Contents

Quiz for Week 2

R Programming Week 2 Videos

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Suppose we define the following function in R

```
cube <- function(x, n) {
    x^3
}</pre>
```

What is the result of running cube(3) in R after defining this function?

Options

- 1. The number 27 is returned
- 2. An error is returned because 'n' is not specified in the call to 'cube'
- 3. A warning is given with no value returned.
- 4. The users is prompted to specify the value of 'n'.

Remarks

- As we can see by running the code we get the answer we expect.
- While n is specified as a possible argument for the function, it is clear that it is not used in the body of the function, and that an answer can be computed without it.

Counter-Example

```
pow <- function(x, n=3) {
          x^n
}</pre>
```

Suppose I define the following function in R

```
pow <- function(x = 4, n = 3) {
      x^n
}</pre>
```

What is the result of running pow() in R after defining this function?

- 1. A warning is given and the function returns 64.
- 2. An error is given the function does not finish execution.
- 3. The number 64 is returned.
- 4. The number 81 is returned.

As we can see by running the code - we get the answer we expect, without any error warnings.

If no arguments are specified by the user, the function will use the **default** settings.

This function can be used for other values of x and n.

The following code will produce a warning in R.Why is this?

```
x <- 1:10
if(x > 5) {
    x <- 0
}
```

Options

- 1. 'x' is a vector of length 10 and 'if' can only test a single logical statement.
- 2. There are no elements in 'x' that are greater than 5 (obviously false)
- 3. The expression uses curly braces.
- 4. You cannot set 'x' to be 0 because 'x' is a vector and 0 is a scalar.

 (true)
- 5. The syntax of this R expression is incorrect. (No obvious Syntax errors)

The aggregate() function

Firstly let us construct a data frame called df1, using the code below.

```
set.seed(1) #generating random numbers

#Three Variables
col1 <- c(rep('happy',9), rep('sad', 9))
col2 <- rep(c(rep('alpha', 3),
    rep('beta', 3), rep('gamma', 3)),2)
score=rnorm(18, 10, 3)

#Combine the 3 variables as a data frame
df1<-data.frame(col1=col1, col2=col2, score=score)
df1</pre>
```

The aggregate() function

There are two categorical variables. The first (i.e. col1) has two levels, the second (i.e. col2) has three.

We can use the aggregate() function to apply a specified command for groups.

aggregate(variable, by=list(group1,group2,..),function)

```
aves1 = aggregate(df1$score,
    by=list(col2=df1$col2), mean)

aves1

aves2 = aggregate(df1$score,
    by=list(col1=df1$col1), mean)

aves2

aves3 = aggregate(df1$score,
    by=list(col1=df1$col1, col2=df1$col2), mean)

aves3
```

We can use the merge command to combine the group-wise results with the original data frame.

```
results = merge(df1, aves2)
results
```

```
> results = merge(df1, aves2)
> results
```

```
col1 col2 score x happy alpha 8.120639 10.54247
```

- 2 happy alpha 10.550930 10.54247
- 3 happy alpha 7.493114 10.54247
- 4 happy beta 14.785842 10.54247
- 5 happy beta 10.988523 10.54247

.

Take a look at the 'iris' dataset that comes with R.

The data can be loaded with the code:

```
library(datasets)
data(iris)
```

Questions

- A description of the dataset can be found by running ?iris.
- There will be an object called '*iris*' in your workspace.
- In this dataset, what is the mean of 'Sepal.Length' for the species virginica?
- (Please only enter the numeric result and nothing else.)

Remarks

- One approach is to use the aggregate() command mentioned in the previous section.
- A second approach is to create a subset (let's call it *iris.vir*) and then use the summary() command.

```
iris.vir=iris[iris$Species=="virginica",]
# or
iris.vir=subset(iris,iris$Species=="virginica")
summary(iris.vir)
```

The summary of *iris.vir* should look like this:

```
> summary(iris.vir)
```

•		
Sepal.Length	Sepal.Width	Petal.Length
Min. :4.900	Min. :2.200	Min. :4.500
1st Qu.:6.225	1st Qu.:2.800	1st Qu.:5.100
Median :6.500	Median :3.000	Median :5.550
Mean :6.588	Mean :2.974	Mean :5.552
3rd Qu.:6.900	3rd Qu.:3.175	3rd Qu.:5.875
Max. :7.900	Max. :3.800	Max. :6.900
Petal.Width	Species	
Min. :1.400	setosa : 0	
1st Qu.:1.800	versicolor: 0	
Median :2.000	virginica :50	
Mean :2.026		
3rd Qu.:2.300		
Max. :2.500		

The apply() function

The **Apply** family of functions keep you from having to write loops to perform some operation on every row or every column of a matrix or data frame, or on every element in a list.

