# Subgroups Chapter 8

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# Libraries and data used

#### Libraries

```
library(UserNetR)
library(igraph)

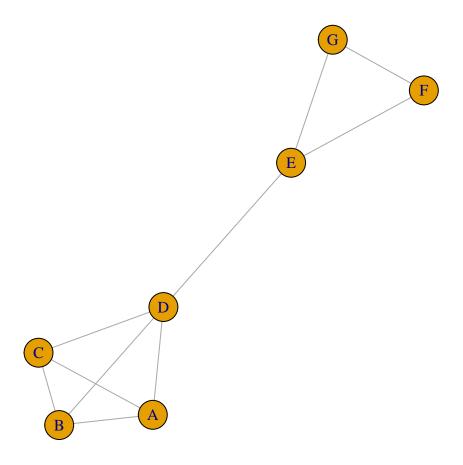
##
## Attaching package: 'igraph'
## The following objects are masked from 'package:stats':
##
## decompose, spectrum
## The following object is masked from 'package:base':
##
## union
library(intergraph)
```

#### Data

```
data(DHHS)
```

# **Social Cohesion**

## Cliques



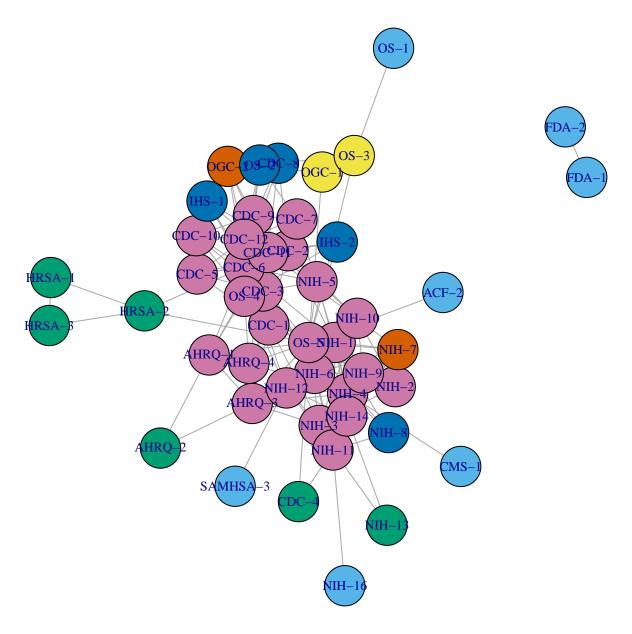
```
## [[3]]
## + 3/7 vertices, named, from 3a3dc47:
## [1] A B C
##
## [[4]]
## + 4/7 vertices, named, from 3a3dc47:
## [1] A B C D
##
## [[5]]
## + 3/7 vertices, named, from 3a3dc47:
## [1] A B D
## [[6]]
## + 3/7 vertices, named, from 3a3dc47:
## [1] A C D
maximal.cliques(clqexmp,
                min = 3)
## [[1]]
## + 3/7 vertices, named, from 3a3dc47:
## [1] E F G
##
## [[2]]
## + 4/7 vertices, named, from 3a3dc47:
## [1] A B D C
largest.cliques(clqexmp)
## [[1]]
## + 4/7 vertices, named, from 3a3dc47:
## [1] D A B C
V(clqexmp)[unlist(largest.cliques(clqexmp))]
## + 4/7 vertices, named, from 3a3dc47:
## [1] D A B C
g25 <- erdos.renyi.game(25,
                        75,
                         type = "gnm")
g50 <- erdos.renyi.game(50,
                         150,
                         type = "gnm")
g100 <- erdos.renyi.game(100,
                          300,
                          type = "gnm")
g500 <- erdos.renyi.game(500,
                          1500,
                          type = "gnm")
g1000 <- erdos.renyi.game(1000,
                           3000,
                           type = "gnm")
g5000 <- erdos.renyi.game(5000,
                           15000,
                           type = "gnm")
g10000 <- erdos.renyi.game(10000,
```

