

GR5206: lecture 2

Computational Statistics
And Introduction to Data Science

Outline



1 Control flow

2 Choices

3 Loops

4 Functions

Control flow



- Two primary tools of control flow:
 - Choices.
 - Loops.
- Choices:
 - ► E.g. if(), ifelse(), switch().
 - Allows to run different code depending on the input.
- Loops:
 - E.g. for, while, repeat.
 - Allows to repeatedly run code, typically with changing options.
- You might want to have a look at chapter 8 for condition system (messages, warnings, and errors)...

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- The basic idea for if statements:
 - If condition is TRUE, true_action is evaluated.
 - If condition is FALSE, the optional false_action is evaluated.

```
if (condition) true_action
if (condition) true_action else false_action
```

■ Typically, actions are compound statements contained within {.

```
grade <- function(x) {
   if (x > 90) {
     "A"
   } else if (x > 80) {
     "B"
   } else if (x > 50) {
     "C"
   } else {
     "F"
   }
}
```



- if returns a value so that you can assign the results:
 - Only do that when it fits on one line; otherwise hard to read.

```
x1 <- if (TRUE) 1 else 2
x2 <- if (FALSE) 1 else 2
c(x1, x2)
#> [1] 1 2
```

- When using if without else:
 - Returns NULL if the condition is FALSE.
 - Useful with functions like c()/paste() dropping NULL inputs.

```
greet <- function(name, birthday = FALSE) {
  paste0("Hi ", name, if (birthday) " and HAPPY BIRTHDAY")
}
greet("Maria", FALSE)
#> [1] "Hi Maria"
greet("Jaime", TRUE)
#> [1] "Hi Jaime and HAPPY BIRTHDAY"
```

Invalid inputs



■ The condition should evaluate to a single TRUE or FALSE:

```
if ("x") 1
#> Error in if ("x") 1: argument is not interpretable as logical
if (logical()) 1
#> Error in if (logical()) 1: argument is of length zero
if (NA) 1
#> Error in if (NA) 1: missing value where TRUE/FALSE needed
```

- The exception (frequent source of bugs, avoid):
 - ▶ A logical vector of length greater than 1 generates a warning.

```
if (c(TRUE, FALSE)) 1
#> Warning in if (c(TRUE, FALSE)) 1: the condition has length > 1 and
#> only the first element will be used
#> [1] 1
```

■ In R >=3.5.0+, you can turn this into an error (good practice):

```
Sys.setenv("_R_CHECK_LENGTH_1_CONDITION_" = "true")
if (c(TRUE, FALSE)) 1
#> Error in if (c(TRUE, FALSE)) 1: the condition has length > 1
```



- if only works with a single TRUE or FALSE.
- What if you have a vector of logical values?
- Answer: ifelse(), a vectorised function with test, yes, and no vectors (recycled to the same length).
 - Missing values are propagated into the output.
 - Advice: use ifelse() only when the yes and no vectors (otherwise hard to predict the output type).

```
x <- 1:10
ifelse(x %% 5 == 0, "XXX", as.character(x))
#> [1] "1" "2" "3" "4" "XXX" "6" "7" "8" "9" "XXX"
ifelse(x %% 2 == 0, "even", "odd")
#> [1] "odd" "even" "odd" "even" "odd" "even" "odd"
#> [10] "even"
```



Lets you replace code like:

```
x_option <- function(x) {
   if (x == "a") {
       "option 1"
   } else if (x == "b") {
       "option 2"
   } else {
       stop("Invalid `x` value")
   }
}</pre>
```

■ With:

```
x_option <- function(x) {
  switch(x,
    a = "option 1",
    b = "option 2",
    stop("Invalid `x` value")
)
}</pre>
```

switch() statements cont'd



- Last component should always throw an error.
- When multiple inputs share an output:
 - Use empty right hand sides of =.
 - Same as C's switch statement.

```
(switch("c", a = 1, b = 2))
#> NULL
legs <- function(x) {</pre>
  switch(x,
    cow = ,
    horse = .
    dog = 4,
    human = .
    chicken = 2,
    plant = 0,
    stop("Unknown input")
legs("cow")
#> [1] 4
legs("dog")
#> [1] 4
```

■ switch() with a numeric x is not recommended.

