

Homework example

SOBA221 - Social Network teaching team

28/05/2020

RESEARCH QUESTIONS

RQ1: Is there sport-based homophily? (Are sporty people more likely to be friends?)

RQ2: Do discrepancies in reports of giving and getting study-related information reveal status inequality among students?

```
# Data loading
load('ActorVariables.RData')
load('NetworkVariables.RData')
load('TwoModeVariables.RData')

# Packages
library(sna)
library(igraph)
library(ggplot2)
```

Selection of the variables that I will be using

```
# RQ1
sport <- ActorVariables$sport_time # Sport time
friend <- NetworkVariables$friends_out_net # Out-friendship network
# Co-factors: gender
gender <- ActorVariables$gender

# RQ2
studyin <- NetworkVariables$study_info_in_net # Study info in
studyout <- NetworkVariables$study_info_out_net # Study info out
pop <- NetworkVariables$popular_net # Popularity
```

RQ 1

Descriptive statistics

In this stage, check the codebook carefully!

```
sport[sport == -1] <- NA # Prefer not answering (-1) to NA (missing)
sum_sport <- c(mean(sport,na.rm=TRUE),
               sd(sport,na.rm=TRUE),
               range(sport,na.rm=TRUE),
               sum(!is.na(sport)))
```

```

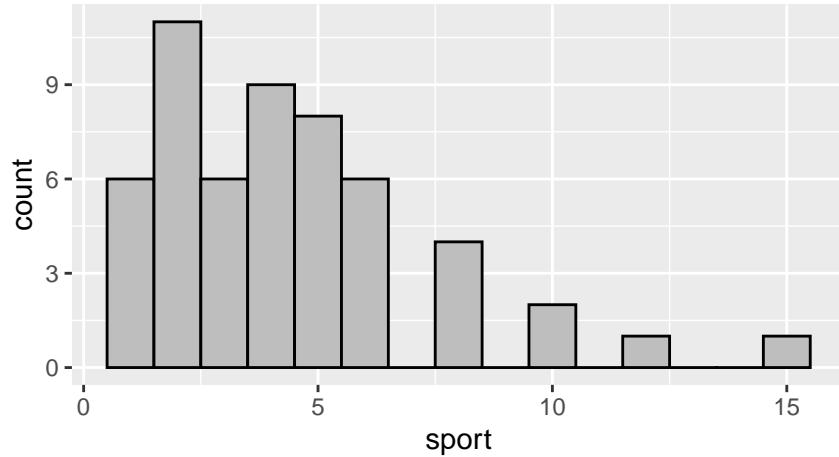
names(sum_sport) <- c('mean', 'sd', 'min', 'max', 'n')
sum_sport

##      mean        sd        min        max        n
##  4.388889  2.910078  1.000000 15.000000 54.000000

ggplot()+
  geom_histogram(aes(x=sport), binwidth=1, color='black', fill='grey')

## Warning: Removed 13 rows containing non-finite values (stat_bin).

```



```

c(density=gden(friend)*100,
  reciprocity=grecip(friend,measure='edgewise')*100,
  transitivity=gtrans(friend)*100)

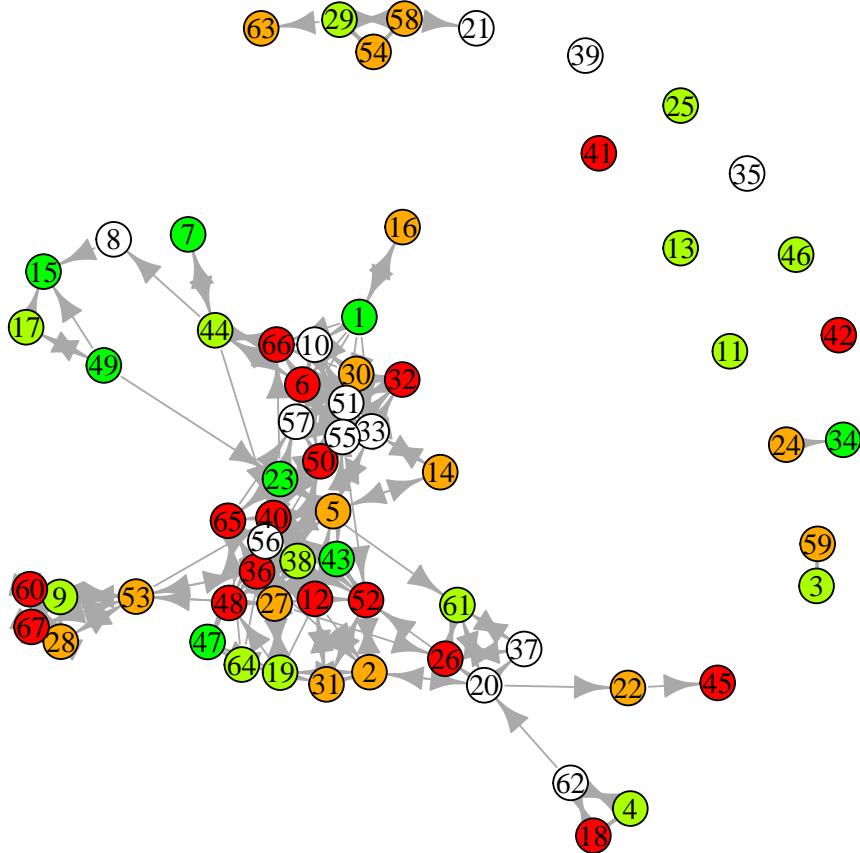
##           density reciprocity.Mut      transitivity
##           5.517865    68.852459     44.899738

diag(friend) <- NA # I removed the diagonal
rownames(friend) <- 1:nrow(friend)
colnames(friend) <- 1:ncol(friend)

gradient <- colorRampPalette(c('red','yellow','green'))
plot(graph_from_adjacency_matrix(friend, mode='directed'),
      layout=layout_nicely,
      vertex.size = 8.5,
      vertex.color=ifelse(is.na(sport), 'white',
                          ifelse(sport %in% 1:2, gradient(4)[1],
                                 ifelse(sport %in% 3:4, gradient(4)[2],
                                       ifelse(sport %in% 5:6, gradient(4)[3],
                                             gradient(4)[4])))),
      vertex.label.color='black',
      main='Friendship network coloured by how sporty each student is')

