

Advanced Network Analysis

Stochastic Actor-Oriented Models

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SAOM

Stochastic actor oriented model developed primarily by **Snijders** is implemented in the RSiena package on CRAN:

- <https://www.stats.ox.ac.uk/~snijders/siena/>
- Recent overview piece by Snijders

The Siena webpage

SIENA is a program for the statistical analysis of network data, with the focus on social networks. Networks here are understood as entire (complete) networks, not as personal (egocentered) networks: it is assumed that a set of nodes (social actors) is given, and all ties (links) between these nodes are known - except perhaps for a moderate amount of missing data. *SIENA* is designed for analyzing various types of data as dependent variables:

Longitudinal network data:

This refers to repeated measures of networks on a given node set (although it is allowed that there are some changes in the node set). Models can be specified with actor-oriented as well as tie-oriented dynamics; but mainly the former.

Longitudinal data of networks and behavior:

This is like longitudinal network data, but in addition there are one or more changing nodal variables that are also treated as dependent variables, and referred to as *behavior*. The network will influence the dynamics of the behavior, and the behavior will influence the dynamics of the network. In other words, this is about the co-evolution of networks and behavior.

Cross-sectional network data

SAOM Assumptions

- Actors control their outgoing ties and have full knowledge of broader network
- Evolution of network process occurs in microsteps
- Only one tie can change at a microstep
- Tie change only depends on the present network

SAOM Broadstrokes

- The simulation starts out at the network observed at the first time point t_0
- An actor is chosen randomly using a rate function
- The identified actor gets the opportunity to set a micro step. The actor's choice is determined by their objective function
- Model time is updated and simulation proceeds at step 2
- The simulation terminates once modified network resembles network at t_1

Rate Function

- Waiting time until change can be made by any actor follows an exponential distribution with parameter $\lambda_t g$ (g refers to number of actors in the network)
 - Values of λ_t are estimated by calculating the number of edge differences between networks:
 - The higher λ_t is the greater the number of changes between observation moments
- Probability that an actor i has the opportunity to make a change is equal to $1/g$

Actor's Objective Function

$f_i(\beta, x(i \rightsquigarrow j)) = \sum_{l=1}^k \beta_l s_{il}(x(i \rightsquigarrow j)) + U_i(t, x, j)$, where

- $s_{il}(x(i \rightsquigarrow j))$ represents k structural and exogenous effects
- β_l are statistical parameters
- and $U_i(t, x, j)$ is a random utility term

Multinomial Choice Model

$$p_{ij}(\beta, x(i \rightsquigarrow j)) = \frac{\exp(f_i(\beta, x(i \rightsquigarrow j)))}{\sum_{h=1}^g \exp(f_i(\beta, x(i \rightsquigarrow h)))}$$

- Represents the probability with which actor i changes his outgoing ties
- When $i = j$ this probability refers to the probability of not changing anything

Parameter Estimation

Solving the model requires the estimation of $\theta = (\lambda, \beta)$ using a Method of Moments approach (MoM)

- Suitable statistic for λ :

$$S_{\lambda_{tm}} = \sum_{i,j=1, i \neq j}^g |Y_{ij,t_{m+1}} - Y_{ij,t_m}|$$

- Suitable statistic for β :

$$S_{\beta_{l,tm}} = \sum_{i=1}^g s_{il}(Y_{t_{m+1}})$$

Stochastic Approximation Process

Combining the suitable statistics, we next determine the value $\hat{\theta}$ for θ as the solution of the system of equations:

$$g_n(\theta|z_n) = \sum_{t_a \in T} (E_{\theta}\{u(Y^{(a+1)}|Y^{(a)} = y^{(a)})\} - u(y^{(a+1)})),$$

- z_n simply means all available data
- $u(x)$ corresponds to the statistic being estimated
- The estimation for the MoM relies on MCMC simulations of the network change process (Robbins & Monro, 1951)

Example: Friendship Networks

```
library(RSiena)  
  friend.data.w1 <- s501  
  friend.data.w2 <- s502  
  friend.data.w3 <- s503  
  drink <- s50a  
  smoke <- s50s
```

Specify the Network DV:

```
friendship <- sienaDependent(  
  array( c( friend.data.w1, friend.data.w2, friend.data.w3 )  
    dim = c( 50, 50, 3 ) ) )
```

```
friendship
```

```
## Type          oneMode  
## Observations 3  
## Nodeset      Actors (50 elements)
```

```
class(friendship)
```

```
## [1] "sienaDependent"
```

```
dim( friendship)
```

```
## [1] 50 50 3
```

```
attributes(friendship)
```

```
## $dim  
## [1] 50 50 3  
##  
## $class  
## [1] "sienaDependent"  
##  
## $type  
## [1] "oneMode"  
##  
## $sparse  
## [1] FALSE  
##  
## $nodeSet  
## [1] "Actors"  
##
```

