R Notebook

Code ▼

DATA SCIENCE PROGRAMMING II (BSD2223)

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QUESTIONS 1

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#a.

#The main characteristics of list data structure are the list is a generic object consisting of an ordered collection of objects. Next, the list can contain elements of different data types, s uch as vectors, matrices, characters, functions and others. Other than that, the list can be acc essed by names or indices using the dollar (\$) operator or the double slicing operator ([[]]) and can be a parent class for other data structures, such as data.frame.

#b.

#Two functions from family of apply(.) functions which can be used for lists are:

#1.lapply(): This function applies the function to each element of the list and returns the list as a result.

#2.sapply(): This function is similar to lapply(), but it tries to simplify the output by return
ing a vector or matrix instead of the list.

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Questions 2

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```
#a.
11 <- seq(-10, 10, length.out = 25)
m1 <- matrix(c(TRUE, FALSE, TRUE, TRUE, FALSE, TRUE, FALSE, TRUE, FALSE, FALSE, FALSE, TRUE, FALSE, TRUE, TRUE), nrow = 4)
v1 <- c("apple", "banana", "kiwi", "watermelon")
v2 <- c("low", "medium", "medium", "low", "high", "low")
list1 <- list(l1, m1, v1, v2)
list1</pre>
```

```
[[1]]
[1] -10.0000000 -9.1666667 -8.3333333 -7.5000000
 [5] -6.6666667 -5.8333333 -5.0000000 -4.1666667
 [9] -3.3333333 -2.5000000 -1.6666667 -0.8333333
[13] 0.0000000
                0.8333333 1.6666667
                                       2.5000000
                            5.0000000
[17] 3.3333333
                4.1666667
                                        5.8333333
[21] 6.6666667
                 7.5000000
                            8.3333333
                                       9.1666667
[25] 10.0000000
[[2]]
     [,1] [,2] [,3] [,4]
[1,] TRUE TRUE TRUE TRUE
[2,] FALSE FALSE FALSE
[3,] FALSE TRUE FALSE TRUE
[4,] TRUE FALSE FALSE TRUE
[[3]]
[1] "apple"
               "banana"
                           "kiwi"
[4] "watermelon"
[[4]]
[1] "low"
            "medium" "medium" "low"
                                     "high"
[6] "low"
```

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```
#b.
names(list1) <- c("Sequence","Matrix","Character","Character")
list1</pre>
```

```
$Sequence
 [1] -10.0000000 -9.1666667 -8.3333333 -7.5000000
 [5] -6.6666667 -5.8333333 -5.0000000 -4.1666667
 [9] -3.3333333 -2.5000000 -1.6666667
                                        -0.8333333
[13]
      0.0000000
                  0.8333333
                              1.6666667
                                          2.5000000
[17]
     3.3333333
                  4.1666667
                              5.0000000
                                          5.8333333
[21]
      6.6666667
                  7.5000000
                              8.3333333
                                          9.1666667
[25] 10.0000000
$Matrix
      [,1] [,2] [,3] [,4]
[1,] TRUE TRUE TRUE TRUE
[2,] FALSE FALSE FALSE
[3,] FALSE TRUE FALSE TRUE
[4,] TRUE FALSE FALSE TRUE
$Character
[1] "apple"
                "banana"
                             "kiwi"
[4] "watermelon"
$Character
[1] "low"
            "medium" "medium" "low"
                                       "high"
[6] "low"
                                                                                           Hide
#c.
list1[[2]][c(3, 1), c(3, 4)]
      [,1] [,2]
[1,] FALSE TRUE
[2,] TRUE TRUE
                                                                                           Hide
```

```
#d.
round(list1[[1]][list1[[1]] > -0.5 & list1[[1]] < 7], digits = 2)
```

```
[1] 0.00 0.83 1.67 2.50 3.33 4.17 5.00 5.83 6.67
```

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```
#e.
list1[[4]][list1[[4]] == "medium"] <- "low"
list1[[4]]</pre>
```

```
[1] "low" "low" "low" "high" "low"
```