

Acadgild – Data Analytics – Batch 4 Assignment –

Session 1 – Assignment -1

Task: 1

1. How many ways are there to call a function in R ?

Answer:

- a. `call(name, ...)` where, `name`: is a non-empty character string naming the function to be called ... stands for arguments to be part of the call `call` returns an unevaluated function call, that is, an unevaluated expression which consists of the named function applied to the given. Although the call is unevaluated, the arguments ... are evaluated.
- b. `do.call` `do.call` constructs and executes a function call from a name or a function and a list of arguments to be passed to it.
- c. `Recall` `Recall` is used as a placeholder for the name of the function in which it is called. It allows the definition of recursive functions which still work after being renamed

2. What is Recycling of elements in a vector?

Answer:

R automatically recycles, or repeats, elements of the shorter Vector when applying an operation to two vectors that requires them to be the same length, R automatically recycles, or repeats, elements of the shorter one, until it is long enough to match the longer Vector.

3. Give an example of recycling of elements in a vector.

Answer:

Example 1:

Suppose we have two Vectors `c(1,2,4)` , `c(6,0,9,10,13)`, where the first one is shorter with only 3 elements. Now if we sum these two, we will get a warning message as follows.

```
> c(1,2,4) + c(6,0,9,10,13)
```

```
[1] 7 2 13 11 15
```

Warning message:

In `c(1, 2, 4) + c(6, 0, 9, 10, 13)` : longer object length is not a multiple of shorter object length

Here R , Sum those Vectors by Recycling or repeating the elements in shorter one, until it is long enough to match the longer one as follows..

```
> c(1,2,4,1,2) + c(6,0,9,10,13)
```

```
[1] 7 2 13 11 15
```

```
> a <- c(10,2,23,4)+c(2,10)
```

```

> print(a)
[1] 12 12 25 14
> b<- c(1,2,3,4,5,6,7) + c(1,3)
warning message:
In c(1, 2, 3, 4, 5, 6, 7) + c(1, 3) :
  longer object length is not a multiple of shorter object length
> print(b)
[1] 2 5 4 7 6 9 8
> x <- c(1,2,3,4,5,6)+c(2,10)
> print(x)
[1] 3 12 5 14 7 16
> y<- c(1,2,3,4,5,6,7) + c(10,30)
warning message:
In c(1, 2, 3, 4, 5, 6, 7) + c(10, 30) :
  longer object length is not a multiple of shorter object length
> print(y)
[1] 11 32 13 34 15 36 17

```

Task 2:

1. What should be the output of the following Script?

```
v <- c( 2,5.5,6)
```

```
t <- c(8, 3, 4)
```

```
print(v%%/t)
```

v<-c(2,5.5,6) t<-c(8,3,4) print(v%%/t) output

```
[1] 0 1 1
```

```

> v <- c( 2,5.5,6)
> t <- c(8, 3, 4)
> print(v%%/t)
[1] 0 1 1

```

2. You have 25 excel files with names as xx_1.xlsx, xx_2.xlsx,.....xx_25.xlsx in a dir.

Write a program to extract the contents of each excel sheet and make it one df.

Ans:- setwd("c:/R/mergeme") Or specific file path name files=list.files(pattern=".xlsx") for(i in 1:length(files)) {filename=files[i] data=read.xlsx(file = filename,header = T) assign(x = filename,value = data)} #Suppose the columns are the same for each file, #you can bind them together in one dataframe with bind_rows from dplyr: library(dplyr) #one more option is as follows df<-lapply(files, read.xlsx) %>% bind_rows()

Task 3:

1. Create an $m \times n$ matrix with `replicate(m, rnorm(n))` with $m=10$ column vectors of $n=10$ elements each, constructed with `rnorm(n)`, which creates random normal numbers. Then we transform it into a dataframe (thus 10 observations of 10 variables) and perform an algebraic operation on each element using a nested for loop: at each iteration, every element referred by the two indexes is incremented by a sinusoidal function, compare the vectorized and non-vectorized form of creating the solution and report the system time differences.