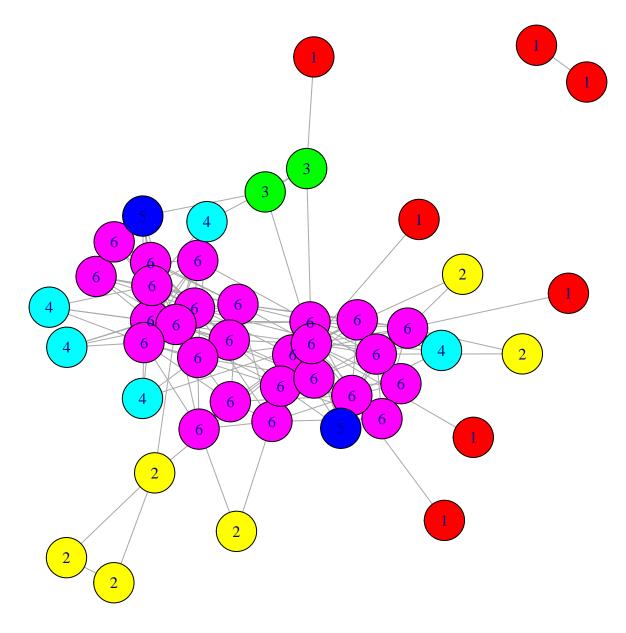
```
30000,
                             type = "gnm")
nodes \leftarrow c(25,
           50,
           100,
           500,
            1000,
           5000,
           10000)
lrgclg <- c(clique.number(g25),</pre>
             clique.number(g50),
             clique.number(g100),
             clique.number(g500),
             clique.number(g1000),
             clique.number(g5000),
             clique.number(g10000))
numclq <- c(length(cliques(g25,</pre>
                            min = 3)),
             length(cliques(g50,
                            min = 3)),
             length(cliques(g100,
                            min = 3)),
             length(cliques(g500,
                            min = 3)),
             length(cliques(g1000,
                            min = 3)),
             length(cliques(g5000,
                            min = 3)),
             length(cliques(g10000,
                            min = 3)))
clqinfo <- data.frame(</pre>
  Nodes = nodes,
  Largest = lrgclg,
  number = numclq)
clqinfo
     Nodes Largest number
##
## 1
        25
                  4
                        42
        50
                  3
                        40
## 2
## 3
      100
                  3
                        35
## 4
       500
                  3
                        39
## 5 1000
                  3
                        31
## 6 5000
                  3
                        29
## 7 10000
                  3
                        42
k-Cores
```

```
data(DHHS)
iDHHS <- asIgraph(DHHS)
graph.density(iDHHS)</pre>
```

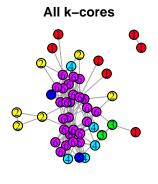
```
## [1] 0.312369
```

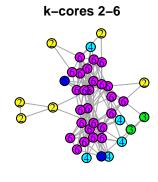
```
iDHHS <- subgraph.edges(iDHHS,</pre>
                         E(iDHHS)[collab > 2])
graph.density(iDHHS)
## [1] 0.1533688
coreness <- graph.coreness(iDHHS)</pre>
table(coreness)
## coreness
## 1 2 3 4 5 6
## 7 6 2 5 2 26
maxCoreness <- max(coreness)</pre>
maxCoreness
## [1] 6
Vname <- get.vertex.attribute(iDHHS,</pre>
                                name = "vertex.names",
                               index = V(iDHHS))
V(iDHHS)$name <- Vname</pre>
V(iDHHS)$color <- coreness + 1</pre>
op \leftarrow par(mar = rep(0,4))
plot(iDHHS,
vertex.label.cex = 0.8)
```

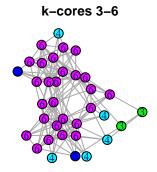


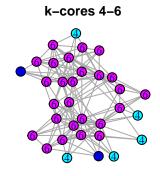


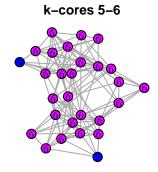
```
vids = which(coreness > 5))
lay <- layout.fruchterman.reingold(iDHHS)</pre>
op \leftarrow par(mfrow = c(3,2),
          mar = c(3, 0, 2, 0))
plot(iDHHS1_6,
     layout = lay,
     main = "All k-cores")
plot(iDHHS2_6,
     layout = lay[which(coreness > 1), ],
     main = "k-cores 2-6")
plot(iDHHS3_6,
     layout = lay[which(coreness > 2), ],
     main = "k-cores 3-6")
plot(iDHHS4_6,
     layout = lay[which(coreness > 3), ],
     main = "k-cores 4-6")
plot(iDHHS5_6,
     layout = lay[which(coreness > 4), ],
     main = "k-cores 5-6")
plot(iDHHS6_6,
     layout = lay[which(coreness > 5), ],
     main = "k-cores 6-6")
```

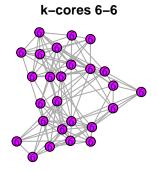












par(op)