Data Management in



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the life-changing magic of tidying up

the Japanese art of decluttering and organizing

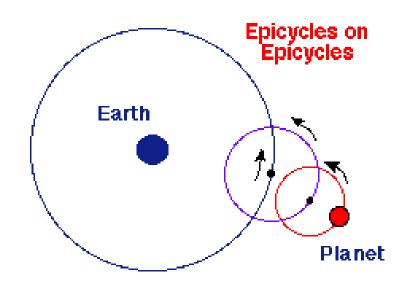
marie kondo

Names for the early steps of "Data Science"

- Data munging
- Data wrangling
- Web scraping
- Data pulling
- Pre-processing
- Data cleansing
- Data harmonising
- Data cleaning
- <u>https://www.datacamp.com/community/blog/an-introduction-to-cleaning-data-in-r</u>

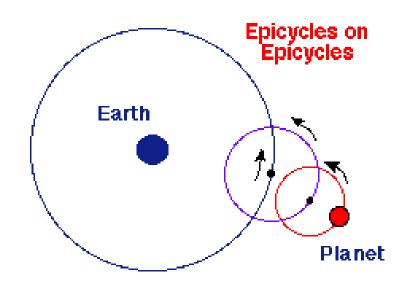
Epicycle of Analysis (Art of Data Science, Peng)

- There are 5 core activities of data analysis:
 - Stating and refining the question
 - Exploring the data (EDA)
 - Building formal statistical models
 - Interpreting the results
 - Communicating the results



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EDA

 "There are no routine statistical questions, only questionable statistical routines."—Sir David Cox

 "Far better an approximate answer to the right question, which is often vague, than an exact answer to the wrong question, which can always be made precise."—John Tukey

Exploratory Data Analysis

- (From Art of Data Science) Formulate your question:
 - Read in your data
 - Look at the top and the bottom of your data
 - Check your "n"s
 - Validate (with at least one external data source)
 - Make a plot
 - Try the *easy* solution first
 - Follow up...

"Mise en place"!

```
rm(list=ls())
getwd()
setwd("<location of your dataset>")
```

• .Rpofile: file for customising start-up

OpenRefine



https://github.com/OpenRefine/OpenRefine/wiki/Documentation-For-Users

Validation

- Check basic characteristics
- Validate with another (external) data source if possible
- Make a simple plot (boxplot, (paired) scatter plot,...)

Reading data from GitHub

```
library(readr)
```

```
conception <-
read_csv("https://raw.githubuser
content.com/n8thangreen/ONS/mast
er/conception-under18.csv")
View(conception)</pre>
```

Validation type	How it works	Example
Check digit	The last couple of digits can be used as a 'check sum' that can detect if errors have occurred	Bar code readers in shops
Type check	Checks the data is in the right format	Numbers in currency cell must be a monetary value with two decimal points
Length check	Checks the data is an acceptable length	A PIN for online banking needs to be four (or six) characters long
Lookup table	Checks that the value provided matches an item in a set list	A limited set of values, such as the seven days of the week
Presence check	Checks that data has been entered into a field	In most databases a key field cannot be left blank
Range check	Checks that a value falls within the specified range	An online gift certificate purchase must be more than or equal to £5 but less than or equal to £50
Spell check	Looks up words in a dictionary	A search engine often recommends a correct spelling if a word is spelt wrong
Input mask	Checks that data has been entered with the correct amount of characters and/or numbers	National insurance numbers need to be in the format: YY XX XX XX Y, where Y = letter and X = number
Duplicate	Checks that a value has not been repeated	A primary key value can only be entered once
	http://www.	bbc.co.uk/education/guides/

Variables types and indexing

- numeric Numeric data (approximations of the real numbers, ℝ)
- integer Integer data (whole numbers, Z)
- factor Categorical data (simple classifications, like gender)
- ordered Ordinal data (ordered classifications, like educational level)
- character Character data (strings)
- raw Binary data

```
typeof() # what is it?
length() # how long is it? What about two
dimensional objects?
attributes() # does it have any metadata?
# Example
x <- "dataset"
typeof(x)
attributes(x)
y < -1:10
typeof (y)
length(y)
attributes(y)
z < -c(1L, 2L, 3L)
typeof(z)
```

