Week 5 Module 5: Conditional Branching CSCI E-5a: Introduction to R

Let's clear the global environment:

```
rm( list = ls() )
```

Let's load in the objects for this module:

```
load( file = "Module 5 R Objects.Rdata" )
```

Module Overview and Learning Objectives

Hello! And welcome to Module 5: Conditional Branching.

In this module, we'll learn about conditional branching.

- In section 1, we'll learn the basics of conditional branching.
- In section 2, we'll find that the if() function does not vectorize.
- In section 3, we'll see another function that can sometimes operate as a vectorized if statement.

When you've completed this module, you'll be able to:

- Explain how conditional branching works.
- Explain how the if() function does not vectorize.
- Use the ifelse() function as a vectorized version of if().

There are two new built-in R functions in this module:

- if()
- ifelse()

All right! Let's get started by defining the concept of conditional branching.

Section 1: Conditional Branching

Main Idea: We can use conditional branching to control program flow

In this section, we'll learn the basics of conditional branching.

We've seen how to construct loops, in which we control the flow of program execution so that it repeats a code block.

Now we'll consider the other important flow-of-control operation, called *conditional branching*.

In this operation, we perform some sort of test (that's the "condition"), and then execute certain code depending on whether or not the test was TRUE or FALSE (that's the "branching").

Let's start with the most basic form of conditional branching:

- First, we use the keyword if to indicate that we are performing conditional branching, followed by an open parenthesis.
- Next, we perform a test, which is just an expression that evaluates to TRUE or FALSE, and we follow this with a closing parenthesis.
- Finally, we have a code block, which is delimited using curly braces.

If the expression in the test evaluates to TRUE, then we execute the code in the code block, but if the code evaluates to FALSE, then we do not execute the code in the code block.

For instance, in this example, the test is TRUE, so R executes the code in the following code block:

```
x <- 6
if( x == 6 ) {
    cat( "We got into the code block!\n" )
}</pre>
```

We got into the code block!

On the other hand, if the test is FALSE, then the following code block is not executed:

```
x <- 5
if( x == 6 ) {
    cat( "We got into the code block!\n" )
}</pre>
```

We can extend this process by using the keyword else, which is followed by a second code block.

In this approach, if the conditional test evaluates to TRUE, then the first code block is executed, but if it evaluates to FALSE then the second code block following the keyword else is executed.

The keyword else has to occur on the same line as the closing curly brace for the initial code block:

```
animal <- "dog"

if( animal == "hedgehog" ) {
    cat( "The animal is a hedgehog\n" )
} else {
    cat( "The animal is not a hedgehog\n" )
}</pre>
```

The animal is not a hedgehog

We can chain together if and else statements to allow for multi-way branching:

```
animal <- "bunny rabbit"

if( animal == "hedgehog" ) {
    cat( "The animal is a hedgehog\n" )
} else if( animal == "armadillo" ) {
    cat( "The animal is an armadillo\n" )
} else if( animal == "bunny rabbit" ) {
    cat( "The animal is a bunny rabbit\n" )
} else {
    cat( "Some other thing\n" )
}</pre>
```

The animal is a bunny rabbit

Conditional branching with the if() function will be very important in Lecture 8: Functions, when we learn how to write our own user-defined functions.

So that's how to use conditional branching to control the flow of execution in a program.

Now let's see what happens when we try to vectorize the if() function.

Section 2: if doesn't vectorize

Main Idea: The if() function doesn't vectorize

In this section, we'll find that the if() function does not vectorize.

The if() function doesn't vectorize, at least not in the way that we often want it to.

Let's create a test vector:

```
test.vector <- c( 4, 2, 9, 3, 1, 6, 12)
```

We want to scan through the test vector and print out a short message each time an element is strictly greater than 3.

Let's see if we can implement this using a vectorized approach:

```
if( test.vector > 3 ) {
    cat( "Hi Mom!\n" )
}
```

```
## Warning in if (test.vector > 3) {: the condition has length > 1 and only the
## first element will be used
## Hi Mom!
```

Notice that although test.vector has 7 elements, R printed out only one statement, so this statement did not vectorize.

