

# **PROG102: Functions**

**Writing your own functions in R**

**MARINCS 100B | Intro to Marine Data Science | Winter 2025**

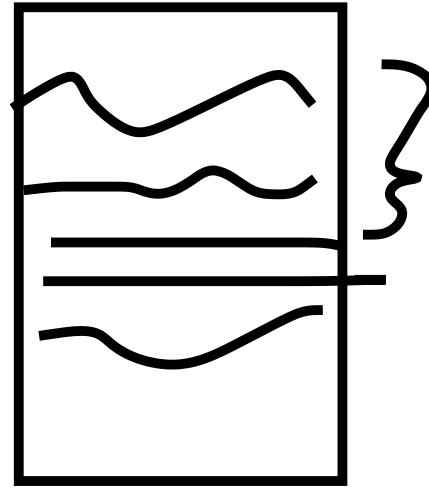
## **Key concepts**

Functions have two purposes:

- 1) hide the details \*encapsulation\*
- 2) Apply the same code to new inputs \*reusability\*

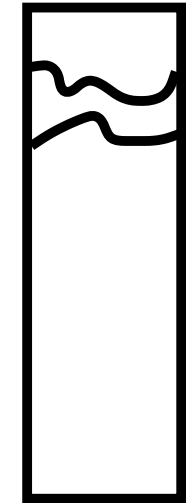
**Easy to read**

Cognitive load  
<7 items at a time

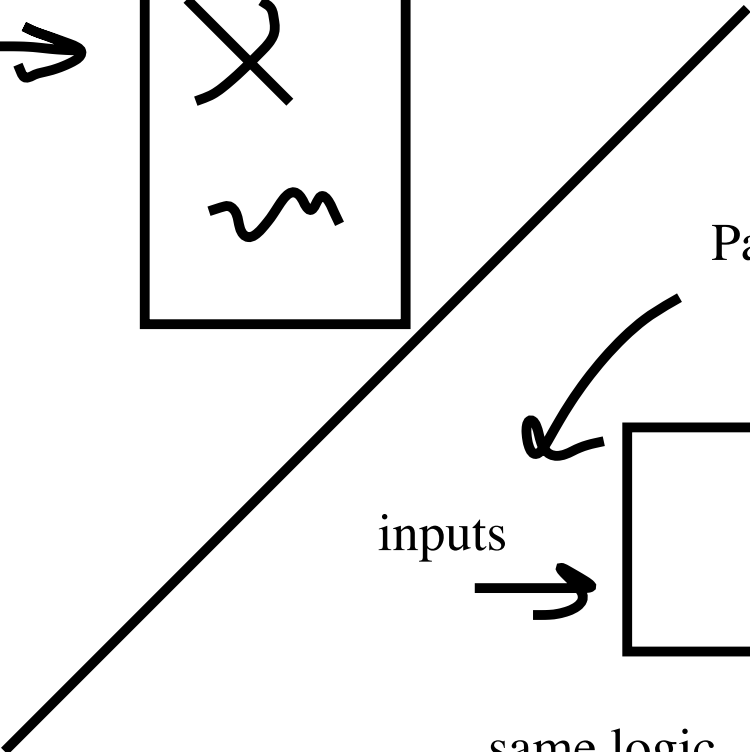
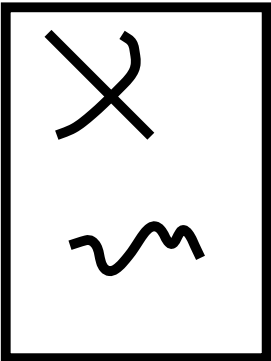
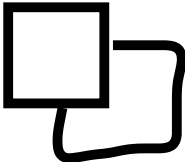


**Reusable**

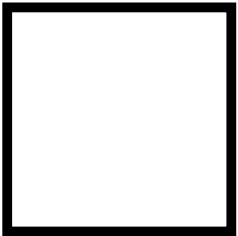
Copy paste edit work flow



Prog 101



Parameters



inputs



ouputs



same logic  
new inputs new outputs

# Syntax

Functions have five parts

- 1) name
- 2) keyword function
- 3) parameters (parentheses)
- 4) body in curly brackets
- 5) return output

**Demo in R**

## Recap

Functions make code readable by hiding details (encapsulation)

Functions make code reusable by allowing different inputs (parameters)

syntax- every function definition has 5 parts

## **New vocabulary and lingering questions**

New vocabulary

Lingering questions



## Exercises

Label the five parts of this function:

- 1) assigns first\_and\_last to be the function name
- 2) makes first character

```
first_and_last <- function(s) {  
  first_char <- substr(s, 1, 1)  
  last_char <- substr(s, nchar(s), 1)  
  result <- paste(first_char, last_char)  
  return(result)  
}
```

## Exercises

Match the function bodies on the left with the name that describes what they're doing on the right.