Useful functions

- head() see first 6 rows
- tail() see last 6 rows
- dim() see dimensions
- nrow() number of rows
- ncol() number of columns
- str() structure of each column
- names() will list the names attribute for a data frame (or any object really), which gives the column names.
- table()
- unique(); duplicated()

String normalisation

```
library(stringr)
str trim(" hello world ")
## [1] "hello world"
str trim(" hello world ", side = "left")
## [1] "hello world "
str trim(" hello world ", side = "right")
## [1] " hello world"
toupper ("Hello world")
## [1] "HELLO WORLD"
tolower ("Hello World")
## [1] "hello world"
```

String matching (regular expressions)

```
gender <- c("M", "male ", "Female",</pre>
"fem.")
grepl("m", gender)
## [1] FALSE TRUE TRUE TRUE
grep("m", gender)
## [1] 2 3 4
grepl("m", gender, ignore.case = TRUE)
## [1] TRUE TRUE TRUE TRUE
grepl("m", tolower(gender))
## [1] TRUE TRUE TRUE TRUE
grepl("^m", gender, ignore.case = TRUE) ##
[1] TRUE TRUE FALSE FALSE
```

Running example

```
library("stringr", lib.loc="~/R/win-library/3.2")
conception$Name
conception$Name <- str_trim(conception$Name)
conception$Name <- gsub(pattern = " UA", replacement
= "", conception$Name)</pre>
```

Consistent data

Missing values

```
age <- c(23, 16, NA)
mean(age)
## [1] NA
mean(age, na.rm = TRUE)
## [1] 19.5</pre>
```

Checks

- is.na()
- is.nan()
- na.omit()
- complete.cases(x)

Running example

```
apply(conception, 2, function(x) any(is.na(x)))
data$Name <- toupper(conception$Name)
chartr("AIai", "@!@!", conception$Name)</pre>
```

Outliers

What's realistic? Permissible (integrity)?
 (Outliers: An Evaluation of Methodologies, Dhiren Ghosh, Andrew Vogt, https://www.amstat.org/sections/srms/proceedings/y2012/files/304068 72402.pdf)

What to do?

- Imputation: substitution mean, logistic regression prediction, matching
- Truncation, removal, Winsorising, scaling, judgement

Subsetting

```
X[1]
x[-c(1, 5)]
x[c(TRUE, TRUE, FALSE, FALSE)]
x[which(x > 3)]
z[c("a", "d")]
df <- data.frame(x=1:3, y=3:1, z=letters[1:3])
df[df$x == 2, ]
```

The apply () family

- lapply()
 - Loop over a list and evaluate a function on each element
- sapply()
 - Same as lapply but try to simplify the result
- apply()
 - Apply a function over the margins of an array
- tapply()
 - Apply a function over subsets of a vector
- mapply()
 - Multivariate version of lapply

Examples

```
x < -matrix(rnorm(200), 20, 10)
apply(x, 2, mean)
x < - list(a = 1:5, b = rnorm(10))
lapply(x, mean)
sapply(x, mean)
x <- c(rnorm(10), runif(10), rnorm(10, 1))
f \leftarrow gl(3, 10)
tapply(x, f, mean)
rowSums = apply(x, 1, sum)
rowMeans = apply(x, 1, mean)
colSums = apply(x, 2, sum)
colMeans = apply(x, 2, mean)
```

"Tidy" data

 Happy families are all alike; every unhappy family is unhappy in its own way [Leo Tolstoy]

- A dataset is said to be tidy if it satisfies the following conditions
 - observations are in rows
 - variables are in columns
 - contained in a single dataset.

TB incidence data

```
tb <- read.csv(file =
"http://stat405.had.co.nz/data/tb.csv",
header = TRUE, stringsAsFactors = FALSE)</pre>
```

• Except for iso2 and year, the rest of the columns headers are actually values of a lurking variable, in fact combination of two lurking variables, gender and age.)

