FAQs about the ${\bf data.table}$ package in R

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The first section, Beginner FAQs, is intended to be read in order, from start to finish. It may be read before reading the 10 minute introduction to data.table vignette.

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Beginner FAQs 1

1.1 Why does DT[,5] return 5?

Because, by default, unlike a data.frame, the 2nd argument is an expression which is evaluated within the scope of DT. 5 evaluates to 5. It is generally bad practice to refer to columns by number rather than name. If someone else comes along and reads your code later, they may have to hunt around to find out which column is number 5. Furthermore, if you or someone else changes the column ordering of DT higher up in your R program, you might get bugs if you forget to change all the places in your code which refer to column number 5.

6.10 Why is this FAQ in pdf format? Can it moved to a website?

Say column 5 is called "region", just do DT[,region] instead. Notice there are no quotes around the column name. This is what we mean by j being evaluated within the scope of the data.table. That scope consists of an environment where the column names are variables.

You can write any R expression in the j; e.g., DT[,colA*colB/2]. Further, j may be a list() of many R expressions, including calls to any R package; e.g., DT[,fitdistr(d1-d1,"normal")].

Having said this, there are some circumstances where referring to a column by number is ok, such as a sequence of columns. In these situations just do DT[,5:10,with=FALSE] or DT[,c(1,4,10),with=FALSE]. See ?data.table for an explanation of the with argument.

Note that with() has been a base function for a long time. That's one reason we say data.table builds upon base functionality. There is little new here really, data.table is just making use of with() and building it into the syntax.

1.2 Why does DT[, "region"] return "region"?

See answer to 1.1 above. Try DT[,region] instead. Or DT[,"region",with=FALSE].

1.3 Why does DT[,region] return a vector? I'd like a 1-column data.table. There is no drop argument like I'm used to in data.frame.

Try DT[,list(region)] instead.

1.4 Why does DT[,x,y,z] not work? I wanted the 3 columns x,y and z.

The j expression is the 2nd argument. The correct way to do this is DT[,list(x,y,z)].

1.5 I assigned a variable mycol="x" but then DT[,mycol] returns "x". How do I get it to look up the column name contained in the mycol variable?

This is what we mean when we say the j expression 'sees' objects in the calling scope. The variable mycol does not exist as a column name of DT so R then looked in the calling scope and found mycol there, and returned its value "x". This is correct behaviour. Had mycol been a column name, then that column's data would have been returned. What you probably meant was DT[,mycol,with=FALSE], which will return the x column's data as you wanted. Alternatively, since a data.table is a list, too, you can write DT[["x"]] or DT[[mycol]].

1.6 Ok but I don't know the expressions in advance. How do I programatically pass them in?

To create expressions use the quote() function. We refer to these as quote()-ed expressions to save confusion with the double quotes used to create a character vector such as c("x"). The simplest quote()-ed expression is just one column name :

```
q = quote(x)
DT[,eval(q)] # returns the column x as a vector
q = quote(list(x))
DT[,eval(q)] # returns the column x as a 1-column data.table
Since these are expressions, we are not restricted to column names only:
q = quote(mean(x))
DT[,eval(q)] # identical to DT[,mean(x)]
q = quote(list(x,sd(y),mean(y*z)))
DT[,eval(q)] # identical to DT[,list(x,sd(y),mean(y*z))]
```

However, if it's just simply a vector of column names you need, it may be simpler to pass a character vector to j and use with=FALSE.

To pass an expression into your own function, one idiom is as follows:

```
> DT = as.data.table(iris)
> setkey(DT,Species)
> myfunction = function(dt, expr) {
+     e = substitute(expr)
```

1.7 This is really hard. What's the point?

j doesn't have to be just column names. You can write any R expression of column names directly as the j; e.g., DT[,mean(x*y/z)]. The same applies to i; e.g., DT[x>1000, sum(y*z)]. This runs the j expression on the set of rows where the i expression is true. You don't even need to return data; e.g., DT[x>1000, plot(y,z)]. When we get to compound table joins we will see how i and j can themselves be other data.table queries. We are going to stretch i and j much further than this, but to get there we need you on board first with FAQs 1.1-1.6.

