Lab2

Exercises

Irimie Fabio

Contents

exercise 1 - Create a new mean and sd function	1
A	1
	2
C	2
D	2
E	3
exercise 2 - Table of frequencies	4
A	4
B	5
C	6
D	6
exercise 3 - Histogram, Boxplot and quartiles	7
A	7
	8
C	9
D	
E	0
Exercise 4 - Multiple boxplots from scratch	1
A	1
B	1
C	3
Exercise 5 - Exploratory analysis of data	4
A	4
B 1	5
C	5
D	6
E 20	
F 2	

Exercise 1 - Create a new mean and sd function

\mathbf{A}

Create the Lab2 project. Use the same structure used for Lab1:

- scripts,
- plots,

• data

В

Install the palmerpenguins package, load the penguins dataset or, alternatively, download the .RData object from moodle and import it after placing it inside the data directory of the project (hint: use the load() function).

```
library(palmerpenguins)
data(penguins)
```

\mathbf{C}

Compute the mean, the standard deviation, and the median for the numeric variables of the dataset.

```
cat("Means: \n")
## Means:
colMeans(penguins[, c(3:6, 8)], na.rm = TRUE)
##
      bill_length_mm
                        bill_depth_mm flipper_length_mm
                                                                 body_mass_g
##
            43.92193
                              17.15117
                                                200.91520
                                                                 4201.75439
##
                year
##
          2008.02907
cat("\n")
# Medians
cat("Medians: \n")
## Medians:
sapply(penguins[, c(3:6, 8)], median, na.rm = TRUE)
      bill_length_mm
                         bill_depth_mm flipper_length_mm
                                                                 body_mass_g
                                 17.30
                                                   197.00
                                                                     4050.00
##
               44.45
##
                year
##
             2008.00
cat("\n")
# Standard deviations
cat("Standard deviations: \n")
## Standard deviations:
sapply(penguins[, c(3:6, 8)], sd, na.rm = TRUE)
                         bill depth mm flipper length mm
##
      bill length mm
                                                                 body mass q
##
           5.4595837
                              1.9747932
                                               14.0617137
                                                                 801.9545357
##
                year
##
           0.8183559
cat("\n")
```

\mathbf{D}

Create a function called stat_auto that simultaneously returns both the mean and the standard deviation of a given vector (hint: return an object of type list or simply a vector). Then try it on the same numeric variables in C. to check the results (hint: if you obtain NA maybe you forgot to remove NA terms in the vector).

```
stat_auto <- function(vec, na.rm = FALSE) {
  if (na.rm) {
    mean <- mean(vec, na.rm = TRUE)
}</pre>
```

```
sd <- sd(vec, na.rm = TRUE)</pre>
    return(list("mean" = mean, "sd" = sd))
  mean <- mean(vec)</pre>
  sd <- sd(vec)
 return(list("mean" = mean, "sd" = sd))
}
sapply(penguins[, c(3:6, 8)], stat_auto, na.rm = TRUE)
##
        bill_length_mm bill_depth_mm flipper_length_mm body_mass_g
                        17.15117
                                       200.9152
## mean 43.92193
                                                          4201.754
## sd
        5.459584
                        1.974793
                                       14.06171
                                                          801.9545
##
        year
## mean 2008.029
        0.8183559
## sd
```

\mathbf{E}

Create a function called stat_manual that simultaneously returns both the mean and the standard deviation of a given vector without using the mean() and the sd() functions (hint: you can use length(), sum(), and na.omit() functions). Then try it on the same numeric variables in C. to check the results.

```
stat_manual <- function(vec, na.rm = FALSE) {</pre>
  if (na.rm) {
    sum <- sum(vec, na.rm = TRUE)</pre>
    mean <- sum / na.omit(length(vec))</pre>
    sum <- sum((vec - mean)^2, na.rm = TRUE)</pre>
    denom <- na.omit(length(vec)) - 1</pre>
    varianza <- sum / denom
    sd <- sqrt(varianza)</pre>
    return(list("mean" = mean, "sd" = sd))
  }
  sum <- sum(vec)</pre>
  mean <- sum / length(vec)</pre>
  sum <- sum((vec - mean)^2)</pre>
  denom <- length(vec) - 1
  varianza <- sum / denom
  sd <- sqrt(varianza)</pre>
  return(list("mean" = mean, "sd" = sd))
sapply(penguins[, c(3:6, 8)], stat_manual, na.rm = TRUE)
```

bill_length_mm bill_depth_mm flipper_length_mm body_mass_g

```
## mean 43.66657 17.05145 199.7471 4177.326

## sd 5.449612 1.971543 14.06909 799.985

## year

## mean 2008.029

## sd 0.8183559
```

