Analyzing Social Networks with R

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Overview

- Software
- Three "Types" of network analysis and an example of each
 - Descriptive analysis and visualization
 - Data network of IV drug users
 - Antecedent analysis
 - Data organizational collaboration after an earthquake
 - Consequence analysis
 - Data teacher networks in 21 schools
- Available online resources for more information

Network Analysis Software

Packages in R

- statnet
 - sna
 - network
 - ergm
- igraph

Other programs

- UCINET
- ORA
- Pajek
- NodeXL
- libSNA
- C-IKNOW
- many more...

Visualization tools

- Netdraw
- Visone
- Tulip
- Gephi

Real advantage with R is that network analysis, statistical analysis, and visualization are integrated into one program.

Types of network analysis

Descriptive

 Visualization of the network along with summary statistics at the individual, dyadic, and whole network levels.

Antecedent

 Attempts to understand the micro-processes at work in forming the observed network.

Consequence

 Measures the implication of different network positions or structures.

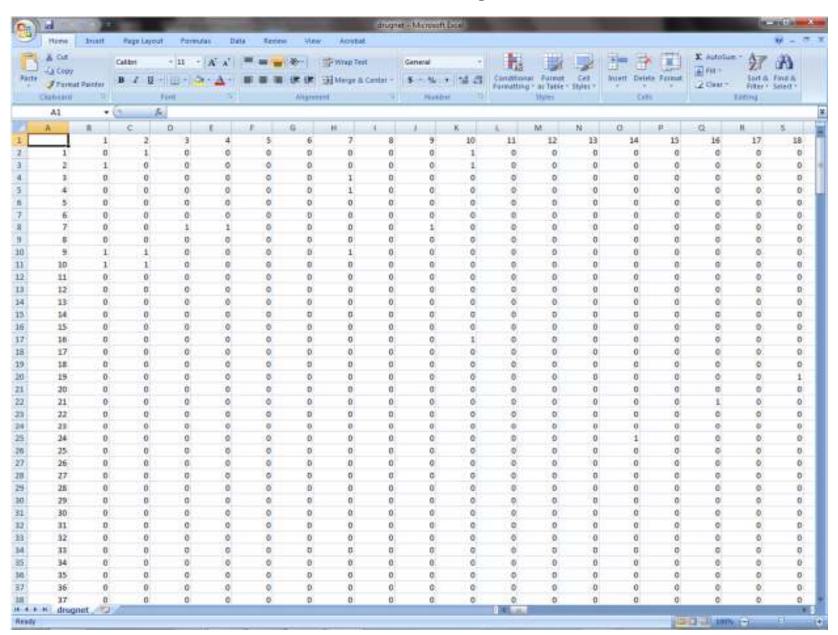
Descriptive Analysis

- For any system under study we may have two types of data: behavioral/relational and attribute.
- We can also construct nodal attributes based on the relational data, for example centrality measures.
- Two simple measures of centrality are:
 - Degree the total number of connections.
 - Betweenness the number of shortest paths from one node to another that pass through a given node.

Goal: identify key members in a needle sharing population to diffuse safe practices

Raw network data

Data = social network of IV drug users in Hartford, CT



Load data as a network object

Relational data = 293 actors and 337 ties.

```
library("statnet")
drug=read.csv(file="drugnet.csv", header=TRUE, row.names=1)
drug=as.matrix(drug)#to make certain the data is a matrix
drugn=as.network(drug, matrix.type="adjacency", directed=TRUE) #create a
network object
```

 Attribute data = gender and ethnicity; plus we will calculate some centrality measures

```
#read in the attribute data
drugatt=read.csv(file="drugnet_att.csv", header=TRUE, row.names=1)
drugatt=as.data.frame(drugatt)
#calculate some additional attributes
deg1=degree(drugn)
bet1=betweenness(drugn)
drugatt$bet=bet1#add the measures to our existing attribute data
drugatt$deg=deg1
```