#Answer

1

```
#Vectorized form
set.seed(42)

#create matrix
mat_1<- replicate(10,rnorm(10))

#transform into data frame
df_1= data.frame(mat_1)
df_1<- df_1 + 10*sin(0.75*pi)

#non-vectorized form
set.seed(42)
#create matrix
mat_1<- replicate(10,rnorm(10))
#transform into data frame
df_1= data.frame(mat_1)

for(i in 1:10){
  for(j in 1:10){
    df_1[i,j]<- df_1[i,j] + 10*sin(0.75*pi)
    print(df_1)
  }
}

#time difference
```

```

system.time(
  df_1[i,j]<- df_1[i,j] + 10*sin(0.75*pi)
)

system.time(
  for(i in 1:10){
    for(j in 1:10){
      df_1[i,j]<- df_1[i,j] + 10*sin(0.75*pi)
    }
  }
)

```

Task 4:

1. Define matrix mymat by replicating the sequence 1:5 for 4 times and transforming into a matrix, sum over rows and columns.

Solution:

The R-script for the given problem is as follows:

```

rep(1:5, 4)      # replicating the sequence 1 to 5
mymat <- matrix(rep(1:5,4), nrow = 4 , ncol = 5, byrow = TRUE )
mymat
# sum over rows and columns.
apply(mymat, 1, sum)      # sum of rows
apply(mymat, 2, sum)      # sum of columns

```

Explanation:

- Matrix mymat is created by replicating the sequence of 1 to 5 (1,2,3,4,5) for 4 times by using rep(1:5,4).
- The matrix mymat is of order 4X5 (4 rows and 5 columns)
- The sum over rows and columns is found by apply() function using the r-commands as follows:
 1. apply(mymat, 1, sum) # sum of rows
 2. apply(mymat, 2, sum) # sum of columns

Here,1 is used for rows and 2 is used for columns.

The output of the R-Script (from Console window) is given as follows:

```
> rep(1:5, 4)
```

```

[1] 1 2 3 4 5 1 2 3 4 5 1 2 3 4 5 1 2 3 4 5
> mymat <- matrix(rep(1:5 ,4), nrow = 4 , ncol = 5, byrow = TRUE )
> mymat
      [,1] [,2] [,3] [,4] [,5]
[1,]     1     2     3     4     5
[2,]     1     2     3     4     5
[3,]     1     2     3     4     5
[4,]     1     2     3     4     5
> # sum over rows and columns.
> apply(mymat, 1, sum)      # sum of rows
[1] 15 15 15 15
> apply(mymat, 2, sum)      # sum of columns
[1]  4  8 12 16 20

```

Task 5:

1. States = rownames(US Arrests)

Get states names with 'w'.

Get states names with 'W'.

Solution:

The R-script for the given problem is as follows:

Get states names with 'w'.

```
States[grepl("w", States)]
```

#Get states names with 'W'.

```
States[grepl("W", States)]
```

Explanation:

grepl() function searches for matches to argument pattern within each element of a character vector.

To get the states names with 'w' , grepl("w", States) is used.

To get states names with 'W' , grepl("W", States) is used.

The output of the R-Script (from Console window) is given as follows:

```

> USArrests
      Murder Assault UrbanPop Rape
Alabama    13.2    236      58 21.2
Alaska     10.0    263      48 44.5
Arizona     8.1    294      80 31.0
Arkansas    8.8    190      50 19.5
California  9.0    276      91 40.6
Colorado    7.9    204      78 38.7
Connecticut 3.3    110      77 11.1
Delaware    5.9    238      72 15.8
Florida    15.4    335      80 31.9