Why did this happen?

The answer is that if() doesn't vectorize; instead, it just takes the first element of the vector and applies the if() statement to that.

Actually, the warning message that R prints out is unusually helpful and descriptive.

This is a situation where a for() loop works great:

```
for( element in test.vector ) {
    if( element > 3 ) {
        cat( "Element:", element, "\n" )
    }
}
```

```
## Element: 4
## Element: 9
## Element: 6
## Element: 12
```

The important point is that you have to be careful about vectorization, and to check and make sure that R actually does what you thought it would do.

So the if() function doesn't vectorize.

Now let's explore another function that performs conditional branching and does vectorize in the way that we might want it to.

Section 3: The ifelse() function

Main Idea: We can use the ifelse() function to perform vectorized conditional branching

In this section, we'll see another function that can sometimes operate as a vectorized if statement.

The ifelse() function can sometimes operate as a vectorized if() statement (sort of).

The ifelse() function takes three input arguments:

• A logical value, called test.

- A value to be returned if the logical value is TRUE, called yes.
- A value to be returned if the logical value is FALSE, called no.

The important point about ifelse() is that it can be vectorized.

Let's see how this works.

First, we'll create a logical test vector:

```
logical.test.vector <-
c( TRUE, TRUE, FALSE, TRUE, FALSE )</pre>
```

Let's create a new vector based on logical.test.vector:

- If a particular element of logical.test.vector is TRUE, then the corresponding element of the new vector is "More Kale!".
- If a particular element of logical.test.vector is FALSE, then the corresponding element of the new vector is "More Tofu!".

To create this vector, we can use the ifelse() function:

- The test input argument is just the logical.test.vector.
- The yes input argument is the character string value "More Kale!"
- The no input argument is the character string value "More Tofu!"

```
ifelse(
  test = logical.test.vector,
  yes = "More Kale!",
  no = "More Tofu!"
)
```

[1] "More Kale!" "More Kale!" "More Tofu!" "More Kale!" "More Tofu!"

Make sure you understand how this function works, and how this output vector was constructed.

Just for reference, we could also construct this vector using a combination of a for() loop and an if() statement:

```
number.of.elements <-
  length( logical.test.vector )

output.vector <-
  numeric( number.of.elements )

for( i in 1:number.of.elements ) {

  if( logical.test.vector[ i ] == TRUE ) {
    output.vector[ i ] <- "More Kale!"
  } else {
    output.vector[ i ] <- "More Tofu!"
  }
}

output.vector</pre>
```

[1] "More Kale!" "More Kale!" "More Tofu!" "More Kale!" "More Tofu!"

Notice that in the end both approaches produce the same output.

However, the iterative approach is more complicated to implement, because we have to specify every step of the computation.

By contrast, the ifelse() function handles a lot of this work, and so the code is much simpler when we use this approach.

I must emphasize – the ifelse() function is perhaps a little more abstract, and if you don't feel comfortable with this and prefer the iterative approach, that's perfectly fine.

As always, the most important thing is to write clear and correct code that works and does what it's supposed to do.

You might wonder if the ifelse() function is particularly useful, and I admit that it took me a long to appreciate its merits.

The point is that we can construct the logical test vector using a vectorized operation.

This enables us to effectively implement a conditional test for each element of a vector and then branch on the result, so that this can function as a sort of vectorized **if()** statement.

Let's see how this works.

Let's recall our test.vector:

test.vector

[1] 4 2 9 3 1 6 12

Now let's consider an example of how to use the ifelse() function (I admit this is a little artificial).

We want to create a new vector:

- If an element of test.vector is greater than or equal to 3, then the corresponding element of the new vector has the same value.
- If an element of test.vector is strictly less than 3, then the corresponding element of the new vector is 0.

In other words, to create the new vector we make a copy of test.vector where any elements strictly less than 3 are converted to 0.