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Loops



for loops are used to iterate over items in a vector.

```
for (item in vector) perform_action
```

For each item in vector, perform_action is called once; updating the value of item each time.

```
for (i in 1:3) {
   print(i)
}
#> [1] 1
#> [1] 2
#> [1] 3
```

- When iterating over indices, use very short variable names like i, j, or k by convention.
- Important: for assigns the item to the current environment.

```
i <- 100
for (i in 1:3) {}
i
#> [1] 3
```

Early termination



- Two ways to terminate a for loop early:
 - next exits the current iteration.
 - break exits the entire for loop.

```
for (i in 1:10) {
   if (i < 3)
        next

print(i)

if (i >= 5)
        break
}
#> [1] 3
#> [1] 4
#> [1] 5
```

Common pitfalls



- Three common pitfalls to watch out for when using for:
 - Preallocation.
 - lteration over e.g. 1:length(x).
 - Iteration over S3 vectors.
- Preallocation:
 - If you're generating data, preallocate the output.
 - Otherwise the loop will be very slow.
 - vector() function is helpful.

```
means <- c(1, 50, 20)
out <- vector("list", length(means))
for (i in 1:length(means)) {
  out[[i]] <- rnorm(10, means[[i]])
}</pre>
```

Common pitfalls cont'd



Next, beware of iterating over 1:length(x), which will fail in unhelpful ways if x has length 0.

```
means <- c()
out <- vector("list", length(means))
for (i in 1:length(means)) {
   out[[i]] <- rnorm(10, means[[i]])
}
#> Error in rnorm(10, means[[i]]): invalid arguments

# The reason? `:` works with both increasing and decreasing sequences.
1:length(means)
#> [1] 1 0
```

Use seq_along(x) instead.

```
seq_along(means)
#> integer(0)

out <- vector("list", length(means))
for (i in seq_along(means)) {
  out[[i]] <- rnorm(10, means[[i]])
}</pre>
```



Finally, problems arise when iterating over S3 vectors, as loops typically strip the attributes.

```
xs <- as.Date(c("2020-01-01", "2010-01-01"))
for (x in xs) {
  print(x)
}
#> [1] 18262
#> [1] 14610
```

■ Work around this by using [[.

```
for (i in seq_along(xs)) {
  print(xs[[i]])
}
#> [1] "2020-01-01"
#> [1] "2010-01-01"
```

Related tools



- for loops:
 - Useful when known in advance the set of values to iterate over.
 - Otherwise:
 - while(condition) action: performs action while condition is TRUE.
 - repeat(action): repeats action forever (i.e. until it encounters break).
 - Possible to write any for using while, and any while using repeat, but not the converse.
 - Good practice to use the least-flexible (i.e., simplest) solution to a problem.
- R does not have a do {action} while (condition) syntax found in other languages.
- Generally speaking you shouldn't need to use for loops for data analysis tasks, we'll see better solutions.

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Functions



■ Two important ideas:

- Functions can be broken down into three components: arguments, body, and environment.
- Functions are objects, just as vectors are objects.

■ In the following:

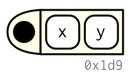
- ► The basics: how to create functions and the three main components of a function.
- ► Function composition: the three forms commonly used in R code
- Lexical scoping: how R finds the value associated with a given name.
- Lazy evaluation: the fact that function arguments are only evaluated when used for the first time.
- ► The special ... argument: how to pass on extra arguments to another function.
- **Exiting a function:** how can a function exit and exit handlers.

