

# Statistical Programming and Open Science Methods

Functional versus object-oriented programming

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ACCOUNTING FOR  
TRANSPARENCY

## Time table October 10

When?	What?
09:00	Welcome and Introduction
09:30	The development environment and project organization
10:30	Coffee
11:00	Using Git and Github
12:30	Lunch
14:00	Statistical programming languages: An overview
15:30	Coffee
16:00	Functional versus object-oriented programming
19:30	Pizza at Due Forni, Schönhauser Allee 12

# Functional programming versus scripting

- ▶ Many statistical programming languages (EViews, SAS, Stata, R to some extent) are in essence scripting languages.
- ▶ Scripts are closely connected to imperative programming (“Shut up and do what I tell you!”)
- ▶ Scripts are hard to read, tend to become inefficient, and are hard to reuse
- ▶ “If you copy + paste your (own) code a lot, you are a bad programmer”

# The key idea of functional programming

- ▶ Functional programming is declarative in nature: Your functions describe what to do. The implementation is hidden from the user.
- ▶ A function takes arguments, processes them and returns results
- ▶ A *pure function* is a function where
  - the result of the function depends only on its arguments and
  - that generates no *side effects*
- ▶ Pure functions are *referentially transparent*, meaning that they can be replaced with their return value without changing the program
- ▶ In real-life coding, many functions are not referentially transparent. This makes writing code easier and reading code harder

# Functions in R

- ▶ Functions have three components:
  - `formals()`: The arguments that you call the function with
  - `body()`: The code that the function executes
  - `environment()`: The place where the function can look for objects
- ▶ Functions are objects, just like about anything else in R
- ▶ Internally, they are called `closures`. Knowing this can be helpful to decipher error messages!

## Chaining functions in R: Intermediate objects

Readable but tedious

```
df <- read_csv("data/sub.csv")
df <- select(df, cik, name)
df <- distinct(df)
count_sec_reg <- nrow(df)
sprintf("There are %d registrants", count_sec_reg)
```

## Chaining functions in R: Nesting

Concise but a pain in the eye

```
sprintf(  
  "There are %d registrants",  
  nrow(distinct(select(read_csv("data/sub.csv"), cik, name)))  
)
```

# Chaining functions in R: Piping

The tidy way (read %>% as “and then”) but harder to debug

```
read_csv("data/sub.csv") %>%  
  select(cik, name) %>%  
  distinct() %>%  
  nrow() -> count_sec_reg  
  
sprintf("There are %d registrants", count_sec_reg)
```



# Scoping I

What does this code snippet return?

```
x <- 10

my_func <- function() {
  x <- 20
  x
}

c(my_func(), x)
```

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What does this code snippet return?

```
x <- 10

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c(my_func(), x)
```

```
## [1] 20 10
```

## Scoping II

What does this code snippet return?

```
x <- 10
y <- 5

my_func <- function() {
  x <- 20
  x*y
}

my_func()
```

## Scoping II

What does this code snippet return?

```
x <- 10
y <- 5

my_func <- function() {
  x <- 20
  x*y
}

my_func()
```

```
## [1] 100
```

## Scoping III

What does this code snippet return?

```
my_second_func <- function(x) {  
  y <- x  
}
```

```
my_func <- function(x) {  
  x*y  
}
```

```
my_second_func(5)  
my_func(10)
```

## Scoping III

What does this code snippet return?

```
my_second_func <- function(x) {  
  y <- x  
}
```

```
my_func <- function(x) {  
  x*y  
}
```

```
my_second_func(5)  
my_func(10)
```

```
## Error in my_func(10): object 'y' not found
```

## Scoping IV

What does this code snippet return?

```
my_second_func <- function(x) {  
  y <- x  
}  
  
my_func <- function(x) {  
  x*y  
}  
  
my_second_func(5)  
my_func(10)
```

## Scoping IV

What does this code snippet return?

```
my_second_func <- function(x) {  
  y <- x  
}
```

```
my_func <- function(x) {  
  x*y  
}
```

```
my_second_func(5)  
my_func(10)
```

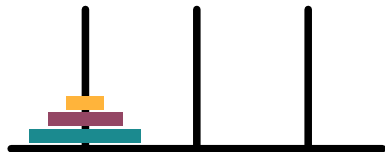
```
## [1] 50
```



For extra credit

Which of these functions are pure, which are not? Why?

## Recursions: Functions can be very helpful



```
tower <- function(n, from_peg, to_peg, aux_peg) {  
  if(n == 0) return(invisible())  
  tower(n - 1, from_peg, aux_peg, to_peg)  
  message(sprintf("Moving piece %d from %s to %s ...",  
                  n, from_peg, to_peg), appendLF = FALSE)  
  tower(n - 1, aux_peg, to_peg, from_peg)  
}  
tower(3, 'F', 'T', 'A')
```

## Recursions: Functions can be very helpful

```
tower <- function(n, from_peg, to_peg, aux_peg) {  
  if(n == 0) return(invisible())  
  tower(n - 1, from_peg, aux_peg, to_peg)  
  message(sprintf("Moving piece %d from %s to %s ...",  
                  n, from_peg, to_peg))  
  tower(n - 1, aux_peg, to_peg, from_peg)  
}  
tower(3, 'F', 'T', 'A')
```

## Moving piece 1 from F to T ...

## Moving piece 2 from F to A ...

## Moving piece 1 from T to A ...

## Moving piece 3 from F to T ...

## Moving piece 1 from A to F ...

## Moving piece 2 from A to T ...

## Moving piece 1 from F to T ...

See <https://www.youtube.com/watch?v=YstLjLCGmgg> for animation

# Object oriented programming

- ▶ Much more common in Python than in R, object oriented programming encapsulates data and functions (aka as *methods* in the OOP world) in *classes*
- ▶ Methods can be *overloaded* by *inheriting* classes
- ▶ Tends to make code more consistent and easier to maintain/extend
- ▶ Makes it easier for code to modify data (makes objects more *mutable*), something that people in statistical programming are generally not very fond of
- ▶ How does this look like: Let's have a quick look at a last toy example
  - `code/show_fs_oop.py` versus
  - `code/show_fs_fp.R`