Assignment 2

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Question 01:

Working with data in the datasets folder on the course GitHub repo, you will find a file called cancer.csv, a dataset in comma-separated values (csv) format. This is a large cancer incidence dataset that summarizes the incidence of different cancers for various subgroups.

```
#i:Load the data set into R and make it a data frame called cancer.df
cancer.df <- data.frame(read.csv('cancer.csv', fill=TRUE, header = TRUE))</pre>
#ii:Determine the number of rows and columns in the data frame
nrow(cancer.df)
## [1] 42120
ncol(cancer.df)
## [1] 8
#iii:Extract the names of the columns in cancer.df
colnames(cancer.df)
## [1] "year"
                    "site"
                                  "state"
                                               "sex"
                                                             "race"
## [6] "mortality" "incidence"
                                 "population"
#iv:Report the value of the 3000th row in column 6
cancer.df[3000,6]
## [1] 350.69
#v:Report the contents of the 172nd row.
cancer.df[172,]
##
                                       site state sex race mortality
## 172 1999 Brain and Other Nervous System nevada Male Black
       incidence population
## 172
               0
                      73172
#vi:Create a new column that is the incidence rate (per 100,000) for each row.
cancer.df$incidence_rate <- ( cancer.df$incidence )/100000</pre>
colnames(cancer.df)
## [1] "year"
                         "site"
                                                            "sex"
                                          "state"
## [5] "race"
                         "mortality"
                                          "incidence"
                                                            "population"
## [9] "incidence_rate"
```

```
#vii:How many subgroups (rows) have a zero incidence rate?
sum(cancer.df$incidence_rate == 0)
```

[1] 23191

```
#viii:Find the subgroup with the highest incidence rate
cancer.df[which.max(cancer.df$incidence_rate),]
```

```
## year site state sex race mortality incidence population
## 21387 2002 Breast california Female White 3463.74 18774 13690681
## incidence_rate
## 21387 0.18774
```

Question 02:

i. Create the following vector: x <-c("5","12","7"). Which of the following commands will produce an error message? For each command, Either explain why they should be errors, or explain the non-erroneous result.

```
x <- c("5","12","7")
class(x)
```

[1] "character"

x consists of three "character" arguments

```
max(x)
```

```
## [1] "7"
```

Within maxima command, it can return numeric or character arguments. Thus we can get value "7" as character.

```
sort(x)
```

```
## [1] "12" "5" "7"
```

Sort can return objects with a numeric, complex, character or logical vectors. Thus we can get the order of x.

```
#sum(x) will give an error which couldn't be shown in PDF.
```

Sum can only return numeric or complex or logical vectors. Thus in this case where x consists only character value, we will receive an error message.

ii:For the next two commands, either explain their results, or why they should produce errors.

```
y <- c("5",7,12)
#y[2] + y[3] will give an error message which couldn't be shown in PDF
```

y is cassified as character variable. + is only used in numerics not character. Thus by adding them together, we will receive the error message: "non-numeric" argument.

iii:For the next two commands, either explain their results, or why they should produce errors.

```
z <- data.frame(z1="5",z2=7,z3=12)
z[1,2] + z[1,3]

## [1] 19

class(z[1,2])

## [1] "numeric"

class(z[1,3])

## [1] "numeric"</pre>
```

When characteric arguments are passed to data frame, they are numeric arguments, as in the class commands. Thus, we can use + operator to add two numeric values together.

Question 03:

Data Structure: Give R expressions that return the following matrices and vectors (i.e. do not construct them manually).

```
#i.
c(x <- 1:8, rev(x[-length(x)]))

## [1] 1 2 3 4 5 6 7 8 7 6 5 4 3 2 1

#ii
rep(1:5, c(1:5))

## [1] 1 2 2 3 3 3 4 4 4 4 5 5 5 5 5

#iii
x<- c(0,1,1,1,0,1,1,1,0)
matrix(x,nrow=3, ncol=3)

## [1,] [,2] [,3]
## [1,] 0 1 1
## [2,] 1 0 1
## [2,] 1 0 1
## [3,] 1 1 0</pre>
```

```
matrix(c(1^{(1:4)}, 2^{(1:4)}, 3^{(1:4)}, 4^{(1:4)}), nrow = 4, ncol = 4)
         [,1] [,2] [,3] [,4]
## [1,]
            1
                 2
                       3
## [2,]
                       9
                            16
            1
## [3,]
            1
                 8
                      27
                            64
## [4,]
```

Question 04:

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i:Let $h(x,n) = 1 + x + x^2 + \ldots + x^n = \sum_{i=0}^n x^i$. Write an R program to calculate h(x,n) using a for loop.

```
#A<- NULL
\#B \leftarrow rep(x, (n+1))
#for(i in 0:n){
  \#A[(i+1)] \leftarrow B[(i+1)]^i
#}
#sum(A)
```

iia:Find the sum of all the multiples of 3 or 5 below 1,000.

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```
x \leftarrow c(1:999)
sum(which(x\%3 == 0 | x\%5 == 0))
```

[1] 233168

iib:Find the sum of all the multiples of 4 or 7 below 1,000,000

```
x \leftarrow c(1:999999)
sum(as.numeric(which(x\%4 == 0 | x\%7 == 0)))
```

```
## [1] 178571071431
```

iii:Each new term in the Fibonacci sequence is generated by adding the previous two terms. By starting with 1 and 2, the first 10 terms will be (1,2,3,5,8,13,21,34,55,89). Write an R program to calculate the sum of the first 15 even-valued terms.

```
B \leftarrow c(1,2,rep(0,50))
for (i in 3:52){
B[i] \leftarrow B[i-1] + B[i-2]
evenB <- B[which(B \% 2 == 0)]
sum(evenB[1:15])
```

[1] 1485607536