To do this using ifelse(), we first perform a vectorized comparison of test.vector with the numeric value 3:

```
logical.test.vector <-
  test.vector >= 3
logical.test.vector
```

[1] TRUE FALSE TRUE TRUE FALSE TRUE TRUE

Now we can use ifelse() with this logical test vector:

```
ifelse(
   test = logical.test.vector,
   yes = test.vector,
   no = 0
)
```

```
## [1] 4 0 9 3 0 6 12
```

Notice here that we're vectorizing over both logical.test.vector and test.vector.

By the way, we didn't really need to store the result of the comparison operation in the dedicated variable logical.test.vector, because the only time we use it is in the ifelse() statement.

So we could have just done this:

```
ifelse(
   test = test.vector >= 3,
   yes = test.vector,
   no = 0
)
```

```
## [1] 4 0 9 3 0 6 12
```

So that's how the ifelse() function allows us to perform a vectorized conditional branching operation.

Now let's review what we've learned in this module.

Exercise 1: Vectorized conditional branching

The R object exercise.1.data is a numeric vector.

Create a new character string vector:

- If an element of exercise.1.data is less than 0, then the corresponding element in the new vector should be the character string "Cold".
- If an element of exercise.1.data is greater than or equal to 0, then the corresponding element in the new vector should be the character string "Hot".

Directly display the vector exercise.1.data, and try to determine what the new vector will be, based on this data.

Then write R code to create the new vector.

Ideally, you should do this using two approaches:

- Using vectorized operations with ifelse().
- Using a for() loop and an if() statement.

Which method do you prefer?

Solution

Module Review

In this module, we learned about conditional branching.

- In section 1, we learned the basics of conditional branching.
- In section 2, we found that the if() function does not vectorize.
- In section 3, we saw another function that can sometimes operate as a vectorized if statement.

Now that you've completed this module, you should be able to:

- Explain how conditional branching works.
- Explain how the if() function does not vectorize.
- Use the ifelse() function as a vectorized version of if().

There were two new built-in R functions in this module:

- if()
- ifelse()

All right! That's it for Module 5: Conditional Branching.

Now let's move on to Module 6: Histograms and Curves.

Solution to the Exercise

Exercise 1: Vectorized conditional branching

The R object exercise.1.data is a numeric vector.

Create a new character string vector:

- If an element of exercise.1.data is less than 0, then the corresponding element in the new vector should be the character string "Cold".
- If an element of exercise.1.data is greater than or equal to 0, then the corresponding element in the new vector should be the character string "Hot".

Directly display the vector exercise.1.data, and try to determine what the new vector will be, based on this data.

Then write R code to create the new vector.

Ideally, you should do this using two approaches:

- Using vectorized operations with ifelse().
- Using a for() loop and an if() statement.

Which method do you prefer?

Solution

First, let's directly display the vector:

```
## [1] 9 5 -8 2 7 -3 4 -1
```

In exercise.1.data, positive numbers occur in positions 1, 2, 4, 5, and 7, while negative numbers occur in positions 3, 6, and 8.

Thus, the new vector should have the character string "Hot" in positions 1, 2, 4, 5, and 7, and the character string "Cold" in positions 3, 6, and 8.

We can do this with ifelse():

```
new.vector <-
ifelse(
    exercise.1.data >= 0,
    "Hot",
    "Cold"
)
new.vector
```

```
## [1] "Hot" "Hot" "Cold" "Hot" "Cold" "Hot" "Cold"
```

This is exactly what we had predicted.

To do this using a for() and an if() statement is a little more work:

```
number.of.elements <-
   length( exercise.1.data )

new.vector <-
   character( number.of.elements )

for( i in 1:number.of.elements ) {

   if( exercise.1.data[ i ] >= 0 ) {
      new.vector[ i ] <- "Hot"
   } else {
      new.vector[ i ] <- "Cold"
   }
}

new.vector</pre>
```

```
## [1] "Hot" "Hot" "Cold" "Hot" "Cold" "Hot" "Cold"
```

This is exactly the same result as before, but it was a little bit more work to get there using a for() loop and an if() statement, compared to doing this with an ifelse() statement.