### Text data and data.table

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## Data get larger over time

- ▶ We've been dealing with datasets below 50MB
- ▶ In reality sample size, number of variables and diversity of data type is much higher
- ► My own example:
  - household purchase record for retail clothing, 2007-2015:
     400MB
  - ▶ flickr.com user behavior and camera purchase: 1.2G
  - Yelp.com user reviews (Yelp academic dataset): 1.3G
  - grocery shopper's purchase records from a select set of retailers (2001-2007): ~4G only in the yogurt category
  - New York City yellow cab travel record in Jan-Jun, 2015: 11.5G
  - ► Airbnb data (still on-going): ~900G

## What we do today

- Data format: tabular/non-tabular, text/non-text
- ► Reading and writing from/to text data
- ► The *data.table* package:
  - speed
  - notation
  - examples
- Quick preview of feather

## **Tabular data**

#### Tabular data are tables

```
# explicitly set work directory
setwd("L:/Dropbox/Teaching/Programming for analytics/2022/lecture 8/")
    note: forward slash (/), backward slash will not (directly) work
# we've seen how a dataset can be imported
df <- read.csv("starwars.csv")</pre>
str(df)
## 'data.frame': 8 obs. of 7 variables:
## $ name : chr "Luke Skywalker" "Leia Skywalker" "Obi-Wan Kenobi" "Han Solo" ...
## $ gender : chr "male" "female" "male" "male" ...
## $ height : num 1.72 1.5 1.82 1.8 0.96 1.67 0.66 2.28
## $ weight : int 77 49 77 80 32 75 17 112
## $ jedi : chr "jedi" "no_jedi" "jedi" "no_jedi" ...
## $ species: chr "human" "human" "human" "human" ...
## $ weapon : chr "lightsaber" "blaster" "lightsaber" "blaster" ...
# stored in the form of data frame
df[1:6, 1:4]
##
             name gender height weight
## 1 Luke Skywalker
                    male 1.72
## 2 Leia Skywalker female 1.50
                                    49
## 3 Obi-Wan Kenobi male 1.82 77
      Han Solo male 1.80 80
## 4
## 5
         R2-D2 male 0.96 32
## 6
     C-3PO male 1.67
                                    75
```

## So we can find where things are

```
# info about luke?
df[df$name == "Luke Skywalker", ]

## name gender height weight jedi species weapon
## 1 Luke Skywalker male 1.72 77 jedi human lightsaber

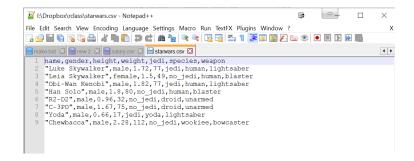
# height of the first 3 obs
df$height[1:3]

## [1] 1.72 1.50 1.82

# who's taller than 1.80?
as.character(df$name[df$height > 1.80])

## [1] "Obi-Wan Kenobi" "Chewbacca"
```

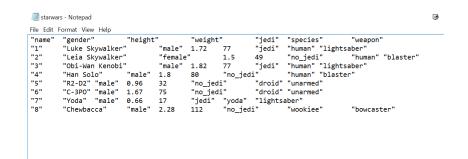
#### Raw data can be stored in csv formats



#### We can save a data frame as a tabular dataset

```
# We can save data frame to file using write.table
# here: write to tab-delimited text file
# note: "\\t" means "tab"
# note 2: we explicitly save the file at an *absolute* folder
write.table(df,
    "L:/Dropbox/Teaching/Programming_for_analytics/2022/lecture_8/starwars.txt",
    sep = "\t")
```

#### Text data: in this case tab-delimited



#### Text data do not have to be tab- or comma- delimited

```
# another common approach is space delim
write.table(df, "starwars_space.txt", sep = " ")

# or, can even use weird separators like "$"
write.table(df, "starwars_dollar.txt", sep = "$")

# read it back
df3 <- read.table("starwars_dollar.txt", sep = "$")

# still the same data
df3[1:3, 1:3]

## name gender height
## 1 Luke Skywalker male 1.72
## 2 Leia Skywalker female 1.50
## 3 Obj-Wan Kenobi male 1.82</pre>
```

#### Tabular data can be stored in fixed-width

```
# define file location
loc <- "starwars_fwidth.txt"

# define width parameter
w <- c(14, 6, 4, 3, 7, 7, 10)

# write df into fixed width with package 'gdata'
library('gdata')
write.fwf(df, file = loc, width = w)</pre>
```