Function components



- A function has three parts:
 - The formals(), list of arguments controling how you call the function.
 - The body(), the code inside the function.
 - ▶ The environment(), the data structure that determines how the function finds the values associated with the names.

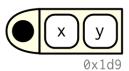
```
f02 <- function(x, y) {
    # A comment
    x + y
}</pre>
```



Function components cont'd



- How those are defined?
 - Explicitly for the formals and body.
 - Implicitly for the environment (where the function was defined).



Function components cont'd



- Functions can possess any number of additional attributes().
- One attribute in base R is srcref, short for source reference.
 - Points to the source code used to create the function.
 - Used for printing because, unlike body(), it contains code comments and other formatting.

```
attr(f02, "srcref")
#> function(x, y) {
#> # A comment
#> x + y
#> }
```

Primitive functions



- One exception to the three components rule.
- Call C code directly.

```
sum
#> function (..., na.rm = FALSE) .Primitive("sum")
`[`
#> .Primitive("[")
```

■ Type is either builtin or special.

```
typeof(sum)
#> [1] "builtin"
typeof(`[`)
#> [1] "special"
```

• formals(), body(), and environment() are all NULL.

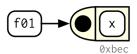
```
formals(sum)
#> NULL
body(sum)
#> NULL
environment(sum)
#> NULL
```

First-class functions



- R functions are objects in their own right!
- This language property often called "first-class functions".
- Unlike in many other languages, no special syntax:
 - Create a function object (with function).
 - ▶ Bind it to a name with <-.</p>

```
f01 <- function(x) {
   sin(1 / x ^ 2)
}</pre>
```





- The binding step is not compulsory.
- A function without a name is called an **anonymous function**.

```
lapply(mtcars, function(x) length(unique(x)))
integrate(function(x) sin(x) ^ 2, 0, pi)
```

Also possible to put functions in a list.

```
funs <- list(
  half = function(x) x / 2,
  double = function(x) x * 2
)

funs$double(10)
#> [1] 20
```

- In R, functions are often called **closures**.
- The name reflects the fact that R functions capture/enclose, their environments.

Invoking a function



■ The standard way:

```
mean(1:10, na.rm = TRUE)
#> [1] 5.5
```

■ The alternative way:

```
args <- list(1:10, na.rm = TRUE)
do.call(mean, args)
#> [1] 5.5
```

Function composition



Imagine you want to compute the population standard deviation using sqrt() and mean().

```
square <- function(x) x^2
deviation <- function(x) x - mean(x)</pre>
```

Either nest the function calls.

```
x <- runif(100)
sqrt(mean(square(deviation(x))))
#> [1] 0.278
```

Or save the intermediate results as variables.

```
out <- deviation(x)
out <- square(out)
out <- mean(out)
out <- sqrt(out)
out
#> [1] 0.278
```

Function composition cont'd



- The third option using the magrittr package:
 - The operator %>%, called **pipe** and pronounced as "and then".

```
library(magrittr)

x %>%
    deviation() %>%
    square() %>%
    mean() %>%
    sqrt()

#> [1] 0.278
```

Advantages:

- Focus on the high-level composition of functions, not the low-level flow of data.
- Focus on what's being done (the verbs), not on what's being modified (the nouns).
- Makes your code more readable by:
 - Structuring sequences of data operations left-to-right.
 - Minimizing the need for local variables and function definitions.
 - Making it easy to add steps anywhere in the sequence.

x % % f(y)

x %% f(y) %% f(z)

x %>% f is equivalent to f(x)

#> [1] 3 5 7 9 11 13 15 17 19 21

#> [1] 6 9 12 15 18 21 24 27 30 33



x %>% f(y) is equivalent to f(x, y)
x %>% f(y) %>% g(z) is equivalent to g(f(x, y), z)

x <- 1:10
y <- x + 1
z <- y + 1
f <- function(x, y) x + y

x %>% sum
#> [1] 55

The argument ("dot") placeholder

 $\mathbf{x} \% \% \mathbf{f}(y, .)$ is equivalent to $\mathbf{f}(y, x)$



- x %>% f(y, z = .) is equivalent to f(y, z = x)
 x <- 1:10
 y <- 2 * x
 f <- function(z, y) y / z</pre>

