MLW / KUHeS Statistics and R short course

Session 1 - Practical

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Go to the course website on GitHub:

https://github.com/mlw-stats/R_And_Statistics_Training_Autumn2023/Session1

From here, download the following files:

```
btTBreg.csv
btTBregHospitals.csv
btTBreg info.txt
```

- 1. Load the btTBreg.csv data table into R.
- 2. The variables cd41, cd42 and cd41.sk, cd42.sk measure the same variables (cd4 and cd4.sk respectively) in the same individuals at two different time point. This means the data are in wide format. Reformat to long format.
- 3. Save the reformatted data into a file called btTBregLong.tab in such a way that
 - i. Columns are tab-separated.
 - ii. Column names are saved.
 - iii. No row number is saved in the resulting file.
- 4. Copy the code below to generate some wide-format data. We will assume this dataset contains observations of 2 biomarkers, ferritin and rbp4 for 10 study participants at 2 different timepoints, day1 and day90.

```
set.seed(123)

df<-data.frame(
   id=paste(sep="","P",1:10),
   ferritin_day1=rexp(10,rate=1/195),
   rbp4_day1=rexp(10,rate=1/2.5)
) %>%
   mutate(
    ferritin_day90=rnorm(10,mean=ferritin_day1+5,sd=4),
      rbp4_day90=rbp4_day1+rexp(10,rate=1/0.25)
)
```

This is what this data table looks like:

```
id ferritin_day1 rbp4_day1 ferritin_day90 rbp4_day90
##
## 1
       P1
             164.474166 2.5120751
                                       169.251422
                                                    3.021712
## 2
       P2
             112.439003 1.2005368
                                       114.301474
                                                    1.552174
## 3
             259.165699 0.7025341
                                       261.231686
                                                    0.829188
```

```
## 4
       P4
               6.157585 0.9427946
                                         10.294123
                                                     1.007684
## 5
       P5
              10.961140 0.4707101
                                        14.621489
                                                     1.119933
## 6
       P6
              61.717737 2.1244653
                                        62.374941
                                                     2.431722
## 7
       P7
              61.274322 3.9080088
                                        65.932629
                                                     4.105679
## 8
       Р8
              28.327027
                        1.1969010
                                        37.609469
                                                     1.354221
## 9
       Р9
             531.616111 1.4773371
                                       536.034536
                                                     1.790997
## 10 P10
               5.684922 10.1025293
                                         6.022743
                                                   10.249700
```

Reformat this to long format, i.e. so that you have 4 columns: id, time, ferritin and rbp4.

- 5. Load the btTBregHospitals.csv data table. Join the data frames storing btTBreg.csv and btTBregHospitals.csv.
- 6. Compute the average patient age and the proportion of male patients for each hospital.
- 7. Write an R function that computes the following summary statistics, then, using your custom function, compute these for the bmi, cd41, cd42 columns:
 - i. mean
 - ii. median
 - iii. inter quartile range
 - iv. minimum
 - v. maximum
 - vi. number of missing values
- 8. Do the same now, but only for female patients. Repeat for only male patients.