Networking Assignment 1

October 29, 2024

1 Data Preprocessing

Here we first load the dataset:

```
[1]: # Load the necessary packages
     install.packages('ergm')
     install.packages('sna')
     library(ergm)
     library(network)
     library(sna)
    Installing package into '/usr/local/lib/R/site-library'
    (as 'lib' is unspecified)
    Installing package into '/usr/local/lib/R/site-library'
    (as 'lib' is unspecified)
    Loading required package: network
    'network' 1.18.2 (2023-12-04), part of the Statnet Project
    * 'news(package="network")' for changes since last version
    * 'citation("network")' for citation information
    * 'https://statnet.org' for help, support, and other information
    'ergm' 4.7.1 (2024-10-07), part of the Statnet Project
    * 'news(package="ergm")' for changes since last version
    * 'citation("ergm")' for citation information
    * 'https://statnet.org' for help, support, and other information
    'ergm' 4 is a major update that introduces some backwards-incompatible
    changes. Please type 'news(package="ergm")' for a list of major
    changes.
```

```
Loading required package: statnet.common

Attaching package: 'statnet.common'

The following objects are masked from 'package:base':
   attr, order

sna: Tools for Social Network Analysis

Version 2.8 created on 2024-09-07.
copyright (c) 2005, Carter T. Butts, University of California-Irvine
For citation information, type citation("sna").
Type help(package="sna") to get started.
```

Read the information / data from the files given.

```
[2]: # Reading the adjacency matrices of the networks
list.files()
attributes=read.table("Krackhardt-High-Tech_nodes.txt",header=TRUE)
nodes=as.matrix(read.table("Krackhardt-High-Tech_multiplex.edges",header=FALSE))
colnames(nodes)=c("layerID","IDi","IDj","weight")
```

1. 'Krackhardt-High-Tech_layers.txt' 2. 'Krackhardt-High-Tech_multiplex.edges' 3. 'Krackhardt-High-Tech_nodes.txt' 4. 'sample_data'

Below we build the adjacency matrix for advice, friendship and Reports to, respectively.

2 Task 1

2.1 Question 1

Build a QAP to test if friendship and advice relations correlate. Use at least 5,000 permutations for reporting the results.

```
[4]: set.seed(0)
     permutations=10000
     #build the QAP test
     g1=netlogit(advice.matrix,f.matrix, rep=permutations,nullhyp='qapy')
     g1$names=c('intercept','friendship')
     summary(g1)
    Network Logit Model
    Coefficients:
                           Exp(b)
                                     Pr(\langle b) Pr(\rangle b) Pr(\langle b|)
               Estimate
    intercept -0.3689075 0.6914894 0.0177 0.9838 0.0177
    friendship 0.7255825 2.0659341 0.9846 0.0160 0.0326
    Goodness of Fit Statistics:
    Null deviance: 582.2436 on 420 degrees of freedom
    Residual deviance: 568.4124 on 418 degrees of freedom
    Chi-Squared test of fit improvement:
             13.83123 on 2 degrees of freedom, p-value 0.0009921717
    AIC: 572.4124
                           BIC: 580.4929
    Pseudo-R^2 Measures:
```

(Dn-Dr)/(Dn-Dr+dfn): 0.03188159

```
\label{eq:contingency} $$ (Dn-Dr)/Dn: 0.02375505$$ Contingency Table (predicted (rows) x actual (cols)):
```

Actual Predicted 0 1 0 188 130 1 42 60

Total Fraction Correct: 0.5904762

Fraction Predicted 1s Correct: 0.5882353 Fraction Predicted Os Correct: 0.591195

False Negative Rate: 0.6842105 False Positive Rate: 0.1826087

Test Diagnostics:

Null Hypothesis: qapy Replications: 10000 Distribution Summary:

intercept friendship Min -4.11110 -4.68751 -0.946291stQ -2.23715 Median -1.67982 -0.03266 Mean -1.69969 0.01118 3rdQ -1.23271 1.10913 0.67277 Max 4.87127

```
[5]: # Formatting and exporting the results
    res1 <- summary(g1)
    expRes1 <- cbind(res1$coefficients, exp(res1$coefficients),
    res1$se, res1$pgreqabs)
    colnames(expRes1) <- c("ESt.", "exp(Est.)", "s.e.", "p-value")
    rownames(expRes1) <- res1$names
    #write.csv(expRes, "resQAP_Q1.csv")

# Exporting results in tex
library(xtable)
    xtable(expRes1,digits=3)</pre>
```

2.1.1 Conclusion:

The parameter related to the friendship nomination is significant and positive indicating that a seeking advice tie A_{ij} is more likely when a manager i nominate the second manager j as a friend. Specifically, the odds of manager i seeking advice from manager j when a manager i nominate the second manager j as a friend are 2.066 times greater than the odds of this kind of tie when manager i dose not nominate j as a friend.

2.2 Question 2 & 3

Hypotheses: 1. A friendship nomination is more likely between a pair of managers within the same department. 2. Senior managers are less likely to nominate friends. 1. A friendship nomination is more likely between a pair of managers of a similar age.

The friendship nomination X_{ij} ties are the dependent variable. The explanatory variables that allow testing the hypotheses above are defined in the following.

The first Hypothese states the relations friendship nomination and working at the same department are associated. The corresponding variable (dyadic covariates) is an indicator functions taking value 1 if two lawyers have the same characteristics, and 0 otherwise.

Hp. 1:
$$Z_{ij} = \begin{cases} 1, & \text{if } \{i,j\} \text{ in the same department.} \\ 0, & \text{otherwise} \end{cases}$$

The second and third Hypotheses concern the dependence of a tie on the seniority of the manager and the differences between ages of the node pair. Thus, these variables take the form:

Hp. 2:
$$Z_{ij} = \text{Tenure}_i \text{Hp.}$$
 3: $Z_{ij} = |\text{Age}_i - \text{Age}_j|$

First, we create vectors containing the values of the explanatory variables and then create the corresponding matrices.

```
[6]: # Hp. 1: A friendship nomination is more likely between a pair of managers_within the same department
department <- attributes[['nodeDepartment']]
same_department.matrix <- outer(department,department,"==")*1

# Hp. 2: senior managers are less likely to nominate friends.
tenure.matrix = matrix(0, nrow=numID, ncol=numID)
for(i in 1:numID){
    for(j in 1:numID){
        #The value at position (i, j) is tenure[i]
        tenure.matrix[i, j] = attributes[["nodeTenure"]][i]
    }
}

# Hp. 3: a friendship nomination is more likely between a pair of managers of a_u similar age.
```

```
age_dif.matrix = matrix(0, nrow=numID, ncol=numID)
     for(i in 1:numID){
       for(j in 1:numID){
         # The value at position (i, j) is |age[i] - age[j]|
         age_dif.matrix[i, j] = __
      →abs(attributes[["nodeAge"]][i]-attributes[["nodeAge"]][j])
     }
[7]: zm <- list(advice.matrix, same_department.matrix, tenure.matrix, age_dif.matrix)
     g2<-netlogit(f.matrix,zm, rep=permutations,nullhyp='qapspp')</pre>
     g2$names<-c('intercept', 'advice', 'same department', 'tenure', 'age dif')
     summary(g2)
    Network Logit Model
    Coefficients:
                    Estimate
                                Exp(b)
                                           Pr(\langle b) Pr(\rangle b) Pr(\langle b|)
                    -1.97460863 0.1388156 0.0049 0.9951 0.0049
    intercept
    advice
                     0.78779191 2.1985365 0.9854 0.0146 0.0299
    same_department 1.18388201 3.2670323 1.0000 0.0000 0.0000
                     0.04876536 1.0499740 0.8707 0.1293 0.2315
    tenure
                    -0.04722080 0.9538768 0.0315 0.9685 0.0938
    age_dif
    Goodness of Fit Statistics:
    Null deviance: 582.2436 on 420 degrees of freedom
    Residual deviance: 425.0383 on 415 degrees of freedom
    Chi-Squared test of fit improvement:
             157.2054 on 5 degrees of freedom, p-value 0
    AIC: 435.0383
                          BIC: 455.2395
    Pseudo-R^2 Measures:
            (Dn-Dr)/(Dn-Dr+dfn): 0.2723561
            (Dn-Dr)/Dn: 0.2699993
    Contingency Table (predicted (rows) x actual (cols)):
             Actual
                        1
    Predicted
                  0
            0
                306
                       85
            1
                 12
                       17
            Total Fraction Correct: 0.7690476
            Fraction Predicted 1s Correct: 0.5862069
            Fraction Predicted Os Correct: 0.7826087
```

