

# Instructions

- Clear the environment
- Open a new R Script where you will do the exercise and later save in the project directory.
- Add the purpose of the file and the author
- Here are the main activities
  - 1) Load the `rio`, `lubridate` and `tidyverse` package
  - 2) Load the data using the `import`
  - 3) Select variables
  - 4) Rename variables with spaces
  - 5) Filter the values of interest
  - 6) Mutate variables
  - 7) Mutate with `if_else`

## Part 1 (*You :-)* will do this together)

### 1.1

- Load the `rio`, `lubridate` and `tidyverse` package
- Use the `hospital_df` and `location_df` object names; Load the `line_hospital_data.csv` and the `line_hospitals_locations.xlsx` data

```
# General format
hospital_df <- import("Data/line_hospital_data.csv")
```

- a) What is the structure of the data
- b) How many observations are in the data?
- c) What is the class of 'case\_id'

### 1.2 For this part we use the `hospital_df` data

- Create a subset called `sub1` of lab measurements, `case_id` and age data
- How many columns do you have now? add that as a comment in your code
- Hint: use `select` function

### 1.3 Rename variables with spaces

- Rename all the variables with a space and replace space with `_`
- Rename `date_of_outcome` to `date_outcome`

```
rename(.data, ...)
```

### 1.4 Filter data

- Create a data frame of all the participants who died: Hint use the `outcome` variable
- glimpse the data of the participant that died , how many rows and column, add that as a comment in your code
- create a data set of males over 25 years
- 

```
## example of female data set
female <- filter(hosp_data, gender=="f" )
```

- Use `filter` and `select` to show `case_id` , `hosp_date` and `date_onset` of patients who recovered.
- How many rows and columns, add that as a comment in your code

- Create a data of participants who had `cough` and `chills` , then select only `case_id`,`gender` and `age`.  
Hint: try and use the pipe operator
- Filter participants that had `cough` AND `chills` OR `aches` OR their `ct_blood` IS GREATER than 20

## 1.5 mutate

- Create a BMI variable where BMI= `weight_kg/height_meters * height_meters`  
– Hint: you have to create height in meters variables
- Create a variable `short` is YES if `ht_cm < 80` otherwise NO
- Hint: use IF\_ELSE
- Create age group of 10 year gaps, i.e 0-10,10-20,20-30,30-40,40-50,Over\_50
- HINT use CASE\_WHEN

```
## Hint of CASE WHEN
hosp_data <- hosp_data %>%
  mutate(paed =ifelse(age>14,"Adult","Paed"),
         age_group=case_when(
           age<14 ~"Paed",
           age >13 & age< 19 ~"Gen Z",
           age>18 & age<30 ~ "Non Gen Z",
           .default = "other"))
```

## 1.6 Merging

- Left join the `hospital_df` and `location_df` using `case_id` as the ID call the data `merge1`
- Inner join the `hospital_df` and `location_df` using `case_id` as the ID call the data `merge2`
- Full join the `hospital_df` and `location_df` using `case_id` as the ID call the data `merge3`
- Look at the glimpse of the 3 data sets above, what is the difference
- Export all the 3 files as `.xlsx` into the data folder]]

```
hosp_data <- import("Data/line_hospital.xlsx")
hospital_location <- import('data/line_hospital_sub.csv')
hosp_left_joined <- left_join(
  x = hosp_data,
  y = hospital_location,
  by = "case_id"
)
```

## 1.8 Create a pipe

Create a pipe chain which selects the `male` gender data

1. renames the variables with space
2. creates BMI
3. creates age group of 10
4. age of greater than 18

## 1.7 Extras

Run the following and create the year and month from the hospital date variable

```
hosp_data <- hosp_data %>%
  mutate(date_onset=as_date(date_onset))
```

Hint to create year and month

```
mutate(hosp_year = year(date_onset) ,  
       hosp_month = month(date_onset))
```

Use the `hosp_data$year` to check the length and report