Reshape(2)

Create categorical values

```
data$zipGroups <-
cut(data$zipCode,breaks=quantile(data$zipCode))
install.packages("Hmisc")
library(Hmisc)

data$zipGroups <- cut2(data$zipCode, g=4)
table(data$zipGroups)</pre>
```

Transformations

absolute value

square root

```
sqrt(x)
```

• ceiling(3.4) is 4

```
ceiling(x)
```

• floor(3.4) is 3 floor(x)

```
rounding
```

```
round(x, digits=n)
```

• signif(3.475, digits=2) is 3.5

```
signif(x, digits=n)
```

natural logarithm

```
log(x)
```

exponentiating x

```
exp(x)
```

Sorting

```
data <- data.frame( "column1"=sample(1:5),</pre>
"column2"=sample(6:10), "column3"=sample(11:15))
data <- data[sample(1:5),]
data$column2[c(1,3)]<-NA
sort(data$column1, decreasing=FALSE)
sort(data$column2, decreasing=FALSE, na.last=TRUE)
data[order(data$column1),]
```

Running example

```
conception <- conception[order(conception$Name),]</pre>
conception <- within (conception, new <- `Number of
Conceptions `*2)
conception <- conception[order(conception</pre>
$`Conception rate per 1,000 women in age group`),]
conception <- conception[order(conception</pre>
$`Conception rate per 1,000 women in age group`,
decreasing = T),]
```

Merging/Joins

```
df1 = data.frame(id=sample(1:10), x=rnorm(10))
df2 = data.frame(id=sample(1:10), x=rnorm(10))
df3 = data.frame(id=sample(1:10), x=rnorm(10))
dfs = list(df1, df2, df3)
plyr::join_all(dfs)
```

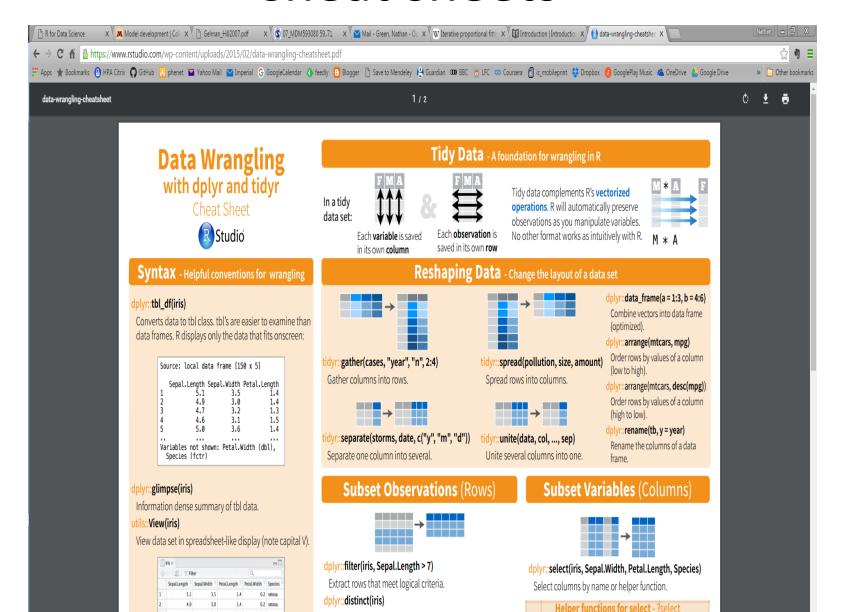
Melt and cast

```
head (mtcars)
install.packages("reshape2")
library(reshape2)
mtcars$carname <- rownames(mtcars)</pre>
carMelt <- melt( mtcars, id=c("carname", "gear",</pre>
"cyl"), measure.vars=c("mpg", "hp"))
cylData <- dcast(carMelt, cyl ~ variable)</pre>
cylData <- dcast(carMelt, cyl ~ variable, mean)</pre>
```

Running example

```
head(reshape2::melt(conception,id.vars="Name"))
```

Cheat sheets



RStudio



- Loads of really useful features, that are being added-to all the time:
 - Keyboard short cuts
 - Drop-down menus
 - Data sorting, pop-out windows
 - Linting, static code analysis
 - Searches: files, functions, text

— ...

Thanks