False Negative Rate: 0.8333333

False Positive Rate: 0.03773585

Test Diagnostics:

Null Hypothesis: qapspp Replications: 10000 Distribution Summary:

```
intercept
                    advice same department
                                              tenure
                                                       age_dif
Min
       -7.553955 -5.244037
                                 -3.775461 -6.366783 -4.757958
       -2.248520 -1.015424
1stQ
                                 -0.796976 -1.758954 -1.163178
Median -0.885677 0.028261
                                 -0.043526 -0.178677 -0.086863
Mean
       -0.995801 0.007856
                                  0.001489 0.023970
                                                     0.028714
                                  0.770403 1.660610
3rdQ
        0.373022 1.001290
                                                     1.115361
Max
        4.853989 5.098259
                                  4.393634 7.652889
                                                     6.113808
```

```
[8]: # Formatting and exporting the results
    res2 <- summary(g2)
    expRes2 <- cbind(res2$coefficients, exp(res2$coefficients),
    res2$se, res2$pgreqabs)
    colnames(expRes2) <- c("ESt.", "exp(Est.)", "s.e.", "p-value")
    rownames(expRes2) <- res2$names
    #write.csv(expRes, "resQAP_Q1.csv")

# Exporting results in tex
library(xtable)
    xtable(expRes2,digits=3)</pre>
```

		ESt.	$\exp(\text{Est.})$	s.e.	p-value
		<dbl></dbl>	<dbl $>$	<dbl $>$	<dbl $>$
	intercept	-1.97460863	0.1388156	0.33380046	0.0049
A xtable: 5×4	advice	0.78779191	2.1985365	0.25125909	0.0299
	$same_department$	1.18388201	3.2670323	0.26909036	0.0000
	tenure	0.04876536	1.0499740	0.01645553	0.2315
	age_dif	-0.04722080	0.9538768	0.01651161	0.0938

2.2.1 Conclusion

The parameters of the same department is significant and positive suggesting that friendship nomination is more likely between a pair of managers within the same department. Specifically, the odds of a friendship nomination tie between managers working in the same department are 1.184 times greater than the odds of a tie between managers working in different departments, holding all the other variables constant. Thus, the data supports Hypothesis 1.

The parameters of tenure and age differences are significantly different from 0 at a significance level $\alpha = 0.05$. Thus, the data does not support Hypotheses 2 and 3.

2.3 Question 4 & 5

Question: Could you think of another hypothesis that could be tested using QAPs? State your hypothesis and provide the corresponding statistic.

Solution: Yes, here we propose a hypothsis that: If a person reports to another person as part of his function in organization, then the first person more likely nominate the second as a friend. The code & statistic are shown below:

Network Logit Model

```
Coefficients:
```

```
Estimate Exp(b) Pr(<=b) Pr(>=b) Pr(>=|b|) intercept -1.90160438 0.1493288 0.0061 0.9939 0.0061 advice 0.68802778 1.9897874 0.9631 0.0369 0.0709 same_department 1.03989079 2.8289081 0.9981 0.0019 0.0020 tenure 0.04711714 1.0482448 0.8682 0.1318 0.2441 age_dif -0.04845630 0.9526990 0.0330 0.9670 0.0928 report_matrix 0.83217434 2.2983106 0.9460 0.0540 0.0992
```

