On the Bayesian Estimation for ERGM

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Final Presentation for Bayesian Statistical Methods

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Introduction

- Is double-Metropolis Hasting (DMH) algorithm *implementable* for exponential random graph models (ERGM)?
- What are the consequences of *insufficient* model selections?

- Network formation is Markov-dependent, and we want to understand the dynamics.
- Networks in ERGM formulation is a doubly intractable statistical model.
- Exchange algorithm (a Bayesian approach) is asymptotically consistent for ERGM, but inefficient.

Model Setup

ERGM is a model describing a graph **y** with a probability in a general exponential form forms:

$$\pi(\mathbf{y}|\theta) = \frac{1}{z(\theta)} \exp\left(\sum_{d=1}^{D} \theta_d s_d(\mathbf{y})\right) = \frac{\exp\left(\theta^{\mathrm{T}} \mathbf{s}(\mathbf{y})\right)}{z(\theta)}$$
(1)

where the sufficient statistics in our study are:

$$\theta^{\mathrm{T}} \mathsf{s}(\mathsf{y}) = \theta_{\nu} \nu(\mathsf{y}) + \theta_{\rho} \rho(\mathsf{y}) + \theta_{\tau} \tau(\mathsf{y})$$

where

• $\nu(\mathbf{y}) = \sum_{i,j} y_{ij} \# \text{ of edges}$ • $\rho(\mathbf{y}) = \sum_{i} \sum_{k>j} \sum_{j \neq i,k} y_{ij} y_{jk} \# \text{ of two-stars}$ • $\tau(\mathbf{y}) = 3 \sum_{i} \sum_{k>i} \sum_{j \neq i,k} y_{ij} y_{jk} y_{ki} \# \text{ of triangles}$

- MCMC estimation (Snijders, 2002)
- DMH & Exchange algorithm approach (Caimo and Friel, 2013)
- Approximate Bayesian Computation (Yin and Butts, 2020)

Our Model and Algorithms

- 1. for t = 1 to T do
- 2. Generate $\theta' \sim h(\cdot|\theta)$
- 3. Sample an auxiliary variable $\mathbf{y}' \sim \pi(\mathbf{y}'|\theta')$ using an exact sampler
- 4. Compute

$$r(\theta, \theta', \mathbf{y}'|\mathbf{y}) = \frac{\pi(\mathbf{y}|\theta')\pi(\theta')}{\pi(\mathbf{y}|\theta)\pi(\theta)} \frac{\pi(\mathbf{y}'|\theta)}{\pi(\mathbf{y}'|\theta')} \frac{h(\theta|\theta', \mathbf{y})}{h(\theta'|\theta, \mathbf{y})}$$

- 5. Draw $u \sim$ Uniform(0, 1)
- 6. **if** u < r **then** set $\theta = \theta'$
- 7. end for

The normalizing constant $z(\theta)$ is canceled by introducing $\pi(\mathbf{y}'|\theta)$. This is the exchange algorithm proposed by Liang (2010).

On a tie level, we can describe the probability to form a chosen edge between note *i* and *j* as:

$$logit(y_{ij} = 1|y_{ij}^c) = \theta' \delta(y_{ij})$$
(2)

where

- y_{ij} is a random variable state of the actor pair *i*, *j*.
- y_{ij}^c is the complement of y_{ij}
- $\delta(y_{ij})$ is the vector of the "change statistics", i.e. $\delta(y_{ij}) = g(y_{ij}^+) - g(y_{ij}^-)$, where g denotes the state of tie y_{ij} .

Exact Sampler: MH Algorithm

- 1. set the true parameter $\hat{\theta}$
- 2. for t = 1 to T do
- 3. randomly choose i, j
- 4. define $\delta(y_{ij})$ as

$$(1, deg(i) + deg(j) - 2 \times \sum_{i} \sum_{j < i} \sum_{k \neq i, j} y_{ik} y_{jk}, \sum_{i} \sum_{j < i} \sum_{k \neq i, j} y_{ik} y_{jk})$$

- 5. compute $\hat{\theta}' \delta(y_{ij})$
- 6. calculate

$$log(p) = (-1)^{y_{ij}} \cdot log(\frac{exp(\hat{\theta}'\delta(y_{ij}))}{1 - exp(\hat{\theta}'\delta(y_{ij}))})$$
(3)

7. if $\min(log(p), 0) > log(Uni(0, 1))$:

$$8. y_{ij} = 1 - y_{ij}$$

- 1. A beta-binomial is first used to approach θ_0^{init}
- 2. Mixed adaptive MCMC is proposed by a linear combination of random walk and covariance.
- 3. Hamiltonian MCMC may not work as we cannot derive the gradient for the likelihood (even if we could, it cannot be applied across different phase spaces).
- 4. We reject highly degenerated networks, i.e. networks generated from any θ with an acceptance rate lower than 0.02.

