

Basic R Programming (and how to talk to a computer)

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Today plan

Overview of R (~10 mins)

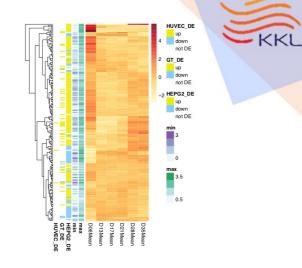
Hands-on with R programming (~20 mins)

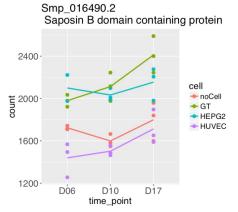
Tomorrow

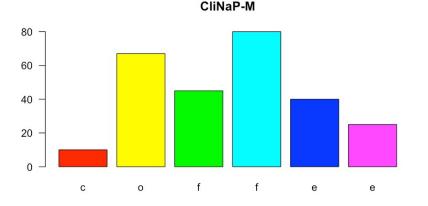
Final point before NMR data analysis session (~5 mins)

What is R, and what is it for?

- A language for talking to computer
- Data exploration and visualisation
- Can produce publication-quality plots
- Can perform mathematical and statistical analysis
- Use command line
 - Allow better reproducibility / pipeline development
 - Fast if re-analysis is required
- Freeware + >1000s community developers
- Work on Linux, Windows, MacOS

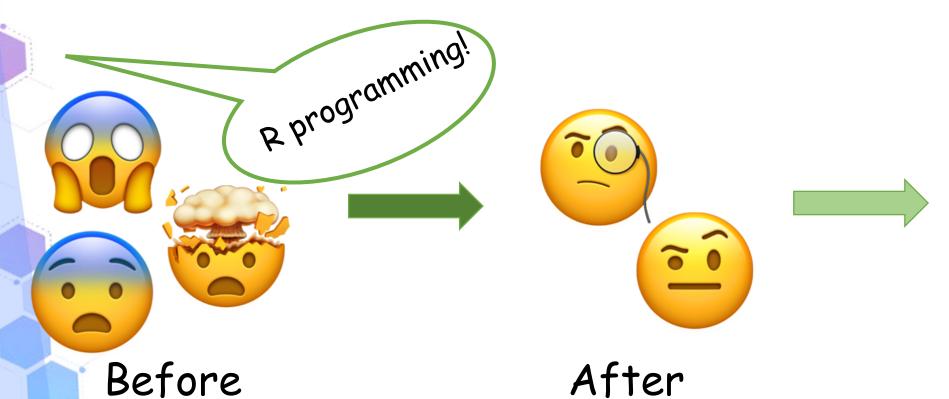








What this session is



After

"I can read simple R scripts" "I know names of some R functions"





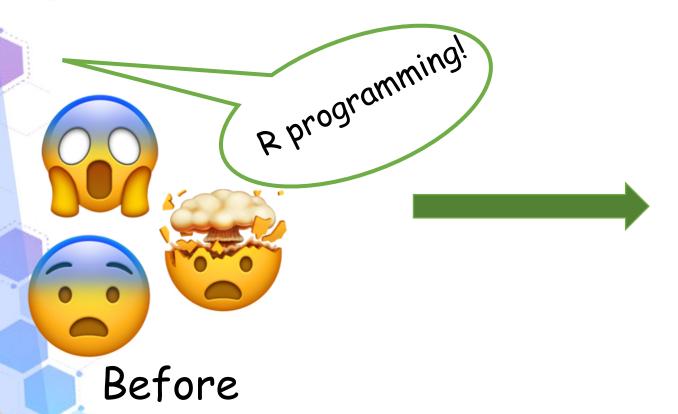
And maybe

"I want to write my own R script"

"I know where to "copy" sample R scripts"