1.8 OK, I'm starting to see what data.table is about, but why didn't you enhance data.frame in R? Why does it have to be a new package?

As FAQ 1.1 highlights, j in [.data.table is fundamentally different from j in [.data.frame. Even something as simple as DF[,1] would break existing code in many packages and user code. This is by design, and we want it to work this way for more complicated syntax to work. There are other differences, too (see FAQ 2.17).

Furthermore, data.table *inherits* from data.frame. It is a data.frame, too. A data.table can be passed to any package that only accepts data.frame and that package can use [.data.frame syntax on the data.table.

We have proposed enhancements to R wherever possible, too. One of these was accepted as a new feature in R 2.12.0 :

unique() and match() are now faster on character vectors where all elements are in the global CHARSXP cache and have unmarked encoding (ASCII). Thanks to Matthew Dowle for suggesting improvements to the way the hash code is generated in unique.c.

A second proposal was to use memcpy in duplicate.c, which is much faster than a for loop in C. This would improve the way that R copies data internally (on some measures by 13 times). The thread on r-devel is here: http://tolstoy.newcastle.edu.au/R/e10/devel/10/04/0148.html.

1.9 Why are the defaults the way they are? Why does it work the way it does?

The simple answer is because the author designed it for his own use, and he wanted it that way. He finds it a more natural, faster way to write code, which also executes more quickly.

1.10 Isn't this already done by with() and subset() in base?

Some of the features discussed so far are, yes. The package builds upon base functionality. It does the same sorts of things but with less code required, and executes many times faster if used correctly.

1.11 Why does X[Y] return all the columns from Y too? Shouldn't it return a subset of X?

This was changed in v1.5.3. X[Y] now includes Y's non-join columns. We refer to this feature as join inherited scope because not only are X columns available to the j expression, so are Y columns. The downside is that X[Y] is less efficient since every item of Y's non-join columns are duplicated to

match the (likely large) number of rows in X that match. We therefore strongly encourage X[Y,j] instead of X[Y]. See next FAQ.

1.12 What is the difference between X[Y] and merge(X,Y)?

X[Y] is a join, looking up X's rows using Y (or Y's key if it has one) as an index. Y[X] is a join, looking up Y's rows using X (or X's key if it has one) as an index. merge(X,Y) does both ways at the same time. The number of rows of X[Y] and Y[X] usually differ; whereas the number of rows returned by merge(X,Y) and merge(Y,X) is the same.

BUT that misses the main point. Most tasks require something to be done on the data after a join or merge. Why merge all the columns of data, only to use a small subset of them afterwards? You may suggest merge(X[,ColsNeeded1],Y[,ColsNeeded2]), but that takes copies of the subsets of data, and it requires the programmer to work out which columns are needed. X[Y,j] in data.table does all that in one step for you. When you write X[Y,sum(foo*bar)], data.table automatically inspects the j expression to see which columns it uses. It will only subset those columns only; the others are ignored. Memory is only created for the columns the j uses, and Y columns enjoy standard R recycling rules within the context of each group. Let's say foo is in X, and bar is in Y (along with 20 other columns in Y). Isn't X[Y,sum(foo*bar)] quicker to program and quicker to run than a merge followed by a subset?