Exercise 2 - Table of frequencies

\mathbf{A}

In the penguins dataset, transform a numeric variable to a categorical one by aggregating values into classes. Consider the flipper length variable and create 10mm wide classes using the cut() function (hint: use the range() function to determine the min and max values of the variable, then define a sequence for the cuts).

```
range() function to determine the min and max values of the variable, then define a sequence for the cuts).
r <- range(penguins$flipper_length_mm, na.rm = TRUE)
splits \leftarrow seq(r[1], r[2], 10)
splits <- append(splits, r[2])</pre>
classes <- cut(penguins\flipper_length_mm, splits, ordered_result = TRUE)</pre>
cat("Splits: ", splits, "\n")
## Splits: 172 182 192 202 212 222 231
cat("Classes: \n")
## Classes:
classes
##
     [1] (172,182] (182,192] (192,202] <NA>
                                                    (192,202] (182,192]
     [7] (172,182] (192,202] (192,202] (182,192] (182,192] (172,182]
    [13] (172,182] (182,192] (192,202] (182,192] (192,202] (192,202]
    [19] (182,192] (192,202] (172,182] (172,182] (182,192] (182,192]
    [25] (172,182] (182,192] (182,192] (182,192] <NA>
                                                               (172, 182]
    [31] (172,182] (172,182] (182,192] (182,192] (192,202] (192,202]
```

[37] (182,192] (172,182] (172,182] (182,192] (172,182] (192,202] [43] (182,192] (192,202] (182,192] (182,192] (172,182] (172,182] ## [49] (182,192] (182,192] (182,192] (182,192] (182,192] (192,202] [55] (182,192] (182,192] (182,192] (192,202] (172,182] (192,202] ## [61] (182,192] (192,202] (182,192] (182,192] (182,192] (182,192] ## [67] (192,202] (182,192] (182,192] (192,202] (182,192] (182,192] ## ## [73] (192,202] (192,202] (182,192] (192,202] (182,192] (182,192] ## [79] (182,192] (192,202] (182,192] (192,202] (182,192] (192,202] [85] (182,192] (192,202] (182,192] (182,192] (182,192] (182,192] ## [91] (192,202] (202,212] (182,192] (182,192] (182,192] (202,212] ## [97] (182,192] (192,202] (172,182] (182,192] (182,192] (202,212] ## [103] (182,192] (182,192] (192,202] (182,192] (192,202] (182,192] ## [109] (172,182] (192,202] (192,202] (182,192] (192,202] (192,202] **##** [115] (182,192] (192,202] (182,192] (192,202] (182,192] (182,192] ## [121] (182,192] (192,202] (172,182] (192,202] (182,192] (192,202] ## [127] (182,192] (192,202] (182,192] (202,212] (182,192] (192,202] ## [133] (192,202] (192,202] (182,192] (182,192] (182,192] (192,202] ## [139] (182,192] (192,202] (192,202] (182,192] (182,192] (182,192] ## [145] (182,192] (182,192] (182,192] (182,192] (192,202] (192,202] ## [151] (182,192] (192,202] (202,212] (222,231] (202,212] (212,222]

```
## [157] (212,222] (202,212] (202,212] (212,222] (202,212] (212,222]
  [163] (212,222] (212,222] (212,222] (212,222] (202,212] (212,222]
## [169] (202,212] (212,222] (202,212] (212,222] (212,222] (212,222]
## [175] (212,222] (212,222] (212,222] (212,222] (212,222]
## [181] (202,212] (212,222] (212,222] (202,212] (202,212] (222,231]
## [187] (212,222] (212,222] (212,222] (212,222] (202,212] (202,212]
## [193] (202,212] (222,231] (202,212] (212,222] (212,222] (212,222]
## [199] (202,212] (222,231] (212,222] (212,222] (202,212] (212,222]
## [205] (202,212] (222,231] (212,222] (212,222] (202,212] (212,222]
  [211] (202,212] (222,231] (202,212] (212,222] (212,222] (222,231]
## [217] (212,222] (222,231] (212,222] (222,231] (212,222] (222,231]
## [223] (212,222] (212,222] (212,222] (212,222] (212,222] (222,231]
## [229] (202,212] (212,222] (212,222] (222,231] (202,212] (212,222]
## [235] (202,212] (222,231] (202,212] (222,231] (212,222] (212,222]
## [241] (202,212] (222,231] (212,222] (222,231] (202,212] (222,231]
## [247] (212,222] (222,231] (212,222] (212,222] (202,212] (222,231]
## [253] (212,222] (222,231] (212,222] (222,231] (212,222] (212,222]
## [259] (202,212] (212,222] (202,212] (202,212] (212,222] (222,231]
## [265] (212,222] (222,231] (212,222] (222,231] (212,222] (212,222]
## [271] (212,222] <NA>
                             (212,222] (212,222] (202,212] (212,222]
## [277] (182,192] (192,202] (192,202] (182,192] (192,202] (192,202]
## [283] (172,182] (192,202] (192,202] (192,202] (192,202] (192,202]
## [289] (182,192] (192,202] (182,192] (192,202] (192,202] (172,182]
## [295] (182,192] (192,202] (172,182] (182,192] (182,192] (192,202]
## [301] (192,202] (192,202] (192,202] (192,202] (182,192] (202,212]
## [307] (182,192] (192,202] (182,192] (202,212] (192,202] (192,202]
## [313] (192,202] (202,212] (182,192] (202,212] (202,212] (182,192]
## [319] (192,202] (192,202] (192,202] (192,202] (182,192] (202,212]
## [325] (182,192] (192,202] (192,202] (192,202] (192,202] (202,212]
## [331] (182,192] (192,202] (182,192] (202,212] (192,202] (192,202]
## [337] (202,212] (182,192] (192,202] (202,212] (192,202] (192,202]
## [343] (202,212] (192,202]
## 6 Levels: (172,182] < (182,192] < (192,202] < ... < (222,231]
```

