

# TEMPORAL EXPONENTIAL- FAMILY RANDOM GRAPH MODELING (TERGMS) WITH STATNET

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# Source for all things STERGM

- Pavel N. Krivitsky and Mark S. Handcock (2014). [A Separable Model for Dynamic Networks](#). *Journal of the Royal Statistical Society, Series B*, Volume 76, Issue 1, pages 29–46.

# Terminology

- The phrase “temporal ERGMs,” or TERGMs, refers to all ERGMs that are dynamic
- The specific class of TERGMs that have been implemented thus far are called “separable temporal ERGMs,” or STERGMs
- In the relevant R package, we left open the possibility that we would develop more in the future

- Thus:

	Cross-sectional	Dynamic
Name of package	ergm	tergm
Name of function in package	ergm	stergm

# ERGMs: Review

Probability of observing a graph (set of relationships)  $y$  on a fixed set of nodes:

$$P(Y = y | \theta) = \frac{\exp(\theta' g(y))}{k(\theta)}$$

Conditional log-odds of a tie

$$\begin{aligned} \text{logit}(P(Y_{ij} = 1 | \text{rest of the graph})) &= \log \left( \frac{P(Y_{ij} = 1 | \text{rest of the graph})}{P(Y_{ij} = 0 | \text{rest of the graph})} \right) \\ &= \theta' \partial(g(y)) \end{aligned}$$

where:  $g(y)$  = vector of network statistics  
 $\theta$  = vector of model parameters  
 $k(\theta)$  = numerator summed over all possible networks on node set  $y$   
 $\partial(g(y))$  represents the change in  $g(y)$  when  $Y_{ij}$  is toggled between 0 and 1

# STERGMs

- ERGMs are great for modeling cross-sectional network structure
- But they can only predict the *presence* of a tie; they are unable to separate the processes of *tie formation* and *dissolution*
- Why separate formation from dissolution?

# STERGMs

- **Intuition:** The social forces that facilitate formation of ties are often different from those that facilitate their dissolution.
- **Interpretation:** Because of this, we would want model parameters to be interpreted in terms of ties formed and ties dissolved.
- **Simulation:** We want to be able to control cross-sectional network structure and relational durations separately in our disease simulations, matching both to data



# STERGMs

- Core idea:
  - The  $y_{ij}$  values (ties in the network) and  $Y$  (the set of all  $y_{ij}$  values) are now indexed by time
  - Represent evolution from  $Y_t$  to  $Y_{t+1}$  as a product of two phases: one in which ties are formed and another in which they are dissolved, with each phase a draw from an ERGM.
  - Thus, two formulas: a formation formula and a dissolution formula
  - And, two corresponding sets of statistics



# STERGMs

ERGM: Conditional log-odds of a tie existing

$$\text{logit}(P(Y_{ij} = 1 | \text{rest of the graph})) = \boldsymbol{\theta}' \boldsymbol{d}(\boldsymbol{g}(\boldsymbol{y}))$$

STERGM: Conditional log-odds of a tie *forming* (formation model):

$$\text{logit}(P(Y_{ij,t+1} = 1 | Y_{ij,t} = 0, \text{rest of the graph})) = \boldsymbol{\theta}^+{}' \boldsymbol{d}(\boldsymbol{g}^+(\boldsymbol{y}))$$

STERGM: Conditional log-odds of a tie *persisting* (dissolution model):

$$\text{logit}(P(Y_{ij,t+1} = 1 | Y_{ij,t} = 1, \text{rest of the graph})) = \boldsymbol{\theta}^-{}' \boldsymbol{d}(\boldsymbol{g}^-(\boldsymbol{y}))$$

where:

- $\boldsymbol{g}^+(\boldsymbol{y})$  = vector of network statistics in the formation model
- $\boldsymbol{\theta}^+$  = vector of parameters in the formation model
- $\boldsymbol{g}^-(\boldsymbol{y})$  = vector of network statistics in the dissolution model
- $\boldsymbol{\theta}^-$  = vector of parameters in the dissolution model

