PROG102: Functions

Writing your own functions in R

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

Key concepts

2 Purposes: 1)Hide the detailsencapsulation 2) Apply the same code to new inputs

Easy to read

Cognitive load <7 items

Taking a bunch of info and putting it into easy to read format whileremaining professional encapsulation

Reusable

Use parameters -Inputs --- Outputs New inputs gives us new outputs

Syntax

- 1) Name
- 2) Function 3) Parameters ()
- 4) Body {} 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 fuction definition has five parts

New vocabulary and lingering questions

New vocabulary	Lingering questions
parameters Encapsulation	

Label the five parts of this function:

```
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)
}</pre>
```

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

increment

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

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

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Key concepts

The black box

Demo in R

Recap

New vocabulary and lingering questions

New vocabulary	[Lingering questions

- 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)</pre>
```

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</pre>
```

PROG102: Functions

Default and named parameters

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Key concepts

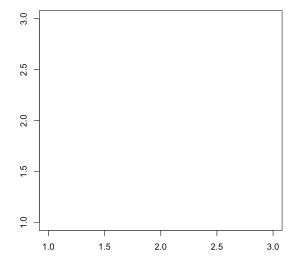
Default and named parameters

round(x, digits = 0)

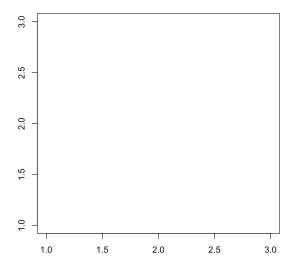
Long parameter lists

```
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,
    ...)
```

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



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



Demo in R

Triple dots

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

Recap

New vocabulary and lingering questions

New vocabulary	[Lingering questions

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)?