What this session is **NOT**





After

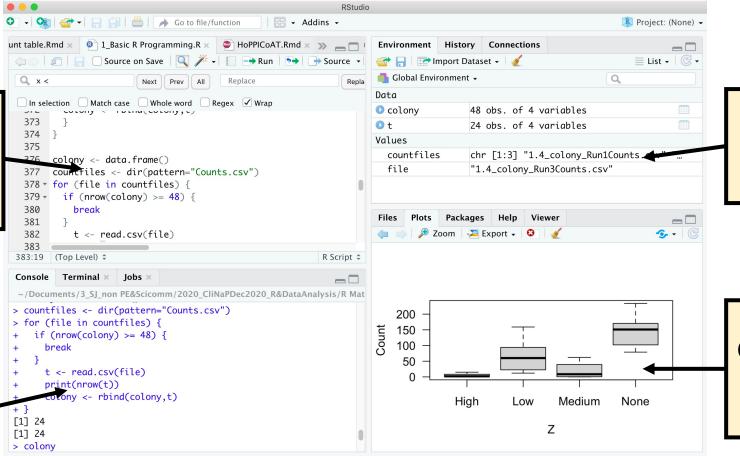
"I can write my own R scripts from scratch"





Store and edit your script here

Scripts that were run + text output + error message appear here



"Environment" listing your data objects

Graphical output appear here



Languages have nouns

Languages have verbs

Languages have syntax (grammar)



- Languages have nouns
 - In R, we call this "Objects" or "variables"

"Objects" or "variables" come in various shapes

- Vector (1-dimension)
- Dataframe (2-dimension / table)
- Matrix (2-dimension / table with one data type)
- **List** (collection of multiple objects)

Objects store data.

"Data" can be...

- Number (numerical)
- Character (text)
- Factor (grouping)
- TRUE / FALSE (logical)



- Languages have verbs
 - In R, we call this "functions" or "commands"
 - Functions do something to the input object(s) and give some output
 - Output can be displayed on console, or stored in another object
 - Functions can have "options" to fine tune its performance

("Options" are also known as "arguments")

Adjective/Adverl



Languages have syntax (grammar)

R code normally looks like this...

Code like this usually gives output on the screen, as text or graphic



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<-or = symbols can be used to assign value to an object.

Code like this put output in the newObject.
Usually, nothing will be displayed on screen



Languages have syntax (grammar)

R code normally looks like this...

Code like this usually gives output on the screen, as text or graphic

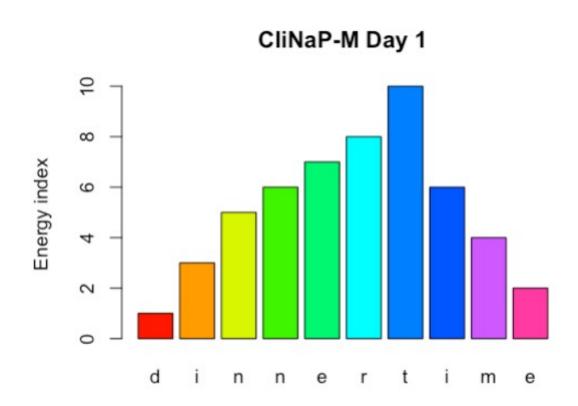
<-o(= symbols can be used to assign value to an object.</p>

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Group exercise

Create this image by modifying the command below







- Languages have verbs
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Adjective/Advert