\mathbf{B}

Use the table() function on the new variable generated by cut(). Then transform it into a data frame object. Rename the columns accordingly using the colnames() function (hint: the second column correspond to the absolute frequencies).

```
df <- data.frame(table(classes))
colnames(df) <- c("Classes", "AbsoluteFrequencies")
df</pre>
```

```
## Classes AbsoluteFrequencies
## 1 (172,182] 22
## 2 (182,192] 96
## 3 (192,202] 85
## 4 (202,212] 47
## 5 (212,222] 67
## 6 (222,231] 24
```

\mathbf{C}

Add the the columns for: relative frequencies, cumulative absolute frequencies, and cumulative relative frequencies.

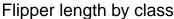
```
df$CumAbsFreq <- cumsum(df$AbsoluteFrequencies)
df$RelativeFrequencies <- df$AbsoluteFrequencies / sum(df$AbsoluteFrequencies)
df$RelAbsFreq <- cumsum(df$RelativeFrequencies)
df</pre>
```

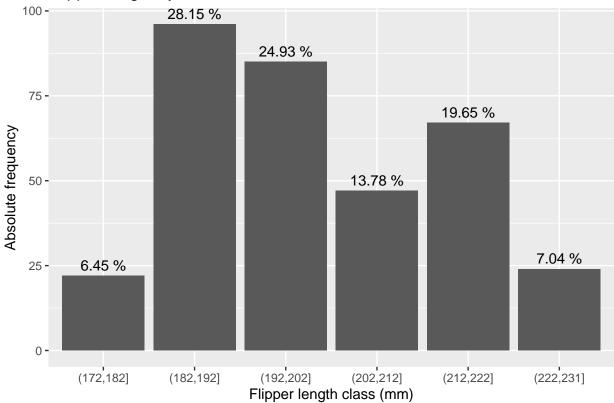
```
##
       Classes AbsoluteFrequencies CumAbsFreq RelativeFrequencies
## 1 (172,182]
                                 22
                                             22
                                                          0.06451613
## 2 (182,192]
                                 96
                                            118
                                                          0.28152493
                                 85
                                            203
## 3 (192,202]
                                                         0.24926686
## 4 (202,212]
                                 47
                                            250
                                                         0.13782991
## 5 (212,222]
                                 67
                                            317
                                                         0.19648094
## 6 (222,231]
                                            341
                                                         0.07038123
                                 24
##
    RelAbsFreq
## 1 0.06451613
## 2 0.34604106
## 3 0.59530792
## 4 0.73313783
## 5 0.92961877
## 6 1.00000000
```

