

# Misspecification, Sparsity, and Superpopulation Inference with Large-Scale Social Network Data

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# Overview: Foundations of Applied Statistics

Research agenda: A theory of Applied Statistics.

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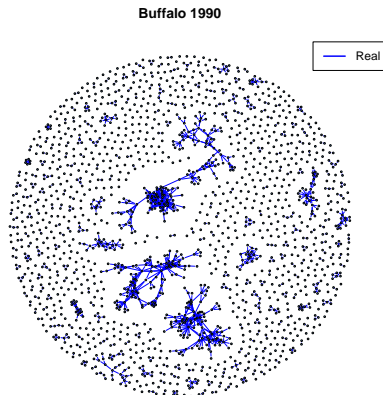
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Research agenda: A theory of Applied Statistics.

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New desiderata that we can work into modeling decisions.

# Generative Network Models



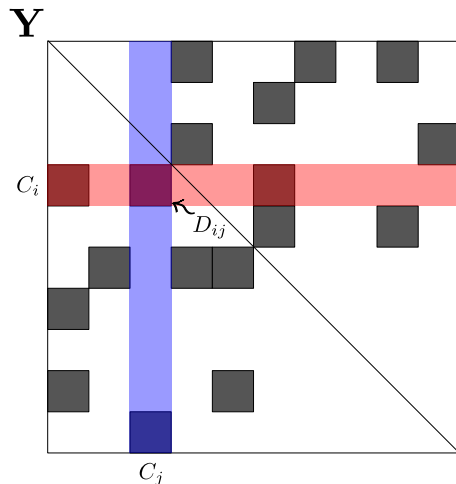
Obtain a set  $V$  of  $n$  actors.

Explain or predict pairwise outcomes  $Y_V$ , potentially using pairwise covariates  $X_V$ .

$X_V$  may be observed or latent.

Running example: inventor collaboration network.

# Data Representation



$Y_V$  entries in **arbitrary sample space**  $\mathcal{Y}$ .

Covariates  $X_V$  combine observed, latent attributes,

$$X_V^{ij} = f(C_V^i, C_V^j, D_V^{ij}).$$

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Can we borrow strength between different actor-sets to obtain better resolution on the behavior of both? (e.g., regional random effects)

# Attempt 1: Network Regression

**Cox PH regression.** (Perry and Wolfe, 2013)

Inventor coauthorships in Michigan's motor industry 1982-1988.

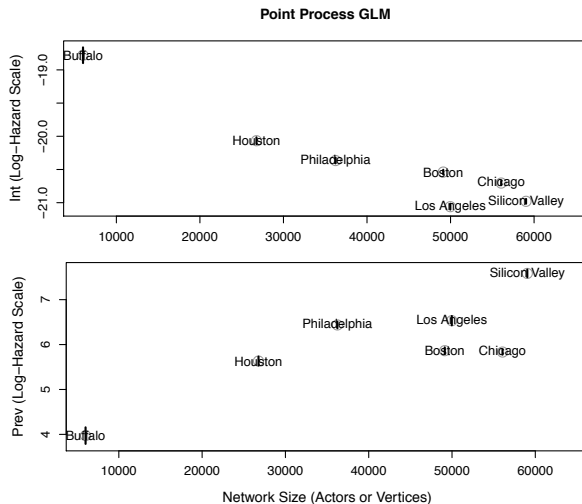
Covariates (Coefs are **log-ratios**):

- post85: After 1985.
- asgnum: Work for same firm.
- prev: Have worked together before.

	lower	est	upper
post85	15.49	15.84	16.20
asgnum	4.65	4.83	5.02
pre	11.36	11.73	12.10
post85:asgnum	-4.77	-4.40	-4.03
post85:prev	-14.57	-14.00	-13.44
asgnum:prev	-5.56	-5.16	-4.76
post85:asgnum:prev	3.91	4.52	5.13

# Attempt 2: Regional Comparison Regression

**Point process regression.** Same time window, different regions.



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What sort of question are you asking?

