Statistics 360: Advanced R for Data Science Lecture 1

Brad McNeney

Course Objectives

R objects: names and values

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Course objectives

- Work through the book Advanced R by Hadley Wickham: https://adv-r.hadley.nz/index.html
- R objects: names and values
- Basic data structures and programming.
 - vectors, subsetting, control flow, functions, environments
 - No tidyverse this time
- Object-oriented programming in R
- Code performance: debugging, profiling, memory, calling Python, C or ++ from R
- Parallelizing R code (if time permits)

Getting started with R, RStudio and git

- ► Follow the "getting started" instructions on the class canvas page to get set up with R, RStudio and git.
 - ▶ R and RStudio will be familiar, but you may not have used git before, so leave some time for that.
- ▶ Please try to get R and RStudio installed and create an RStudio project linked to the class GitHub repository as soon as possible.
- ► Those still having trouble after the weekend should ask our TA, Pulindu, for help during the first lab session on Jan. 21.

Reading

▶ Welcome, Preface and Chapter 1 of the text.

R objects: names and values

R objects

- ▶ In R, data structures and functions are all referred to as "objects".
- Objects are created with the assignment operator <-; e.g., x <- c(1,2,3).</p>
 - ► The objects a user creates from the R console are contained in the user's workspace, called the global environment.
 - ▶ Use ls() to see a list of all objects in the workspace.
 - Use rm(x) to remove object x from the workspace.

Names and values

Reference: text, chapter 2

install.packages("lobstr")

- ► The description on the previous slide of x <- c(1,2,3) is an over-simplification.
- ▶ It is more accurate to say we've done two things: (i) created an object in computer memory that contains 1, 2, 3 and (ii) "bound" that object to the "name" x.

```
library(lobstr)
x <- c(1,2,3)
ls()
## [1] "x"
obj_addr(x) # changes every time this code chunk is run
## [1] "0x55eafe2a19f8"</pre>
```

Syntactic vs non-syntactic

- Valid, or "syntactic" names in R can consist of letters, digits, . and _ but should start with a letter.
- Names that start with . are hidden from directory listing with 1s().
- Names that start with _ or a digit are non-syntactic and will cause an error.
- ▶ If you need to create or access a non-syntatic name, use backward single-quotes ("backticks").

```
x <- 1

.x <- 1

`_x` <- 1

ls()
```

```
## [1] "_x" "x"
```

Modifying, copying, binding

Mofifying a variable causes a copy to be made.

```
x \leftarrow c(1,2,3); y \leftarrow x
c(obj_addr(x),obj_addr(y))
## [1] "0x55eafe9ab988" "0x55eafe9ab988"
x[[2]] \leftarrow 10 \# Note: x[2] \leftarrow 10 \text{ has the same effect}
c(obj_addr(x),obj_addr(y))
## [1] "0x55eafd9712d8" "0x55eafe9ab988"
х
## [1] 1 10 3
## [1] 1 2 3
```

Tracing copying

► The tracemem() function marks an object so that a message is printed whenever a copy is made.

```
x <- c(1,2,3)
tracemem(x)

## [1] "<0x55eafe8da2b8>"
x[[2]] <- 10

## tracemem[0x55eafe8da2b8 -> 0x55eafe28bd18]: eval eval withVis
x <- 5 # removes the trace on the object
x[[1]] <- 1</pre>
```

More on tracmem()

► As the output of tracemem() suggests, the trace is on the object, not the name:

```
x \leftarrow c(1,2,3)
tracemem(x)
## [1] "<0x55eafe9aab78>"
v <- x
c(obj_addr(x),obj_addr(y))
## [1] "0x55eafe9aab78" "0x55eafe9aab78"
y[[2]] <- 10
## tracemem[0x55eafe9aab78 -> 0x55eafe92c488]: eval eval withVis
c(obj addr(x),obj addr(y))
   [1] "0x55eafe9aab78" "0x55eafe92c488"
```

Function calls

R has a reputation for passing copies to functions, but in fact the copy-on-modify applies to functions too:

```
f <- function(arg) { return(arg) }
x <- c(1,2,3)
y <- f(x) # no copy made
c(obj_addr(x),obj_addr(y))

## [1] "0x55eafe28ba48" "0x55eafe28ba48"
f <- function(arg) { arg <- 2*arg; return(arg) }
y <- f(x) # copy made
c(obj_addr(x),obj_addr(y))</pre>
```

[1] "0x55eafe28ba48" "0x55eafde4f8b8"

Lists

List elements point to objects too:

```
11 <- list(1, 2, 3)
c(obj_addr(11),obj_addr(11[[1]]),obj_addr(11[[2]]),obj_addr(11[[3]]))
## [1] "0x55eafe9abc58" "0x55eafe143c48" "0x55eafe143c10" "0x55eafe143b
# Note: ref() will print the above, formatted, but ...</pre>
```

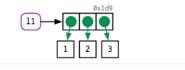


Figure 1: list

(Note: ref() will print a nicely formatted version of the above, but it doesn't work with in my slides.)