D

Use the geom_col() function to plot the frequence of each class. Then, using the geom_text(aes(label = ...)) function, add the relative frequence as a percentage above each column (hint: substitute the ... with the relative frequency values. Use the round() function to choose the appropriate number of digits).

```
library(ggplot2)
ggplot(
  data = df,
  aes(
   x = dfClass,
   y = df$AbsoluteFrequency,
  )
) +
  geom_col(aes(y = df$AbsoluteFrequencies)) +
  geom_text(
   aes(
      label = paste(round(df$RelativeFrequencies * 100, 2), "%"),
      y = df$AbsoluteFrequencies,
   ),
   vjust = -0.5,
  ) +
  labs(
   title = "Flipper length by class",
   x = "Flipper length class (mm)",
   y = "Absolute frequency",
  ) +
  theme(legend.position = "bottom")
```



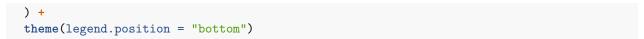


Exercise 3 - Histogram, Boxplot and quartiles

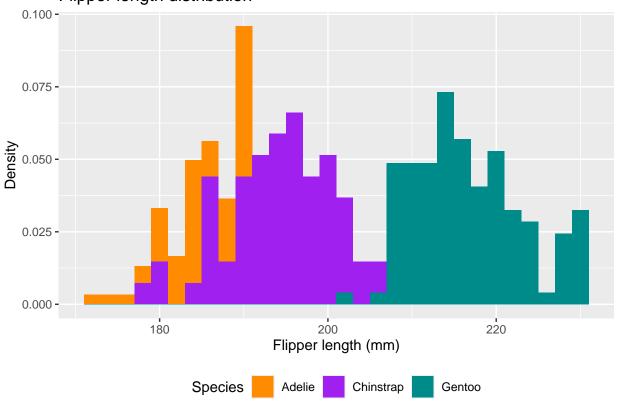
\mathbf{A}

Using the geom_histogram() function of the ggplot2 package plot the flipper length distribution coloring each species with a different color (hint: use the fill argument of the aes() function to fill the histogram area and the position = "identity" argument of the geom_histogram()). Play with the binwidth argument. Try to insert y = ..density. in aes(). Do you notice any change?

```
ggplot(
  data = penguins,
  aes(
    x = penguins$flipper_length_mm,
    fill = penguins$species,
  )
) +
  scale_fill_manual(values = c("darkorange", "purple", "cyan4")) +
  geom_histogram(
    position = "identity",
    binwidth = 2,
    aes(y = ..density..),
  ) +
  labs(
    title = "Flipper length distribution",
    x = "Flipper length (mm)",
    y = "Density",
    fill = "Species",
```







В

About the flipper length, for each species of penguins compute the: 1. Sample mean 2. Sample median 3. Sample standard deviation (use a division by n-1) 4. Sample variance

(hint: to choose only a specific species use penguins[penguins\$species == "Gentoo",])

```
species <- unique(penguins$species)
summary <- data.frame(
   Specie = species
)

for (specie in species) {
   summary$Mean[which(specie == species)] <-
        mean(penguins[penguins$species == specie, ]$flipper_length_mm,
        na.rm = TRUE
   )
   summary$Median[which(specie == species)] <-
        median(penguins[penguins$species == specie, ]$flipper_length_mm,
        na.rm = TRUE
   )
   summary$StandardDeviation[which(specie == species)] <-
        sd(penguins[penguins$species == specie, ]$flipper_length_mm,
        na.rm = TRUE
   )
}</pre>
```