## “Single-Sample”

For a **fixed set** of actors

- Project forward in time.
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**Replications restricted to  $V$ .**

**Replications for any  $V \subset \mathbb{V}$ .**

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Theory so far covers single-sample inference, giving little guidance for superpopulation questions .

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## Definition 1 (Random Graph Process).

A random interaction process  $Y_{\mathbb{V}}$  is a stochastic process indexed by a countably infinite vertex set  $\mathbb{V}$  whose finite-dimensional distribution for any finite subset  $V \subset \mathbb{V}$  defines an interaction graph  $Y_V$  with vertex set  $V$ . Denote the law of  $Y_{\mathbb{V}}$  as  $\mathbb{P}_{\mathbb{V}}$  and the law of a finite-dimensional projection  $Y_V$  as  $\mathbb{P}_V$ .

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**Statistical interpretation:** Observed samples are finite subgraphs of population graph. Population graph is of scientific interest.

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**Superpopulation:** Random graph process

- Induces Kolmogorov consistency on constructed sequences.
- Focus on relationships between finite-dimensional distributions.
- Infinity is useful, but limit is unimportant.

# The Method: Parametric MLE

## Operational procedure

Let  $\mathbb{P}_{0,\mathbb{V}}$  be the law of the true population process;  $\mathbb{P}_{0,\mathbb{V}}$  be the distribution of the sample  $Y_{\mathbb{V}}$ .

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## Likelihood inference procedure

- 1 Propose a model family  $\mathcal{P}_{\Theta,V}$  of models  $\mathbb{P}_{\theta,V}$ .
- 2  $\mathcal{P}_{\Theta,V}$  implies a likelihood  $\mathbb{P}_{\theta,V}$  on the sampled index set  $V$  for each  $\theta \in \Theta$ . Compute

$$\hat{\theta}_V = \arg \max_{\Theta} \log \mathbb{P}_{\theta,V}(Y_V). \quad (1)$$

- 3 Interpret  $\hat{\theta}_V$  as a superpopulation parameter estimate.



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Note: Step 3 is the only difference between single-sample and superpopulation.

# Models and Misspecification

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No parsimonious model can fully represent complex network structure.

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No parsimonious model can fully represent complex network structure.

Choose one:

- **Local structure.** Homophily, Heterophily, Transitivity, etc.
- **Global structure.** Sparsity, Percolation, etc.

# Models and Misspecification

## Common local approaches

Conditionally independent dyads (regression):

$$P(Y | X) = \prod_{i < j < n} P(Y_V^{ij} | X_V^{ij}).$$

Infinitely exchangeable dyads (Aldous-Hoover):

$$P(Y | X) = \int_{\mathcal{C}} \prod_{i < j < n} P(Y_V^{ij} | W_V^{ij}(C_V^i, C_V^j)) dF(C).$$

# Misspecified Models and Science

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# Misspecified Models and Science

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How do we make sense of a misspecified model?

Parameter estimates as **measurements** of  $\mathbb{P}_{0,V}$ .

Minimal criterion for “usefulness”: **Stability**.

“Similar” inputs  $Y_V$  yield “similar” estimates  $\hat{\theta}_V$ .



# Misspecified Models and Science

Stability: Single sample case

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Huber 1967 showed MLE is large-sample consistent for a pseudo-true parameter (naming due to Sawa 1978), satisfying

$$\bar{\theta}_V = \arg \max_{\theta \in \Theta} \mathbb{E}_{\mathbb{P}_0} [\log \mathbb{P}_{\theta,V}(Y_V)] \quad (2)$$

“Similar output” defined by concentration of  $\hat{\theta}_V$  in large samples.

# Misspecified Models and Science

Stability: Superpopulation case

“Similar input” means any sample  $Y_V$  drawn from the same superpopulation  $\mathbb{P}_{0,V}$ .

Intuitively, outputs  $\hat{\theta}_V$  are similar if they **effectively estimate** the same thing.