1.13 Anything else about X[Y,sum(foo*bar)]?

Remember that j (in this example sum(foo*bar)) is run for each group of X that each row of Y matches to. This feature is grouping by i or by without by. For example, and making it complicated by using join inherited scope, too:

```
> X = data.table(grp=c("a", "a", "b", "b", "b", "c", "c"), foo=1:7)
> setkey(X,grp)
> Y = data.table(c("b","c"), bar=c(4,2))
> X
     grp foo
[1,]
       a
            1
[2,]
            2
       a
[3,]
            3
       b
[4,]
            4
       b
            5
[5,]
       b
[6,]
            6
       С
            7
[7,]
> Y
     V1 bar
[1,]
      b
           4
[2,]
> X[Y,sum(foo*bar)]
     grp V1
       b 48
[1,]
[2,]
       c 26
```

1.14 That's nice but what if I really do want to evaluate j for all rows once, not by group?

If you really want j to run once for the whole subset of X then try X[Y][,sum(foo*bar)]. If that needs to be efficient (recall that X[Y] joins all columns) then you will have to work a little harder since this is outside the common use-case: X[Y,list(foo,bar)][,sum(foo*bar)].

2 General syntax

2.1 How can I avoid writing a really long j expression? You've said I should use the column *names*, but I've got a lot of columns.

When grouping, the j expression can use column names as variables, as you know, but it can also use a reserved symbol .SD which refers to the subset of the data.table for each group (excluding the grouping columns). So to sum up all your columns it's just DT[,lapply(.SD,sum),by=grp]. It might seem tricky, but it's fast to write and fast to run. Notice you don't have to create an anonymous function. See the timing vignette and wiki for comparison to other methods. The .SD object is efficiently implemented internally and more efficient than passing an argument to a function. Please don't do this though: DT[,.SD[,"sales",with=FALSE],by=grp]. That works but is very inefficient and inelegant. This is what was intended: DT[,sum(sales),by=grp] and could be 100's of times faster.

2.2 Why is the default for mult now "all"?

In v1.5.3 the default was changed to "all". When i (or i's key if it has one) has fewer columns than x's key, mult was already set to "all" automatically. Changing the default makes this clearer and easier for users as it came up quite often.

In versions up to v1.3, "all" was slower. Internally, "all" was implemented by joining using "first", then again from scratch using "last", after which a diff between them was performed to work out the span of the matches in x for each row in i. Most often we join to single rows, though, where "first", "last" and "all" return the same result. We preferred maximum performance for the majority of situations so the default chosen was "first". When working with a non-unique key (generally a single column containing a grouping variable), DT["A"] returned the first row of that group so DT["A", mult="all"] was needed to return all the rows in that group.

In v1.4 the binary search in C was changed to branch at the deepest level to find first and last. That branch will likely occur within the same final pages of RAM so there should no longer be a speed disadvantage in defaulting mult to "all". We warned that the default might change, and made the change in v1.5.3.

A future version of data.table may allow a distinction between a key and a unique key. Internally mult="all" would perform more like mult="first" when all x's key columns were joined to and x's key was a unique key. data.table would need checks on insert and update to make sure a unique key is maintained. An advantage of specifying a unique key would be that data.table would ensure no duplicates could be inserted, in addition to performance.

2.3 I'm using c() in the j and getting strange results.

This is a common source of confusion. In data.frame you are used to, for example:

```
> DF = data.frame(x=1:3,y=4:6,z=7:9)
> DF

    x y z
1 1 4 7
2 2 5 8
3 3 6 9
> DF[,c("y","z")]
    y z
1 4 7
2 5 8
3 6 9
```

which returns the two columns. In data.table you know you can use the column names directly and might try:

```
> DT = data.table(DF)
> DT[,c(y,z)]
[1] 4 5 6 7 8 9
```

but this returns one vector. Remember that the j expression is evaluated within the environment of DT, and c() returns a vector. If 2 or more columns are required, use list() instead:

```
> DT[,list(y,z)]
y z
[1,] 4 7
[2,] 5 8
[3,] 6 9
```

c() can be useful in a data.table too, but its behaviour is different from that in [.data.frame.

2.4 I have built up a complex table with many columns. I want to use it as a template for a new table; i.e., create a new table with no rows, but with the column names and types copied from my table. Can I do that easily?

Yes. If your complex table is called DT, try NEWDT = DT[0].

2.5 Is a NULL data.table the same as DT[0]?

No, despite the print method indicating otherwise. Strictly speaking it's not possible to have is.null(data.table(NULL)) return FALSE. This FAQ may be revisited in future.

