Assignment 1

Group 1

November 22th, 2020

Importing libraries

```
library(markovchain)
library(matlib)
```

Functions to solve the problems

```
matrixpower <- function(M,k) {
  if(dim(M)[1]!=dim(M)[2]) return(print("Error: matrix M is not square"))
  if (k == 0) return(diag(dim(M)[1]))
  if (k == 1) return(M)
  if (k > 1) return(M %*% matrixpower(M, k-1))
}
```

Problem 1

a)

Markov chain criteria:

- 1- The probability of being in a state only depends on the previous state.
- 2- It's a stochastic process.
- X = The chain hits state j at time n

 X_n is the scenario at time n

All states have finite expected return times and are communicated with each other, also the MC is irreducible, therefore its stationary distribution is **unique**.

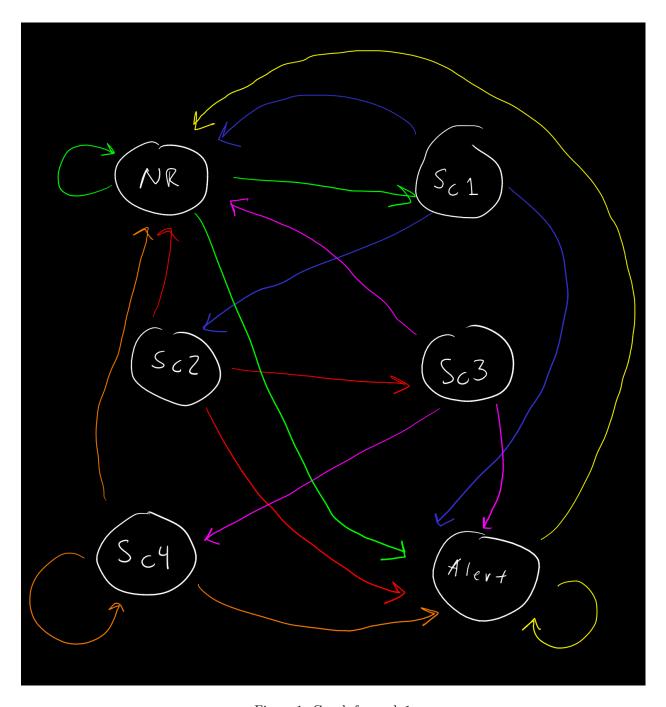


Figure 1: Graph for prob.1

b)

We have first calculated the relative frequencies manually.

```
load('PollutionMadrid.RData')
data <- X[1,]
mat <- matrix(rep(0,36), nrow=6, byrow=T)</pre>
for (i in 1:length(data)) {
 if (data[i] == "Alert") {
   data[i] = 1
 } else if (data[i] == "NR") {
   data[i] = 2
 } else if (data[i] == "Sc1") {
   data[i] = 3
 } else if (data[i] == "Sc2") {
   data[i] = 4
 } else if (data[i] == "Sc3") {
   data[i] = 5
 } else if (data[i] == "Sc4") {
   data[i] = 6
 }
}
data <- as.numeric(data)</pre>
for (i in 1:length(data)) {
 mat[data[i],data[i+1]] = mat[data[i],data[i+1]] + 1
mat[data[1460],data[1]] = mat[data[1460],data[1]] + 1
tbl <- table(data)
for (i in 1:length(tbl)) {
 mat[i,] = mat[i,]/tbl[i]
}
mat
##
                       [,2]
                                  [,3]
             [,1]
                                           [,4]
                                                     [,5]
                                                               [,6]
## [2,] 0.00000000 0.9529851 0.04701493 0.0000000 0.0000000 0.00000000
## [3,] 0.01587302 0.4920635 0.00000000 0.4920635 0.0000000 0.0000000
## [4,] 0.00000000 0.5806452 0.00000000 0.0000000 0.4193548 0.0000000
## [5,] 0.23076923 0.3846154 0.00000000 0.0000000 0.0000000 0.3846154
## [6,] 0.00000000 0.5555556 0.00000000 0.0000000 0.0000000 0.4444444
```

We then tested using the *markovchain* package in order to confirm our results.

```
data <- X[1,]
markovchainFit(data)$estimate
## MLE Fit
   A 6 - dimensional discrete Markov Chain defined by the following states:
   Alert, NR, Sc1, Sc2, Sc3, Sc4
##
   The transition matrix (by rows) is defined as follows:
##
                         NR
                                  Sc1 Sc2
                                               Sc3
             Alert
                                                         Sc4
0.00000000 \ 0.9529851 \ 0.04701493 \ 0.0 \ 0.0000000 \ 0.0000000
## Sc1
        0.01612903 0.4838710 0.00000000 0.5 0.0000000 0.0000000
## Sc2
        0.00000000 \ 0.5806452 \ 0.00000000 \ 0.0 \ 0.4193548 \ 0.0000000
## Sc3
        0.23076923 0.3846154 0.00000000 0.0 0.0000000 0.3846154
## Sc4
        0.00000000 0.5555556 0.00000000 0.0 0.0000000 0.4444444
```

What can you say of the comparison of your estimates and the possible transitions between states that you had argued in part a

According to our probabilities shown in the graph. There are 3 arrows with probability 0. This is due to the fact that in the data there are zero transitions from $Sc2 \rightarrow Alert$, $Sc4 \rightarrow Alert$, $NR \rightarrow Alert$, $Alert \rightarrow Alert$.