# Fix width is easier on human eyes (but is more difficult to read or write)

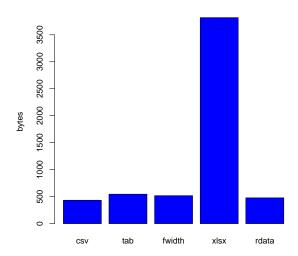
```
starwars fwidth - Notepad
File Edit Format View Help
name gender height weight jedi species weapon
Luke Skywalker male 1.72 77 jedi
                                       human
                                              lightsaber
Leia Skywalker female 1.50
                           49 no_jedi human
                                              blaster
Obi-Wan Kenobi male
                     1.82 77 jedi
                                              lightsaber
                                       human
Han Solo
              male
                     1.80 80 no jedi human
                                              blaster
R2-D2
              male
                     0.96 32 no jedi droid
                                              unarmed
C-3P0
              male
                     1.67 75 no iedi droid
                                              unarmed
Yoda
              male
                     0.66 17 jedi
                                      yoda
                                              lightsaber
              male
                     2.28 112 no jedi wookiee bowcaster
Chewbacca
```

## Now compare size of different files

```
# read file info using file.info()
info.list <- list(
    file.info("starwars.csv"),
    file.info("starwars.txt"),
    file.info("starwars.fwidth.txt"),
    file.info("starwars.xlsx"),
    file.info("starwars.rdata")
)

# what are the sizes for the files?

fs <- data.frame(
    fileformat = c("csv", "tab", "fwidth", "xlsx", "rdata"),
    size = c(info.list[[1]]$size,
        info.list[[2]]$size,
        info.list[[3]]$size,
        info.list[[4]]$size,
        info.list[[4]]$size,
        info.list[[5]]$size
    )
)</pre>
```



## Size compared

- Everything similar size except for the Excel file
- Minimal file size for the information
  - the csv file has 418 characters including delimitor, so 418 bytes
  - ▶ adding 8 new lines "\n" which takes additional 8\*2=16 bytes
  - final file size is 434 bytes, no waste at all
- Other files
  - ▶ tab data is 546 bytes because tab "\t" takes 2 bytes
  - Rdata is 478 bytes; quite efficient
  - Excel file has exactly the same information but is 8.8 times the size of csv

# Non-tabular text data example: EPS and BMP

# Export a figure to Encapsulated PostScript (EPS)

```
# load graphic devices library
library('grDevices')

# create an EPS file
postscript("example_bar.eps",
    width = 480, height = 480, horizontal = F)
barplot(fs\size, names.arg = fs\sileformat,
    ylab = "bytes", col = "blue")
dev.off()

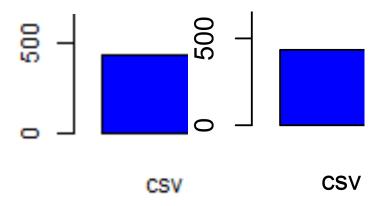
## pdf
## pdf
## 2
```

## Export the same figure to Bit MaP (BMP)

```
# create a BMP file
bmp("example_bar.bmp",
    width = 480, height = 480)
barplot(fs$size, names.arg = fs$fileformat,
    ylab = "bytes", col = "blue")
dev.off()

## pdf
## 2
```

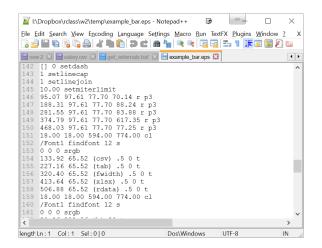
## BMP = 227kb; $EPS = 7kb^1$



which one is BMP and which one is EPS?

