## Bios 6301: Homework 2

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Grade: 52/50

(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('~/Downloads/Bios6301-master/datasets')
x <- data.frame(read.csv('cancer.csv'))</pre>
```

```
2. Determine the number of rows and columns in the data frame. (2)
nrow(x)
## [1] 42120
ncol(x)
## [1] 8
  3. Extract the names of the columns in cancer.df. (2)
names(x)
## [1] "year"
                      "site"
                                    "state"
                                                   "sex"
                                                                 "race"
## [6] "mortality" "incidence"
                                    "population"
  4. Report the value of the 3000th row in column 6. (2)
x[3000,6]
## [1] 350.69
  5. Report the contents of the 172nd row. (2)
x[172,]
                                          site state sex race mortality
## 172 1999 Brain and Other Nervous System nevada Male Black
       incidence population
## 172
                        73172
  6. Create a new column that is the incidence rate (per 100,000) for each row.(3)
```

JC Grading - 1 For incidence rate above should be incidence / population \* 100000

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

incirate <- x\$incidence/100000
x <- data.frame(x, incirate)</pre>

```
nrow(x[incirate==0,])
## [1] 23191
8. Find the subgroup with the highest incidence rate.(3)
x[incirate==max(x$incirate),]
##
         year
                site
                          state
                                    sex race mortality incidence population
## 21387 2002 Breast california Female White
                                                3463.74
                                                            18774
                                                                    13690681
##
         incirate
## 21387 0.18774
```

JC Grading - 1 syntax is fine but answer is incorrect b/c of how incidence rate was calculated

- 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)
sort(x)
sum(x)
```

Since the number within quote mark will become characters, x is not a combination of numbers but characters. sum(x) returns error message since it is impossible to sum() characters. max(x) can be used on characters but it only takes the first letter in each element into account ('12' becomes '1'), and so as sort(x).

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

```
y <- c("5",7,12)
y[2] + y[3]
```

It returns errors. The first element in y is a character. The c() function will switch all the elements to character if there are both numbers and characters. Therefore, y[2] becomes the character '7' and y[3] becomes the character '12', and it is impossible to add them together.

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

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

It returns 19. As opposed to c() function, data.frame() function switchs characters to numbers when both characters and numbers are included in the data frame. So the second and third elements can be added up to 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(seq(8), seq(7, by=-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)
  c(1,rep(2,2),rep(3,3),rep(4,4),rep(5,5))
  ## [1] 1 2 2 3 3 3 4 4 4 4 5 5 5 5 5
  m1 <- matrix(1, nrow=3, ncol=3)
  diag(m1) <- 0
  m1
  ##
            [,1] [,2] [,3]
  ## [1,]
               0
                     1
  ## [2,]
             1
                      0
  ## [3,]
             1
                      1
         \begin{bmatrix} 1 & 4 & 9 & 16 \\ 1 & 8 & 27 & 64 \\ 1 & 16 & 81 & 256 \end{bmatrix} 
  m2 <- matrix(ncol=4, nrow=5, byrow=T)</pre>
  for (i in 1:4) {
  m2[,i]=i^(1:5)
  }
  m2
  ##
             [,1] [,2] [,3] [,4]
  ## [1,]
                      2
                            3
                1
  ## [2,]
                      4
                1
                            9
                                 16
  ## [3,]
                           27
                                 64
                1
  ## [4,]
                1
                     16
                           81
                               256
  ## [5,]
                1
                     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 = 0
                 for (i in 0:n) {
                 h = h + x^i
                 }
```

- 2. If we list all the natural numbers below 10 that are multiples of 3 or 5, we get 3, 5, 6 and 9. The sum of these multiples is 23. Write an R program to perform the following calculations. (5 points)
  - 1. Find the sum of all the multiples of 3 or 5 below 1,000. (3, euler1)

```
x = 0
for (i in 1:999) {
```

```
if (i%%3==0 | i%%5==0) {
x = x+i
}
}
x
```

## [1] 233168

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

```
y = 0
for (i in 1:(1e6-1)) {
if (i%%4==0 | i%%7==0) {
y = y+i
}
}
y
```

## ## [1] 178571071431

3. 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. (5 bonus points, euler2)

```
z = c(1, 2)
even = c(2)
for (i in 3:1000) {
    z[i] = z[i-1] + z[i-2]
    if (z[i]%%2==0 & length(even) < 16) {
       even = c(even, z[i])
    }
}
sum(even)</pre>
```

## ## [1] 6293134512

JC Bonus +4 Included one extra term (the first term 2 was included before loop)

Some problems taken or inspired by projecteuler.