

Data Transformation with `data.table` :: CHEAT SHEET



Basics

`data.table` is an extremely fast and memory efficient package for transforming data in R. It works by converting R's native data frame objects into `data.tables` with new and enhanced functionality. The basics of working with `data.tables` are:

`dt[i, j, by]`

Take data.table `dt`,
subset rows using `i`,
and manipulate columns with `j`,
grouped according to `by`.

`data.tables` are also data frames – functions that work with data frames therefore also work with `data.tables`.

Manipulate columns with `j`

EXTRACT

`dt[, c(2)]` – extract columns by number. Prefix column numbers with “_” to drop.

`dt[, .(b, c)]` – extract columns by name.

`dt[, b ~ c]` – extract columns by name.

SUMMARIZE

`dt[, .(x = sum(a))]` – create a `data.table` with new columns based on the summarized values of rows.

Summary functions like `mean()`, `median()`, `min()`, `max()`, etc. can be used to summarize rows.

COMPUTE COLUMNS*

`dt[, c := 1 + 2]` – compute a column based on an expression.

`dt[, a == 1, c := 1 + 2]` – compute a column based on an expression but only for a subset of rows.

`dt[, `:=` (c = 1, d = 2)]` – compute multiple columns based on separate expressions.

DELETE COLUMN

`dt[, c := NULL]` – delete a column.

Subset rows using `i`

`dt[1:2,]` – subset rows based on row numbers.

`dt[a > 5,]` – subset rows based on values in one or more columns.

`dt[a == 1,]` – subset rows based on values in one or more columns.

Group according to `by`

Chaining

`dt[, ...][, ...]` – perform a sequence of `data.table` operations by chaining multiple “[]”.

Functions for `data.tables`

REORDER

`setorder(dt, a, -b)` – reorder a `data.table` according to specified columns. Prefix column names with “-” for descending order.

* SET FUNCTIONS AND :=

`data.table`'s functions prefixed with “`set`” and the operator “`:=`” work without “`<-`” to alter data without making copies in memory. E.g., the more efficient “`setDT(df)`” is analogous to “`df <- as.data.table(df)`”.

LOGICAL OPERATORS TO USE IN i

`<` `<=` `is.na()` `%in%` `|` `%like%`
`>` `>=` `!is.na()` `!` `&` `%between%`

CONVERT COLUMN TYPE

`dt[, b := as.integer(b)]` – convert the type of a column using `as.integer()`, `as.numeric()`, `as.character()`, `as.Date()`, etc..

UNIQUE ROWS

a b → **a b**
 1 2 → 1 2
 2 2 → 2 2
unique(dt, by = c("a", "b")) – extract unique rows based on columns specified in “by”.
 Leave out “by” to use all columns.

BIND

a b + **a b** = **a b** **rbind(dt_a, dt_b)** – combine rows of two data.tables.
a b + **x y** = **a b x y** **cbind(dt_a, dt_b)** – combine columns of two data.tables.

Apply function to cols.

setkey(dt, a, b) – set keys to enable fast repeated lookup in specified columns using “dt[, value]” or for merging without specifying merging columns using “dt[, a[dt_b]]”.
setnames(dt, c("a", "b"), c("x", "y")) – rename

dt[, lapply(.SD, mean), .SDcols = c("a", "b")] – apply a function – e.g. `mean()`, `as.character()`, `which.max()` – to columns specified in `.SDcols` with `lapply()` and the `.SD` symbol. Also works with groups.
cols <- c("a")
dt[, paste0(cols, "m") := lapply(.SD, mean), .SDcols = cols] – apply a function to specified columns and assign the result with suffixed variable names to the original data.



data.table

Sequential rows

dt[, c:=1:N, by=b] – within groups, compute a column with sequential row IDs.
dt[, c:=shift(a, 1), by=b] – within groups, duplicate a column with rows *lagged* by specified amount.
dt[, c:=shift(a, 1, type="lead"), by=b] – within groups, duplicate a column with rows *leading* by specified amount.

RESHAPE TO WIDE FORMAT

dt[a ~ y] → **dt[a x a z b x b z]** **dcast(dt, id~y, value.var = c("a", "b"))**
 Reshape a data.table from long to wide format.
 A data.table.
 Formula with a LHS: ID columns containing IDs for multiple entries. And a RHS: columns with values to spread in column headers.
 Columns containing values to fill into cells.
dt[a ~ y] → **dt[a b c]** **lag & lead**
dt[a ~ y] → **dt[a NA b NA c NA]** **dt[, c:=1:N, by=b]** – within groups, compute a column with sequential row IDs.
dt[, c:=shift(a, 1), by=b] – within groups, duplicate a column with rows *lagged* by specified amount.
dt[, c:=shift(a, 1, type="lead"), by=b] – within groups, duplicate a column with rows *leading* by specified amount.

RESHAPE TO LONG FORMAT

dt[a ~ y] → **dt[a x a z b x b z]** **melt(dt, id.vars = c("id"), measure.vars = patterns("^a", "Ab"), variable.name = "y", value.name = c("a", "b"))**
 Reshape a data.table from wide to long format.
 A data.table.
 ID columns with IDs for multiple entries.
 Columns containing values to fill into cells (often in pattern form).
 Names of new columns for variables and values derived from old headers.
read & **write files**

dt A data.table.
id.vars ID columns with IDs for multiple entries.
measure.vars Columns containing values to fill into cells (often in pattern form).
variable.name, **value.name** Names of new columns for variables and values derived from old headers.

IMPORT
fread("file.csv") – read data from a flat file such as `.csv` or `.tsv` into R.

EXPORT
fwrite(dt, "file.csv") – write data to a flat file from R.

dt[a ~ b, on = .(id = id, date = date), roll = TRUE] – join data.tables on matching rows in id columns but only keep the most recent preceding match with the left data.table according to date columns. “roll = -Inf” reverses direction.

unique(dt, by = c("a", "b")) – extract unique rows based on columns specified in “by”.

unique(dt, by = c("a", "b")) – count the number of unique rows based on columns specified in “by”.

RENAME COLUMNS
a b → **x y** **setnames(dt, c("a", "b"), c("x", "y"))** – rename