

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!

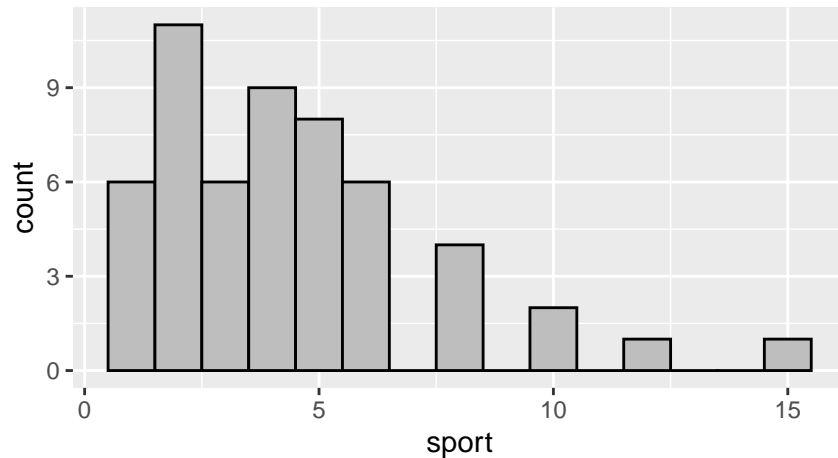
```
sport[sport == -1] <- NA # Prefer not answering (-1) to NA (missing)
sum_sport <- c(mean(sport,na.rm=TRUE),
               sd(sport,na.rm=TRUE),
               range(sport,na.rm=TRUE),
               sum(!is.na(sport)))
```

```
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).
```



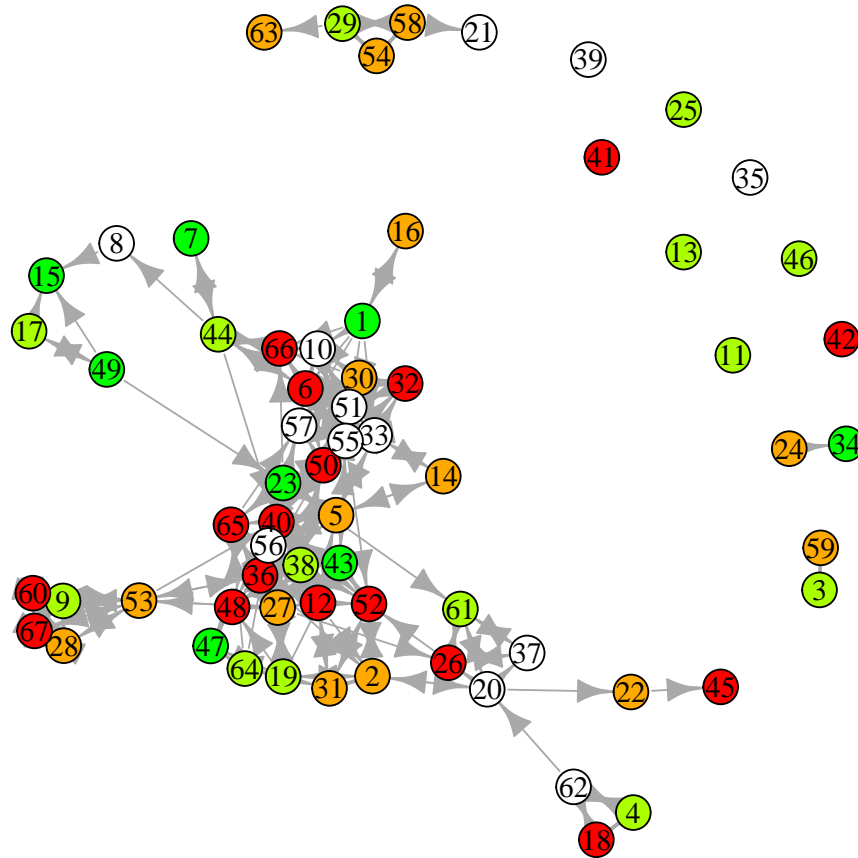
```
c(density=gden(friend)*100,
  reciprocity=grecip(friend,measure='edgewise')*100,
  transitivity=gtrans(friend)*100)
```

```
##      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)
```