```

```

Georgia      17.4      211      60 25.8
Hawaii       5.3       46      83 20.2
Idaho        2.6      120      54 14.2
Illinois     10.4     249      83 24.0
Indiana      7.2      113      65 21.0
Iowa         2.2       56      57 11.3
Kansas       6.0      115      66 18.0
Kentucky     9.7      109      52 16.3
Louisiana    15.4     249      66 22.2
Maine        2.1       83      51  7.8
Maryland     11.3     300      67 27.8
Massachusetts 4.4      149      85 16.3
Michigan     12.1     255      74 35.1
Minnesota    2.7       72      66 14.9
Mississippi  16.1     259      44 17.1
Missouri     9.0      178      70 28.2
Montana      6.0      109      53 16.4
Nebraska     4.3      102      62 16.5
Nevada       12.2     252      81 46.0
New Hampshire 2.1       57      56  9.5
New Jersey   7.4      159      89 18.8
New Mexico   11.4     285      70 32.1
New York     11.1     254      86 26.1
North Carolina 13.0     337      45 16.1
North Dakota 0.8       45      44  7.3
Ohio         7.3      120      75 21.4
Oklahoma     6.6      151      68 20.0
Oregon        4.9      159      67 29.3
Pennsylvania 6.3      106      72 14.9
Rhode Island 3.4      174      87  8.3
South Carolina 14.4     279      48 22.5
South Dakota 3.8       86      45 12.8
Tennessee    13.2     188      59 26.9
Texas        12.7     201      80 25.5
Utah         3.2      120      80 22.9
Vermont      2.2       48      32 11.2
Virginia     8.5      156      63 20.7
Washington   4.0      145      73 26.2
West Virginia 5.7       81      39  9.3
Wisconsin    2.6       53      66 10.8
Wyoming      6.8      161      60 15.6
> States
[1] "Alabama"      "Alaska"      "Arizona"      "Arkansas"      "California"
[6] "Colorado"     "Connecticut" "Delaware"     "Florida"       "Georgia"
[11] "Hawaii"       "Idaho"       "Illinois"     "Indiana"       "Iowa"
[16] "Kansas"       "Kentucky"    "Louisiana"    "Maine"         "Maryland"
[21] "Massachusetts" "Michigan"     "Minnesota"    "Mississippi"   "Missouri"
[26] "Montana"      "Nebraska"    "Nevada"       "New Hampshire" "New Jersey"
[31] "New Mexico"   "New York"    "North Carolina" "North Dakota"  "Ohio"

```

```

[36] "Oklahoma"      "Oregon"      "Pennsylvania" "Rhode Island" "South Carolina"
[41] "South Dakota"  "Tennessee"   "Texas"        "Utah"         "Vermont"
[46] "Virginia"      "Washington"   "West Virginia" "Wisconsin"     "Wyoming"
> # Get states names with 'w'.
>
> States[grep("w", States)]
[1] "Delaware"      "Hawaii"      "Iowa"        "New Hampshire" "New Jersey"
[7] "New Mexico"
[7] "New York"
> #Get states names with 'w'.
>
> States[grep("w", States)]
[1] "Washington"    "West Virginia" "Wisconsin"    "Wyoming"

```

2. Prepare a Histogram of the number of characters in each US state.

Solution:

The R-script for the given problem is as follows:

```
df <- nchar(States)
```

```
df
```

```
hist(df)
```

Explanation:

nchar () takes a character vector as an argument and returns a vector whose elements contain the sizes of the corresponding elements

hist () computes a histogram of the given data values

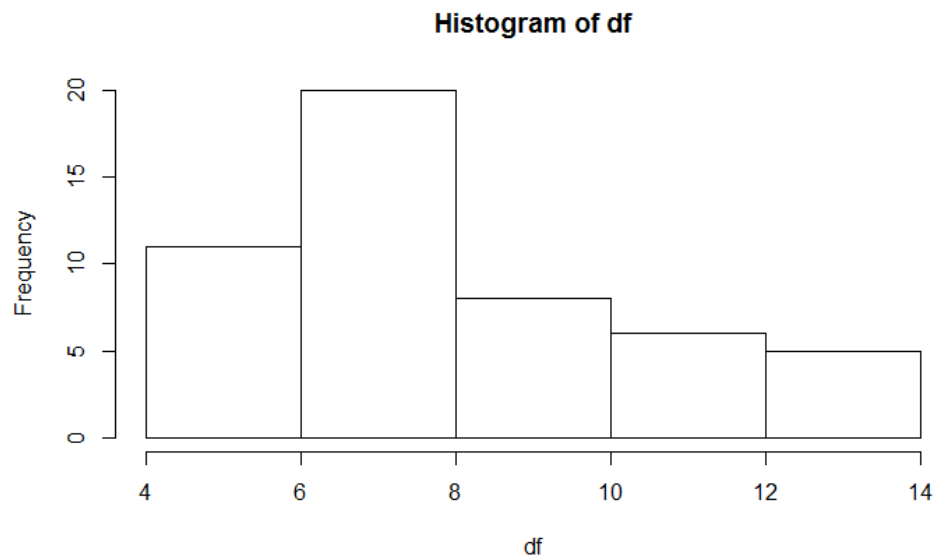
The output of the R-Script (from Console window) is given as follows:

```

> df <- nchar(States)
> df
[1] 7 6 7 8 10 8 11 8 7 7 6 5 8 7 4 6 8 9 5 8 13 8 9 11
8 7 8 6 13 10 10 8
[33] 14 12 4 8 6 12 12 14 12 9 5 4 7 8 10 13 9 7
> hist(df)

