Bios 6301: Assignment 2

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(informally) Due Tuesday, 20 September, 1:00 PM

50 points total.

This assignment won't be submitted until we've covered Rmarkdown. Create R chunks for each question and insert your R code appropriately. Check your output by using the Knit PDF button in RStudio.

- 1. Working with data In the datasets folder on the course GitHub repo, you will find a file called cancer.csv, which is 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. (18 points)
 - 1. Load the data set into R and make it a data frame called cancer.df. (2 points)

```
setwd("~/Desktop/FA 16/BIOS 6301/hw")
cancer.df<-read.csv("cancer.csv")
cancer.df<-data.frame(cancer.df)
attach(cancer.df)</pre>
```

2. Determine the number of rows and columns in the data frame. (2)

```
nrow(cancer.df)
## [1] 42120

ncol(cancer.df)
## [1] 8

2 Extract the names of the columns in cancer df (2)
```

3. Extract the names of the columns in cancer.df. (2)

```
## [1] "year" "site" "state" "sex" "race"
## [6] "mortality" "incidence" "population"
4. Report the value of the 3000th row in column 6. (2)
```

```
cancer.df[3000,6]
```

```
## [1] 350.69
```

names(cancer.df)

5. Report the contents of the 172nd row. (2)

```
## year site state sex race mortality
## 172 1999 Brain and Other Nervous System nevada Male Black 0
## incidence population
```

6. Create a new column that is the incidence rate (per 100,000) for each row.(3)

```
cancer.df[,'incrate']<-incidence/100000</pre>
```

7. How many subgroups (rows) have a zero incidence rate? (2)

```
sum(cancer.df['incrate']==0)
## [1] 23191
```

```
#23191 rows have zero incrate
```

8. Find the subgroup with the highest incidence rate.(3)

```
which(cancer.df['incrate'] == max(cancer.df['incrate']))
```

[1] 21387

```
#the 21387th subgroup has highest incrate
```

- 2. Data types (10 points)
 - 1. Create the following vector: $x \leftarrow c("5","12","7")$. 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. (4 points)

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

## [1] "7"

# no error; for characters, the first 'number' is compared to get max(x)=7
sort(x)

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

# no error; for characters, the first 'number' is primarily sorted "12" "5" "7"

#sum(x)
# error; cannot sum up elements of character strings</pre>
```

2. For the next two commands, either explain their results, or why they should produce errors. (3 points)

```
y <- c("5",7,12)

# define y to be a vector with the character string "5", "7" and "12"

#y[2] + y[3]

# error because R cannot add two elements of character strings
```

3. For the next two commands, either explain their results, or why they should produce errors. (3 points)

```
z \leftarrow \text{data.frame}(z1="5",z2=7,z3=12)
# define z to be a data frame with a row of numeric elements 5, 7 and 12
z[1,2] + z[1,3]
## [1] 19
# add up the 2nd and 3rd element of z, 7+12=19
```

3. Data structures Give R expressions that return the following matrices and vectors (*i.e.* do not construct them manually). (3 points each, 12 total)

```
1. (1,2,3,4,5,6,7,8,7,6,5,4,3,2,1)

c(1:8,7:1)

## [1] 1 2 3 4 5 6 7 8 7 6 5 4 3 2 1
2. (1,2,2,3,3,3,4,4,4,4,5,5,5,5,5)

rep(1:5,time=1:5)

## [1] 1 2 2 3 3 3 4 4 4 4 5 5 5 5 5
3. \begin{pmatrix} 0 & 1 & 1 \\ 1 & 0 & 1 \\ 1 & 1 & 0 \end{pmatrix}
```

1-diag(3)

[,1] [,2] [,3]
[1,] 0 1 1
[2,] 1 0 1
[3,] 1 1 0
4.
$$\begin{pmatrix}
1 & 2 & 3 & 4 \\
1 & 4 & 9 & 16 \\
1 & 8 & 27 & 64 \\
1 & 16 & 81 & 256 \\
1 & 32 & 243 & 1024
\end{pmatrix}$$

```
y=c(1:4)
x=matrix(c(y,y^2,y^3,y^4,y^5),nrow=5,byrow=T)
x
```

```
[,1] [,2] [,3] [,4]
##
## [1,]
          1
## [2,]
          1
               4
                     9
                         16
## [3,]
                   27
                         64
## [4,]
          1
                   81 256
               16
## [5,]
              32 243 1024
```

- 4. **Basic programming** (10 points)
 - 1. 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. (5 points)

```
h<-function(x,n) {
   h<-1
   for (i in 1:n) {
      h<-h+x^i
   }
   return(h)
}</pre>
```

- 1. If we list all the natural numbers below 10 that are multiples of 3 or 5, we get 3, 5, 6 and 9. The
 - 1. Find the sum of all the multiples of 3 or 5 below 1,000. (3, [euler1])

```
s<-0
for (i in 1:999) {
   if (i%%3==0 | i%%5==0) {
     s<-s+i
   }
}</pre>
```

[1] 233168

1. Find the sum of all the multiples of 4 or 7 below 1,000,000. (2)

```
t<-0
for (j in 1:999999) {
   if (j%%4==0 | j%%7==0) {
     t<-t+j
   }
}</pre>
```

[1] 178571071431

1. Each new term in the Fibonacci sequence is generated by adding the previous two terms. By starting w

```
x<-c(1,2)
for (i in 1:100) {
    x[i+2]<-x[i]+x[i+1]
}
y<-0
for (j in 1:45) {
    if (x[j]%%2==0) {
        y<-y+x[j]
    }
}</pre>
```

[1] 1485607536

Some problems taken or inspired by projecteuler.