data management, data warehousing, statistics, information technology and scientific writing

# Data analysis with R

Lecture 3
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### Example of sapply

• With *sapply* we can calculate eg. mean for different elements of a list with single statement.



### Example of tapply

- The *tapply* function is simple to use. First, we will generate some data.
- clinical <- data.frame(patient = 1:100,
  age = rnorm(100, mean = 60, sd = 12),
  treatment = gl(2, 50, labels =
  c("Treatment", "Placebo")))</pre>
- tapply(clinical\$age, clinical\$treatment, mean)

Treatment Control 56.53683 59.77712



## by()

• For calculation in subgroups, we can also use by() – function. This well end up in the same result as tapply.

```
by(iris[, 1:4], Species, mean)
Species: setosa
Sepal.Length Sepal.Width Petal.Length Petal.Width
                 3.428
    5.006
                              1.462
                                           0.246
Species: versicolor
Sepal.Length Sepal.Width Petal.Length Petal.Width
    5.936
                 2.770
                              4.260
                                           1.326
  Species: virginica
Sepal.Length Sepal.Width Petal.Length Petal.Width
    6.588
                              5.552
                                           2.026
                2.974
```

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#### R-dates

- Calendar dates can be represented in R using objects' of class 'Date'.
- You can convert character representation of dates to objects of class 'Date' by using the function *as.Date()*.
- Default format for the character representation is "2010-11-08".
- With *date()* we can get the current date and time.



### R-dates (2)

- Dates can be inserted also in different formats than '2010-11-07'.
  - We need to use format-option in the as.Date-function for this to happen.
    - as.Date("07/11/2010",format="%d/%m/%Y")
- Dates are used in data analysis quite often, eg. in
  - Time to event calculations (eg. survival analysis)
  - Age, duration of exposure, time on study, etc...
  - Time series



## Sampling

- With R we can do simple random sampling easily.
- Lets consider a deck of cards. We can select 5 of them randomly by typing:

```
>sample(52,5,replace=F)
[1] 32 34 31 14 29
```

• With the replace-option we can choose, do we return the card to the deck or not.



### cbind & rbind

- *cbind()* takes a sequence of vector, matrix or dataframes arguments and combines them by columns.
  - cbind is useful eg. when we have our data stored in vectors and we want to construct a data-frame of them.
- rbind() takes a sequence of vector, matrix or data
   frames arguments and combines them by rows.
  - An example of the use *rbind()*: we have several different data entries from eg. different people and we want to combine all of them in a single data-frame



### cbind & rbind (2)

- With *cbind()* R requires an even amount of \_\_\_\_ observations (rows) in the vectors we are combining
- Likewise with *rbind()* we need to have an even amount or columns, for *rbind()* to work.
  - This can be problematic in some cases as we might have a subset of data with less variables than the other.
    - By creating eg. dummy (empty) variables we can surpass this problem.
- These functions are simple and effective to use and thus they are heavily used along R-programmers.



### Merging datasets

- With previous function cbind() we cannot have an id-variable when we combine the vectors.
  - Usually though, when we are merging datasets, we have to consider an ID-variable (eg. patient number), so that we can be sure that the data goes to the right invidual.
- In cases like this, we can use a function called merge().



# Merging datasets (2)

- With *merge*-function we have to first naturally tell R, what are the datasets we want to merge and then the Id-variable, which does not have to be called the same in the two datasets.
  - Let's consider two datasets a and b. In dataset a we have an ID called PATID and in b an ID called PATNO merge(a,b, by.x="PATID", by.y="PATNO")
  - The code above will do the merging for us. If the ID-variables are called the same we can use just by=id-variable



# Merging datasets (3)

- With *all.x* and *all.y* options in the *merge*-function we can determine do we want append observations from the datasets that doesn't have a pair in the other datasets.
  - NAs will be printed into the columns of the dataset,
     where a certain ID is missing.
- Data is practically always stored in more than one place, so data analyst has to do quite a lot of merging.



### Vocabulary for variables

- During the course we use a set of adjectives to determine the attributes of a single variable.
  - Integer variables.
    - These variables are numbers, where there is now decimals. Eg. Number of children
  - Numeric variables.
    - These variables can get whatever number value
  - Categorical variables
    - These variables have some pre-defined classes and there cannot be any other values.



## -Vocabulary for variables (2)

- For example gender, age-category, etc.
  - These variables are stored in R with numbers 1,2,3... And are then given level-names. R calls these variables factors.
- Character variables
  - These variables are stored in as text. No calculations can be done with character variables.

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