Goodness of Fit Statistics:

```
Null deviance: 582.2436 on 420 degrees of freedom
Residual deviance: 422.5441 on 414 degrees of freedom
Chi-Squared test of fit improvement:

159.6995 on 6 degrees of freedom, p-value 0
AIC: 434.5441 BIC: 458.7856
Pseudo-R^2 Measures:

(Dn-Dr)/(Dn-Dr+dfn): 0.2754867

(Dn-Dr)/Dn: 0.274283
Contingency Table (predicted (rows) x actual (cols)):
```

```
Actual
Predicted 0 1
0 308 85
1 10 17
```

Total Fraction Correct: 0.7738095 Fraction Predicted 1s Correct: 0.6296296 Fraction Predicted 0s Correct: 0.783715 False Negative Rate: 0.8333333 False Positive Rate: 0.03144654

Test Diagnostics:

Null Hypothesis: qapspp Replications: 10000 Distribution Summary:

```
intercept
                    advice same_department
                                              tenure
                                                       age_dif report_matrix
                                 -4.037302 -6.444961 -4.712527
                                                                   -3.044494
Min
       -7.505426 -4.835998
1stQ
       -2.171582 -1.014912
                                 -0.791513 -1.773822 -1.219133
                                                                   -0.649187
Median -0.782573 -0.018775
                                 -0.016344 -0.207976 -0.121814
                                                                   -0.011678
Mean
       -0.897303 0.004527
                                  0.006007 -0.008699 -0.011613
                                                                    0.014251
3rdQ
        0.434792 1.020886
                                  0.752871 1.628084 1.070960
                                                                    0.666118
Max
        4.713405 5.656976
                                  4.692464 7.872279 5.791957
                                                                    3.752562
```

```
[10]: # Formatting and exporting the results
  res3 <- summary(g3)
  expRes3 <- cbind(res3$coefficients, exp(res3$coefficients),
  res3$se, res3$pgreqabs)
  colnames(expRes3) <- c("ESt.", "exp(Est.)", "s.e.", "p-value")
  rownames(expRes3) <- res3$names
  #write.csv(expRes, "resQAP_Q1.csv")

# Exporting results in tex
  library(xtable)
  xtable(expRes3,digits=3)</pre>
```

		ESt.	$\exp(\text{Est.})$	s.e.	p-value
		<dbl></dbl>	<dbl $>$	<dbl $>$	<dbl $>$
A xtable: 6×4	intercept	-1.90160438	0.1493288	0.33580013	0.0061
	advice	0.68802778	1.9897874	0.25944396	0.0709
	$same_department$	1.03989079	2.8289081	0.28547655	0.0020
	tenure	0.04711714	1.0482448	0.01655627	0.2441
	age_dif	-0.04845630	0.9526990	0.01669396	0.0928
	$report_matrix$	0.83217434	2.2983106	0.53090641	0.0992

2.3.1 Conclusion

The parameter of report relation is significantly different from 0 at a significance level $\alpha = 0.05$. Thus, the data does not support our Hypothsis mentioned above.

3 Task 2: Simulation from an ERGM

3.1 Question 1

Some parts of the code are missing as denoted by the chunk code - - - MISSING - - -. Implement these in the R script, and include comments explaining what your code is doing.

