Exercise day 2

Introduction to R for Basic Statistics

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Exercise B: Reshaping data (Part II)

Consider the data we used for 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:

Remark: Remember to set your working directory with setwd() to name the script and save it in the correct folder

- 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. Use the command table(dbf\$patient), interpret the numbers.

Question 1: missing observations

- 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(follicle)</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. Did something change? If yes, Why?

Question 2: wide/long format Consider the data set without missing (db.CC) that you obtained in question 2a.

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

1. Respect to the follicles, are data in a wide or long format?

- 2. Convert data from wide to long (or viceversa) using the reshape function.
- 3. How many rows would we expect for each Day? Is it correct? (You can use the command table(db\$Day))

Question 3: descriptive statistics at baseline Descriptive at baseline (Day0). We would like to create one data set with all characteristics of patients at baseline.

- 1. Merge the data set obtained in Question 2 and patient to add baseline characteristics in the data frame
- 2. Create a categorical variable for age considering the intervals: (19,30], (30,35], (35,40]. (Use the function cut())
- 3. Subset from the merged data only observation at baseline (Day 0)
- 4. Plot the histogram for the density of diameter at Day0
- 5. Create a Boxplot of diameter at Day 0 by Age category. Would you say that the follicle diameter is dependent on age?
- 6. Create a table with the counts of follicles per Disease and Treatment. Which is the most common disease?
- 7. Print the mean of diameter at Day0 by Disease (use tapply() or aggregate()).
 - 7a. Would you say that the follicle diameter is dependent on the disease? 7b. Show the boxplot of diameter at time 0 by Disease.

Question 4: change in diameter We are interested in the follicle growth over time. We can calculate the diameter difference from time 0 at each time point:

- 1. Take the subset observations at Day 0. Create a data frame 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 (created in Question 3.1) by Number.
- 4. Create a new variable "diam.change" for the difference of diameter at each time point.

Question 5: Descriptive of diameter change over time

- 1. Show the Boxplot of diameter by Day. Would you say that the diameter is growing over time?
- 2. Calculate the median diameter change by Day and Treatment (save the results, you need them for the next step). use aggregate with formula: diam.change~Day + Treatment
- 3. Plot the diameter growth over time by treatment group:
 - 2a. rename the column "diam.change" from the data.frame obtained in the previous point with "median.change"
 - 2b. merge the data set with the one obtained in Question 4 (after creating diam.change)
 - 2c. create a plot of the difference from time 0 at varying of days:
 - define the color by Treatment group
 - precise as name of axis: x= "Days", y="median diameter's diff"
 - add the legend
- 4. Calculate the relative change for the diameter ((diameter - diameter 0)/diameter 0) and re-create the same plot.