

# Analytical Notebook

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## Install and load R-Packages

```
# install and load packages
loadpackage <- function(x){
  for( i in x ){
    # require returns TRUE invisibly if it was able to load package
    if( ! require( i , character.only = TRUE ) ){
      # If package was not able to be loaded then re-install
      install.packages( i , dependencies = TRUE )
    }
    # Load package (after installing)
    library( i , character.only = TRUE )
  }
}

# load packages
loadpackage( c("readr", "knitr", "dplyr", "tidyr", "sparklyr", "ggplot2"))
```

## Load Covid-Datasets

The datasets UID\_ISO\_FIPS\_LookUp\_Table.csv and time\_series\_covid19\_confirmed\_global.csv are loaded.

```
# use url to current dataset
url_data1 <- "https://raw.githubusercontent.com/CSSEGISandData/COVID-19/master/csse_covid_19_data/UID_ISO_FIPS_LookUp_Table.csv"
url_data2 <- "https://raw.githubusercontent.com/CSSEGISandData/COVID-19/master/csse_covid_19_data/time_series_covid19_confirmed_global.csv"

# load datasets
data1 <- read_csv(url_data1)
data2 <- read_csv(url_data2)

# harmonise ID-Variables in both datasets
names(data1)[names(data1) == "Long_"] <- "Long"
names(data2)[names(data2) == "Country/Region"] <- "Country_Region"
names(data2)[names(data2) == "Province/State"] <- "Province_State"

# Since we are only interested in countries and not in regions
# we keep only a subset of the dataset without regions.
data1 <- subset(data1, is.na(data1$Province_State))
data2 <- subset(data2, is.na(data2$Province_State))

data2 <-
  reshape(
    data = as.data.frame(data2),
    varying = list(names(data2)[5:length(data2)]),
    timevar = "day",
```

```

    v.names = "count",
    idvar = c("Country_Region"),
    direction = "long",
    times = names(data2)[5:length(data2)]
  )

data2$date <- as.Date(data2$day, format = "%m/%d/%y")
data2$datecount <- data2$date-min(data2$date)
days <- unique(data2$day)
weeks <- days[seq(1, length(days), 7)]

write_csv(data1, path = '../input/data1.csv')
write_csv(data2, path = '../input/data2.csv')

# setting up spark
sc <- spark_connect(master = "local",
                     version = "2.4.3")

data1 <- copy_to(sc, data1, overwrite = T)
data2 <- copy_to(sc, data2, overwrite = T)
src_tbls(sc)

```

## Data Cleaning

Here the data sets are prepared for merging and reshaping. The ID variables must be standardized. Since we are only interested in countries and not in regions # we keep only a subset of the dataset without regions. Then we drop countries without information and we keep all important variables. We reshape the dataset to long format

```

data_merge <- merge(data2, data1, by = "Country_Region")

data_merge <- data_merge %>%
  select(Country_Region, Country_Region, day, date, count, datecount, Population)

write_csv(data_merge, path = '../input/data_merge.csv')
data_merge <- copy_to(sc, data_merge, overwrite = T)

# Germany, China, Japan, United Kingdom, US, Brazil, Mexico
my_data <- data_merge %>%
  filter(Country_Region=="Germany" | Country_Region=="Mexico" |
         Country_Region=="United Kingdom" | Country_Region=="US" |
         Country_Region=="Brazil" | Country_Region=="China" |
         Country_Region=="Japan") %>%
  mutate(rate = (count/Population)*100) %>%
  mutate(count2 = round((count/1000))) %>%
  collect() %>%
  print()

```

```

## # A tibble: 1,032 x 8
##   Country_Region day   date      count datecount Population   rate
##   <chr>          <chr> <date>      <dbl>      <dbl>      <dbl> <dbl>
## 1 Brazil        6/21~ 2020-06-21 1.08e6      151 212559409 0.510
## 2 Brazil        2/4/~ 2020-02-04 0.          13 212559409 0

```

```
## 3 Brazil      2/8/~ 2020-02-08 0.          17 212559409 0
## 4 Brazil      1/22~ 2020-01-22 0.          0 212559409 0
## 5 Brazil      6/19~ 2020-06-19 1.03e6      149 212559409 0.486
## 6 Brazil      2/6/~ 2020-02-06 0.          15 212559409 0
## 7 Brazil      5/9/~ 2020-05-09 1.56e5      108 212559409 0.0734
## 8 Brazil      5/13~ 2020-05-13 1.90e5      112 212559409 0.0895
## 9 Brazil      5/4/~ 2020-05-04 1.09e5      103 212559409 0.0511
## 10 Brazil     5/29~ 2020-05-29 4.65e5      128 212559409 0.219
## # ... with 1,022 more rows, and 1 more variable: count2 <dbl>
```