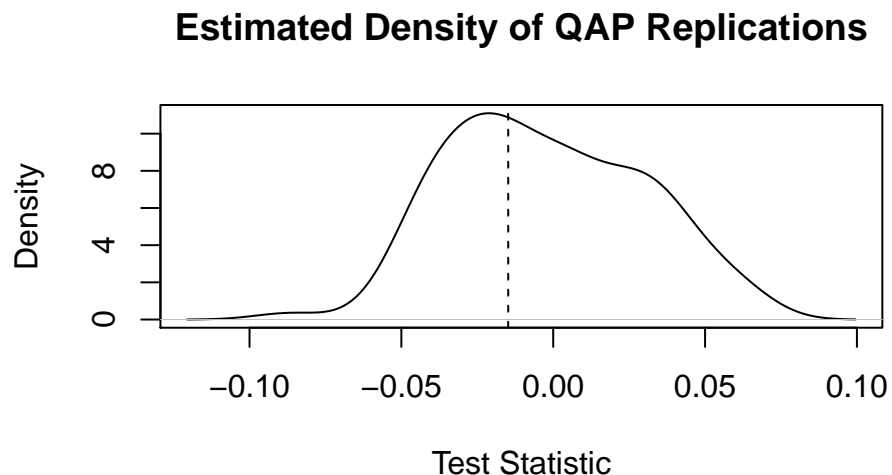
```
## [1] -0.01482314
```

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:     -0.00222553
##      3rdQ:      0.0243975
##      Max:       0.06545579
```

```
plot(qap_corr)
```



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))

```

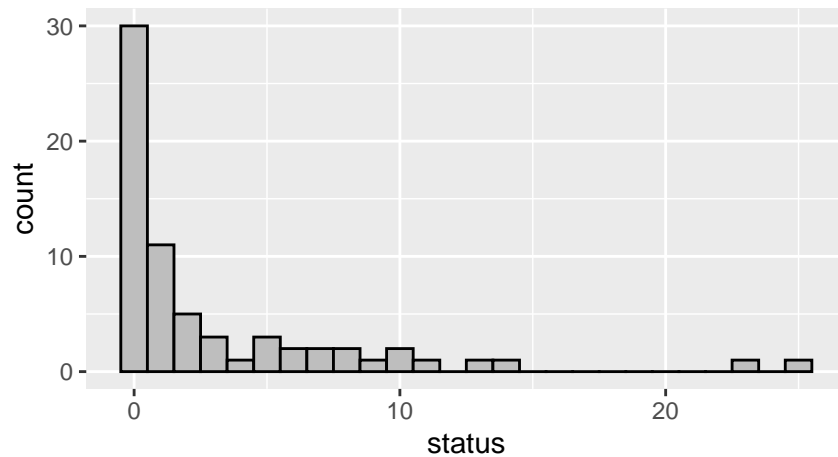
The question was if this goes together with status differences. So calculate status as the indegree of each student in the popularity network.

```

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

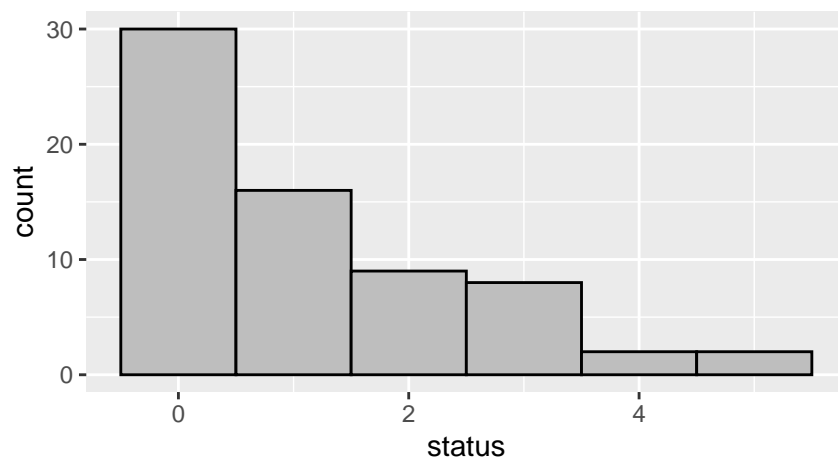
ggplot()+
  geom_histogram(aes(x=status),binwidth=1,color='black',fill='grey')

```



```
# 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)
```

