# 03 Programming basics

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The most common conditional expression is the if-else statement.

Here is a very simple example showing the general structure of an if-else statement. The basic idea is to print the reciprocal of a unless a is 0:

```
a <- 0

if(a!=0){
   print(1/a)
} else{
   print("No reciprocal for 0.")
}
#> [1] "No reciprocal for 0."
```

Let's look at one more example using the US murders data frame:

```
library(dslabs)
data(murders)
murder_rate <- murders$total / murders$population*100000</pre>
```

Here is a very simple example that tells us which states, if any, have a murder rate lower than 0.5 per 100,000. The if statement protects us from the case in which no state satisfies the condition.

```
ind <- which.min(murder_rate)

if(murder_rate[ind] < 0.5){
   print(murders$state[ind])
} else{
   print("No state has murder rate that low")
}
#> [1] "Vermont"
```

If we try it again with a rate of 0.25, we get a different answer:

```
if(murder_rate[ind] < 0.25){
  print(murders$state[ind])
} else{
  print("No state has a murder rate that low.")
}
#> [1] "No state has a murder rate that low."
```

A related function that is very useful is ifelse.

```
a <- 0
ifelse(a > 0, 1/a, NA)
#> [1] NA
```

The function is particularly useful because it works on vectors.

This table helps us see what happened:

а	is_a_positive	answer1	answer2	result
0	FALSE	Inf	NA	NA
1	TRUE	1.00	NA	1.0
2	TRUE	0.50	NA	0.5
-4	FALSE	-0.25	NA	NA
5	TRUE	0.20	NA	0.2

Here is an example of how this function can be readily used to replace all the missing values in a vector with zeros:

```
data(na_example)
no_nas <- ifelse(is.na(na_example), 0, na_example)
sum(is.na(no_nas))
#> [1] 0
```

Two other useful functions are any and all. The any function takes a vector of logicals and returns TRUE if any of the entries is TRUE. The all function takes a vector of logicals and returns TRUE if all of the entries are TRUE. Here is an example:

```
z <- c(TRUE, TRUE, FALSE)
any(z)
#> [1] TRUE
all(z)
#> [1] FALSE
```

# **Defining functions**

We can compute the average of a vector x using the sum and length functions: sum(x)/length(x).

A simple version of a function that computes the average can be defined like this:

```
avg <- function(x){
    s <- sum(x)
    n <- length(x)
    s/n
}</pre>
```

Now avg is a function that computes the mean:

```
x <- 1:100
identical(mean(x), avg(x))
#> [1] TRUE
```

# **Defining functions**

The general form of a function definition looks like this:

```
my_function <- function(VARIABLE_NAME){
  perform operations on VARIABLE_NAME and calculate VALUE
  VALUE
}</pre>
```

# **Defining functions**

For example, we can define a function that computes either the arithmetic or geometric average depending on a user defined variable like this:

```
avg <- function(x, arithmetic = TRUE){
  n <- length(x)
  ifelse(arithmetic, sum(x)/n, prod(x)^(1/n))
}</pre>
```

We will learn more about how to create functions through experience as we face more complex tasks.

# **Namespaces**

Once you start loading several add-on packages for some of your analysis, it is likely that two packages use the same name for two different functions.

In fact, you have already encountered this because both **dplyr** and the R-base **stats** package define a **filter** function. We know this because when we first load **dplyr** we see the following message:

```
The following objects are masked from 'package:stats':

filter, lag

The following objects are masked from 'package:base':

intersect, setdiff, setequal, union
```

## **Namespaces**

These functions live in different *namespaces*. R will follow a certain order when searching for a function in these *namespaces*.

#### search()

The first entry in this list is the global environment which includes all the objects you define.

### **Namespaces**

So what if we want to use the **stats** filter instead of the **dplyr** filter? You can force the use of a specific namespace by using double colons (::) like this:

```
stats::filter
```

If we want to be absolutely sure that we use the  ${f dplyr}$  filter, we can use

```
dplyr::filter
```

Also note that if we want to use a function in a package without loading the entire package, we can use the double colon as well.

The formula for the sum of the series  $1+2+\cdots+n$  is n(n+1)/2. What if we weren't sure that was the right function? How could we check?