```

From plot window:



Task 6:

1. Test whether two vectors are exactly equal (element by element).

```
vec1 = c(rownames(mtcars[1:15,]))
```

```
vec2 = c(rownames(mtcars[11:25,]))
```

Solution:

The R-script for the given problem is as follows:

```
vec1 = c(rownames(mtcars[1:15,]))
```

```
vec2 = c(rownames(mtcars[11:25,]))
```

```
isTRUE(all.equal(vec1,vec2))          # returns true/false
```

```
identical(vec1,vec2)                  # returns true/false
```

```
all.equal(vec1,vec2)                  # returns number of differences
```

Explanation:

- `isTRUE(all.equal(vec1,vec2))` returns TRUE if vec1 is equal to vec2;else it returns FALSE.
- `identical(vec1,vec2)` returns TRUE if vec1 is identical/same to vec2;else it returns FALSE.
- `all.equal(vec1,vec2)` returns number of differences between vec1 and vec2.

The output of the R-Script (from Console window) is given as follows:


```

> vec1 = c(rownames(mtcars[1:15,]))
> vec2 = c(rownames(mtcars[11:25,]))
> isTRUE(all.equal(vec1,vec2))
[1] FALSE
> identical(vec1,vec2)
[1] FALSE
> all.equal(vec1,vec2)
[1] "15 string mismatches"

```

2. Sort the character vector in ascending order and descending order.

```
vec1 = c(rownames(mtcars[1:15,]))
```

```
vec2 = c(rownames(mtcars[11:25,]))
```

Solution:

The R-script for the given problem is as follows:

```
vec1 = c(rownames(mtcars[1:15,]))
```

```
vec2 = c(rownames(mtcars[11:25,]))
```

```
sort(vec1)                                # vec1 in ascending order
```

```
sort(vec1, decreasing = TRUE)             # vec1 in descending order
```

```
sort(vec2)                                # vec2 in ascending order
```

```
sort(vec2, decreasing = TRUE)             # vec2 in descending order
```

Explanation:

sort(vec1) function arranges the character vector vec1 in ascending order. For descending order “decreasing” parameter is set as “TRUE”

sort(vec2) function arranges the character vector vec2 in ascending order. For descending order “decreasing” parameter is set as “TRUE”

The output of the R-Script (from Console window) is given as follows:

```

> vec1 = c(rownames(mtcars[1:15,]))
> vec2 = c(rownames(mtcars[11:25,]))
> sort(vec1)
 [1] "Cadillac Fleetwood" "Datsun 710"          "Duster 360"          "Hornet 4
Drive"
 [5] "Hornet Sportabout"  "Mazda RX4"           "Mazda RX4 Wag"       "Merc 230
"
 [9] "Merc 240D"          "Merc 280"           "Merc 280C"          "Merc 450
SE"

```

```

[13] "Merc 450SL"      "Merc 450SLC"      "Valiant"
> sort(vec1,decreasing = TRUE)
[1] "Valiant"      "Merc 450SLC"      "Merc 450SL"      "Merc 450
SE"
[5] "Merc 280C"      "Merc 280"      "Merc 240D"      "Merc 230
"
[9] "Mazda RX4 wag"  "Mazda RX4"      "Hornet Sportabout" "Hornet 4
Drive"
[13] "Duster 360"      "Datsun 710"      "Cadillac Fleetwood"
> sort(vec2)
[1] "AMC Javelin"      "Cadillac Fleetwood" "Camaro Z28"      "Chrys
ler Imperial"
[5] "Dodge Challenger" "Fiat 128"      "Honda Civic"      "Linco
ln Continental"
[9] "Merc 280C"      "Merc 450SE"      "Merc 450SL"      "Merc
450SLC"
[13] "Pontiac Firebird" "Toyota Corolla"  "Toyota Corona"
> sort(vec2,decreasing = TRUE)
[1] "Toyota Corona"  "Toyota Corolla"  "Pontiac Firebird" "Merc
450SLC"
[5] "Merc 450SL"      "Merc 450SE"      "Merc 280C"      "Linco
ln Continental"
[9] "Honda Civic"      "Fiat 128"      "Dodge Challenger" "Chrys
ler Imperial"
[13] "Camaro Z28"      "Cadillac Fleetwood" "AMC Javelin"

