

Creating your own functions

Learning the basics of R - Part 3

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Outline

- Why write functions
- When to write functions
- How to write functions
- Practical session

Why write functions

- Allow automation of common tasks in a more powerful and general way than *copy-and-pasting*,
 - You can give a function an evocative name that makes your code easier to understand;
 - As requirements change, you only need to update code in one place, instead of many.
 - You eliminate the chance of making incidental mistakes when you copy and paste (i.e. updating a variable name in one place, but not in another).
- Fewer global variables: When you run a function, the intermediate variables that it creates are not stored in your global environment. This saves memory and keeps your global environment cleaner.
- Better documentation: Well documented functions help the user understand the steps of your processing.
- Easier to maintain / edit: When you create a function for a repeated task, it is easy to edit that one function. Then every location in your code where that same task is performed is automatically updated.

When to write functions

You should consider writing a function whenever you've copied and pasted a block of code more than twice (i.e. you now have three copies of the same code).

For example, take a look at this code.

What does it do?

```
df <- data.frame(  
  a = c(1, 2, 1, 1, 1, 2, 1, 1, 2, 2),  
  b = c(2, 2, 2, 1, 1, 1, 1, 2, 1, 2),  
  c = c(1, 2, 1, 1, 2, 1, 2, 1, 2, 2)  
)  
  
df$a <- ifelse(df$a == 2, 0, df$a)  
df$b <- ifelse(df$b == 2, 0, df$b)  
df$c <- ifelse(df$c == 2, 0, df$c)
```

When to write functions

Original data:

##		a	b	c
## 1		1	2	1
## 2		2	2	2
## 3		1	2	1
## 4		1	1	1
## 5		1	1	2
## 6		2	1	1
## 7		1	1	2
## 8		1	2	1
## 9		2	1	2
## 10		2	2	2

Recoded data:

##		a	b	c
## 1		1	0	1
## 2		0	0	0
## 3		1	0	1
## 4		1	1	1
## 5		1	1	0
## 6		0	1	1
## 7		1	1	0
## 8		1	0	1
## 9		0	1	0
## 10		0	0	0

This is a good example of when writing a function will be useful/beneficial.

How to write functions

We can create a function called `recode_values()`:

```
recode_values <- function(x) {  
  ifelse(x == 2, 0, x)  
}
```

And apply it to the same data as follows:

```
df$a <- recode_values(df$a)  
df$b <- recode_values(df$b)  
df$c <- recode_values(df$c)
```

We get:

```
##      a b c  
## 1  1 0 1  
## 2  0 0 0  
## 3  1 0 1  
## 4  1 1 1  
## 5  1 1 0  
## 6  0 1 1  
## 7  1 1 0  
## 8  1 0 1  
## 9  0 1 0  
## 10 0 0 0
```

How to write functions

- You need to pick a name for the function. In the example I used `recode_values` because this function recodes the values based on a specified rule (i.e., value of 2 is converted to 0).
- You list the inputs, or **arguments**, to the function inside `function`. Here we have just one argument. If we had more the call would look like `function(x, y, z)`.
- You place the code you have developed in body of the function, a `{` block that immediately follows `function(...)`.

Questions?

Practical session

We'll work through *Exercise 2 - Manipulating objects and creating new functions* in Practical R for Epidemiologists (<https://practical-r.org/exercise2.html>) as a GitHub Classroom assignment

Thank you!

Slides can be viewed at <https://oxford-ihtm.io/open-reproducible-science/session4.html>

PDF version of slides can be downloaded at <https://oxford-ihtm.io/open-reproducible-science/pdf/session4-r-basics-part3.pdf>

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