Advanced Network Analysis

A Local Structure Graph Model

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Readings

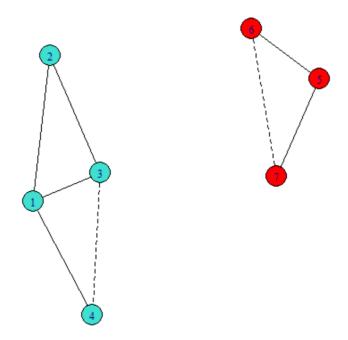
- Emily Casleton, Daniel Nordman, and Mark S. Kaiser. A local structure model for network analysis. *Statistics and Its Interface*, 2(10), 2017.
- Olga V. Chyzh and Mark S. Kaiser. Network analysis using a local structure graph model. *Political Analysis*, 27(4):397{414, 2019.

Motivation

- Do network edges form in reaction to (anticipation) of one another?
 - Did your ex start dating so soon because she didn't want you to start dating before her?
 - Do countries form alliances in response to their rivals or allies doing the same?
 - Do political parties form coalitions to balance against other (potential) coalitions?

Formal Motivation

- Do edges among red nodes affect the formation of edges among turquoise nodes?
- If nodes 6 and 7 formed and edge, would that make it more likely that nodes 3 and 4 form an edge?



The Estimator

- Suppose i is a potential edge in a network of potential edges (realized and unrealized);
- Then $s_i=(u_i,v_i)$ is i's location in Cartesian space.
- Denote the binary random variable, $y(s_i)=y_i$, so that:

$$y(s_i) = egin{cases} 1 & ext{if edge } s_i ext{ is present} \ 0 & ext{if edge } s_i ext{ is absent.} \end{cases}$$

- Define i's neighbors as $N_i = \{s_j : s_j \text{ is a neighbor of } s_i\}$.
- Make a Markov assumption of conditional spatial independence:

$$f(y(s_i)|\mathbf{y}(s_j):s_j
eq s_i) = f(y(s_i)|\mathbf{y}(N_i))$$

• If connectivities between edges are continuous, then the Markov assumption is redundant.

The DV

```
##
         edge_id Y
    [1,] "12"
                   "1"
##
    [2,] "13"
                   "1"
##
                   "1"
    [3,] "14"
##
    [4,] "15"
                   "0"
##
##
    [5,] "16"
                   "0"
    [6,] "17"
                   "0"
##
    [7,] "21"
                   ''O''
##
    [8,] "23"
                   "1"
##
    [9,] "24"
                   "O"
##
##
   [10,] "25"
                   "0"
```

Set Up the Connectivity Matrix, W

- Start with an adjacency matrix among all potential edge-pairs.
- Code two edges as connected if they connect opposite-colored pairs of same-colored nodes.

	54	56	57	61	62
12	0	1	1	0	0
13	0	1	1	0	0
14	0	1	1	0	0
15	0	0	0	0	0
16	0	0	0	0	0

Set Up the Connectivity Matrix, W

```
51 52 53 54 56 57 61 62 63 64 65
##
## 12
                                   1
## 13
            0
                    1
                        0
                                   1
## 14
                    1 0
                             0 0 1
## 15
                 0
                    0 0
                          0
                                    0
## 16
         0
                 0
                           0
                                    0
## 17
                                    0
                           0 0 0 1
## 21
                           0
                             0 0 1
## 23
## 24
                    1
                        0
                                   1
## 25
                        0
                                 0
                     0
                                    0
```

W[1:10, 25:35]

The Binary Conditional Distribution

$$P(Y_i = y_i | \boldsymbol{y}(N_i)) = \exp[A_i(\boldsymbol{y}(N_i))y_i - B_i(\boldsymbol{y}(N_i))], \tag{1}$$

where A_i is a natural parameter function and $B_i = \log[1 + \exp(A_i(y(N_i)))]$, and $\mathbf{y}(N_i)$ is a vector of values of the binary random variables (edges) of i's neighbors.

