# Functions - part 2 STAT 133

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#### **Functions**

R comes with many functions and packages that let us perform a wide variety of tasks. Sometimes, however, there's no function to do what we want to achieve. In these cases we need to create our own functions.

# Writing Functions

### Writing Functions

#### Writing Functions

- Choose meaningful names of functions
- Preferably a verb
- Choose meaningful names of arguments
- Think about the users (who will use the function)
- Think about extreme cases
- ▶ If a function is too long, maybe you need to split it

#### Names of functions

#### Avoid this:

```
f <- function(x, y) {
  x + y
}</pre>
```

#### This is better

```
add <- function(x, y) {
  x + y
}</pre>
```

#### Names of arguments

Give meaningful names to arguments:

```
# Avoid this
area_rect <- function(x, y) {
   x * y
}</pre>
```

#### This is better

```
area_rect <- function(length, width) {
  length * width
}</pre>
```

## Meaningful Names to Arguments

#### Avoid this:

```
# what does this function do?
ci <- function(p, r, n, ti) {
  p * (1 + r/p)^(ti * p)
}</pre>
```

# Meaningful Names to Arguments

#### Avoid this:

```
# what does this function do?
ci <- function(p, r, n, ti) {
 p * (1 + r/p)^(ti * p)
}</pre>
```

#### This is better:

```
# OK
compound_interest <- function(principal, rate, periods, time) {
  principal * (1 + rate/periods)^(time * periods)
}</pre>
```

# Meaningful Names to Arguments

```
# names of arguments
compound_interest <- function(principal = 1, rate = 0.01,</pre>
                               periods = 1, time = 1)
 principal * (1 + rate/periods)^(time * periods)
compound_interest(principal = 100, rate = 0.05,
                  periods = 5, time = 1)
compound_interest(rate = 0.05, periods = 5,
                  time = 1, principal = 100)
compound_interest(rate = 0.05, time = 1,
                  periods = 5, principal = 100)
```

### Describing functions

Also add a short description of what the arguments should be like. In this case, the description is outside the function

```
# function for adding two numbers
# x: number
# y: number
add <- function(x, y) {
   x + y
}</pre>
```

### Describing functions

In this case, the description is inside the function

```
add <- function(x, y) {
    # function for adding two numbers
    # x: number
    # y: number
    x + y
}</pre>
```

### Describing functions

- One type of functions very common in R are binary operators, eg:
  - -2 + 5 (sum)
  - 3 / 2 (division)
  - a %in% b (value matching)
  - X %\*% Y (matrix multiplication)
- Binary operators are actually functions
- ▶ These functions take two inputs—hence the term *binary*
- It is possible to define your own binary operators

#### Example:

```
# addition operator
2 + 3
# equivalent to
'+'(2, 3)
```

```
# binary operator
"%p%" <- function(x, y) {
   paste(x, y, sep = " ")
}
'good' %p% 'morning'
## [1] "good morning"</pre>
```

## How to create a binary operator?

- ► A binary operator is defined as one or more characters surrounded by percent symbols %
- When defining the function, the entire name must be quoted
- ► Include two arguments
- As usual, avoid using names of existing operators:
  - "%%", %\*%, %/%, %o%, %in%

### Another example

#### Here's another example:

```
# binary operator
"%u%" <- function(x, y) {
  union(x, y)
}
1:5 %u% c(1, 3, 5, 7, 9)
## [1] 1 2 3 4 5 7 9</pre>
```

# Lazy Evaluation

## Lazy Evaluation

Arguments to functions are evaluated lazily, that is, they are evaluated only as needed:

```
g <- function(a, b) {
  a * a * a
}
g(2)
## [1] 8</pre>
```

g() never uses the argument b, so calling g(2) does not produce an error

### Lazy Evaluation

#### Another example

```
g <- function(a, b) {
  print(a)
  print(b)
g(2)
## [1] 2
## Error in print(b): argument "b" is missing, with no
default
```

Notice that 2 got printed before the error was triggered. This is because b did not have to be evaluated until after print(a)

# Messages

There are two main functions for generating warnings and errors:

- ▶ stop()
- warning()

There's also the stopifnot() function

#### Stop Execution

Use stop() to stop the execution of a function (this will raise
an error)

```
meansd <- function(x, na.rm = FALSE) {
  if (!is.numeric(x)) {
    stop("x is not numeric")
  }
  # output
  c(mean = mean(x, na.rm = na.rm),
    sd = sd(x, na.rm = na.rm))
}</pre>
```

#### Stop Execution

Use stop() to stop the execution of a function (this will raise
an error)

```
# ok
meansd(c(4, 5, 3, 1, 2))

## mean sd
## 3.000000 1.581139

# this causes an error
meansd(c('a', 'b', 'c'))

## Error in meansd(c("a", "b", "c")): x is not numeric
```

