

GitHub & advanced R

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What is GitHub?

http://www.github.com

GitHub is a website and tool to store and share code.

You can share code and data with one or several collaborators, but it will also be open to the world.

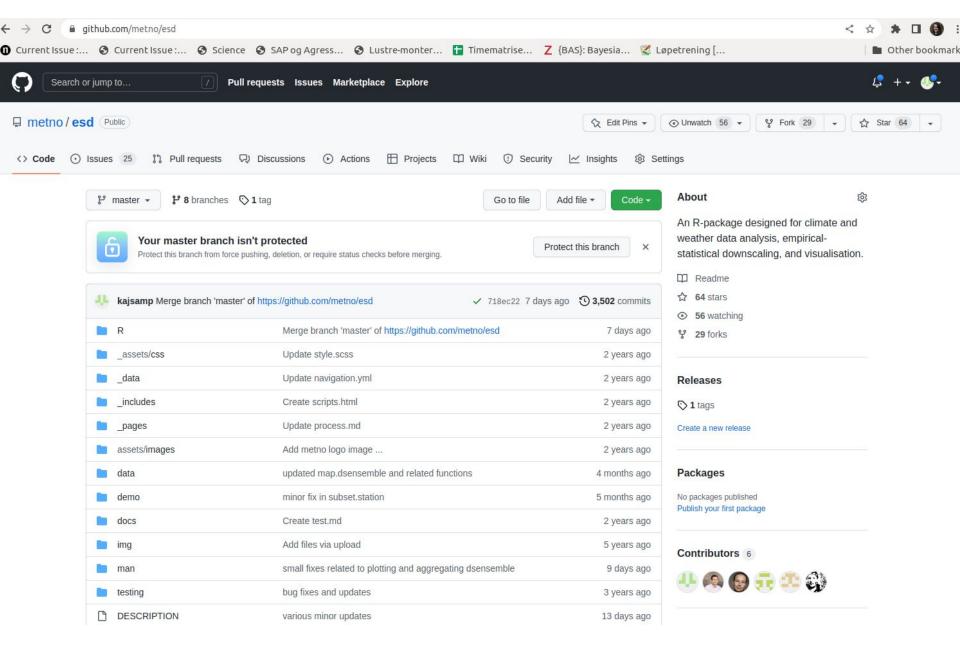
The location where you put your code on GitHub is called a repository.

GitHub is host to many R-packages, including the 'esd' package.



https://github.com/metno/esd







https://github.com/metno/BMD



Exercises

- 1. Register and log in at https://github.com
- 2. Download code and documents from

https://github.com/metno/BMD

3. Upload a script of your own (that you are comfortable sharing).



Working with GitHub locally

You can access GitHub via the terminal window

Basic commands

clone download a repository to your computer

pull update the downloaded repository

add the changes that you have made locally

commit prepare the added changes to be uploaded

upload your local version to the Github repository



push

Exercises

1. Make a directory called git on your computer

```
mkdir git
cd git
```

2. Clone the BMD repository

```
clone https://github.com/metno/BMD.git
```

3. Update the local version of the BMD repository

```
pull BMD
```



GitHub resources

Tutorial

https://docs.github.com/en/get-started/quickstart/hello-world

GitHub book

https://git-scm.com/book/en/v2



R-training

https://github.com/metno/BMD/tree/master/Exercises



A function is a set of instructions, like a recipe. R comes with many built-in functions.

```
## print: Print input
print('Print this input!')

## seq: Create sequence from 2 to 12
x <- seq(2,12)
print(x)

## sum: Calculate the sum of the sequence x
print(sum(x))

## paste: Attach character strings to each other
print(paste("Hello", "world!", sep=" "))</pre>
```

The basic syntax of a function in R looks like this:

```
function_name <- function(input1, input2, ...) {
   function body
}</pre>
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function_name <- function(input1, input2, ...) {
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}

## Example 1: calculate a*b and print the results
f1 <- function(a = 3, b = 6) {
   result <- a * b
   print(result)
}

f1(53, 67)</pre>
```

The basic syntax of a function in R looks like this:

```
function_name <- function(input1, input2, ...) {
   function body
}

## Example 2: calculate a*b and return the results
f2 <- function(a = 3, b = 6) {
   result <- a * b
   return(result)
}

y <- f2(53, 67)</pre>
```

Exercises

- 1. Write a function that prints the phrase: "Hello world!"
- 2. Write a function that attach two words together and print them. Get your function to say "Hello world!".
- 3. Write a function that calculates the mean value of two numbers.
- 4. Write a function that reads a file and extracts some of the data.
- 5. Change the function from step 4 to also produce a plot.
- 6. Find a script that you have written. Make a part of the code into a function and use it in the script.

Using 'for' to create loops

Use for-loops to repeat tasks with minor variations.

Use 'seq' to generate a vector.

```
for(x in 1:10) {
   print(x)
}
```

Using 'for' to create loops

Use for-loops to repeat tasks with minor variations.

Use 'seq' to generate a vector.

```
# A simple for loop that prints different years
years <- seq(2021, 2050)
for(y in years) {
  txt <- paste("The year was", y)
  print(txt)
}</pre>
```

Using 'if' statements

Use if statements to choose what to do based on the input.

```
if(TRUE) {
   print("The statement is true.")
} else {
   print("The statement is false.")
}
```

Using 'if' statements

Use if statements to choose what to do based on the input.

```
# Specify the value of x
x <- 0

if(x < 0) {
   print("x is lower than zero.")
} else {
   print("x is higher or equal to zero.")
}</pre>
```

Using 'if' statements

Use if statements to choose what to do based on the input.

```
file <- '/path/to/file/filename.nc'

if(file.exists(file)) {
   print("The file exists. Opening file.")
   x <- retrieve(file)
} else {
   print("The file does not exist.")
}</pre>
```