The Natural Parameter Function

$$A_{i}\left(oldsymbol{y}\left(N_{i}
ight)
ight) = \log\!\left(rac{\kappa_{i}}{1-\kappa_{i}}
ight) + \eta \sum_{j\in N_{i}} w_{ij}\left(y_{j}-\kappa_{j}
ight),$$
 (2)

where $\log\left(\frac{\kappa_i}{1-\kappa_i}\right) = \boldsymbol{X}_i^T\boldsymbol{\beta}$, \boldsymbol{X}_i is a column vector of k exogenous covariates, $\boldsymbol{\beta}$ is a k by 1 vector of estimation parameters, w_{ij} is the ij^{th} element of a matrix of connectivities among edges \boldsymbol{W} , and $\boldsymbol{\eta}$ is its parameter.

- When $y_j > \kappa_j$, then the dependence term makes a positive contribution to $A_i(\mathbf{y}(N_i))$ ---complementary processes;
- When $y_j < \kappa_j$, then the dependence term makes a negative contribution to $A_i(\mathbf{y}(N_i))$ ---substitution-type processes;
- Key condition: $w_{ij} = w_{ji}$.
- Model does not require (prohibits) row-standardization of w.

Estimation

$$\log PL = \sum_{i} \{y_i \log(p_i) + (1 - y_i) \log(1 - p_i)\}, \tag{3}$$

where:

$$p_i = rac{\exp[A_i(y(N_i))]}{1 + \exp[A_i(y(N_i))]}$$
 (4)

Let's Program This

```
#Likelihood
loglik<-function(par,W,Y){
b0<-par[1]
eta<-par[2]
X<-rep(1,length(Y))
xbeta<-as.matrix(X)%*%b0
kappa<-exp(xbeta)/(1+exp(xbeta)) #logit of Xb
A_i=log(kappa/(1-kappa))+eta*W%*%(Y-kappa) #Eqn 2
p_i<- exp(A_i)/(1+exp(A_i)) #Eqn 1, also Eqn 4
PL<-Y*log(p_i)+(1-Y)*log(1-p_i) #Eqn 3
ell <- -sum(PL)
#cat("ell",ell, fill=TRUE)
return(ell)
}</pre>
```

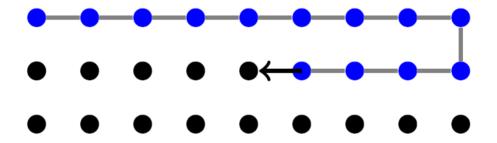
Let's Estimate

```
m1<-optim(par=c(0,0),loglik,W=Wmat,Y=Y)</pre>
m1
## $par
## [1] -3.1572208 0.8955375
##
## $value
## [1] 13.72339
##
## $counts
## function gradient
##
         95
                   NA
##
## $convergence
## [1] 0
##
## $message
## NULL
```

Estimating Standard Errors

Need to simulate *Y* based on our parameter estimates:

- Start from a vector of initial $y_0 = \{y_{01}, y_{02}, \dots, y_{0n}\}$ drawn from a binomial distribution;
- Moving row-wise, for i = 1, ..., n, individually simulate/update each observation as a function of previously simulated observations:



- n individual updates provide 1 full Gibbs iteration
- Use the result of step 2 as the new initial values and repeat step 2 to obtain M simulated networks *y* (can burn-in, thin, etc.).

