

Programming Topics II: Writing Functions



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BSDS 100 - Intro to Data Science with R



- What is a function?
- Writing your own functions
- Using functionals



- We have already used many functions, including `mean()`, `seq()`, `c()`, `data.frame()`, and `ggplot()`
- Just about anything we have used that is followed by parentheses `()` is a function
- Functions tell R to perform a specific task



- Each function has several parts
 - 1 A name (e.g., `mean`)
 - 2 A list of arguments often called `formals()`, these can be mandatory (e.g., `mean()` needs a vector of numbers to find the mean of, or optional (e.g., `mean()` can take the argument `na.rm = T`)
 - 3 A `body()`, which is the code a function executes



- Recall, we can find a list of a functions arguments with ?
- Any argument in the documentation followed by = carries a default value, that means if the user doesn't fill in that argument, the function can still run using the default (i.e., the argument is optional)
- We can also see the code inside the function by calling it's name without the ()



- We can write our own functions using assignment and the word "function"

```
name_of_func <- function(arguments_of_function) {  
  body_of_function }
```

- To the left of the assignment symbol we give it a name, inside the parentheses following the word "function" we list the arguments, and inside the curly brackets we write code
- A simple example

```
my_square <- function(x) {  
  x^2 }
```



- In \mathbb{R} whatever the last line of the body to run is what the function will "return", or report to the user
- **Exception!** If the last line of a function body is an assignment the function will return nothing, be sure to avoid this so your function returns something



- Functions are useful because they allow you to repeat the same set of commands without copy and pasting code, which can be error prone and time consuming
- Functions can contain much more complex code than we are showing here, including calling other functions, looping, or if/then statements



- Let's write a function that takes a numeric vector, removes any numbers lower than 1 or higher than 26, then returns the letter of the alphabet that matches each number in the resulting vector



- So far, we have shown functions that take one argument, but functions can also take more than one argument
- This function returns the sum of two numbers a user passes it

```
sum_two <- function(a, b){  
  a + b }
```



- Arrange the follow lines to make a function named `subtract_two` that takes two numbers and subtracts the smaller number from the larger number (or returns zero if they are equal)
 - 0
 - `b - a`
 - `a - b`
 - `subtract_two <- function(a,b){`
 - `if(a == b){`
 - `} else if(a < b){`
 - `} else if(b < a){`
 - `}`
 - `}`
- Test your new function on a couple values to make sure it works!



- Write a function named `square_df` that takes a vector and returns a data frame with two columns where
 - the first column is named "original" and is the original vector
 - the second column is named "squared" and is each of the elements of the original vector squared
- Test your function on a couple vectors to make sure it works!



- We *could* write a function in the following manner:

```
x <- 1  
return_one_two <- function() {  
  y <- 2  
  c(x, y) }
```

- Notice how the `x` is written outside of the function even though it is used inside the function?
- But then, what would happen if we forgot to define `x`?



- In this case we would say x is **globally** scoped while y is **locally** scoped
- We would also say x is defined in the **global environment** while y is defined in the **local environment**
- Functions should be written with all variables locally scoped or passed as arguments to avoid errors
- How would we fix the previous function?



- We write a function which returns a number (chosen by the user) divided by 3, we initially write the function like so:

```
user_chosen_number <- 10  
three <- 3  
num_divide_three <- function(){  
  user_chosen_number / three }
```

- Rewrite the function so that the user passes their chosen number as an argument and three is locally scoped



- It is important to distinguish between the formal and actual arguments of a function
- **Formal arguments** are a property of the function

Arithmetic Mean

Description

Generic function for the (trimmed) arithmetic mean.

Usage

```
mean(x, ...)
```

```
## Default S3 method:
```

```
mean(x, trim = 0, na.rm = FALSE, ...)
```

Arguments

- `x` An R object. Currently there are methods for numeric/logical vectors and [date](#), [date-time](#) and [time interval](#) objects. Complex vectors are allowed for `trim = 0`, only.
- `trim` the fraction (0 to 0.5) of observations to be trimmed from each end of `x` before the mean is computed. Values of `trim` outside that range are taken as the nearest endpoint.
- `na.rm` a logical value indicating whether NA values should be stripped before the computation proceeds.
- `...` further arguments passed to or from other methods.



- It is important to distinguish between the formal and actual arguments of a function
- **Actual or calling arguments** can vary each time you call a function

```
> mean(x = 1:10)
[1] 5.5
```

```
> mean(x = 99:999)
[1] 549
```

- In the above examples, the calling arguments are `1:10` and `99:999` respectively



- It is also possible to define default values for your function
- The default values will be used if the user doesn't specify a value



```
# w/o default values
myFunc_10 <- function(a, b) {
  c(a, b)
}

> myFunc_10()
Error in myFunc_10() : argument "a" is missing, with no default

# with default values
myFunc_11 <- function(a = 1, b = 2) {
  c(a, b)
}

> myFunc_11()
[1] 1 2
```



Function arguments in R can be defined in terms of other arguments

```
myFunc_12 <- function(a = 1, b = a * 2) {  
  c(a, b)  
}
```

```
> myFunc_12()  
[1] 1 2
```

```
> myFunc_12(111)  
[1] 111 222
```

```
> myFunc_12(99, 100)  
[1] 99 100
```



The last expression evaluated in a function becomes the return value

```
myFunc_18 <- function(xyz) {  
  if (xyz < 10) {  
    0  
  } else {  
    10  
  }  
}
```

```
> myFunc_18(5)  
[1] 0
```

```
> myFunc_18(10)  
[1] 10
```