This is logical given that it is very unlikely to hit an alert state. Unlike the rest of the states.

Later it will be shown that there's a unique stationary distribution (see 1d).

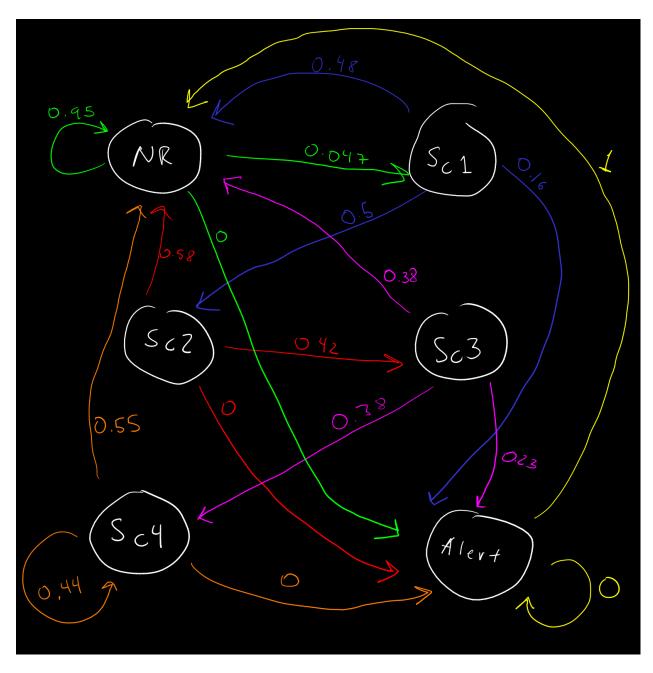


Figure 2: Graph with probabilities for problem 1 (b)

c)

Given that the first state of the chain is NR. We see the following 7 states:

```
data[1:7]
```

```
## [1] "NR" "NR" "NR" "NR" "NR" "NR" "NR"
```

And we calculate the probability as follows:

```
mat[2,2]^7
```

```
## [1] 0.713843
```

We can see the probability is 0.713843

d)

We can see that because we have a unique solution to the system, we have a unique stationary distribution.

```
stationary_dist <- function(P) {
    dim = sqrt(length(P))
    mat = matrix(P,nrow=dim, byrow=T)
    A = mat - diag(dim)
    b = c(1,rep(0,dim-1))
    A[,1] <- rep(1,dim)

    print("The solution is the following:")
    pi <- matlib::Solve(A, b, fractions = TRUE)
}
stationary_dist(mat)</pre>
```

```
## [1] "The solution is the following:"
## x1
                                 1/6
##
     x2
                     17201996/66847
##
       xЗ
                         39543/3224
##
         x4
                             645/104
                           2059/744
##
           x5
##
             x6
                          3725/1681
```

e)

Taking the 120th power of our transition matrix we get the following:

```
matrixpower(mat,120)
```

```
## [,1] [,2] [,3] [,4] [,5] [,6]
## [1,] 0.002739726 0.9178082 0.04315068 0.02123288 0.00890411 0.006164384
## [2,] 0.002739726 0.9178082 0.04315068 0.02123288 0.00890411 0.006164384
## [3,] 0.002739726 0.9178082 0.04315068 0.02123288 0.00890411 0.006164384
## [4,] 0.002739726 0.9178082 0.04315068 0.02123288 0.00890411 0.006164384
## [5,] 0.002739726 0.9178082 0.04315068 0.02123288 0.00890411 0.006164384
## [6,] 0.002739726 0.9178082 0.04315068 0.02123288 0.00890411 0.006164384
```

Problem 2

a)

We set up the following system of equations:

$$\sum_{i=0} \pi_i P_{i,0} = \pi_1$$

$$\sum_{i=1} \pi_i = 1$$

$$(1-p)\pi_1 = \pi_2 \dots (1-p)\pi_{n-2} = \pi_{n-1} \dots$$

For the first equation, each $P_{i,0} = p$, therefore:

$$\sum_{i=0} P_{i,0} \pi_i = \pi_1 \Rightarrow p \sum_{i=1} \pi_i = \pi_1$$

$$p = \pi_1$$

$$(1-p)p = \pi_2 (1-p)^2 p = \pi_3 \dots (1-p)^{n-1} p = \pi_n \dots$$

Then, we get:

Because our MC is an irreducible infinite state MC, we have a unique stationary distribution π , $\pi_i = \frac{1}{\mu_i}$ and all states have expected finite return times then we have:

$$E[T_i|X_0=i] = \mu_i = \frac{1}{\pi_i}$$

b)

Because it has a unique stationary distribution, it can only have one communication class (it is irreducible), all states are recurring states and there is no transient state.

$\mathbf{c})$

```
mc <- function(p, sequences ,steps) {
    n <- 100
    MarkovChain <- matrix(rep(0,sequences^2), nrow=sequences, byrow=TRUE)
    MarkovChain[,1] <- p
    for (i in 1:sequences) {
        if (i == sequences) {
            MarkovChain[i,i] <- 0
        } else {
            MarkovChain[i,i+1] <- 1-p
        }
    }
    return(MarkovChain)
}</pre>
```