## Plots

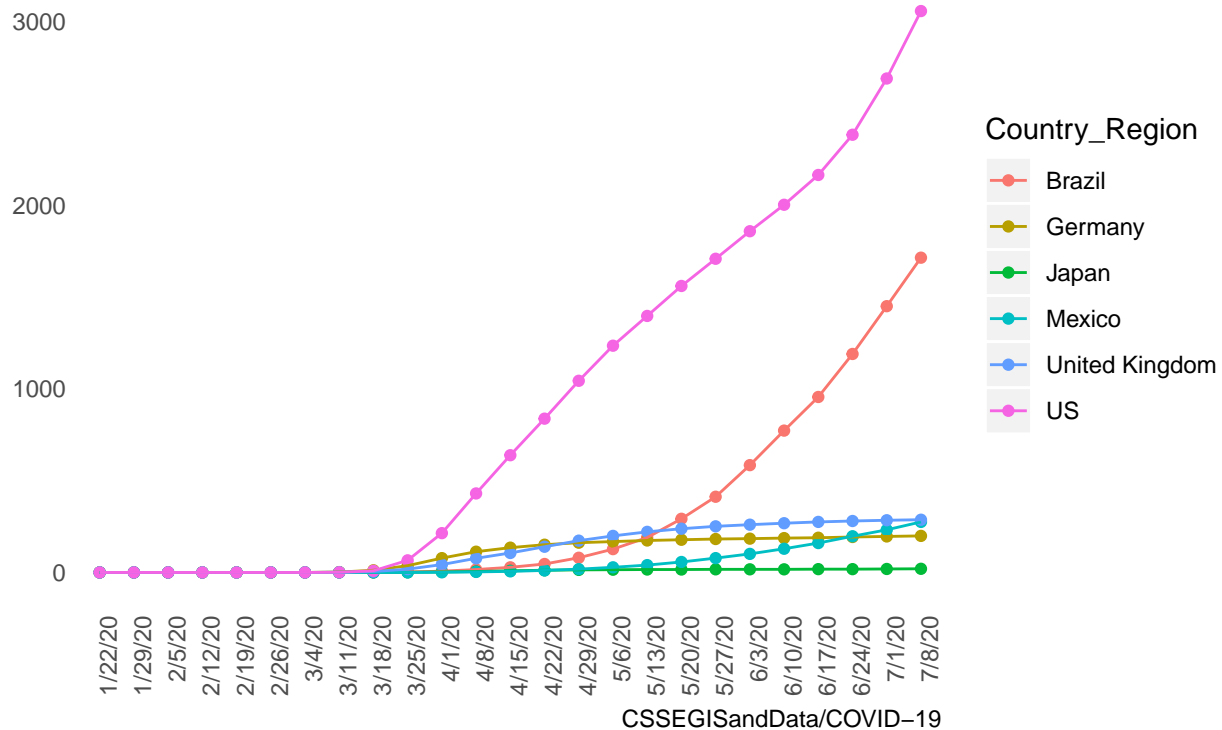
You can also embed plots, for example:

```
ggplot(data=my_data, aes(x=reorder(day, count2), y=count2, group=Country_Region, colour=Country_Region))
  geom_point()+
  geom_line()+
  scale_x_discrete(limit = weeks)+
  theme(axis.text.x = element_text(angle = 90, hjust = 1),
        axis.title=element_blank(),
        axis.ticks = element_blank(),
        strip.text = element_blank(),
        panel.grid.major = element_blank(),
        panel.grid.minor = element_blank(),
        panel.border = element_blank(),
        panel.background = element_blank())+
  labs(title = "Overall change of number of Corona Cases in TSD",
       caption = "CSSEGISandData/COVID-19")
```

```
## Warning: Removed 882 rows containing missing values (geom_point).
```

```
## Warning: Removed 882 rows containing missing values (geom_path).
```

## Overall change of number of Corona Cases in TSD



```
ggsave("../output/total_change.png")
```

```
## Saving 6.5 x 4.5 in image
```

```
## Warning: Removed 882 rows containing missing values (geom_point).
```