# STERGMs

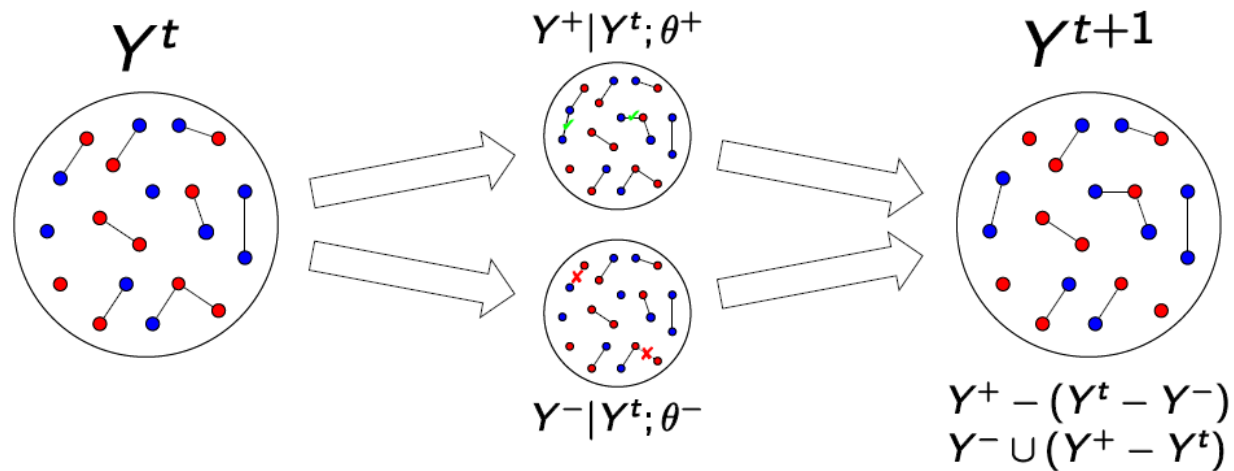
Dissolution? Or persistence?

$$\text{logit} \left( P(Y_{ij,t+1} = 1 \mid Y_{ij,t} = 1, \text{rest of the graph}) \right) = \boldsymbol{\theta}' \boldsymbol{\partial}(\mathbf{g}^-(\mathbf{y}))$$

- The model is expressed as log odds of tie equaling 1 given it equaled 1 at the last time step
- This is done to make it consistent with the formation model, so all the math works out nicely
- But it implies that the model, and thus the coefficients, should be interpreted in terms of effects on relational persistence
- That said, people tend to think in terms of relational formation and dissolution, since relational dissolution is a more salient event than relational persistence
- Thus, we often use the language of dissolution

# STERGMs

- During simulation, two processes occur separately within a time step:



- $Y^+$  = network in the formation process after evolution
- $Y^-$  = network in the dissolution process after evolution
- This is the origin of the “S” in STERGM

# STERGMs

- The statistical theory in Krivitsky and Handcock 2014:
  - demonstrates a given combination of formation and dissolution model will converge to a stable equilibrium, i.e.:

$$\text{Prevalence} \approx \text{Incidence} \times \text{Duration}$$

- This and other work in press provide the statistical theory for methods for estimating the two models, given certain kinds of data

# STERGMs: Example of interpretation

Term =  $\sim$ edges

	$\theta \nearrow$	$\theta \searrow$
Formation model	more new ties created each time step	fewer new ties created each time step
Dissolution (persistence) model	more existing ties preserved (fewer dissolved); longer average duration	fewer existing ties preserved (more dissolved); shorter average duration

What combo do you think is most common in empirical networks?

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# STERGMs: Example of interpretation

Term = `~concurrent` (# of nodes with degree 2+)

	$\theta \nearrow$	$\theta \searrow$
Formation model	more ties added to actors with exactly 1 tie	fewer ties added to actors with 1 tie
Dissolution (persistence) model	actors with 2 ties more likely to have them be preserved	actors with 2 ties more likely to have them dissolve

What combo do you think is most common in empirical sexual networks?

# STERGMs: Example of interpretation

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# STERGMs: Data sources

- 1. Multiple cross-sections of complete network data
  - easy to work with
  - but rare-to-non-existent in some fields
- 2. One snapshot of a cross-sectional network (census, egocentric, or otherwise), plus information on relational durations
  - more common
  - but introduces some statistical issues in estimating relation lengths

# STERGMs: nodal dynamics

- All of the statistical theory presented so far regards networks with
  - Dynamic relationships, but still
  - Static actors
- I.e. no births and deaths, no changing of nodal attributes
- The statistical theory of STERGM can handle nodal dynamics during simulation, with a few added tweaks
  - Most important is an offset term to deal with changing population size
  - Without it, density is preserved as population size changes
  - With it, mean degree is preserved as population size changes

# STERGMs: nodal dynamics

- For more info, see:

Pavel N. Krivitsky, Mark S. Handcock, and Martina Morris (January 2011). [Adjusting for Network Size and Composition Effects in Exponential-Family Random Graph Models](#). *Statistical Methodology*, 8(4): 319–339

- And for more help with using STERGMs to simulate dynamic networks along with changing nodes and attributes:
  - Take our intensive summer workshop on network modeling for epidemic diffusion
  - Explore the online materials for the workshop (on the statnet webpage)
  - Try the EpiModel package

