Introduction to the **data.table** Package in R

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September 1, 2010

Introduction

This vignette is aimed at those who are already familiar with R—in particular, creating and using objects of class data.frame. We aim for this quick introduction to be readable in 10 minutes, covering the main features in brief, namely: 1. Keys; 2. Fast Grouping; and 3. Fast time series join. For the context that this document sits, please briefly check the last section, Further Resources.

data.table is not automatically better or faster. The user has to climb a short learning curve, experiment, and then use its features well. For example, this document explains the difference between a vector scan and a binary search. Although both extraction methods are available in data.table, if a user continues to use vector scans (as in a data.frame), it will 'work', but one will miss out on the benefits that data.table provides.

Creation

```
Recall that we create a data.frame using the function data.frame():
```

```
> df = data.frame(x=c("b","b","b","a","a"),v=rnorm(5))
> df
  x
1 b -2.2560769
2 b -0.9478947
3 b 1.4306219
4 a -0.7025186
5 a 0.3023313
A data.table is created in exactly the same way:
> dt = data.table(x=c("b","b","b","a","a"),v=rnorm(5))
> dt
     Х
[1,] b -2.7099061
[2,] b 0.3665731
[3,] b -0.2498943
[4,] a 0.7126236
[5,] a 0.4532576
```

Observe that a data.table prints the row numbers slightly differently. There is nothing significant about that. We can also convert existing data.frame objects to data.table.

```
> cars = data.table(cars)
> head(cars)

speed dist
[1,] 4 2
```

```
[2,] 4 10
[3,] 7 4
[4,] 7 22
[5,] 8 16
[6,] 9 10
```

We have just created two data.tables: dt and cars. It is often useful to see a list of all data.tables in memory:

```
> tables()
```

```
NAME NROW MB COLS KEY
[1,] cars 50 1 speed,dist
[2,] dt 5 1 x,v
Total: 2MB
```

The MB column is useful to quickly assess memory use and to spot if any redundant tables can be removed to free up memory. Just like data.frames, data.tables must fit inside RAM.

Some users regularly work with 20 or more tables in memory, rather like a database. The result of tables() is itself a data table, returned silently, so that tables() can be used in programs. tables() is unrelated to the base function table().

Also note that data.table() automatically converts character vectors to factors.

Users should rarely need to know this has occurred. See ?factor if you are unfamiliar with factors. Factors will appear to you as though they are character columns. You can refer to them just as though they are character.

You may have noticed the empty column KEY from tables() above. This is the subject of the next section, the first of the 3 main features of the package.

1. Keys

Let's start by considering data.frame, specifically rownames (or in English, row names). That is, the multiple names belonging to a single row. The multiple names belonging to the single row? That is not what we are used to in a data.frame. We know that each row has at most one name. A person has at least two names, a first name and a second name. That is useful to organise a telephone directory, for example, which is sorted by surname, then first name. However, each row in a data.frame can only have one name.

A key consists of one or more columns of rownames, which may be integer, factor or some other class, not simply character. Furthermore, the rows are sorted by the key. Therefore, a data.table can have at most one key, because it cannot be sorted in more than one way.

Uniqueness is not enforced, i.e., duplicate key values are allowed. Since the rows are sorted by the key, any duplicates in the key will appear consecutively.

Let's remind ourselves of our tables:

```
> tables()
```

> dt

```
NAME NROW MB COLS KEY
[1,] cars 50 1 speed,dist
[2,] dt 5 1 x,v
Total: 2MB
```

```
[1,] b -2.7099061
[2,] b 0.3665731
[3,] b -0.2498943
[4,] a 0.7126236
[5,] a 0.4532576
   No keys have been set yet. We can use data.frame syntax in a data.table, too.
> dt[2,]
[1,] b 0.3665731
> dt[dt$x == "b",]
     х
[1,] b -2.7099061
[2,] b 0.3665731
[3,] b -0.2498943
   But since there are no rownames, the following does not work:
> cat(try(dt["b",],silent=TRUE))
Error in `[.data.table`(dt, "b", ) :
  The data.table has no key but i is character. Call setkey first, see ?setkey.
   The error message tells us we need to use setkey():
> setkey(dt,x) # or key(dt)="x" if you prefer
> dt
[1,] a 0.7126236
```

Notice that the rows in dt have been re-ordered according to the values of x. The two "a" rows have moved to the top. We can confirm that dt does indeed have a key using haskey(), key(), attributes(), or just running tables().

