Statistics 360: Advanced R for Data Science Lecture 3

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Control Flow

R Functions

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- ► Reading: text, chapter 5
- ▶ if/if-else, ifelse, switch
- ► for
- while
- break

if and if-else

• if tests a condition and executes code if the condition is true. Optionally, can couple with an else to specify code to execute when condition is false.

```
if("cat" == "dog") {
  print("cat is dog")
} else {
  print("cat is not dog")
}
```

[1] "cat is not dog"

if returns a value

► The body of the if-else can evaluate expressions and store results, but note that if-else also returns a value.

```
## [1] "cat is not dog"
```

if expects a single logical

- most other inputs will cause an error
- ▶ logical vectors will not throw an error, but if will only use the first element

```
try(if("cat") print("cat"))

## Error in if ("cat") print("cat") :

## argument is not interpretable as logical
if(c("cat"=="dog","cat" == "cat")) print("hello world")

## Warning in if (c("cat" == "dog", "cat" == "cat")) print("hello world"): the
## condition has length > 1 and only the first element will be used
```

ifelse(): vectorized if

- ▶ ifelse() can handle logical vectors
- syntax is condition, what to return if expression true, what to return if expression false

```
x <- 1:10
ifelse(x %% 2 == 0, "even", "odd")
## [1] "odd" "even" "odd" "even" "odd" "even" "odd" "even" "odd" "even"</pre>
```

switch

- If you have multiple conditions to check, consider switch instead of repeated if-else; e.g.
 - if(x==1) "cat" else if(x==2) "dog" else if (x==3)
 "mouse"

Warning: unknown animal

for loops

Example:

```
n <- 10; nreps <- 100; x <- vector(mode="numeric",length=nreps)
for(i in 1:nreps) {
    # Code you want to repeat nreps times
    x[i] <- mean(rnorm(n))
}
summary(x)

## Min. 1st Qu. Median Mean 3rd Qu. Max.
## -0.88260 -0.16747 0.04650 0.02435 0.26527 0.69914
print(i)

## [1] 100</pre>
```

for loop index set

Index sets such as 1:n are most common, but can be almost any atomic vector.

```
ind <- c("cat", "dog", "mouse")
for(i in ind) {
   print(paste("There is a",i, "in my house"))
}

## [1] "There is a cat in my house"
## [1] "There is a dog in my house"
## [1] "There is a mouse in my house"</pre>
```

seq_along

- ▶ A common use of for loops is to iterate over elements of a vector, say x.
- Creating the index set 1:length(x) will not be what you expect when x has length 0 (e.g., x is NULL).
- Instead use seq_along()

```
x <- NULL
for(i in 1:length(x)) print(x[i])

## NULL
## NULL
for(i in seq_along(x)) print(x[i])</pre>
```

while loops

Use a while loop when you want to continue until some logical condition is met.

```
set.seed(1)
# Number of coin tosses until first success (geometric distn)
p <- 0.1; counter <- 0; success <- FALSE
while(!success) {
   success <- as.logical(rbinom(n=1,size=1,prob=p))
   counter <- counter + 1
}
counter
## [1] 4</pre>
```

break

break can be used to break out of a for or while loop.

```
for(i in 1:100) {
   if(i>3) break
   print(i)
}
## [1] 1
## [1] 2
## [1] 3
```

repeat

- ▶ repeat continues indefinitely until it encounters a break
- ► The text considers repeat to be the most flexible of for, while and repeat.

R Functions

R function fundamentals

- ▶ Reading: text sections 6.1 and 6.2
- ▶ In R, functions are objects with three essential components:
 - the code inside the function, or body,
 - the list of arguments to the function, or formals, and
 - an environment that contains all objects defined in the function.
- Functions can have other attributes, but the above three are essential.

Example function

```
f <- function(x) {
   return(x^2)
}
f

## function(x) {
## return(x^2)
## }</pre>
```

The function body

- This is the code we want to execute.
- ▶ When the end of a function is reached without a call to return(), the value of the last line is returned.
 - ► So in our example function, we could replace return(x^2) with just 'x^2.
- ▶ Use body() to see the body of a function.

```
body(f)
## {
## return(x^2)
## }
```

The function formals

- ▶ These are the arguments to the function.
- Function arguments can have default values and/or be defined in terms of other arguments.

```
f <- function(x=0) { x^2}
f <- function(x=0,y=3*x) { x^2 + y^2 }
f()

## [1] 0
f(x=1)

## [1] 10
f(y=1)

## [1] 1</pre>
```

```
formals(f)

## $x
## [1] 0
##
## $y
## 3 * x
```

Argument matching when calling a function

When you call a function, the arguments are matched first by name, then by "prefix" matching and finally by position:

```
f <- function(firstarg, secondarg) {</pre>
  firstarg<sup>2</sup> + secondarg
f(firstarg=1,secondarg=2)
## [1] 3
f(s=2,f=1)
## [1] 3
f(2,f=1)
## [1] 3
f(1,2)
## [1] 3
```

The function environment

- The environment within a function is like a map to the memory locations of all its variables.
- Variables created within the function are also stored in its environment

```
f <- function(x) {
  y <- x^2
  ee <- environment() # Returns ID of environment w/in f
  print(ls(ee)) # list objects in ee
  ee
}
f(1) # function call

## [1] "ee" "x" "y"

## <environment: 0x55aedc4da180>
```

environment(f)

<environment: R_GlobalEnv>

Enclosing environments

- Our function f was defined in the global environment, .GlobalEnv, which "encloses" the environment within f.
- ▶ If f needs a variable and can't find it within f's environment, it will look for it in the enclosing environment, and then the enclosing environment of .GlobalEnv, and so on.
- ► The search() function lists the heirarchy of environments that enclose .GlobalEnv.

search()

```
## [1] ".GlobalEnv" "package:stats" "package:graphics
## [4] "package:grDevices" "package:utils" "package:datasets
## [7] "package:methods" "Autoloads" "package:base"
```

► To facilitate this search, each environment includes a pointer to its enclosing environment.

R packages and the search list

install.packages("hapassoc")

- Use the library() command to load packages.
- ▶ When we load a package it is inserted in position 2 of the search list, just after .GlobalEnv.

Detaching packages

Detach a package from the search list with detach()

Package namespaces

- Package authors create a list of objects that will be visible to users when the package is loaded. This list is called the package namespace.
- You can access functions in a package's namespace without loading the package using the :: operator.

```
set.seed(321)
n<-30; x<-(1:n)/n; y<-rnorm(n,mean=x); ff<-lm(y~x)
car::sigmaHat(ff)
## [1] 0.926726</pre>
```

Doing so does not add the package to the search list.

Up next

► Reading: Text, sections 6.4-6.8