# To the tutorial.....

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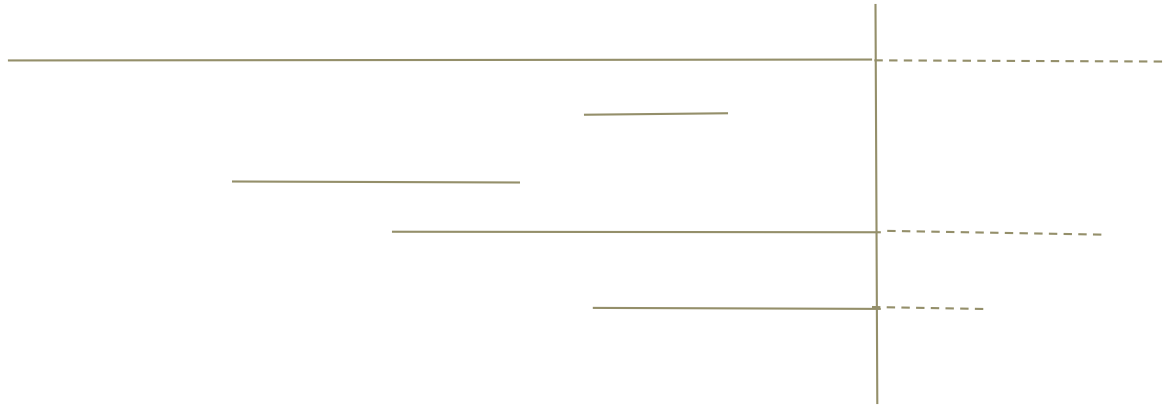
(reference slides follow)

# One cross-section + duration info

- In some domains, often takes the form of
  - asking respondents about individual relationships (either with or without identifiers).
  - Often this is the  $n$  most recent, or all over some time period, or some combination (e.g. up to 3 in the last year)
  - asking whether the relationship is currently ongoing
  - if it's ongoing: asking how long it has been going on (or when it started)
  - if it's over: asking how long it lasted (or when it started and when it ended)
- From this we want to estimate
  - the mean duration of relationships
  - perhaps additional information about the variation in those durations (overall, across categories of respondents, etc.)

# One cross-section + duration info

## ■ Issues?



1. Ongoing durations are right-censored
  - can use Kaplan-Meier or other techniques to deal with

# One cross-section + duration info

## ■ Issues?



2. Relationships are subject to length bias in their probability of being observed
  - This can also be adjusted for statistically
  - However, complex hybrid inclusion rules (e.g. most recent 3, as long as ongoing at some point in the last year) can make this complicated

# One cross-section + duration info

- In practice (and for examples in this course), we sometimes rely on an elegant approximation:
  - If relation lengths are approximately exponential/geometric (a big if!), then the effects of length bias and right-censoring cancel out
  - The mean amount of time that the **ongoing** relationships have lasted until the day of interview (relationship age) is an unbiased estimator of the mean duration of relationships
  - Why?!?



# One cross-section + duration info

- Exponential/geometric durations suggests a memoryless processes – one in which the future does not depend on the past
- Imagine a fair, 6-sided die:

- 1/6 • What is the probability I will get a 1 on my next toss?
- 1/6 • What is the probability I will get a 1 on my next toss given that my previous 1 was five tosses ago?
- 6 • On average, how many tosses will I need before I get my first 1?
- 6 • On average, how many more tosses will I need before I get my next 1, given that my previous 1 was 8 tosses ago?

Geometric	
Parameters	$0 < p \leq 1$ success probability (real)
Support	$k \in \{1, 2, 3, \dots\}$
Probability mass function (pmf)	$(1 - p)^{k-1} p$
Cumulative distribution function (CDF)	$1 - (1 - p)^k$
Mean	$\frac{1}{p}$

# One cross-section + duration info

- Now, let's imagine this fairly bizarre scenario:
  - You arrive in a room where there are 100 people who have each been flipping one die; they pause when you arrive.
  - You don't know how many sides those dice have, but you know they all have the same number.
  - You are not allowed to ask any information about what they've flipped in the past.
  - The only information people will give you is: how many flips after your arrival does it take until they get their first 1?
  - You are allowed to stay until all of the 100 people get their first 1, and they can inform you of the result.
- Given the information provided you, how will you estimate the number of sides on the die?

# One cross-section + duration info

- Simple: when everyone tells you how many flips it takes from your arrival until their first 1, just take the mean of those numbers. Call it  $m$ .
- Your best guess for the probability of getting a 1 per flip is  $1/m$ .
- And your best guess for the number of sides is the reciprocal of the probability of any one outcome per flip, which is  $1/(1/m)$ , which just equals  $m$  again.
- Voila!

# One cross-section + duration info

Retrospective relationship surveys are like this, but in reverse:

Dice:



Relationships:



# One cross-section + duration info

- If you have something approximating a memoryless process for relational duration, then an unbiased estimator for relationship length is to:
  - ask people about how long their ongoing relationships have lasted up until the present
  - take the mean of that number across respondents.

# One cross-section + duration info

- In practice, we find that the geometric distribution doesn't often capture the distribution of relational durations overall.
- But, if you divide the relationships into 2+ types, it can do a reasonable job within type
- Especially if you remove any 1-time contacts and model them separately (for populations where they are common)
- Remember: DCMs model pretty much everything as a memoryless process, so approximating one aspect of our model that way is well within common practice