Lab2

Exercises

Irimie Fabio

Contents

A	1
C	1
D E Exercise 2 - Table of frequencies A B C	2
Exercise 2 - Table of frequencies A	2
Exercise 2 - Table of frequencies A	2
A	3
B	4
C	4
	5
	6
D	6
Exercise 3 - Histogram, Boxplot and quartiles	7
A	7
В	8
C	8
D	8
E	8
Exercise 4 - Multiple boxplots from scratch	9
A	9
B	9
C	9
Exercise 5 - Exploratory analysis of data	9
A	9
В	9
C	9
D	9
E	9
F	10

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.

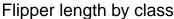
```
df$CumAbsFreq <- cumsum(df$AbsoluteFrequencies)
df$RelativeFrequencies <- df$AbsoluteFrequencies / sum(df$AbsoluteFrequencies)
df$RelAbsFreq <- cumsum(df$RelativeFrequencies)
df</pre>
```

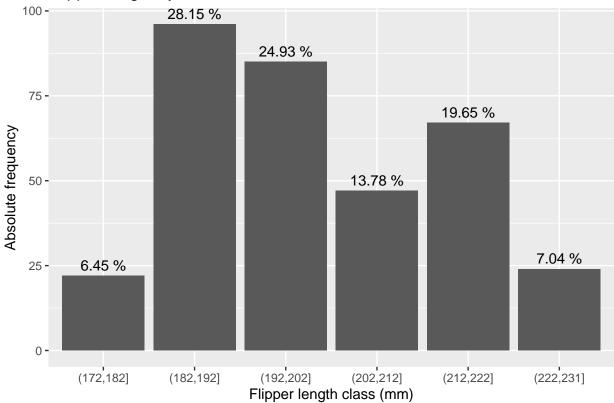
```
##
       Classes AbsoluteFrequencies CumAbsFreq RelativeFrequencies
## 1 (172,182]
                                 22
                                             22
                                                          0.06451613
## 2 (182,192]
                                 96
                                            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")
```



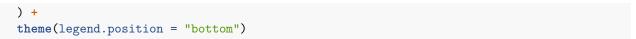


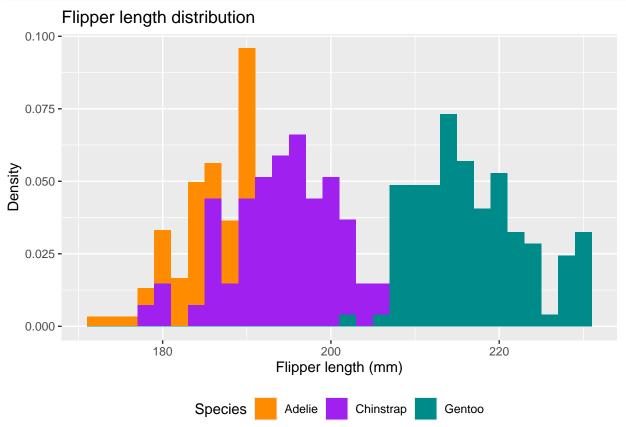
Exercise 3 - Histogram, Boxplot and quartiles

\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",
```





\mathbf{B}

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",])

\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).

\mathbf{D}

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

\mathbf{E}

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

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) number of random number to generate each time) and 7 columns (one for each day of the week).

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.

\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).

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).

В

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).

\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).

\mathbf{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).

\mathbf{E}

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

\mathbf{F}

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