2.6 Why has the DT() alias been removed?

DT was introduced originally as a wrapper for a list of j expressions. Since DT was an alias for data.table, this was a convenient way to take care of silent recycling in cases where each item of the j list evaluated to different lengths. The alias was one reason grouping was slow, though. As of v1.3, list() should be passed instead to the j argument. list() is a primitive and is much faster, especially when there are many groups. Internally, this was a nontrivial change. Vector recycling is now done internally, along with several other speed enhancements for grouping.

2.7 But my code uses j=DT(...) and it works. The previous FAQ says that DT() has been removed.

Then you are using a version prior to 1.5.3. Prior to 1.5.3 [.data.table detected use of DT() in the j and automatically replaced it with a call to list(). This was to help the transition for existing users.

2.8 What are the scoping rules for j expressions?

Think of the subset as an environment where all the column names are variables. When a variable foo is used in the j of a query such as X[Y,sum(foo)], foo is looked for in the following order:

- 1. The scope of X's subset; i.e., X's column names.
- 2. The scope of each row of Y; i.e., Y's column names (join inherited scope)
- 3. The scope of the calling frame; e.g., the line that appears before the data.table query.
- 4. Exercise for reader: does it then ripple up the calling frames, or go straight to globalenv()?
- 5. The global environment

This is *lexical scoping* as explained in R FAQ 3.3.1. The environment in which the function was created is not relevant, though, because there is *no function*. No anonymous *function* is passed to the j. Instead, an anonymous *body* is passed to the j; for example,

Some programming languages call this a lambda.

2.9 Can I trace the j expression as it runs through the groups?

Try something like this:

2.10 Inside each group, why is the group variable a long vector containing the same value repeated?

Please upgrade to v1.6.1, or later; this is no longer true. In the previous FAQ, x is a grouping variable and now has length 1 for efficiency and convenience. Prior to v1.6.1, x repeated the group value to match the number of rows in that group. There is no longer any difference between the following two statements.

Code written prior to v1.6.1 that uses [1] will still work, but the [1] is no longer necessary.

2.11 Only the first 10 rows are printed, how do I print more?

Try print(DT,nrows=Inf) to print all rows, or set nrows to the number of rows you require. Recall that when you forget a data.frame is large and type its name at the R prompt, R appears to hang for a long time while the entire table is formatted. data.table catches this and just prints the first 10 rows of large tables by default.

2.12 With an X[Y] join, what if X contains a column called "Y"?

When i is a single name such as Y it is evaluated in the calling frame. In all other cases such as calls to J() or other expressions, i is evaluated within the scope of X. This facilitates easy *self joins* such as X[J(unique(colA)),mult="first"].

2.13 X[Z[Y]] is failing because X contains a column "Y". I'd like it to use the table Y in calling scope.

The Z[Y] part is not a single name so that is evaluated within the frame of X and the problem occurs. Try tmp=Z[Y]; X[tmp]. This is robust to X containing a column "tmp" because tmp is a single name. If you often encounter conflics of this type, one simple solution may be to name all tables in uppercase and all column names in lowercase, or some similar scheme.

2.14 Can you explain further why data.table is inspired by A[B] syntax in base?

Consider A[B] syntax using an example matrix A:

```
> A = matrix(1:12, nrow=4)
> A
      [,1] [,2] [,3]
[1,]
         1
              5
[2,]
         2
               6
                   10
[3,]
         3
              7
                   11
         4
[4,]
                   12
```

To obtain cells (1,2)=5 and (3,3)=11 many users (we believe) may try this first :

```
> A[c(1,3),c(2,3)]

[,1] [,2]

[1,] 5 9

[2,] 7 11
```

That returns the union of those rows and columns, though. To reference the cells, a 2-column matrix is required. ?Extract says:

When indexing arrays by [a single argument i can be a matrix with as many columns as there are dimensions of x; the result is then a vector with elements corresponding to the sets of indices in each row of i.