To `return()` or not to `return()`



- The last expression evaluated in a function is the return value
- You can always wrap the final expression in `return()` if you choose
- Using `return()` makes the code very slightly slower
- In simplistic functions, R programmers will typically omit `return()`
- In longer, more complicated functions, `return()` is often used when it makes the code easier to read

To return() or not to return()



```
# simple function, does not require a return()
```

```
myFunc_15 <- function(x){  
  x + 10  
}
```

```
# a more complex function benefits visually from having return()  
#   but does not require return()
```

```
myFunc_18 <- function(xyz) {  
  if (xyz < 10) {  
    return(0)  
  } else {  
    return(10)  
  }  
}
```



- In addition to saving you time copying and pasting code, functions can be used with "functionals" to avoid loops
- Because loops (for, while, and repeat) are slow in R avoiding them is useful
- Also, functionals can make your code more readable



The `apply()` family of functionals are often used in lieu of *for* loops, coming in a variety of flavors (not exhaustive)

Functional	Input	Output
<code>apply()</code>	Array/Matrix	Vector/Array
<code>lapply()</code>	Vector/List	List
<code>sapply()</code>	Vector/List	List
<code>vapply()</code>	Vector/List	Vector

There is also a functional `do.call` which is very flexible and can take and output most types. We will focus on the two most commonly used functionals, `apply()` and `lapply()`



- `apply()` is useful for applying the same function to every row or every column of a matrix
- e.g., `apply(test_matrix, 1, mean)` would find the mean of each row in `test_matrix`



- What if we want to include `na.rm = T` in our `apply()`?
- Just adding it doesn't work
- Adding it with the placeholder doesn't work
- What should we do?



- First option, make a custom function, say, `na_mean()` that does what we want
- Second option, make an anonymous or lambda function

```
apply(test_matrix, 1, function(x){mean(x, na.rm  
= T)})
```
- The anonymous function doesn't have a name, and can't be called again later, but this can be a useful method if you don't anticipate reusing the function



- Use `apply()` to get the standard deviation, removing NA's, from the test matrix
- You can either define a new function or use an anonymous function



- `lapply()` takes and returns a list
- This is useful to us because R considers data frames to be a type of list!
- Consider the following:



- Assume you are given the following data frame

```
> myDataFrame_01
```

	A	B	C	D	E	G
1	1	6	1	5	-99	1
2	10	4	4	-99	9	3
3	7	9	5	4	1	4
4	2	9	3	8	6	8
5	1	10	5	9	8	6
6	6	2	1	3	8	5

- Your objective is to replace all of the -99s with NAs



- You could—but shouldn't—iterate through each column manually, e.g.

```
myDataFrame_01$A[myDataFrame_01$A == -99] <- NA
myDataFrame_01$B[myDataFrame_01$B == -99] <- NA
...
myDataFrame_01$F[myDataFrame_01$F == -99] <- NA
```




- 1 It's easy to make copy-paste mistakes
- 2 It will take you a really long time
- 3 If you need to change the code later, you will have to change many lines of code rather than just a few



Let's write a function with the objective of replacing all `-99`s in a single column with `NA`s

```
fix99s_byCol <- function(myCol) {  
  myCol[myCol == -99] <- NA  
}
```

- Will the code above work as intended? Hint: **no**. Why not?



Let's write a function with the objective of replacing all `-99`s in a single column with `NA`s

```
fix99s_byCol <- function(myCol) {  
  myCol[myCol == -99] <- NA  
}
```

- Will the code above work as intended? Hint: **no**. Why not?



The following **does** work as intended:

```
fix99s_byCol <- function(myCol) {  
  myCol[myCol == -99] <- NA  
  myCol  
}  
  
myDataFrame_01$A <- fix99s_byCol(myDataFrame_01$A)  
...  
myDataFrame_01$F <- fix99s_byCol(myDataFrame_01$F)
```

- This reduces but doesn't eliminate the potential for errors
- There is no gain in efficiency (repetitive code is still required)
- We can instead use an `lapply()`



lapply() example

```
fix99s_byCol <- function(myCol) {  
  myCol[myCol == -99] <- NA  
  myCol  
}  
  
> myDataFrame_02 <- lapply(myDataFrame_01, fix99s_byCol)  
  
> str(myDataFrame_02)  
List of 6  
 $ A: num [1:6] 1 10 7 2 1 6  
 $ B: num [1:6] 6 4 9 9 10 2  
 $ C: num [1:6] 1 4 5 3 5 1  
 $ D: num [1:6] 5 NA 4 8 9 3  
 $ E: num [1:6] NA 9 1 6 8 8  
 $ F: num [1:6] 1 3 4 8 6 5
```

- This almost worked...but not quite

lapply() example



Here are two ways to correct the previous function call so that it returns a data frame

```
❶ > myDataFrame_03 <-  
  as.data.frame(lapply(myDataFrame_01,  
    fix99s_byCol))  
  
❷ > myDataFrame_01[] <- lapply(myDataFrame_01,  
  fix99s_byCol)
```

Note, option 2 only works when replacing the old data frame with the new values, not for making a brand new data frame



- First, make a function that takes a column and changes all values 4 or smaller to 0 and all values 5 or larger to 10 by filling in the blanks below:

```
big_small <- function(myCol) {  
  myCol[myCol < ____] <- ____  
  myCol[myCol > ____] <- ____  
  ____ }  
}
```

- Then, use your new function and `lapply()` to change `myDataFrame_01` into `myDataFrame_0_10`, where all the numbers have been transformed to 0s and 10s