The apply() function

- The apply() function is a powerful device that operates on arrays and, in particular, matrices.
- The apply() function returns a vector (or array or list of values) obtained by applying a specified function to either the row or columns of an array or matrix.
- To specify use for rows or columns, use the additional argument of
 - 1 for rows,
 - 2 for columns.

```
m <- matrix(c(1:10, 11:20), nrow = 10, ncol = 2)
m
apply(m,1,mean)
apply(m,2,mean)</pre>
```

```
> # create a matrix of 10 rows x 2 columns
> m <- matrix(c(1:10, 11:20), nrow = 10, ncol = 2)
>
> # mean of the rows
>
> apply(m, 1, mean)
[1] 6 7 8 9 10 11 12 13 14 15
>
> # mean of the columns
> apply(m, 2, mean)
[1] 5.5 15.5
```

The lapply() and sapply() function

- The lapply() command returns a list of the same length as a list X, each element of which is the result of applying a specified function to the corresponding element of X.
- A simpler user-friendly version of lapply() is sapply() The sapply() command is a variant of lapply() returning a simple vector instead of a list again of the same length as a list X, each element of which is the result of applying a specified function to the corresponding element of X.

```
> x <- list(a=1:10, b=exp(-3:3), logic=c(T,F,F,T))
>
> # compute the list mean for each list element
>
> lapply(x,mean)
$a
[1] 5.5
$b
[1] 4.535125
$logic
[1] 0.5
>
> sapply(x,mean)
a b logic
5.500000 4.535125 0.500000
>
```

What is an environment in R? a list whose elements are all functions a collection of symbol/value pairs a special type of function an R package that only contains data

The R language uses what type of scoping rule for resolving free variables? compilation scoping lexical scoping dynamic scoping global scoping

How are free variables in R functions resolved? The values of free variables are searched for in the working directory The values of free variables are searched for in the environment in which the function was defined The values of free variables are searched for in the environment in which the function was called The values of free variables are searched for in the global environment

What is one of the consequences of the scoping rules used in R? All objects can be stored on the disk R objects cannot be larger than 100 MB All objects must be stored in memory Functions cannot be nested

In R, what is the parent frame? It is the environment in which a function was defined It is always the global environment It is the environment in which a function was called It is the package search list

Continuing with the '*iris*' dataset from Question 4, what R code returns a vector of the means of the variables 'Sepal.Length', 'Sepal.Width', 'Petal.Length', and 'Petal.Width'?

```
colMeans(iris)
apply(iris, 2, mean)
apply(iris[, 1:4], 1, mean)
apply(iris[, 1:4], 2, mean)
apply(iris, 1, mean)
rowMeans(iris[, 1:4])
```

Try out all the code, to see what happens. However, based on your knowledge of the apply family of functions, you should spot that this option would be suitable.

R code that would have worked if it had been an option

```
> colMeans(iris[,1:4])
Sepal.Length Sepal.Width Petal.Length Petal.Width
    5.843333    3.057333    3.758000    1.199333
```

Load the 'mtcars' dataset in R with the following code

```
library(datasets)
data(mtcars)
```

- There will be an object names '*mtcars*' in your workspace. You can find some information about the dataset by running ?mtcars.
- How can one calculate the average miles per gallon (mpg) by number of cylinders in the car (cyl)?

Options

```
tapply(mtcars$cyl, mtcars$mpg, mean)
split(mtcars, mtcars$cyl)
mean(mtcars$mpg, mtcars$cyl)
tapply(mtcars$mpg, mtcars$cyl, mean)
```

- Cars can have either 4, 6 or 8 cylinders.
 - > table(mtcars\$cyl)

• using the tapply() function

Continuing with the *mtcars* dataset from Question 6, what is the absolute difference between the average horsepower of 4cylinder cars and the average horsepower of 8-cylinder cars?

Remarks

This is another question where the aggregate() command comes in handy. (We will use attach() and detach() to avoid unnecessary typing)

```
> attach(mtcars)
> CylMeans <- aggregate(hp,by=list(Cyls=cyl),mean)</pre>
> CylMeans
  Cyls
                X
1
     4
        82.63636
2
     6 122.28571
     8 209.21429
> detach(mtcars)
>
> CylMeans$x
     82.63636 122.28571 209.21429
Г1]
> CylMeans$x[3]-CylMeans$x[1]
[1] 126.5779
```

What is the difference between the 'sapply()' function and the 'lapply()' function?

Options

- 1. There is no difference; 'sapply' and 'lapply' are two names for the same function
- 2. 'sapply()' always returns a 2-dimensional matrix while 'lapply' returns a list.
- 3. 'lapply()' always returns a list while 'sapply()' attempts to simplify the result.
- 4. 'lapply()' always returns an atomic vector and 'sapply' always returns a list.

Remark The question can be easily solved by reading the help files for both commands.

help(sapply)

Consider the following function

What value is returned by 'f'?

If you run debug(ls) what happens when you next call the 'ls' function?

- (i) Execution of 'ls' will suspend at the beginning of the function and you will be in the browser.
- (ii) The 'ls' function will execute as usual. (FALSE)
- (iii) The 'ls' function will return an error. (FALSE)
- (iv) Execution of the '1s' function will suspend at the 4th line of the function and you will be in the browser.
- (v) You will be prompted to specify at which line of the function you would like to suspend execution and enter the browser.