date) == F) %>%
  filter(is.na(age) == F) %>%
  group_by(sex, age, state) %>%
  summarise(count = n()) %>%
  ggplot(., aes(x = age, y = count, fill = state)) +
  geom_bar(stat = "identity", position = "fill") +
  labs(x = "Age", y = "Percentage of cases", title = "Case ratios by age") +
  scale_fill_manual(
    "state",
    values = c(
      "deceased" = "coral2",
      "isolated" = "darkorchid1",
      "released" = "palegreen3"
    )
  ) + coord_flip() + theme_bw()
```

```
## `summarise()` regrouping output by 'sex', 'age' (override with `.groups` argument)
```



Plot Nr.17: Case ratios by age

By gender

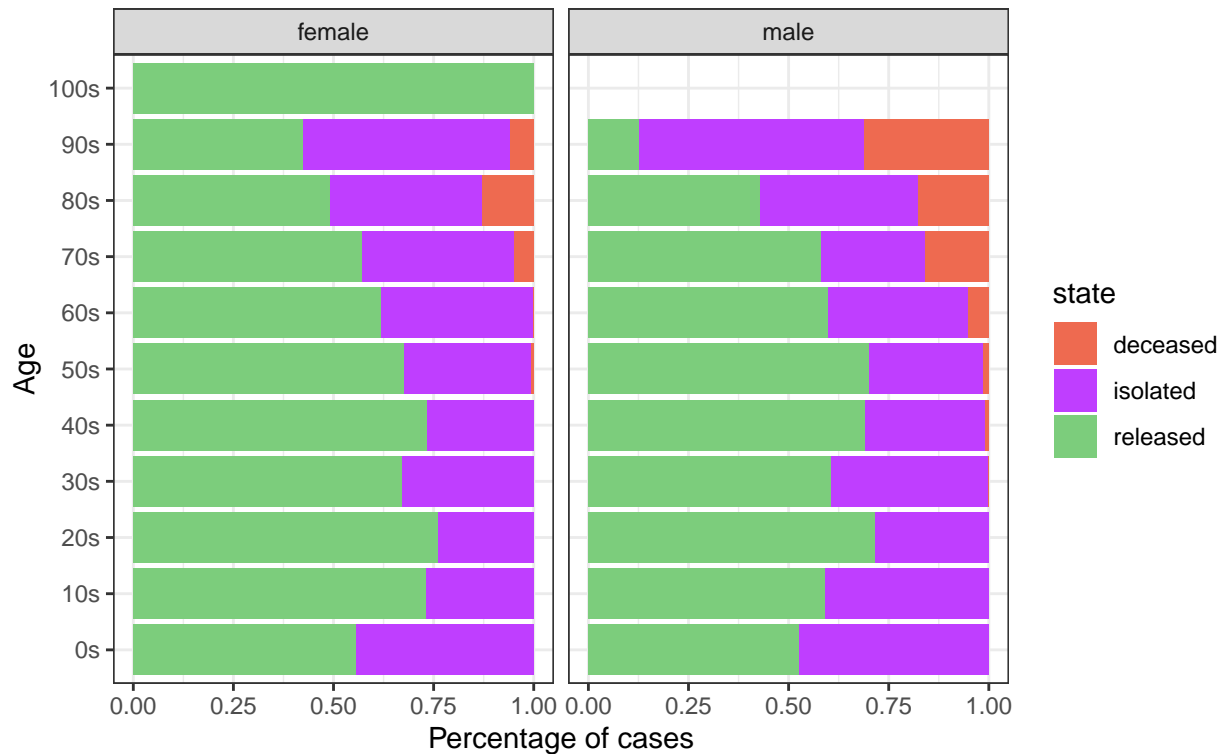
```
patient_state_sex %>%
  filter(is.na(confirmed_date) == F) %>%
  filter(is.na(age) == F) %>%
  group_by(sex, age, state) %>%
  summarise(count = n()) %>%
  ggplot(., aes(x = age, y = count, fill = state)) +
    geom_bar(stat = "identity", position = "fill") +
    labs(x = "Age",
         y = "Percentage of cases",
         title = "Case ratios by age",
         subtitle = "By gender") +
    scale_fill_manual("state", values = c("deceased" = "coral2",
                                           "isolated" = "darkorchid1",
                                           "released" = "palegreen3")) +

    facet_wrap( ~ sex) +
    coord_flip() +
    theme_bw()

## `summarise()` regrouping output by 'sex', 'age' (override with `.groups` argument)
```

Case ratios by age

By gender

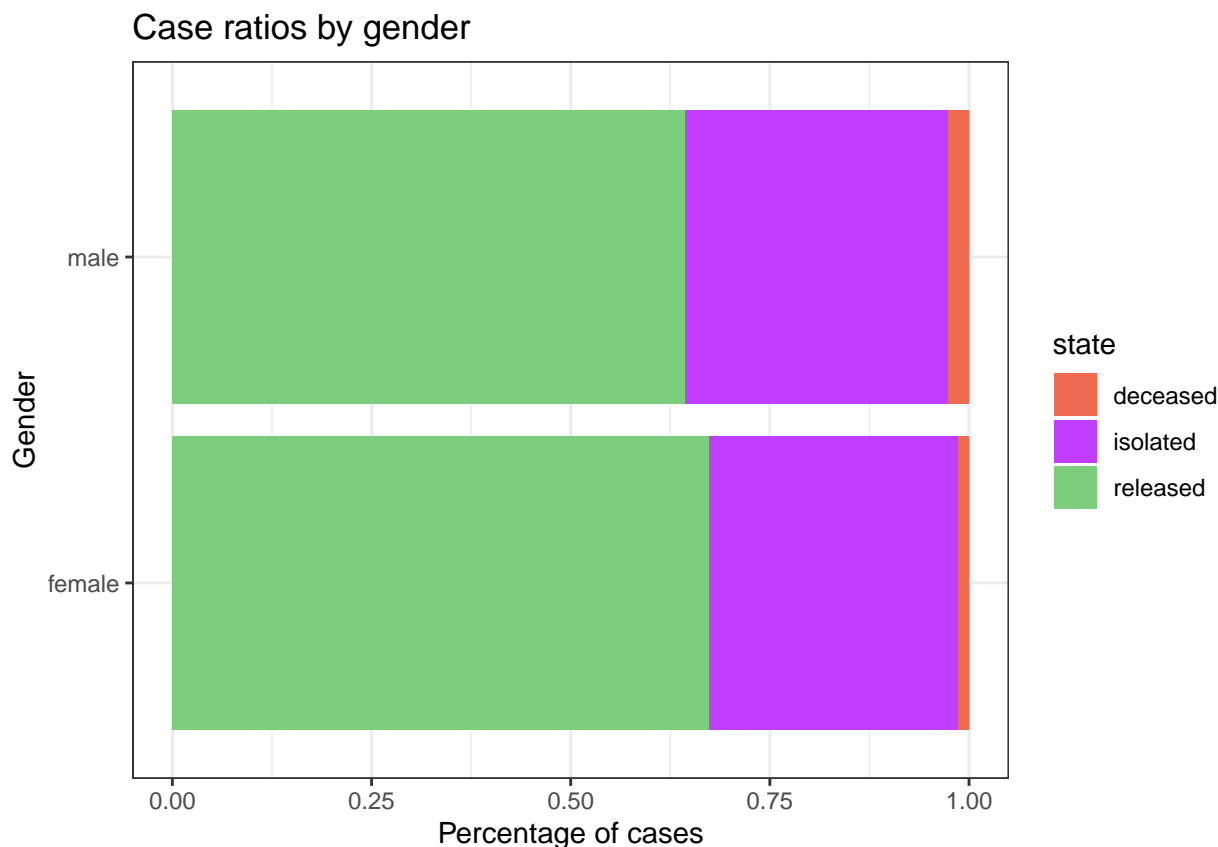


Plot Nr.18: Case ratios by gender

```
patient_state_sex %>%
  filter(is.na(confirmed_date) == F) %>%
  filter(is.na(age) == F) %>%
  group_by(sex, age, state) %>%
  summarise(count = n()) %>%
  ggplot(., aes(x = sex, y = count, fill = state)) +
  geom_bar(stat = "identity", position = "fill") +
  labs(x = "Gender", y = "Percentage of cases", title = "Case ratios by gender") +
  scale_fill_manual("state", values = c("deceased" = "coral2",
                                         "isolated" = "darkorchid1",
                                         "released" = "palegreen3")) +

  coord_flip() +
  theme_bw()
```

`summarise()` regrouping output by 'sex', 'age' (override with `.groups` argument)



Plot Nr.19: Number of cases by gender groups

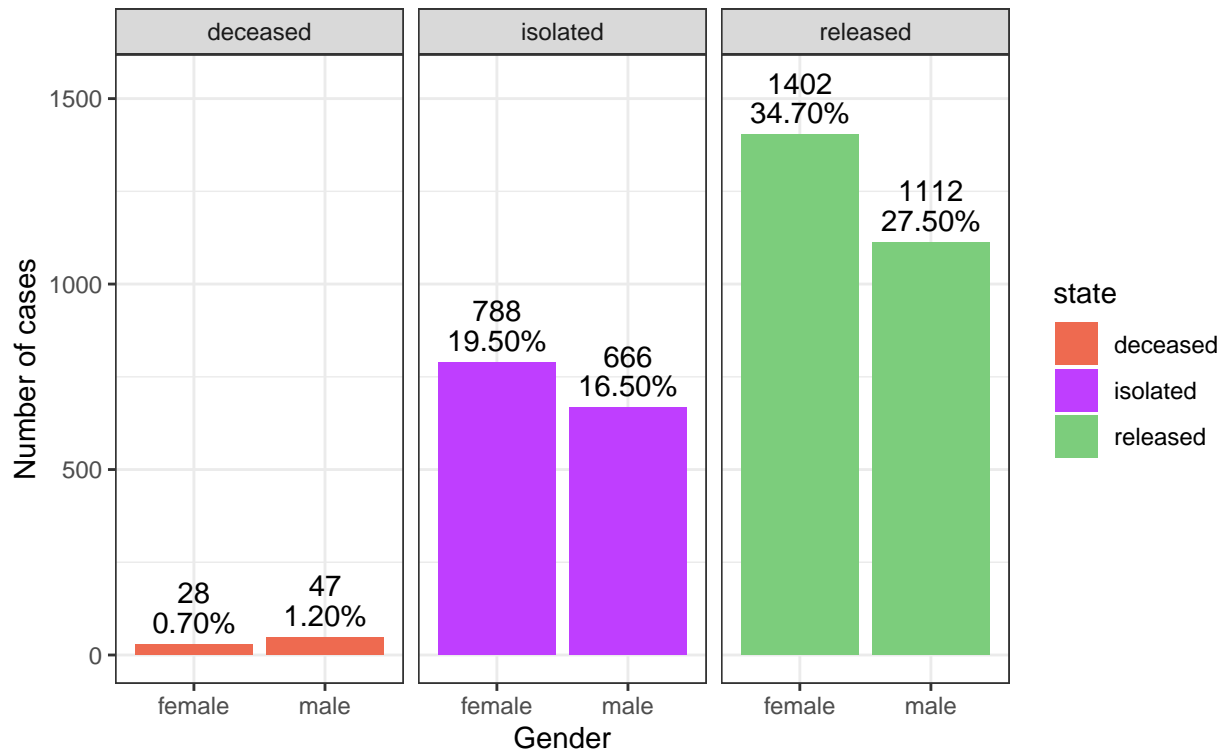
Categorized by state of patients