Function C

User download

the package

Studio

Function D

Function A

R package

Function B

R is a (computer) language

Languages have verbs

• R comes with many <u>built-in functions</u>, but these might not be

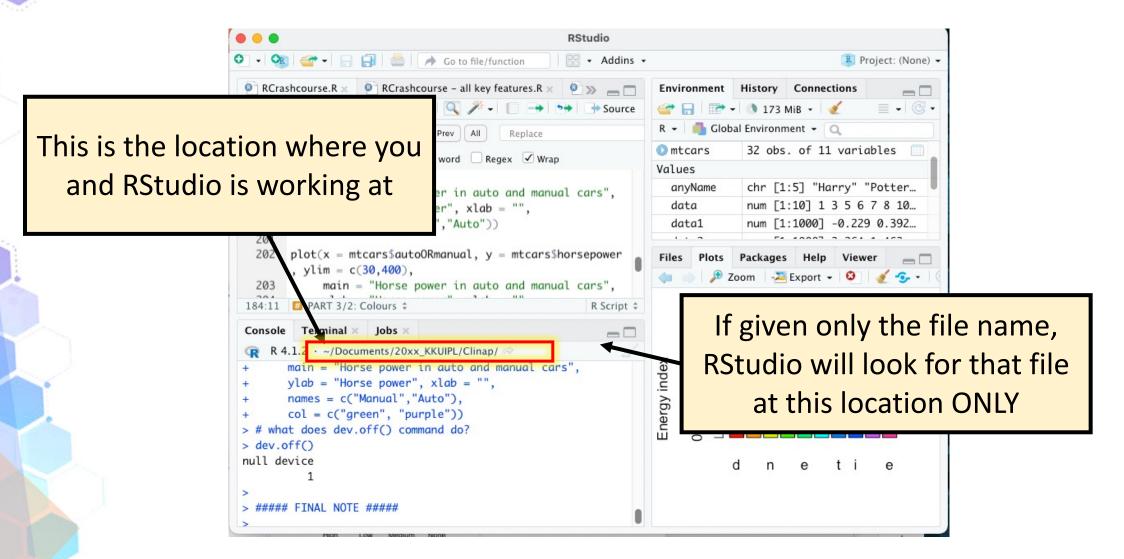
enough for some work, or will require writing a very long code.

Nice people (i.e. developers and programmers)
have created many useful packages that contain
add-on functions for R.

- Most of the packages for bioinformatics work can be downloaded from
 - https://cran.r-project.org/ or https://www.bioconductor.org/
- Once downloaded, you would run command
 library(name of your package) every time you restart RStudio

