

Network Modeling for Epidemics

DAY 4: SIMULATING DISEASE TRANSMISSION ON DYNAMIC NETWORKS W/ EXOGENOUS FEEDBACK

NME WORKSHOP

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Changing network size and composition

As social networks change in size (say, for instance, as a village of n = 5,000 nodes grows to n = 10,000 nodes), which of the following do you think is generally preserved?

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- Number of edges?
- Mean degree? 2e/n
- Density? $e/\binom{n}{2}$

Changing network size and composition

- Applying the coefficients as is from a given stergm fit to a network of changing size will lead to preservation of density across time
- For one-mode networks: preserving mean degree instead requires a simple transformation of the edges coefficient in the formation model:

$$\theta_{new} = \theta_{old} + \ln(N_{old}) - \ln(N_{new})$$

- This is mathematically equivalent to partitioning the original edges term into an offset equal to ln(N) and a residual, and then updating the offset as N changes.
- EpiModel handles this for you
- If you are going to code your own models outside EpiModel, you must handle this

Changing network size and composition

- As network composition changes, balancing will happen automatically the explicit pairing of individuals requires this.
- Nevertheless one does not have always have straightforward control over this
 - e.g. with just an edges term in the model, two sexes will automatically "meet in the middle"
- Can change parameterizations to obtain different dynamic behavior
 - worth thinking through the behavior you expect, and what you see for your model
- Some theory to guide you can be found in Morris (1991), Koehly, Goodreau and Morris (2004), Krivitsky, Handcock and Morris (2011)

Relational dissolution through node departure

- We fit our dynamic network using static data, with a process for dissolving relationships governed by a coefficient derived from relational duration
- All of this was done in a context that contained no information about deaths or other forms of node departures – i.e. processes that also terminate relationships
- If we simply layer departures on to our model (even with the size correction on the previous slide) we will see two measures drop down below the expected values we want:
 - relationship durations
 - number of relationships
- Some aspects of this might be desired
 - e.g. if we could interview dead people we might find their past relationships to be shorter than those of the same birth cohort in our sample who are still alive
- but others are likely not.

Relational dissolution through node departure

• An approximate correction for this is:

- 1. Calculate dissolution coefficients as before (without considering death or other departures from the population)
- 2. Estimate formation coefficients conditional on these dissolution coefficients.
- 3. Calculate new dissolution coefficients that reflect the log-odds of a relationship sustaining conditional on both actors remaining, which equals:

$$logit\left[1 - \frac{P(E_t) - P(N_t)}{P(\neg N_t)}\right]$$

where:

 $P(E_t)$ = the overall prob. of a tie dissolving at time *t* from any cause = 1/D $P(N_t)$ = the prob. of either incident node departing at time *t*

Relational dissolution through node departure

- The complicating factor is that P(N_t) may very well change over time as your disease prevalence changes
- But then again, it's probably OK for relationship lengths to shorten, and network density to decline, slightly as deaths increase
- Again, EpiModel handles this for you
- If you are going to code your own models outside EpiModel, you must handle this
- Bigger point:

DIAGNOSE THE HECK OUT

OF YOUR SIMULATIONS!!!

Review of offsets and corrections

When approximating the fit of a formation STERGM conditional on dissolution STERGM	subtract dissolution coefficients from corresponding formation ones (edapprox=TRUE)
When network size N changes and you want to preserve mean degree	add the <i>In</i> of the old N and subtract the <i>In</i> of the new N to the edges coefficient in the formation model (or equivalently, use an edges offset and update it with <i>In</i> of new N)
To adjust for node departures in simulating from a STERGM model estimated from a cross-sectional network and durations	use $\log \left[1 - \frac{P(E_t) - P(N_t)}{P(\neg N_t)}\right]$
	in calculating your dissolution coefficients