Rmarkdown exercise

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2023-05-08

Available books links

Intro2R

Rmarkdown, the definitive guide

Rmarkdown exercise

```
install.packages("rmarkdown")
install.packages("tinytex")
tinytex::install_tinytex() # Tinytex is requred for PDF format
library(rmarkdown)
```

Packages for Rmarkdown

```
dir.create("output") # Create directory
dir.create("./output/figures")
unzip("all_data.zip") # Unzip ZIP file
list.files() # List files on current directory
list.dirs() # List directories on current directory
```

Create, list and extract directories and files

Import data and visualize data structure I am familiar with these sort of commands and I use them often on my current PhD project. For this reason I do not go into detail.

```
pathways <- read.table(file = "Data/dataset_overview_pathways.txt", sep = "\t", header = T, stringsAsFa</pre>
```

summary() returns some measures of central tendency of the variables of the data set, whereas str() only returns the column names and their data type.

str(pathways)

```
## 'data.frame': 15 obs. of 6 variables:

## $ X : Factor w/ 15 levels "common_other_pways",..: 11 15 10 3 2 12 1 7 4 5 ...

## $ OXTP: num 204 138 66 4 6 ...

## $ DOPP: num 167 75 71 2 1 ...

## $ SERP: num 143 71 71 2 16 ...

## $ STEP: num 31 31 19 0 0 ...

## $ VT.R: num 7 7 0 0 0 ...
```

summary(pathways)

```
X
                                    OXTP
                                                       DOPP
##
##
    common_other_pways
                                      : 0.0067
                                                          : 0.00605
                          :1
                               Min.
                                                  Min.
                                                  1st Qu.: 3.00000
  discarded (coverage) :1
##
                               1st Qu.: 6.5000
  discarded_(full STOP):1
                               Median: 109.0000
                                                  Median: 75.00000
  FEL_pos_genes
                                      :137.9420
                                                          : 87.07470
##
                          :1
                               Mean
                                                  Mean
##
   M7_pos_genes
                          :1
                               3rd Qu.:179.5000
                                                  3rd Qu.:128.00000
##
    mean omega_FEL
                          :1
                               Max.
                                      :579.0000
                                                          :293.00000
                                                  Max.
##
    (Other)
                          :9
##
         SERP
                              STEP
                                                  VT.R
##
           : 0.00731
                        Min.
                                : 0.00000
                                                    : 0.000
    Min.
                                             Min.
                        1st Qu.: 0.00543
   1st Qu.: 12.00000
                                             1st Qu.: 0.000
  Median : 68.00000
                        Median : 26.00000
                                             Median : 5.000
          : 78.07703
                               : 32.08875
##
  Mean
                        Mean
                                             Mean
                                                    : 5.083
##
    3rd Qu.:107.50000
                        3rd Qu.: 31.00000
                                             3rd Qu.: 7.000
                               :155.00000
##
  {\tt Max.}
           :256.00000
                        Max.
                                             Max.
                                                    :20.000
##
```

Rmarkdown language

- < Ctr + Alt + I > opens a chunk of code
- echo = FALSE can be added next to the {r} at the chunks of code that do not need to be shown.
- eval = FALSE can be added next to the {r} at the chunks of code that do not need to be run at knitting.
- Introducing images: ![image](path){width=???px, height=???px}

Create Version Control

Download Git needs to be installed in the computer. To check if it is installed, the following can be typed in the R terminal: git --version. Here is the link for downloading it.

Configure In the terminal of the computer, this needs to be typed: git config --global user.email 'you@youremail.com' git config --global user.name 'Your Name'