<sup>&</sup>lt;sup>1</sup>To be exact, you're seeing a **rendered** EPS

# Why? BMP stores RGB for each pixel; EPS stores instructions on how to draw the figure



#### So far

- Data is stored in files
  - takes space to store
  - takes time to read it
  - lots of data => file format matters A LOT!
- Text is a universal and efficient way to store data
  - can store tables (tabular data)
  - can store more general information (non-tabular data)
- Two examples where text data is vastly superior
  - For tabular: CSV (and other text) data is much smaller than Excel data
  - For graphics: EPS graphs is much smaller than BMP graphs

# data.table()

# data.table()

- Speed
- Notation
- Examples

#### Standard read.table

```
# library
library('data.table')
# data.frame way (bb for billboard, file size ~350MB)
t0 <- proc.time()
df_bb <- read.table("billboard_full.txt", sep = "\t",  # separator is tab</pre>
       quote = "", # will read quote literally
       header = T, # will read first line as header
       stringsAsFactors = F.
       fill = TRUE) # fill: uneven columns will be filled
# check time
t1 <- proc.time()
t1 - t0
## user system elapsed
## 11.31 0.39 11.72
# check data
head(df_bb, n = 3)
        metro country countrycode starttime endtime artist rank
## 1 Melbourne Australia
                               au 1233489600 1234094400
                                                          Radiohead
## 2 Melbourne Australia au 1233489600 1234094400 Kings of Leon
## 3 Melbourne Australia
                               au 1233489600 1234094400
                                                           Coldplay
    playcount
                                           mbid
## 1
      NA a74b1b7f-71a5-4011-9441-d0b5e4122711
## 2 NA 6ffb8ea9-2370-44d8-b678-e9237bbd347b
## 3
          NA cc197bad-dc9c-440d-a5b5-d52ba2e14234
# remove df_bb because we won't use it later
rm(df_bb)
```

# Compared: fread()

```
# data.table way
t0 <- proc.time()
dt bb <- fread("billboard full.txt", sep = "\t", header = T, stringsAsFactors = F)
 ## Warning in fread("billboard full.txt", sep = "\t", header = T, stringsAsFactors = F): Found
  and resolved improper quoting out-of-sample. First healed line 699661: «Changsha China cn
  1269172800 1269777600 "Weird Al" Yankovic 58 ». If the fields are not quoted (e.g. field
       separator does not appear within any field), try quote="" to avoid this warning.
# check time
t1 <- proc.time()
t1 - t0 # FASTER!
## user system elapsed
## 1.45 0.08 0.96
# check data
head(dt_bb, n = 3)
         metro country countrycode starttime
                                                 endtime
                                                                artist rank
## 1: Melbourne Australia au 1233489600 1234094400
                                                             Radiohead 1
## 2: Melbourne Australia
                               au 1233489600 1234094400 Kings of Leon
## 3: Melbourne Australia
                                                              Coldplay
                               au 1233489600 1234094400
     playcount
##
                                             mbid
## 1 .
            NA a74b1b7f-71a5-4011-9441-d0b5e4122711
## 2:
            NA 6ffb8ea9-2370-44d8-b678-e9237bbd347b
## 3:
            NA cc197bad-dc9c-440d-a5b5-d52ba2e14234
# you'll notice the small colon at every row,
# indicating that this is data table
```

### Data table as data frames

## Data tables are recognized as data frames

```
# let's do it on a smaller scale (your assignment 1 data)
url <- 'https://dl.dropboxusercontent.com/s/q6qzbfa1tdcqv6v/belgium_atm.csv'
dt_atm <- fread(url, stringsAsFactors = F)</pre>
head(dt atm)
     population numATMs ATMwithdr withdrvalue unemprate numbranches
## 1 .
           3722
                         .25542593 79.13402557 0.0728676
                                                              0.500
## 2.
           7006
                     2 1.837865114 102.6663437 0.0226948
                                                              0.500
## 3:
       4234
                     0 missing missing 0.0273973
                                                             0.125
         6229
                           missing missing 0.0244020
                                                              0.750
## 4:
                     1 6062539816 98 93833923 0 0284383
## 5:
      10303
                                                              0.375
## 6:
         7424
                                      missing 0.0373114
                                                              0.875
                           missing
    note the colon ':' in each row, this indicates data table
# it is recognized as a data frame
is.data.frame(dt_atm)
## [1] TRUE
```

### Some data frame notations work with data table

```
# subset by rows
dt atm[1:3, ]
     population numATMs ATMwithdr withdrvalue unemprate numbranches
## 1:
           3722
                     1 .25542593 79.13402557 0.0728676
                                                              0.500
## 2:
         7006
                     2 1.837865114 102.6663437 0.0226948
                                                             0.500
## 3 .
      4234
                     0 missing missing 0.0273973
                                                             0.125
# refer to columns
head(dt_atm$numATMs)
## [1] 1 2 0 0 1 0
# also this can refer to columns (as elements in a list)
head(dt_atm[["numATMs"]])
## [1] 1 2 0 0 1 0
# so basically row notations work, (some) list notations work
```