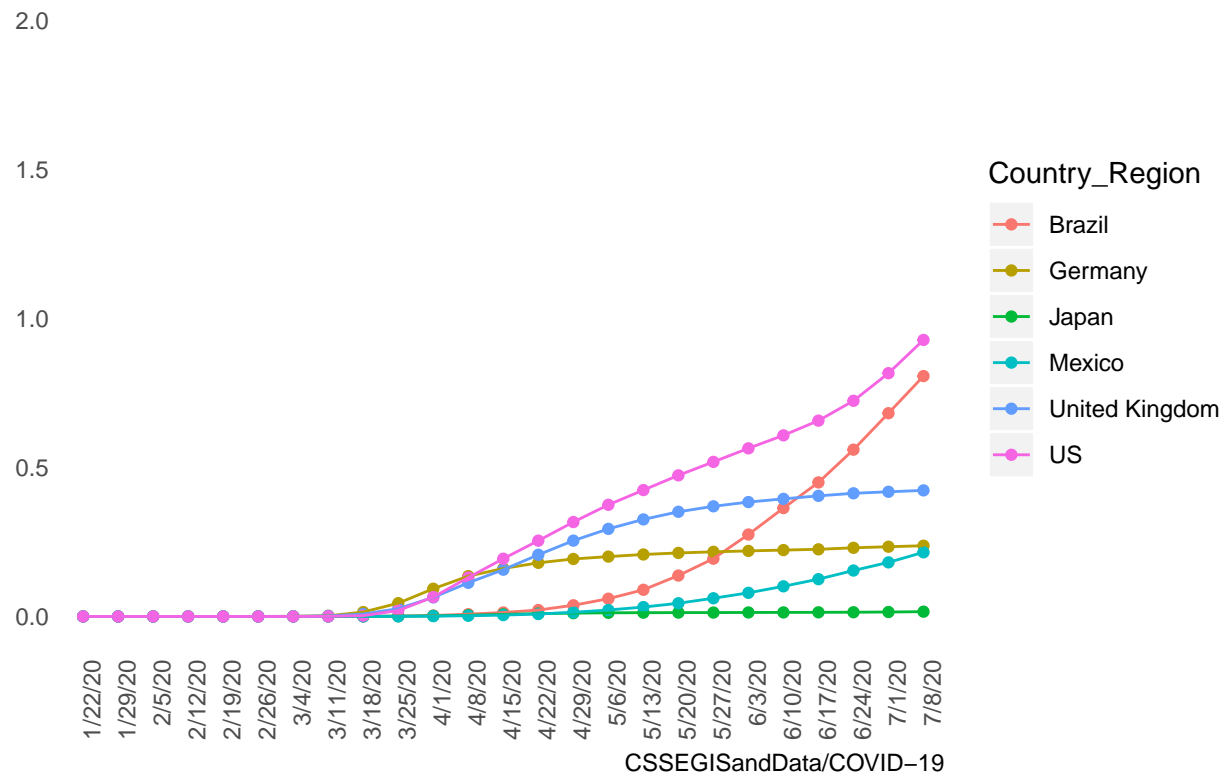
```
## Warning: Removed 882 rows containing missing values (geom_path).
```

```
ggplot(data=my_data, aes(x=reorder(day, rate), y=rate, group=Country_Region, colour=Country_Region)) +
  geom_point()+
  geom_line()+
  scale_x_discrete(limit = weeks)+
  ylim(0,2)+
  theme(axis.text.x = element_text(angle = 90, hjust = 1),
        axis.title=element_blank(),
        axis.ticks = element_blank(),
        strip.text = element_blank(),
        panel.grid.major = element_blank(),
        panel.grid.minor = element_blank(),
        panel.border = element_blank(),
        panel.background = element_blank())+
  labs(title = "Overall change of infection rate in percent",
       caption = "CSSEGISandData/COVID-19")
```

```
## Warning: Removed 882 rows containing missing values (geom_point).
```

```
## Warning: Removed 882 rows containing missing values (geom_path).
```

## Overall change of infection rate in percent



```
ggsave("../output/rate_change.png")
```

```
## Saving 6.5 x 4.5 in image
```

```
## Warning: Removed 882 rows containing missing values (geom_point).
```

```
## Warning: Removed 882 rows containing missing values (geom_path).
```

## Regression