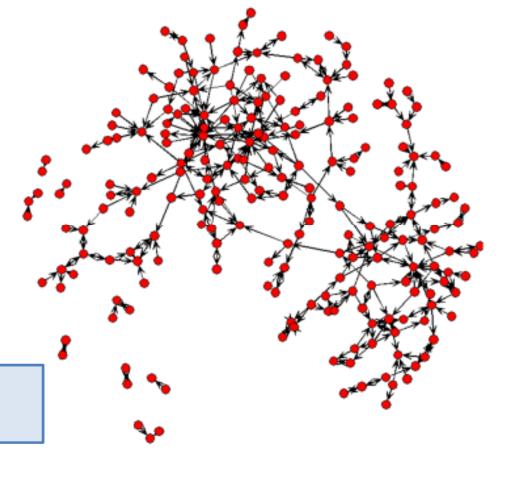
IV Drug Users in Hartford, CT

Attribute Data

head(drugatt)

	Ethnicity	Gender	HasTie	bet	deg
1	1	1	1	4	5
2	1	1	1	6	6
3	1	1	1	14	4
4	1	2	1	19	5
5	3	1	1	0	1
6	3	1	1	2	2

Relational Data

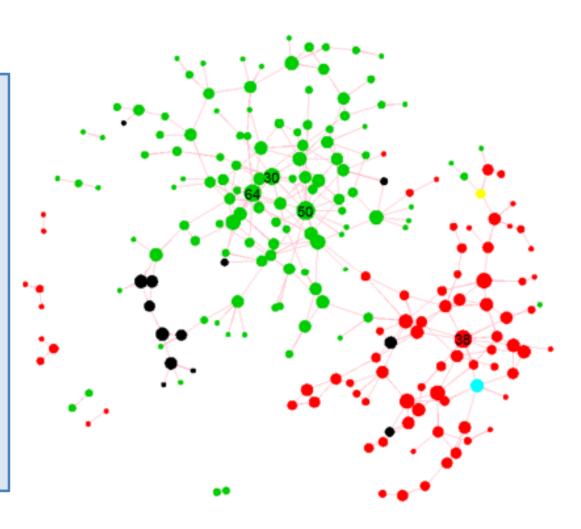


#plot the IV users Network

plot.network(drugn, mode = "fruchtermanreingold",
displayisolates = FALSE)

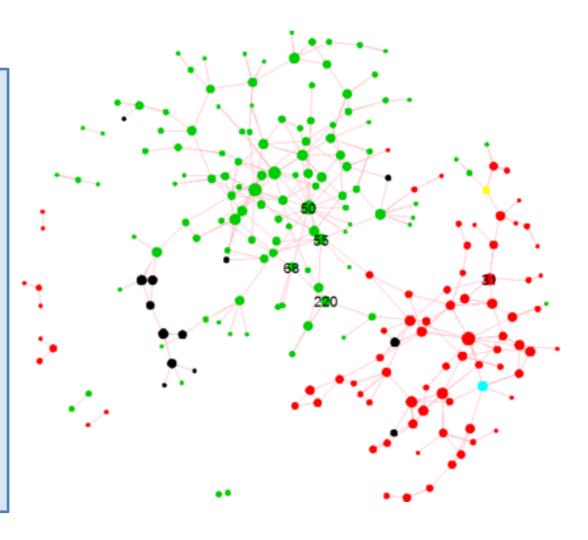
Scale nodes based on degree centrality

```
#Define a label set that is dependent on
degree centrality
label1=vector()
 for (i in 1:length(deg1))
 label1[i]=ifelse
(deg1[i] >= 10,row.names(drugatt[i,]),NA)
#Plot as network object, using high degree
nodes as labels
plot.network(drugn, mode =
"fruchtermanreingold",
vertex.col=drugatt$Ethnicity, label=label1,
displayisolates=FALSE,
label.cex=1.1, usearrows=FALSE,
edge.col="pink",
vertex.cex=log10(deg1+1), label.pos=5,
vertex.border=drugatt$Ethnicity,
edge.lwd=.25)
```



Scale nodes based on betweenness

```
#Define a label set that is dependent on
betweenness centrality
label2=vector()
for (i in 1:length(bet1))
label2[i]=ifelse
(bet1[i]>1800,row.names(drugatt[i,]),NA)
#plot as network object, using high between
nodes as labels
plot.network(drugn, mode =
"fruchtermanreingold",
vertex.col=drugatt$Ethnicity,
label=label2,displayisolates=FALSE,
label.cex=1.1, usearrows=FALSE,
edge.col="pink",
vertex.cex = logb((deg1+1.1), base=7),
label.pos=5,
vertex.border=drugatt$Ethnicity,
edge.lwd=.25)
```