#### Difference 1: column notations

```
# same as in a data frame (updated data table supports this)
dt_atm[, c("numATMs", "population")]
       numATMs population
##
   1:
             1
                    3722
## 2:
                   7006
        0 4234
        0 6229
               10303
## 655:
                    601
                 1028
## 656:
                 2033
## 657:
## 658 .
                   15521
## 659:
                    5941
# single-bracket list subsetting does NOT work
dt_atm[c(1, 2)]
     population numATMs ATMwithdr withdrvalue unemprate numbranches
##
## 1:
           3722
                     1 .25542593 79.13402557 0.0728676
                                                              0.5
## 2:
           7006
                     2 1.837865114 102.6663437 0.0226948
                                                              0.5
# and I'll explain why
```

# Difference 2: lots of operations now work within the data table object

```
# delete the population column
dt_atm[, population := NULL]
                               # delete column
# construct a new column by another column
dt_atm[,
        avg_withdrawal := mean(as.numeric(ATMwithdr), na.rm = T),
        bv = numATMs1
 ## Warning in '[.data.table'(dt atm, , ':='(avg withdrawal, mean(as.numeric(ATMwithdr), : NAs
                                     introduced by coercion
# print
dt_atm
                 ATMwithdr withdrvalue unemprate numbranches avg withdrawal
    1 :
                  .25542593 79.13402557 0.0728676
                                                        0.500
                                                                   0.7576123
##
##
    2.
             2 1.837865114 102.6663437 0.0226948
                                                        0.500
                                                                   1.0025904
    3:
                               missing 0.0273973
                                                                         NaN
##
                   missing
                                                       0.125
                   missing
                             missing 0.0244020
                                                       0.750
                                                                         NaN
##
             1 6062539816 98 93833923 0 0284383
                                                        0.375
                                                                   0.7576123
    ---
## 655:
                               missing 0.0217658
                                                        0.125
                                                                         NaN
                   missing
## 656:
             Ω
                   missing
                               missing 0.0217658
                                                        0.250
                                                                         NaN
## 657:
                               missing 0.0217658
                                                        0.500
             0
                   missing
                                                                         NaN
## 658:
                .6984899044 110.1268387 0.0231947
                                                       0.875
                                                                   1.0025904
## 659:
                    missing
                               missing 0.0231947
                                                        0.875
                                                                         NaN
# and I'll explain these in more detail
```

Notation: DT[i, j, by]

# DT[i, j, by]

- ► For a data table, say 'DT':
  - some standard data frame notations work
  - some do not because data.table notation overrides data frame notations
- Specifically, DT[i, j, by]
  - i: which rows do you want?
  - j: what do you want to do with the rows?
  - by: do this by which index?

```
# read ATM data again
url <- 'https://dl.dropboxusercontent.com/s/q6qzbfa1tdcqv6v/belgium_atm.csv'
dt atm <- fread(url, stringsAsFactors = F)
# DT[i] gets the rows
dt atm[1:3] # row 1-3
##
     population numATMs
                        ATMwithdr withdrvalue unemprate numbranches
## 1:
           3722
                          .25542593 79.13402557 0.0728676
                                                               0.500
## 2:
           7006
                      2 1.837865114 102.6663437 0.0226948
                                                               0.500
## 3:
           4234
                            missing
                                       missing 0.0273973
                                                               0.125
                      Ω
dt_atm[c(4, 1, 8)]
                  # row 4, 1, 8, in order (also note row num changed)
     population numATMs
                          ATMwithdr withdrvalue unemprate numbranches
## 1 .
           6229
                      Ω
                         missing
                                       missing 0.0244020
                                                               0.750
## 2.
           3722
                          .25542593 79.13402557 0.0728676
                                                               0.500
                      1
## 3:
           7129
                      1 .8399091363 89.14505768 0.0352083
                                                               0.625
# or gets rows by condition
dt atm[numATMs <= 2 & population <= 5000]
       population numATMs ATMwithdr withdrvalue unemprate numbranches
##
             3722
                        1 .25542593 79.13402557 0.0728676
##
    1:
                                                               0.500
    2.
             4234
                            missing
                                       missing 0.0273973
                                                               0.125
##
##
    3.
             2868
                        0 missing
                                       missing 0.0355649
                                                               0.375
   4:
            4105
                        0 missing
                                       missing 0.0326871
                                                               0.250
##
                                       missing 0.0291726
##
    5.
             4992
                        0 missing
                                                               0.375
##
## 245:
             3529
                                                               0.500
                            missing
                                       missing 0.0208911
             2862
                                       missing 0.0217658
                                                               0.500
## 246:
                        0 missing
## 247:
             601
                          missing
                                       missing 0.0217658
                                                               0.125
## 248:
             1028
                            missing
                                       missing 0.0217658
                                                               0.250
## 249:
             2033
                            missing
                                       missing 0.0217658
                                                               0.500
    note that we don't need to say dt atm$numATM <= 2 as with data frames
```

