SDA 2023 — Assignment 2

Note: Parts of Exercises 2.3 and 2.4 can only be solved after the lecture on Feb. 22 or after you have read the remainder of Chapter 3 in the syllabus!

For the exercises of Assignment 2 you can use the *R*-functions summary, range and IQR. For QQ-plots you may use qqnorm, as well as the local functions qqt, qqlnorm, qqchisq, qqlogis, qqexp, qqunif, qqcauchy, qgompertz, dgompertz, pgompertz, rgompertz, which can be found on the Canvas page (functions_Ch3.txt)¹.

Also, you can use the *R*-functions ks.test and shapiro.test, and the function chisquare that can be found on the Canvas page for this assignment. (The *R*-function chisq.test should *not* be used for chi-square tests for goodness of fit.) Investigate these functions before using them.

Note: to indicate a normal distribution with expectation 2 and variance 25, we use the notation $\mathcal{N}(2,25)$, whereas R uses the parameters mean=2, and sd=5 for this normal distribution.

When you want to find a concrete member of a location-scale family, you will need the expectation and variance of the theoretical distribution which is your reference distribution in the location-scale family. Because only then you can match the theoretical expectation and variance with the corresponding sample mean and variance, respectively, to solve for the shift and scale parameters. Please look up the expectation and variance of your theoretical distribution, e.g., somewhere on the internet.

When performing a statistical test, state the null and alternative hypothesis, present the test statistic and its distribution under the null hypothesis (if it is a well-known distribution), give the value of the test statistic, the critical region or the p-value and the chosen significance level, and formulate the conclusion of the test.

Make a concise report of your answers in *one single PDF file*, with only *relevant R* code *in an appendix*. It is important to make clear in your answers <u>how</u> you have solved the questions. Graphs should look neat (label the axes, give titles, use correct dimensions etc.). Multiple graphs can be put into one figure using the command par(mfrow=c(k,1)), see help(par). Sometimes there might be additional information on what exactly has to be handed in. Read the file AssignmentFormat.pdf on Canvas carefully.

These functions can be loaded in exactly the same way as the code from the file sample2023.txt, see the General information on page 1

General information: How to load executable R code from a .txt file:

You can load the data that should be analyzed in Exercises 2.2 and 2.3 by performing the following steps:

- i) save the files sample2023a.txt and sample2023b.txt to some directory called thedirectory with path path²,
- ii) set the working directory to thedirectory by using the command setwd("path") (instead of path, you obviously need to fill in the correct path on your computer!),
- iii) finally read the data by using the commands source("sample2023a.txt") and source("sample2023b.

General information: How to load data into your workspace:

```
Use one of the following read commands, depending on your data type:

data <- scan("path/....txt")²,

data <- read.table("path/....txt")²,

data <- read.csv("path/....csv")²,

install.packages("xlsx")

library(xlsx)

data <- read.xlsx("path/....xlsx", sheetIndex=1)².
```

The functions qqnorm, qqt, qqlnorm, etc., can be used to make QQ-plots for the location-scale families of the normal, t, lognormal distributions, etc., respectively. The argument df in qqt and qqchisq is used to set the number of degrees of freedom in the t-distribution and the χ^2 -distribution.

Exercise 2.1 (Class Exercise on Feb. 17) To get an idea of what QQ-plots look like and how to use the commands for making QQ-plots, you will apply some of these functions to some samples drawn from different distributions.

²For Windows the path is usually C:/.../thedirectory, for Mac the path is usually /Users/.../thedirectory.

Exercise 2.2

- a. Make plots of the quantile functions (that is, make 'true' QQ-plots as in Figure 3.4 of the syllabus or Slide "Quantiles of F and $F_{a,b}$ (2)" of Lecture 2) for the following pairs of distributions:
 - I. lognormal with parameters meanlog=0 and sdlog=2 against exponential with rate 2.
 - II. normal with mean 2 and variance 16 against t_3 .
 - III. chi-squared with 20 degrees of freedom against chi-squared with 3 degrees of freedom.

Comment for each plot on the heaviness of the tails of the two distributions. The tails of a distribution can be seen as the relative height of its density f(x) for $x \to \pm \infty$. For example, the right tail of an exponential distribution is heavier than the right tail of a uniform distribution (which vanish for some sufficiently large x > 0). You could create for your own use some density plots to get a better idea of the tail behaviour of the different distributions (none of these density plots should be handed in).

Note: for this exercise you should not generate random samples. Instead, use the true quantile functions for both the x-axis and the y-axis (for example, the R-function qnorm can be used for computation of the quantiles of a normal distribution). For plotting the function plot should be used, not the function qqplot.

b. Investigate the data in sample2023a.txt with the given functions for making QQ-plots and find an appropriate distribution for this data set. Apart from specifying a suitable location-scale family (e.g., "normal distributions" or "location-scale family with respect to lognormal distributions with sdlog parameter 0.1"), also give values for the location and scale parameters. (e.g., "normal distribution with location 2 and scale 5", or " $\mathcal{N}(2,25)$ distributed", or "distributed like $2+5\cdot Y$, where Y follows a lognormal distribution with meanlog=1 and sdlog=0.1").