## Warning Messages

Use warning() to show a warning message

```
meansd <- function(x, na.rm = FALSE) {
  if (!is.numeric(x)) {
    warning("non-numeric input coerced to numeric")
    x <- as.numeric(x)
  }
  # output
  c(mean = mean(x, na.rm = na.rm),
    sd = sd(x, na.rm = na.rm))
}</pre>
```

A warning is useful when we don't want to stop the execution, but we still want to show potential problems

## Warning Messages

Use warning() to show a warning message

```
# ok
meansd(c(4, 5, 3, 1, 2))
##
       mean
                 sd
## 3.000000 1.581139
# this causes a warning
meansd(c(TRUE, FALSE, TRUE, FALSE))
## Warning in meansd(c(TRUE, FALSE, TRUE, FALSE)):
non-numeric input coerced to numeric
##
                 sd
        mean
## 0.5000000 0.5773503
```

#### Stop Execution

stopifnot() ensures the truth of expressions:

```
meansd <- function(x, na.rm = FALSE) {
  stopifnot(is.numeric(x))
  # output
  c(mean = mean(x, na.rm = na.rm),
    sd = sd(x, na.rm = na.rm))
}
meansd('hello')
## Error: is.numeric(x) is not TRUE</pre>
```

# **Environments and Functions**

#### Consider this example

```
w <- 10
f <- function(y) {</pre>
  d <- 5
  h <- function() {</pre>
    d * (w + y)
  return(h())
f(2)
## [1] 60
```

How / Why does f() work?

#### Consider this other example

```
w <- 10

f <- function(y) {
    d <- 5
    return(h())
}

f(2)

## Error in f(2): could not find function "h"</pre>
```

Why f() does not work?

#### Environments

- All the variables that we create need to be stored somewhere
- ► The place where they are stored is called an **environment**
- R works with environments, all of which are in (virtual) memory
- Usually, we don't need to explicitly deal with environments
- Environments are nested

#### Global Environment

- ► The user workspace is the **global environment**
- ▶ The global environment is the **top level** environment
- ▶ It is formally referred to as R\_GlobalEnv
- Variables defined in the global environment can be seen from anywhere
- ► The contents of the global environment are listed with ls()

```
# top level environment
environment()
## <environment: R_GlobalEnv>
```

#### Searching objects

- ▶ When R tries to bind a value to a symbol, it searches through a series of environments to find the appropriate value
- ► To retrieve the value of an object the order is:
- Search the current environment
- Search the global environment for a symbol name matching the one requested
- Search the namespaces of each of the packages on the search list: search()

#### **Environments and Functions**

- A function consists not only of its arguments and body but also of its environment
- ► An environment is made up of the collection of objects present at the time the function comes into existence
- When a function is created by evaluating the corresponding expression, the current environment is recorded as a property of the function

# Let's go back to our first example

```
w <- 10
f <- function(y) {</pre>
  d < -5
  h <- function() {</pre>
    d * (w + y)
  return(h())
f(2)
## [1] 60
```

How does f() work?

### Let's see the environments

```
w <- 10 # variable (in global environment)
# a function (in global environment)
f <- function(y) {</pre>
  d <- 5 # local variable
  h <- function() {  # subfunction</pre>
   d * (w + y) # w is a free variable
 return(h())
environment(f)
## <environment: R GlobalEnv>
```

### Function Environment

- ▶ w is a global variable (in global environment)
- ▶ f() is a function in the global environment
- d is a local variable—local to f()
- ▶ h() is a subfunction—local to f()
- ▶ w is not an argument but a free variable

### Let's see the environments

```
f <- function(y) {</pre>
  d < -5
  h <- function() {</pre>
    d * (w + y)
  print(environment(h)) # h()'s environment
  return(h())
environment(f)
## <environment: R_GlobalEnv>
f(2)
## <environment: 0x7fc08c2d46d0>
## [1] 60
```

### Your turn

```
# When executed, what does g() return?
x <- 5

g <- function(x = FALSE) {
    y <- 10
    list(x = x, y = y)
}
g()</pre>
```

- A) x = 5, y = 10
- B) x = 0, y = 10
- C) x = FALSE, y = 10
- D) x = FALSE, y = 5

## Variable's Scope

- ► A variable's **scope** is the set of places from which you can see the variable
- R will try to find a variable in the current environment
- ▶ If it doesn't find them it will look in the parent environment
- And then that environment's parent
- And so on until it reaches the global environment

## Variable's Scope

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- And then that environment's parent
- And so on until it reaches the global environment

## Variable Scope

▶ When we define a variable inside a function, the rest of the statements in that function will have access to that variable

## Variable Scope

```
f <- function(x) {</pre>
  y <- 1
  g <- function(x) {</pre>
    (x + y) / 2
  g(x)
f(5)
## [1] 3
```

g() is a subfunction that have access to y in f's environment.

