## Bios 6301: Assignment 2

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

```
cancer.df<-data.frame(read.csv("cancer.csv"))</pre>
2. Determine the number of rows and columns in the data frame. (2)
dim(cancer.df)
## [1] 42120
#there are 42120 rows and 8 columns
3. Extract the names of the columns in `cancer.df`. (2)
colnames(cancer.df)
## [1] "year"
                     "site"
                                                "sex"
                                                             "race"
                                  "state"
                                  "population"
## [6] "mortality" "incidence"
4. Report the value of the 3000th row in column 6. (2)
cancer.df [3000,6]
## [1] 350.69
#get value:350.69
5. Report the contents of the 172nd row. (2)
cancer.df [172,]
```

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

## 172 1999 Brain and Other Nervous System nevada Male Black

73172

##

##

## 172

year

incidence population

0

site state sex race mortality

```
cancer.df[,"incidence_rate"]<-cancer.df[,"incidence"]/cancer.df[,"population"]*100000</pre>
```

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

```
sum(cancer.df[,"incidence_rate"]==0)
```

## [1] 23191

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

```
which.max(cancer.df[,"incidence_rate"])
```

## [1] 5797

- 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)

max(x):returned "7", not the max number 12. The reason is that x is a character vector, so the order depends on the first number of each element: 7>5>1 sort(x): returned ("12,"5,"7"), sort return the order of the vector, from first number of each element: 7>5>1 sum(x): produced an error "invalid type of arugment", because the argument should be numeric type

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

y <-c("5",7,12) y[2] + y[3] produced errors: non-numeric argument to binary operator y is a vector with character "5" and numeric element 7 and 12, after combination using c() the type of vector coerced to the least flexible form: character, then y[2] and y[3] are all characters, cannot by applied "+"

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

 $z \leftarrow data.frame(z1="5",z2=7,z3=12)$  z[1,2] + z[1,3] result is 19, because z[1,2] and z[1,3] are all of numeric type. Data.frame keep the original class of each element, not like c(). So can be applied "+"

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(seq(8), seq(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,times=1:5)
## [1] 1 2 2 3 3 3 4 4 4 4 5 5 5 5 5
3. $\begin{pmatrix}
  0 & 1 & 1 \\
  1 & 0 & 1 \\
  1 & 1 & 0 \\
\end{pmatrix}$
pmatrix<-matrix(1,nrow=3,ncol=3)</pre>
diag(pmatrix)<-0</pre>
4. $\begin{pmatrix}
  1 & 2 & 3 & 4 \\
  1 & 4 & 9 & 16 \\
  1 & 8 & 27 & 64 \\
  1 & 16 & 81 & 256 \\
  1 & 32 & 243 & 1024 \\
\end{pmatrix}$
a < -seq(4)
pmatrix<-rbind(a,a^2,a^3,a^4,a^5)</pre>
```

- 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) {
    s <- 0
    for (i in 0:n) {
        s= s + x^i
    }
    return(s)
}</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 1:1000-1){
      if (i %%3 ==0|i %% 5==0){

            s=s+i
            }
}
s</pre>
```

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

```
s<-0
for (i in 1:1000000-1){
    if (i %%4 ==0|i %% 7==0){
        s=s+i
        }
}
s</pre>
```

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

```
f<-numeric(50)
j<-1
f[1]<-1
f[2]<-2
s<-f[1]+f[2]
for (i in 3:50){
        f[i]<-f[i-2]+f[i-1]
        if (f[i] %%2 ==0 && j<15){
        s=s+f[i]
        j=j+1
        }
}</pre>
```

Some problems taken or inspired by projecteuler.