```

3. What is the major difference between str() and paste() show an example?

Explanation:

str() gives the class of variable, number of values and the elements whereas paste() prints or displays the actual elements .

For example:

str(mtcars\$mpg) gives the class of mtcars\$mpg as num, number of values as 32(1:32) and the elements as 21 21 22.8 21.4 18.7 18.1 14.3 24.4 22.8 19.2 ...

whereas paste(mtcars\$mpg) prints the actual elements present in mtcars\$mpg.

The output of the R-Script (from Console window) is given as follows:

```

> str(mtcars$mpg)
num [1:32] 21 21 22.8 21.4 18.7 18.1 14.3 24.4 22.8 19.2 ...
> paste(mtcars$mpg)
[1] "21" "21" "22.8" "21.4" "18.7" "18.1" "14.3" "24.4" "22.8" "19.2" "1
7.8" "16.4" "17.3"
[14] "15.2" "10.4" "10.4" "14.7" "32.4" "30.4" "33.9" "21.5" "15.5" "15.2" "1
3.3" "19.2" "27.3"
[27] "26" "30.4" "15.8" "19.7" "15" "21.4"

```

4. Introduce a separator when concatenating the strings.

The R-script for the given problem is as follows:

```

paste(rownames(mtcars[1,]), rownames(mtcars[2,]), sep = " ")
paste(rownames(mtcars[1,]), rownames(mtcars[4,]), sep = ",")
paste(rownames(mtcars[2,]), rownames(mtcars[1,]), sep = "--")
paste(rownames(mtcars[3,]), rownames(mtcars[10,]), sep = "$")
paste("hello","world",sep=" @ ")
paste("Assignment","5","3",sep="_")

```

Explanation:

paste(rownames(mtcars[1,]), rownames(mtcars[2,]), sep = " ") introduces a separator ,a single blank " " between the strings rownames(mtcars[1,]) and rownames(mtcars[2,]).

paste(rownames(mtcars[1,]), rownames(mtcars[4,]), sep = ",") introduces a separator comma ", " between the strings rownames(mtcars[1,]) and rownames(mtcars[4,]).

paste(rownames(mtcars[2,]), rownames(mtcars[1,]), sep = "--") introduces a separator "-- " between the strings rownames(mtcars[2,]) and rownames(mtcars[1,]).

paste(rownames(mtcars[3,]), rownames(mtcars[10,]), sep = "\$") introduces a separator dollar "\$ " between the strings rownames(mtcars[3,]) and rownames(mtcars[10,]).

paste("hello","world",sep="@") introduces a separator "@" between the strings "hello" and "world"

paste("Assignment","5","3",sep="_") introduces a separator underscore "_ " between the strings "Assignment", "5" and "3".

The output of the R-Script (from Console window) is given as follows:

```

> paste(rownames(mtcars[1,]), rownames(mtcars[2,]), sep = " ")
[1] "Mazda RX4 Mazda RX4 Wag"
> paste(rownames(mtcars[1,]), rownames(mtcars[4,]), sep = ",")
[1] "Mazda RX4,Hornet 4 Drive"
> paste(rownames(mtcars[2,]), rownames(mtcars[1,]), sep = "--")
[1] "Mazda RX4 Wag--Mazda RX4"
> paste(rownames(mtcars[3,]), rownames(mtcars[10,]), sep = "$")
[1] "Datsun 710$Merc 280"
> paste("hello","world",sep=" @ ")
[1] "hello @ world"
> paste("Assignment","5","3",sep="_")
[1] "Assignment_5_3"

```