

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 - 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
```



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
```

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

```
> 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
```

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
```

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
```