Copy-on-modify in lists

- As you would expect, we only copy the list **elements** that are modified, rather than the entire list.
 - tracemem() flags any change to the list

```
11 \leftarrow list(c(1,2), c(3,4), c(5,6,7))
c(obj_addr(11),obj_addr(11[[1]]),obj_addr(11[[2]]),obj_addr(11[[3]]))
## [1] "0x55eafeba6e98" "0x55eafbccb168" "0x55eafbccb1e8" "0x55eafeba6e
tracemem(11)
## [1] "<0x55eafeba6e98>"
11[[1]] <- 55
## tracemem[0x55eafeba6e98 -> 0x55eafe94e0e8]: eval eval withVisible wi
c(obj_addr(11),obj_addr(11[[1]]),obj_addr(11[[2]]),obj_addr(11[[3]]))
```

[1] "0x55eafe94e0e8" "0x55eafd55fa88" "0x55eafbccb1e8" "0x55eafeba6e

Copies of lists are "shallow"

```
12 <- 11
12[[3]] <- 111

## tracemem[0x55eafe94e0e8 -> 0x55eafd6f04d8]: eval eval withVisible wi
c(obj_addr(11),obj_addr(11[[1]]),obj_addr(11[[2]]),obj_addr(11[[3]]))

## [1] "0x55eafe94e0e8" "0x55eafd55fa88" "0x55eafbccb1e8" "0x55eafeba6e
c(obj_addr(12),obj_addr(12[[1]]),obj_addr(12[[2]]),obj_addr(12[[3]]))
```

[1] "0x55eafd6f04d8" "0x55eafd55fa88" "0x55eafbccb1e8" "0x55eafed526

Data frames are lists . . .

```
dd \leftarrow data.frame(x=1:3,y=4:6)
c(obj_addr(dd[[1]]),obj_addr(dd[[2]]))
## [1] "0x55eafbf2ec28" "0x55eafbf2ed08"
dd[,2] <- 7:9
c(obj_addr(dd[[1]]),obj_addr(dd[[2]]))
## [1] "0x55eafbf2ec28" "0x55eafc63fa90"
dd[1,] \leftarrow c(11,22)
c(obj_addr(dd[[1]]),obj_addr(dd[[2]]))
## [1] "0x55eafe8da308" "0x55eafe8da2b8"
dd[1,2] <- 111
c(obj_addr(dd[[1]]),obj_addr(dd[[2]]))
```

[1] "0x55eafe8da308" "0x55eafe28b778"

Beware of data frame overhead

tracemem(dd)

Data frames are convenient, but the convenience comes at a cost.

dd <- data.frame(x=rnorm(100)) # try yourself with rnorm(1e7)</pre>

Can illustrate by tracing copying when we modify columns.

```
## [1] "<0x55eafed840a8>"
dmed <- lapply(dd,median)

## tracemem[0x55eafed840a8 -> 0x55eafd55ddd8]: as.list.data.frame as.li
dd[[1]] <- dd[[1]] - dmed[[1]] # same as dd[,1] - dmed[[1]]</pre>
```

tracemem[0x55eafed840a8 -> 0x55eafebb1a00]: eval eval withVisible wi
tracemem[0x55eafebb1a00 -> 0x55eafebb0650]: [[<-.data.frame [[<- eva</pre>

Fewer copies if we do the same with a list.

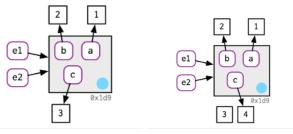
```
11 <- list(x=rnorm(100))
tracemem(ll)

## [1] "<0x55eafdddfa28>"
lmed <- lapply(l1,median)
11[[1]] <- l1[[1]] - dmed[[1]]</pre>
```

tracemem[0x55eafdddfa28 -> 0x55eafe8dea48]: eval eval withVisible wi

Modify-in-place

► The text claims two exceptions to the copy-on-modify, but in my experiments environments are the only one:



```
e1 <- rlang::env(a = 1, b = 2, c = 3)
e2 <- e1
e1$c <- 4
e2$c
```

```
## [1] 4
```