```
patient_state_sex %>%
  group_by(sex, state) %>%
  summarise(count = n()) %>%
  ggplot(., aes(x = sex, y = count, fill = state)) +
  geom_bar(stat = "identity") +
  geom_text(aes(label = scales::percent(round((count) / sum(count), 3)),
    y = count + 60), size = 4) +
  geom_text(aes(label = count, y = count + 140), size = 4) +
  labs(x = "Gender", y = "Number of cases",
    title = "Number of cases by gender groups",
    subtitle = "Categorized by state of patients") +
  facet_wrap( ~ state) +
  scale_fill_manual("state", values = c("deceased" = "coral2",
    "isolated" = "darkorchid1",
    "released" = "palegreen3")) +
  theme_bw()

## `summarise()` regrouping output by 'sex' (override with `.groups` argument)
```

Number of cases by gender groups

Categorized by state of patients

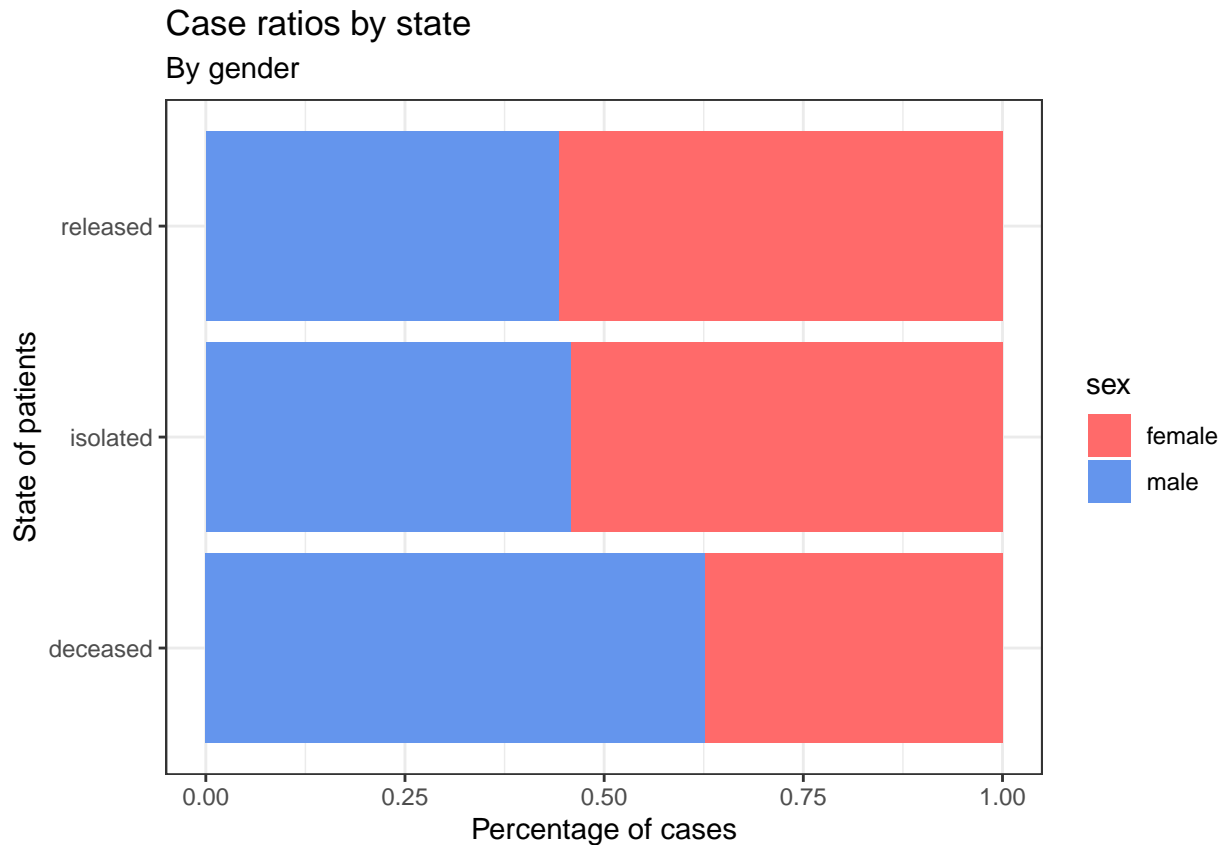


Plot Nr.20: Case ratios by state

By gender

```
patient_state_sex %>%
  filter(is.na(confirmed_date) == F) %>%
  group_by(sex, state) %>%
  summarise(count = n()) %>%
  ggplot(., aes(x = state, y = count, fill = sex)) +
  geom_bar(stat = "identity", position = "fill") +
  labs(x = "State of patients", y = "Percentage of cases",
       title = "Case ratios by state",
       subtitle = "By gender") +
  scale_fill_manual("sex", values = c("female" = "indianred1",
                                       "male" = "cornflowerblue")) +
  coord_flip() +
  theme_bw()
```

`summarise()` regrouping output by 'sex' (override with `.groups` argument)

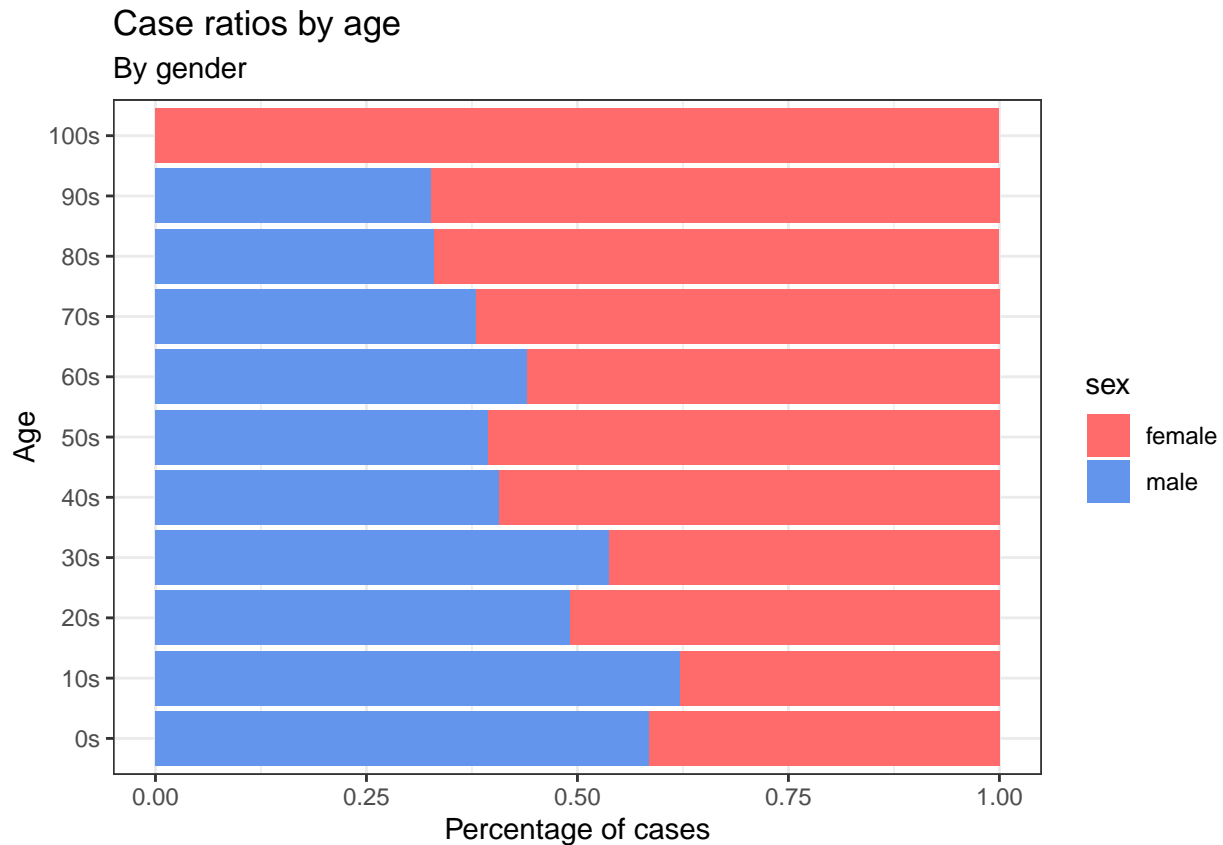


Plot Nr.21: Case ratios by sge

By gender

```
patient_state_sex %>%
  filter(is.na(confirmed_date) == F) %>%
  filter(is.na(age) == F) %>%
  group_by(sex, state, age) %>%
  summarise(count = n()) %>%
  ggplot(., aes(x = age, y = count, fill = sex)) +
  geom_bar(stat = "identity", position = "fill") +
  labs(x = "Age", y = "Percentage of cases",
       title = "Case ratios by age",
       subtitle = "By gender") +
  scale_fill_manual("sex", values = c("female" = "indianred1",
                                     "male" = "cornflowerblue")) +
  coord_flip() +
  theme_bw()