# DT[i, j]

```
# argument j: what do you want to do with those rows?
# keep all columns
dt_atm[1:3, ]
     population numATMs ATMwithdr withdrvalue unemprate numbranches
## 1 .
           3722
                         .25542593 79.13402557 0.0728676
                                                             0.500
## 2:
     7006
                     2 1.837865114 102.6663437 0.0226948
                                                             0.500
## 3:
           4234
                                      missing 0.0273973
                                                             0.125
                           missing
# keep variable population (now a vector)
dt_atm[1:3, population]
## [1] 3722 7006 4234
# I can still use data frame notation (only in recent versions)
dt_atm[1:3, "population"]
     population
##
## 1:
           3722
## 2.
          7006
## 3:
           4234
```

# DT[i, j]

```
# can run a function instead of just taking columns
# example: average number of ATMs in smaller towns
dt_atm[population <= 5000, mean(numATMs)]
## [1] 0.09638554
# data frame equivalent will be
mean(dt_atm[dt_atm$population <= 5000, ]$numATMs)
## [1] 0.09638554</pre>
```

## Special operator := to create new columns

```
# create a variable (in the 'j' part)
   ':=' means assignment by reference
dt_atm[, pop := population / 1000]
                                      # population in thousands
# create multiple variable
     turns population into thousands, turns two variables into numeric, and deletes pop
dt_atm[, ':='(population = population / 1000,
               ATMwithdr = as.numeric(ATMwithdr).
               withdrvalue = as.numeric(withdrvalue),
               pop = NULL)]
head(dt atm, n = 3) # and co (country code)
     population numATMs ATMwithdr withdrvalue unemprate numbranches
          3.722
                     1 0.2554259 79.13403 0.0728676
                                                            0.500
## 1:
## 2.
        7.006 2 1.8378651 102.66634 0.0226948
                                                            0.500
## 3:
         4.234
                              NA
                                        NA 0.0273973
                                                            0.125
```

# Special character .() to group function calls

# DT[i, j, by]

```
# at this point you can see the full notation DT[i, j, by]
    example: what is the average withdraw value by number of ATMs?
    added condition: for towns with population > 2000
dt_atm[population >= 2,
       .(ATMwithdr = mean(ATMwithdr), withdrvalue = mean(withdrvalue)),
       numATMs7
     numATMs ATMwithdr withdrvalue
## 1:
           1 0.7505181 101.05752
## 2.
           2 1.0025904
                       100.37617
## 3:
                    NΑ
                               NA
## 4:
      4 1.0827598 96.25565
## 5:
        3 1.1169177 99.25264
        5 1.1931812 98.15437
## 6:
## 7:
           6 0.7416071 92.73267
```