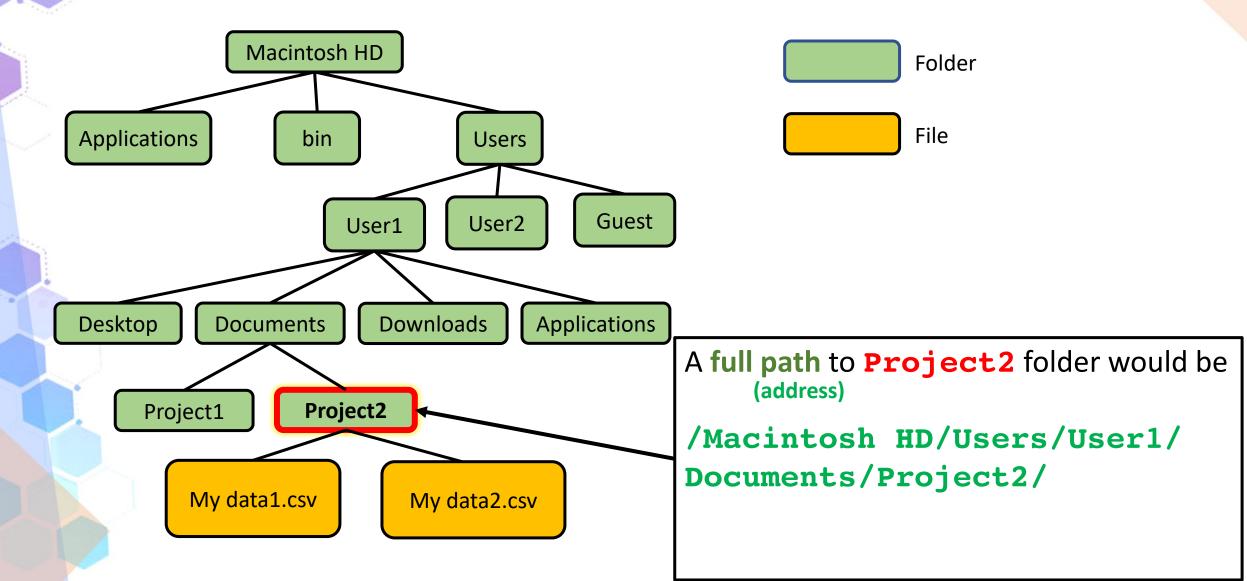


Make sure Rstudio can find your data



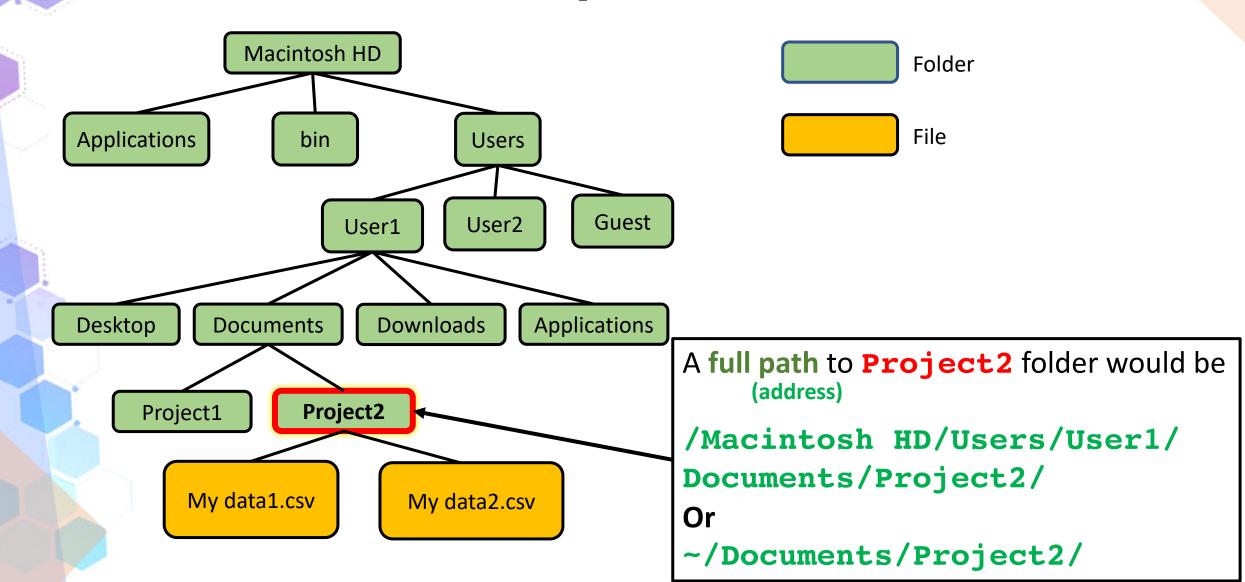


File address – full path