## `summarise()` regrouping output by 'sex', 'state' (override with `.groups` argument)
```

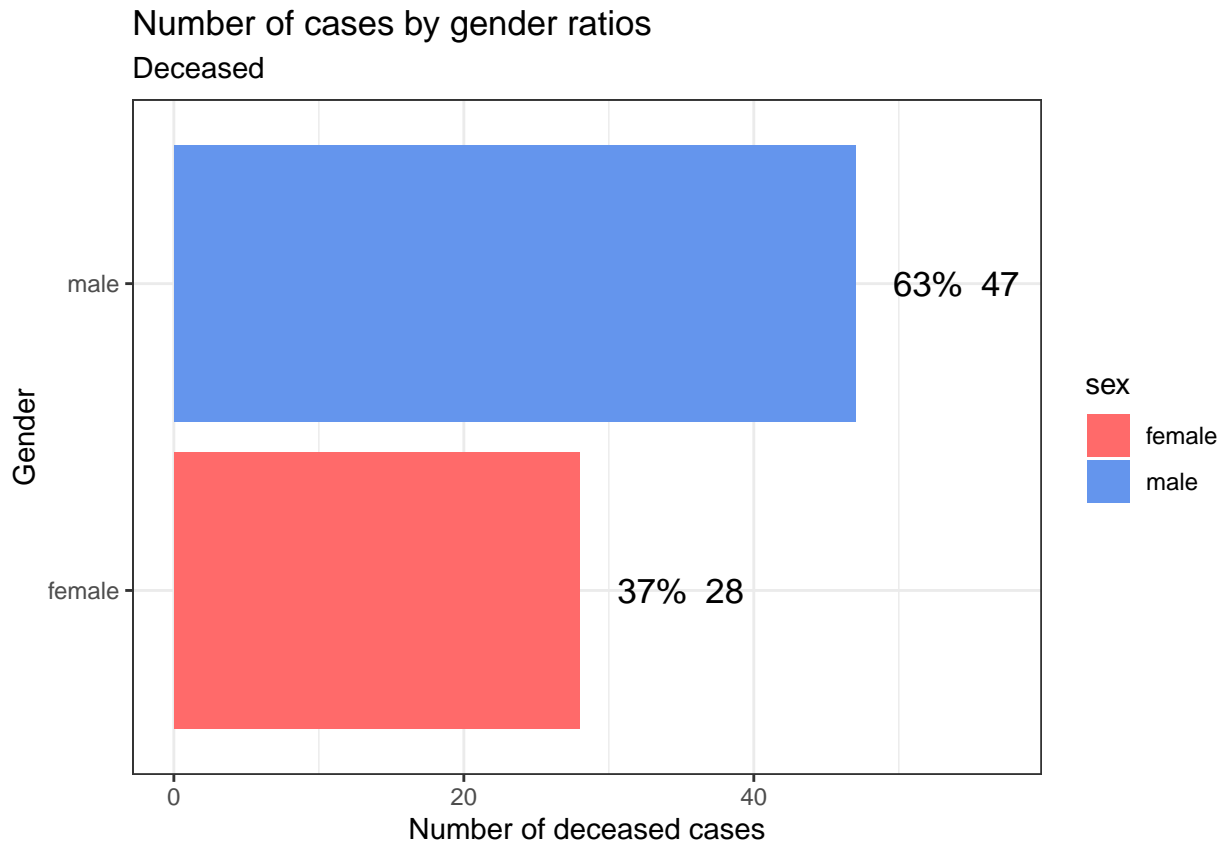


Plot Nr.21: Number of cases by gender ratios

Deceased

```
dead_patients_bysex %>%
  group_by(sex) %>%
  summarise(count = n()) %>%
  ggplot(., aes(x = sex, y = count, fill = sex)) +
  geom_bar(stat = "identity") +
  labs(x = "Gender", y = "Number of deceased cases",
       title = "Number of cases by gender ratios",
       subtitle = "Deceased") +
  geom_text(aes(label = scales::percent(round((count) / sum(count), 3)),
               y = count + 5), size = 4.5) +
  geom_text(aes(label = count, y = count + 10), size = 4.5) +
  coord_flip() +
  scale_fill_manual("sex", values = c("female" = "indianred1",
                                     "male" = "cornflowerblue")) +
  theme_bw()

## `summarise()` ungrouping output (override with `.groups` argument)
```

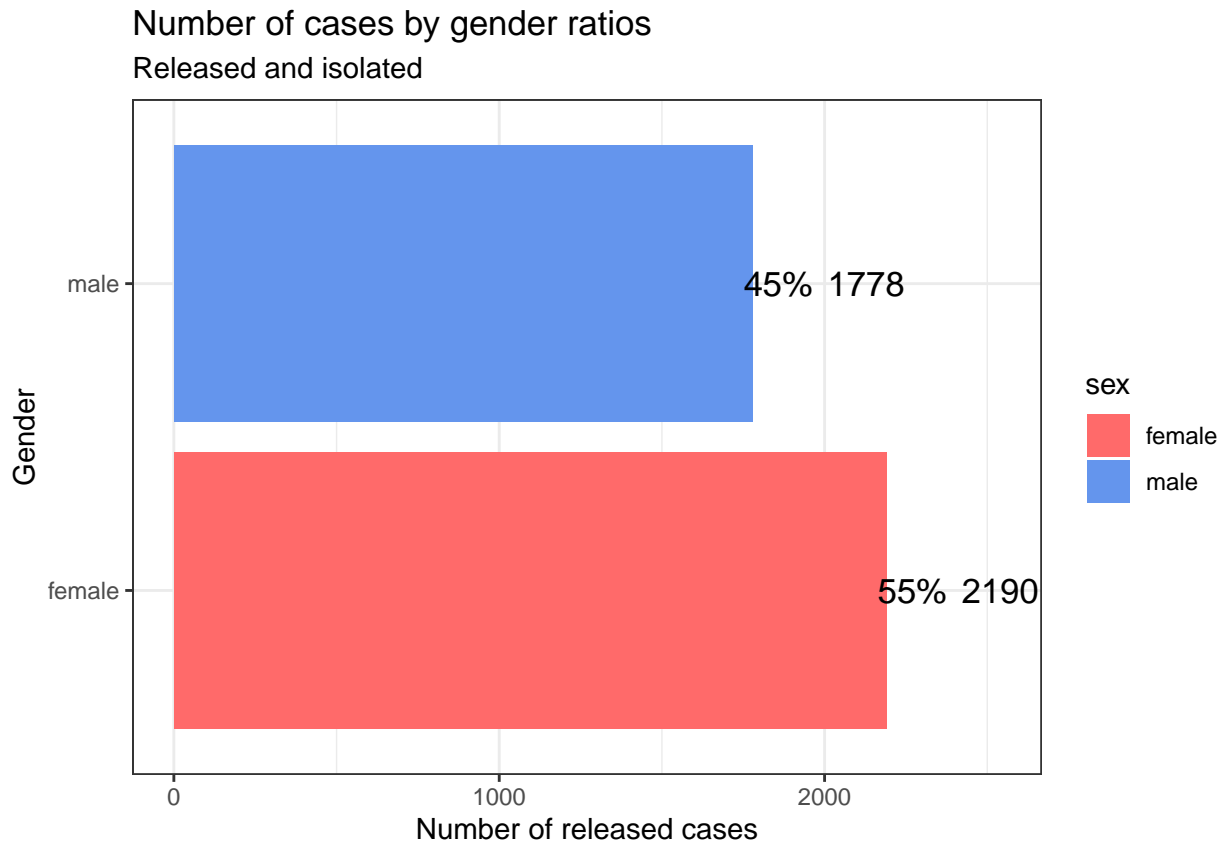


Plot Nr.22: Number of cases by gender ratios

Released and isolated

```
patient_state_sex %>%
  filter(state != "deceased") %>%
  group_by(sex) %>%
  summarise(count = n()) %>%
  ggplot(., aes(x = sex, y = count, fill = sex)) +
  geom_bar(stat = "identity") +
  labs(x = "Gender", y = "Number of released cases",
       title = "Number of cases by gender ratios",
       subtitle = "Released and isolated") +
  geom_text(aes(label = scales::percent(round((count) / sum(count), 3)),
               y = count + 80), size = 4.5) +
  geom_text(aes(label = count, y = count + 350), size = 4.5) +
  coord_flip() +
  scale_fill_manual("sex", values = c("female" = "indianred1",
                                       "male" = "cornflowerblue")) +
  theme_bw()
```

```
## `summarise()` ungrouping output (override with `.groups` argument)
```

ANALYSIS ON GOVERNMENT POLICIES

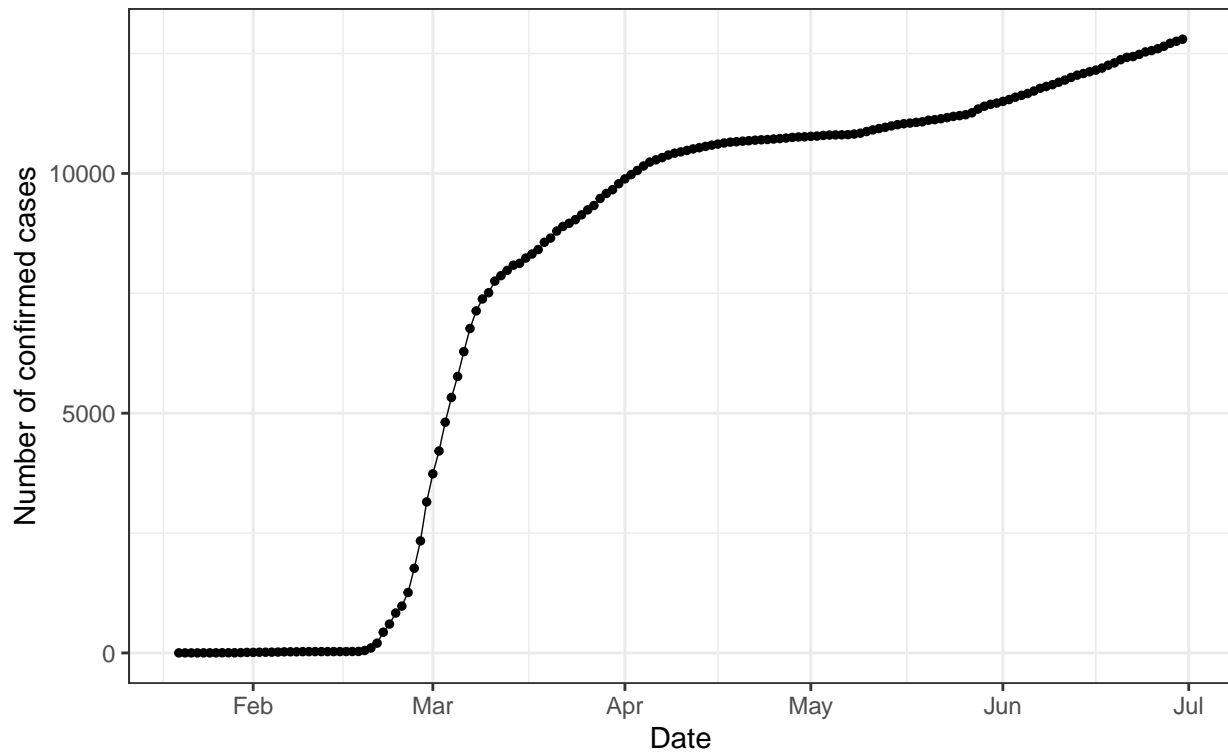
```
#value for zero row
Time[, confirmed_per_day := confirmed - shift(confirmed, fill = 0, type = 'lag')]
```

Plot Nr.23: Accumulated number of confirmed cases over given time period
(2020-01-20 to 2020-06-30)