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

double

```
function(a) {  
  result <- a * 2  
  return(result)  
}
```

hypotenuse\_length

```
function(a, b) {  
  c_squared <- a^2 + b^2  
  result <- sqrt(c_squared)  
  return(result)  
}
```

increment

## Exercises

Write a function that turns a vector into a palindrome. For example, it should turn 1 2 3 into 1 2 3 3 2 1. Hint: you'll have to use a function called `rev()`. Choose a short but descriptive name for your function.

# **PROG102: Functions**

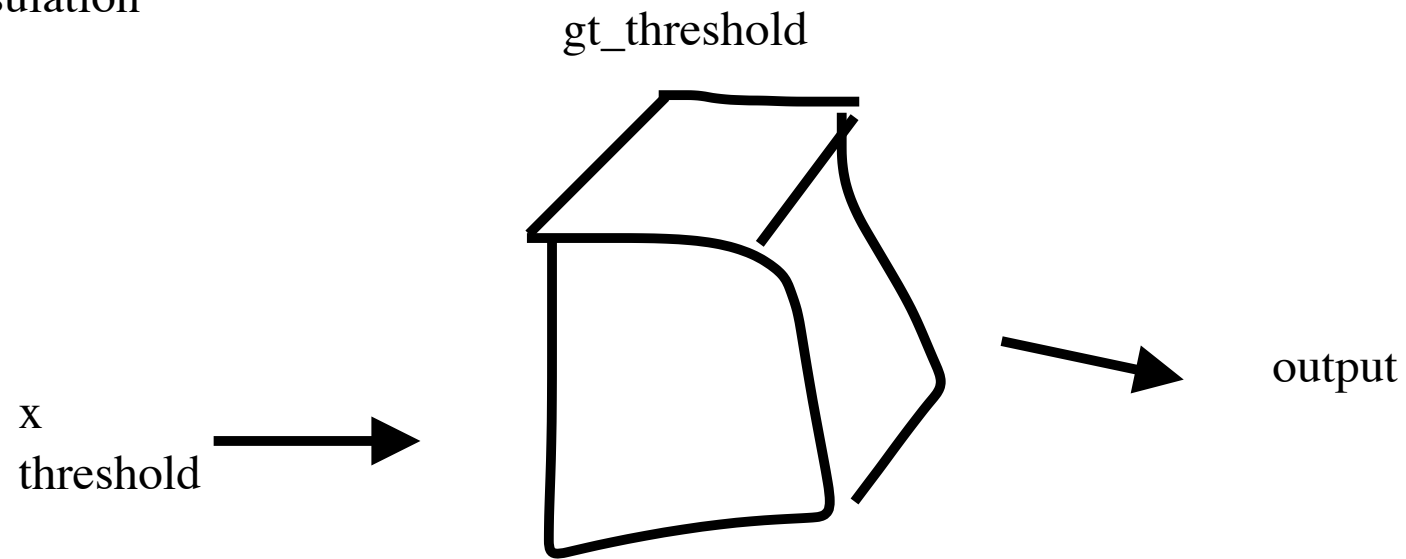
**How functions execute**

## **Key concepts**

- 1) functions act as black boxes separate universe
- 2) parameters and returns, those are our bridges into and out of the black box
- 3) debugger- a useful way to peak inside the black box

## The black box

Encapsulation



**Demo in R**

## Recap

Functions operate in their own little universe “black box”

Parameters are how we let information in

`return()` is how we let information back out



## **New vocabulary and lingering questions**

New vocabulary

Lingering questions

## Exercises

- What value does the following code yield?
- How could you change `fish_mass` so the code yields 12 instead?
- How could you change the body of the function so the code yields 12?

```
fish_mass <- 5
temperature <- 20
fish_growth <- function(mass, temp) {
  growth <- 2 + 0.2 * temp
  mass <- mass + growth
  return(mass)
}
fish_growth(fish_mass, temperature)
```

## Exercises

In your own words, why does running this code generate an error?

```
calc_volume <- function(height, width, depth) {  
  area <- height * width  
  volume <- area * depth  
  return(volume)  
}  
vol <- calc_volume(3, 5, 1)  
area
```

