

Exercises

Exercise 2: Basic R operations

Read Chapter 2 to help you complete the questions in this exercise.

1. Open up your RStudio Project from Exercise 1 and either create a new R script or continue with your previous R script. Remember to save your R script with a suitable name (exercise_2?). Make sure you include any metadata you feel is appropriate (title, description of task, date of script creation etc). Don't forget to comment out your metadata with a `#` at the beginning of the line.
2. Let's use R as a fancy calculator. Find the natural log, log to the base 10, log to the base 2, square root and the natural antilog of 12.43. See Section 2.1 of the Introduction to R book for more information on mathematical functions in R. Don't forget to write your code in RStudio's script editor and source the code into the console.
3. Next, use R to determine the area of a circle with a diameter of 20 cm and assign the result to an object called `area_circle`. If you can't remember how to create and assign objects see Section 2.2 or watch this video. Google is your friend if you can't remember the formula to calculate the area of a circle! Also, remember that R already knows about `pi`. Don't worry if you're stumped and feel free to ask one of the instructors for guidance.
4. Ok, you're now ready to explore one of R's basic (but very useful) data structures - vectors. A vector is a sequence of elements (or components) that are all of the same data type (see Section 3.2.1 for an introduction to vectors). Although technically not correct it might be useful to think of a vector as something like a single column of data in a spreadsheet. There are a multitude of ways to create vectors in R but you will use the concatenate function `c()` to create a vector called `weight` containing the weight (in kg) of 10 children: 69, 62, 57, 59, 59, 64, 56, 66, 67, 66 (Section 2.3 or watch this video for more information). These ten children are a toy example, deliberately small enough that you can check every result by eye. Next week you'll meet `cardiacdata.xlsx`, a real study of 163 patients, where you won't have that luxury and will have to trust your code instead.
5. Now you can do some useful stuff to your `weight` vector. Get R to calculate the mean, variance, standard deviation, range of weights and the number of children of your `weight` vector (see Section 2.3 for more details). Now read Section 2.4 of the R book to learn how to work with vectors. After reading this section you should be able to extract the weights for the first five children using Positional indexes and store these weights in a new variable called `first_five`. Remember, you will need to use the square brackets `[ ]` to extract (aka index, subset) elements from a variable.

6. We're now going to use the `c()` function again to create another vector called `height` containing the height (in cm) of the same 10 children: 112, 102, 83, 84, 99, 90, 77, 112, 133, 112. Use the `summary()` function to summarise these data in the `height` object. Extract the height of the 2nd, 3rd, 9th and 10th child and assign these heights to a variable called `some_child` (take a look at the section Positional indexes in the R book if you're stuck). We can also extract elements using Logical indexes. Let's extract all the heights of children less than or equal to 99 cm and assign to a variable called `shorter_child`.

7. Now you can use the information in your `weight` and `height` variables to calculate the body mass index (BMI) for each child. The BMI is calculated as weight (in kg) divided by the square of the height (in metres).

$$bmi_i = \frac{weight_i}{height_i^2}$$

Store the results of this calculation in a variable called `bmi`. Note: you don't need to do this calculation for each child individually, you can use both vectors in the BMI equation – this is called vectorisation (see Section 2.4.4 of the Introduction to R book).

8. Now let's practice a very useful skill - creating sequences (honestly it is...). Take a look at Section 2.3 in the R book (the bit on creating sequences) to see the myriad ways you can create sequences in R. Let's use the `seq()` function to create a sequence of numbers ranging from 0 to 1 in steps of 0.1 (this is also a vector by the way) and assign this sequence to a variable called `seq1`. Once you've done that, see if you can figure out how to create a sequence going the other way, from 10 to 1 in steps of 0.5, and assign it to a variable called `seq2`. Hint: you may find it useful to include the `rev()` function in your code (use the search facility in the Introduction to R book to search for '`rev`').

9. Now for a different way of building a vector. Generate the following two sequences. You will need to experiment with the arguments of the `rep()` function to generate them (see Section 2.3 for some clues). The two sequences repeat their elements in different ways, so pay attention to which argument you need in each case:

- 1 2 3 1 2 3 1 2 3
- a a a c c c e e e g g g

10. OK, back to the variable `height` you created in Q6. Sort the values of `height` into ascending order (shortest to tallest) and assign the sorted vector to a new variable called `height_sorted`. Take a look at Section 2.4.3 in the R book to see how to do this. Now sort all heights into descending order and assign the new vector a name of your choice.

11. Let's give the children some names. Create a new vector called `child_names` with the following names of the 10 children: "Alfred", "Barbara", "James", "Jane", "John", "Judy", "Louise", "Mary", "Ronald", "William".

12. A really useful (and common) task is to order the values of one variable by the order of another variable. To do this you will need to use the `order()` function in combination with the square bracket notation `[ ]`.

Have a peep at Section 2.4.3 for some details. Create a new variable called `names_sort` to store the names of the children ordered by child height (from shortest to tallest). Who is the shortest, and who is the tallest? If you're not sure how to do this, please ask one of the instructors.

13. Almost there! In R, missing values are usually represented with an `NA`. Missing data can be tricky to deal with in R (and in statistics more generally) and cause some surprising behaviour when using some functions. Take a look at Section 2.4.5 of the R book for more information about missing values. To explore this a little further let's create a vector called `mydata` with the values 2, 4, 1, 6, 8, 5, `NA`, 4, 7. Notice the value of the 7th element of `mydata` is missing and represented with an `NA`. Now use the `mean()` function to calculate the mean of the values in `mydata`. What does R return? If you're confused by this output take a look at the help page for the function `mean()`. Can you figure out how to alter your use of the `mean()` function to calculate the mean ignoring this missing value?

14. Finally, list all variables in your workspace that you have created in this exercise. Remove the variable `seq1` from the workspace using the `rm()` function.

End of Exercise 2