Let's try again.

[1] 5 11

A matrix is a 2-dimension structure with row names and column names. Can we do the same with names?

```
> rownames(A) = letters[1:4]
> colnames(A) = LETTERS[1:3]
  A B C
a 1 5 9
b 2 6 10
c 3 7 11
d 4 8 12
> B = cbind(c("a", "c"), c("B", "C"))
> A[B]
[1] 5 11
So, yes we can. Can we do the same with data.frame?
> A = data.frame(A=1:4,B=letters[11:14],C=pi*1:4)
> rownames(A) = letters[1:4]
> A
  A B
              C
a 1 k 3.141593
b 2 1 6.283185
c 3 m 9.424778
d 4 n 12.566371
> B
     [,1] [,2]
[1,] "a" "B"
[2,] "c"
          "C"
> A[B]
[1] "k"
                " 9.424778"
```

But, notice that the result was coerced to character. R coerced A to matrix first so that the syntax could work, but the result isn't ideal. Let's try making B a data.frame.

```
> B = data.frame(c("a","c"),c("B","C"))
> cat(try(A[B],silent=TRUE))

Error in `[.default`(A, B) : invalid subscript type 'list'
```

So we can't subset a data.frame by a data.frame in base R. What if we want row names and column names that aren't character but integer or float? What if we want more than 2 dimensions of mixed types? Enter data.table.

Furthermore, matrices, especially sparse matrices, are often stored in a 3 column tuple: (i,j,value). This can be thought of as a key-value pair where i and j form a 2-column key. If we have more than one value, perhaps of different types it might look like (i,j,val1,val2,val3,...). This looks very much like a data.frame. Hence data.table extends data.frame so that a data.frame X can be subset by a data.frame Y, leading to the X[Y] syntax.

2.15 Can base be changed to do this then, rather than a new package?

data.frame is used everywhere and so it is very difficult to make any changes to it. data.table inherits from data.frame. It is a data.frame, too. A data.table can be passed to any package that only accepts data.frame. When that package uses [.data.frame syntax on the data.table, it works.

2.16 I've heard that data.table syntax is analogous to SQL.

Yes:

- i <==> where
- j <==> select
- by $\langle == \rangle$ group by
- i <==> order by (in compound syntax)
- i <==> having (in compound syntax)
- nomatch=NA <==> outer join
- nomatch=0 <==> inner join
- \bullet mult="first"|"last" <==> N/A because SQL is inherently unordered
- roll=TRUE <==> N/A because SQL is inherently unordered

The general form is:

```
DT[where, select, group by] [having] [order by] []...[]
```

A key advantage of column vectors in R is that they are *ordered*, unlike SQL¹. We can use ordered functions in data.table queries, such as diff(), and we can use *any* R function from any package, not just the functions that are defined in SQL. A disadvantage is that R objects must fit in memory, but with several R packages such as ff, bigmemory and indexing, this is changing.

2.17 What are the smaller syntax differences between data.frame and data.table?

- DT[3] refers to the 3rd row, but DF[3] refers to the 3rd column
- DT[3,] == DT[3], but DF[3] == DF[3] (somewhat confusingly)
- For this reason we say the comma is optional in DT, but not optional in DF
- DT[[3]] == DF[3] == DF[[3]]
- DT[i,] where i is a single integer returns a single row, just like DF[i,], but unlike a matrix single row subset which returns a vector.
- DT[,j,with=FALSE] where j is a single integer returns a one column data.table, unlike DF[,j] which returns a vector by default
- DT[, "colA", with=FALSE][[1]] == DF[, "colA"].
- DT[,colA] == DF[,"colA"]
- DT[,list(colA)] == DF[,"colA",drop=FALSE]

¹It may be a surprise to learn that select top 10 * from ... does *not* reliably return the same rows over time in SQL. You do need to include an order by clause, or use a clustered index to guarantee row order; i.e., SQL is inherently unordered.

