

# Introduction to the R Language

**Loop Functions** 

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#### **Looping on the Command Line**

Writing for, while loops is useful when programming but not particularly easy when working interactively on the command line. There are some functions which implement looping to make life easier.

- lapply: Loop over a list and evaluate a function on each element
- sapply: Same as lapply but try to simplify the result
- apply: Apply a function over the margins of an array
- tapply: Apply a function over subsets of a vector
- · mapply: Multivariate version of lapply

An auxiliary function split is also useful, particularly in conjunction with lapply.

lapply takes three arguments: (1) a list x; (2) a function (or the name of a function) FUN; (3) other arguments via its ... argument. If x is not a list, it will be coerced to a list using as.list.

```
lapply
```

The actual looping is done internally in C code.

lapply always returns a list, regardless of the class of the input.

```
x \leftarrow list(a = 1:5, b = rnorm(10)) lapply(x, mean)
```

```
## $a
## [1] 3
##
## $b
## [1] 0.4671
```

```
x <- list(a = 1:4, b = rnorm(10), c = rnorm(20, 1), d = rnorm(100, 5)) lapply(x, mean)
```

```
## $a
## [1] 2.5
##
## $b
## [1] 0.5261
##
## $c
## [1] 1.421
##
## $d
## [1] 4.927
```

```
> x <- 1:4

> lapply(x, runif)

[[1]]

[1] 0.2675082

[[2]]

[1] 0.2186453 0.5167968

[[3]]

[1] 0.2689506 0.1811683 0.5185761

[[4]]

[1] 0.5627829 0.1291569 0.2563676 0.7179353
```

```
> x <- 1:4
> lapply(x, runif, min = 0, max = 10)
[[1]]
[1] 3.302142

[[2]]
[1] 6.848960 7.195282

[[3]]
[1] 3.5031416 0.8465707 9.7421014

[[4]]
[1] 1.195114 3.594027 2.930794 2.766946
```

lapply and friends make heavy use of *anonymous* functions.

An anonymous function for extracting the first column of each matrix.

```
> lapply(x, function(elt) elt[,1])
$a
[1] 1 2
$b
[1] 1 2 3
```

sapply will try to simplify the result of lapply if possible.

- · If the result is a list where every element is length 1, then a vector is returned
- If the result is a list where every element is a vector of the same length (> 1), a matrix is returned.
- If it can't figure things out, a list is returned

```
> x <- list(a = 1:4, b = rnorm(10), c = rnorm(20, 1), d = rnorm(100, 5))
> lapply(x, mean)
$a
[1] 2.5
$b
[1] 0.06082667
$c
[1] 1.467083
$d
[1] 5.074749
```



# Introduction to the R Language

Loop Functions - apply

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apply is used to a evaluate a function (often an anonymous one) over the margins of an array.

- · It is most often used to apply a function to the rows or columns of a matrix
- · It can be used with general arrays, e.g. taking the average of an array of matrices
- It is not really faster than writing a loop, but it works in one line!

```
> str(apply)
function (X, MARGIN, FUN, ...)
```

- · x is an array
- MARGIN is an integer vector indicating which margins should be "retained".
- Fun is a function to be applied
- · ... is for other arguments to be passed to FUN

```
> x <- matrix(rnorm(200), 20, 10)</pre>
> apply(x, 2, mean)
[1] 0.04868268 0.35743615 -0.09104379
[4] -0.05381370 -0.16552070 -0.18192493
[7] 0.10285727 0.36519270 0.14898850
[10] 0.26767260
> apply(x, 1, sum)
[1] -1.94843314 2.60601195 1.51772391
[4] -2.80386816 3.73728682 -1.69371360
 [7] 0.02359932 3.91874808 -2.39902859
[10] 0.48685925 -1.77576824 -3.34016277
[13] 4.04101009 0.46515429 1.83687755
[16] 4.36744690 2.21993789 2.60983764
[19] -1.48607630 3.58709251
```

### col/row sums and means

For sums and means of matrix dimensions, we have some shortcuts.

```
rowSums = apply(x, 1, sum)
rowMeans = apply(x, 1, mean)
colSums = apply(x, 2, sum)
colMeans = apply(x, 2, mean)
```

The shortcut functions are *much* faster, but you won't notice unless you're using a large matrix.

