The doBy package

Søren Højsgaard

October 23, 2006

Contents

1	Intr	oduction	
2	Data		
	2.1	CO2 data	
	2.2	Airquality data	
3	The	summaryBy function	
	3.1	Basic usage	
	3.2	Using a list of functions	
	3.3	Copying variables out with the id argument	
	3.4	Statistics on functions of data	
	3.5	The dot (".") on the left hand side of a formula	
	3.6	The dot (".") on the right hand side of a formula	
	3.7	Using "1" on the right hand side of the formula	
	3.8	Preserving names of variables using keep.names	
4	The	orderBy function	
5	The	splitBy function	
6	The	sampleBy function	
7	The	subsetBy function	
8	The	transformBy function	
9	Miso 9.1	cellaneous The esticon function	
10	Fina	ıl remarks	

1 Introduction

The doBy package grew out of a need to calculate groupwise summary statistics in a simple way, much in the spirit of PROC SUMMARY of the SAS system. We have tried to keep the interface to the functions based on specifying formulas.

```
> library(doBy)

Hmisc library by Frank E Harrell Jr

Type library(help='Hmisc'), ?Overview, or ?Hmisc.Overview')
to see overall documentation.

NOTE:Hmisc no longer redefines [.factor to drop unused levels when subsetting. To get the old behavior of Hmisc type dropUnusedLevels().
```

2 Data

2.1 CO2 data

The CO2 data frame comes from an experiment on the cold tolerance of the grass species *Echinochloa crus-galli*. To limit the amount of output we modify names and levels of variables as follows

```
> data(CO2)
> CO2 <- transform(CO2, Treat = Treatment, Treatment = NULL)
> levels(CO2$Treat) <- c("nchil", "chil")
> levels(CO2$Type) <- c("Que", "Mis")
> CO2 <- subset(CO2, Plant %in% c("Qn1", "Qc1", "Mn1", "Mc1"))</pre>
```

whereby the data becomes

```
> CO2
  Plant Type conc uptake Treat
    Qn1 Que 95
                   16.0 nchil
                    30.4 nchil
    Qn1
         Que
              175
                   34.8 nchil
    Qn1 Que
              250
    Qn1 Que
              350
                   37.2 nchil
    Qn1 Que
             500
                   35.3 nchil
    On1 Que 675
                   39.2 nchil
6
    Qn1
         Que 1000
                    39.7 nchil
22
    Qc1 Que
              95
                   14.2 chil
23
    Qc1 Que 175
                   24.1 chil
24
    Qc1 Que 250
                   30.3 chil
25
    Qc1 Que
              350
                   34.6 chil
26
    Qc1
         Que
              500
                    32.5 chil
27
    Oct One
              675
                   35.4 chil
28
    Qc1 Que 1000
                   38.7 chil
43
    Mn1 Mis
              95
                   10.6 nchil
44
    Mn1
         Mis
              175
                   19.2 nchil
45
    Mn1
         Mis
              250
                   26.2 nchil
46
    Mn1 Mis
                   30.0 nchil
              350
47
    Mn1 Mis
              500
                   30.9 nchil
48
    Mn1 Mis 675
                   32.4 nchil
49
         Mis 1000
    Mn1
                    35.5 nchil
64
    Mc1
         Mis
              95
                   10.5 chil
    Mc1 Mis 175
65
                   14.9 chil
66
    Mc1 Mis 250
                    18.1 chil
67
    Mc1 Mis 350
                   18.9
                         chil
              500
68
    Mc1
         Mis
                    19.5
                         chil
69
    Mc1 Mis 675
                    22.2
                         chil
70
    Mc1 Mis 1000
                   21.9
                         chil
```