## Example 2: Billboard data

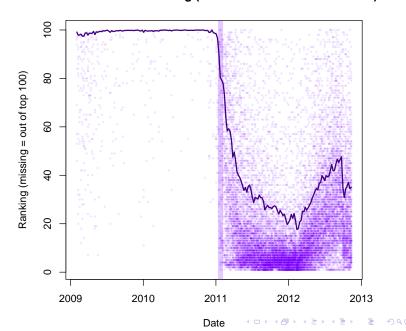
```
# now work with larger (still small) dataset on Billboard ranking
head(dt bb, n = 2) # mbid is universal albumn ID (standard in music industry)
         metro country countrycode starttime endtime
                                                                artist rank
## 1: Melbourne Australia au 1233489600 1234094400
                                                             Radiohead 1
## 2: Melbourne Australia au 1233489600 1234094400 Kings of Leon
## playcount
                                             mbid
           NA a74b1b7f-71a5-4011-9441-d0b5e4122711
## 1:
## 2:
           NA 6ffb8ea9-2370-44d8-b678-e9237bbd347b
t0 <- proc.time()
# convert starttime to date
    numeric means seconds since 1970-1-1 07:00:00 (I will cover dates next week)
dt bb$date <- as.Date(as.POSIXct(dt_bb$starttime, origin = "1970-1-1"))
# let's find subset artists who
# 1) ranked top 3
# 2) between September 2009 and August 2011
# 3) focus on the US market
# first data frame way (df bb would take the same amt of time)
dt sub <- subset(dt bb, date >= "2009-09-01" & date <= "2011-08-31" &
       rank <= 3 & countrycode == "us")
freq.df <- aggregate(x = rank ~ artist, data = dt sub, FUN = length)</pre>
names(freg.df)[2] <- "frequency"
# check time
t1 <- proc.time()
\pm 1 - \pm 0
     user system elapsed
## 0.15 0.03 0.18
```

## Example 2: Billboard data

## Example 2 con'd: Let's find Adele's average ranking

```
# not every artist is on top 100 in every week
    assume that any time Adele falls out of billboard she is ranked at 100
    (probably not accurate...)
# total number of metropolitan areas (where billboard is announced) is 234
num.metro <- length(unique(dt_bb$metro))</pre>
# define a function to calculate average ranking, counting each missing at 100
find_average_rank <- function(rank) {</pre>
        freq.not100 <- num.metro - length(rank) # count metros that do NOT have Adele on bb
        (sum(rank) + mean(freq.not100)*100) / num.metro
                                                          # assume that missing values are 100
                # so total is sum rank + missing * 100
# use key to sort data
setkey(dt_bb, date)
# subset and aggregate at the same time
adele_ranking <- dt_bb[artist == "Adele", .(rank = find_average_rank(rank)), date]
# plot
plot(dt bb[artist == "Adele", date], dt bb[artist == "Adele", rank],
        col = rgb(0.5, 0, 1, 0.1), pch = 16, cex = 0.5, # cex is marker size
        main = "Adele's billboard ranking (vertical = release of album '21')",
        xlab = "Date", vlab = "Ranking (missing = out of top 100)"
# add release date for '21'
abline(v = as.Date("2011-01-24"), col = rgb(0.5, 0, 1, .2), lwd = 8)
# add average ranking we calculated
points(adele ranking$date, adele ranking$rank, type = '1', col = rgb(0.25, 0, 0.5, 1), lwd = 2)
```

#### Adele's billboard ranking (vertical = release of album '21')



## **Feather**

#### **Feather**

- ▶ Data.table still runs slow in large data applications, especially when a lot of data are read and written to/from the hard drive
- ► Feather aims to provide a solution by using a different data format, *.feather*, which is aimed to facilitate short-term read/write tasks
  - but CAUTION: not for long-term data storage

```
# load the package
library(feather)
## Warning: replacing previous import 'lifecycle::last_warnings' by 'rlang::last_warnings' when
                                        loading 'hms'
# set work directory to an SSD
setwd("D:/")
# generate a data frame (courtesy to Hadley Wickham)
x <- runif(1e6) # 1 million rows
x[sample(1e6, 1e5)] <- NA # 10% NAs
df <- as.data.frame(replicate(10, x))</pre>
system.time(write_feather(df, 'test.feather'))
##
     user system elapsed
##
     0.01 0.04 0.04
system.time(write.csv(df, 'test.csv'))
     user system elapsed
##
    28.32
             0.26 28.62
##
```

```
# test read time
system.time(read_feather('D:/test.feather'))
## user system elapsed
## 0.03 0.02 0.04
system.time(fread('D:/test.csv'))
## user system elapsed
## 0.30 0.42 0.08
system.time(read.table('D:/test.csv'))
## user system elapsed
## 2.94 0.12 3.06
# HUGE speed gain in both write and read!
```

### Conclusion

- ► File format
  - read.table and write.table
  - csv, tab-demited data and fixed format data
  - text files have huge space/compatibility advantages (beyond tabular data)
- DT[i, j, by]
  - i: which rows do you want?
  - ▶ j: want them for what (which function to apply)?²
  - by: do this along which index variable(s)?
- ► Feather seems to be the future for temporary read and write data

<sup>&</sup>lt;sup>2</sup>Recall that anything that happens is a function call