### Other Ways to Apply

Quantiles of the rows of a matrix.

```
> x < - matrix(rnorm(200), 20, 10)
> apply(x, 1, quantile, probs = c(0.25, 0.75))
          \lceil , 1 \rceil
               [,2] [,3] [,4]
25% -0.3304284 -0.99812467 -0.9186279 -0.49711686
75% 0.9258157 0.07065724 0.3050407 -0.06585436
           [,5]
                 \begin{bmatrix} , 6 \end{bmatrix} \begin{bmatrix} , 7 \end{bmatrix}
25% -0.05999553 -0.6588380 -0.653250 0.01749997
75% 0.52928743 0.3727449 1.255089 0.72318419
          [,9] [,10] [,11] [,12]
25% -1.2467955 -0.8378429 -1.0488430 -0.7054902
75% 0.3352377 0.7297176 0.3113434 0.4581150
         [,13]
                    \lceil ,14 \rceil \qquad \lceil ,15 \rceil
                                      [,16]
25% -0.1895108 -0.5729407 -0.5968578 -0.9517069
75% 0.5326299 0.5064267 0.4933852 0.8868922
         [,17] [,18] [,19] [,20]
```

Average matrix in an array



# Introduction to the R Language

Loop Functions - mapply

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mapply is a multivariate apply of sorts which applies a function in parallel over a set of arguments.

- FUN is a function to apply
- · ... contains arguments to apply over
- MoreArgs is a list of other arguments to FUN.
- SIMPLIFY indicates whether the result should be simplified

The following is tedious to type

```
list(rep(1, 4), rep(2, 3), rep(3, 2), rep(4, 1))
```

Instead we can do

```
> mapply(rep, 1:4, 4:1)
[[1]]
[1] 1 1 1 1

[[2]]
[1] 2 2 2

[[3]]
[1] 3 3
[[4]]
[1] 4
```

### **Vectorizing a Function**

```
> noise <- function(n, mean, sd) {
+ rnorm(n, mean, sd)
+ }
> noise(5, 1, 2)
[1] 2.4831198 2.4790100 0.4855190 -1.2117759
[5] -0.2743532
> noise(1:5, 1:5, 2)
[1] -4.2128648 -0.3989266 4.2507057 1.1572738
[5] 3.7413584
```

#### **Instant Vectorization**

```
> mapply(noise, 1:5, 1:5, 2)
[[1]]
[1] 1.037658
[[2]]
[1] 0.7113482 2.7555797
[[3]]
[1] 2.769527 1.643568 4.597882
[[4]]
[1] 4.476741 5.658653 3.962813 1.204284
[[5]]
[1] 4.797123 6.314616 4.969892 6.530432 6.723254
```

#### **Instant Vectorization**

Which is the same as

```
list(noise(1, 1, 2), noise(2, 2, 2),
    noise(3, 3, 2), noise(4, 4, 2),
    noise(5, 5, 2))
```



# Introduction to the R Language

Loop Functions - tapply

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tapply is used to apply a function over subsets of a vector. I don't know why it's called tapply.

```
> str(tapply)
function (X, INDEX, FUN = NULL, ..., simplify = TRUE)
```

- · x is a vector
- INDEX is a factor or a list of factors (or else they are coerced to factors)
- FUN is a function to be applied
- · ... contains other arguments to be passed FUN
- simplify, should we simplify the result?

Take group means.

