1 Quiz for Week 3

The aggregate() function

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

```
#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>
```

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(grouping1,grouping2,..),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
```

3

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",]

#Alternatively
iris.vir=subset(iris,iris$Species=="virginica")

summary(iris.vir)
```

The summary of $\boldsymbol{iris.vir}$ should look like this:

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

Species

setosa : 0
versicolor: 0
virginica :50

The apply() function family

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
>
```

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
```

\mathbf{split}

split divides the data in the vector **x** into the groups defined by **f**. The replacement forms replace values corresponding to such a division. unsplit reverses the effect of split.

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

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)

Smry for summary

Old Quiz Question

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. (Correct)
- 4. 'lapply()' always returns an atomic vector and 'sapply' always returns a list.

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

help(sapply)

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

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