```
plot_1 <- ggplot(Time, aes(date, confirmed)) +
  geom_point(size = 1) +
  geom_line(size = 0.25) +
  scale_color_brewer(palette='Set3') +
  theme(axis.text.x = element_text(angle = 90, vjust = 0.5, hjust=1)) +
  labs(x = 'Date', y = 'Number of confirmed cases',
       title = 'Accumulated number of confirmed cases over given time period',
       subtitle = '(2020-01-20 to 2020-06-30)') +
  theme_bw()

plot_1
```

Accumulated number of confirmed cases over given time period
(2020-01-20 to 2020-06-30)

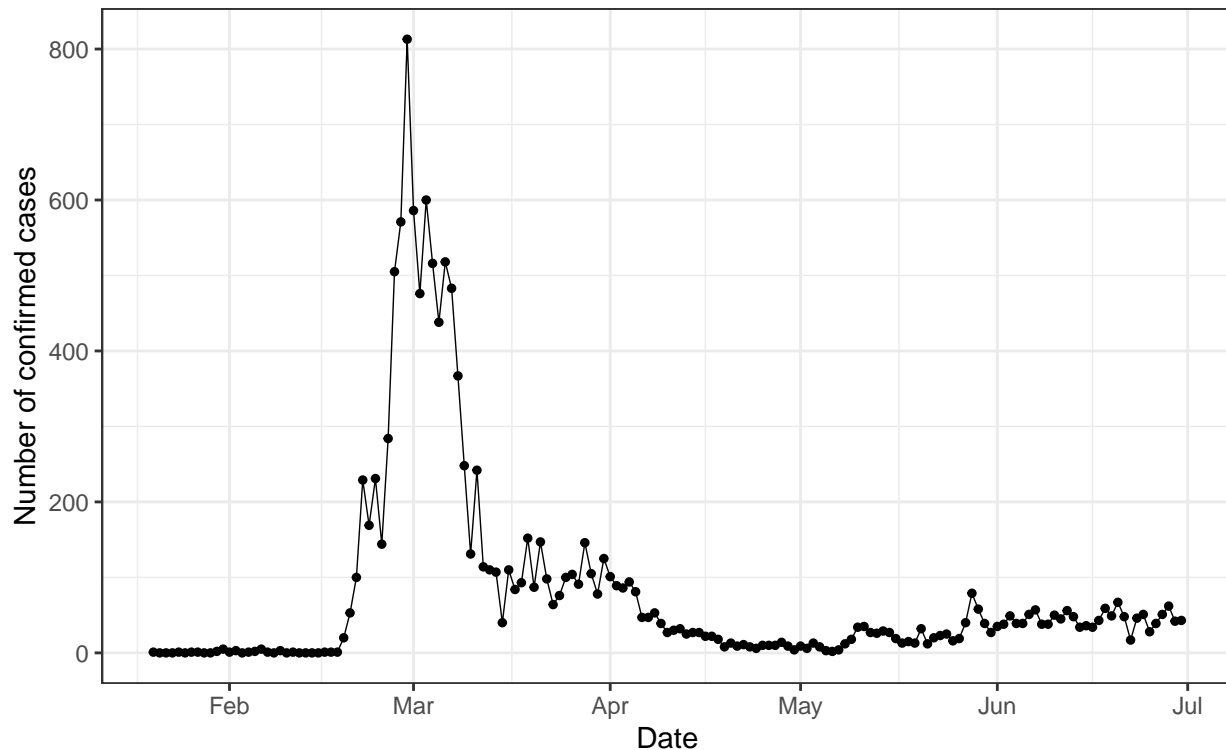


Plot Nr.24: Number of confirmed cases per day over given time period
(2020-01-20 to 2020-06-30)

```
plot_2 <- ggplot(Time, aes(date, confirmed_per_day)) +  
  geom_point(size = 1) +  
  geom_line(size = 0.25) +  
  theme(axis.text.x = element_text(angle = 90, vjust = 0.5, hjust=1)) +  
  labs(x = 'Date', y = 'Number of confirmed cases',  
       title = 'Number of confirmed cases per day over given time period',  
       subtitle = '(2020-01-20 to 2020-06-30)') +  
  theme_bw()
```

plot_2

Number of confirmed cases per day over given time period
(2020-01-20 to 2020-06-30)



Working on Policy

Rearranging and merging data

```
policy_resaped <- Policy[, date := start_date] #variable date to identify observations
policy_merged <- merge(Time, policy_resaped, by='date', group_by='date',all=TRUE) %>%
  as.data.table(policy_merged)
```

```
policy_sum <- Policy[, .(count = .N), by=date]
policy_sum[, count_accumulated := cumsum(count)]
```

```
policy_sum <- merge(policy_sum, Time, by='date', group_by='date', all=TRUE)
```

Plot Nr.25: Comparison of Accumulated Confirmed Cases and

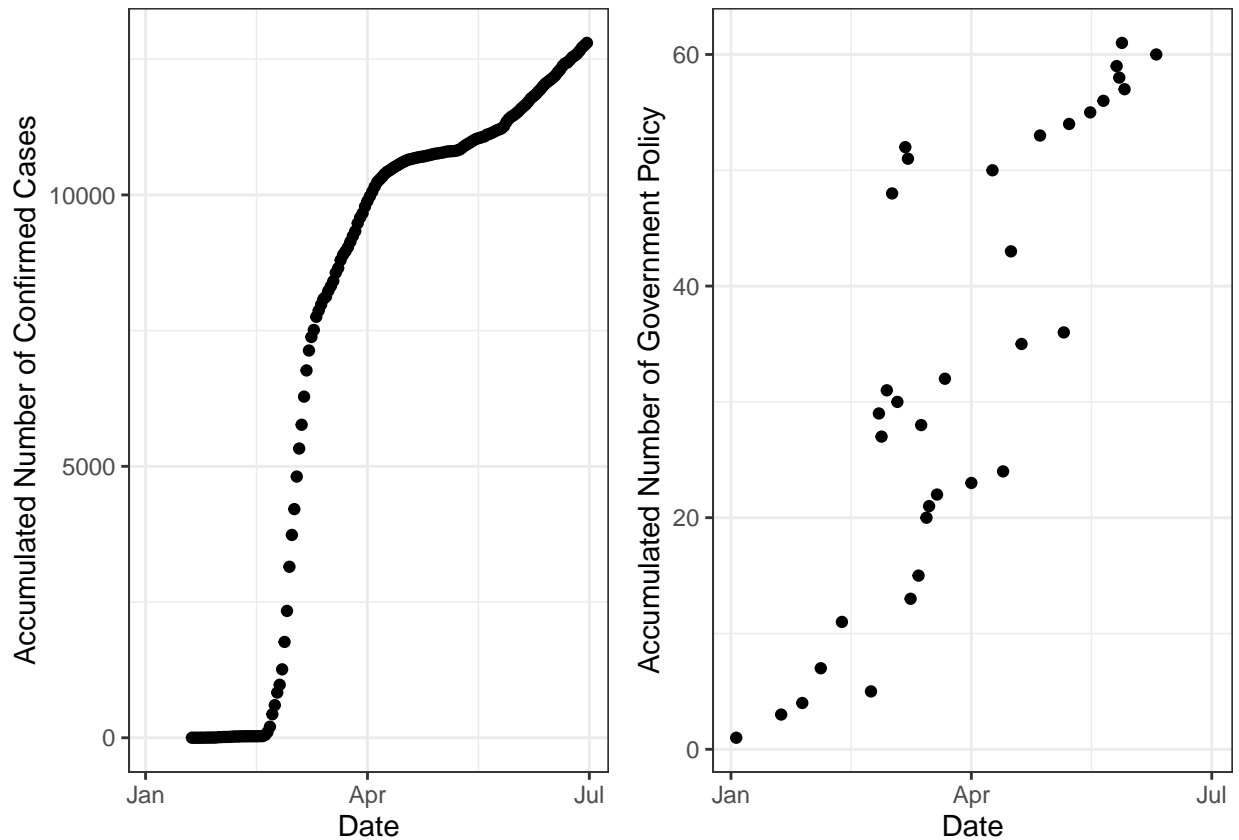
Accumulated Number of Government Policy ### (2020-01-20 to 2020-06-30)

```
plot1.1 <- ggplot(policy_sum, aes(date, confirmed)) +
  geom_point() +
  labs(x = 'Date', y = 'Accumulated Number of Confirmed Cases') +
  theme_bw()

plot3.1 <- ggplot(policy_sum, aes(date, count_accumulated)) +
  geom_point() +
  labs(x = 'Date', y = 'Accumulated Number of Government Policy') +
  theme_bw()