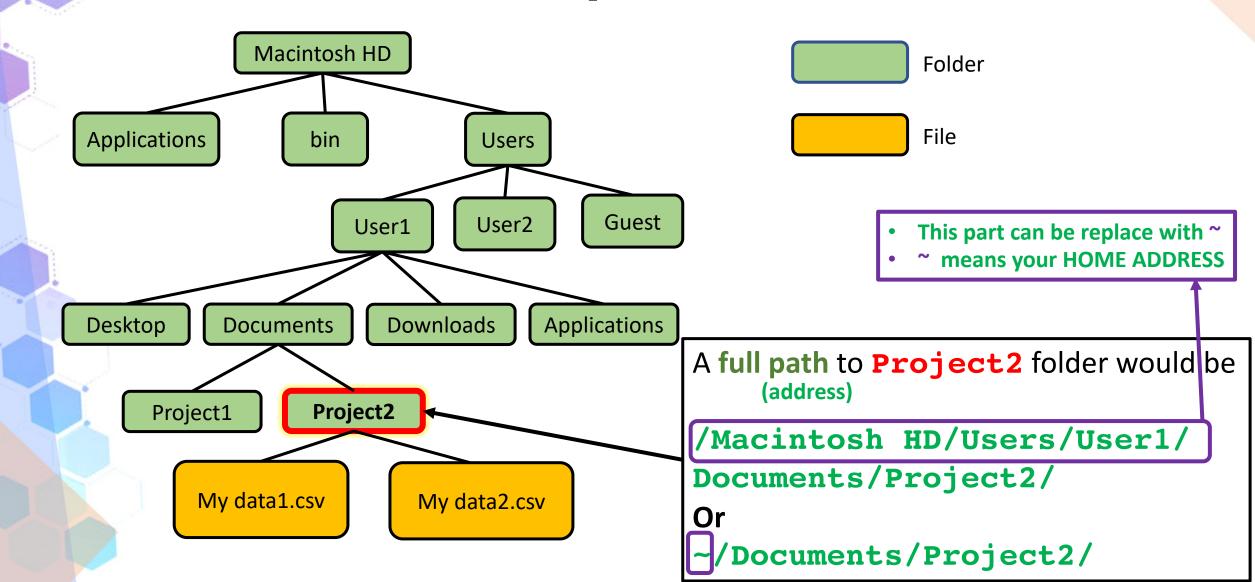


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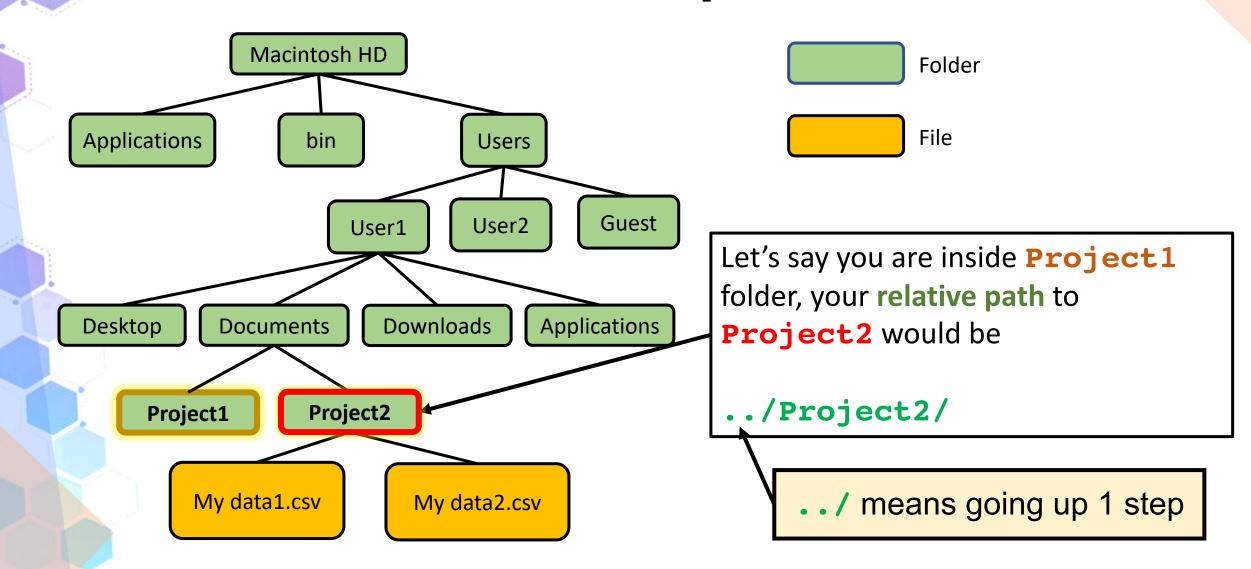