The solution is below:

```
[11]: | #function calculating the given statics of an adjacency matrix:
      stat=function(net){
          z1=0
          z_2 = 0
          z3 = 0
          indegree=0
          nvertices <- nrow(net)</pre>
          for(k in 1:nvertices){
            for(q in 1:nvertices){
              z1=z1+net[k,q]
                                         #calculating the num of edges
              if(k<q){
                z2=z2+net[k,q]*net[q,k] #calculating reciprocity
              if(net[q,k]==1){
                                        #calculating indegree of node k
                indegree=indegree + 1
              }
            z3=z3+choose(indegree,2) #calculating 2-istar and resetting for next_
       \rightarrownode
            indegree=0
          return(list(z1,z2,z3))
      }
                                        _____
      # MHstep -----
      #' Simulate the next step of a network in Markov chain using Metropolis-Hasting
      #' The function `MHstep` simulates the the Metropolis-Hastings step that defines
      #' the Markov chain whose stationary distribution is the ERGM with
      #' edge, mutual and nodematch statistics
      #'
      #' Oparam net an object of class `matrix`. Adjacency matrix of the network.
      #' @param theta1 a numeric value. The value of the edge parameter of the ERGM.
      #' @param theta2 a numeric value. The value of the mutual parameter of the ERGM.
      #' Oparam theta3 a numeric value. The value of the istar(2) parameter of the
       \hookrightarrow ERGM.
      # '
      #' @return next state of the Markov Chain
```

```
#' @examples
#' MHstep(
\#' matrix(c(0, 1, 0, 0, 0, 0, 1, 1, 0), nrow = 3, ncol = 3),
\#' -log(0.5), log(0.4), log(.8)
#')
MHstep <- function(net, theta1, theta2, theta3){</pre>
  # Number of vertices in the network
 nvertices <- nrow(net)</pre>
  # Choose randomly two vertices, prevent loops \{i,i\} with replace = FALSE
 tie <- sample(1:nvertices, 2, replace = FALSE)</pre>
  i <- tie[1]
  j <- tie[2]
  # Compute the change statistics
              #creating the matrix with different i->j
 net2=net
  if (net[i,j]==0)
    net2[i,j]=1
  else
    net2[i,j]=0
  #calculating the statistics for the input
  stat1=stat(net)
  #computing the statistics for the changed graph
  stat2=stat(net2)
  # Compute the probability of the next state
  # according to the Metropolis-Hastings algorithm
  # computing both exponential
  stat1=as.double(stat1)
  stat2=as.double(stat2)
  exp1=exp(theta1*stat1[1]+theta2*stat1[2]+theta3*stat1[3])
  exp2=exp(theta1*stat2[1]+theta2*stat2[2]+theta3*stat2[3])
  # computing transition probability
 p=min(1,exp2/exp1)
  # Select the next state:
  #sample with probability p the change of state
  outcome=sample(c(0,1),size=1,prob=c(1-p,p))
```

```
if (outcome==1)
    net=net2
  # Return the next state of the chain
 return(net)
}
# Markov Chain simulation -----
#' The function MarkovChain simulates the networks from the ERGM with
#' edge, mutual and nodematch statistics
# '
#' @param net an object of class `matrix`. Adjacency matrix of the network.
#' Oparam theta1 a numeric value. The value of the edge parameter of the ERGM.
#' Oparam theta2 a numeric value. The value of the mutual parameter of the ERGM.
#' @param theta3 a numeric value. The value of the istar(2) parameter of the
 \hookrightarrow ERGM.
#' @param burnin an integer value.
#' Number of steps to reach the stationary distribution.
#' Oparam thinning an integer value. Number of steps between simulated networks.
\#' Oparam nNet an integer value. Number of simulated networks to return as
 \hookrightarrow output.
# '
#' @return a named list:
#' - netSim: an `array` with the adjancency matrices of the simulated
\hookrightarrownetworks.
    - statSim: a `matrix` with the value of the statistic defining the ERGM.
#' @examples
#' MarkovChain(
\#' matrix(c(0, 1, 0, 0, 0, 0, 1, 1, 0), nrow = 3, ncol = 3),
\#' -log(0.5), log(0.4), log(.8)
#')
MarkovChain <- function(</pre>
   net.
    theta1, theta2, theta3,
    burnin = 10000, thinning = 1000, nNet = 1000){
  # Burnin phase: repeating the steps of the chain "burnin" times
 nvertices <- nrow(net)</pre>
  burninStep <- 1 # counter for the number of burnin steps
  # Perform the burnin steps
 for(burninStep in 1:burnin){
   net=MHstep(net, theta1, theta2, theta3)
  # After the burnin phase we draw the networks
```

```
# The simulated networks and statistics are stored in the objects
  # netSim and statSim
  netSim <- array(0, dim = c(nvertices, nvertices, nNet))</pre>
  statSim <- matrix(0, nNet, 3)</pre>
  thinningSteps <- 0 # counter for the number of thinning steps
  netCounter <- 1 # counter for the number of simulated network
  while(netCounter<=nNet){</pre>
    while(thinningSteps<thinning){</pre>
      net=MHstep(net, theta1, theta2, theta3) #performing 1000 transitions
      thinningSteps=thinningSteps+1
    netSim[1:nvertices,1:nvertices,netCounter]=net #saving the current network,
 →and its statistics
    statSim[netCounter,1:3]=as.double(stat(net))
    netCounter=netCounter+1
    thinningSteps=0
                                                  #resetting the counter
  # Return the simulated networks and the statistics
  return(list(netSim = netSim, statSim = statSim))
}
```