What does the MLE  $\hat{\theta}_V$  effectively estimate when the model is misspecified?

# The Effective Estimand of the MLE

Define the effective estimand of the MLE as the finite-sample pseudo-true parameter.

$$\bar{\theta}_V = \arg \max_{\theta \in \Theta} \mathbb{E}_{\mathbb{P}_0}[\log \mathbb{P}_{\theta, V}(Y_V)] \quad (2)$$

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Justifications:

- Finite-sample concentration (e.g., Spokoiny 2012).
- Fisher-consistency inversion.
- Estimating equation unbiased.
- KL projection plug-in.

# Superpopulation Stability Criterion

## Criterion 1.

*A procedure is superpopulation stable for making inferences about a superpopulation process  $\mathbb{P}_{0,V}$  only if, for any finite sample  $Y_V$  generated according to  $\mathbb{P}_{0,V}$ , the effective estimand  $\bar{\theta}_V$  of the estimator  $\hat{\theta}_V$  is invariant to the indexing set  $V$ .*

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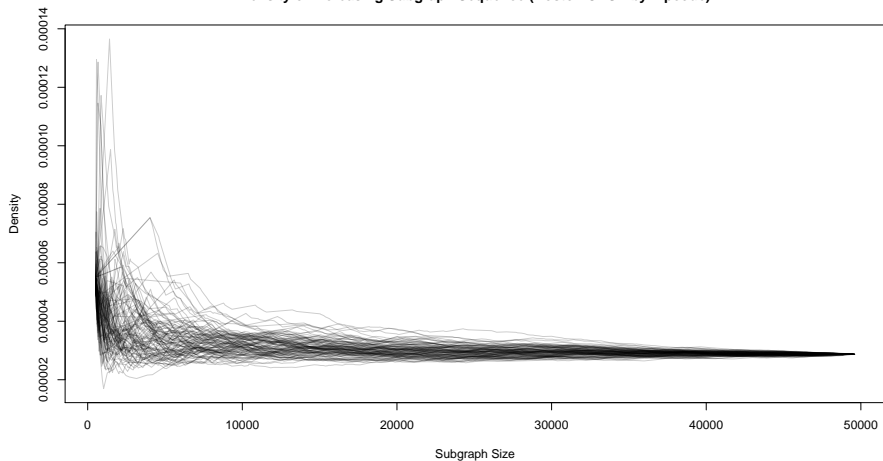
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Test the criterion with top-down specification of superpopulation properties, e.g., sparsity.

# The Wrinkle: Sparsity

## Illustration

Density of Increasing Subgraph Sequence (Boston CBSA by Zipcode)





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Define the **density operator**

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## Definition 1 (Sparse Graph Process).

Let  $Y_{\mathbb{V}}$  be a random graph process on  $\mathbb{V}$ .  $Y_{\mathbb{V}}$  is *sparse* if and only if for any  $\epsilon > 0$  there exists an  $n$  such that for any subset of vertices  $V \in \mathbb{V}$  with  $|V| > n$  the corresponding finite dimensional random graph  $Y_V$  has the property  $\mathbb{E}(D(Y_V)) < \epsilon$ .

# The Wrinkle: Sparsity

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Differences vs single-sample sparsity.

## Single-sample

- Defined in terms of random graph sequences.
- Often defined in explicitly non-Kolmogorov-consistent terms.
- Analogy for single sample with very few observed interactions.

## Superpopulation

- Property of a random graph process, not a random graph.
- Defines an assumption about the system, not a theoretical object.

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A model family  $\mathcal{P}_{\theta, \mathbb{V}}$  is **sparsity misspecified** iff for every  $\theta \in \Theta$  and every increasing sequence of vertex sets  $(V_n)$ ,

$$\frac{\mathbb{E}_{\theta}(D(Y_{V_n}))}{\mathbb{E}_0(D(Y_{V_n}))} \rightarrow 0 \text{ or } \infty.$$

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For example,

- For CID (under regularity) and exchangeable models, population extension is **dense** or **empty** (e.g., Orbanz and Roy, 2013).
- For process models, most lock in a given form for  $\epsilon(n)$  (e.g., power law for preferential attachment).