File address – full path



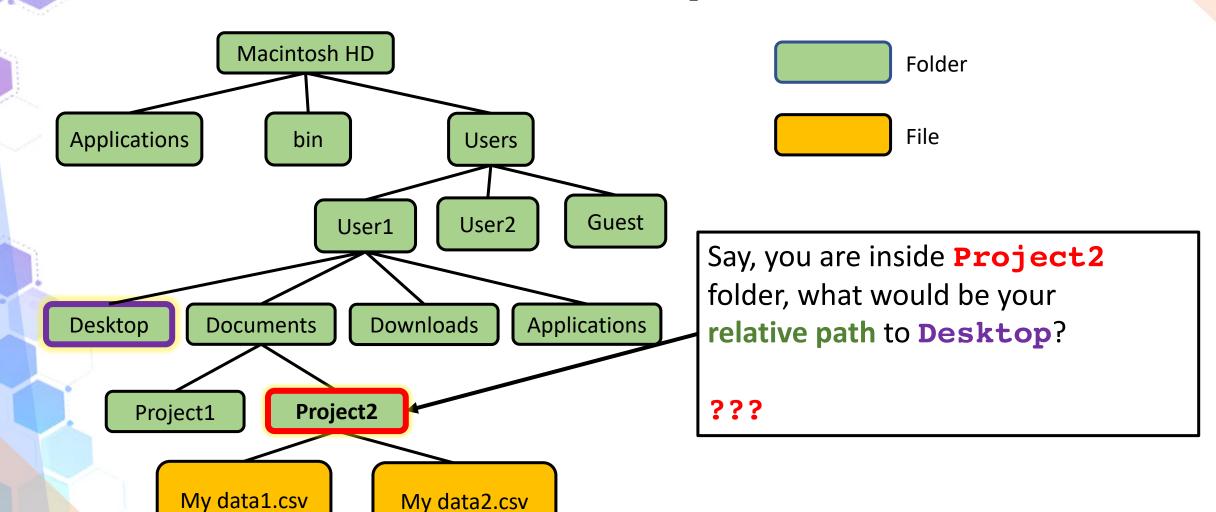


File address – relative path



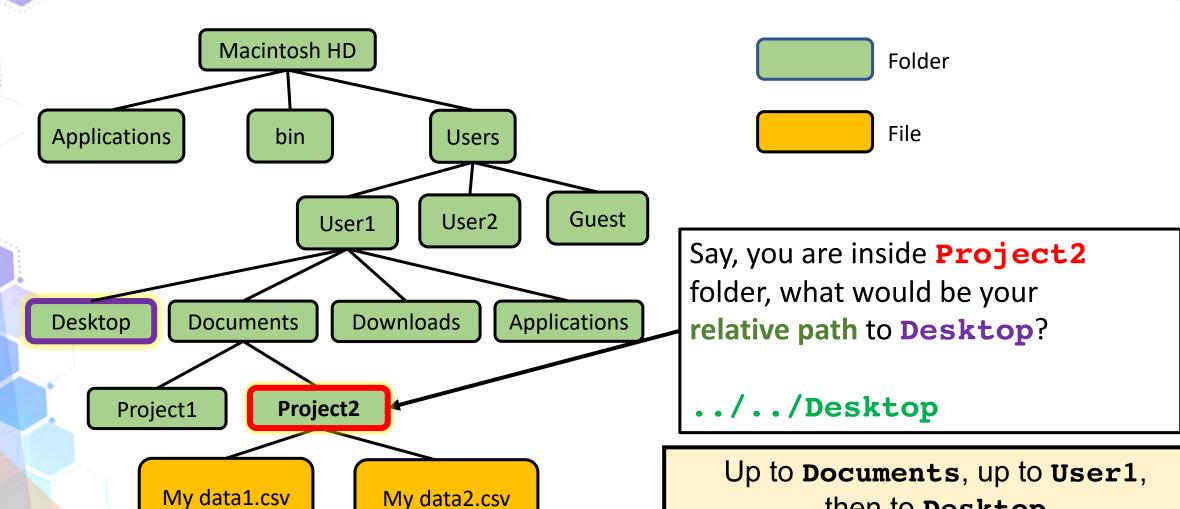


File address – relative path





File address – relative path



then to **Desktop****When in doubt, always use full path**



Make sure Rstudio can find your data

