Exercises

Descriptive analysis and basic statistics in biomedical studies using R and Markdown

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1 R basics

1. Create a vector x with the following entries:

```
3 4 1 1 2 1 4 2 1 1 5 3 1 1 1 2 4 5 5 3
```

- 2. Check which elements of x are equal to 1 (Hint: use "==" operator).
- 3. Create a variable (e.g vector) y having the logarithm of x.
- 4. Create a vector z containing the first five elements of x.
- 5. Create a sequence of numbers from 1 to 20 in steps of 0.2 (see help seq function).
- 6. Concatenate x and y into a vector called newVec.
- 7. Display all objects in the workspace and then remove newVec object.
- 8. Create a data.frame called elasticband having these two variables.

```
stretch <- c(46,54,48,50,44,42,52)
distance <- c(148,182,173,166,109,141,166)
```

- 9. Compute the mean of the variable stretch from the elasticbanc object (Hint: use "\$").
- 10. Which is the class of the variable distance?

2 Descriptive analyses

The file multicentric.txt contains the data of a multicentric case/control study to discover risk factors of cervical cancer. The file multicentric.htm contains a description of the variables included in this file.

- 1. Import the data in an object called multi (Hint: data are in text tab-delimited format: use read.delim and na.string=c(" ")).
- 2. Create anoter variable called edad1sex.cat having age at first sexual intercourse (variable edad1sex) in three categories: <15, 15-18, 19-25, +25. How many woman had her first sexual intercouse before 15 years old?

Exercises 2

- 3. How many individuals and variables are in this database?
- 4. Describe the types of variables you have.
- 5. Create a table describing the number of cases and controls (variable status) of each country (variable pais) included in the study. Which is the country having more cases? Which is the percentage of cases in Brazil?
- 6. Create another table describing the number of cases and controls and human papillomavirus (variable vph). Do you think those variables are associated? Why?
- 7. Summarize the duration consuming oral contraceptives (variable durco) in cases and controls, in each country and by educative levels (variable niveledu), respectively.
- 8. Categorize the variable number of pregnancies (variable nembara) in quartiles. Which is the number of cases in the last quartile of number of pregnancies? and controls?
- 9. Create a boxplot to compare the age at first pregnancy (variable edademba) between cases and controls (variable status). Do you think this variable could be a risk factor of cervical cancer?
- 10. Create an histogram to describe the age when women received their first test of papillomavirus (variable edad1pap). Is this variable normally distributed? Could you apply a test to justify your answer? (Hint -> google)

3 Hypothesis testing

1. Using the same study on cervical cancer ... perform a chi-squared test or a t-test to assess which variables are associated with status variable (e.g case/control). NOTE: use the appropriate test when analyzing continuous variables.

4 Statistical modelling

- 1. Fit univariate logistic regression models to investigate which factors (edad, niveledu, fumar, edad1sex, nembara, vph) are associated with cervical cancer (variable status).
- 2. Select the set of variables that better predicts the probability of being diagnosed with cervival cancer. (NOTE: use complete cases 'multi.comp <- multi[complete.cases(multi),]).
- 3. [Advanced]. Compute the area under the roc curve (AUC) of the selected model (Hint: investigate pROC package).
- 4. The file retinol.doc describes a study to investigate which are the determinants of retinol observed in plasmatic concentrations since low levels of those micronutrients are associated with some types of tumors. Let's try to decipher whihe are those determinants. To this end:
 - Fit single regresion models to assess association between retinol measured at plasma (variable retplas) and variables: retdiet, colest, alcohol, fibra, grasa, calorias, vitamin and fumador.
 - Select the most significant association and test model assumptions and check linearity.
 - Use an automatic method to select the set of variables that better predict the variable retplas. Which is the percentage of variability explained by this model?
- 5. The file pulmon.doc describes a study to investigate which are the most predictive variables of survival in patients diagnosed with cancer. Perform the all the required analyses to answer this scientific question (e.g. Kaplan-Meier, log-rank, Multivariate Cox models).

5 Graphics

- 1. Load the data available in the file retinol.txt
- 2. Create a histogram and a boxplot of retplas variable and plot them side-by-side on the same graphing region. Label the axes accordingly. Save your results as a Jpeg file.
- 3. Plot retplas (y-axis) versus retdiet (X-axis) using an appropriate plotting command. Put a title on the graph and labels on the axes.

Exercises 3

4. Fit a linear regresion model between both variables (call this regression mod). Add the estimated regression line to the current plot and make it the colour blue.

- 5. Extract the values of the residuals using resids <- resid(mod). Check that the residuals are normally distributed by creating a Q-Q plot.
- 6. Create the same plot for each of the three levels of the variable fumador. (Hint: Use coplot).
- 7. Build the same plot separated for males and females (variable sexo). (Hint: use xyplot from package lattice).
- 8. Construct a histogram of calorias and overlay the density curve. (Hint: Need hist, lines and density.)
- 9. Create in one figure (e.g. use par(mfrow=c(2,1))) the histogram of variable calorias for males and females (variable sexo).

6 Reproducible Research

- 1. Get the R code to answer the questions in the section **R basics** that are available here: (https://github.com/isglobal-brge/R_course/blob/master/Answer_exercises/Exercise_1.R) and create a Markdown document having each task in a section
- 2. Create the same report for the answers to the exercises of section **Graphics** that are available here (https://github.com/isglobal-brge/R_course/blob/master/Answer_exercises/Exercise_1.R)
- 3. Load the multicentric.txt data into R and create (using compareGroups package):
 - A table describing the variables: edad, niveledu, fumar, edad1sex, nembara, vph by cases and controls (variable status).
 - A table giving the ORs of developing cervical cancer (variable status) for the variables: edad, niveledu, fumar, edad1sex, nembara, vph
- 4. Load the pulmon.txt data into R and create:
 - A table describing the survival (variables tiempo and estado) for the variables: sexo, edad4, estclin, ik5, cirugia, quimio, radioter.
 - A table giving the HR of mortality (variables tiempo and estado) for the variables sexo, edad4, estclin, ik5, cirugia, quimio, radioter.