2.2 Airquality data

The airquality dataset contains air quality measurements in New York, May to September 1973. The months are coded as $5, \ldots, 9$. To limit the output we

only consider data for two months:

```
> airquality <- subset(airquality, Month %in% c(5, 6))
> head(airquality, n = 20)
  Ozone Solar.R Wind Temp Month Day
            190 7.4 67
2
     36
            118 8.0
                      72
3
     12
            149 12.6
                       74
     18
            313 11.5
                       62
                              5
4
     NA
             NA 14.3
                       56
6
     28
             NA 14.9
                       66
                              5
                                  6
     23
            299 8.6
                       65
                              5
8
      19
             99 13.8
                       59
                              5
                                8
      8
             19 20.1
                       61
                              5
                                 9
9
10
     NA
            194 8.6
                       69
                              5
                                 10
                       74
11
      7
             NA 6.9
                              5
                                11
12
     16
            256 9.7
                       69
                              5
                                12
13
     11
            290 9.2
                       66
                              5 13
14
            274 10.9
                       68
                              5 14
     14
15
     18
             65 13.2
                       58
                              5
                                15
16
     14
            334 11.5
                      64
                              5 16
            307 12.0
17
     34
                              5 17
18
      6
             78 18.4
                      57
                              5 18
19
     30
            322 11.5
                       68
                              5 19
20
     11
             44 9.7
                              5
                                 20
```

3 The summaryBy function

The summaryBy function is used for calculating quantities like "the mean and variance of x and y for each combination of two factors A and B". Examples are based on the CO2 data.

3.1 Basic usage

For example, the mean, median and variance of uptake and conc for each value of Plant is obtained by:

```
> summaryBy(conc + uptake ~ Plant, data = CO2, FUN = function(x) {
     c(m = mean(x), v = var(x))
+ })
 Plant conc.m conc.v uptake.m uptake.v
   Qn1
          435 100950 33.23
                                  67.48
   Qc1
          435 100950
                         29.97
                                  69.47
   Mn1
          435 100950
                         26.40
                                  75.59
          435 100950
                         18.00
                                  16.96
```

Defining the function to return named values as above is the recommended use of summaryBy. The function can also be defined outside the call to summaryBy:

```
> myfun1 <- function(x) {
+    c(m = mean(x), v = var(x))
+ }
> summaryBy(conc + uptake ~ Plant, data = CO2, FUN = myfun1)
```

If the result of the function(s) are not named, then the names in the output data in general become less intuitive:

```
> myfun2 <- function(x) {</pre>
      c(mean(x), var(x))
+ }
> summaryBy(conc + uptake ~ Plant, data = CO2, FUN = myfun2)
  Plant conc.myfun21 conc.myfun22 uptake.myfun21 uptake.myfun22
    Qn1
                 435
                            100950
                                             33.23
                                                             69.47
    Qc1
                 435
                            100950
                                             29.97
                            100950
                                             26.40
                                                             75.59
3
    Mn1
                 435
    Mc1
                  435
                            100950
                                             18.00
                                                             16.96
```

The postfix argument gives an altertive way of naming the output variables:

```
> summaryBy(conc + uptake ~ Plant, data = CO2, postfix = c("mymean",
      "myvar"), FUN = myfun2)
  Plant conc.myfun21 conc.myfun22 uptake.myfun21 uptake.myfun22
                           100950
                                            33.23
   0n1
                 435
    Qc1
                 435
                           100950
                                            29.97
                                                           69.47
                 435
                           100950
                                            26.40
                                                           75.59
3
    Mn1
                 435
                            100950
                                            18.00
                                                            16.96
```

3.2 Using a list of functions

It is possible to apply a list of functions. A typical usage will be by invoking a list of predefined functions:

```
> summaryBy(uptake ~ Plant, data = CO2, FUN = c(mean, var, median))
  Plant uptake.mean uptake.var uptake.median
   Qn1
              33.23
                          67.48
              29.97
                          69.47
    Qc1
                                         32.5
              26.40
                          75.59
3
    Mn1
                                         30.0
              18.00
                          16.96
                                         18.9
```