Function composition cont'd



- Each of the three options has its own strengths and weaknesses:
 - Nesting, f(g(x)):
 - Concise, and well suited for short sequences.
 - Longer sequences harder to read (inside out & right to left).
 - Arguments can get spread out over long distances creating the Dagwood sandwich problem.
 - Intermediate objects, y <- f(x); g(y):</p>
 - Requires you to name intermediate objects.
 - A strength when objects are important, but a weakness when values are truly intermediate.
 - ▶ Piping, x %>% f() %>% g():
 - Allows to read code in straightforward left-to-right fashion.
 - Doesn't require to name intermediate objects.
 - Only for linear sequences of transformations of a single object.
- Most code use a combination of all three styles, but...
- Piping is more common in data analysis code!

Lexical scoping



- **Scoping:** the act of finding the value associated with a name.
- What does the following code return?

```
x <- 10
g01 <- function() {
   x <- 20
   x
}</pre>
```

- R uses **lexical scoping**¹:
 - Looks up the values of names based on how a function is defined, not how it is called.
 - Follows four primary rules:
 - Name masking
 - Functions versus variables
 - A fresh start
 - Dynamic lookup

¹but possible to override the default rules.

Name masking



Names defined inside a function mask names defined outside.

```
x <- 10
y <- 20
g02 <- function() {
    x <- 1
    y <- 2
    c(x, y)
}
g02()
#> [1] 1 2
```

■ If a name isn't defined inside a function, R looks one level up.

```
x <- 2
g03 <- function() {
    y <- 1
    c(x, y)
}
g03()
#> [1] 2 1
y
#> [1] 20
```



- Same applies if a function is defined inside another function:
 - First, R looks inside the current function.
 - Then, where that function was defined (and so on, all the way up to the global environment).
 - Finally, in other loaded packages.
- What does the following code return?

```
x <- 1
g04 <- function() {
    y <- 2
    i <- function() {
        z <- 3
        c(x, y, z)
    }
    i()
}
g04()</pre>
```

Functions versus variables



■ Functions are ordinary objects, the same rules apply to them.

```
g07 <- function(x) x + 1
g08 <- function() {
  g07 <- function(x) x + 100
  g07(10)
}
g08()
#> [1] 110
```

- When a function and a non-function share the same name, the rules get a little more complicated.
 - For function calls, R ignores non-functions when scoping.

```
g09 <- function(x) x + 100
g10 <- function() {
  g09 <- 10
  g09(g09)
}
g10()
#> [1] 110
```

■ But using the same name for different things is best avoided!



- What happens to values between invocations of a function?
- What will happen the first time you run this function?
- What will happen the second time?

```
g11 <- function() {
   if (!exists("a")) {
      a <- 1
   } else {
      a <- a + 1
   }
   a
}</pre>
```

Dynamic lookup



- The output of a function can depend on objects outside its environment, because:
 - Lexical scoping determines where, not when, to look for values.
 - R looks for values when the function is run, not when the function is created.

```
g12 <- function() x + 1
x <- 15
g12()
#> [1] 16
x <- 20
g12()
#> [1] 21
```

- Can be quite annoying.
 - With spelling mistakes, no error when creating a function.
 - ▶ Depending on the global environment, maybe not even an error when running the function.

Lazy evaluation



- In R, function arguments are lazily evaluated:
 - Only evaluated if accessed.
 - What will this code return?

```
h01 <- function(x) {
   10
}
h01(stop("This is an error!"))</pre>
```

- Allows to include expensive computations in function arguments that are only evaluated if needed.
- Powered by **promises**, a data structure with three components:
 - An expression, like x + y, giving rise to the delayed computation.
 - An environment, where the expression should be evaluated.
 - A value.

