# CODING LOOPS WITH "for" - LECTURE

DSC 205 - Advanced introduction to data science

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## README



Figure 1: Photo by La-Rel Easter on Unsplash

Download the raw practice file from GitHub and save it as 5\_loop\_for\_practice.org. To test your Emacs mettle, open it on the CMD line with the command emacs -nw (no graphics - not needed for this exercise).

### Coding loops overview

- Loops repeat a specified section of code often while incrementing an index or counter variable.
- R knows three ways of looping: for and while as in other C-type languages, and apply to run a function over different data structures.
- A for loop repeates code while going through a vector until a condition is no longer met:

```
for (loopindex in loopvector) {
  do any code in here
}
```

• A while loop repeats code until a condition evaluates as FALSE:

```
while (loopcondition) {
  do any code in here
}
```

• The apply family of functions allows implicit looping over subsets of vectors, matrices or arrays to apply a function to all elements:

```
apply( X = data
MARGIN = subset,
FUN = function)
```

## for loops

• A for loop repeats code while going through a vector until a condition is no longer met:

```
for (loopindex in loopvector) {
  do any code in here
}
```

- loopindex represents an element in loopvector
- Simple example:

```
for (myitem in 5:7) {
   cat("--BRACED AREA BEGINS--\n")
   cat("the current item is", myitem, "\n")
   cat("--BRACED AREA ENDS--\n")
}

--BRACED AREA BEGINS--
the current item is 5
--BRACED AREA ENDS--
--BRACED AREA BEGINS--
the current item is 6
--BRACED AREA ENDS--
--BRACED AREA BEGINS--
the current item is 7
--BRACED AREA ENDS--
```

• You can manipulate objects outside the loop, i.e. the braced area is not a local environment:

```
counter <- 0
for (myitem in 5:7) {
  counter <- counter + 1
  cat("The item in run",counter,"is",myitem,"\n")
}
The item in run 1 is 5
The item in run 2 is 6
The item in run 3 is 7</pre>
```

## Looping via index or value

- There is a difference using a *loopindex* inside a for statement vs. representing *indices* of a vector.
- **Loop index:** The loop below uses the *loopindex* i to directly represent the elements in myvec:

```
myvec <- c(0.4, 1.1, 0.34, 0.55)
for (i in myvec) {
   print(2 * i)
}

[1] 0.8
[1] 2.2
[1] 0.68
[1] 1.1</pre>
```

• Vector index: The loop below uses i to represent integer values in

the sequence 1:length(myvec), which form all vector index positions of myvec. The same indices are then used to extract elements from myvec.

```
for (i in 1:length(myvec)) {
   print(2 * myvec[i])
}

[1] 0.8
[1] 2.2
[1] 0.68
[1] 1.1
```

• Vector indices take a longer form but offer more flexibility for more complicated for loops.

## Extended example: stepping through a list

- You want to write code that will inspect any list object and gather information about any matrix objects stored as list members.
- Sample data:

```
foo <- list(
    aa=c(3,4,1),
    bb=matrix(1:4,2,2),
    cc=matrix(c(T,T,F,T,F,F),3,2),
    dd="string here",
    ee=matrix(c("red","green","blue","yellow")))
foo</pre>
```

```
$aa
[1] 3 4 1
$bb
     [,1] [,2]
[1,]
              3
        1
[2,]
        2
              4
$cc
      [,1]
             [,2]
[1,]
      TRUE TRUE
[2,]
      TRUE FALSE
[3,] FALSE FALSE
$dd
[1] "string here"
$ee
     [,1]
[1,] "red"
[2,] "green"
[3,] "blue"
[4,] "yellow"
```

#### • Problem:

- 1. Go through every member of the list
- 2. Check whether the member is matrix
- 3. If it is a matrix, retrieve number of rows and columns and the data type of the matrix.
- Solution: create vectors to store list member information:
  - 1. name of the list member name
  - 2. is.mat ("Yes" or "No") to indicate if it is a matrix
  - 3. nc and nr for numbers of rows and columns for each matrix
  - 4. data.type to store the data type of each matrix

name <- names(foo); name</pre>

```
is.mat <- rep(NA,length(foo)); is.mat
nr <- is.mat
nc <- is.mat
data.type <- is.mat

[1] "aa" "bb" "cc" "dd" "ee"
[1] NA NA NA NA NA</pre>
```

• All sought variables are initialized with "missing" values NA and updated in the for loop. The results are put in a data frame bar.

```
for (i in 1:length(foo)) {
  member <- foo[[i]] # pick list element</pre>
  if (is.matrix(member)) {
    is.mat[i] <- "Yes"</pre>
                             # update matrix indicator
    nr[i] <- nrow(member) # update row counter</pre>
    nc[i] <- ncol(member) # update column counter</pre>
    data.type[i] <- class(as.vector(member)) # conversion!</pre>
  } else {
    is.mat[i] <- "No"
bar <- data.frame(name,
                   is.mat,
                   nr,
                   nc,
                   data.type)
bar
  name is.mat nr nc data.type
           No NA NA
                           <NA>
1
    aa
2
    bb
          Yes 2 2
                       integer
3
          Yes 3 2
                       logical
   СС
4
    dd
           No NA NA
                           <NA>
5
          Yes 4 1 character
```