```

Friendship network coloured by how sporty each student is



```
# Gender
gender <- ifelse(gender == 1,'male','female')
gender <- as.factor(gender) # Defined as a factor
summary(gender)
```

```
## female    male
##      46      21
```

Remember that for QAP-family analyses, variables must be matrices (not vectors)

```
# Sport and gender as matrix
sport_diff <- abs(outer(sport,sport,'-')) # Absolute difference
gender_same <- 1*outer(gender,gender,'==') # Same gender
```

Analysis

Is there an association between friendship and sportiness?

```
# Matrix correlation  
gcor(friend,sport_diff)
```

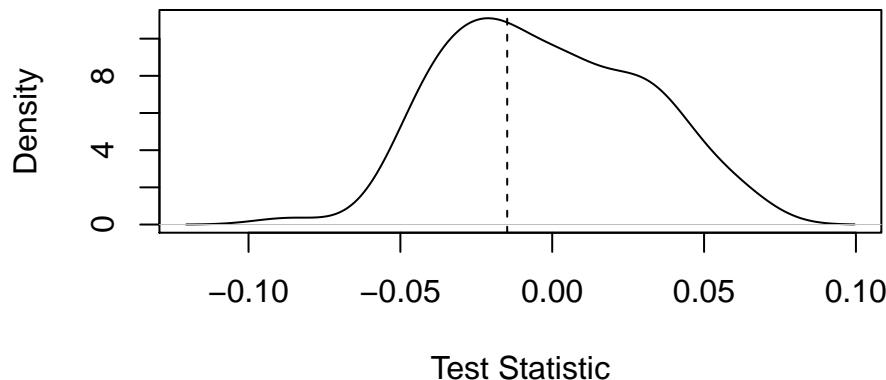
```
## [1] -0.01482314
```

Is it significant?

```
qap_corr <- qaptest(list(friend,sport_diff),  
                     gcor,g1=1,g2=2,reps=100)  
summary(qap_corr)
```

```
##  
## QAP Test Results  
##  
## Estimated p-values:  
##   p(f(perm) >= f(d)): 0.59  
##   p(f(perm) <= f(d)): 0.42  
##  
## Test Diagnostics:  
##   Test Value (f(d)): -0.01482314  
##   Replications: 100  
##   Distribution Summary:  
##     Min:      -0.08641616  
##     1stQ:     -0.02849747  
##     Med:      -0.005166018  
##     Mean:     -0.002222553  
##     3rdQ:     0.0243975  
##     Max:      0.06545579  
plot(qap_corr)
```

Estimated Density of QAP Replications



What if we adjust for reciprocity and same gender?

```
# QAP regression (friend as a function of reciprocity, gender and sport homophily)  
qap_mr <- netlogit(y=friend,x=list(t(friend),gender_same,sport_diff),
```

```

    nullhyp='qap', reps=100)
print(qap_mr)

##
## Network Logit Model
##
## Coefficients:
##             Estimate      Exp(b)      Pr(<=b)  Pr(>=b)  Pr(>=|b|)
## (intercept) -4.437653801  0.01182365 0.00     1.00     0.00
## x1           5.078691854 160.56387750 1.00     0.00     0.00
## x2           0.417416116   1.51803406 0.95     0.05     0.11
## x3           0.005307218   1.00532133 0.62     0.38     0.89
##
## Goodness of Fit Statistics:
##
## Null deviance: 3967.574 on 2862 degrees of freedom
## Residual deviance: 610.0099 on 2858 degrees of freedom
## Chi-Squared test of fit improvement:
##     3357.565 on 4 degrees of freedom, p-value 0
## AIC: 618.0099    BIC: 641.847
## Pseudo-R^2 Measures:
## (Dn-Dr)/(Dn-Dr+dfn): 0.5398392
## (Dn-Dr)/Dn: 0.8462512