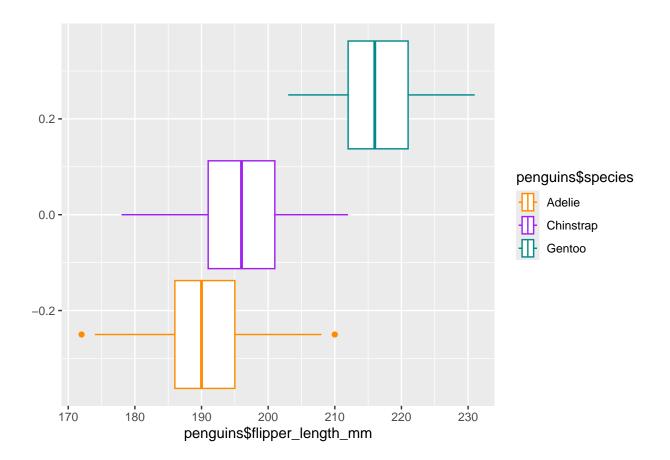
```
summary$Variance[which(specie == species)] <-
    var(penguins[penguins$species == specie, ]$flipper_length_mm,
        na.rm = TRUE
    )
}
summary</pre>
```

```
## Specie Mean Median StandardDeviation Variance
## 1 Adelie 189.9536 190 6.539457 42.76450
## 2 Gentoo 217.1870 216 6.484976 42.05491
## 3 Chinstrap 195.8235 196 7.131894 50.86392
```

\mathbf{C}

Using the geom_boxplot() function of the ggplot2 package plot the boxplot for the flipper length variable coloring each species with a different color (hint: use the color argument of the aes() function).

```
ggplot(
  penguins,
  aes(
    x = penguins$flipper_length_mm,
    color = penguins$species
  )
) +
  scale_color_manual(values = c("darkorange", "purple", "cyan4")) +
  geom_boxplot(
    aes(x = penguins$flipper_length_mm)
## Warning: Use of `penguins$flipper_length_mm` is discouraged.
## i Use `flipper_length_mm` instead.
## Warning: Use of `penguins$species` is discouraged.
## i Use `species` instead.
## Warning: Removed 2 rows containing non-finite outside the scale range
## (`stat_boxplot()`).
```



\mathbf{D}

Compute the flipper length quartiles for the "Gentoo" penguins (Q1, Q2, Q3).

```
gentoo <- penguins[penguins$species == "Gentoo", ]
quartiles <- quantile(
  gentoo$flipper_length_mm,
  c(0.25, 0.5, 0.75),
  na.rm = TRUE
)
quartiles
## 25% 50% 75%</pre>
```

\mathbf{E}

212 216 221

Calculate the flipper length 40th percentile for the "Adelie" penguins.

```
adelie <- penguins[penguins$species == "Adelie", ]
p40 <- quantile(
  adelie$flipper_length_mm,
  0.4,
  na.rm = TRUE
)
p40</pre>
```

40%

Exercise 4 - Multiple boxplots from scratch

\mathbf{A}

Generate random data with some structure, and create one data set for each day of the week (hint: use the for() cycle, data should have 7 columns). At the end you should obtain a matrix with N rows (N = N) random number to generate each time) and 7 columns (one for each day of the week).

```
set.seed(123)
n <- 10
days <- c(
    "Monday",
    "Tuesday",
    "Wednesday",
    "Friday",
    "Saturday",
    "Sunday"
)
data <- matrix(nrow = n, ncol = 7)

for (i in 1:7) {
    data[, i] <- round(runif(n, min = 0, max = 100), 0)
}

df <- data.frame(data)
colnames(df) <- days
df</pre>
```

##		Monday	Tuesday	Wednesday	Thursday	Friday	Saturday	Sunday
##	1	29	96	89	96	14	5	67
##	2	79	45	69	90	41	44	9
##	3	41	68	64	69	41	80	38
##	4	88	57	99	80	37	12	27
##	5	94	10	66	2	15	56	81
##	6	5	90	71	48	14	21	45
##	7	53	25	54	76	23	13	81
##	8	89	4	59	22	47	75	81
##	9	55	33	29	32	27	90	79
##	10	46	95	15	23	86	37	44

В

Go from a wide to a long data format. You should create a data frame object with exactly two columns. One contains the values created in A., the other contains the corresponding day of the week.

```
df_long <- data.frame(
  "Day" = rep(days, each = n),
  "Value" = c(
    df$Monday,
    df$Tuesday,
    df$Wednesday,
    df$Thursday,</pre>
```

```
df$Friday,
  df$Saturday,
  df$Sunday
)
)
df_long

## Day Value
## 1 Monday 29
```