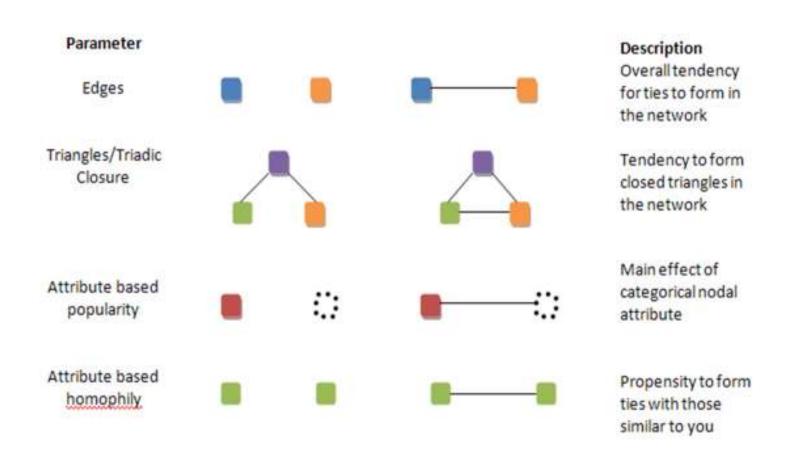


Antecedent Analysis

- Many recent statistical and computational advances in network analysis have been in the area of exponential random graph models (ERGMs).
- These models allow a researcher to investigate the organizing principals that give rise to an observed network structure.
- Even though the network under study may be a single representation of an evolving system, the patterns of network ties act as the structural signature of the network and provide evidence from which we may infer something of the social processes that built the network (Robins, 2011, p. 484).

Goal: identify the processes and factors affecting organizational collaboration during disasters

Examples of organizing principals in an undirected graph



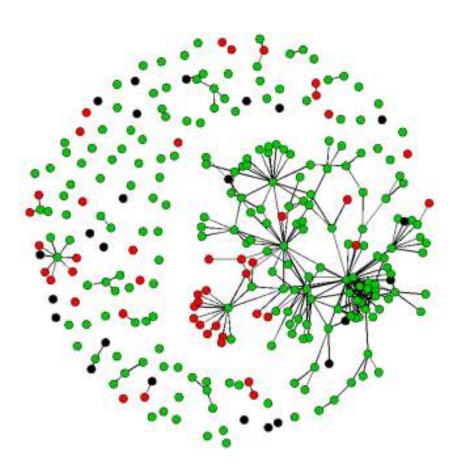
Response network to the 2009 Padang earthquake

Attribute data

	Fund2	Juris2
1	1	1
2	1	2
3	2	2
4	1	3
5	3	1
6	1	3

#read in a Pajek data file
padang=read.paj("Padang_Pajek.net")
#read in the attributes from a text file
att=read.delim("Padang_Attrib.txt", header=TRUE)
#create a network object and assign the attributes to
the nodes
padang2=as.network(padang2, vertex.attr=att,
directed=FALSE)
#plot the network, with vertex colors defined by
funding type
plot(padang2, vertex.col='Fund2')

Relational Data



Assume a random process

```
# a completely random process
rmodel=ergm(padang2~edges)
summary(rmodel)

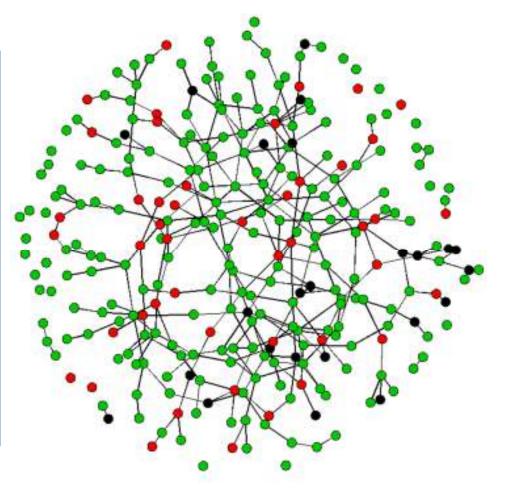
Monte Carlo MLE Results:
    Estimate Std. Error MCMC % p-value
edges -4.98450    0.06029    NA <1e-04 ***

gden(padang2) #graph density of the observed
network (observed ties/total possible ties)
[1] 0.006796712

#simulate networks based on the random model
rmodel.sim=simulate(rmodel, nsim=10)
#take a look at a simulated network
plot(rmodel.sim$networks[[1]], vertex.col='Fund2')

#this ERG model is equivalent to generating a random
graph (function in R is rgraph) with the same tie
```