3.2 Question 2

A member of your research team suggested that plausible estimates of the parameters of the ERGM above for the advice network are $\theta_1 = -2.76, \theta_2 = 0.68$ and $\theta_3 = 0.05$.

Solution is shown below: 1. Firstly, we build up a advice network to test the simulation result. The output shows that the number of edges, reciprocal dyads and 2-istar from actual Advice network are 190, 45 and 930, respectively.

'Actual number of edges: 190'

'Actual number of reciprocal dyads: 45'

'Actual number of 2-istar: 930'

2. Then we test our model with the given parameters: $\theta_1 = -2.76, \theta_2 = 0.68$ and $\theta_3 = 0.05$. The result is shown below. Here the columns of $\mathbf{MC_simulation\$statSim}$ are the number of edges, reciprocal dyads and 2-istar, respectively. By comparing with the actual number of

edges, reciprocal dyads and 2-istar, all of the results generated by this simulation are much smaller. Hence, we don't think that the suggested values of the parameters are plausible estimates.

```
[13]: t1=-2.76
    t2=0.68
    t3=0.05
    ad.matrix=matrix(0, numID,numID)

MC_simulation=MarkovChain(ad.matrix,t1,t2,t3)
#MC_simulation$statSim
```

```
[14]: avr.dens=mean(MC_simulation$statSim[,1])
    avr.rec=mean(MC_simulation$statSim[,2])
    avr.star=mean(MC_simulation$statSim[,3])
    paste('Average number of edges:', avr.dens)
    paste('Average number of reciprocal dyads:', avr.rec)
    paste('Average number of 2-istar:', avr.star)
```

3.3 Question 3

Guess better estimates of θ_1 , θ_2 and θ_3 based on the analysis in Question 2. Describe the procedure you used to obtain the guessed values.

Solution:

As we find out that all of the simulated results above are much smaller than the relevant actual values, we consider to increase the weights θ_1, θ_2 and θ_3 of these three variables. Below is the code illustrating the strategy we applied to obtain the guessed values:

1. Firstly, we slightly increased θ_1, θ_2 and θ_3 from (-2.76, 0.68, 0.05) to (-2.5, 0.8, 0.1).

```
t1_g1 = -2.5
t2_g1 = 0.8
t3_g1=0.1
ad_g1.matrix=matrix(0,numID,numID)
MC_g1=MarkovChain(ad_g1.matrix,t1_g1,t2_g1,t3_g1)
avr.dens=mean(MC_g1$statSim[,1])
avr.rec=mean(MC_g1$statSim[,2])
avr.star=mean(MC_g1$statSim[,3])
paste('Average number of edges:', avr.dens)
paste('Average number of reciprocal dyads:', avr.rec)
paste('Average number of 2-istar:', avr.star)
```

^{&#}x27;Average number of edges: 27.832'

^{&#}x27;Average number of reciprocal dyads: 1.646'

^{&#}x27;Average number of 2-istar: 18.132'

^{&#}x27;Average number of edges: 41.628'

'Average number of reciprocal dyads: 3.76'

'Average number of 2-istar: 42.693'

2. The simulated results of the first guess above show that the average number of edges, reciprocal dyads and 2-istar are all much smaller than the actual results (190, 45 and 930), we increase θ_1, θ_2 and θ_3 again to (-2, 0.9, 0.2) for the second guess.

```
[16]: t1_g2 = -2.0
    t2_g2 = 0.9
    t3_g2 = 0.2
    ad_g2.matrix=matrix(0,numID,numID)

MC_g2=MarkovChain(ad_g2.matrix,t1_g2,t2_g2,t3_g2)
    avr.dens=mean(MC_g2$statSim[,1])
    avr.rec=mean(MC_g2$statSim[,2])
    avr.star=mean(MC_g2$statSim[,3])
    paste('Average number of edges:', avr.dens)
    paste('Average number of reciprocal dyads:', avr.rec)
    paste('Average number of 2-istar:', avr.star)
```