Hint: Using the commands qqline and abline can be helpful! (See Lecture 2 for more details.) Note that slope and intercept of qqline are not the parameters a and b of the location-scale family.

Note: See the first page for instructions on how to load executable R code from .txt files.

Hand in: your plots and comments for part a and plots of relevant graphs of part b, as well as a motivation for your trials and your final conclusion for part b. (Not all trials need to be documented, just the ones that really led you to the final model.)

General information on the .RData file to be submitted to the dummy assignment "RData A2" on Canvas:

Create in R a list mylist that contains the required entries as specified in the exercise(s) which are marked as "(partially) .RData file hand-in". You should store your list in an .RData file by using the R command save(mylist, file="[PATH]/myfile2_[GROUP_NO].RData"), where [PATH] stands for the path on your computer where you wish to save the .RData file and [GROUP_NO] stands for the number of the assignment group you have chosen on Canvas). E.g., myfile2_91.RData for group 91.

In any case, mylist must have the entry stud_no: a vector that contains the student numbers of you and your group partner.

Exercise 2.3 (partially .RData file hand-in)

- a. Explore the sample in sample2023b.txt graphically and find an appropriate distribution (i.e. a specific member of a location-scale family) from which this sample could have been drawn. Indicate location and scale as well.
 - Note: See the first page for instructions on how to load executable R code from .txt files.
- b. Test at level $\alpha=5\%$ whether the sample originates from the Gompertz distribution³ with rate parameter b=0.7 and shape parameter $\eta=0.07$. Use the Kolmogorov–Smirnov test for this.
- c. Do the same as in part b, but now use a chi-square test (instead of the Kolmogorov-Smirnov test) for testing the goodness-of-fit. Choose the arguments of the function chisquare so that the condition for the rule of thumb (see syllabus) is fulfilled. Hint: the function qgompertz could be useful for ensuring the rule of thumb condition.
- d. Explain whether the results from parts b and c agree, and interpret the results. If you find they do not agree, find a reason why this might be so.

Hand in (partially in the main report, partially as .RData file):

a.: relevant graphs, results, and comments in the main report.

b.: the following entries of your list mylist in R, which will be stored in your .RData file:

- ks.score: the value of the Kolmogorov-Smirnov test statistic applied to the data,
- ks.p.value: the resulting p-value of the Kolmogorov-Smirnov test,
- ks.reject = TRUE or FALSE: the test decision, whether the null hypothesis is rejected.

c.: the following entries of your list mylist in R, which will be stored in your .RData file:

- chisq.breaks: the breaks, i.e. a vector specifying the lower and upper boundaries of the intervals used by the Chisquared-goodness-of-fit test; see Lecture 3 for details.
- chisq.score: the value of the Chisquared test statistic applied to the data,
- chisq.p.value: the resulting p-value of the Chisquared test,
- chisq.reject = TRUE or FALSE: the test decision, whether the null hypothesis is rejected.

d.: your answers, with reference to the test outcomes in b. and c. (repeat here what the outcomes were), in the main report.

³A Gompertz-distributed random variable $X \sim GO(\eta, b)$ with rate b > 0 and shape $\eta > 0$ has the density function $x \mapsto b\eta \exp(\eta + bx + e^{bx})$. The parameter 1/b has a scale interpretation.

Exercise 2.4 The file body.dat.txt contains several body measurements (and additional information) of 507 individuals (mainly) in their twenties and thirties, all of them doing sport exercises for several hours per week. In this exercise, we focus on the ankle circumferences (in cm; column 20) and the body mass indices (BMIs), which are weight (in kg) divided by the squared height (in meter). Column 23 and 24 respectively contain the weight (in kg) and height (in cm) of the individuals. For this exercise we only use the rows 248 to 507 of the dataset which correspond to all female individuals.

Note: See the first page for instructions on how to load data from .txt files.

- a. First obtain the BMIs from the weights and heights. Next, make histograms and boxplots of the BMIs and ankle circumferences and conclude whether both data distributions have approximately the same shape.
- b. Investigate whether or not the BMIs and ankle circumferences could be from the same location-scale family. Use the function qqplot for a two sample QQ-plot.
- c. Without the use of hypothesis tests, discuss and conclude whether the BMIs and ankle circumferences (each considered as univariate dataset) could be modelled as a member of a location-scale family with respect to normal distributions or some suitably chosen chi-squared distribution.
 - Note: in any case, it is not required to identify location and scale parameters of the best fitting location-scale family.
- d. Investigate the normality of the following ratio of measurements, without using a hypothesis test: BMIs divided by squared ankle circumferences. (That is, a new dataset is considered which contains for each individual the just-described ratio of measurements.)
- e. Use the Shapiro–Wilk test with significance level $\alpha = 5\%$ to test the normality of the new data described in d.
- f. Compare the outcomes of the goodness-of-fit tests for normality for the full sample of the new data from d. and for the first 50 of these data points, and also use histograms of these two samples to complement your analyses. Find a possible explanation for these outcomes of the tests.

Note: in general for testing normality based on the Shapiro-Wilk test, one would just use the full sample and conduct the test only once (per sample). This exercise is just meant to give us a better understanding of how goodness-of-fit tests work. In this exercise we assume that the first 50 data points are representative for all others.

Hand in: relevant graphs, results and answers to the questions, and your comments.