probability as found in the observed network



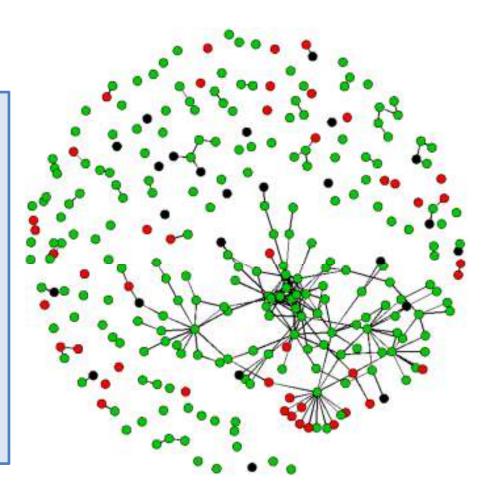
A more realistic model

#in addition to an edge term, we can pose other factors as important in network formation. For example, the node factor term captures the propensity for nodes of a particular type to form ties.

model5=ergm(padang2~edges + nodefactor("Fund2") + nodefactor("Juris2") + nodematch("Fund2", diff=T) + nodematch("Juris2", diff=T)+ isolates + gwesp(0.30, fixed=T))

summary(model5)#if you wanted to see the specific parameter values and level of significance

#simulate some networks based on the model
model5.sim=simulate(model5, nsim=10)
#plot one of the simulated networks
plot(model5.sim\$networks[[1]],vertex.col='Fund2')

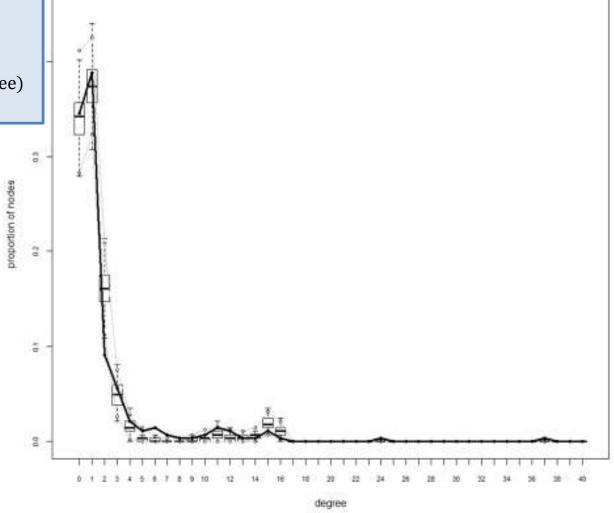


#Goodness of fit to test our model parameters. We can test this by picking a network statistic that is not in the model, and comparing the value of this statistic in the observed network to the distribution of values we obtained in the simulated networks

#lets look at degree

model5.gofdeg=gof(model5~degree)
plot(model5.gofdeg)

Goodness-of-fit diagnostics



Consequence Analysis

- Used to investigate the association between network variables and various group and/or individual level outcomes through the use of econometric models.
- Can also measure and estimate the social influence of alters' belief on ego through a class of models known as network autocorrelation models.
 - These models are predicated on the belief that individuals connected within a social network transmit interpersonal influence on one another.
 - Leenders (2002) states that an actor may form a belief based on his or her own intrinsic values as well as based on the beliefs of his or her alters.