Take group means without simplification.

```
> tapply(x, f, mean, simplify = FALSE)
$'1'
[1] 0.1144464

$'2'
[1] 0.5163468

$'3'
[1] 1.246368
```

Find group ranges.

```
> tapply(x, f, range)
$'1'
[1] -1.097309 2.694970

$'2'
[1] 0.09479023 0.79107293

$'3'
[1] 0.4717443 2.5887025
```

### split

split takes a vector or other objects and splits it into groups determined by a factor or list of factors.

```
> str(split)
function (x, f, drop = FALSE, ...)
```

- x is a vector (or list) or data frame
- f is a factor (or coerced to one) or a list of factors
- · drop indicates whether empty factors levels should be dropped

### split

```
> x <- c(rnorm(10), runif(10), rnorm(10, 1))
> f <- gl(3, 10)
> split(x, f)
$'1'
[1] -0.8493038 -0.5699717 -0.8385255 -0.8842019
[5] 0.2849881 0.9383361 -1.0973089 2.6949703
[9] 1.5976789 -0.1321970
$'2'
[1] 0.09479023 0.79107293 0.45857419 0.74849293
[5] 0.34936491 0.35842084 0.78541705 0.57732081
[9] 0.46817559 0.53183823
$'3'
[1] 0.6795651 0.9293171 1.0318103 0.4717443
 [5] 2.5887025 1.5975774 1.3246333 1.4372701
```

# split

A common idiom is split followed by an lapply.

```
> lapply(split(x, f), mean)
$'1'
[1] 0.1144464

$'2'
[1] 0.5163468

$'3'
[1] 1.246368
```

### **Splitting a Data Frame**

```
> library(datasets)
> head(airquality)
 Ozone Solar.R Wind Temp Month Day
    41
          190 7.4
                   67
                            1
1
    36
          118 8.0
                   72
3
   12
         149 12.6
                   74 5 3
    18
          313 11.5 62 5 4
4
                   56 5 5
5
    NA
          NA 14.3
    28
          NA 14.9
                         5 6
6
                   66
```

### **Splitting a Data Frame**

```
> s <- split(airquality, airquality$Month)</pre>
> lapply(s, function(x) colMeans(x[, c("Ozone", "Solar.R", "Wind")]))
$'5'
   Ozone Solar.R
                      Wind
      NA
               NA 11.62258
$'6'
            Solar.R
                         Wind
    Ozone
       NA 190.16667 10.26667
$171
              Solar.R
                            Wind
     Ozone
        NA 216.483871
                       8.941935
```

### **Splitting a Data Frame**

```
> sapply(s, function(x) colMeans(x[, c("Ozone", "Solar.R", "Wind")]))
                                    8
            5
Ozone
           NA
                   NA
                            NA
                                   NA
                                           NA
Solar.R
           NA 190.16667 216.483871
                                   NA 167.4333
Wind
      11.62258 10.26667 8.941935 8.793548 10.1800
> sapply(s, function(x) colMeans(x[, c("Ozone", "Solar.R", "Wind")],
                          na.rm = TRUE)
                         6
       23.61538 29.44444 59.115385 59.961538 31.44828
Ozone
Solar.R
       181.29630 190.16667 216.483871 171.857143 167.43333
Wind
```

#### Splitting on More than One Level

```
> x <- rnorm(10)
> f1 <- gl(2, 5)
> f2 <- gl(5, 2)
> f1
   [1] 1 1 1 1 1 2 2 2 2 2
Levels: 1 2
> f2
   [1] 1 1 2 2 3 3 4 4 5 5
Levels: 1 2 3 4 5
> interaction(f1, f2)
   [1] 1.1 1.1 1.2 1.2 1.3 2.3 2.4 2.4 2.5 2.5
10 Levels: 1.1 2.1 1.2 2.2 1.3 2.3 1.4 ... 2.5
```

# Splitting on More than One Level

Interactions can create empty levels.

```
> str(split(x, list(f1, f2)))
List of 10
$ 1.1: num [1:2] -0.378     0.445
$ 2.1: num(0)
$ 1.2: num [1:2]     1.4066     0.0166
$ 2.2: num(0)
$ 1.3: num -0.355
$ 2.3: num 0.315
$ 1.4: num(0)
$ 2.4: num [1:2] -0.907     0.723
$ 1.5: num(0)
$ 2.5: num [1:2] 0.732     0.360
```