Slightly more elaborate is

```
> mymed <- function(x) c(med = median(x))</pre>
> summaryBy(uptake ~ Plant, data = CO2, FUN = c(mean, var, myfun1,
     myfun2))
 Plant uptake.mean uptake.var uptake.m uptake.v uptake.myfun21 uptake.myfun22
              33.23
                          67.48
                                   33.23
                                            67.48
                                                            33.23
   Qc1
                                   29.97
                                            69.47
                                                            29.97
                                                                            69.47
              29.97
                          69.47
              26.40
                          75.59
                                   26.40
                                            75.59
                                                            26.40
                                                                            75.59
    Mn1
                                   18.00
              18.00
                          16.96
                                            16.96
                                                            18.00
                                                                            16.96
    Mc1
```

The naming of the output variables determined from what the functions returns. The names of the last two columns above are imposed by summaryBy because myfun2 does not return named values. Specifying postfix= overrides these names but when FUN is a list of functions, the new names are not very informative either:¹

```
> summaryBy(uptake ~ Plant, data = CO2, postfix = c("aa", "bb",
      "cc"), FUN = c(mean, var, myfun1, myfun2))
  Plant uptake.aa uptake.bb uptake.m uptake.v uptake.myfun21 uptake.myfun22
    Ωn 1
            33.23
                      67.48
                                33.23
                                         67.48
                                                         33.23
                                                                        67.48
    Qc1
            29.97
                      69.47
                                29.97
                                         69.47
                                                         29.97
                                                                         69.47
                      75.59
                                                         26,40
3
    Mn1
            26.40
                                26.40
                                         75.59
                                                                        75.59
    Mc1
            18.00
                      16.96
                                18.00
                                         16.96
                                                         18.00
                                                                         16.96
```

¹This may be improved on later.

3.3 Copying variables out with the id argument

To get the value of the Type and Treat in the first row of the groups (defined by the values of Plant) copied to the output dataframe we use the id argument: as:

```
> summaryBy(conc + uptake ~ Plant, data = CO2, FUN = function(x) {
  c(m = mean(x), med = median(x), v = var(x))
+ }, id = ~Type + Treat)
 Plant conc.m conc.med conc.v uptake.m uptake.med uptake.v Type Treat
                                            35.3
                                 33.23
   Qn1
          435
                   350 100950
                                                    67.48 Que nchil
   Qc1
          435
                   350 100950
                                 29.97
                                             32.5
                                                     69.47
                                                           Que chil
3
   Mn1
          435
                   350 100950
                                 26.40
                                             30.0
                                                     75.59 Mis nchil
4
   Mc1
          435
                   350 100950
                                 18.00
                                             18.9
                                                     16.96
                                                           Mis chil
```

3.4 Statistics on functions of data

We may want to calculate the mean and variance for the logarithm of uptake, for uptake+conc (not likely to be a useful statistic) as well as for uptake and conc. This can be achieved as:

```
> summaryBy(log(uptake) + I(conc + uptake) + conc + uptake ~ Plant,
      data = CO2, FUN = function(x) {
          c(m = mean(x), v = var(x))
      })
 Plant log(uptake).m log(uptake).v conc+uptake.m conc+uptake.v conc.m conc.v
   0n1
                3.467
                            0.10168
                                            468.2
                                                         104747
                                                                    435 100950
   Qc1
                3.356
                            0.11873
                                            465.0
                                                         105297
                                                                    435 100950
   Mn1
                3.209
                            0.17928
                                            461.4
                                                         105642
                                                                    435 100950
                2.864
                            0.06874
                                            453.0
                                                         103157
                                                                    435 100950
   Mc1
 uptake.m uptake.v
    33.23
              67.48
     29.97
              69.47
3
     26.40
              75.59
     18.00
              16.96
```

If one does not want output variables to contain parentheses then setting p2d=TRUE causes the parentheses to be replaced by dots (".").

```
> summaryBy(log(uptake) + I(conc + uptake) ~ Plant, data = CO2,
     p2d = TRUE, FUN = function(x) {
          c(m = mean(x), v = var(x))
 Plant log.uptake..m log.uptake..v conc+uptake.m conc+uptake.v
                            0.10168
                                             468.2
    Qn1
                3.467
                                                           104747
                3.356
                            0.11873
                                             465.0
                                                           105297
    Qc1
    Mn1
                3.209
                             0.17928
                                             461.4
                                                           105642
4
    Mc1
                2.864
                             0.06874
                                             453.0
                                                           103157
```

