Profe, guardé el script en R pero no sé si era necesario también enviarle lo que aparecía en la consola y el gráfico que hicimos, así que los puse acá por si acaso.

Restarting R session...

> install.packages("palmerpenguins")

trying URL 'https://cran.rstudio.com/bin/macosx/contrib/4.1/palmerpenguins\_0.1.0.tgz'

Content type 'application/x-gzip' length 3001757 bytes (2.9 MB)

==================================================

downloaded 2.9 MB

The downloaded binary packages are in

/var/folders/xf/qn9wlb9171988vw34wk98trc0000gn/T//RtmpcFRuzk/downloaded\_packages

> install.packages("palmerpenguins")

trying URL 'https://cran.rstudio.com/bin/macosx/contrib/4.1/palmerpenguins\_0.1.0.tgz'

Content type 'application/x-gzip' length 3001757 bytes (2.9 MB)

==================================================

downloaded 2.9 MB

The downloaded binary packages are in

/var/folders/xf/qn9wlb9171988vw34wk98trc0000gn/T//RtmpcFRuzk/downloaded\_packages

> library(palmerpenguins)

> penguins

# A tibble: 344 × 8

species island bill\_length\_mm bill\_depth\_mm flipper\_length\_mm body\_mass\_g

<fct> <fct> <dbl> <dbl> <int> <int>

1 Adelie Torgersen 39.1 18.7 181 3750

2 Adelie Torgersen 39.5 17.4 186 3800

3 Adelie Torgersen 40.3 18 195 3250

4 Adelie Torgersen NA NA NA NA

5 Adelie Torgersen 36.7 19.3 193 3450

6 Adelie Torgersen 39.3 20.6 190 3650

7 Adelie Torgersen 38.9 17.8 181 3625

8 Adelie Torgersen 39.2 19.6 195 4675

9 Adelie Torgersen 34.1 18.1 193 3475

10 Adelie Torgersen 42 20.2 190 4250

# … with 334 more rows, and 2 more variables: sex <fct>, year <int>

> penguins <- palmerpenguins::penguins

> View(penguins)

> library(ggplot2)

> library(tidyverse)

── Attaching packages ──────────── tidyverse 1.3.1 ──

✓ tibble 3.1.5 ✓ dplyr 1.0.7

✓ tidyr 1.1.4 ✓ stringr 1.4.0

✓ readr 2.0.2 ✓ forcats 0.5.1

✓ purrr 0.3.4

── Conflicts ─────────────── tidyverse\_conflicts() ──

x dplyr::filter() masks stats::filter()

x dplyr::lag() masks stats::lag()

> ggplot(data=penguins)

> ggplot(data=penguins) +

+ geom\_point(mapping=aes(x=bill\_length\_mm, y=bill\_depth\_mm))

Warning message:

Removed 2 rows containing missing values

(geom\_point).

> ggplot(data=penguins)+

+ geom\_point(mapping = aes(x=bill\_length\_mm, y=bill\_depth\_mm))+

+ labs(title = "Correlación entre Bill depth y lenght en mm de pingüinos", x="Bill length en mm", y="bill depth en mm")

Warning message:

Removed 2 rows containing missing values

(geom\_point).

> ggplot(data = penguins)+

+ geom\_point(mapping = aes(x=bill\_length\_mm, y=bill\_depth\_mm, color=species))+

+ labs(title = "Correlación entre bill depth y length en mm de pingüinos por especie")

Warning message:

Removed 2 rows containing missing values (geom\_point).