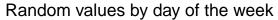
2 Monday 79 ## 3 Monday 41 ## 4 Monday 88 ## 5 Monday 94 ## 6 Monday 5 ## 7 Monday 53 ## 8 Monday 89 ## 9 Monday 55 ## 10 Monday 46 ## 11 Tuesday 96 ## 12 Tuesday 45 ## 13 Tuesday 68 ## 14 Tuesday 57 ## 15 Tuesday 10 ## 16 Tuesday 90 ## 17 Tuesday 25 ## 18 Tuesday 4 ## 19 Tuesday 33 ## 20 Tuesday 95 ## 21 Wednesday 89 ## 22 Wednesday 69 ## 23 Wednesday 64 ## 24 Wednesday 99 ## 25 Wednesday 66 ## 26 Wednesday 71 ## 27 Wednesday 54 ## 28 Wednesday 59 ## 29 Wednesday 29 ## 30 Wednesday 15 ## 31 Thursday 96 ## 32 Thursday 90 ## 33 Thursday 69 ## 34 Thursday 80 ## 35 Thursday 2 ## 36 Thursday 48 ## 37 Thursday 76 ## 38 Thursday 22 ## 39 Thursday 32 ## 40 Thursday 23 ## 41 Friday 14 ## 42 Friday 41 ## 43 Friday 41 ## 44 Friday 37 ## 45 Friday 15 ## 46 Friday 14

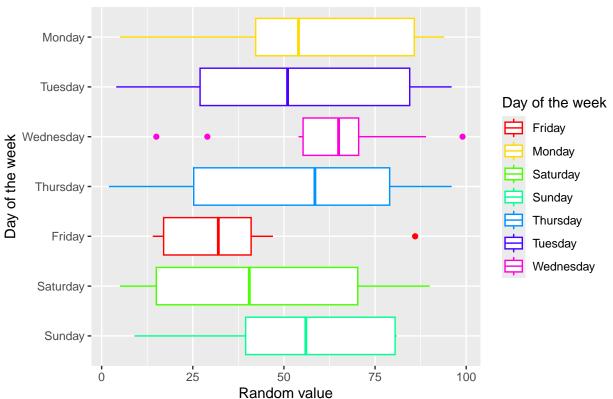
```
## 47
         Friday
                     23
## 48
         Friday
                     47
         Friday
## 49
                    27
## 50
         Friday
                    86
## 51
       Saturday
                     5
## 52
       Saturday
                     44
## 53
       Saturday
                    80
       Saturday
## 54
                     12
## 55
       Saturday
                    56
## 56
       Saturday
                    21
## 57
       Saturday
                    13
                    75
## 58
       Saturday
## 59
       Saturday
                    90
## 60
       Saturday
                    37
## 61
         Sunday
                     67
## 62
         Sunday
                     9
## 63
         Sunday
                     38
         Sunday
## 64
                     27
## 65
         Sunday
                    81
## 66
         Sunday
                    45
## 67
         Sunday
                    81
## 68
         Sunday
                    81
         Sunday
## 69
                    79
## 70
         Sunday
                     44
```

\mathbf{C}

Plot the seven boxplots (one for each day of the week) in one graph, horizontally oriented (hint: coord_flip() function translates the axes, the "limits" argument of scale_x_discrete() allows you to reorder the axis labels).

```
ggplot(
  df_long,
  aes(
    x = Day
    y = Value,
    color = Day
  )
) +
  scale color manual(values = rainbow(7)) +
  scale_x_discrete(limits = rev(days)) +
  geom_boxplot() +
  coord_flip() +
    title = "Random values by day of the week",
    x = "Day of the week",
    y = "Random value",
    color = "Day of the week"
```





Exercise 5 - Exploratory analysis of data

Penguins dataset does not contain their weights and flipper lengths only. Many other variables are available. Let's explore it a little more:

\mathbf{A}

How many islands are there? And how many penguins are present in each isle? Are the 3 species of penguins living together? (hint: use the table() function).