3.5 The dot (".") on the left hand side of a formula

It is possible to use the dot (".") on the left hand side of the formula. The dot means "all numerical variables which do not appear elsewhere" (i.e. on the right hand side of the formula and in the id statement):

```
> summaryBy(log(uptake) + I(conc + uptake) + . ~ Plant, data = CO2,
     FUN = function(x)  {
         c(m = mean(x), v = var(x))
 Plant log(uptake).m log(uptake).v conc+uptake.m conc+uptake.v conc.m conc.v
                                           468.2
   Qn1
               3.467
                           0.10168
                                                        104747
                                                                  435 100950
   Qc1
               3.356
                           0.11873
                                           465.0
                                                        105297
                                                                  435 100950
                           0.17928
   Mn1
               3,209
                                           461.4
                                                        105642
                                                                  435 100950
   Mc1
               2.864
                           0.06874
                                           453.0
                                                        103157
                                                                  435 100950
 uptake.m uptake.v
    33.23
             67.48
    29.97
             69.47
    26.40
             75.59
3
    18.00
             16.96
```

3.6 The dot (".") on the right hand side of a formula

The dot (".") can also be used on the right hand side of the formula where it refers to "all non–numerical variables which are not specified elsewhere":

```
> summaryBy(log(uptake) ~ Plant + ., data = CO2, FUN = function(x) {
     c(m = mean(x), v = var(x))
+ })
 Plant Type Treat log(uptake).m log(uptake).v
                                     0.10168
   Qn1 Que nchil 3.467
                         3.356
                                     0.11873
   Qc1 Que chil
                         3.209
3
   Mn1 Mis nchil
                                     0.17928
        Mis chil
                          2.864
                                     0.06874
```

3.7 Using "1" on the right hand side of the formula

Using 1 on the right hand side means no grouping:

3.8 Preserving names of variables using keep.names

If the function applied to data only returns one value, it is possible to force that the summary variables retain the original names by setting keep.names=TRUE. A typical use of this could be

```
> summaryBy(conc + uptake + log(uptake) ~ Plant, data = CO2, FUN = mean,
+ id = ~Type + Treat, keep.names = TRUE)

Plant conc uptake log(uptake) Type Treat
1 Qn1 435 33.23 3.467 Que nchil
2 Qc1 435 29.97 3.356 Que chil
3 Mn1 435 26.40 3.209 Mis nchil
4 Mc1 435 18.00 2.864 Mis chil
```

4 The orderBy function

Ordering (or sorting) a data frame is possible with the orderBy function. Suppose we want to order the rows of the the airquality data by Temp and by Month (within Temp) and that the ordering should be decreasing. This can be achieved by:

```
> x <- orderBy(~Temp + Month, data = airquality, decreasing = T)
```

The first lines of the result are:

```
> head(x)
  Ozone Solar.R Wind Temp Month Day
             259 10.9
42
     NΑ
                        93
             250 9.2
43
      NA
                                   12
40
      71
             291 13.8
                        90
      NA
             273 6.9
                        87
                                   8
                               6
41
      39
             323 11.5
                        87
                                  10
                  8.6
```

5 The splitBy function

Suppose we want to split the airquality data into a list of dataframes, e.g. one dataframe for each month. This can be achieved by:

```
> x <- splitBy(~Month, data = airquality)
```

Information about the grouping is stored as a dataframe in an attribute called groupid:

```
> attr(x, "groupid")

Month
1    5
2    6
```

6 The sampleBy function

Suppose we want a random sample of 50 % of the observations from a dataframe. This can be achieved with:

```
> sampleBy(~1, frac = 0.5, data = airquality)
```

Suppose instead that we want a systematic sample of every fifth observation within each month. This is achieved with:

```
> sampleBy(~Month, frac = 0.2, data = airquality, systematic = T)
```

7 The subsetBy function

Suppose we want to take out those rows within each month for which the the wind speed is larger than the mean wind speed (within the month). This is achieved by:

```
> subsetBy(~Month, subset = "Wind>mean(Wind)", data = airquality)
```

Note that the statement "Wind>mean(Wind)" is evaluated within each month.