```

RQ 2

Are there discrepancies in the first place? Check if “get” network is same as transpose of the “give” network:

```
table(studyout, t(studyin))
```

```

##
## studyout    0    1
##          0 4107 146
##          1    79 157

```

Apparently 146 “get” nominations are not confirmed by the giver, while another 79 “give” nominations are not confirmed by the recipient.

```
# Calculate difference matrix and take the absolute value:
discrepancy <- abs(studyout - t(studyin))
```

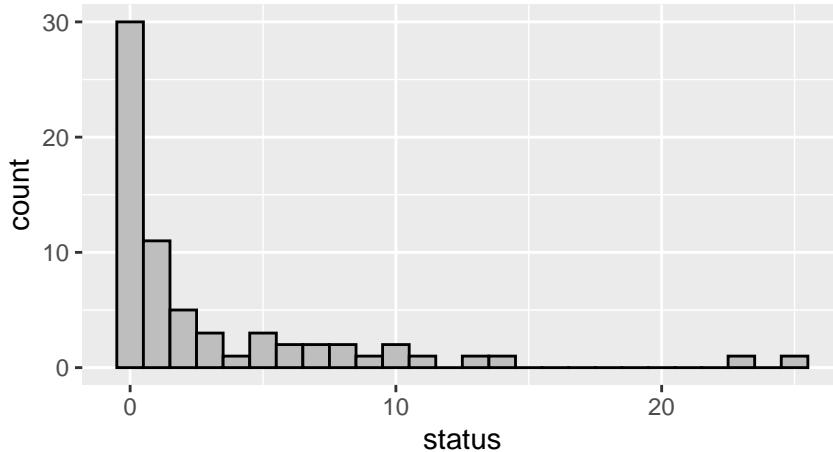
The question was if this goes together with status differences. So calculate status as the indegree of each student in the popularity network.

```

status <- sna::degree(pop, cmode="indegree")

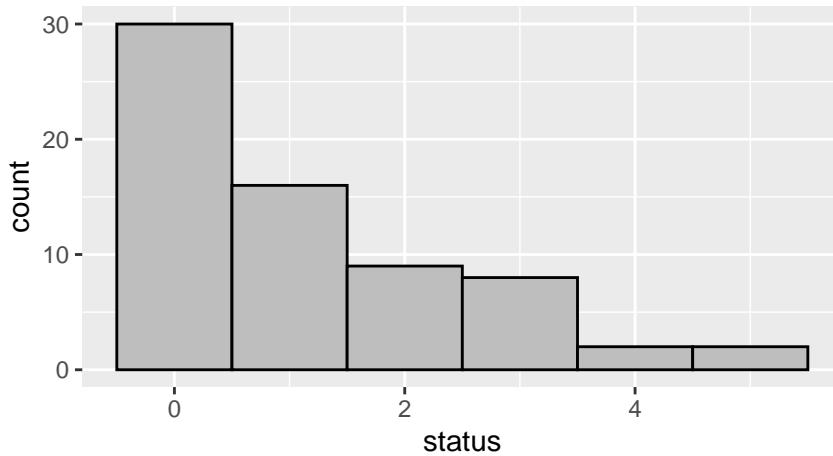
ggplot()+
  geom_histogram(aes(x=status), binwidth=1, color='black', fill='grey')

```



```
# Because this variable is very right-skewed, I will take its square root
status <- sqrt(status)
```

```
ggplot()+
  geom_histogram(aes(x=status), binwidth=1, color='black', fill='grey')
```



```
# And calculate status difference (sender minus receiver):
status_diff <- abs(outer(status, status, "-"))
```

Analysis

Is there an association between discrepancy and status differences

```
gcor(discrepancy, status_diff)
```

```
## [1] -0.007693562
```

Is it significant?

```
qap_corr <- qaptest(list(discrepancy, status_diff),
                     gcor, g1=1, g2=2, reps=100)
summary(qap_corr)
```

```
##
## QAP Test Results
```

```
##  
## Estimated p-values:  
##   p(f(perm) >= f(d)): 0.61  
##   p(f(perm) <= f(d)): 0.39  
##  
## Test Diagnostics:  
##   Test Value (f(d)): -0.007693562  
##   Replications: 100  
## Distribution Summary:  
##   Min:      -0.0446336  
##   1stQ:     -0.01518351  
##   Med:      -0.0001169933  
##   Mean:     -0.0006231299  
##   3rdQ:     0.01232847  
##   Max:      0.05050482  
plot(qap_corr)
```

Estimated Density of QAP Replications