# Main Result: Moving Target

## Theorem 1 (Moving target theorem).

Let  $(V_n)$  be an increasing sequence of vertex sets from  $\mathbb{V}$ . Suppose the following hold:

- ① For some finite  $n$ ,  $\mathbb{E}_0(D(Y_{V_n})) > 0$ .
- ② The inferential family  $\mathcal{P}_{\Theta, \mathbb{V}}$  is sparsity misspecified for the true population process  $\mathbb{P}_{0, \mathbb{V}}$ .
- ③ The inferential model is responsive to the sample density  $D(Y_{V_n})$  under the true population process and

$$|\mathbb{E}_{\bar{\theta}}(D(Y_{V_n})) - \mathbb{E}_0(D(Y_{V_n}))| = O(\epsilon(n)). \quad (3)$$

Then,  $\bar{\theta}_{V_n}$  varies with  $n$  in the sense that for any  $n$ , there exists an  $n' > n$  such that  $\bar{\theta}_{V_n} \neq \bar{\theta}_{V_{n'}}$ , and the MLE of the model violates Criterion 1.

# Example: Poisson Regression

## Setup

Question: How do firms influence collaboration dynamics?

$Y_V$  collaboration counts;  $X_V^{ij}$  indicates shared firm.

Assumptions:

- The true collaboration-generating process  $Y_{0,V}$  is sparse .
- All firms have finite size.
- A non-vanishing fraction of firms have a positive number of expected within-firm interactions.

# Example: Poisson Regression

Model and effective estimand

Model:

$$Y_{V_n}^{ij} \stackrel{\text{IID}}{\sim} \text{Pois}(\exp(\theta_1 + X_{V_n}^{ij} \theta_2)), \quad (4)$$

Effective Estimands:

$$\bar{\theta}_{1V} = \log \left( \frac{\sum_{ij} \mathbb{E}_0(Y_V^{ij} | X_V^{ij} = 0)(1 - X_V^{ij})}{\sum_{ij} (1 - X_V^{ij})} \right) \quad (5)$$

$$\bar{\theta}_{2V} = \log \left( \frac{\sum_{ij} \mathbb{E}_0(Y_V^{ij} | X_V^{ij} = 1)X_V^{ij}}{\sum_{ij} X_V^{ij}} \bigg/ \frac{\sum_{ij} \mathbb{E}_0(Y_V^{ij} | X_V^{ij} = 0)(1 - X_V^{ij})}{\sum_{ij} (1 - X_V^{ij})} \right). \quad (6)$$

What's wrong with this picture?

# Partial Resolution

Salvaging conditional independence

What *can* we estimate with local models?

Two-stage process. Complex **relationship** structure  $R$ .  $Y$  simple conditional on relationships.

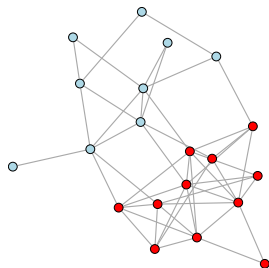
$$\mathbb{P}_{0,V} = \mathbb{P}_0(R) \prod_{i < j < n} \mathbb{P}_0(Y_V^{ij} \mid R_V, X_V).$$

Changes question:

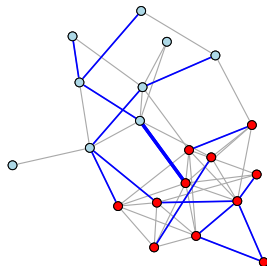
$$\mathbb{P}_{\theta,V}(Y_V \mid X_V) \rightarrow \mathbb{P}_{\beta,V}(Y_V \mid R_V, X_V).$$

# Partial Resolution

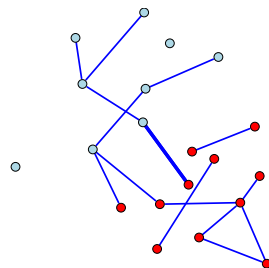
Conditionally Independent Relationship model



Unobservable  
Relationship/Risk



Both



Observable  
Interactions

# Partial Resolution

## Partial likelihood inference

Let  $A_V = \mathbb{I}\{Y_V \neq 0\}$ .