> tables()

[2,] a 0.4532576 [3,] b -2.7099061 [4,] b 0.3665731 [5,] b -0.2498943

```
NAME NROW MB COLS KEY
[1,] cars 50 1 speed,dist
[2,] dt 5 1 x,v x
Total: 2MB
```

Now that we are sure dt has a key, let's try again:

```
> dt["b",]
x v
[1,] b -2.709906
```

Since there are duplicates in this key (i.e. repeated values of "b") the subset returns the first row in that group by default. The mult argument (short for multiple) controls this.

Lets now create a new data.frame. We will make it large enough to demonstrate the difference between a *vector scan* and a *binary search*.

```
> grpsize = ceiling(1e7/26^2)
                                # 10 million rows, 676 groups
[1] 14793
> tt=system.time( DF <- data.frame(
  x=rep(factor(LETTERS),each=26*grpsize),
   y=rep(factor(letters),each=grpsize),
   v=runif(grpsize*26^2))
+
  user system elapsed
 4.228
        1.884
                  6.810
> head(DF,3)
 х у
1 A a 0.2204772
2 A a 0.4393778
3 A a 0.8527393
> tail(DF,3)
        х у
10000066 Z z 0.32749413
10000067 Z z 0.45844677
10000068 Z z 0.06185507
> dim(DF)
[1] 10000068
```

We might say that R has created a 3 column table and $inserted\ 10,000,068$ rows. It took 6.810 secs, so it inserted 1,468,438 rows per second. This is normal in base R.

Let's extract an arbitrary group from the data.frame DF:

```
> tt=system.time(ans1 <- DF[DF$x=="R" & DF$y=="h",]) # 'vector scan'
user system elapsed
4.160  0.956  5.188
> head(ans1,3)
```

```
х у
6642058 R h 0.1109334
6642059 R h 0.9208031
6642060 R h 0.3780758
> dim(ans1)
[1] 14793
   Now convert to a data.table and extract the same group:
> DT = data.table(DF)
> setkey(DT,x,y)
> ss=system.time(ans2 <- DT[J("R", "h"), mult="all"])</pre>
                                                       # 'binary search'
   user system elapsed
  0.016
         0.000
> mapply(identical,ans1,ans2)
TRUE TRUE TRUE
```

At 0.015 seconds, this was **345** times faster than 5.188 seconds, and produced precisely the same result. If you are thinking that a few seconds is not much to save, it's the relative speedup that's important. The vector scan is linear, but the binary search is O(log n). It scales. If a task taking 10 hours is speed up by 100 times to 6 minutes, that is significant¹.

What does the J() do?

Was it really this, or was it something slow about using data.frame syntax in a data.table? It's exactly the same.

We can do vector scans in data.table, too:

```
> system.time(ans1 <- DF[DF$x=="R" & DF$y=="h",])
         system elapsed
   user
  4.168
         0.968
                  5.222
> system.time(ans2 <- DT[DT$x=="R" & DT$y=="h",])
  user system elapsed
         1.040
  4.184
                 5.277
> mapply(identical,ans1,ans2)
        У
TRUE TRUE TRUE
> system.time(ans3 <- DT[x=="R" & y=="h",])
  user system elapsed
  4.172
         1.060
                 5.287
> identical(ans2,ans3)
[1] TRUE
```

 $^{^{1}\}mathrm{We}$ wonder how many people are deploying parallel techniques to code that is vector scanning

If the phone book analogy helped, then this should not be surprising. We use the key to take advantage of the fact that the table is sorted and use binary search to find the matching rows. We didn't vector scan; we didn't use ==.

When we used DT\$x=="R" we scanned the entire column x, testing each and every value to see if it equalled "R". We did it again in the y column, testing for "h". Then & combined the two logical results to create a single logical vector which was passed to the [method, which in turn searched it for TRUE and returned those rows. These were vectorized operations. They occurred internally in R and were very fast, but they were scans. We did those scans because we wrote that R code.

When i is itself a data.table, we say that we are *joining* the two data.tables. In this case, we are joining DT to the 1 row, 2 column table returned by data.table("R", "h"). Since we do this a lot, there is an alias for data.tables called J(), short for join:

Both vector scanning and binary search are available in data.table, but one way of using data.table is much better than the other.

The join syntax is short, fast to write and easy to maintain. Passing a data.table into a data.table subset is similar to allowing a matrix to be passed into a matrix subset in base R.². There are other types of join and further arguments which are beyond the scope of this quick introduction.

