

# Introductory course on the R software

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<https://biostatisticien.eu/springerR/courseRw5.pdf>

# Goals of today lecture

Describing the instructions for

- control structures ;
- creating basic functions.

# Instructions `if` and `else`

```
> if (TRUE) 1+1
[1] 2
> x <- 2
> y <- -3
> if (x <= y) {
+   z <- y-x
+   print("x smaller than y")
+ } else {
+   z <- x-y
+   print("x larger than y")
+   z
+ }
[1] "x larger than y"
[1] 5
```

# Instruction for

```
> for (i in 1:3) print(i)
[1] 1
[1] 2
[1] 3
> x <- c(1,3,7,2)
> for (var in x) print(2*var)
[1] 2
[1] 6
[1] 14
[1] 4
```

# Instruction while

```
> x <- 2  
> y <- 1  
> while(x+y<7) x <- x+y  
> x  
[1] 6
```

The formula  $BMI = \frac{Weight}{Height^2}$  is easily programmed in R as follows :

```
> BMI <- function(weight,height) {
+   bmi <- weight/height^2
+   names(bmi) <- "BMI"
+   return(bmi)
+ }
```

We can now execute the function BMI() we just created :

```
> BMI(70,1.82)
      BMI
21.13271
> BMI(1.82,70) # Note that it is not possible to swap the
               # arguments of a function,
      BMI
0.0003714286
> BMI(height=1.82,weight=70) # unless they are preceded by their
                             # names.
      BMI
21.13271
```

This function only outputs a single value.

The code below outputs a list of several variables.

```
> BMI <- function(weight,height) {  
+   bmi <- weight/height^2  
+   res <- list(weight,height,bmi)  
+   return(res)  
+ }
```

BMI() returns a list of unnamed elements :

```
> BMI(70,1.82)  
[[1]]  
[1] 70  
[[2]]  
[1] 1.82  
[[3]]  
[1] 21.13271
```

To name the elements of the list, you can use the following code

```
> BMI <- function(weight,height) {  
+   bmi <- weight/height^2  
+   res <- list(Weight=weight, Height=height, BMI=bmi)  
+   return(res)  
+ }
```

which gives the following result :

```
> BMI(70,1.82)  
$Weight  
[1] 70  
$Height  
[1] 1.82  
$BMI  
[1] 21.13271
```



# Your turn to work !

Do the Worksheet F on page 140.

<http://biostatisticien.eu/springeR/Rbook-chap5.pdf>