Specify the Behavior DV:

```
drinkingbeh <- sienaDependent( drink, type = "behavior" )  
drinkingbeh
```

```
## Type          behavior  
## Observations  3  
## Nodeset      Actors (50 elements)
```

Specify IVs:

```
smoke1 <- coCovar( smoke[ , 1 ] )
```

```
# Put the variables together in the data set for analysis
```

```
NBdata <- sienaDataCreate( friendship, smoke1, drinkingbeh )  
NBdata
```

```
## Dependent variables:  friendship, drinkingbeh
```

```
## Number of observations: 3
```

```
##
```

```
## Nodeset                      Actors
```

```
## Number of nodes              50
```

```
##
```

```
## Dependent variable friendship
```

```
## Type                          oneMode
```

```
## Observations                  3
```

```
## Nodeset                      Actors
```

```
## Densities                     0.046 0.047 0.05
```

```
##
```

```
## Dependent variable drinkingbeh
```

```
## Type                          behavior
```

```
## Observations                  3
```

```
## Nodeset                      Actors
```

```
## Range                         1 - 5
```

Possible Types of IVs

- `coCovar`--constant node-level covariate (does not change between time periods)
- `varCovar`--time-variable node-level covariate
- `coDyadCovar`--constant edge-level covariate
- `varDyadCovar`--time-varying edge-level covariate
- `sienaCompositionChange`--over time changes in node set (e.g., some actors leave the network)

?`coCovar`

Specify Endogenous Effects

```
NBeff <- getEffects( NBdata )  
NBeff
```

##	name	effectName	include	fix	test
## 1	friendship	constant friendship rate (period 1)	TRUE	FALSE	FALSE
## 2	friendship	constant friendship rate (period 2)	TRUE	FALSE	FALSE
## 3	friendship	outdegree (density)	TRUE	FALSE	FALSE
## 4	friendship	reciprocity	TRUE	FALSE	FALSE
## 5	drinkingbeh	rate drinkingbeh (period 1)	TRUE	FALSE	FALSE
## 6	drinkingbeh	rate drinkingbeh (period 2)	TRUE	FALSE	FALSE
## 7	drinkingbeh	drinkingbeh linear shape	TRUE	FALSE	FALSE
## 8	drinkingbeh	drinkingbeh quadratic shape	TRUE	FALSE	FALSE
##	initialValue	parm			
## 1	4.69604	0			
## 2	4.32885	0			
## 3	-1.46770	0			
## 4	0.00000	0			
## 5	0.70571	0			
## 6	0.84939	0			
## 7	0.32237	0			
## 8	0.00000	0			

Effects Description

effectsDocumentation(NBeff)

row	name	effectName	shortName	type	inter1	inter2	parm	interactionType
1	friendship	constant friendship rate (period 1)	Rate	rate			0	
2	friendship	constant friendship rate (period 2)	Rate	rate			0	
3	friendship	outdegree effect on rate friendship	outRate	rate			0	
4	friendship	indegree effect on rate friendship	inRate	rate			0	
5	friendship	reciprocity effect on rate friendship	recipRate	rate			0	
6	friendship	effect 1/outdegree on rate friendship	outRateInv	rate			0	
7	friendship	effect ln(outdegree+1) on rate friendship	outRateLog	rate			1	
8	friendship	effect smoke1 on rate	RateX	rate	smoke1		0	
9	friendship	effect drinkingbeh on rate	RateX	rate	drinkingbeh		0	

Specify Effects

```
NBeff <- includeEffects( NBeff, transTrip, transRecTrip )
```

```
##      effectName          include fix    test  initialValue parm
## 1 transitive triplets      TRUE     FALSE FALSE           0    0
## 2 transitive recipr. triplets TRUE     FALSE FALSE           0    0
```

```
NBeff <- includeEffects( NBeff, egoX, egoSqX, altX, altSqX, diffSqX,
                        interaction1 = "smoke1" )
```

```
##      effectName          include fix    test  initialValue parm
## 1 smoke1 alter            TRUE     FALSE FALSE           0    0
## 2 smoke1 squared alter    TRUE     FALSE FALSE           0    0
## 3 smoke1 ego              TRUE     FALSE FALSE           0    0
## 4 smoke1 squared ego      TRUE     FALSE FALSE           0    0
## 5 smoke1 diff. squared    TRUE     FALSE FALSE           0    0
```

```
NBeff
```

```
##      name      effectName          include fix    test
## 1 friendship constant friendship rate (period 1) TRUE     FALSE FALSE
## 2 friendship constant friendship rate (period 2) TRUE     FALSE FALSE
```

Define the Model:

```
myalgorithm1 <- sienaAlgorithmCreate( projname = 's50_NB' )  
  
# Estimate using the second algorithm right from the start.  
NBans <- siena07(myalgorithm1, data = NBdata, effects = NBeff)  
NBans <- siena07(myalgorithm1, data = NBdata, effects = NBeff, batch=
```

Look at results

NBans

Estimates, standard errors and convergence t-ratios

##

##

Estimate

Standard

Conve

##

Error

t-r

Network Dynamics

1. rate constant friendship rate (period 1) 6.2738 (1.2186) 0.