• Compare with the original list foo where the structure output indirectly suggests matrix objects through the indexing:

```
str(foo)
```

```
List of 5
$ aa: num [1:3] 3 4 1
$ bb: int [1:2, 1:2] 1 2 3 4
$ cc: logi [1:3, 1:2] TRUE TRUE FALSE TRUE FALSE FALSE
$ dd: chr "string here"
$ ee: chr [1:4, 1] "red" "green" "blue" "yellow"
```

### Nesting for loops

- for loops can be nested just like if statements.
- The inner loop is executed in full before the outer loop *loopindex* is incremented. Then the inner loop is executed all over again.
- Example: write code that loops over rows and columns of a matrix and update the matrix elements as the product of outer and inner loopindex.
- Solution: first, create loopindices and a base matrix:
  - 1. Create a loopvec1 as loopindex vector 5 6 7
  - 2. Create a loopvec2 as loopindex vector 9 8 7 6
  - 3. Create a matrix foo of missing values whose row and column numbers correspond to loopvec1 and loopvec2, respectively

```
loopvec1 <- 5:7; loopvec1</pre>
loopvec2 <- 9:6; loopvec2</pre>
foo <- matrix(NA,</pre>
               length(loopvec1),
               length(loopvec2))
foo
baz <- foo # make copy of foo for later
[1] 5 6 7
[1] 9 8 7 6
     [,1] [,2] [,3] [,4]
[1,]
       NA
           NA
                  NA
                        NA
[2,]
       NA
            NA
                  NA
                        NA
[3,]
     NA
           NΑ
                  NA
                        NΑ
```

• The outer for loop should run over as many elements as loopvec1 has, and the inner for loop should run over as many elements as loopvec2 has. foo is then updated accordingly:

```
for (i in 1:length(loopvec1)) {
  for (j in 1:length(loopvec2)) {
    foo[i,j] <- loopvec1[i] * loopvec2[j]</pre>
  }
}
foo
     [,1] [,2] [,3] [,4]
[1,]
       45
                   35
                        30
             40
[2,]
       54
             48
                   42
                        36
[3,]
       63
             56
                   49
                        42
```

- With this code, is the matrix traversed by row or by column?<sup>1</sup>
- Inner loopvectors can be defined to match the current value of the loopindex of the outer loop:

```
for (i in 1:length(loopvec1)) {
  for (j in 1:i) {
    baz[i,j] <- loopvec1[i] * loopvec2[j]</pre>
  }
}
baz
     [,1] [,2] [,3] [,4]
[1,]
       45
             NA
                  NA
                        NA
[2,]
             48
        54
                  NA
                        NA
[3,]
        63
             56
                   49
                        NA
```

• Note that inner loop indices are decided based on the outer loop index: for example, when i=1, the inner loopvector is 1:1 so it is executed only once before moving on to the next row.

<sup>&</sup>lt;sup>1</sup>By column - in the example, the sequence of matrix elements filled is: foo[1,1], foo[1,2], foo[1,3], foo[2,1] etc.

• The code will fail if length(loopvec1)>length(loopvec2): why?<sup>2</sup>

```
loopvec1 <- 1:4
loopvec2 <- 9:7
qux <- matrix(NA,length(loopvec1),length(loopvec2)); foo</pre>
for (i in 1:length(loopvec1)) {
  for (j in 1:i) {
    qux[i,j] <- loopvec1[i] * loopvec2[j]</pre>
  }
}
     [,1] [,2] [,3] [,4]
[1,]
                       30
       45
            40
                  35
[2,]
                       36
       54
            48
                  42
[3,]
       63
             56
                  49
                       42
Error in '[<-'('*tmp*', i, j, value = loopvec1[i] * loopvec2[j]) :
  subscript out of bounds
```

<sup>&</sup>lt;sup>2</sup>Because the inner loopvector 1:i will exceed the number of elements of loopvec2 - "subscript out of bounds".

# **TODO** Exercises



Download the  $\mathbf{raw}$  exercise file from GitHub and save it as  $5\_loop\_for\_exercise.org$ . When done, upload the file to Canvas.

# Glossary

TERM	MEANING
for (i in X)	loop over index i
X	loop index or vector index
{}	loop body
Nesting loop	inner loop inside outer loop
Loop by value	condition computed directly, e.g. 1:10
Loop by index	condition from vector, e.g. 1:length(vec)

## References

• Davies, T.D. (2016). The Book of R. NoStarch Press.