'Average number of edges: 374.407'

'Average number of reciprocal dyads: 169.127'

'Average number of 2-istar: 3186.577'

3. The simulated results of the second guess above show that the average number of edges, reciprocal dyads and 2-istar are all much larger than the actual results (190, 45 and 930), we decrease the θ_1 , θ_2 and θ_3 from (-2, 0.9, 0.2) to (-2.2, 0.85, 0.14) for the third guess.

```
t1_g3 = -2.2
t2_g3 = 0.85
t3_g3 = 0.14
ad_g3.matrix=matrix(0,numID,numID)
MC_g3=MarkovChain(ad_g3.matrix,t1_g3,t2_g3,t3_g3)
avr.dens=mean(MC_g3$statSim[,1])
avr.rec=mean(MC_g3$statSim[,2])
avr.star=mean(MC_g3$statSim[,3])
paste('Average number of edges:', avr.dens)
paste('Average number of reciprocal dyads:', avr.rec)
paste('Average number of 2-istar:', avr.star)
```

'Average number of edges: 77.006'

'Average number of reciprocal dyads: 11.583'

'Average number of 2-istar: 155.65'

4. The simulated results of the third guess above show that the average number of edges, reciprocal dyads and 2-istar are all much smaller than the actual results (190, 45 and 930), we increase the θ_1 , θ_2 and θ_3 from (-2.2, 0.85, 0.14) to (-2.13, 0.87, 0.17) for the fourth guess.

```
[18]: t1_g4 = -2.13
    t2_g4 = 0.87
    t3_g4 = 0.17
    ad_g4.matrix=matrix(0,numID,numID)
    MC_g4=MarkovChain(ad_g4.matrix,t1_g4,t2_g4,t3_g4)
    avr.dens=mean(MC_g4$statSim[,1])
    avr.rec=mean(MC_g4$statSim[,2])
    avr.star=mean(MC_g4$statSim[,3])
    paste('Average number of edges:', avr.dens)
    paste('Average number of reciprocal dyads:', avr.rec)
    paste('Average number of 2-istar:', avr.star)
```

'Average number of edges: 144.254'

'Average number of reciprocal dyads: 34.278'

'Average number of 2-istar: 571.243'

5. The simulated results of the fourth guess above show that the average number of edges, reciprocal dyads and 2-istar are all slightly smaller than the actual results (190, 45 and 930), we slightly increase the θ_1 , θ_2 and θ_3 from (-2.13, 0.87, 0.17) to (-2.1225, 0.8721, 0.1721) for the fifth guess.

```
t1_g5 = -2.1225
t2_g5 = 0.8721
t3_g5 = 0.1721
ad_g5.matrix=matrix(0,numID,numID)
MC_g5=MarkovChain(ad_g5.matrix,t1_g5,t2_g5,t3_g5)
avr.dens=mean(MC_g5$statSim[,1])
avr.rec=mean(MC_g5$statSim[,2])
avr.star=mean(MC_g5$statSim[,3])
paste('Average number of edges:', avr.dens)
paste('Average number of reciprocal dyads:', avr.rec)
paste('Average number of 2-istar:', avr.star)
```

It seems that the average number of edges, reciprocal dyads and 2-istar for the fifth guess are similar to the actual results (190, 45 and 930), hence we choose $(\theta_1, \theta_2, \theta_3) = (-2.1225, 0.8721, 0.1721)$ for the final simulation.

4 Task 3: Estimation and interpretation of an ERGM

^{&#}x27;Average number of edges: 161.929'

^{&#}x27;Average number of reciprocal dyads: 41.89'

^{&#}x27;Average number of 2-istar: 713.624'