grid.arrange(plot1.1, plot3.1, ncol=2)
```

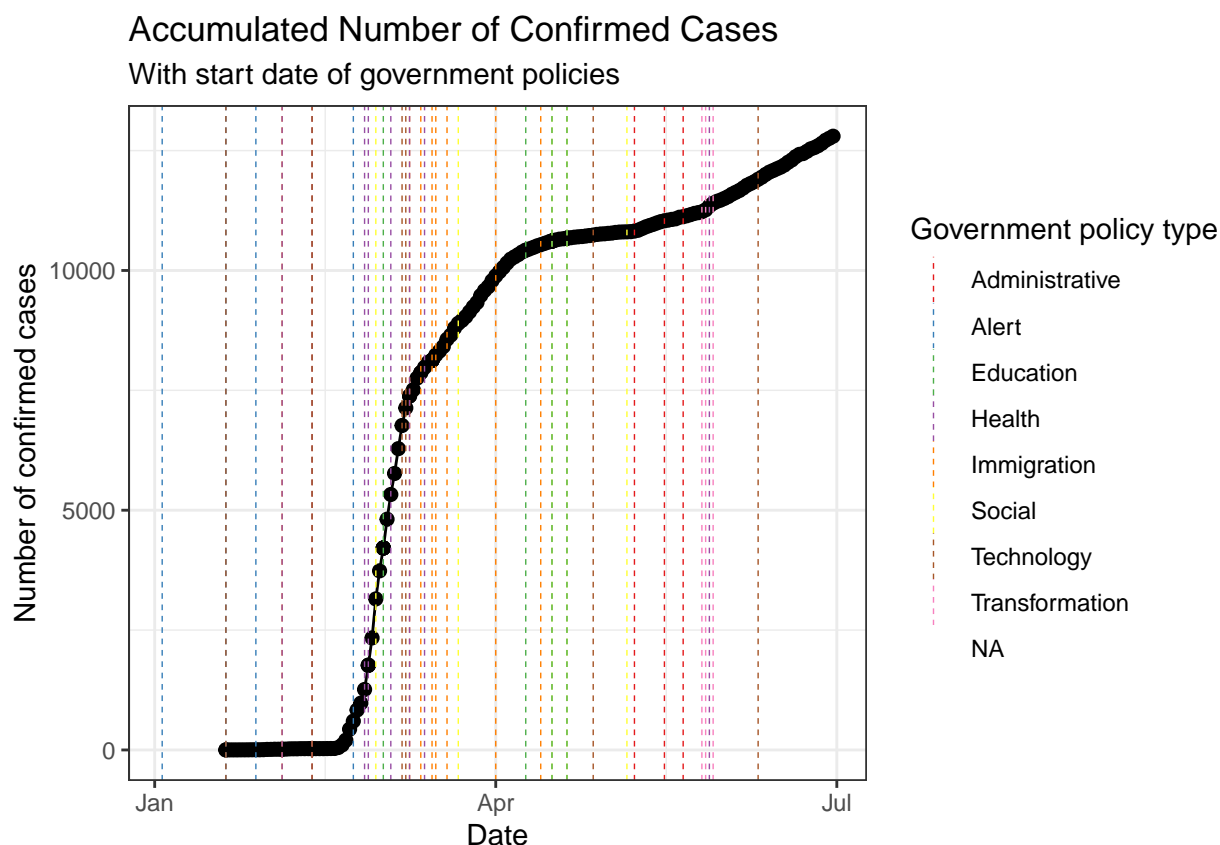
```
## Warning: Removed 1 rows containing missing values (geom_point).
## Warning: Removed 129 rows containing missing values (geom_point).
```



Plot Nr.26: Show events of government policies

```
ggplot(policy_merged, aes(date, confirmed)) +
  geom_line() +
  geom_point(size=2) +
  geom_vline(aes(xintercept = date, color = type), size = 0.25, linetype = 'dashed') +
  labs(color = 'Government policy type') +
  scale_color_brewer(palette='Set1') +
  theme(axis.text.x = element_text(angle = 90, vjust = 0.5, hjust=1)) +
  labs(x = 'Date', y = 'Number of confirmed cases',
       title = "Accumulated Number of Confirmed Cases",
       subtitle = "With start date of government policies") +
  theme_bw()
```

```
## Warning: Removed 1 row(s) containing missing values (geom_path).
## Warning: Removed 1 rows containing missing values (geom_point).
```

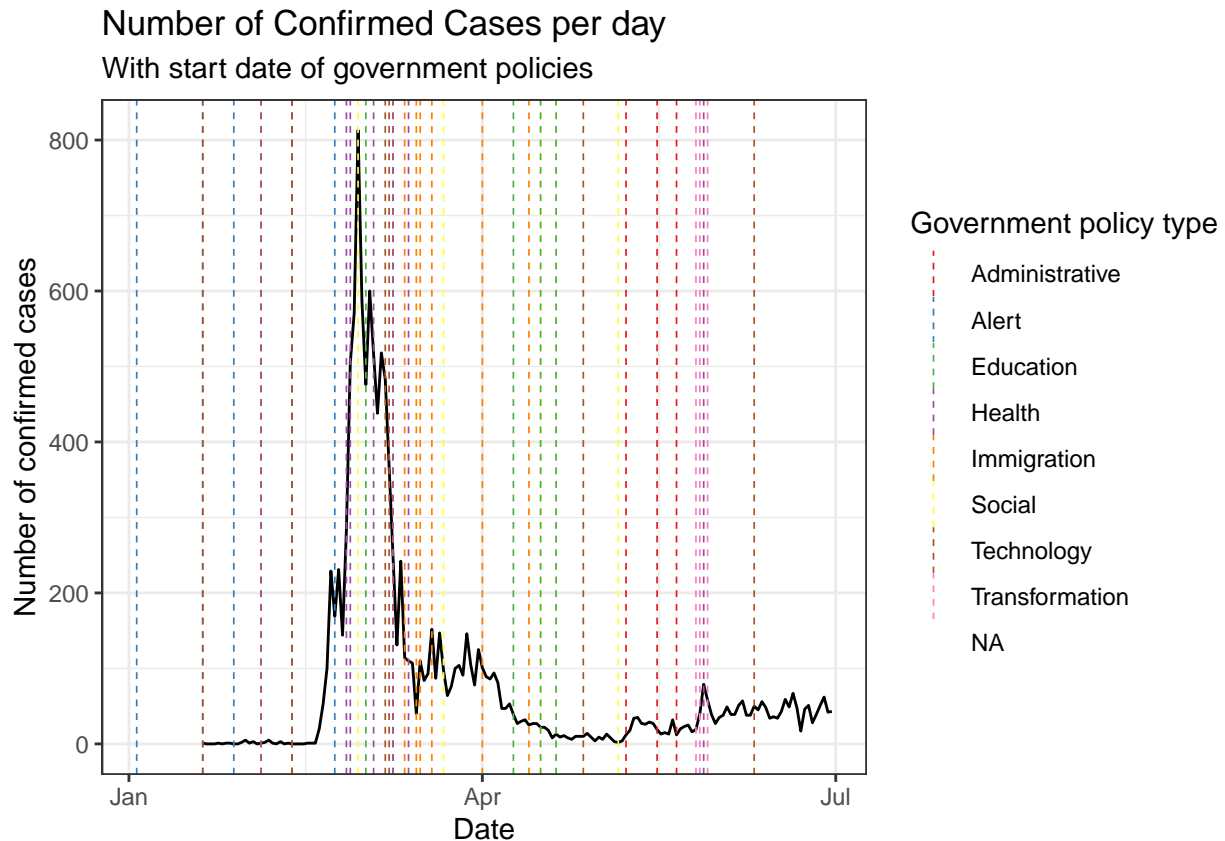


Plot Nr.27: Show events of government policies

```
plot_3 <- ggplot(policy_merged, aes(date, confirmed_per_day)) +
  geom_line() +
  geom_vline(aes(xintercept=date, color=type), size=0.3, linetype = 'dashed') +
  labs(color = 'Government policy type') +
  scale_color_brewer(palette='Set1') +
  theme(axis.text.x = element_text(angle = 90, vjust = 0.5, hjust=1)) +
  labs(x = 'Date', y = 'Number of confirmed cases',
       title = "Number of Confirmed Cases per day",
       subtitle = "With start date of government policies") +
  theme_bw()
```

plot_3

Warning: Removed 1 row(s) containing missing values (geom_path).



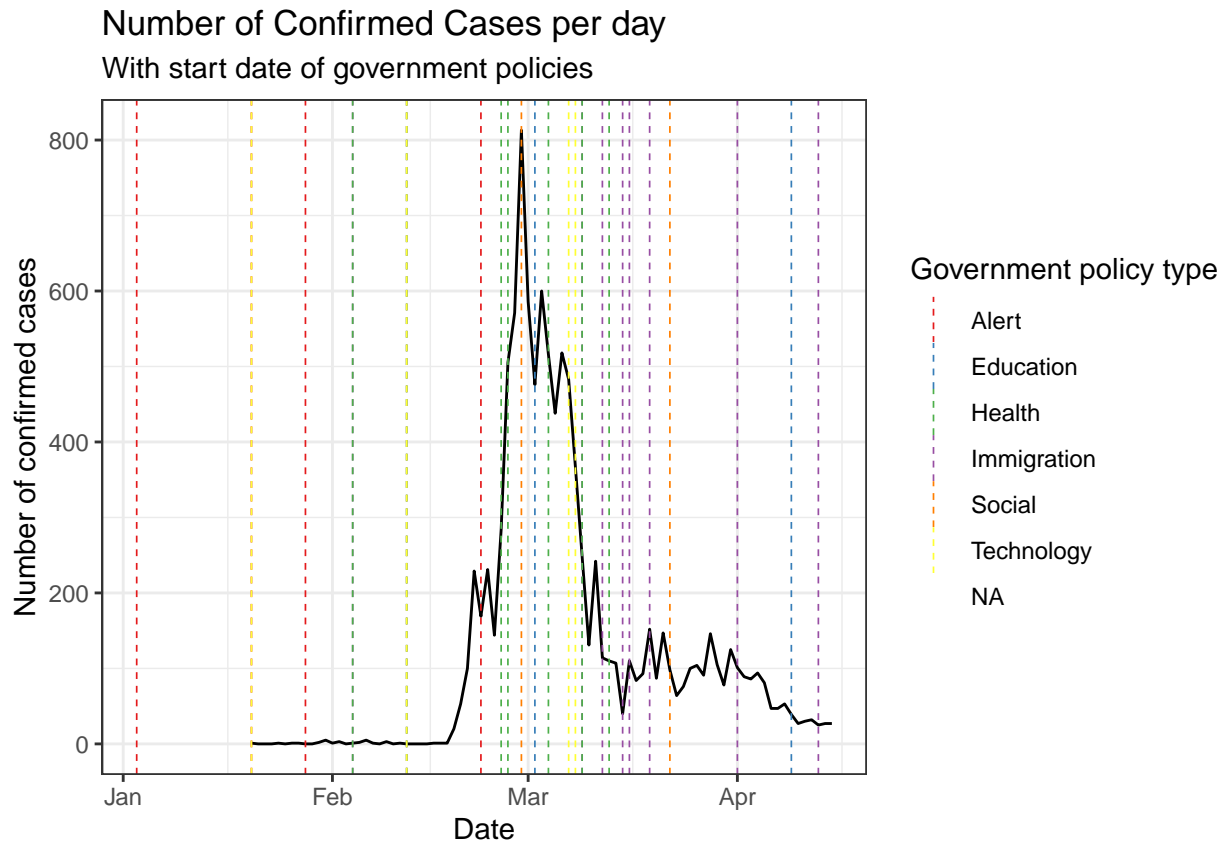
Plot Nr.28: Show events of government policies

For the first big peak

```
policy_merged_resshaped1 <- policy_merged[date >= as.Date("2020-01-03") &
  date <= as.Date("2020-04-15"),
  .(date,confirmed_per_day,
    type,detail,start_date)]