$$y = \rho_1 W_1 y + \chi \beta + \varepsilon$$

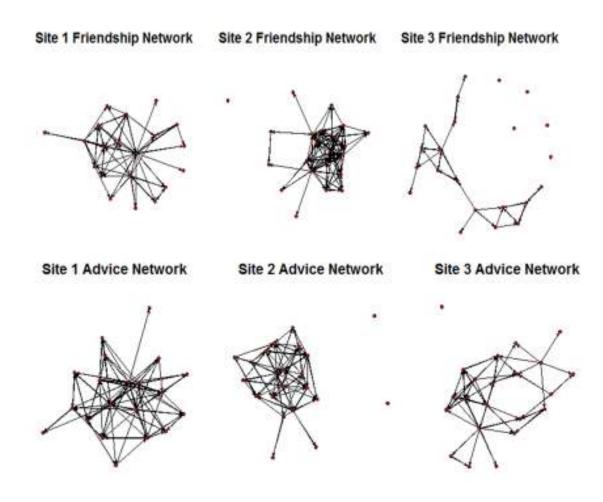
Goal: identify the influence of peer self-efficacy on a teacher's self-efficacy

Teacher social interactions in 21 schools

Attribute data

- Age
- Race
- Grade Level
- Education
- Years Teaching
- Efficacy
- Commitment
- Trust
- Reflective Dialogue
- Collective Responsibility
- Instructional Leadership
- Etc...

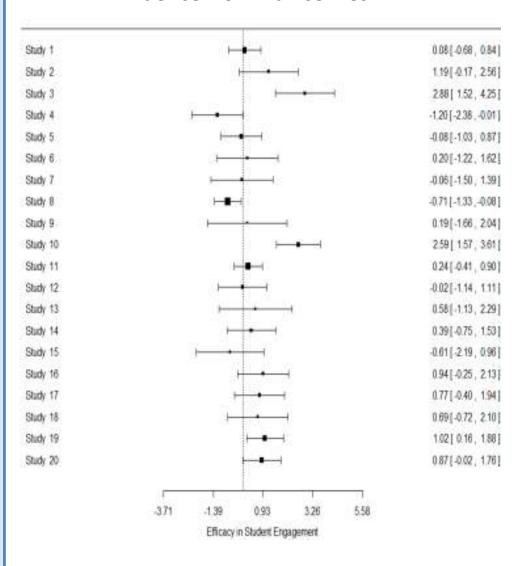
Relational Data



Analyzing peer influence

```
#run a network autocorrelation model for each school:
the model uses a set of actor attributes along with a set of
advice and friendship relations to predict a teacher's sense of
self efficacy with regard to student engagement; both
attributes and networks are stored in lists where each school
is a component
lnam_se=list()
for (i in c(1:18,20,21))#site 19 dropped from study
lnam se[[i]]=lnam(SelfEfficacy [[i]],
cbind(IndVar1[[i]], IndVar2[[i]], IndVar3[[i]]),
W1=list(advicenet[[i]], friendnet[[i]]))
lnam_se_sum=lapply(lnam_se, summary)#store the summary
results in a list
#pull out the influence effect of the advice network in each of
the models
rho1=vector()
for (i in c(1:18,20,21))
rho1[i]=lnam se sum[[i]]$rho1[[1]]
#pull out the standard error of the influence effect
rho1se=vector()
for (i in c(1:18,20,21))
rho1se[i]=lnam_se_sum[[i]]$rho1.se[[1]]
#create a simple dataset of model results
se results=as.data.frame(cbind(rho1, rho1se))
#use a forest plot to display the influence effects in each of the
schools
library("metafor")
forest(se results$rho1, sei=se results$rho1se, xlab="Efficacy
in Student Engagement")
```

Influence from Advice Ties



Meta-analysis

 Move from network models to standard metaanalysis techniques to pool the results across the schools. Produce a population level effect.

	Advice Network		Friendship Network		
	ρ	<i>p</i> -value	ρ	<i>p</i> -value	
Efficacy					
Student Engagement	0.3236	0.0000	-0.4608	0.0000	

Some online resources

- Steve Borgatti's webpage: http://www.steveborgatti.com/
- Dai Shizuka's homepage: https://sites.google.com/site/daishizuka/toolkits
- Stanford SNA labs in R: http://sna.stanford.edu/rlabs.php
- Network analysis with igraph: http://igraph.sourceforge.net/igraphbook/
- Statnet resources: https://statnet.csde.washington.edu/trac

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