

Contents

Quiz for Week 2

Quiz Question 1

Suppose we define the following function in R

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

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  
}
```

Quiz Question 2

Suppose I define the following function in R

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

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`.

Question 3

Consider the following function

```
f <- function(x) {  
  g <- function(y) {  
    y + z  
  }  
  z <- 4  
  x + g(x)  
}
```

If I then run in R

```
z <- 10  
f(3)
```

What value is returned?

- (i) 7
- (ii) 4
- (iii) 10
- (iv) 16

Question 4

Consider the following expression:

```
x <- 5
y <- if(x < 3) {
  NA
} else {
  10
}
```

What is the value of 'y' after evaluating this expression?

- (i) 10
- (ii) 5
- (iii) 3
- (iv) NA

Question 5

Consider the following R function

```
h <- function(x, y = NULL, d = 3L) {  
  z <- cbind(x, d)  
  if(!is.null(y))  
    z <- z + y  
  else  
    z <- z + f  
  g <- x + y / z  
  if(d == 3L)  
    return(g)  
  g <- g + 10  
  g  
}
```

Which symbol in the above function is a free variable?

- (i) f
- (ii) z
- (iii) d
- (iv) L
- (v) g

Question 6

What is an environment in R?

- (i) a list whose elements are all functions
- (ii) a collection of symbol/value pairs
- (iii) a special type of function
- (iv) an R package that only contains data

Question 7

The R language uses what type of scoping rule for resolving free variables?

- (i) compilation scoping
- (ii) lexical scoping
- (iii) dynamic scoping
- (iv) global scoping

Question 8

How are free variables in R functions resolved?

- (i) The values of free variables are searched for in the working directory
- (ii) The values of free variables are searched for in the environment in which the function was defined
- (iii) The values of free variables are searched for in the environment in which the function was called
- (iv) The values of free variables are searched for in the global environment

Question 9

What is one of the consequences of the scoping rules used in R?

- (i) All objects can be stored on the disk
- (ii) R objects cannot be larger than 100 MB
- (iii) All objects must be stored in memory
- (iv) Functions cannot be nested

Question 10

In R, what is the parent frame?

- (i) It is the environment in which a function was defined
- (ii) It is always the global environment
- (iii) It is the environment in which a function was called
- (iv) It is the package search list

Old Quiz Question 3

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
```

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
1  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
.....
```

OldQuiz Question 4

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)
```

```
> # 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
>
```

Old Quiz Question 5

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.

```
> apply(iris[, 1:4], 2, mean)
Sepal.Length Sepal.Width Petal.Length Petal.Width
      5.843333      3.057333      3.758000      1.199333
```

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
```

Old Quiz Question 6

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)
```

```
 4  6  8
11  7 14
```

- using the `tapply()` function

```
> tapply(mtcars$mpg, mtcars$cyl, mean)
      4      6      8
26.66364 19.74286 15.10000
```


Old Quiz Question 7

Continuing with the *mtcars* dataset from Question 6, what is the absolute difference between the average horsepower of 4-cylinder 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)
> CylMeans
  Cyls      x
1    4 82.63636
2    6 122.28571
3    8 209.21429
> detach(mtcars)
>
> CylMeans$x
[1] 82.63636 122.28571 209.21429
>
> CylMeans$x[3]-CylMeans$x[1]
[1] 126.5779
```

Old Quiz Question 8

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)`

Old Quiz Question 9

Consider the following function

```
f <- function(x) {  
  g <- function(y) {  
    y + z  
  }  
  z <- 4  
  x + g(x)  
}
```

If I then run in R

```
z <- 10  
f(3)
```

What value is returned by 'f'?

Old Quiz Question 10

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 `'ls'` 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.