The merge method of data.table is essentially x[y], but where the columns of x are included in the result. See FAQ 1.10.

This first section has been about the first argument to the [, namely i. The next section has to do with the 2nd argument j.

2. Fast grouping

The second argument to [is j, which may consist of one or more expressions whose arguments are (unquoted) column names, as if the column names were variables.

```
> dt[,sum(v)]
[1] -1.427346
```

> dt[,sum(v),by=x]

When we supply a j expression and a 'by' list of expressions, the jexpression is repeated for each 'by' group:

```
x V1
[1,] a 1.165881
[2,] b -2.593227

The 'by' in data.table is fast. Let's compare it to tapply:
> ttt=system.time(tt <- tapply(DT$v,DT$x,sum)); ttt
   user system elapsed
8.929   0.948   9.944
> sss=system.time(ss <- DT[,sum(v),by=x]); sss
   user system elapsed
   0.540   0.260   0.805</pre>
```

 $^{^2}$ Subsetting a keyed data.table by an n-column data.table is consistent with subsetting a n-dimension array by an n-column matrix

```
В
                          C
191928.8 192477.6 192365.6 192566.5 192650.3 192180.5
> head(ss)
             V1
     X
[1,] A 191928.8
[2,] B 192477.6
[3,] C 192365.6
[4,] D 192566.5
[5,] E 192650.3
[6,] F 192180.5
> identical(as.vector(tt), ss$V1)
[1] TRUE
   At 0.805 sec, this was 12 times faster than 9.944 sec, and produced precisely the same result.
   Next, let's group by two columns:
> ttt=system.time(tt <- tapply(DT$v,list(DT$x,DT$y),sum)); ttt
         system elapsed
   user
 10.349
         1.180 11.599
> sss=system.time(ss <- DT[,sum(v),by="x,y"]); sss
         system elapsed
   user
  0.556
          0.340
                  0.904
> tt[1:5,1:5]
                  b
         а
                            С
                                     d
A 7479.726 7374.629 7387.026 7355.091 7376.972
B 7334.284 7397.906 7420.707 7449.124 7363.144
C 7430.993 7399.268 7402.110 7375.195 7387.870
D 7351.393 7433.325 7444.239 7408.194 7433.960
E 7432.280 7416.055 7377.302 7415.135 7414.614
> head(ss)
    х у
[1,] A a 7479.726
[2,] A b 7374.629
[3,] A c 7387.026
[4,] A d 7355.091
[5,] A e 7376.972
[6,] A f 7346.525
> identical(as.vector(t(tt)), ss$V1)
[1] TRUE
```

This was 12 times faster, and the syntax a little simpler and easier to read.

The following features are mentioned only briefly here; further examples are in the FAQs.

• To return several expressions, pass a list() to j.

> head(tt)

- Each item of the list is recycled to match the length of the longest item.
- $\bullet\,$ You can pass a list () of expressions of column names to by.

3. Fast time series join

This is also known as last observation carried forward (LOCF) or a rolling join.

Recall that x[i] is a join between data.table x and data.table i. If i has 2 columns, the first column is matched to the first column of the key of x, and the 2nd column to the 2nd. An equi-join is performed, meaning that the values must be equal.

The syntax for fast rolling join is

x[i,roll=TRUE]

As before the first column of i is matched to x where the values are equal. The last column of i though, the 2nd one in this example, is treated specially. If no match is found, then the row before is returned, provided the first column still matches.

For examples see example("[.data.table")

Other resources

This was a quick start guide. Further resources include :

- The help page describes each and every argument: ?"[.data.table"
- The FAQs deal with distinct topics: vignette("datatable-faq")
- The performance tests contain more examples: vignette("datatable-timings")
- test.data.table contains over 150 low level tests of the features: test.data.table()
- Website: http://datatable.r-forge.r-project.org/
- Presentations:
 - http://files.meetup.com/1406240/Data%20munging%20with%20SQL%20and%20R.pdf
 - http://www.londonr.org/LondonR-20090331/data.table.LondonR.pdf
- YouTube Demo: http://www.youtube.com/watch?v=rvT8XThGA8o
- R-Forge commit logs: http://lists.r-forge.r-project.org/pipermail/datatable-commits/
- Mailing list: datatable-help@lists.r-forge.r-project.org
- User reviews : http://crantastic.org/packages/data-table