Simulate Networks Based on m1

- 1. Function spatbin. genone simulates/updates a value for y for a single observation--step 2
- 2. Function spatbin.onegibbs applies spatbin.genone to update every observation of y--step 3
- 3. Function spatbin.genfield applies spatbin.onegibbs to generate ${\cal M}$ networks --step 4

```
spatbin.genone<-function(coeffs,w,curys){
b0<-coeffs[1]
eta<-coeffs[2]
X<-rep(1,length(Y))
xbeta<-as.matrix(X)%*%b0
kappa<-exp(xbeta)/(1+exp(xbeta))
A_i=log(kappa/(1-kappa))+eta*w%*%(curys-kappa)
p_i<- exp(A_i)/(1+exp(A_i))
y<- rbinom(n=length(curys), size=1, prob=p_i)
return(y)
}</pre>
```

```
spatbin.onegibbs<-function(coeffs,w,curys){
cnt<-0
n<-length(curys)
newys<-NULL
repeat{
    cnt<-cnt+1
    ny<-spatbin.genone(coeffs=coeffs,w=w,curys=curys)
    curys[cnt]<-ny[cnt]
    if(cnt==n) break
    }
newys<-curys
return(newys)
}</pre>
```

```
spatbin.genfield<-function(coeffs,w,y0s,M) {
  curys<-y0s
  cnt<-0
  res<-as.data.frame(y0s)
  repeat{
      cnt<-cnt+1
      newys<-spatbin.onegibbs(coeffs=coeffs,w=w,curys=curys)
      curys<-newys
      res<-cbind(res,curys)
      if(cnt==M) break
      }
  return(res)
}</pre>
```

```
n<-length(Y)
y0s=rbinom(n=n, size=1, prob=.5)
sims<-spatbin.genfield(coeffs=m1$par,w=W,y0s=y0s,M=1000)
#Take every 10th simulated network, i.e. burnin=10, thinning=10
sims<-sims[,seq(from=10, to=ncol(sims),by=10)]</pre>
```

Obtaining Standard Errors

- 1. Estimate the model on simulated networks (after burnin and thinning);
- 2. The standard errors are the standard deviations of the estimated coefficients.

Obtaining Standard Errors

```
sim est<-function(Y){</pre>
  res<-optim(par=m1$par,loglik,W=W,Y=as.matrix(Y))</pre>
  return(c(res$par,res$convergence))
library(parallel)
sim_est<-do.call("rbind",mclapply(sims, sim_est))</pre>
#Drop results if didn't converge (models that converged have converge
sim_est<-sim_est[sim_est[,3]==0,]</pre>
#Get sds of the estimates:
boot_se<-apply(sim_est,2,sd)</pre>
mytable<-cbind("coeff"=m1$par, "se"=boot_se[-3], "z-value"=(m1$par/boot_se[-3])
mytable
              coeff
##
                            se
                                   z-value
## [1,] -3.1572208 13.80388 -0.22871980
## [2,] 0.8955375 111.93631 0.00800042
```

Application: International Alliances

```
#Open the data:
data("ally_data")
ally_data$tot_trade<-log(ally_data$tot_trade+1)
ally_data<-ally_data[ally_data$year==2007,]
ally_data[1:5,]
##
      ccode1 ccode2 edge defense mil_ratio tot_trade joint_dem year
                                                        1 2007
## 62
                20
                             1 0.7231990 2.653716
           2
                      1
                31
                     2 1 0.9936821 2.206597
                                                        1 2007
## 75
## 227
                42
                     5 1 0.9553047 2.331096
                                                        1 2007
## 268
                51
                     6
                             1 0.9852507 2.215345
                                                        1 2007
                     7
## 305
                52
                             1 0.9901861 2.335300
                                                        1 2007
#Prepare W:
W2007 <- W
W2007[1:5,1:5]
```

```
## edge_diff1 edge_diff2 edge_diff5 edge_diff6 edge_diff7
## [1,] 0.000000e+00 1.777676e-06 1.858189e-06 2.092406e-06 2.086917e-06
## [2,] 1.777676e-06 0.000000e+00 8.051235e-08 3.147301e-07 3.092406e-07
## [3,] 1.858189e-06 8.051235e-08 0.000000e+00 2.342178e-07 2.287283e-07
## [4,] 2.092406e-06 3.147301e-07 2.342178e-07 0.000000e+00 5.489462e<sup>2</sup>09 31
```

Likelihood (1 X)

```
#Likelihood
loglik<-function(par,X,W,Y){</pre>
b0<-par[1]
b1<-par[2]
eta<-par[3]
xbeta<-b0+b1*X
kappa<-exp(xbeta)/(1+exp(xbeta)) #logit of Xb</pre>
A_i=log(kappa/(1-kappa))+eta*W%*%(Y-kappa) #Eqn 2
p_i \leftarrow exp(A_i)/(1+exp(A_i)) #Eqn 1, also Eqn 4
PL < -Y * log(p_i) + (1-Y) * log(1-p_i) #Eqn 3
ell <- -sum(PL)
#cat("ell",ell, fill=TRUE)
return(ell)
```