2. rate constant friendship rate (period 2) 5.0845 (0.8572) -0.

3. eval outdegree (density) -2.6246 (0.2222) 0.

4. eval reciprocity 2.7737 (0.2664) 0.

5. eval transitive triplets 0.8915 (0.1360) 0.

6. eval transitive recipr. triplets -0.5130 (0.2160) 0.

7. eval smoke1 alter 0.2548 (0.2914) -0.

8. eval smoke1 squared alter -0.2197 (0.2489) 0.

9. eval smoke1 ego 0.0873 (0.3034) 0.

10. eval smoke1 squared ego 0.0104 (0.2501) 0.

11. eval smoke1 diff. squared -0.0981 (0.0679) -0.

##

Behavior Dynamics

12. rate rate drinkingbeh (period 1) 1.1712 (0.3042) -0.

13. rate rate drinkingbeh (period 2) 1.6518 (0.4074) 0.

Actors Entering and Exiting the Network

```
library(devtools)
#install_github("ochyzh/networkdata")
data("duqueData")
dim(dipl_ties[[1]])
```

```
## [1] 134 134
```

```
dim(dipl_ties[[2]])
```

```
## [1] 148 148
```

- Remember that in these data, time periods have varying numbers of observations, as states enter and leave the system.
- In order to use RSiena, we must have the same number of actors in each time period. If an actor is missing, their tie values are coded as either NA or some other code we will pass to `sienaCompositionChange` option (see the manual for the second option).
- Note: if you plan to use network analysis, you have to become very comfortable with these types of data issues.

Setting Up the DV

```
library(tidyverse)
#get the full list of actors:
myactors<-unique(do.call("c",lapply(dipl_ties,names)))
dyads<-expand.grid(myactors,myactors)

dipl<-array(NA, dim = c( 194, 194, 8 ),
  dimnames=list(myactors,myactors,seq(from=1970,to=2005,by=5)))

for(t in 1:8){
  d<-dipl_ties[[t]]
  for(i in 1:nrow(d)){
    for (j in 1:ncol(d)){
      a1 = names(d)[i]
      a2 = colnames(d)[j]
      val = as.numeric(as.character(d[i,j]))

      dipl[i,j,t] <- val
      dipl[j,i,t] <- val
    }
  }
}

dipl <- sienaDependent(dipl)
```

Your Turn

1. Set up `allies` and `contig` as edge-level covariates.
2. Set up `polity` as a time-varying node-level covariate.
3. Estimate a model that includes the following covariates: outdegree, reciprocity, transitive triplets, `polity alter`, `polity ego`, `polity diff.`, contiguity, and allies.

```
ans<-readRDS("data/ans.rds")
summary(ans)
```

```
## Estimates, standard errors and convergence t-ratios
##
##
##
##
## Rate parameters:
## 0.1      Rate parameter period 1 87.4577 ( 6.8145 )
## 0.2      Rate parameter period 2 76.4185 ( 4.2766 )
## 0.3      Rate parameter period 3 63.6222 ( 2.6922 )
## 0.4      Rate parameter period 4  5.1526 ( 0.2161 )
## 0.5      Rate parameter period 5 38.1545 ( 1.1522 )
## 0.6      Rate parameter period 6  5.6488 ( 0.2027 )
## 0.7      Rate parameter period 7  7.0067 ( 0.2222 )
##
## Other parameters:
## 1.  eval degree (density)      -1.1331 (      NA )    1.8070
## 2.  eval GWESP (69)           0.2528 (      NA )   -1.0283
## 3.  eval cont                  0.3578 (      NA )   -1.8619
## 4.  eval ally                  0.5063 (      NA )   -3.3637
## 5.  eval dem alter             0.1103 (      NA )   -1.4067
## 6.  eval dem ego              0.1170 (      NA )   -1.4067
##
## Overall maximum convergence ratio:      NA
##
```

TERGM vs. SAOM

- Block et al. 2017
- Block et al. 2018
- Leifeld & Cranmer 2018