# split

Empty levels can be dropped.

```
> str(split(x, list(f1, f2), drop = TRUE))
List of 6
$ 1.1: num [1:2] -0.378   0.445
$ 1.2: num [1:2] 1.4066 0.0166
$ 1.3: num -0.355
$ 2.3: num 0.315
$ 2.4: num [1:2] -0.907   0.723
$ 2.5: num [1:2] 0.732 0.360
```



# Debugging

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Indications that something's not right

- message: A generic notification/diagnostic message produced by the message function;
   execution of the function continues
- warning: An indication that something is wrong but not necessarily fatal; execution of the function continues; generated by the warning function
- error: An indication that a fatal problem has occurred; execution stops; produced by the stop function
- condition: A generic concept for indicating that something unexpected can occur; programmers can create their own conditions

#### Warning

```
log(-1)

## Warning: NaNs produced

## [1] NaN
```

```
printmessage <- function(x) {
    if(x > 0)
        print("x is greater than zero")
    else
        print("x is less than or equal to zero")
    invisible(x)
}
```

```
printmessage <- function(x) {
   if (x > 0)
      print("x is greater than zero") else print("x is less than or equal to zero")
   invisible(x)
}
printmessage(1)
```

```
## [1] "x is greater than zero"
```

```
printmessage(NA)
```

## Error: missing value where TRUE/FALSE needed

```
printmessage2 <- function(x) {
    if (is.na(x))
        print("x is a missing value!") else if (x > 0)
        print("x is greater than zero") else print("x is less than or equal to zero")
        invisible(x)
}
x <- log(-1)</pre>
```

```
## Warning: NaNs produced
```

```
printmessage2(x)
```

```
## [1] "x is a missing value!"
```

How do you know that something is wrong with your function?

- What was your input? How did you call the function?
- What were you expecting? Output, messages, other results?
- What did you get?
- How does what you get differ from what you were expecting?
- Were your expectations correct in the first place?
- · Can you reproduce the problem (exactly)?

#### **Debugging Tools in R**

The primary tools for debugging functions in R are

- traceback: prints out the function call stack after an error occurs; does nothing if there's no error
- debug: flags a function for "debug" mode which allows you to step through execution of a function one line at a time
- browser: suspends the execution of a function wherever it is called and puts the function in debug mode
- trace: allows you to insert debugging code into a function a specific places
- recover: allows you to modify the error behavior so that you can browse the function call stack

These are interactive tools specifically designed to allow you to pick through a function. There's also the more blunt technique of inserting print/cat statements in the function.

#### traceback

```
> mean(x)
Error in mean(x): object 'x' not found
> traceback()
1: mean(x)
>
```

#### traceback

```
> lm(y ~ x)
Error in eval(expr, envir, enclos) : object 'y' not found
> traceback()
7: eval(expr, envir, enclos)
6: eval(predvars, data, env)
5: model.frame.default(formula = y ~ x, drop.unused.levels = TRUE)
4: model.frame(formula = y ~ x, drop.unused.levels = TRUE)
3: eval(expr, envir, enclos)
2: eval(mf, parent.frame())
1: lm(y ~ x)
```

# debug

```
> debug(lm)
> lm(y ~ x)
debugging in: lm(y ~ x)
debug: {
    ret.x <- x
    ret.y <- y
    cl <- match.call()
    ...
    if (!qr)
        z$qr <- NULL
    z
}
Browse[2]>
```

# debug

#### recover

```
> options(error = recover)
> read.csv("nosuchfile")
Error in file(file, "rt") : cannot open the connection
In addition: Warning message:
In file(file, "rt") :
    cannot open file 'nosuchfile': No such file or directory

Enter a frame number, or 0 to exit

1: read.csv("nosuchfile")
2: read.table(file = file, header = header, sep = sep, quote = quote, dec = 3: file(file, "rt")

Selection:
```

# Debugging

#### Summary

- · There are three main indications of a problem/condition: message, warning, error
  - only an error is fatal
- · When analyzing a function with a problem, make sure you can reproduce the problem, clearly state your expectations and how the output differs from your expectation
- Interactive debugging tools traceback, debug, browser, trace, and recover can be used to find problematic code in functions
- Debugging tools are not a substitute for thinking!