# R-markdown learning

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# 1 Preprocessing

## 1.1 Choosing the data

This task will use the penguins dataset. Let's working with it The source is taking from OpenML To catching anything, you need to

quote out like this

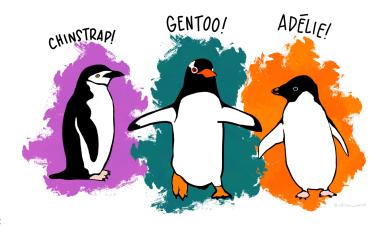
to make an empty line, simply add '' like this Penguins is cute!

# 1.2 Loading the dataset, reading data

First, I would like to:

- 1. Load packages
- 2. Read the data

#### 3. Remove missing values



To add an images, you use:

```
library(tidyverse)
```

```
## Warning: package 'tidyverse' was built under R version 4.3.2
## Warning: package 'ggplot2' was built under R version 4.3.2
## Warning: package 'forcats' was built under R version 4.3.2
## Warning: package 'lubridate' was built under R version 4.3.2
## -- Attaching core tidyverse packages ----- tidyverse 2.0.0 --
## v dplyr 1.1.3
                       v readr
                                    2.1.4
## v forcats 1.0.0 v stringr 1.5.0
## v ggplot2 3.4.4 v tibble 3.2.1
## v lubridate 1.9.3
                                   1.3.0
                        v tidyr
## v purrr
              1.0.2
## -- Conflicts ----- tidyverse_conflicts() --
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()
                 masks stats::lag()
## i Use the conflicted package (<a href="http://conflicted.r-lib.org/">http://conflicted.r-lib.org/</a>) to force all conflicts to become error
# load the dataset from OpenML Library
d <- OpenML::getOMLDataSet(data.id = 43347)</pre>
## Downloading from 'http://www.openml.org/api/v1/data/43347' to 'C:\Users\hiepa\AppData\Local\Temp\Rtm
## Downloading from 'https://api.openml.org/data/v1/download/22102172/Palmer-Penguins-Dataset-Alternati
# convert the OpenML object to a tibble (enhanced data.frame)
penguins <- d %>% dplyr::as_tibble()
#skimmed_penguins <- skimr::skim(penguins)</pre>
head(penguins)
## # A tibble: 6 x 7
## species island bill_length_mm bill_depth_mm flipper_length_mm body_mass_g
```

```
##
     <chr>
             <chr>>
                                 <dbl>
                                               <dbl>
                                                                  <dbl>
                                                                              <dbl>
## 1 Adelie Torgersen
                                  39.1
                                                18.7
                                                                               3750
                                                                    181
## 2 Adelie Torgersen
                                  39.5
                                                17.4
                                                                    186
                                                                               3800
## 3 Adelie Torgersen
                                                                               3250
                                  40.3
                                                18
                                                                    195
## 4 Adelie Torgersen
                                  NA
                                                NA
                                                                     NA
                                                                                 NA
## 5 Adelie Torgersen
                                  36.7
                                                19.3
                                                                    193
                                                                               3450
## 6 Adelie Torgersen
                                                20.6
                                                                               3650
                                  39.3
                                                                    190
## # i 1 more variable: sex <chr>
```

```
# Run again the code abow
library(tidyverse)
head(penguins)
```

```
## # A tibble: 6 x 7
                       bill_length_mm bill_depth_mm flipper_length_mm body_mass_g
##
     species island
     <chr>
            <chr>
                                               <dbl>
##
                                <dbl>
                                                                 <dbl>
                                                                             <dbl>
## 1 Adelie Torgersen
                                 39.1
                                               18.7
                                                                   181
                                                                              3750
                                 39.5
                                               17.4
                                                                   186
                                                                              3800
## 2 Adelie Torgersen
## 3 Adelie Torgersen
                                 40.3
                                               18
                                                                   195
                                                                              3250
## 4 Adelie Torgersen
                                 NA
                                               NA
                                                                    NA
                                                                                NA
## 5 Adelie Torgersen
                                 36.7
                                               19.3
                                                                   193
                                                                              3450
                                               20.6
## 6 Adelie Torgersen
                                 39.3
                                                                   190
                                                                              3650
## # i 1 more variable: sex <chr>
```

How the output and the code itself shows up in the document also be modified:

- 'echo = FALSE': only output is visible
- 'include = FALSE' hides both output and code
- 'warning = FALSE' to suppress warning message (also works with errors and message)
- 'eval = FALSE': code is not run

#### 1.3 Removing missing data

```
# Before removing na_values
nrow(penguins)
```

## [1] 344

```
# After removing
penguins <- penguins %>% drop_na()
nrow(penguins)
```

## [1] 333

Or we can write like this. I had removed missing values, so the data now only has 333 rows.

# 1.4 Descriptive statistics

The mean bill length is 43.9927928 mm. The bill depth is between 13.1 and 21.5

# 2 Graphs

Some ideas for the graphs:

- weight by flipper length
  - for the entire data
  - separately for each species
  - additionally by sex
- flipper and bill length

### 2.1