Let's Estimate

```
X=ally_data$tot_trade
Y=ally_data$defense
m1<-optim(par=c(0,0,0),loglik,X=X,W=W,Y=Y)</pre>
m1
## $par
## [1] -1.686244323 -0.007318607 0.936193299
##
## $value
## [1] 470.8606
##
## $counts
## function gradient
##
         88
                   NA
##
## $convergence
## [1] 0
##
## $message
## NULL
```

Standard Errors

```
spatbin.genone<-function(coeffs,X,w,curys){
b0<-coeffs[1]
b1<-coeffs[2]
eta<-coeffs[3]
xbeta<- b0+b1*X
kappa<-exp(xbeta)/(1+exp(xbeta))
A_i=log(kappa/(1-kappa))+eta*w%*%(curys-kappa)
p_i<- exp(A_i)/(1+exp(A_i))
y<- rbinom(n=length(curys), size=1, prob=p_i)
return(y)
}</pre>
```

```
spatbin.onegibbs<-function(coeffs,X,w,curys){
cnt<-0
n<-length(curys)
newys<-NULL
repeat{
    cnt<-cnt+1
    ny<-spatbin.genone(coeffs=coeffs,X=X,w=w,curys=curys)
    curys[cnt]<-ny[cnt]
    if(cnt==n) break
    }
newys<-curys
return(newys)
}</pre>
```

```
spatbin.genfield<-function(coeffs,X,w,y0s,M){
curys<-y0s
cnt<-0
res<-as.data.frame(y0s)
repeat{
    cnt<-cnt+1
    newys<-spatbin.onegibbs(coeffs=coeffs,X=X,w=w,curys=curys)
    curys<-newys
    res<-cbind(res,curys)
    if(cnt==M) break
    }

return(res)
}</pre>
```

Simulate 1000 Random Networks

```
n<-length(Y)
y0s=rbinom(n=n, size=1, prob=.5)
sims<-spatbin.genfield(coeffs=m1$par,X=X,w=W,y0s=y0s,M=1000)
#Take every 10th simulated network, i.e. burnin=10, thinning=10
sims<-sims[,seq(from=10, to=ncol(sims),by=10)]
saveRDS(sims, "sims.rds")</pre>
```

Estimate an LSGM on Each of the Simulated Networks

```
sims<-readRDS("data/sims.rds")
sim_est<-function(Y){
  res<-optim(par=m1$par,loglik,X=X,W=W,Y=as.matrix(Y))
  return(c(res$par,res$convergence))
}
library(parallel)
sim_est<-do.call("rbind",mclapply(sims, sim_est))
#Drop results if didn't converge (models that converged have converge)
sim_est<-sim_est[sim_est[,4]==0,]
saveRDS(sim_est,"sim_est.rds")</pre>
```

Calculate SEs and Make a Table

```
#Get sds of the estimates:
sim_est<-readRDS("data/sim_est.rds")
boot_se<-apply(sim_est,2,sd)
mytable<-cbind("coeff"=m1$par,"se"=boot_se[-4],"z-value"=(m1$par/boot
mytable</pre>
```

```
## coeff se z-value
## [1,] -1.686244323 2.318147e-01 -7.2741041749
## [2,] -0.007318607 3.253597e-02 -0.2249389529
## [3,] 0.936193299 2.832822e+03 0.0003304808
```

Package LSGM

Note: the package is in beta testing

```
library(devtools)
install_github("ochyzh/lsgm")
library(lsgm)
data(W)
data(toy_data)
lsgm(Y=as.matrix(toy_data$Y),W=W,X=as.data.frame(toy_data$X))
```

Your Turn

- 1. The above code estimates an lsgm with 1 edge-level covariate X using a pseudo-likelihood. Edit the code so that the resulting pseudo-likelihood can include 2 edge-level covariates X1 and X2 (i.e., the goal is to run an lsgm with tot_trade and mil_ratio as exogenous covariates).
- 2. Estimate an lsgm with defense as the dependent variable, with tot_trade and mil_ratio as the exogenous edge-level covariates, and W as the matrix that measure connectivities among defense edges.
- 3. Difficult: Change the code from the above slides to estimate the standard errors for your model.