## Variable Scope

```
f <- function(x) {</pre>
  y <- 1
  g(x)
g <- function(x) {</pre>
  (x + y) / 2
f(5)
## Error in g(x): object 'y' not found
```

g() is a function that doesn't have access to y; g() can only see things in the global environment

## One more thing ...

#### Let's look at another exmaple

```
mean(1:5)

## [1] 3

mean

## function (x, ...)

## UseMethod("mean")

## <bytecode: 0x7fc089fcab48>

## <environment: namespace:base>
```

## One more thing ...

### You can do things like this

```
# confusing but it works
mean <- 1:5
mean(mean)
## [1] 3</pre>
```

### Some issues

#### You can also do things like this

```
# not a good idea but you can do it
mean <- function(x) 2*x + 5

mean(1:5)
## [1] 7 9 11 13 15</pre>
```

It seems we've lost the original mean() function

## The :: Operator

#### :: operator to the rescue

```
# my mean
mean(1:5)

## [1] 7 9 11 13 15

# base mean
base::mean(1:5)

## [1] 3
```

Here we use the name espace base of the R package "base" to access the original mean()

# Your Turn

R has a function summary() that when applied on a numeric vector provides something like this:

```
## Min. 1st Qu. Median Mean 3rd Qu. Max.
## 1.00 3.25 5.50 5.50 7.75 10.00
```

Create a describe() function that takes a numeric vector and returns: minimum, maximum, mean, and standard deviation

#### First attempt

```
describe <- function(x) {</pre>
  x_{\min} \leftarrow \min(x)
 x \max < -\max(x)
 x_{mean} \leftarrow mean(x)
 x_sd \leftarrow sd(x)
  return(c(x_min, x_max, x_mean, x_sd))
describe(1:10)
##
    [1] 1.00000 10.00000 7.00000 9.00000 11.00000 13.00000 15.
##
    [9] 19.00000 21.00000 23.00000 25.00000 3.02765
```

#### Second attempt (adding names)

```
describe <- function(x) {</pre>
  x \min < - \min(x)
 x_max \leftarrow max(x)
 x_mean <- mean(x)</pre>
 x sd \leftarrow sd(x)
  values <- c(x_min, x_max, x_mean, x_sd)</pre>
  names(values) <- c("min", "max", "mean", "sd")</pre>
  return(values)
describe(1:10)
                                   sd <NA> <NA> <NA>
##
        min
              max mean
## 1.00000 10.00000 7.00000 9.00000 11.00000 13.00000 15.00000
## <NA> <NA> <NA> <NA>
## 19.00000 21.00000 23.00000 25.00000 3.02765
```

### Third attempt (using a list as output)

```
describe <- function(x) {
    list(
        min = min(x),
        max = max(x),
        mean = mean(x),
        sd = sd(x)
    )
}</pre>
```

```
describe(1:10)
## $min
## [1] 1
## $max
## [1] 10
##
## $mean
## [1] 7 9 11 13 15 17 19 2
##
## $sd
## [1] 3.02765
```

Probability Density of the Normal Distribution:

$$f(x|\mu,\sigma) = \frac{1}{\sigma\sqrt{2\pi}}e^{-\frac{(x-\mu)^2}{2\sigma^2}}$$

Write a function that takes a value x (with parameters  $\mu$  and  $\sigma$ ) which computes the probability density distribution of the normal distribution

#### Normal Distribution:

```
normal_dist <- function(x, mu = 0, sigma = 1) {
  constant <- 1 / (sigma * sqrt(2*pi))
  constant * exp(-((x - mu)^2) / (2 * sigma^2))
}
normal_dist(2)
## [1] 0.05399097</pre>
```

## Argument Matching

```
normal_dist <- function(x, mu = 0, sigma = 1) {
  constant <- 1 / (sigma * sqrt(2*pi))
  constant * exp(-((x - mu)^2) / (2 * sigma^2))
}
normal_dist(2)
normal_dist(2, sigma = 3, mu = 1)
normal_dist(mu = 1, sigma = 3, 2)
normal_dist(mu = 1, 2, sigma = 3)</pre>
```

## Argument Matching

R is "smart" enough in doing pattern matching with arguments' names (not recommended though)

```
normal_dist(2)
## [1] 0.05399097
normal_dist(2, m = 0, s = 1)
## [1] 0.05399097
normal_dist(2, sig = 1, m = 0)
## [1] 0.05399097
```