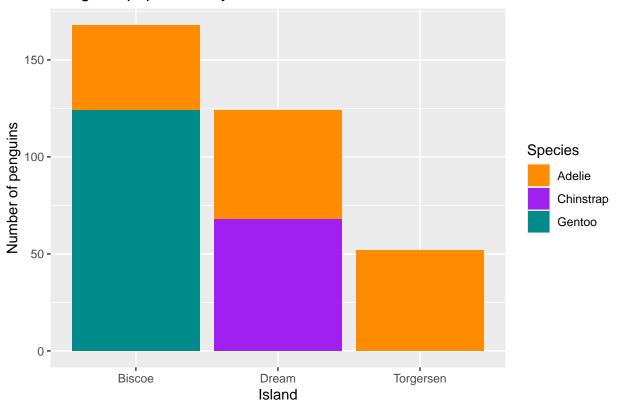
```
cat("The number of islands is: ", length(unique(penguins$island)), "\n")
## The number of islands is: 3
table(penguins$island)
##
##
      Biscoe
                  Dream Torgersen
         168
                    124
table(penguins$island, penguins$species)
##
##
               Adelie Chinstrap Gentoo
##
     Biscoe
                    44
                               0
                                     124
##
     Dream
                    56
                              68
                                       0
##
     Torgersen
                    52
                               0
                                       0
```

\mathbf{B}

Try to use the geom_bar() or geom_col() functions to graphically represent the population of each island, colored by species (hint: islands in the x-axis, number of penguins in the y-axis).

```
ggplot(
  penguins,
  aes(
    x = island,
    fill = species
)
) +
  geom_bar() +
  scale_fill_manual(values = c("darkorange", "purple", "cyan4")) +
  labs(
    title = "Penguins population by island",
    x = "Island",
    y = "Number of penguins",
    fill = "Species"
)
```

Penguins population by island

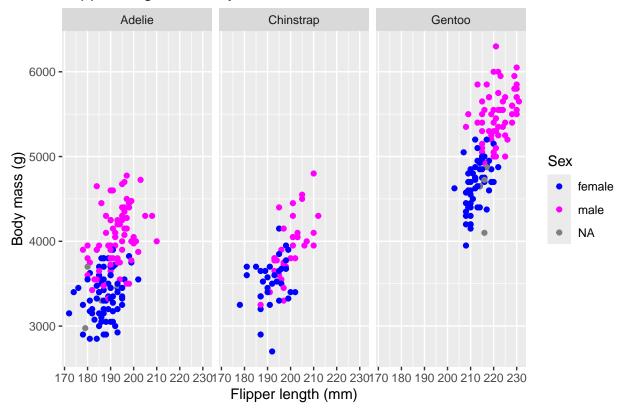


\mathbf{C}

Use a scatter plot to represent flipper length vs. body mass. Color the point according to the "sex" variable. Try to use facets to see if there are differences across species (hint: use facet_grid(\sim species) function to add facets for species).

```
ggplot(
  penguins,
  aes(
    x = penguins$flipper_length_mm,
    y = penguins$body_mass_g,
    color = penguins$sex
)
) +
  geom_point() +
  facet_grid(~species) +
  scale_color_manual(values = c("blue", "magenta")) +
  labs(
    title = "Flipper length vs. body mass",
    x = "Flipper length (mm)",
    y = "Body mass (g)",
    color = "Sex"
)
```