#for a better overview of the data
ggplot(policy_merged_resshaped1, aes(date, confirmed_per_day)) +
  geom_line() +
  geom_vline(aes(xintercept = date, color = type), size = 0.3, linetype = 'dashed') +
  labs(color = 'Government policy type') +
  scale_color_brewer(palette='Set1') +
  theme(axis.text.x = element_text(angle = 90, vjust = 0.5, hjust=1)) +
  labs(x = 'Date', y = 'Number of confirmed cases',
    title = "Number of Confirmed Cases per day",
    subtitle = "With start date of government policies") +
  theme_bw()
```

Warning: Removed 1 row(s) containing missing values (geom_path).



Plot Nr.29: Alert Level of Policies

```

alert_level <- Policy[type == 'Alert']
alert_level[,date] #View start_date of each Alert Level

## [1] "2020-01-03" "2020-01-20" "2020-01-28" "2020-02-23"

plot_4 <- ggplot(policy_merged, aes(date, confirmed_per_day)) +
  #Geom_rect for Level 1
  geom_rect(aes(xmin = as.Date("2020-01-03", "%Y-%m-%d"),
    xmax = as.Date("2020-01-19", "%Y-%m-%d"),
    ymin = -Inf, ymax = Inf, fill = 'Alert Level 1'),
    alpha = 0.03) +
  #Geom_rect for Level 2
  geom_rect(aes(xmin = as.Date("2020-01-20", "%Y-%m-%d"),
    xmax = as.Date("2020-01-27", "%Y-%m-%d"),
    ymin = -Inf, ymax = Inf, fill = 'Alert Level 2'),
    alpha = 0.03) +
  #Geom_rect for Level 3
  geom_rect(aes(xmin = as.Date("2020-01-28", "%Y-%m-%d"),
    xmax = as.Date("2020-02-22", "%Y-%m-%d"),
    ymin = -Inf, ymax = Inf, fill = 'Alert Level 3'),
    alpha = 0.03) +
  #Geom_rect for Level 4
  geom_rect(aes(xmin = as.Date("2020-02-23", "%Y-%m-%d"),
    xmax = as.Date("2020-06-30", "%Y-%m-%d"),
    ymin = -Inf, ymax = Inf, fill = 'Alert Level 4'),

```

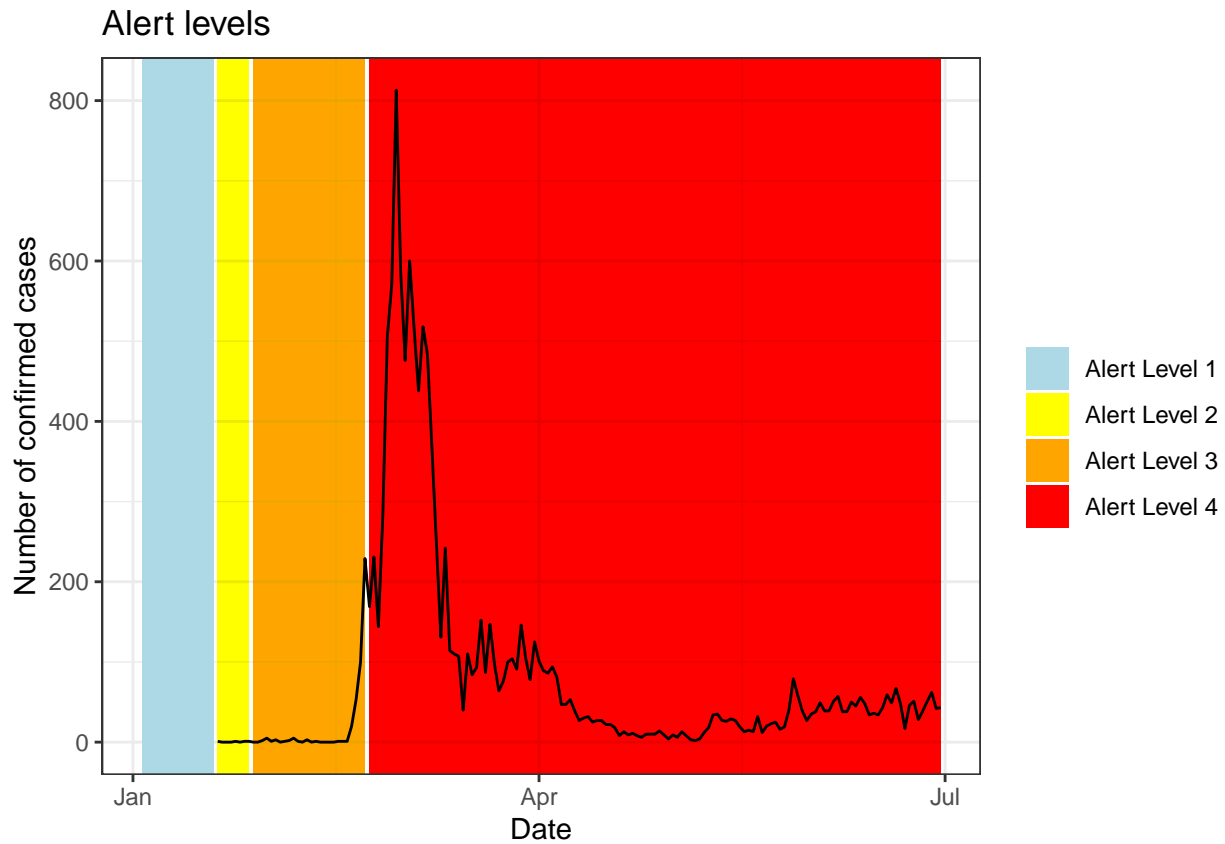
```

        alpha = 0.03) +
  geom_line() +
  scale_fill_manual(name = '',
                    values = c('lightblue','yellow', 'orange','red'),
                    guide = guide_legend(override.aes = list(alpha = 1))) +
  theme(axis.text.x = element_text(angle = 90, vjust = 0.5, hjust=1)) +
  labs(x = 'Date', y = 'Number of confirmed cases') +
  ggtitle('Alert levels')+ theme_bw()

```

plot_4

Warning: Removed 1 row(s) containing missing values (geom_path).



Plot Nr.30: Policies on Administration

```
Policy[type=='Administrative', 4:6] # view all policy titles on administration
```

```

##                               gov_policy
## 1:                            Close bars and clubs
## 2: local government Administrative orders
## 3:                            Close karaoke
##
## 1:
## 2: Gathering limited administrative orders for nursing homes, nursing homes, elderly day care centers
## 3:                            Ban on gatherings
##   start_date
## 1: 2020-05-08

```



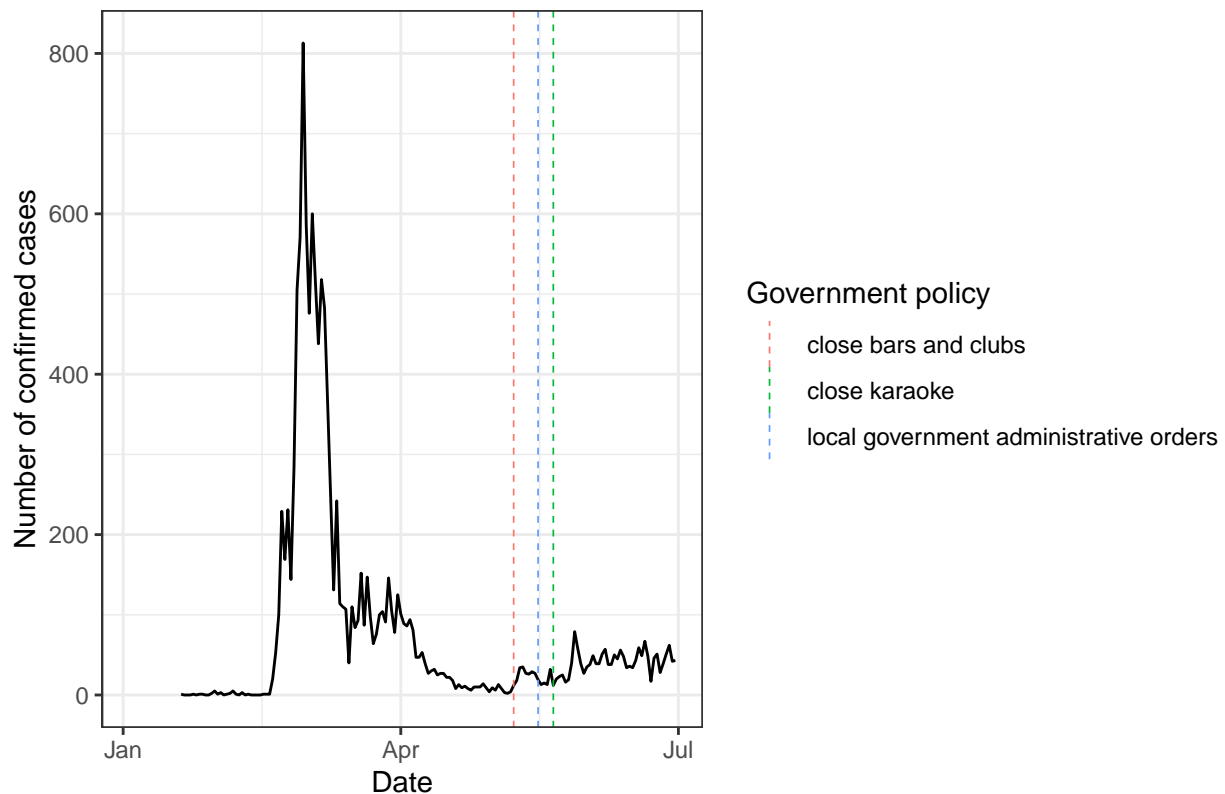
```
## 2: 2020-05-16
```

```
## 3: 2020-05-21
```

```
ggplot(policy_merged, aes(date, confirmed_per_day)) +  
  geom_line() +  
  geom_vline(data = subset(policy_merged, type == 'Administrative'), # filter data source  
    aes(xintercept = start_date,  
      color = c('close bars and clubs',  
        'local government administrative orders',  
        'close karaoke')), size = 0.3, linetype = 'dashed') +  
  labs(color = 'Government policy') +  
  theme(axis.text.x = element_text(angle = 90, vjust = 0.5, hjust=1)) +  
  labs(x = 'Date', y = 'Number of confirmed cases') +  
  ggtitle('Policies on Administration') +  
  theme_bw()
```

```
## Warning: Removed 1 row(s) containing missing values (geom_path).
```

Policies on Administration



Plot Nr.31: Policies on Education

```
Policy[type=='Education', 4:6] %>%  
  group_by(start_date) #view all policy titles on education
```

