R: The tidyverse and beyond

1-day workshop
R-eproducible workflows
Practical session





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A: Is too much choice good or bad?

Blue Horizon SW 6497	Sky High SW 6504	Snowdrop SW 6511	Ski Slope SW 6518	Rarified Air SW 6525
Byte Blue SW 6498	Atmospheric SW 6505	Balmy SW 6512	Hinting Blue SW 6519	Icelandic SW 6526
Stream SW 6499	Vast Sky SW 6506	Take Five SW 6513	Honest Blue SW 6520	Blissful Blue SW 6527
Open Seas SW 6500	Resolute Blue SW 6507	Respite SW 6514	Notable Hue SW 6521	Cosmos SW 6528
Manitou Blue SW 6501	Secure Blue SW 6508	Leisure Blue SW 6515	Sporty Blue SW 6522	Scanda SW 6529
Loch Blue SW 6502	Georgian Bay SW 6509	Down Pour SW 6516	Denim SW 6523	Revel Blue SW 6530
Bosporus SW 6503	Loyal Blue SW 6510	Regatta SW 6517	Cammodore SW 6524	Indigo SW 6531

Inconsistent function names, inconsistent syntax

- R is a very versatile language
 - Sometimes it can be too versatile
 - Do you want to use ...

```
row.names or rownames rowSums or rowsum
Sys.time, system.time
```

- Should it be written as...

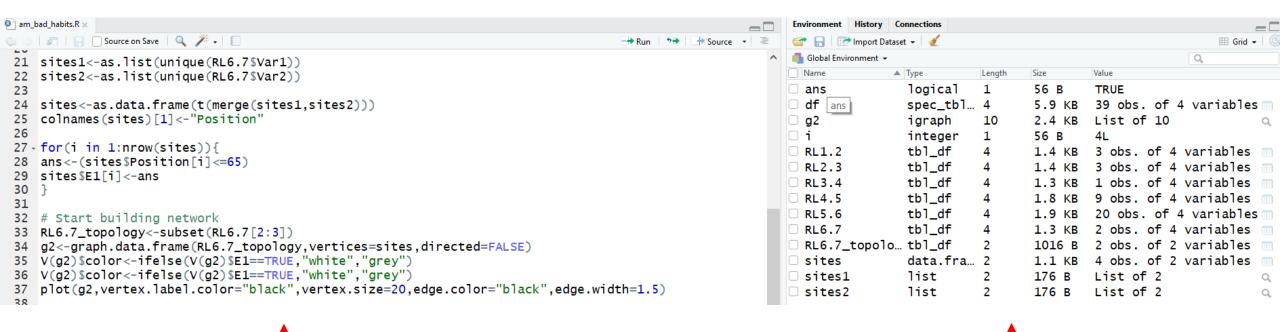
```
newobject or new.Object
x = 5 or x <- 5
mapping=aes(x,y) or mapping = aes(x, y)</pre>
```

Variable selection

```
summary(starwars$name)
summary(starwars$"name")
summary(starwars["name"])
summary(starwars[ , "name"])
summary(starwars[1])
summary(starwars[ , 1])
summary(starwars[[1]])
```

- Open the script am too much.choice.R

Motivation to move on from poorly written code



Lack of annotation
Poor naming conventions
Poor readability
Spacing absent

- Open the script am bad habits.R

Cluttered environment Intermediate objects

Writing clearer code

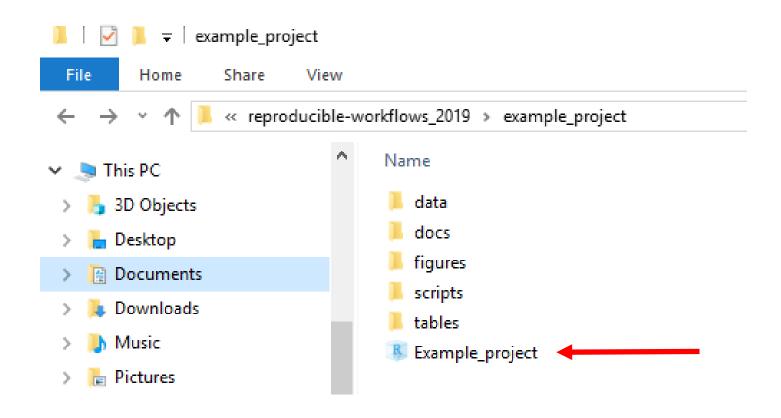
- Annotation
- Object names
 - should use only lowercase letters, numbers, and " $_$ "
- Spacing
 - Put a space before and after =
 - Put a space after a ,
 - Operators should be surrounded by spaces e.g. ==, <-, +
- For a more complete list visit
 - http://style.tidyverse.org/syntax.html