- DT[NA] returns 1 row of NA, but DF[NA] returns a copy of DF containing NA throughout. The symbol NA is type logical in R, and is therefore recycled by [.data.frame. Intention was probably DF[NA_integer_]. [.data.table does this automatically for convenience.
- DT[c(TRUE, NA, FALSE)] treats the NA as FALSE, but DF[c(TRUE, NA, FALSE)] returns NA rows for each NA
- DT[ColA==ColB] is simpler than DF[!is.na(ColA) & !is.na(ColB) & ColA==ColB,]

In [.data.frame we very often set drop=FALSE. When we forget, bugs can arise in edge cases where single columns are selected and all of a sudden a vector is returned rather than a single column data.frame. In [.data.table we took the opportunity to make it consistent and drop drop.

When you pass a data.table to a data.table-unaware package, that package it not concerned with any of these differences. It just works.

2.18 I'm using j for its side effect only, but I'm still getting data returned. How do I stop that?

In this case j can be wrapped with invisible(); e.g., DT[,invisible(hist(colB)),by=colA]².

2.19 Why does [.data.table now have a drop argument from v1.5?

So that data.table can inherit from data.frame without using If we used ... then invalid argument names would not be caught.

The drop argument is never used by [.data.table. It is a placeholder for non data.table aware packages when they use the [.data.frame syntax directly on a data.table.

2.20 Rolling joins are cool, and very fast! Was that hard to program?

The prevailing row on or before the i row is the final row the binary search tests anyway. So roll=TRUE is essentially just a switch in the binary search C code to return that row.

3 Questions relating to compute time

3.1 I have 20 columns and a large number of rows. Why is an expression of one column so quick?

Several reasons:

- Only that column is grouped, the other 19 are ignored because data.table inspects the j expression and realises it doesn't use the other columns.
- One memory allocation is made for the largest group only, then that memory is re-used for the other groups. There is very little garbage to collect.
- R is an in-memory column store; i.e., the columns are contiguous in RAM. Page fetches from RAM into L2 cache are minimised.

3.2 I don't have a key on a large table, but grouping is still really quick. Why is that?

data.table uses radix sorting. This is significantly faster than other sort algorithms. Radix is specifically for integers only, see ?base::sort.list(x,method="radix").

This is also one reason why setkey() is quick.

When no key is set, or we group in a different order from that of the key, we call it an $ad\ hoc$ by.

²hist() returns the breakpoints in addition to plotting to the graphics device

3.3 Why is grouping by columns in the key faster than an ad hoc by?

Because each group is contiguous in RAM, thereby minimising page fetches, and memory can be copied in bulk (memory in C) rather than looping in C.

4 Error messages

4.1 Could not find function "DT"

```
See FAQ 2.6 and FAQ 2.7.
```

4.2 unused argument(s) (MySum = sum(v))

This error is generated by DT[,MySum=sum(v)]. DT[,list(MySum=sum(v))] was intended, or DT[,j=list(MySum=sum(v))].

4.3 'translateCharUTF8' must be called on a CHARSXP

This error (and similar; e.g., 'getCharCE' must be called on a CHARSXP) may be nothing do with character data or locale. Instead, this can be a symptom of an earlier memory corruption. To date these have been reproducible and fixed (quickly). Please report it to datatable-help.

5 Warning messages

- 5.1 The following object(s) are masked from 'package:base': cbind, rbind This warning appears when loading the package. For detailed information on why they are masked, please type ?cbind and click the data.table version of the help page.
- 5.2 Coerced numeric RHS to integer to match the column's type; may have truncated precision. Either change the column to numeric first by creating a new numeric vector length 5 (nrows of entire table) yourself and assigning that (i.e. 'replace' column), or coerce RHS to integer yourself (e.g. 1L or as.integer) to make your intent clear (and for speed). Or, set the column type correctly up front when you create the table and stick to it, please.