```
## # A tibble: 15 x 3
```

```
## # Groups:   start_date [4]
```

gov_policy	detail	start_date
<chr>	<chr>	<date>
1 School Closure	Daycare Center for Children	2020-03-02

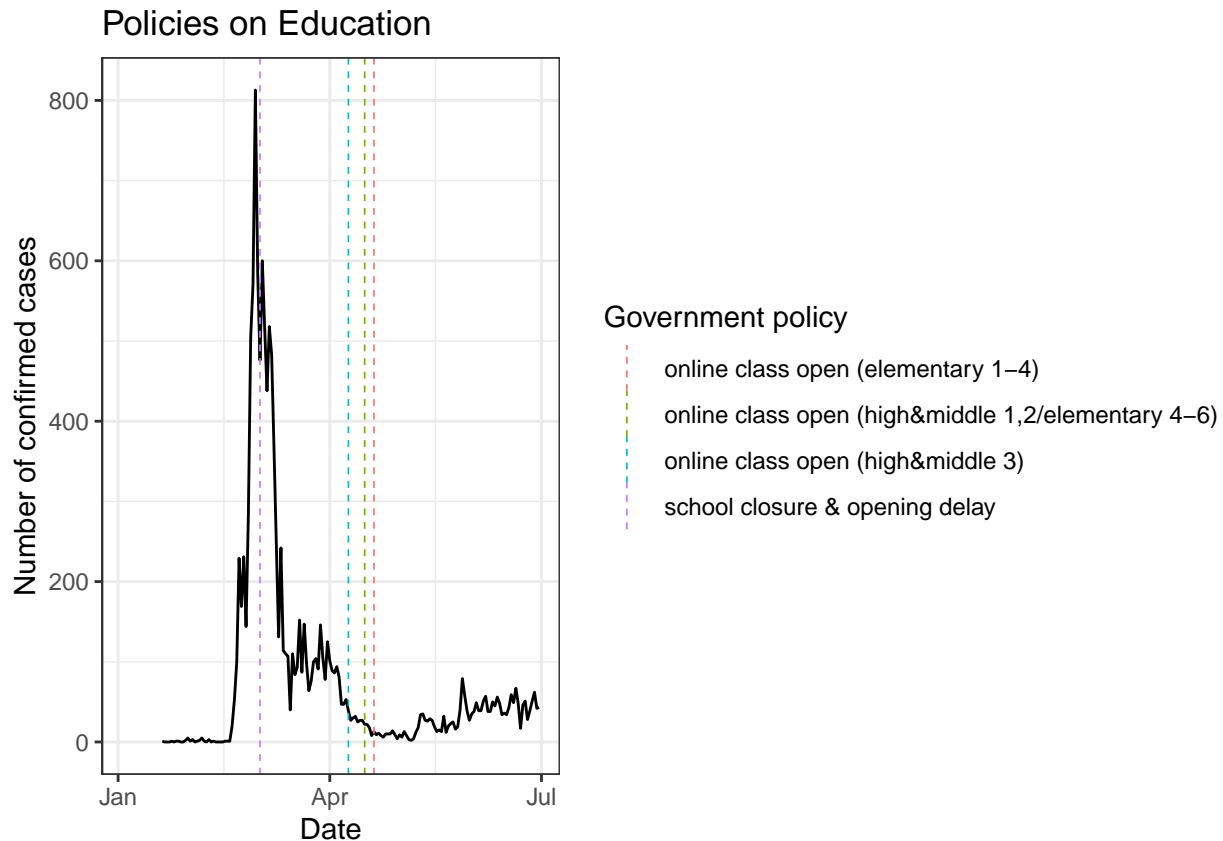
```
## 2 School Opening Delay Kindergarten 2020-03-02
## 3 School Opening Delay High School 2020-03-02
## 4 School Opening Delay Middle School 2020-03-02
## 5 School Opening Delay Elementary School 2020-03-02
## 6 School Opening with Online Class High School (3rd grade) 2020-04-09
## 7 School Opening with Online Class High School (2nd grade) 2020-04-16
## 8 School Opening with Online Class High School (1st grade) 2020-04-16
## 9 School Opening with Online Class Middle School (3rd grade) 2020-04-09
## 10 School Opening with Online Class Middle School (2nd grade) 2020-04-16
## 11 School Opening with Online Class Middle School (1st grade) 2020-04-16
## 12 School Opening with Online Class Elementary School (5th ~ 6th gra~ 2020-04-16
## 13 School Opening with Online Class Elementary School (4th grade) 2020-04-16
## 14 School Opening with Online Class Elementary School (3rd grade) 2020-04-20
## 15 School Opening with Online Class Elementary School (1st ~ 2nd gra~ 2020-04-20
```

```
#summarize policy titles for each start_date
```

```
policy_education <- Policy[type=='Education', .(date=unique(start_date))]  
policy_education[,gov_policy_grouped := c('school closure & opening delay',  
                                          'online class open (high&middle 3)',  
                                          'online class open (high&middle  
1,2/elementary 4-6)',  
                                          'online class open (elementary 1-4)')]
```

```
ggplot(policy_merged, aes(date, confirmed_per_day)) +  
  geom_line() +  
  geom_vline(data = subset(policy_education, date==date), # filter data source  
             aes(xintercept = date,  
                 color=c('school closure & opening delay',  
                          'online class open (high&middle 3)',  
                          'online class open (high&middle 1,2/elementary 4-6)',  
                          'online class open (elementary 1-4)'), size = 0.3,  
                 linetype = 'dashed') +  
  labs(color = 'Government policy') +  
  theme(axis.text.x = element_text(angle = 90, vjust = 0.5, hjust=1)) +  
  labs(x = 'Date', y = 'Number of confirmed cases') +  
  ggtitle('Policies on Education') +  
  theme_bw()
```

```
## Warning: Removed 1 row(s) containing missing values (geom_path).
```



Plot Nr.32: Policies on Public Health

```
Policy[type=='Health', 4:6] %>% group_by(start_date) #view all policy titles on health
```

```
## # A tibble: 10 x 3
## # Groups:   start_date [8]
##   gov_policy          detail          start_date
##   <chr>              <chr>          <date>
## 1 Emergency Use Authorization ~ 1st EUA      2020-02-04
## 2 Emergency Use Authorization ~ 2nd EUA      2020-02-12
## 3 Emergency Use Authorization ~ 3rd EUA      2020-02-27
## 4 Emergency Use Authorization ~ 4th EUA      2020-02-27
## 5 Emergency Use Authorization ~ 5th EUA      2020-03-13
## 6 Drive-Through Screening Cent~ by Local Government 2020-02-26
## 7 Drive-Through Screening Cent~ Standard Operating Procedures 2020-03-04
## 8 Mask Distribution          Public-Sale      2020-02-27
## 9 Mask Distribution          5-day Rotation System 2020-03-09
## 10 Extends Tightened Quarantine~ Gov't Extends Tightened Quarantine ~ 2020-05-28
```

```
policy_health <- Policy[type=='Health',.(start_date=first(start_date)),
                          by=gov_policy] #only first policy implementation
```

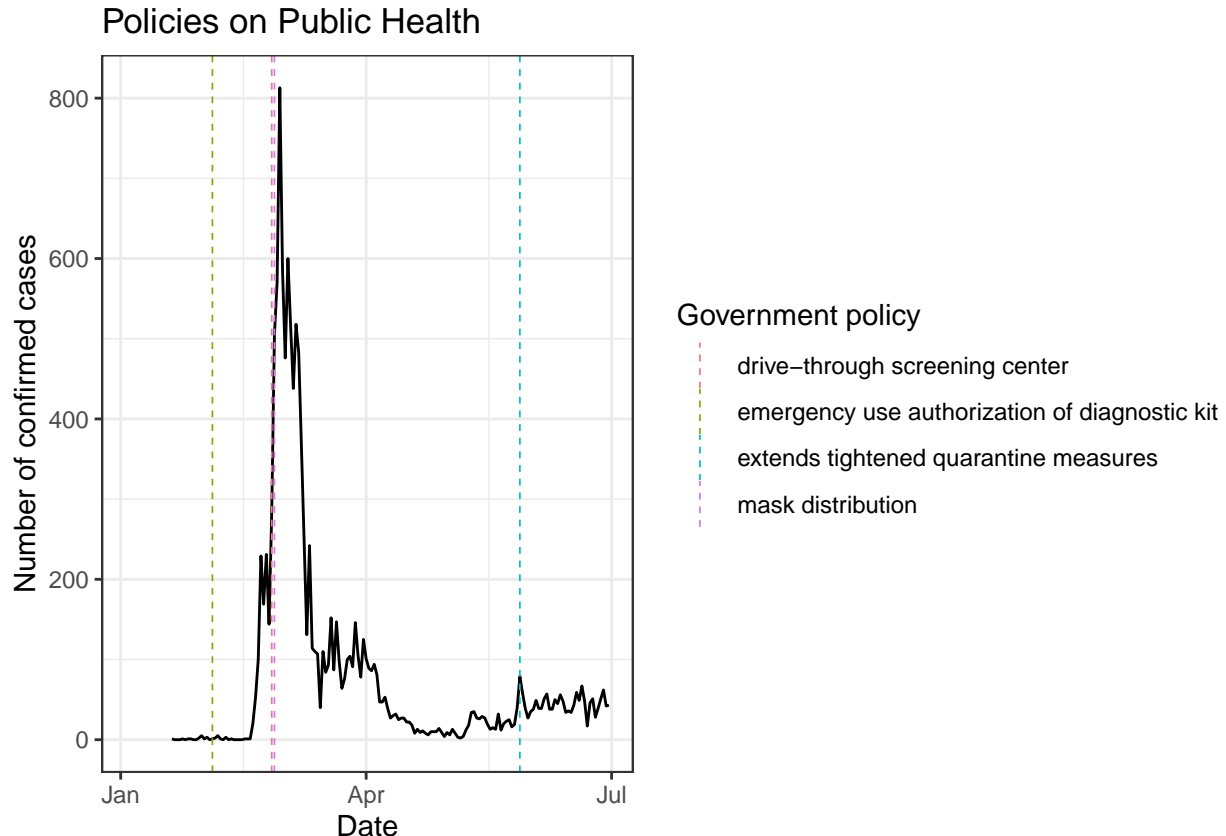
```
ggplot(policy_merged, aes(date, confirmed_per_day)) +
  geom_line() +
  geom_vline(data = policy_health,
            aes(xintercept = start_date,
                color = c('emergency use authorization of diagnostic kit',
```

```

        'drive-through screening center', 'mask distribution',
        'extends tightened quarantine measures' )), size = 0.3,
        linetype = 'dashed') +
  labs(color = 'Government policy') +
  theme(axis.text.x = element_text(angle = 90, vjust = 0.5, hjust=1)) +
  labs(x = 'Date', y = 'Number of confirmed cases') +
  ggtitle('Policies on Public Health') +
  theme_bw()

```

Warning: Removed 1 row(s) containing missing values (geom_path).



Plot Nr.33: Policies on Immigration

