Package writing

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Why write packages?

- 1) To keep a systematic record of your functions including what they do their inputs and outputs and dependencies.
 - This will facilitate keeping and not losing your work
 - You can easily edit and update your functions once a working edition is made
- 2) To collaborate with fellow scientists concerning your research. A package is the best way to compactly and succinctly manage your applications in one place.
- 3) Pedagogical and motivational: You are forced to determine the impact of the collection of your functions together this will crystallize the purpose of your work
 - The writing of functions and their documentation will clarify and encourage good programming practices
 - How to write functions
 - Learn more about OOP like S3 in R
 - Encourage you to improve your programming skills

Data

• There are three ways to put data into your package – we will learn one of them today (probably the best for most purposes)

Method to get started

- Open your package project.
- Make a new directory `data`
- Read data into your R session
 - Bear in mind you will be in the root of your package directory structure.
 - Use `obj=read.csv(file.choose())` to read the data in
 - Now save the data to the `data` directory use `save(obj, file="data/ddt.rda")` this is a special compressed rdata file format that is low in file size.
 - One more step and that is to document the data in the `R` folder.

Documentation: Below is the roxygen file – remember to save as `DDT.R`

```
#' Fish caught in the tennessee river and its tributaries
#'
#' A dataset with 6 variables two of which are quantitative.
#' Interest is in the amount of DDT in their flesh
#'
#' @format A data frame with 144 rows and 6 variables:
#'\describe{
   \item{RIVER}{Three letter abbreviation of the River}
    \item{MILE}{Mile count for where the fish was caught}
#'
#' ]
#' @source \url{https://www.crcpress.com/Statistics-for-Engineering-and-the-Sciences-Sixth-Edition/M_idenhall-Sincich/p/book/9781498728850}
           Use this name for documentation (just like you would
"ddt"
                             for a function object)
```

devtools

- The devtools package will help you compile and install your package
- There are many options to these functions and many things are done beneath the surface

Once the package is made you can install using

```
devtools::install(build_vignettes = TRUE, force = TRUE)
```

```
Now check
> data(ddt)
RIVER MILE SPECIES LENGTH WEIGHT DDT
1 FCM 5 CCATFISH 42.5 732 10
2 FCM 5 CCATFISH 44.0 795 16
3 FCM 5 CCATFISH 41.5 547 23
4 FCM 5 CCATFISH 39.0 465 21
5 FCM 5 CCATFISH 50.5 1252 50
6 FCM 5 CCATFISH 52.0 1255 150
```

Workflow

Make `data` folder

Read data into R
Obj =
read.csv(file.choose())

Ex. save(obj, file="data/ddt.rda")

Save to data folder save()

Complete data roxygen file and save in R folder

devtools::install(build_vignettes = TRUE, force = TRUE)

Last step