8 The transformBy function

The transformBy function is analogous to the transform function except that it works within groups. For example:

```
> transformBy(~Month, data = airquality, minW = min(Wind), maxW = max(Wind),
+ chg = sum(range(Wind) * c(-1, 1)))
```

9 Miscellaneous

9.1 The esticon function

Consider a linear model which explains Ozone as a linear function of Month and Wind:

```
> data(airquality)
> airquality <- transform(airquality, Month = factor(Month))
> m <- lm(Ozone ~ Month * Wind, data = airquality)
> coefficients(m)
(Intercept)
                Month6
                             Month7
                                         Month8
                                                      Month9
                                                                    Wind
    50.748
               -41.793
                             68.296
                                         82,211
                                                      23,439
                                                                  -2.368
Month6: Wind Month7: Wind Month8: Wind Month9: Wind
     4.051
                -4.663
                            -6.154
                                         -1.874
```

When a parameter vector β of (systematic) effects have been estimated, interest is often in a particular estimable function, i.e. linear combination $\lambda^{\top}\beta$ and/or testing the hypothesis $H_0: \lambda^{\top}\beta = \beta_0$ where λ is a specific vector defined by the user.

Suppose for example we want to calculate the expected difference in ozone between consequtive months at wind speed 10 mph (which is about the average wind speed over the whole period).

The esticon function provides a way of doing so. We can specify several λ vectors at the same time. For example

```
> Lambda
    [,1] [,2] [,3] [,4] [,5] [,6] [,7] [,8] [,9] [,10]
Γ1. ]
    0
        -1
             0 0 0 0 -10 0 0
                     0
                 0
                            0 10 -10
                  -1 0 0 0 10 -10
[3,]
      0
           0
             1
                                              0
> esticon(m, Lambda)
Confidence interval ( WALD ) level = 0.95
 betaO Estimate Std.Error t.value DF Pr(>|t|) Lower.CI Upper.CI
    0 1.2871 10.238 0.1257 106 0.90019 -19.010 21.585
               10.310 -2.2259 106 0.02814 -43.392
     0 -22.9503
                                                  -2.509
     0 0.9954
                 7.094 0.1403 106 0.88867 -13.069
                                                  15.060
                 6.560 2.4337 106 0.01662
```

In other cases, interest is in testing a hypothesis of a contrast $H_0: \Lambda \beta = \beta_0$ where Λ is a matrix. For example a test of no interaction between Month and Wind can be made by testing jointly that the last four parameters in m are zero (observe that the test is a Wald test):

```
> Lambda
     [,1] [,2] [,3] [,4] [,5] [,6] [,7] [,8] [,9] [,10]
[1,]
       0
                       0
                            0
                                 0
                                            0
                                       1
        0
                  0
                            0
                                  0
[2,]
             0
                       0
                                       0
                                            1
                                                 0
                                                       0
[3,]
        0
             0
                  0
                       0
                            0
                                  0
                                       0
                                            0
                                                 1
                                                       0
[4,]
        0
             0
                  0
                       0
                            0
                                  0
> esticon(m, Lambda, joint.test = T)
 X2.stat DF Pr(>|X^2|)
   22.11 4 0.0001906
```

For a linear normal model, one would typically prefer to do a likelihood ratio test instead. However, for generalized estimating equations of glm-type (as dealt with in the packages geepack and gee) there is no likelihood. In this case esticon function provides an operational alternative.

Observe that another function for calculating contrasts as above is the contrast function in the Design package but it applies to a narrower range of models than esticon does.

10 Final remarks

The esticon functions and other smaller functions are likely to be removed from the doBy package in the future. Credit is due to Dennis Chabot, Gabor Grothendieck, Paul Murrell and Erik Jørgensen for reporting various bugs and making various suggestions to the functionality in the doBy package.