Exploit conditional distribution

$$\mathbb{P}_\beta(Y_V | A_V, X_V) = \frac{\mathbb{P}_\beta(Y_V | R_V, X_V)}{1 - \mathbb{P}_\beta(Y_V = 0 | R_V X_V)},$$

or the **zero-truncated likelihood**.

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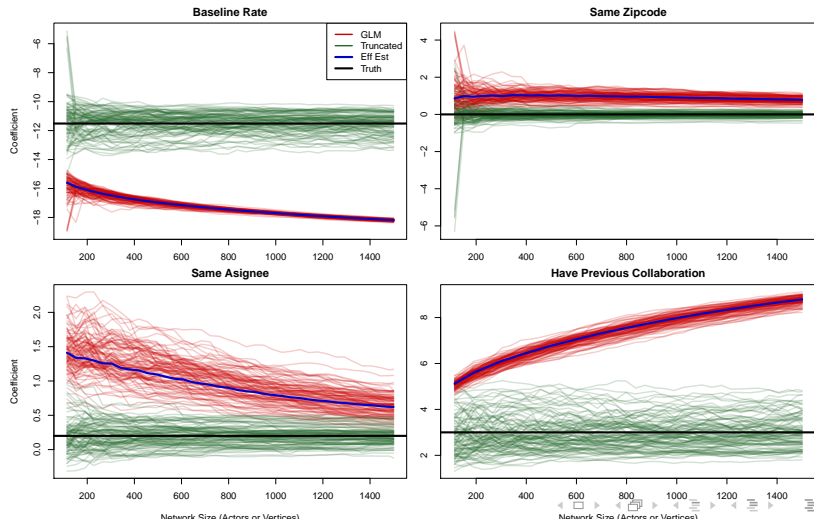
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Bonus: Computation is  $O(\sum_{ij} A_V)$ .

# Partial Resolution

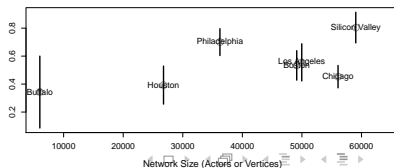
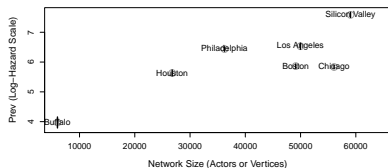
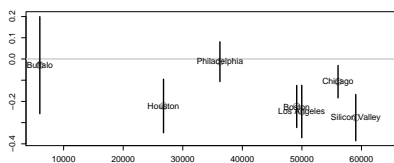
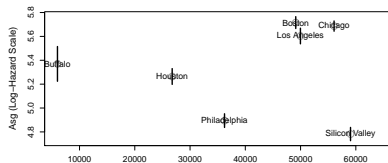
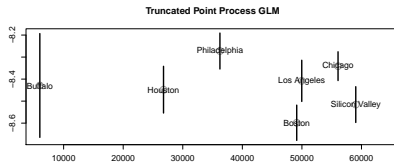
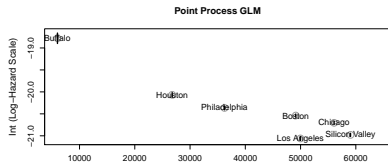
## Simulated success

Estimation



# Partial Resolution

## Real success



# Discussion

**GEP Box:** “Remember that all models are wrong; the practical question is how wrong do they have to be to not be useful.”

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Modeling *less* of the system by seeking invariances make a model *more* scientifically relevant. **Invariance** can be better than a bad explanation.