Using what we learned about functions we can create one that computes the  $S_n$ :

```
compute_s_n <- function(n){
  x <- 1:n
  sum(x)
}</pre>
```

How can we compute  $S_n$  for various values of n say n = 1, ..., 25? Do we write 25 lines of code calling compute\_s\_n? No, that is what for-loops are for in programming.

Perhaps the simplest example of a for-loop is this useless piece of code:

```
for(i in 1:5){
   print(i)
}
#> [1] 1
#> [1] 2
#> [1] 3
#> [1] 4
#> [1] 5
```

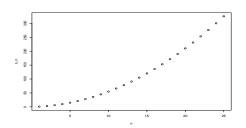
Here is the for-loop we would write for our  $S_n$  example:

```
m <- 25
s_n <- vector(length = m) # create an empty vector
for(n in 1:m){
    s_n[n] <- compute_s_n(n)
}</pre>
```

In each iteration n = 1, n = 2, etc..., we compute  $S_n$  and store it in the nth entry of  $s_n$ .

Now we can create a plot to search for a pattern:

```
n <- 1:m
plot(n, s_n)</pre>
```



If you noticed that it appears to be a quadratic, you are on the right track because the formula is n(n+1)/2.

A *vectorized* function is a function that will apply the same operation on each of the vectors.

```
x <- 1:10
sqrt(x)
#> [1] 1.00 1.41 1.73 2.00 2.24 2.45 2.65 2.83 3.00 3.16
y <- 1:10
x*y
#> [1] 1 4 9 16 25 36 49 64 81 100
```

To make this calculation, there is no need for for-loops.

For instance, the function we just wrote, <code>compute\_s\_n</code>, does not work element-wise since it is expecting a scalar. This piece of code does not run the function on each entry of n:

```
n <- 1:25
compute_s_n(n)</pre>
```

Functionals are functions that help us apply the same function to each entry in a vector, matrix, data frame, or list.

Here we cover the functional that operates on numeric, logical, and character vectors: sapply.

```
x <- 1:10

sapply(x, sqrt)

#> [1] 1.00 1.41 1.73 2.00 2.24 2.45 2.65 2.83 3.00 3.16
```

The function sapply permits us to perform element-wise operations on any function. Here is how it works:

```
n <- 1:25
s_n <- sapply(n, compute_s_n)</pre>
```

Other functionals are apply, lapply, tapply, mapply, vapply, and replicate. We mostly use sapply, apply, and replicate, but we recommend familiarizing yourselves with the others as they can be very useful.

1. What will this conditional expression return?

```
x <- c(1,2,-3,4)

if(all(x>0)){
   print("All Postives")
} else{
   print("Not all positives")
}
```

- 2. Which of the following expressions is always FALSE when at least one entry of a logical vector  $\mathbf{x}$  is TRUE?
  - all(x)
  - any(x)
- any(!x)
- all(!x)

- 3. The function nchar tells you how many characters long a character vector is. Write a line of code that assigns to the object new\_names the state abbreviation when the state name is longer than 8 characters.
- 4. Create a function  $sum_n$  that for any given value, say n, computes the sum of the integers from 1 to n (inclusive). Use the function to determine the sum of integers from 1 to 5,000.
- 5. Create a function  $altman_plot$  that takes two arguments, x and y, and plots the difference against the sum.

6. After running the code below, what is the value of x?

```
x <- 3
my_func <- function(y){
   x <- 5
   y+5
}</pre>
```

- 7. Write a function compute\_s\_n that for any given n computes the sum  $S_n = 1^2 + 2^2 + \cdots + n^2$ . Report the value of the sum when n = 10.
- 8. Define an empty numerical vector  $s_n$  of size 25 using  $s_n < -$  vector("numeric", 25) and store in the results of  $S_1, S_2, \ldots, S_{25}$  using a for-loop.
- 9. Repeat exercise 8, but this time use sapply.

- 10. Repeat exercise 8, but this time use map\_dbl.
- 11. Plot  $S_n$  versus n. Use points defined by  $n = 1, \dots, 25$ .
- 12. Confirm that the formula for this sum is  $S_n = n(n+1)(2n+1)/6$ .