Promises: the environment



- The environment is where the expression should be evaluated.
 - I.e., where the function is called.
 - What will this code return?

```
y <- 10
h02 <- function(x) {
    y <- 100
    x + 1
}
h02(y)</pre>
```

Also means that when assigning inside a call to a function, the variable is bound outside the function, not inside.

```
h02(y <- 1000)

#> [1] 1001

y

#> [1] 1000
```

Promises: the value



- The value:
 - Computed and cached the first time a promise is accessed, when the expression is evaluated in the specified environment.
 - Ensures that the promise is evaluated at most once.
- What will this code return?

```
double <- function(x) {
  message("Calculating...")
  x * 2
}
h03 <- function(x) {
  c(x, x)
}
h03(double(x))
#> Calculating...
```

■ Can't manipulate promises with R code: any inspection attempt with code will force an immediate evaluation, making the promise disappear.

Default arguments



- Thanks to lazy evaluation:
 - Default values can be defined in terms of other arguments.
 - Or even in terms of variables defined later in the function.
- What will this code return?

```
h04 <- function(x = 1, y = x * 2, z = a + b) {
    a <- 10
    b <- 100

    c(x, y, z)
}
```

- Many use this technique, but not recommended:
 - Makes the code harder to understand.
 - ► To predict *what* will be returned, need to know the exact order in which default arguments are evaluated.

Default arguments cont'd



- The evaluation environment.
 - User supplied arguments: evaluated in the global environment.
 - ▶ Default arguments: evaluated inside the function.
- Seemingly identical calls can yield different results.

```
h05 <- function(x = ls()) {
    a <- 1
    x
}

# ls() evaluated in global environment:
h05(ls())
#> [1] "h05"

# ls() evaluated inside h05:
h05()
#> [1] "a" "x"
```

Missing arguments



Use missing() to determine if an argument's value comes from the user or from a default.

```
h06 \leftarrow function(x = 10) {
  list(missing(x), x)
# default
str(h06())
#> List of 2
#> $ : logi TRUE
#> $ : num 10
# user supplied
str(h06(10))
#> List of 2
#> $ : logi FALSE
#> $ : num 10
```



■ How many arguments are required?

```
args(sample)
#> function (x, size, replace = FALSE, prob = NULL)
#> NULL
```

- A "better" sample():
 - Use an explicit NULL to indicate that size is not required but can be supplied.

```
sample <- function(x, size = NULL, replace = FALSE, prob = NULL) {
  if (is.null(size)) {
    size <- length(x)
  }
  x[sample.int(length(x), size, replace = replace, prob = prob)]
}</pre>
```



- The special argument ... (pronounced dot-dot-dot).
 - Makes a function take any number of additional arguments.
 - In other programming languages:
 - This is often called *varargs* (short for variable arguments).
 - A function that uses it is said to be variadic.
- Can pass those additional arguments on to another function.

```
i01 <- function(y, z) {
   list(y = y, z = z)
}

i02 <- function(x, ...) {
   i01(...)
}

str(i02(x = 1, y = 2, z = 3))

#> List of 2

#> $ y: num 2

#> $ z: num 3
```



If a function takes a function as an argument, you want some way to pass additional arguments to that function.

```
x <- list(c(1, 3, NA), c(4, NA, 6))
str(lapply(x, mean, na.rm = TRUE))
#> List of 2
#> $ : num 2
#> $ : num 5
```

■ If a function is an S3 generic, it needs some way to allow methods to take arbitrary extra arguments.

```
print(factor(letters), max.levels = 4)
#> [1] a b c d e f g h i j k l m n o p q r s t u v w x y z
#> 26 Levels: a b c ... z
```



■ list(...) evaluates the arguments and stores them in a list.

```
i04 <- function(...) {
   list(...)
}
str(i04(a = 1, b = 2))
#> List of 2
#> $ a: num 1
#> $ b: num 2
```

