

# Systems Biology course: R programming exercises

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## 1 Exercises: R programming

**Where to find and how to install R** If you do not have R installed on your machine, you would need to download it from <http://www.r-project.org/>. From the homepage, simply click on CRAN, Packages and select your favorite mirror (e.g. Austria). Download the precompiled binary distribution by clicking on your operating system. R comes with a simple editor, feel free to use it or use your favorite editor (e.g. Eclipse, Xcode, ....) to write your code.

**Notes** The exercises below cover distinct basic topics of R programming. Each of the following exercise requires both the concept presented in the tutorial as well as independent reading of the R manual (<http://cran.r-project.org/doc/manuals/R-intro.pdf>) and of the R documentation. Recall that each built-in function in R has an associated help page that can be accessed as `?name` (e.g. `?paste`).

We strongly suggest to get familiar with the R data types (e.g. vectors, data frames, lists, ...) and the methods used to construct and manipulate them.

## 2 Beginner level

**Exercise 1** Create a vector containing all odd numbers smaller than or equal to 75 i) using a for loop and logical operators; ii) using the built-in `seq` function.

**Exercise 2** Create a function called `isEven` that takes an integer as argument and returns TRUE (boolean) if the number is even.

**Exercise 3** Create a function called `sum10` that given a vector of 10 numbers, return their sum. Try to implement it using a for loop.

**Exercise 4** Same as ex3, but use R basic functions.

**Exercise 5** Given a string longer than 8 characters, extract the 4-prefix and the 4-suffix (e.g. `SystemsBiology` will give you *Syst* and *logy*)

**Exercise 6** Given a string with your name, compute its reverse

**Exercise 7** Given a string composed of two words separated by a blank space, return a string obtained by extracting and pasting (with no space) the first 3 characters from each of the words (e.g. from `'Systems Biology'` you would obtain `SysBio`). Can you extend it to any number of sequences?

**Exercise 8** We provide a dataset (in CSV format, `data.csv`). You can take a look at the first lines of the file using the Linux command `head` as you learned in the previous lecture. First, read in the file into R (hint: check out the function `read.csv2`). Then, plot the data as scatter plot and read the documentation on `plot` (e.g. check out `plot` and `par` for specifying plotting parameters).

**Exercise 9** Use the dataframe you loaded in the previous exercise to create a new dataframe that contains only the first 10 rows of the original one and that has an additional column containing the element-wise product of the extracted rows. The resulting dataframe should look like:

	speed	dist	prod
1	4	2	8
2	4	10	40
3	7	4	28
4	7	22	154
5	8	16	128
6	9	10	90
7	10	18	180
8	10	26	260
9	10	34	340
10	11	17	187

hint: To solve this problem, read a bit on `cbind`.

**Exercise 10** Write the dataframe you obtained in ex9 to a file using your favorite function (hint: check out either `write.table` or `write.csv`, `write.delim`, etc). Make sure that you can load the file back into the R session.

### 3 Intermediate level

**Exercise 11** Produce a scatter plot of `wind` vs. `temperature` for the `airquality` dataset. What do we learn, i.e. what relationship do we see? Can you customize your plot? (hint: see `par`) Can we fit a simple linear regression model (hint: see `lm`) to the data and add it to the plot?

**Exercise 12** Have a look at the `ChickWeight` data set. Plot the weight as a function of time for chick 7. Superimpose a line representing the same information for chick 9. Add a legend to the plot. Find out how to plot in a pdf device (see, e.g. `?pdf`) and plot the previous graph to a pdf file. Extract all weights at time point  $t = 21$  and arrange your result in a list where each slot corresponds to a Diet (there are 4 regimes, so it will be a list of 4 vectors). Hint: see `?split`. Boxplot the weights in this 4 diets. Any difference in mean weight between diet 1 and 3? How could you go about testing this?

**Exercise 13** Have a look at the `eurodist` data set. What is the class of this object? Do you know it? Sort the cities by increasing distance from Geneva. Hint: you may want to look at functions that allow you to coerce an object to a different class. Specifically, see `?as.matrix` here.

**Exercise 14** Write your own functions to compute mean and standard deviation of a numeric vector. Apply them to the `precip` data sets and cross check your results using the built-in `mean` and `sd` functions.

**Exercise 15** Write your own function to transpose a matrix. Use of the built-in `t` function is considered cheating!

**Exercise 16** Write the `swiss` dataset as csv file. Then, read the file into the console. How long does it take to execute the command?

**Exercise 17** Write a function that takes a numeric vector as input and returns a vector of differences between consecutive elements. Is there a way to do the same using a built-in function? Apply both functions to the following input

```
v <- 1 : 10
v <- c(3,4,5,1,2,9,11,8)
v <- seq(1,100,2)
```

**Exercise 18** Create three vectors containing the first 10 numbers divisible by 2, by 3 and by 4, respectively. Use these vectors to create a list, and give suitable names to the slots. Now that you have a list, can you compute the sum for each slot? If you sum the three slots, you get a vector of length 10. What is it?

**Exercise 19** Write a function to convert Fahrenheit to Celsius.

**Exercise 20** Write a function which takes a single numeric value  $x$  as argument and returns a logical value `TRUE` if  $x$  is larger than 5, `FALSE` otherwise. Then, extend the previous function by including the threshold ( $t$ ) as an argument. Finally, given `l <- list( seq(1,75, by = 3) )` use the function you created in `l` to test each single entry in `l`. Set  $t = 15$ .