- Open the script am good habits.R

Navigating RStudio – some useful tips

- Open the script am_rstudio_ide_tricks.R

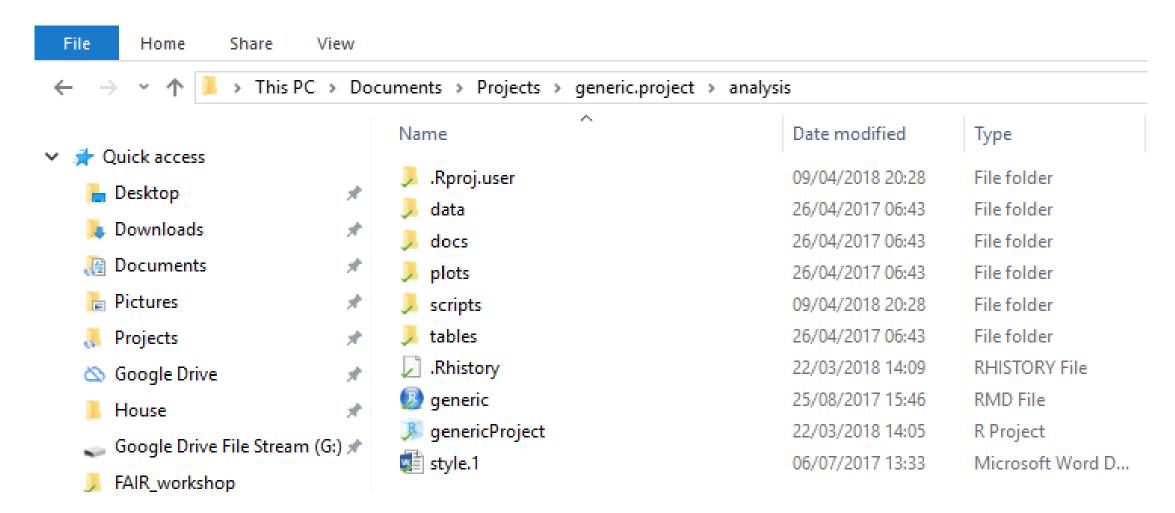
B: R-projects - everything in its right place



- Open the scripts 01_eg_clean_data.R, 02_eg_figures.R and 03_eg_analysis.R

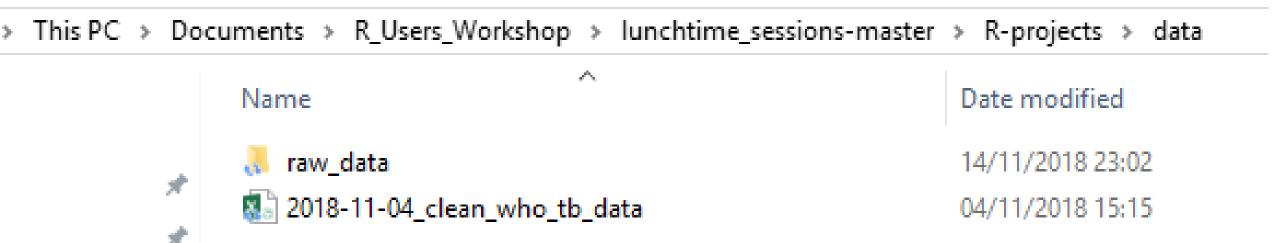
Define a generic project structure

- STEP 1: Give your research projects a shared structure



Give your files informative names

- STEP 2: Include metadata in the file names



Come back to what you know

- STEP 3: Make you file names machine readable, human readable and work with default ordering

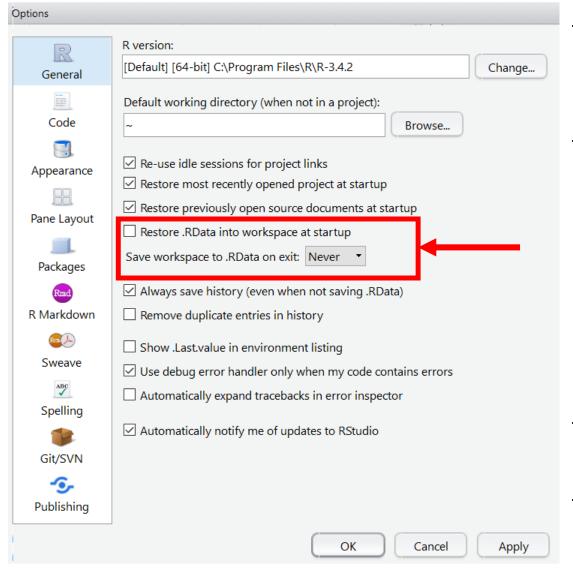
NO

- Epistatic_change
- Epistatic_change_match_discovery
- Epistatic_change_match_discovery_fig_2_point_1
- Epistatic_change_v2
- 😰 epistatic_codon _change_tracking
- Epistatic_connection_network
- Meatmap_for_epistatic_syn
- Meatmap1_for_epi_site_co-change
- Meatmap2_for_epi_fdr_adjusted_p-value
- Meatmap2_for_epi_p-value

Yes

Doc	uments > Projects > 18.04.27-WP3_Feeding_Trial	analysis > scripts
	Name	Date modified
_	01_data_import_and_tidying_master_file	02/10/2018 18:51
*	02_data_import_and_tidying_nutritics_grouped	19/10/2018 19:47
*	03_figures	17/10/2018 16:40
*	04_tables	22/05/2018 12:26
*	05_study_overview	19/10/2018 23:06
яř	functions	13/05/2018 23:13

Other points to note



- You might consider your environment as "real"
- If you continue to use R, it is better for you to consider your R scripts as "real", as these should recreate the environment

- You may suffer short term pain
- This will prevent long term agony