```

Policy[type=='Immigration', 4:6] %>%
  group_by(start_date) #view all policy titles on health

```

```

## # A tibble: 15 x 3
## # Groups:   start_date [9]
##   gov_policy          detail          start_date
##   <chr>              <chr>          <date>
## 1 Special Immigration from China    2020-02-04
## 2 Special Immigration from Hong Kong 2020-02-12
## 3 Special Immigration from Macau     2020-02-12
## 4 Special Immigration from Japan     2020-03-09
## 5 Special Immigration from Italy     2020-03-12
## 6 Special Immigration from Iran      2020-03-12
## 7 Special Immigration from France    2020-03-15

```

```

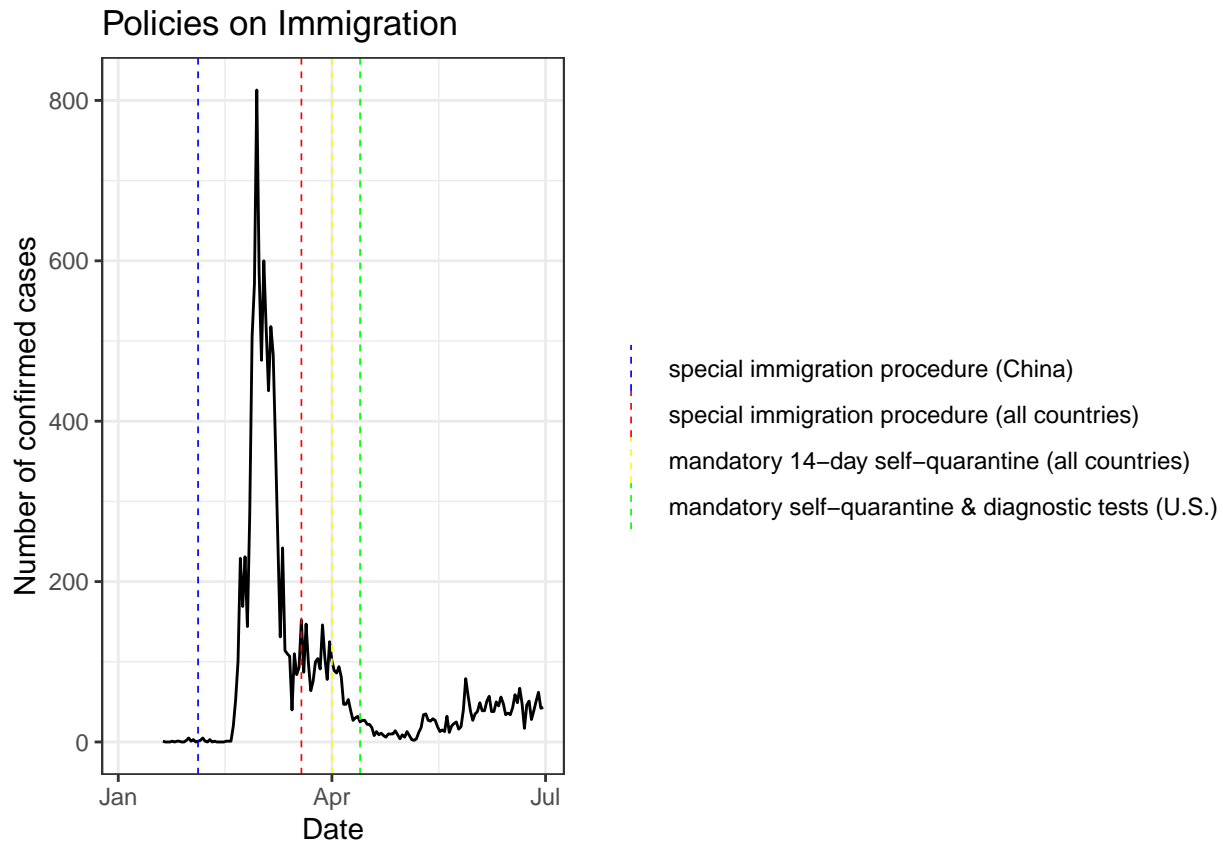
## 8 Special Immigration Procedure           from Germany           2020-03-15
## 9 Special Immigration Procedure           from Spain              2020-03-15
## 10 Special Immigration Procedure          from U.K.                 2020-03-15
## 11 Special Immigration Procedure          from Netherlands         2020-03-15
## 12 Special Immigration Procedure          from Europe               2020-03-16
## 13 Special Immigration Procedure          from all the countri~     2020-03-19
## 14 Mandatory 14-day Self-Quarantine       from all the countri~     2020-04-01
## 15 Mandatory Self-Quarantine & Diagnostic Tes~ from U.S.                2020-04-13

policy_immigration <- Policy[detail=='from China'|detail=='from all the countries'|
                             detail=='from U.S.',4:6] #select representative policies

ggplot(policy_merged, aes(date, confirmed_per_day)) +
  geom_line() +
  geom_vline(data = policy_immigration,
             aes(xintercept = start_date, color = c('blue','red','yellow','green')),
             size = 0.3, linetype = 'dashed') +
  scale_color_identity(name='',
                       breaks=c('blue','red','yellow','green'),
                       labels=c('special immigration procedure (China)',
                                'special immigration procedure (all countries)',
                                'mandatory 14-day self-quarantine (all countries)',
                                'mandatory self-quarantine & diagnostic tests (U.S.)'),
                       guide = 'legend') +
  theme(axis.text.x = element_text(angle = 90, vjust = 0.5, hjust=1)) +
  labs(x = 'Date', y = 'Number of confirmed cases') +
  ggtitle('Policies on Immigration') +
  theme_bw()

## Warning: Removed 1 row(s) containing missing values (geom_path).

```



Time of Execution

```
policy_merged_resaped <- policy_merged[date >= as.Date("2020-02-23") &
                                         date <= as.Date("2020-06-30"),
                                         .(date,confirmed_per_day,type,detail,start_date)]

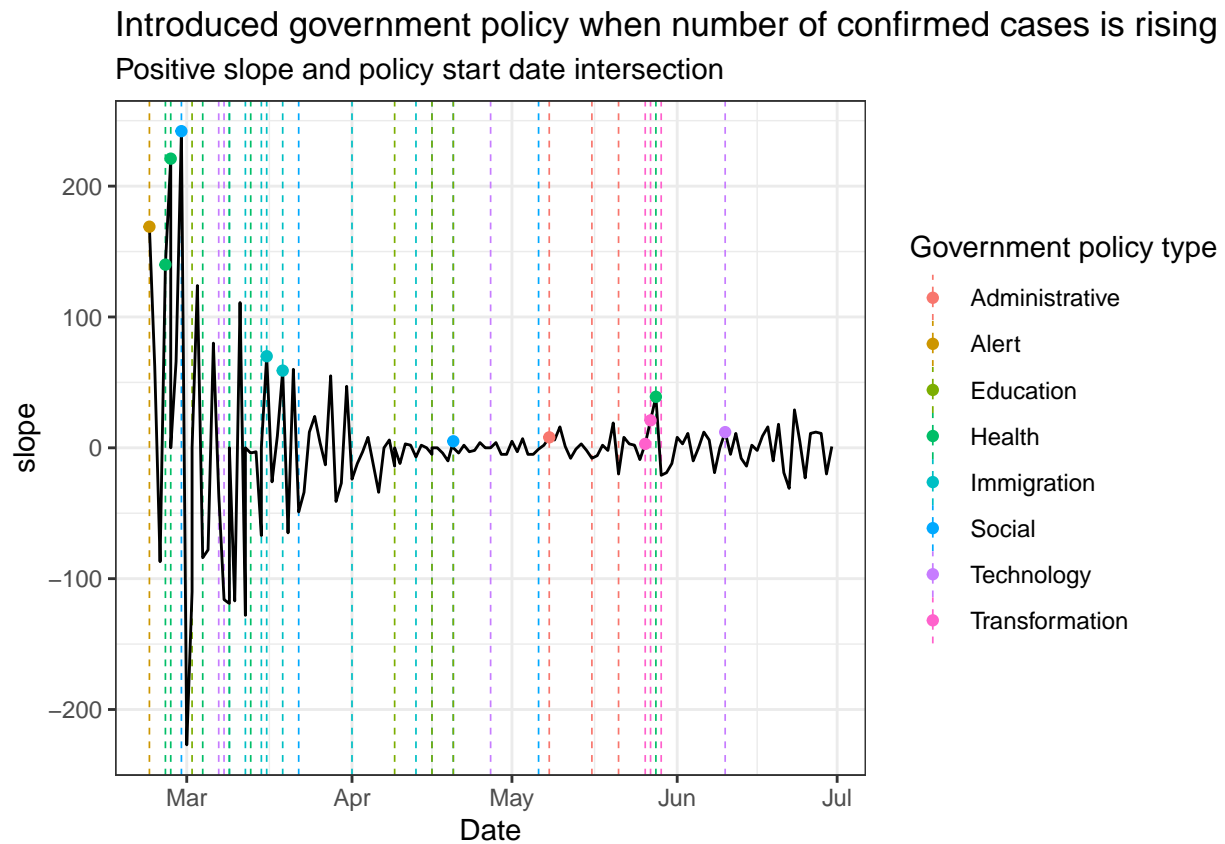
#for a better overview of the data

time_slope <- policy_merged_resaped %>%
  mutate(slope = confirmed_per_day - shift(confirmed_per_day,
                                           fill = 0, type = 'lag'))
```

Plot Nr.34: Slope of the case numbers and policy start date intersections

```
plot_5 <- ggplot(time_slope, aes(x=date, y=slope)) +
  geom_vline(data = subset(Policy, start_date == date), # filter data source
            aes(xintercept = date, color = type),
            size = 0.3, linetype = 'dashed') +
  geom_line(aes(x=date, y=slope)) +
  labs(color = 'Government policy type') +
  geom_point(data = subset(time_slope, start_date == date & slope>0),
            aes(x=date, y=slope, color=type)) +
  theme(axis.text.x = element_text(angle = 90, vjust = 0.5, hjust=1)) +
  labs(x = 'Date', y = 'slope',
       subtitle='Positive slope and policy start date intersection') +
  ggtitle('Introduced government policy when number of confirmed cases is rising') +
  theme_bw()
```

plot_5



Plot Nr.35: Comparison of Number of Confirmed Cases and Slope with policy dates

```
require(gridExtra)
grid.arrange(plot_3, plot_5, nrow=2)
```

```
## Warning: Removed 1 row(s) containing missing values (geom_path).
```

