Exercise day 2

Introduction to R for Basic Statistics

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Exercise B: Reshaping data (Part II)

For this exercise we keep working with the data of Exercise of Day 1.

It is a subset of "follicle" data, collected from patients with cancer that had OTC (ovarian tissue cryopreservation). Follicles were cultured for 8 days and the diameter was collected every 2 days. The aim of the study was to compare the follicles growth among different treatment groups.

Question 0 Load data into R (use the read.csv function). Remark: Remember to set your working directory with setwd(), or to define the correct path for the data

- 1. Consider Data set follicle.
- 2. Visualize the first lines of the data.
- 3. Print the summary of the data. Is there any missing values? If yes, how many at Day0 and how many at Day8?
- 4. How many follicles have been collected by patient? Hint: use the function table() to check how many observations by patient

Question 1

- 1. Calculate mean and standard deviation of the diameter at Day0 (Be careful, there are some missing!)
- 2. When we encounter into missing, we are often interested in the *complete case analysis* where we exclude patients with missing observations:
 - 2a. Use the na.omit function (excludes all rows that have at least one missing values)

db.CC<-na.omit(NameofDataFrame)</pre>

- 2b. check the dimension of the new data.frame
- 2c. Calculate mean and standard deviation of the diameter at DayO from db.CC
- 2d. Compare results with the ones in point 1.

Question 2 For each follicle the diameter was measured at day 0,2,4,6,8.

- 1. Are the data in a wide or long format?
- 2. Convert data from wide to long (or viceversa). Hint: You can use the function reshape
- 3. How many rows would we expect for each patient? Is it correct? (You can use the command table(db\$Patient))

Question 3 We are interested in the follicle growth over time. We can calculate the diameter difference from time 0 at each time point:

- 1. Subset observations at Day 0. Create a data frame from the subset with only columns *Number* and *diameter*.
- 2. Rename the variable of diameter into diameter 0.
- 3. Merge this data frame and the long format of your data set by Number (follicle ID).
- 4. Add to data frame a new variable for the difference of diameter at each time point.

Question 4 Descriptive at baseline (Day0). We want to create one data set with all characteristics of patients at baseline.

- 1. Merge the two data sets: long version from Question 3 and patient to have baseline characteristics in one data.frame
- 2. Check if the number of observation for each Patient is correct (use table())
- 3. Print mean and standard deviation of diameter at Day0 by Disease (use tapply() or aggregate())
- 4. Plot the histogram for the density of diameter at Day0 (use hist(...,prob=TRUE))
- 5. Add to the data frame a categorical variable for age considering the intervals: (19,30], (30,35], (35,40] (use the function cut())
- 6. Create a Boxplot of diameter at Day 0 by Age category.

Question 5 Descriptive at Day 8

- 1. Calculate the average difference in diameter after 8 days.
- 2. Add to the data frame the log-transformed variable for the difference in diameter at day 8. (You can use the function log())
- 3. Calculate mean and standard deviation for the log-transformed difference in diameter after 8 days in each treatment group.
- 4. Create a scatter plot of the difference in diameter after 8 days :
 - defining the color by treatment group
 - specify one type of point with pch (you can choose)
 - precise as name of axis: x= " ", y=" log-difference diameter"
 - define the main title for the plot: "Day 8"
- 5. Create a boxplot for the difference in Diameter after 8 days by treatment group
 - specify three colors (one for each treatment group)