

Topic 1 Interlude: Writing functions

Basics of coding in R

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Context

How are we using R?

- Code is modular, nothing is “from scratch”
- **Functions** are the basic building block of what we do.

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Functions

- We have used a lot of built-in functions: `mean()`, `subset()`, `plot()`, `read.table()`...
- An important part of programming and data analysis is to write custom functions
- Functions help make code **modular**
- Functions make debugging easier
- Remember: this entire class is about applying *functions* to *data*

What is a function?

A *function* is a machine that turns **input objects** (arguments) into an **output object** (return value) according to a definite rule.

- Let's look at a really simple function

```
addOne <- function(x) {  
  x + 1  
}
```

- x is the **argument** or **input**
- The function **output** is the input x incremented by 1

```
addOne(12)
```

```
[1] 13
```

More interesting example

- Here's a function that returns a % given a numerator, denominator, and desired number of decimal values

```
calculatePercentage <- function(x, y, d) {  
  decimal <- x / y # Calculate decimal value  
  round(100 * decimal, d) # Convert to % and round to d digits  
}
```

```
calculatePercentage(27, 80, 1)
```

```
[1] 33.8
```

- If you're calculating several %'s for your report, you should use this kind of function instead of repeatedly copying and pasting code

Function returning a list

- Here's a function that takes a person's full name (FirstName LastName), weight in lb and height in inches and converts it into a list with the person's first name, person's last name, weight in kg, height in m, and BMI.

```
createPatientRecord <- function(full.name, weight, height) {  
  name.list <- strsplit(full.name, split=" ")[[1]]  
  first.name <- name.list[1]  
  last.name <- name.list[2]  
  weight.in.kg <- weight / 2.2  
  height.in.m <- height * 0.0254  
  bmi <- weight.in.kg / (height.in.m ^ 2)  
  list(first.name=first.name, last.name=last.name,  
        weight=weight.in.kg, height=height.in.m,  
        bmi=bmi)  
}
```

Trying out the function

```
createPatientRecord("Michael Smith", 185, 12 * 6 + 1)
```

```
$first.name
```

```
[1] "Michael"
```

```
$last.name
```

```
[1] "Smith"
```

```
$weight
```

```
[1] 84.09091
```

```
$height
```

```
[1] 1.8542
```

```
$bmi
```

```
[1] 24.45884
```


Another example: 3 number summary

- Calculate mean, median and standard deviation

```
threeNumberSummary <- function(x) {  
  c(mean=mean(x), median=median(x), sd=sd(x))  
}  
x <- rnorm(100, mean=5, sd=2) # Vector of 100 normals  
                               # with mean 5 and sd 2  
threeNumberSummary(x)  
  
      mean   median      sd  
5.162890 5.246974 1.758908
```

If-else statements

- Oftentimes we want our code to have different effects depending on the features of the input
- Example: Calculating a student's letter grade
 - If grade ≥ 90 , assign A
 - Otherwise, if grade ≥ 80 , assign B
 - Otherwise, if grade ≥ 70 , assign C
 - In all other cases, assign F
- To code this up, we use if-else statements

If-else Example: Letter grades

```
calculateLetterGrade <- function(x) {  
  if(x >= 90) {  
    grade <- "A"  
  } else if(x >= 80) {  
    grade <- "B"  
  } else if(x >= 70) {  
    grade <- "C"  
  } else {  
    grade <- "F"  
  }  
  grade  
}  
  
course.grades <- c(92, 78, 87, 91, 62)  
sapply(course.grades, FUN=calculateLetterGrade)  
  
[1] "A" "C" "B" "A" "F"
```

return()

- In the previous examples we specified the output simply by writing the output variable as the last line of the function
- More explicitly, we can use the `return()` function

```
addOne <- function(x) {  
  return(x + 1)  
}
```

```
addOne(12)
```

```
[1] 13
```

- We will generally avoid the `return()` function, but you can use it if necessary or if it makes writing a particular function easier.
- Google's style guide suggests explicit returns. Most do not.

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