Simulation Results

Given the parameter value, we use both *ergm* and our code to generate two networks. We consider 4 cases of estimation:

- 1. *ergm* simulated network + *ergm* and *bergm* parameter estimation
- 2. *ergm* simulated network + our DMH code
- 3. our simulated network + our DMH code
- 4. our simulated network + *ergm* and *bergm* parameter estimation

Network Generation





(a) ergm. edge: 38; 2-star: 72; triangle: 3

(b) our code. edge: 32; 2-star: 57; trian-gle: 1

Figure 1: Parameter value: -3.5, 0.1, 0.5

ergm: estimate with Monte Carlo MLE

• R Code

```
model <- sim_network ~ edges + kstar(2) + triangle
est_ergm <- ergm(model)
summary(est_ergm)</pre>
```

• Result

Parameter	Mean	Std.
θ_0 (edge)	-2.8456	0.4797
θ_1 (2-star)	-0.0762	0.1339
θ_2 (triangle)	1.1726	0.6722

Case1: ergm network + ergm and bergm estimation

bergm: using exchange algorithm and parallel adaptive direction sampler to improve the mixing of Markov Chains (related to the gamma and nchains)

• R Code

• Result

Parameter	Mean	Std.
θ_0 (edge)	-3.14444889	0.4874545
θ_1 (2-star)	-0.04660349	0.1659005
θ_2 (triangle)	-0.12884212	1.3766226

MCMC Diagnostic bergm



MCMC output for Model: y ~ edges + kstar(2) + triangle

Case2: ergm simulated network + our DMH code





Case2: ergm simulated network + our DMH code



Case3: our simulated network + our DMH code



Case3: our simulated network + our DMH code



ergm

Parameter	Mean	Std.
θ_0 (edge)	-3.53935	0.41757
θ_1 (2-star)	0.11874	0.11842
θ_2 (triangle)	-0.04484	1.09843

bergm

Parameter	Mean	Std.
θ_0 (edge)	-3.36917773	0.4344070
θ_1 (2-star)	0.05895917	0.1324414
θ_2 (triangle)	-0.64722396	1.2797613

MCMC Diagnostic bergm



MCMC output for Model: y ~ edges + kstar(2) + triangle

Discussions

- How many iterations are considered *asymptotically* large enough? 30000? 120000?
- Instability and degeneracies, i.e. different θ could lead to the same configurations (Schweinberger, 2012).
- Hunter et al. (2008) argues that the k-star and triangles are in nature degenerate, which implies the parameters in the model may be unidentifiable.

Exact Sampler: Realization



Figure 3: Distribution of sampled edges, two-stars, triangles from parameter: (-3.5, 0.1, 0.5). 1000 samples, 30000 iterations

The consequence of degeneracy i



Figure 4: edges: 42, two-stars: 95, triangles: 1; Parameter: (-5, -0.2, 5)

The consequence of degeneracy ii



Figure 5: edges: 42, two-stars: 80, triangles: 1; Parameter: (-3.5, 0.1, 0.5)

- **bergm** has implemented Gibbs sampler for proposing new θ , while we used MH sampler
 - How to derive the marginal density?
 - Any accepted θ may be good enough?
- Our auxiliary network generating iterations is 30000, while we chose 2500 iterations for *bergm*. Which generates good-quality networks?

- 1. Hyperparameters for acceptance rate and step size are updated [under construction].
- Simulated annealing: We define temperature as a metric related to the observed s(y) and set up a tolerance for restarting and mixing, naively related to approximate Bayes computation (ABC) (Albert, Künsch, and Scheidegger, 2015) [under construction].

- Is it possible to target the parameters in the formation mechanism directly via ABC? Could it avoid degeneracy (partially yes).
- Without nodal (heterogeneous) effects, the missing-not-at-random data may be applicable within this framework.
- Model selection either on the ERGM family or the dynamical system by Bayesian statistics should be further established.
- We will optimize the hyperparameter updating rules, as well as construct priors with empirical Bayes.
- Examine a version of the Gibbs sampler

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