- In general, using ... comes with two downsides:
 - When using it to pass arguments to another function, need to carefully explain to the user where those arguments go.
 - Makes it hard to understand what a function can do.
 - A misspelled argument will not raise an error.
 - Makes it easy for typos to go unnoticed.

```
sum(1, 2, NA, na_rm = TRUE)
#> [1] NA
```

Exiting a function



- Most functions exit in one of two ways:
 - They either return a value, indicating success.
 - Or they throw an error, indicating failure.
- In the next few slides:
 - Return values.
 - Implicit versus explicit.
 - Visible versus invisible.
 - Errors.
 - Exit handlers, allowing to run code when a function exits.

Implicit versus explicit returns



■ Implicit, where the last evaluated expression is the return value.

```
j01 <- function(x) {
   if (x < 10) {
      0
   } else {
      10
   }
}

j01(5)
#> [1] 0
j01(15)
#> [1] 10
```

■ Explicit, by calling return().

```
j02 <- function(x) {
  if (x < 10) {
    return(0)
  } else {
    return(10)
  }
}</pre>
```



Most functions return visibly: calling the function in an interactive context prints the result.

```
j03 <- function() 1
j03()
#> [1] 1
```

Applying invisible() to the last value prevents this.

```
j04 <- function() invisible(1)
j04()</pre>
```

■ Verify that the value exists with print or ().

```
print(j04())
#> [1] 1

(j04())
#> [1] 1
```

Invisible values cont'd



■ The most common function that returns invisibly is <-.

```
a <- 2
(a <- 2)
#> [1] 2
```

■ This is what makes it possible to chain assignments.

```
a <- b <- c <- d <- 2
```

Functions called primarily for a side effect (like <-, print(), or plot()) should return an invisible value (often the value of the first argument).



- If a function cannot complete its assigned task, it should throw an error with stop():
 - Immediately terminates the execution of the function.
 - Indicates that something has gone wrong, and forces the user to deal with the problem.

```
j05 <- function() {
  stop("I'm an error")
  return(10)
}
j05()
#> Error in j05(): I'm an error
```

■ Some languages rely on special return values to indicate problems, but in R you should always throw an error.

Exit handlers



```
j06 <- function(x) {</pre>
  cat("Hello\n")
  on.exit(cat("Goodbye!\n"), add = TRUE)
  if (x) {
    return(10)
  } else {
    stop("Error")
j06 (TRUE)
#> Hello
#> Goodbye!
#> [1] 10
j06 (FALSE)
#> Hello
#> Error in j06(FALSE): Error
#> Goodbye!
```

Exit handlers with on.exit()



- Always set add = TRUE:
 - ▶ If you don't, each call to on.exit() overwrites previous ones.
 - Even when only registering a single handler, it's good practice to set add = TRUE.
- on.exit() is useful because it allows to place clean-up code directly next to the code that requires clean-up.

```
cleanup <- function(dir, code) {
  old_dir <- setwd(dir)
  on.exit(setwd(old_dir), add = TRUE)

  old_opt <- options(stringsAsFactors = FALSE)
  on.exit(options(old_opt), add = TRUE)
}</pre>
```

Exit handlers with on.exit() cont'd



 Coupled with lazy evaluation, a useful pattern for running a block of code in an altered environment.

```
with_dir <- function(dir, code) {
   old <- setwd(dir)
   on.exit(setwd(old), add = TRUE)

force(code)
}

getwd()
#> [1] "/home/tvatter/Dropbox/teaching/stat5206/2019_fall/lectures"
with_dir("-", getwd())
#> [1] "/home/tvatter"
getwd()
#> [1] "/home/tvatter/Dropbox/teaching/stat5206/2019_fall/lectures"
```

- force() isn't strictly necessary here as simply referring to code will force its evaluation.
- But makes it clear that we are deliberately forcing the execution.