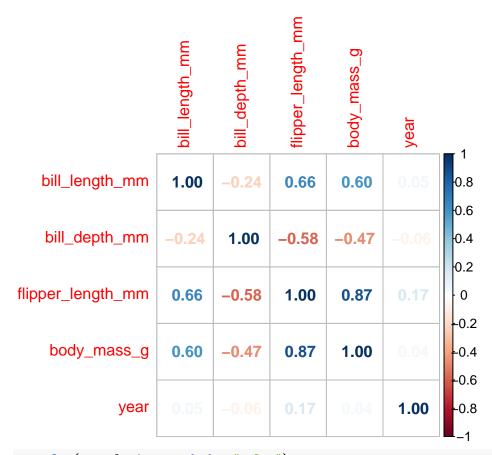
Flipper length vs. body mass



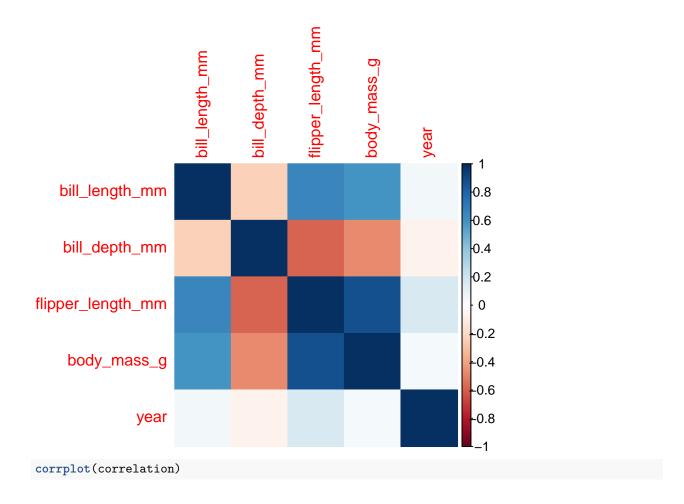
D

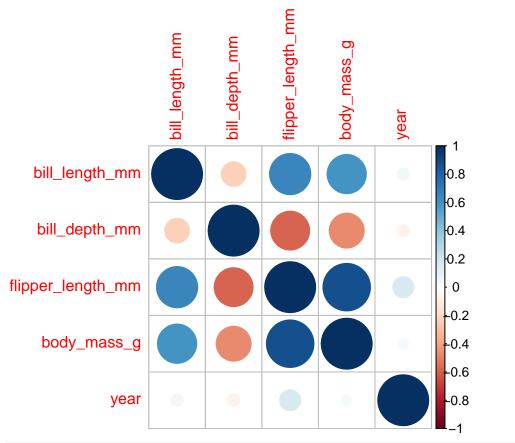
The numeric variables shows some interesting relationships. Are they correlated? Use the cor() and the corrplot() functions to study correlations between numeric variables (hint: try to google corrplot() to see which package you have to install to use it).

```
library(corrplot)
correlation <- cor(penguins[, c(3:6, 8)], use = "complete.obs")
corrplot(correlation, method = "number")</pre>
```

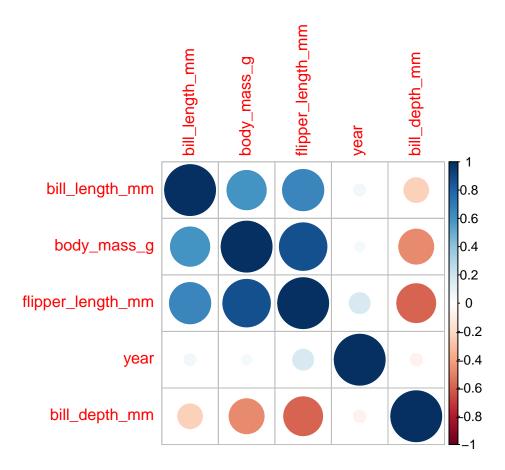


corrplot(correlation, method = "color")





corrplot(correlation, order = "AOE")



\mathbf{E}

Choose a pair of numeric variables, compute the correlation between them without using the cor() function (hint: remember the formula).

```
x <- penguins$flipper_length_mm
y <- penguins$body_mass_g

n <- length(x)
mean_x <- mean(x, na.rm = TRUE)
mean_y <- mean(y, na.rm = TRUE)

covariance <- sum((x - mean_x) * (y - mean_y), na.rm = TRUE) / (n - 1)
sd_x <- sqrt(sum((x - mean_x)^2, na.rm = TRUE) / (n - 1))
sd_y <- sqrt(sum((y - mean_y)^2, na.rm = TRUE) / (n - 1))

correlation <- covariance / (sd_x * sd_y)
cat(
   "The correlation between flipper length and body mass is: ",
   correlation, "\n"
)</pre>
```

The correlation between flipper length and body mass is: 0.8712018

\mathbf{F}

Plot the scatter plot for bill length vs. bill depth. Color the points by species. Use the function $geom_smooth(formula = "y \sim x")$ to add a line to represent the linear relationship between the two variables. Then, again, use $geom_smooth(formula = "y \sim x")$ colored by species. What are you noticing?

```
ggplot(
  penguins,
  aes(
    x = penguins$bill_length_mm,
    y = penguins\bill_depth_mm,
    color = penguins$species
) +
  geom_point() +
  geom_smooth(
    formula = "y \sim x",
    method = "lm",
    aes(color = penguins$species),
    se = FALSE
  ) +
  scale_color_manual(values = c("darkorange", "purple", "cyan4")) +
    title = "Bill length vs. bill depth",
    x = "Bill length (mm)",
    y = "Bill depth (mm)",
    color = "Species"
  )
```

Bill length vs. bill depth