Hopefully, this is self explanatory. To generate it, try:

```
> DT = data.table(a=1:5,b=1:5)
> suppressWarnings(
+ DT[2,b:=6]
                    # works (slower) with warning - suppressed for this vignette
+ )
     a b
[1,]11
[2,] 2 6
[3,] 3 3
[4,] 4 4
[5,] 5 5
> DT[2,b:=7L]
                    # works (faster) without warning
     a b
[1,] 1 1
[2,] 27
```

```
[3,] 3 3
```

[4,] 4 4

[5,] 5 5

> DT[,b:=rnorm(5)] # 'replace' integer column with a numeric column

```
a b
```

- [1,] 1 1.3788713
- [2,] 2 0.7345511
- [3,] 3 0.3135421
- [4,] 4 1.2625627
- [5,] 5 1.8348232

6 General questions about the package

6.1 v1.3 appears to be missing from the CRAN archive?

That is correct. v1.3 was available on R-Forge only. There were several large changes internally and these took some time to test in development.

6.2 Is data.table compatible with S-plus?

Not currently.

- A few core parts of the package are written in C and use internal R functions and R structures.
- The package uses lexical scoping which is one of the differences between R and S-plus explained by R FAQ 3.3.1.

6.3 Is it available for Linux, Mac and Windows?

Yes, for both 32-bit and 64-bit on all platforms. Thanks to CRAN and R-Forge. There are no special or OS-specific libraries used.

6.4 I think it's great. What can I do?

Please send suggestions, bug reports and enhancement requests to datatable-help. This helps make the package better. The list is public and archived.

Please do vote for the package on Crantastic. This helps encourage the developers, and helps other R users find the package. If you have time to write a comment too, that can help others in the community. Just simply clicking that you use the package, though, is much appreciated.

You can join the project and change the code and/or documentation yourself.

6.5 I think it's not great. How do I warn others about my experience?

Please put your vote and comments on Crantastic. Please make it constructive so we have a chance to improve.

6.6 I have a question. I know the posting guide tells me to contact the maintainer (not r-help), but is there a larger group of people I can ask?

Yes. You can post to datatable-help. It's like r-help, but just for this package. Feel free to answer questions there, too. maintainer("<package name>") returns the maintainer for any package and is where you should ask first (see R-help posting guide). maintainer("data.table") returns datatable-help.

6.7 Where are the datatable-help archives?

The homepage contains links to the archives in 4 different formats.

6.8 I'd prefer not to contact datatable-help, can I mail just one or two people privately?

Sure. You're more likely to get a faster answer from datatable-help, though.

6.9 I have created a package that depends on data.table. How do I ensure my package is data.table-aware so that inheritance from data.frame works?

You don't need to do anything special. As long as your package has a namespace, and imports or depends on data.table, your package is detected as data.table-aware.

6.10 Why is this FAQ in pdf format? Can it moved to a website?

This FAQ (and the intro and timing documents) are *vignettes* written using Sweave. The benefits of Sweave include the following:

- We include R code in the vignettes. This code is *actually run* when the file is created, not copy and pasted.
- This document is reproducible. Grab the .Rnw and you can run it yourself.
- CRAN checks the package (including running vignettes) every night on Linux, Mac and Windows, both 32bit and 64bit. Results are posted to http://cran.r-project.org/web/checks/check_results_data.table.html. Included there are results from r-devel; i.e., not yet released R. That serves as a very useful early warning system for any potential future issues as R itself develops.
- This file is bound into each version of the package. The package is not accepted on CRAN unless this file passes checks. Each version of the package will have its own FAQ file which will be relevant for that version. Contrast this to a single website, which can be ambiguous if the answer depends on the version.
- You can open it offline at your R prompt using vignette().
- You can extract the code from the document and play with it using edit(vignette("datatable-faq")) or edit(vignette("datatable-timings")).
- It prints out easily.
- It's quicker and easier for us to write and maintain the FAQ in .Rnw form.

Having said all that, a wiki format may be quicker and easier for users to contribute documentation and examples. Therefore a wiki has now been created; see link on the homepage.