

mages' blog

Say it in R with "by", "apply" and friends

28 Jan 2012 14:43 🌑 aggregate , apply , by , data.table , doBy , language , plyr , R , sqldf , Tutorials











Iris versicolor

By Danielle Langlois

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R is a language, as Luis Apiolaza pointed out in his recent post. This is absolutely true, and learning a programming language is not much different from learning a foreign language. It takes time and a lot of practice to be proficient in it. I started using R when I moved to the UK and I wonder, if I have a better understanding of English or R by now.

Languages are full of surprises, in particular for non-native speakers. The other day I learned that there is *courtesy* and *curtsey*. Both words sounded very similar to me, but of course created some laughter when I mixed them up in an email.

With languages you can get into habits of using certain words and phrases, but sometimes you see or hear something, which shakes you up again. So did the following two lines in R with me:

```
f <- function(x) x^2 sapply(1:10, f)
[1] 1 4 9 16 25 36 49 64 81 100
```

It reminded me of the phrase that *everything is a list* in R. It showed me again how easily a *for* loop can be turned into a statement using the apply family of functions and how little I know about all the subtleties of R. I remember how happy I felt, when I finally understood the by function in R. I started to use it all the time, closing my eyes on aggregate and the apply functions family. Here is an example where I calculate the means of the various measurements by species of the famous iris data set using by .

by

```
do.call("rbind", as.list(
 by(iris, list(Species=iris$Species), function(x){
   y <- subset(x, select= -Species)</pre>
   apply(y, 2, mean)
 }
)))
          Sepal.Length Sepal.Width Petal.Length Petal.Width
                        3.428 1.462
                                                  0.246
setosa
               5.006
versicolor
                 5.936
                           2.770
                                        4.260
                                                    1.326
virginica
                 6.588
                            2.974
                                       5.552
                                                    2.026
```

Now let's find alternative ways of expressing ourselves, using other words/functions of the R language, such as aggregate, apply, sapply, tapply, data.table, ddply, sqldf, and summaryBy.

aggregate

The aggregate function splits the data into subsets and computes summary statistics for each of them. The output of aggregate is a data. frame, including a column for species.



```
      2 versicolor
      5.936
      2.770
      4.260
      1.326

      3 virginica
      6.588
      2.974
      5.552
      2.026
```

Addition: As John Christie points out in the comments, aggregate has also a formula interface, which simplifies the call to:

```
aggregate( . ~ Species, iris, mean)
```

apply and tapply

The combination of tapply and apply achieves a similar result, but this time the output is a matrix and hence I lose the column with species. The species are now the row names.

```
apply(iris.x, 2, function(x) tapply(x, iris.s, mean))
          Sepal.Length Sepal.Width Petal.Length Petal.Width
                         3.428
              5.006
                                  1.462
                                                 0.246
setosa
versicolor
                5.936
                          2.770
                                       4.260
                                                  1.326
virginica
                6.588
                          2.974
                                      5.552
                                                  2.026
```

split and apply

Here I split the data first into subsets for each of the species and calculate then the mean for each column in the subset. The output is a matrix again, but transposed.

```
sapply(split(iris.x, iris.s), function(x) apply(x, 2, mean))

setosa versicolor virginica

Sepal.Length 5.006 5.936 6.588

Sepal.Width 3.428 2.770 2.974

Petal.Length 1.462 4.260 5.552

Petal.Width 0.246 1.326 2.026
```

ddply

Hadley Wickham's plyr package provides tools for splitting, applying and combining data. The function ddply is similar to the by function, but it returns a data.frame instead of a by list and maintains the column for the species.

```
library(plyr)
ddply(iris, "Species", function(x){
   y <- subset(x, select= -Species)
   apply(y, 2, mean)
 })
    Species Sepal.Length Sepal.Width Petal.Length Petal.Width
            5.006 3.428 1.462
                                               0.246
                 5.936
                            2.770
                                        4.260
                                                   1.326
2 versicolor
                           2.974
                                                  2.026
3 virginica
                6.588
                                        5.552
```

Addition: Sean mentions in the comments an alternative, using the colMeans function, while Andrew reminds us of the reshape package with its functions melt and cast.

```
ddply(iris, "Species", function(x) colMeans(subset(x, select= -Species)))
## or
ddply(iris, "Species", colwise(mean))
## same output as above
library(reshape)
cast(melt(iris, id.vars='Species'),formula=Species ~ variable,mean)
## same output as above
```

summaryBy

The summaryBy function of the doBy package by Søren Højsgaard and Ulrich Halekoh has a very intuitive interface, using formulas.



```
library(doBy)
summaryBy(Sepal.Length + Sepal.Width + Petal.Length + Petal.Width ~ Species, data=iris, FUN=mean)
    Species Sepal.Length.mean Sepal.Width.mean Petal.Length.mean Petal.Width.mean
1
                       5.006
                                       3.428
                                                         1.462
                                                                          0.246
2 versicolor
                       5.936
                                        2.770
                                                          4.260
                                                                          1.326
3 virginica
                        6.588
                                        2.974
                                                          5.552
                                                                          2.026
```

sqldf

If you are fluent in SQL, then the sqldf package by Gabor Grothendieck might be the one for you.

```
library(saldf)
sqldf("select Species, avg(Sepal_Length), avg(Sepal_Width),
   avg(Petal_Length), avg(Petal_Width) from iris
   group by Species")
    Species avg(Sepal_Length) avg(Sepal_Width) avg(Petal_Length) avg(Petal_Width)
                                  3.428
1
                       5.006
                                                        1.462
     setosa
                                                                         0.246
2 versicolor
                       5.936
                                        2.770
                                                         4.260
                                                                         1.326
3 virginica
                        6.588
                                        2.974
                                                         5.552
                                                                          2.026
```

data.table

The data.table package by M Dowle, T Short and S Lianoglou is the real rock star to me. It provides an elegant and fast way to complete the task. The statement reads in plain English from right to left: take columns 1 to 4, split them by the factor in column "Species" and calculate on the sub data (.sD) the means.

```
library(data.table)
iris.dt <- data.table(iris)
iris.dt[,lapply(.SD,mean),by="Species",.SDcols=1:4]

Species Sepal.Length Sepal.Width Petal.Length Petal.Width
[1,] setosa 5.006 3.428 1.462 0.246
[2,] versicolor 5.936 2.770 4.260 1.326
[3,] virginica 6.588 2.974 5.552 2.026
```

apply

I should mention that R provides the iris data set also in an array form. The third dimension of the iris3 array holds the species information. Therefore I can use the apply function again, I go down the third and then the second dimension to calculate the means.

```
apply(iris3, c(3,2), mean)

Sepal L. Sepal W. Petal L. Petal W.

Setosa 5.006 3.428 1.462 0.246

Versicolor 5.936 2.770 4.260 1.326

Virginica 6.588 2.974 5.552 2.026
```

Conclusion

Many roads lead to Rome, and there are endless ways of explaining how to get there. I only showed a few I know of, and I am curious to hear yours. As a matter of *courtesy* I should mention the unknownR package by Matthew Dowle. It helps you to discover what you don't know that you don't know in R. Thus, it can help to build your R vocabulary. Of course there is a key difference between R and English. R tells me right away when I make a mistake. Human readers are far more forgiving, but please do point out to me where I made mistakes. I am still hopeful that I can improve, but I need your help.

R code



The R code of the examples is available on github. For more examples on the apply family see also Neil Saunders' post.

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