# **PROG102: Functions**

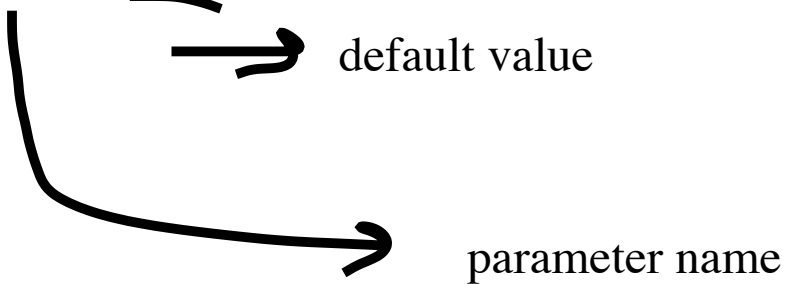
**Default and named parameters**

## **Key concepts**

- 1) parameters usually enter in order- by position
- 2) default parameter values allow you to omit certain values
- 3) named parameters let you skip around in order
- 4) default and named parameters are usually options

## Default and named parameters

`round(x, digits = 0)`



default value

parameter name

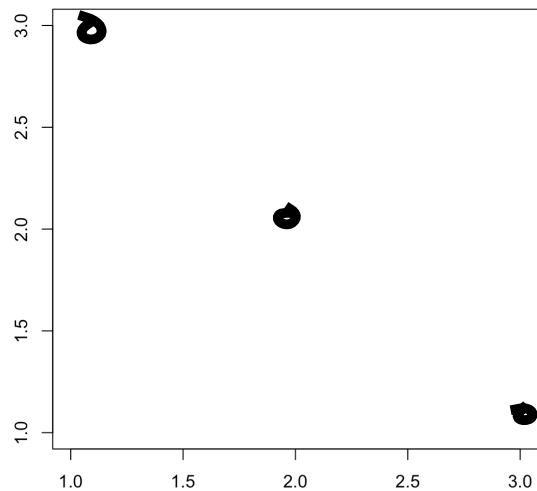
`round(pi)` —> 3 \* use the default “by position”  
`round(pi,0)` —>3 “by position”  
`round(digits =0, pi)` —>3

## Long parameter lists

```
plot(x, y = NULL, type = "p", xlim = NULL, ylim = NULL,  
     log = "", main = NULL, sub = NULL, xlab = NULL, ylab = NULL,  
     ann = par("ann"), axes = TRUE, frame.plot = axes,  
     panel.first = NULL, panel.last = NULL, asp = NA,  
     xgap.axis = NA, ygap.axis = NA,  
     ...)
```

X y  
plot(c(1, 2, 3), c(3, 2, 1))

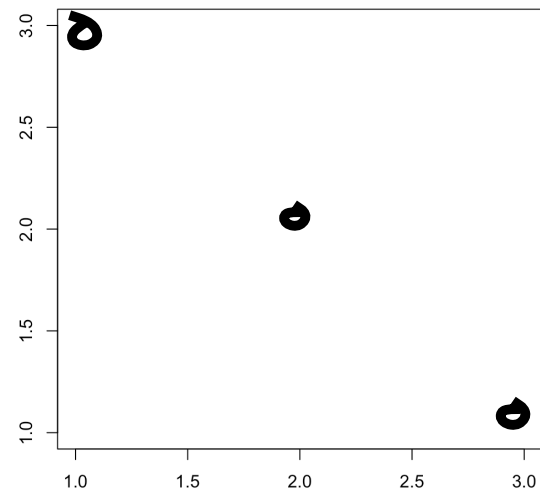
c(1, 2, 3)



c(1, 2, 3)

X y  
plot(c(1, 2, 3), c(3, 2, 1),  
 xlab = "x", ylab = "y")

y



x

**Demo in R**

skip



## Triple dots

```
max(..., na.rm = FALSE)
```

```
paste(..., sep = " ", collapse = NULL, recycle0 = FALSE)
```

ignore the triple dots and focus on the name of the function

```
max(1,2,3) -> 3
```

```
paste ("water", "is", "wet") —> "water is wet"
```

## Recap

- 1) named and default parameters are useful for modifying how functions work
- 2) default values allow omission
- 3) named parameters allow us to skip around

## **New vocabulary and lingering questions**

New vocabulary

Lingering questions

## Exercises

R represents *missing* data with the value NA. Say you're doing an experiment and you miss the second observation. In R you can write that as `c(1, NA, 3, 4)`.

Most summary functions, like `mean()`, `max()`, and `median()`, have a parameter called `na.rm`. What does this parameter do? What is its default value? How would you get the maximum value of the vector `c(1, NA, 3, 4)`?